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Report on Singapore's Antimicrobial Resistance Research Landscape 2009- 2019

ANTIMICROBIAL RESISTANCE COORDINATING OFFICE
NATIONAL CENTRE FOR INFECTIOUS DISEASES
SINGAPORE

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EXECUTIVE SUMMARY

Antimicrobial resistance occurs when microorganisms such as bacteria, viruses, fungi and parasites evolve under selection pressure and develop mechanisms to circumvent the action of medicines. A key driver of resistance is the misuse of antimicrobials in humans, animals and agriculture worldwide. As a result of increasing resistance, infections caused by antimicrobial resistant microorganisms are more difficult to treat and pose a significant challenge to healthcare systems. Without concrete actions to mitigate the threat of antimicrobial resistance, our lives and economy would be affected, with an estimated 10 million dying from drug-resistant infections per year by 2050 [1].

The accelerated development of antimicrobial resistance and the concomitant lower return in investments for the development of antimicrobials by pharmaceutical companies had led to slower rates of research and development into newer antimicrobials. In the development of new antibiotics, most research is now undertaken by academic institutions [2].

In 2015, the Global Action Plan on Antimicrobial Resistance published by the World Health Organization set clear objectives to guide national and international efforts against antimicrobial resistance. Among these objectives was to strengthen our knowledge and evidence base through surveillance and research that would guide our actions and investments towards AMR.

An understanding of the current state of antimicrobial resistance research in Singapore is needed to develop clear rationales for national actions and research investments. Therefore, a review of the antimicrobial resistance research in Singapore was conducted.

A systematic literature review approach was undertaken to retrieve relevant research articles from databases and assembled to review the trends in research. Although antimicrobial resistance research in Singapore had shown an upward trend since 2009 (including the contribution of samples and data to regional research efforts and participation in international collaborations), some domains of research remained less popular. These included research in antimicrobial resistance transmission, the social and economic impact of antimicrobial resistance, and on the knowledge, attitudes and practices of the population on antimicrobial resistance. Furthermore, it identified a need to strengthen our knowledge and evidence base from the animal, environment and food sectors, due to the disproportionate number of research articles identified.

Through this review, research that may have contributed towards national guidelines or institutional practice were also identified, as well as the contribution of existing research towards Singapore's National Strategic Action Plan on Antimicrobial Resistance. Opportunities and recommendations for future research were also proposed based on the gaps identified from this review. These include the development of a national antimicrobial resistance research agenda in One Health, ensuring relevance of future antimicrobial resistance research to the National Strategic Action Plan, adopting a Delphi consensus to identify and prioritise research topics of significance to Singapore, dedicated grant calls for antimicrobial resistance research proposals from the One Health sectors, and ensuring sustained funding to drive the national antimicrobial resistance research agenda in One Health.

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REPORT ON SINGAPORE'S ANTIMICROBIAL RESISTANCE RESEARCH LANDSCAPE FROM 2009 TO 2019

(I) AIM

1. To provide an overview of the types of antimicrobial resistance (AMR) research conducted in Singapore during the period of January 2009 to December 2019.

(II) BACKGROUND

2. AMR is a globally recognised threat to human health, animals and the environment. The May 2015 World Health Assembly adopted the Global Action Plan on AMR with five objectives to address this global crisis [3].
3. In response, Singapore's National Strategic Action Plan (NSAP) on AMR was published on 1 Nov 2017, setting the framework for a national response to AMR, especially bacterial resistance to antibiotics. The plan aims to reduce the emergence and prevent the spread of drug resistant organisms through five core strategies: (1) Education; (2) Surveillance and Risk Assessment; (3) Research; (4) Prevention and Control of Infection; and (5) Optimisation of Antimicrobial Use [4].
4. The third core strategy, Research, aims to provide a better understanding of AMR and to provide information and evidence to support the other core strategies [4].
5. A priority area of action in AMR research highlighted in the NSAP is to understand the AMR research landscape in Singapore. This would identify and prioritise the research needs to ensure that key questions and evidence gaps are addressed to drive the national AMR agenda [4]. The review of Singapore's AMR research landscape was performed by the Antimicrobial Resistance Coordinating Office (AMRCO) at the National Centre for Infectious Diseases (NCID). The findings are presented in this report.

(III) METHODOLOGY

6. A comprehensive search strategy was adopted to retrieve articles from databases that index AMR research articles. Details on the identification of the final list of research articles, including the study selection and eligibility criteria, are described in [Annex A](#). Each research article was read in full to enable its assignment to the appropriate sector and research domain, and for a summary of the research conducted in each domain to be included in this report.
7. Research domains were determined through an expert consultation conducted in 2017. Domain definitions were further defined and refined throughout the process of this landscape review ([Textbox 1](#)). When research could potentially span multiple domains, they were assigned by consensus to the most appropriate single domain that represented its main research focus.

8. The research articles were also assigned to the appropriate One Health sector that the research was from (i.e. animal, environment, food or human). Assignment was based on the source of the isolate/sample or the application of the research. Articles were assigned to multiple sectors where suitable and appropriate.
9. The research articles were also assigned to the relevant NSAP core strategy where they contributed further information to advance the AMR agenda.

Textbox 1. Definition of research domains.

Knowledge, Attitudes, Practices [5, 6]

- Studies that investigated and/or discussed the level of understanding of AMR and antimicrobial use (AMU) by different populations, the individual perception and reaction towards AMR and AMU, and the actions that they take as a result.

Social and Economic Impacts

- Studies that analysed the impact of AMR on the community, the impact of AMR on healthcare costs, and the cost effectiveness of interventions against AMR. Social determinants of health were also considered.

Transmission [7]

- Studies that investigated the spread of AMR microorganisms or genes.

Microbiology

- Laboratory-based studies that investigated a specific aspect of the resistant microorganism that contributed to its resistance profile, increasing existing insights and knowledge.

Diagnostics [8]

- Studies that developed, investigated, or validated methods for identifying the presence of AMR or antimicrobials, or monitored the effect of interactions of antimicrobials with resistant microorganisms.

Intervention

- Studies that developed tools or methods to curb the spread or transmission of AMR pathogens and/or investigated their efficacy.

Surveillance

- Studies that reported the prevalence of AMR (microorganisms or genes), or presence of antimicrobials detected and their quantities. Studies that reported antimicrobial consumption are also included.

Public Health and Epidemiology

- Studies that performed systematic or data-driven analyses of the distribution (frequency, pattern) and determinants (causes, risk factors) of AMR in specific populations; Studies that analysed the impact of AMR on the health of a population as a whole.

Therapeutics

- Studies that developed and investigated new therapy(ies) for treating AMR, including but not limited to:
 - Drug repurposing;
 - Screening libraries for potential lead compounds;
 - Synthesis of compound series with antimicrobial activity against AMR microorganisms;
 - Investigating the mechanisms of action of the therapeutic / mechanisms of resistance mounted by the microorganisms;
 - Comparison of treatments in patients with conditions or infections caused by AMR microorganisms; or
 - Clinical trials.

(IV) SINGAPORE'S AMR RESEARCH LANDSCAPE

10. The review of Singapore's AMR research landscape identified a total of 741 articles that could be divided into three main categories:
 - a. Research studies conceived and led by investigators in Singapore (N=551);
 - b. Research studies conceived and led by investigators from foreign institutions, with a Singapore-based researcher involved (N=128); or
 - c. Research studies where Singapore contributed samples or data to increase knowledge across regions about AMR or AMU (N=62).
11. [Annex B](#) lists all articles identified from this review. Each article was assigned to the relevant research domain, the One Healthⁱ sector, and the core strategy to which the research findings contributed knowledge or evidence.

(A) AMR Research from Singapore, 2009-2019

Overview of Research Trends

12. The number of research articles conceived and led by investigators in Singapore increased from 17 in 2009 to 73 in 2019 (Annex C, [Figure 1](#)).
13. In terms of domain distribution of the research conducted, *Therapeutics* had the most number of research articles, followed by *Public Health and Epidemiology*, and *Surveillance*; Domains with the least articles were *Transmission, Social and Economic Impacts (SEI)* and *Knowledge, Attitudes, Practices (KAP)* (Annex B, Figure 2). All other research domains also saw regular annual publications except for the latter three (Annex C, [Figure 2](#)).
14. For research from a single sector, the *Human* sector (N=490) had the highest number of research articles, followed by the *Environment* sector (N=27). For cross-sector studies, research that straddled the *Human* and *Environment* sectors had the most articles (N=14) (Annex C, [Figure 3A](#)).
15. An increasing trend in the number of articles published from the *Human* sector was observed, from 15 in 2009 to 57 in 2019. AMR research published from the other sectors was sparse, until 2017, when more articles from the *Environment* and *Food* sectors were published (Annex B, [Figure 3B](#)).

ⁱ The One Health High-Level Expert Panel recently updated the definition of One Health as “an integrated, unifying approach that aims to sustainably balance and optimize the health of people, animals, and ecosystems. It recognizes the health of humans, domestic and wild animals, plants, and the wider environment (including ecosystems) are closely linked and interdependent. The approach mobilises multiple, disciplines, and communities at varying levels of society to work together to foster well-being and tackle threats to health and ecosystems, while addressing the collective need for healthy food, water, energy, and air, taking action on climate change and contributing to sustainable development. In Singapore, the One Health sectors include the human, animal, food and environment sectors.

Microorganisms of Interest, 2009-2019

16. AMR bacteria and their related genes were most studied compared to other microorganisms, with more gram-negative than gram-positive bacteria in the studies published between 2009 to 2019. Other AMR microorganisms (fungi, virus or parasites) had fewer studies across the same period under review (Annex C, [Figure 4](#)).
17. The World Health Organization (WHO) priority pathogen list for therapeutics highlighted three categories of AMR microorganisms to be prioritised for the research and development of novel therapeuticsⁱⁱ. Articles in the *Therapeutics* domain showed that there was already interest by researchers from Singapore to prioritise the development of novel therapeutics targeting the above categories. (Annex C, [Table 1](#)). Most of the *Therapeutics* studies were on carbapenem-resistant *Pseudomonas aeruginosa*, or carbapenem-resistant, extended-spectrum beta-lactamase (ESBL)-producing Enterobacteriales in the Priority 1 category, and on MRSAⁱⁱⁱ and VISA/VRSA^{iv} in the Priority 2 category.

AMR Research Output by Institutions, 2009-2019

18. There was significant research conducted by clinicians from public healthcare institutions (PHIs) between 2009 to 2019 on AMR, with most publications from the tertiary hospitals (Singapore General Hospital, National University Hospital, Tan Tock Seng Hospital), and some from the regional general hospitals (Changi General Hospital, Ng Teng Fong General Hospital, Khoo Teck Puat Hospital, KK Women's and Children's Hospital) (Annex C, [Table 2](#)).
19. Research on AMR from institutes of higher learning (IHLs) was dominated by National University of Singapore and Nanyang Technological University, and a small number of publications from the Duke-NUS Medical School, Singapore Institute of Technology, Singapore University of Technology & Design, and the polytechnics (Ngee Ann Polytechnic, Republic Polytechnic and Singapore Polytechnic). Research institutes (RIs) and ministries/agencies in Singapore also conducted AMR research, but had fewer publications compared to the PHIs and IHLs (Annex C, [Table 2](#)).
20. Research from the respective institutions were reflective of their research interests and fields of expertise. PHIs published more research that were in the *Surveillance* and *Public Health and Epidemiology* domains, while IHLs and RIs had strong expertise in *Therapeutics* and *Microbiology* research. PHIs and IHLs showed similar levels of expertise or interests in *Intervention* research (Annex C, [Figure 5](#)).
21. Lastly, most of the AMR research between 2009 and 2019 were funded by grants from or managed by ministries/agencies. Other sources of funding included grants from healthcare clusters, IHLs, or were funded internally (Annex C, [Table 3](#)).

ⁱⁱ "WHO publishes list of bacteria for which new antibiotics are urgently needed." 27 February 2017.
Link: <https://www.who.int/news/item/27-02-2017-who-publishes-list-of-bacteria-for-which-new-antibiotics-are-urgently-needed>

ⁱⁱⁱ *Staphylococcus aureus*, methicillin-resistant

^{iv} *Staphylococcus aureus*, vancomycin-intermediate or vancomycin-resistant

22. A summary of the research from each domain is elaborated in [Annex D](#).

(B) Involvement in Multinational Research Efforts

23. Singapore was also involved in multinational research efforts to address AMR, either through the contribution of samples or data, both in the region or globally, or in international collaborations, providing their expertise in research led by their foreign counterparts. When compared with research published by investigators from Singapore, Singapore's involvement in multinational research also increased (Annex E, [Figure 1](#)).

Contributing Samples to Multinational Studies

24. Between 2009 and 2019, Singapore contributed samples or data to 62 studies. No trend was observed in the studies published between 2009 and 2019 (Annex E, [Figure 1](#) “Sample contribution”; [Annex F](#)).

25. There were 57 studies from the human sector, four studies from the environment sector, and one human-animal cross-sector study.

26. Majority of the articles identified (N=55) were assigned to the *Surveillance* and *Public Health and Epidemiology* domains. The remaining articles were distributed across *Therapeutics* (N=2), *Transmission* (N=2), *Diagnostics* (N=2) and *Microbiology* (N=1).

International Collaborations on AMR, 2009-2019

27. Investigators based in Singapore participated in 128 collaborations between 2009 to 2019, and the trend in collaborations showed an increase from three studies in 2009 to 25 studies in 2019 (Annex E, [Figure 1](#) “International collaborations”).

28. International collaborations on AMR research showed a similar distribution in terms of the research domain assignment with research from Singapore (Annex E, [Figure 2](#)).

29. Researchers in Singapore collaborated with research groups from the high- (17 countries), upper middle- (4 countries) and lower middle- (6 countries) income countries. Countries with more than 10 research collaborations included Australia, the United States, the United Kingdom, Denmark and China (Annex E, [Figure 3](#)).

(V) AMR RESEARCH RELEVANT TO POLICY OR PRACTICE

30. The review of Singapore's AMR research landscape identified the following research topics that were relevant to Singapore's public health policies and practices. Some of the findings provided information on pneumococcal immunisation, and for improving practices in the hospital to prevent the spread of infections caused by AMR microorganisms.

(A) Pneumococcal Immunisation

31. There were three research articles that studied the pneumococcal disease burden and antibiotic resistance rates, as well as how the introduction of PCV-7, and subsequently PCV-13 and PPSV-23 impacted the disease burden.

32. Overall findings showed that the introduction of PCV-7 was followed by concomitant replacement of vaccine serotypes by non-vaccine serotypes circulating in the population. Furthermore, baseline resistance rates were high. Penicillin non-susceptible serotypes, especially in 19A, was observed following introduction of PCV-7. Resistance to erythromycin was also found to be gradually increasing [9].

33. In addition, mortality rates due to pneumococcal disease were high, at 19% and 3% of hospitalised elderly and children, respectively. Of the serotypes found in the study by Martinez-Vega et al. [9], 30% of serotypes identified were not covered by vaccines that were licensed in Singapore at the time of the study in 2013.

34. The period of the three studies coincided with the time at which PCV-7 and/or PCV-13 were introduced. However, it was unclear whether these studies directly contributed to the decision to have the PCV-7 immunisation programme implemented, and subsequently replaced by the newly approved and licensed PCV-13. Global experiences and data gathered from the US and Europe may have had some impact, since these countries reported the emergence of non-vaccine serotypes after introducing PCV-7, especially the high prevalence of the 19A serotype [10].

35. A nationwide pneumococcus surveillance programme was started in 2009 with the National Public Health Laboratory, along with mandatory reporting of invasive pneumococcal disease by clinicians [11]. This facilitated the evaluation of the effectiveness of the pneumococcal immunisation programme in reducing pneumococcal disease burden.

36. The overall high mortality rates also reflected the need to increase vaccine coverage for pneumococcal disease. PCV-13 is included in the National Childhood Immunisation Schedule. In the elderly, PCV-13 and PPSV-23 are recommended.

(B) Interventions in Healthcare Institutions

37. Several studies identified from this review focused on the impact of interventions implemented at PHIs. These could be divided into (1) interventions aimed at improving hand hygiene as part of infection prevention and control (IPC)

practices; and (2) engineering interventions to reduce the spread of AMR transmission when the source was identified.

Hygiene Practices at Healthcare Institutions

38. There were four studies that sought to assess the impact of institutional workflows that aimed to reduce hospital-associated infections by AMR microorganisms. These included the addition of intranasal octenidine to daily antiseptic bathing [12], alcohol hand-rubbing practices [13, 14] and the use of ward level targets to improve hand hygiene compliance rates [15]. The outcomes from the inclusion of these institutional practices reduced infection rates; hand hygiene compliance rates also improved with sustained results.
39. In a study that investigated *Elizabethkingia meningoseptica* infections, clinical staff were found to misuse hand hygiene sinks to dispose patient secretions or rinse re-usable patient care items. An urgent education programme was instituted to change the practice with temporary improvement reported. Room design and staff workflows were recommended to be optimised for patient safety as such lapses in procedures could put patients at risk [16].

Engineering Interventions at Healthcare Institutions

40. Three articles included in the review were on interventions aimed at removing the environmental source to curb AMR transmissions. The changes that were implemented were either already part of institutional workflows, or implemented after findings showed that the changes were beneficial:
 - a. Absence of carbapenem-resistant *Acinetobacter baumannii* (CRAB) found outside patients' wards showed that appropriate cleaning of high- and low-touch surfaces, provision of dedicated blood pressure cuffs for patients and increased accessibility to alcohol-based hand rubs were effective. A pilot project to use real-time whole genome sequencing (WGS) combined with epidemiological linkage analysis to track CRAB as they emerged also facilitated early institution of infection control measures to reduce further transmission [17].
 - b. Molecular typing analysis of MRSA and *Acinetobacter* spp. showed clonally related strains circulated between patients, the hands of healthcare professionals and the environment. As enhanced cleaning demonstrated reduced MRSA infections, systematic cleaning programmes were recommended to evaluate the hygiene standards in healthcare environments [18].
 - c. Following a *Pseudomonas aeruginosa* outbreak in a haematology ward, investigations showed that sinks with multiple grooves could harbour biofilms and were difficult to disinfect adequately. Changing the sink drainage system to one without grooves successfully terminated the outbreak. Thus, the institution replaced sinks in all clinical areas that housed immunosuppressed patients [19].

(VI) IDENTIFYING OPPORTUNITIES FOR AMR RESEARCH

41. This review of Singapore's AMR research landscape was the first to include the One Health sectors, in addition to the human sector. Besides providing the first overall glimpse into the current research emphasis, it also highlighted the areas where more research could be conducted.

(A) Research Relevant to the NSAP Core Strategies to Further Singapore's AMR Agenda

42. Besides Research, Singapore's NSAP on AMR consists of four other core strategies: Education; Surveillance and Risk Assessment; Prevention and Control of Infection; and Optimisation of Antimicrobial Use. Through this landscape review, we identified research currently in progress with potential to contribute relevant knowledge to these core strategies. At the same time, it also allowed us to propose research that would be needed but have yet to be conducted in Singapore ([Annex G](#)).

Education

43. Findings from research articles assigned to the *Knowledge, Attitudes, Practices* domain contributed important insights, as well as identified appropriate education materials that were needed to improve the population's knowledge and to correct the current misconceptions pointed out by the studies.
44. Information from relevant *Surveillance* or *Public Health and Epidemiology* studies could also provide a proxy measure on whether the relevant education interventions had any positive impact on AMR or AMU.
45. The development of content and use of the appropriate modality for education or public awareness campaigns is an important process, and robust and comprehensive user design and testing prior to launch are recommended. There were studies on interventions to increase knowledge but these were few; the mixed or poor outcomes reported also indicated that more work was required [20, 21]. Social and communication science approaches, which would consider how information is perceived by a different audiences, could be applied when developing the content in public outreach campaigns, as suggested by Huttner et al [22]. Based on this review, such studies were not identified.

Surveillance and Risk Assessment

46. Surveillance is an important part of public health efforts for prompt response to potential threats to public, animal and environmental health, to mitigate and prevent detrimental impacts, and for measuring the outcomes of programmes and initiatives launched as part of the NSAP. Almost all research articles that were assigned to the *Surveillance* and *Public Health and Epidemiology* domains were relevant to this core strategy. Studies in the *Transmission* domain could also contribute information to risk assessments, through provision of updated information by which resistant pathogens may spread.

47. Publications from the animal, environment and food sectors should be encouraged due to the low numbers identified in this review and to demonstrate the inter-relationship of AMR in One Health. From the environment and food sectors in particular, research to identify potential sources of AMR would be necessary, as their roles as reservoirs of *de novo* antimicrobial resistance development are not fully elucidated.
48. Integrated surveillance of AMR across the One Health sectors is a priority highlighted in Singapore's NSAP. In relation to this, researchers could further develop or harmonise existing diagnostic tools to aid integrated surveillance. Where possible, the development of diagnostic tools should also move from single sector application to potential cross-sector applications. Methodological studies for sample collection should also be followed-up, as one research had demonstrated that different combinations of anatomical sites gave rise to different sensitivities for the identification of target AMR pathogens [23]. Such lessons learnt from one sector could be applied to other sectors.

Research

49. Research within the NSAP is framed with the objective of improving our understanding of AMR. The research articles identified from this review largely satisfied this objective as the findings contributed towards our knowledge and understanding of AMR from multiple dimensions.
50. Moving forward, knowledge and evidence base acquired through such research should be applied and translated by the research community. An example would be the further development of promising novel treatment or intervention strategies that target drug-resistant microorganisms for practical applications.

Prevention and Control of Infection

51. Prevention and control of infection minimise the incidences and associated risks of emergence and spread of AMR. Most research studies in the *Intervention* domain provided information or engineered novel methods relevant to this core strategy. Studies from *Social and Economic Impacts*, such as a review of screening and isolation programme, also provided evidence of their effectiveness [24].
52. However, studies on vaccines to reduce infection by AMR pathogens were few. There were three studies that correlated the emergence of AMR *Streptococcus pneumoniae* following the introduction of the relevant vaccines. Although timely, they were unlikely to impact policies on vaccine implementation, but provided data that supported them.
53. There was also a lack of research identified from the animal sector. While guidelines were recently developed for pet vaccination and husbandry, and for prudent use of antimicrobials in companion animals (to include references NParks 2020 and 2021), their impact on AMR in animals are still unknown. Therefore, monitoring the uptake of pet vaccination and AMU rates in the animal sector are recommended. Coupled with surveillance of AMR in animals, data correlation would provide useful information that contribute to advancing Singapore's agenda on AMR.

Optimisation of Antimicrobial Use

54. Although research relevant to this core strategy could be identified from almost all research domains, majority of the research were also from the human sector. Research to optimise AMU in the animal sector could be strengthened, with similar studies conducted in the human sector replicated in animals where appropriate.
55. A regulatory framework for veterinary drug registration and supply chain control is under development. When implemented, AMU could be monitored to contribute towards surveillance of the usage trends of antimicrobials, as well as encourage a parallel optimal prescription and usage by veterinarians and pet owners.
56. A cornerstone of improving AMU would be through education. However, only two studies were identified from this review. Poor or mixed outcomes were obtained from current educational efforts to improve AMU. Though there were limitations mentioned in the studies conducted, they provided avenues for improvement of similar studies in the future.

(B) Opportunities in One Health AMR Research

57. Three research domains that had the least research output or articles provided opportunities for future research. These were *Transmission, Social and Economic Impacts, and Knowledge, Attitudes, Practices* on AMR. Another observation made was the small number of research articles from the animal, environment and food sectors, as well as the number of cross-sector research. While AMR research from the environment, food and animal sectors were increasingly being published from 2017, this was only 7.8% of the articles from the human sector. Research from the other sectors should be encouraged to increase our understanding and provide evidence-based approaches for addressing AMR. Cross-sector research is also needed to increase our knowledge of the interrelatedness of AMR across the One Health sectors.

(C) Others

58. Separately, some of the research articles also highlighted existing challenges that researchers faced in order to drive research to address AMR. These challenges provide opportunities for development into future research questions or topics. Such development could take the approach by Tudor Car et al. [25], where challenges identified from a review of the literature were formulated into potential research questions to highlight the areas of need to the research community.

(VII) RECOMMENDATIONS

59. Recommendations proposed are aligned with the opportunities identified from Section (VI), which echo opportunities proposed by others [1, 26]

(A) National Research Agenda on AMR in One Health

60. **A national AMR research agenda that incorporates One Health could be formulated to advance Singapore's AMR agenda.** This landscape review not only identified research that were relevant to each of the NSAP core strategies in Section VI (A), it also identified opportunities. A list of research questions or topics could be formulated to address the gaps; some of these research questions are currently being collated from this review ([Annex G](#)), and additional questions will be obtained through expert solicitation from the One Health sectors for comprehensiveness. The final list will contribute to the development of Singapore's AMR research agenda with emphasis on the priority topics to be funded for research.

61. **Future AMR research in Singapore should also consider their relevance to the NSAP core strategies.** This review showed that research articles identified could contribute valuable information and insights to multiple NSAP core strategies ([Annex B](#)). Therefore, the planning for future research relevant to the advancement of Singapore's AMR agenda should take this aspect into consideration to maximise the potential of each research proposal.

62. **A Delphi consensus [27] could be adopted as part of a priority setting exercise to identify research topics that are of significance to Singapore's One Health AMR research agenda.** The Tripartite collaboration between WHO, Food and Agricultural Organization of the United Nations and the World Organisation for Animal Health is working towards the development of a global research agenda on AMR, and had invited stakeholders to participate in a survey on research questions for OH AMR research that was launched from Jul – Sep 2021 [28]. Singapore could take guidance from this approach to identify relevant research questions that are suitable for Singapore's needs.

(B) Funding for AMR Research from the One Health Sectors

63. Most of the current research addresses AMR from a human perspective. To emphasise the interrelatedness of all One Health sectors and AMR, **grant calls dedicated to AMR research proposals from the One Health sectors should continue.** AMR genes from the environment known to impact human health include ESBL and New Delhi metallo-beta-lactamase (NDM-1) genes [29, 30]. These two genes show that there is a feedback loop between infection, antibiotic use and the development of antibiotic resistance [31]. Identifying and understanding the transmission of such genes from the environment, food and animal sectors would be necessary for planning mitigation measures. Such research is needed to expand our knowledge.

64. Further funding to drive the national AMR research agenda in One Health is needed. The One Health AMR Research Programme (OHARP)^v was conceived to drive cross-sectoral research collaborations on AMR and to identify and dedicate research funding to the priority domains identified through an expert consultation conducted in 2017. Through grant calls launched in 2021 and 2022, four grants were awarded to specific projects that addressed cross-sector transmission of AMR, and knowledge, attitudes and practices of Singaporeans towards AMR. Separately, a social and economic impact study was also commissioned in 2022. With the success achieved, further funding would also be needed to invest in the priority research questions or topics identified from this review.

^v One Health Antimicrobial Research Programme (OHARP) Grant. Website link:
[https://www.nmrc.gov.sg/grants/competitive-research-grants/one-health-antimicrobial-resistance-research-programme-\(oharp\)-grant](https://www.nmrc.gov.sg/grants/competitive-research-grants/one-health-antimicrobial-resistance-research-programme-(oharp)-grant)

(VIII) CONCLUSION

65. This report provides the first comprehensive review of Singapore's AMR research landscape on the types of AMR research conducted in Singapore, and their findings, between 2009 and 2019. Recommendations made were based on the gaps and opportunities identified from current research and presented for adoption to drive the national AMR research agenda.
66. Being the first time that the AMR research in Singapore was surveyed, this review used key word search of relevant databases to identify as many of the relevant research or review articles as possible. With increasing emphasis of the One Health approach in AMR, Singapore's AMR research landscape is expected to evolve as the research focus further develops and matures. Thus, AMRCO will conduct regular reviews and provide updates based on the information collated. Future reviews may also expand and improve the current processes used to increase comprehensiveness where needed.
67. Lastly, as we limited the review to published articles from Singapore, some areas of research that have the potential to advance Singapore's AMR agenda may have been excluded. An example is the development of viable alternatives to reduce overall use of antimicrobials in livestock, which is an area with limited research in Singapore. Future reviews should also compare with AMR research conducted globally and consider how these research could impact on Singapore's NSAP on AMR.

(IX) List of Annexes:

- Annex A Details of identification of research articles.
- Annex B List of relevant research articles identified from this landscape review.
- Annex C Figures and tables describing overview of AMR research from Singapore.
- Annex D Summary of AMR research from Singapore, grouped according to research domains.
- Annex E Figures describing involvement of researchers in multinational research efforts to address AMR.
- Annex F Summary of the types of AMR research that Singapore contributed samples or data to.
- Annex G Mapping research relevant to NSAP core strategies and opportunities for future research.

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Annex A. Details of identification of research articles ([Section III](#))

Search strategy

Search strategy for NCBI Pubmed, and adapted accordingly for Embase (Ovid), Scopus, Cumulative Index to Nursing and Allied Health Literature (CINAHL, Ebsco), Global Health (Ovid), Medline (Ovid) and Web of Science.

- 1 (((((anti*) OR (anti-b*)) OR (anti-m*)) OR (anti-f*)) OR (anti-v*)) OR (microb*)) OR (antimicrob*)) OR (anti-microb*)
- 2 ((resistant) OR (resistance)) OR (resist*)
- 3 #1 OR #2
- 4 (Singapore) OR (Singaporean)
- 5 #3 AND #4
- 6 #5 AND ((“2009/01/01”[Date – Publication] : “2019/12/31”[Date – Publication]))

Eligibility criteria

The table lists the criteria when considering an article for inclusion in this landscape review.

	Include	Exclude
Year of publication	2009 to 2019, inclusive	Preprints and Epub ahead of print between 2009 to 2019
Content type	Research articles Review articles	Editorials Poster/Conference abstracts Commentary Opinion pieces Guidelines Reports Book or Book Chapters
Country	Studies conducted by authors whose primary affiliation is with a Singapore/local institution Studies with samples or data from Singapore	Studies not conducted or involving authors whose primary affiliation is with a Singapore/local institution
Research scope	Research focused on addressing AMR, including but not limited to the following: <ul style="list-style-type: none">- Identifying methods to reduce/prevent AMR- Treating infections due to AMR microorganisms- Developing methods to improve the removal of antimicrobials- Understanding the levels of AMU or AMR	Studies describing research on general infectious diseases without an AMR focus

Annex B. Relevant research articles identified from review of Singapore's antimicrobial resistance research landscape, 2009-2019

Table 1. List of AMR research and review articles published by researchers from Singapore between 2009 and 2019.

Articles were assigned to the most relevant research domain, sector(s) and their contribution to the NSAP core strategy(ies).

Article ID	Sector(s)	Research Domain	Core Strategy(ies)
2009, Chan KS [32]	Human	Surveillance	Surveillance & Risk Assessment
2009, Chuwa EWL [33]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2009, Deepak RN [34]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2009, Donaldson AD [35]	Human	Therapeutics	Research
2009, Fan C [36]	Environment	Microbiology	Research
2009, Fong RKC [37]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2009, Ho YM [38]	Human	Microbiology	Research
2009, Koh TH [39]	Human, Animal	Public Health and Epidemiology	Surveillance & Risk Assessment
2009, Lee CC [40]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2009, Lim T-P [41]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2009, Liu L [42]	Human	Therapeutics	Research
2009, Luan J [43]	Human	Diagnostics	Research; Optimisation of Antimicrobial Use
2009, Prabakaran M [44]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2009, Tan TY [45]	Human	Diagnostics	Research
2009, Teo JWP [46]	Human	Microbiology	Research
2009, Tin S [47]	Human	Therapeutics	Research
2009, Wang S-Q [48]	Human	Therapeutics	Research
2010, Cheow WS [49]	Human	Therapeutics	Research
2010, Cheow WS [50]	Human	Therapeutics	Research
2010, Ch'ng J-H [51]	Human	Microbiology	Research
2010, Ding C [52]	Environment	Public Health and Epidemiology	Surveillance & Risk Assessment
2010, Donaldson AD [53]	Human	Surveillance	Surveillance & Risk Assessment
2010, Gan LSH [54]	Human	Diagnostics	Surveillance & Risk Assessment; Research
2010, Ho J [55]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment; Research
2010, Hsu L-Y [56]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2010, Hsu L-Y [57]	Human	Therapeutics	Research
2010, Husain N [58]	Human	Therapeutics	Research
2010, Inoue M [59]	Human	Therapeutics	Research
2010, Jayaraman P [60]	Human	Therapeutics	Research
2010, Koh TH [61]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2010, Koh TH [62]	Human	Surveillance	Surveillance & Risk Assessment
2010, Kurup A [63]	Human	Intervention	Surveillance & Risk Assessment; Prevention & Control of Infection
2010, Lim LG [64]	Human	Diagnostics	Research
2010, Lim PL [65]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2010, Liu R [66]	Human	Diagnostics	Research
2010, Ng ES-T [67]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2010, Nzila A [68]	Human	Therapeutics	Research
2010, Ong DCT [69]	Human	Diagnostics	Research
2010, Rottmann M [70]	Human	Therapeutics	Research
2010, Samy RP [71]	Human	Therapeutics	Research
2010, Soe WM [72]	Human	Therapeutics	Research
2010, Soe WM [73]	Human	Therapeutics	Research
2010, Sun Y-J [74]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2010, Tan TY [75]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2010, Tan TY [76]	Human	Surveillance	Surveillance & Risk Assessment
2010, Vasoo S [10]	Human	Surveillance	Surveillance & Risk Assessment
2010, Wijaya L [77]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2010, Wong L [78]	Human	Therapeutics	Research
2010, Zhou C [79]	Human	Therapeutics	Research
2011, Bahadin J [80]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2011, Chan HLE [81]	Human	Surveillance	Surveillance & Risk Assessment
2011, Chanratita N [82]	Human	Therapeutics	Research

Article ID	Sector(s)	Research Domain	Core Strategy(ies)
2011, Cheow WS [83]	Human	Therapeutics	Research
2011, Ch'ng J-H [84]	Human	Therapeutics	Research
2011, Fan C [85]	Environment	Microbiology	Research
2011, Hsu L-Y [86]	Human	Surveillance	Surveillance & Risk Assessment
2011, Husain N [87]	Human	Therapeutics	Research
2011, Kyaw BM [88]	Human	Therapeutics	Research
2011, Kyaw BM [89]	Human	Therapeutics	Research
2011, Lee CY [90]	Human	Therapeutics	Research
2011, Lee HK [91]	Human	Diagnostics	Research; Optimisation of Antimicrobial Use
2011, Leung GYC [92]	Human	Therapeutics	Research
2011, Liew YX [93]	Human	Surveillance	Surveillance & Risk Assessment
2011, Liew YX [94]	Human	Diagnostics	Research; Optimisation of Antimicrobial Use
2011, Liew YX [95]	Human	Surveillance	Surveillance & Risk Assessment
2011, Lim PL [96]	Human	Surveillance	Surveillance & Risk Assessment
2011, Lim T-P [97]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2011, Lim T-P [98]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2011, Lingegowda PB [99]	Human	Therapeutics	Research
2011, Mirza H [100]	Human, Animal	Diagnostics	Research; Optimisation of Antimicrobial Use
2011, Mok S [101]	Human	Therapeutics	Research
2011, My NH [102]	Human	Therapeutics	Research
2011, Nederberg F [103]	Human	Therapeutics	Research
2011, Ng LSY [104]	Human, Environment	Intervention	Research; Prevention & Control of Infection
2011, Ong DCT [105]	Human	Diagnostics	Research
2011, Pada SMK [106]	Human	Social and Economic Impacts	Research
2011, Phua CK [107]	Human	Therapeutics	Research
2011, Saeidi N [108]	Human	Therapeutics	Research
2011, Samy RP [109]	Human	Therapeutics	Research
2011, Sim SH [110]	Human, Environment	Therapeutics	Research
2011, Soe WM [111]	Human	Therapeutics	Research
2011, Suhaila M [112]	Human	Therapeutics	Research
2011, Tan TY [113]	Human	Diagnostics	Research
2011, Teo BW [114]	Human	Intervention	Surveillance & Risk Assessment; Prevention & Control of Infection
2011, Teo BW [115]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2011, Teo JWP [116]	Human	Surveillance	Surveillance & Risk Assessment
2011, Vasoo S [11]	Human	Surveillance	Surveillance & Risk Assessment
2011, Xing B [117]	Human	Therapeutics	Research
2012, Bai Y [118]	Human	Therapeutics	Research
2012, Baln MND [119]	Human	Surveillance	Surveillance & Risk Assessment
2012, Cai Y [120]	Human	Surveillance	Surveillance & Risk Assessment
2012, Chen C [121]	Human, Environment	Intervention	Research; Prevention & Control of Infection
2012, Chen T [122]	Human	Intervention	Research; Prevention & Control of Infection
2012, Chen Y-T [123]	Human	Surveillance	Surveillance & Risk Assessment
2012, Cheong CSJ [124]	Human	Surveillance	Surveillance & Risk Assessment
2012, Chew KK [125]	Human	Diagnostics	Research; Optimisation of Antimicrobial Use
2012, Chien JMF [126]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2012, Choudhury S [127]	Human	Surveillance	Surveillance & Risk Assessment
2012, Choudhury S [128]	Human	Diagnostics	Research; Optimisation of Antimicrobial Use
2012, Chow ALP [23]	Human	Diagnostics	Surveillance & Risk Assessment; Research
2012, Chow ALP [129]	Human	Intervention	Prevention & Control of Infection
2012, Cui Y [130]	Human	Therapeutics	Research
2012, Du H [131]	Human	Microbiology	Research
2012, Fukushima K [132]	Human	Therapeutics	Research
2012, Han N [133]	Human	Therapeutics	Research
2012, Huang Y [134]	Human	Therapeutics	Research
2012, Kanagarajan V [135]	Human	Therapeutics	Research
2012, Koh TH [136]	Human	Surveillance	Surveillance & Risk Assessment
2012, Koh TH [137]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment

Article ID	Sector(s)	Research Domain	Core Strategy(ies)
2012, Kyaw BM [138]	Human	Therapeutics	Research
2012, Win M-K [139]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2012, Lee ASG [140]	Human	Diagnostics	Research; Optimisation of Antimicrobial Use
2012, Lee CK [141]	Human	Diagnostics	Surveillance & Risk Assessment; Research
2012, Liew YX [142]	Human	Intervention	Optimisation of Antimicrobial Use
2012, Liew YX [143]	Human	Intervention	Optimisation of Antimicrobial Use
2012, Ling ML [14]	Human	Intervention	Prevention & Control of Infection
2012, Lye DCB [144]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2012, Ng ES-T [145]	Human	Social and Economic Impacts	Research
2012, Poon LM [146]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2012, Sarathy JP [147]	Human	Microbiology	Research
2012, Shao Q [148]	Human	Intervention	Research; Prevention & Control of Infection
2012, Teo JQM [149]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2012, Teo JQM [150]	Human	Intervention	Optimisation of Antimicrobial Use
2012, Teo JWP [151]	Human	Surveillance	Surveillance & Risk Assessment
2012, Vasoo S [152]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2012, Venkatachalam I [153]	Human	Surveillance	Surveillance & Risk Assessment
2012, Verrall AJ [154]	Human	Therapeutics	Research
2012, Wozniak M [155]	Human	Therapeutics	Research
2012, Yeo C-L [156]	Human	Intervention	Optimisation of Antimicrobial Use
2012, Zou G [157]	Human	Microbiology	Research
2013, Ang TL [158]	Human	Therapeutics	Research
2013, Balm MND [159]	Human	Surveillance	Surveillance & Risk Assessment
2013, Balm MND [160]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2013, Balm MND [161]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2013, Balm MND [16]	Human	Transmission	Surveillance & Risk Assessment; Prevention & Control of Infection
2013, Chee CBE [162]	Human	Transmission	Surveillance & Risk Assessment
2013, Chin W [163]	Human	Therapeutics	Research
2013, Ch'ng J-H [164]	Human	Therapeutics	Research
2013, Chong C-W [165]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2013, Chow WL [166]	Environment	Intervention	Research; Prevention & Control of Infection
2013, Chua SL [167]	Human	Therapeutics	Research
2013, Fisher DA [168]	Human	Intervention	Prevention & Control of Infection
2013, Fukushima K [169]	Human	Therapeutics	Research
2013, Goh S [170]	Human	Transmission	Research
2013, Grant D [171]	Human	Transmission	Surveillance & Risk Assessment
2013, Heng YK [172]	Human	Therapeutics	Research
2013, Hon PY [173]	Human	Diagnostics	Research
2013, Koh J-J [174]	Human	Therapeutics	Research
2013, Koh TH [175]	Human	Surveillance	Surveillance & Risk Assessment
2013, Lee GH [176]	Human	Therapeutics	Research
2013, Lee GH [177]	Human	Therapeutics	Research
2013, Lee LK [178]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2013, Li Y [179]	Human	Intervention	Research; Prevention & Control of Infection
2013, Liew YX [180]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2013, Lim CLL [181]	Human	Public Health	Surveillance & Risk Assessment; Optimisation of Antimicrobial Use
2013, Lim S-G [182]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2013, Ling ML [19]	Human, Environment	Transmission	Surveillance & Risk Assessment; Prevention & Control of Infection
2013, Marimuthu K [183]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2013, Molton JS [184]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2013, Ng VWL [185]	Human	Therapeutics	Research
2013, Ong ZY [186]	Human	Intervention	Research; Prevention & Control of Infection
2013, Qiu G [187]	Environment	Intervention	Research
2013, Russell B [188]	Human	Diagnostics	Research
2013, Samy RP [189]	Human	Therapeutics	Research
2013, Samy RP [190]	Human	Therapeutics	Research
2013, Sarathy JP [191]	Human	Microbiology	Research
2013, Sarathy JP [192]	Human	Microbiology	Research
2013, Seah J [193]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use

Article ID	Sector(s)	Research Domain	Core Strategy(ies)
2013, Shao Q [194]	Human	Diagnostics	Research
2013, Sim JHC [195]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2013, Soon MML [196]	Human	Social and Economic Impacts	Research
2013, Tan PS [197]	Human	Therapeutics	Research
2013, Tan SY-Y [198]	Human	Microbiology	Research
2013, Tan TY [18]	Human, Environment	Transmission	Surveillance & Risk Assessment
2013, Teo JQM [199]	Human	Surveillance	Surveillance & Risk Assessment
2013, Teo JWP [200]	Human	Surveillance	Surveillance & Risk Assessment
2013, Teo JWP [201]	Human	Surveillance	Surveillance & Risk Assessment
2013, Vasudevan A [202]	Human	Therapeutics	Research
2013, Vasudevan A [203]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2013, Verrall AJ [204]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2013, Win M-K [205]	Human	Diagnostics	Research
2013, Xia E [206]	Human	Surveillance	Surveillance & Risk Assessment
2013, Xiong P [207]	Human	Intervention	Research; Prevention & Control of Infection
2013, Yang Y [208]	Human	Therapeutics	Research
2013, Yokokawa F [209]	Human	Therapeutics	Research
2013, Yuan X [210]	Human	Intervention	Research; Prevention & Control of Infection
2013, Zou H [211]	Human	Therapeutics	Research
2014, Bandyopadhyay S [212]	Human	Therapeutics	Research
2014, Bayen S [213]	Environment	Diagnostics	Surveillance & Risk Assessment; Research
2014, Castillo CFG [214]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2014, Chen HH [215]	Human	Therapeutics	Research
2014, Ch'ng J-H [216]	Human	Therapeutics	Research
2014, Chow JY [217]	Human	Intervention	Research; Prevention & Control of Infection
2014, Coady DJ [218]	Human	Therapeutics	Research
2014, Deng C-L [219]	Human	Therapeutics	Research
2014, Deng Y [220]	Human	Therapeutics	Research
2014, Gopal P [221]	Human	Therapeutics	Research
2014, Hon PY [222]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2014, Jauneikaite E [223]	Human	Surveillance	Surveillance & Risk Assessment
2014, Khara JS [224]	Human	Therapeutics	Research
2014, Koh TH [225]	Human	Surveillance	Surveillance & Risk Assessment
2014, La M-V [226]	Human	Surveillance	Surveillance & Risk Assessment
2014, Lescar J [227]	Human	Therapeutics	Research
2014, Li X [228]	Human	Intervention	Research; Prevention & Control of Infection
2014, Liu S [229]	Human	Therapeutics	Research
2014, Loh CCY [230]	Human	Diagnostics	Research
2014, Mani V [231]	Human	Diagnostics	Research; Optimisation of Antimicrobial Use
2014, Marimuthu K [232]	Human	Surveillance	Surveillance & Risk Assessment
2014, Marimuthu K [233]	Human	Intervention	Prevention & Control of Infection
2014, Marimuthu K [234]	Human	Intervention	Prevention & Control of Infection
2014, Mok S [235]	Human	Therapeutics	Research
2014, Ng TM [236]	Human	Intervention	Optimisation of Antimicrobial Use
2014, Ng TM [237]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2014, Ng VWL [238]	Human	Therapeutics	Research
2014, Ong ZY [239]	Human	Therapeutics	Research
2014, Samy RP [240]	Human	Therapeutics	Research
2014, Seah XVF [241]	Human	Intervention	Optimisation of Antimicrobial Use
2014, Shekar S [242]	Human	Therapeutics	Research
2014, Singhal A [243]	Human	Therapeutics	Research
2014, Song M [244]	Human	Therapeutics	Research
2014, Tan MW [245]	Human	Public Health and Epidemiology	Research; Optimisation of Antimicrobial Use
2014, Tan TT [246]	Human, Environment	Transmission	Surveillance & Risk Assessment
2014, Tan YE [247]	Human	Diagnostics	Surveillance & Risk Assessment; Research
2014, Tang SS [248]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2014, Teo JWP [249]	Human, Environment	Transmission	Surveillance & Risk Assessment
2014, Teo JWP [250]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment

Article ID	Sector(s)	Research Domain	Core Strategy(ies)
2014, Vasudevan A [251]	Human	Diagnostics	Research; Optimisation of Antimicrobial Use
2014, Venkatachalam I [252]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2014, Wang Y [253]	Human	Therapeutics	Research
2014, Wen H [254]	Human	Microbiology	Research
2014, Wozniak M [255]	Human	Diagnostics	Research
2014, Wu Z [256]	Human	Microbiology	Research
2014, Yeoh LY [257]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2014, Young BE [258]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2014, Zhou YP [259]	Human	Transmission	Surveillance & Risk Assessment
2014, Zhou YP [260]	Human	Therapeutics	Research
2015, Anusha S [261]	Human	Therapeutics	Research
2015, Cai Y [262]	Human	Diagnostics	Research; Optimisation of Antimicrobial Use
2015, Chee CBE [263]	Human	Transmission	Surveillance & Risk Assessment; Research
2015, Cheng J [264]	Human	Therapeutics	Research
2015, Cherng BPZ [265]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2015, Ch'ng JH [266]	Human	Microbiology	Research
2015, Choudhury S [267]	Human	Therapeutics	Research
2015, Chow ALP [268]	Human	Knowledge, Attitudes, Practices	Education; Optimisation of Antimicrobial Use
2015, Chow ALP [269]	Human	Intervention	Optimisation of Antimicrobial Use
2015, Chua AP-G [270]	Human	Therapeutics	Research
2015, Chua NGS [271]	Human	Therapeutics	Research
2015, Chung SJ [272]	Human	Therapeutics	Research
2015, Duan R [273]	Human	Therapeutics	Research
2015, Feng G [274]	Human	Therapeutics	Research
2015, Gopal P [275]	Human	Therapeutics	Research
2015, Harris PNA [276]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2015, Harris PNA [277]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2015, Harris PNA [278]	Human	Intervention	Prevention & Control of Infection
2015, Haver HL [279]	Human	Therapeutics	Research
2015, Ho HJ [13]	Human	Intervention	Prevention & Control of Infection
2015, Hsu L-Y [280]	Human	Transmission	Surveillance & Risk Assessment
2015, Koh J-J [281]	Human	Therapeutics	Research
2015, Koh J-J [282]	Human	Therapeutics	Research
2015, Koh TH [283]	Human, Environment	Surveillance	Surveillance & Risk Assessment
2015, Kumar A [284]	Human, Environment	Microbiology	Research
2015, Lau QY [285]	Human	Therapeutics	Research
2015, Lau QY [286]	Human	Therapeutics	Research
2015, Lau QY [287]	Human	Therapeutics	Research
2015, Lee HK [288]	Human	Surveillance	Surveillance & Risk Assessment
2015, Lew KY [289]	Human	Intervention	Optimisation of Antimicrobial Use
2015, Li J [290]	Human	Therapeutics	Research
2015, Liew YX [291]	Human	Intervention	Optimisation of Antimicrobial Use
2015, Liew YX [292]	Human	Social and Economic Impacts	Research
2015, Lim K [293]	Human	Intervention	Research; Prevention & Control of Infection
2015, Lim T-P [294]	Human	Therapeutics	Research
2015, Lim T-P [295]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2015, Ling ML [296]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2015, Loo LW [297]	Human	Intervention	Optimisation of Antimicrobial Use
2015, Lu S [298]	Human	Intervention	Research; Prevention & Control of Infection
2015, Manjunatha UH [299]	Human	Therapeutics	Research
2015, Ng C [300]	Environment	Surveillance	Surveillance & Risk Assessment
2015, Ng LSY [301]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2015, Ng PS [302]	Human	Therapeutics	Research
2015, Ng TM [303]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2015, Paton NI [304]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2015, Phoon YW [305]	Human	Surveillance	Surveillance & Risk Assessment
2015, Samy RP [306]	Human	Therapeutics	Research

Article ID	Sector(s)	Research Domain	Core Strategy(ies)
2015, Samy RP [307]	Human	Therapeutics	Research
2015, Seneviratne CJ [308]	Human	Surveillance	Surveillance & Risk Assessment
2015, Tan S [309]	Human	Therapeutics	Research
2015, Tang SSL [310]	Human	Surveillance	Surveillance & Risk Assessment
2015, Teng CB [311]	Human	Intervention	Optimisation of Antimicrobial Use
2015, Teo JWP [312]	Human	Surveillance	Surveillance & Risk Assessment
2015, Vasoo S [313]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2015, Vasoo S [314]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2015, Vasudevan A [315]	Human	Social and Economic Impacts	Research
2015, Win M-K [24]	Human	Social and Economic Impacts	Research; Prevention & Control of Infection
2015, Yi X [316]	Environment	Diagnostics	Surveillance & Risk Assessment; Research
2015, Yoon BK [317]	Human	Therapeutics	Research
2016, Ang TL [318]	Human	Surveillance	Surveillance & Risk Assessment
2016, Ariyasu S [319]	Human	Diagnostics	Research
2016, Arora S [320]	Human	Therapeutics	Research
2016, Ashajyothi C [321]	Human	Therapeutics	Research
2016, Aung KT [322]	Food	Surveillance	Surveillance & Risk Assessment
2016, Aung TT [323]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2016, Boudhar A [324]	Human	Therapeutics	Research
2016, Boudhar A [325]	Human	Therapeutics	Research
2016, Cai B [326]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2016, Cai Y [327]	Human	Intervention	Optimisation of Antimicrobial Use
2016, Cai Y [328]	Human	Diagnostics	Research; Optimisation of Antimicrobial Use
2016, Cai Y [329]	Human	Therapeutics	Research
2016, Cao Y [330]	Human	Therapeutics	Research
2016, Chen YT [331]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2016, Chen Y-T [332]	Human	Transmission	Research
2016, Chia G [333]	Human	Intervention	Prevention & Control of Infection
2016, Chow ALP [334]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2016, Chow ALP [335]	Human	Intervention	Optimisation of Antimicrobial Use
2016, Chua SL [336]	Human	Therapeutics	Research
2016, Fisher DA [337]	Human, Environment	Intervention	Prevention & Control of Infection
2016, Ghode P [338]	Human	Therapeutics	Research
2016, Gopal P [339]	Human	Therapeutics	Research
2016, Hemu X [340]	Human	Therapeutics	Research
2016, Ho HJ [341]	Human	Transmission	Surveillance & Risk Assessment
2016, Husain N [342]	Human	Therapeutics	Research
2016, Isenman H [343]	Human	Intervention	Prevention & Control of Infection
2016, Isenman H [344]	Human	Intervention	Prevention & Control of Infection
2016, Khara JS [345]	Human	Therapeutics	Research
2016, Khong WX [346]	Human	Transmission	Surveillance & Risk Assessment
2016, Khong WX [347]	Human	Transmission	Surveillance & Risk Assessment; Research
2016, Koh J-J [348]	Human	Therapeutics	Research
2016, Kong YL [349]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2016, Lakshminarayanan R [350]	Human	Therapeutics	Research
2016, Le T-H [351]	Environment	Surveillance	Surveillance & Risk Assessment
2016, Lee SH [352]	Human	Therapeutics	Research
2016, Li M [353]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2016, Lim EJZ [354]	Food	Surveillance	Surveillance & Risk Assessment
2016, Lim MY-X [355]	Human	Therapeutics	Research
2016, Lim WS [356]	Human	Therapeutics	Research
2016, Low A [357]	Environment	Surveillance	Surveillance & Risk Assessment
2016, Mandakhalikar KD [358]	Human	Intervention	Research; Prevention & Control of Infection
2016, Marimuthu K [359]	Human	Intervention	Prevention & Control of Infection
2016, Ng TM [360]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2016, Pan DST [361]	Human	Knowledge, Attitudes, Practices	Education
2016, Seneviratne CJ [362]	Human	Microbiology	Research
2016, Song CT [363]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment

Article ID	Sector(s)	Research Domain	Core Strategy(ies)
2016, Su CT-T [364]	Human	Therapeutics	Research
2016, Teng CP [365]	Human	Therapeutics	Research
2016, Teo JQM [366]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2016, Teo JWP [367]	Human	Surveillance	Surveillance & Risk Assessment
2016, Teo JWP [368]	Human	Diagnostics	Research
2016, Tran NH [369]	Environment	Diagnostics	Surveillance & Risk Assessment; Research
2016, Tran NH [370]	Environment	Intervention	Research
2016, Truong T [371]	Human	Therapeutics	Research
2016, Tun ZM [372]	Human	Knowledge, Attitudes, Practices	Education
2016, Wee KB [373]	Human	Therapeutics	Research
2016, Wong EHH [374]	Human	Therapeutics	Research
2016, Wong JGX [375]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2016, Teo JQM [376]	Human	Surveillance	Surveillance & Risk Assessment
2017 Lee MHM [20]	Human	Intervention	Education
2017 Zhang J [377]	Food, Environment	Intervention	Research
2017, Ang MLT [378]	Human	Therapeutics	Research
2017, Aung KT [379]	Food, Environment	Surveillance	Surveillance & Risk Assessment
2017, Aw J [380]	Human	Diagnostics	Research
2017, Aziz DB [381]	Human	Therapeutics	Research
2017, Cai Y [382]	Human	Surveillance	Surveillance & Risk Assessment; Optimisation of Antimicrobial Use
2017, Cai Y [383]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2017, Chee CBE [384]	Human	Therapeutics	Research
2017, Chen S [385]	Human	Diagnostics	Research
2017, Chew KL [386]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2017, Chew KL [387]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2017, Chew KL [388]	Human	Public Health	Surveillance & Risk Assessment
2017, Chew KL [389]	Human	Diagnostics	Surveillance & Risk Assessment; Research
2017, Chin JSF [390]	Human	Diagnostics	Research
2017, Chong SM [391]	Food, Animal	Surveillance	Surveillance & Risk Assessment
2017, Chow ALP [392]	Human	Transmission	Surveillance & Risk Assessment
2017, Gopal P [393]	Human	Therapeutics	Research
2017, Gopal P [394]	Human	Therapeutics	Research
2017, Hou Z [395]	Human	Therapeutics	Research
2017, Hsu L-Y [396]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2017, Hsu L-Y [397]	Human, Animal	Transmission	Surveillance & Risk Assessment
2017, Khara JS [398]	Human	Therapeutics	Research
2017, Kityo Cissy [399]	Human	Therapeutics	Research
2017, Lee TH [400]	Human	Knowledge, Attitudes, Practices	Education
2017, Li PQ [401]	Human	Diagnostics	Research
2017, Lin S [402]	Human	Therapeutics	Research
2017, Lin S [403]	Human	Therapeutics	Research
2017, Lin S [404]	Human	Therapeutics	Research
2017, Liu S [405]	Human	Therapeutics	Research
2017, Loo LH [406]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2017, Marimuthu K [407]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2017, Ng C [408]	Environment	Surveillance	Surveillance & Risk Assessment
2017, Ng SMS [409]	Human	Therapeutics	Research
2017, Ng SMS [410]	Human	Therapeutics	Research
2017, Ng SMS [411]	Human	Therapeutics	Research
2017, Paton NI [412]	Human	Diagnostics	Research
2017, Peng J [413]	Human	Therapeutics	Research
2017, Pu Y [414]	Human	Therapeutics	Research
2017, Rashid R [415]	Human	Microbiology	Research
2017, Samy RP [416]	Human	Therapeutics	Research
2017, Seneviratne CJ [417]	Human	Therapeutics	Research
2017, Tan JH [418]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2017, Tan JPK [419]	Human	Therapeutics	Research
2017, Tan TY [420]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2017, Tan TY [421]	Human	Diagnostics	Research; Optimisation of Antimicrobial Use
2017, Teo JQM [422]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2017, Venkatesh M [423]	Human	Therapeutics	Research
2017, Wang A [424]	Human	Microbiology	Research

Article ID	Sector(s)	Research Domain	Core Strategy(ies)
2017, Wang J [425]	Human	Therapeutics	Research
2017, Wang K [426]	Human	Microbiology	Research
2017, Yee M [427]	Human	Therapeutics	Research
2017, Yew PYM [428]	Human	Therapeutics	Research
2017, Yi X [429]	Environment	Intervention	Research
2017, Zhong G [430]	Human	Therapeutics	Research
2017, Zhou C [431]	Human	Intervention	Research; Prevention & Control of Infection
2018, Aziz DB [432]	Human	Therapeutics	Research
2018, Baek J-S [433]	Human, Environment	Therapeutics	Research; Optimisation of Antimicrobial Use
2018, Balne PK [434]	Human	Intervention	Research; Prevention & Control of Infection
2018, Budigi Y [435]	Human	Therapeutics	Research
2018, Cai Q [436]	Environment	Intervention	Research
2018, Cai Y [437]	Human	Diagnostics	Surveillance & Risk Assessment; Research; Optimisation of Antimicrobial Use
2018, Chan HL [438]	Human	Diagnostics	Research; Optimisation of Antimicrobial Use
2018, Chan LY [439]	Human	Diagnostics	Research; Optimisation of Antimicrobial Use
2018, Chan MKL [440]	Human	Diagnostics	Surveillance & Risk Assessment; Research
2018, Chew KL [441]	Human	Diagnostics	Surveillance & Risk Assessment; Optimisation of Antimicrobial Use
2018, Chew KL [442]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2018, Chiang RZ-H [443]	Human	Therapeutics	Research
2018, Chiew CJ [444]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment; Prevention & Control of Infection
2018, Chilambi GS [445]	Human	Therapeutics	Research
2018, Chin W [446]	Human	Therapeutics	Research
2018, Chow ALP [12]	Human	Intervention	Prevention & Control of Infection
2018, Chuang L [447]	Human	Therapeutics	Optimisation of Antimicrobial Use
2018, Ding Y [448]	Human	Transmission	Research
2018, Ding Y [449]	Food	Microbiology	Research
2018, Gao J [450]	Human	Diagnostics	Research
2018, Haller L [451]	Environment	Surveillance	Surveillance & Risk Assessment
2018, Hartantyo SHP [452]	Animal	Surveillance	Surveillance & Risk Assessment
2018, Heng ST [453]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2018, Ho ZJM [454]	Human	Transmission	Surveillance & Risk Assessment
2018, Htun HL [455]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2018, Hu B [456]	Human	Therapeutics	Research
2018, Kalimuddin S [457]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2018, Koh J-J [458]	Human	Therapeutics	Research
2018, Koh TH [459]	Human	Microbiology	Research
2018, Lakshmarayanan R [460]	Human	Therapeutics	Research
2018, Lau QY [461]	Human	Therapeutics	Research
2018, Le T-H [462]	Environment	Intervention	Research
2018, Lee GH [463]	Human	Diagnostics	Surveillance & Risk Assessment; Research
2018, Li J [464]	Human	Therapeutics	Research
2018, Lim CLL [465]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2018, Lim T-P [466]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2018, Lim YH [467]	Human	Therapeutics	Research
2018, Lou W [468]	Human	Therapeutics	Research
2018, Mendez AR [469]	Human, Environment	Intervention	Research; Prevention & Control of Infection
2018, Ng C [470]	Environment	Surveillance	Surveillance & Risk Assessment
2018, Ng C [471]	Environment	Surveillance	Surveillance & Risk Assessment
2018, Ng C [472]	Environment	Microbiology	Surveillance & Risk Assessment
2018, Ng DHL [17]	Human, Environment	Transmission	Surveillance & Risk Assessment
2018, Ng SMS [473]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2018, Ng SMS [474]	Human	Therapeutics	Research
2018, Obuobi S [475]	Human	Therapeutics	Research

Article ID	Sector(s)	Research Domain	Core Strategy(ies)
2018, Ong CH [476]	Human	Diagnostics	Surveillance & Risk Assessment; Optimisation of Antimicrobial Use
2018, Parmar A [477]	Human	Therapeutics	Research
2018, Rocamora FM [478]	Human	Microbiology	Research
2018, Selcuk A [479]	Human	Surveillance	Surveillance & Risk Assessment
2018, Sinha S [480]	Human	Therapeutics	Research
2018, Su CT-T [481]	Human	Therapeutics	Research
2018, Su W [482]	Human	Microbiology	Research
2018, Tan BH [483]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2018, Tan D [484]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment; Optimisation of Antimicrobial Use
2018, Tan YE [485]	Human	Surveillance	Surveillance & Risk Assessment
2018, Tang YW [486]	Human	Therapeutics	Research
2018, Teo JWP [487]	Human	Microbiology	Research; Optimisation of Antimicrobial Use
2018, Teo JWP [488]	Human	Diagnostics	Research; Optimisation of Antimicrobial Use
2018, Tong JX [489]	Human	Therapeutics	Research
2018, Xu HV [490]	Human	Intervention	Research; Prevention & Control of Infection
2018, Yang SS [491]	Human	Therapeutics	Surveillance & Risk Assessment; Research; Optimisation of Antimicrobial Use
2018, Yason JA [492]	Human	Therapeutics	Research
2018, Yeo CK [493]	Human	Therapeutics	Research
2018, Yuan W [494]	Food	Intervention	Research; Prevention & Control of Infection
2018, Zhang Z [495]	Human	Diagnostics	Research
2018, Zhu L [496]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2018, Zwe YH [497]	Food	Surveillance	Surveillance & Risk Assessment
2019, Arfan G [498]	Human	Therapeutics	Research
2019, Aung KT [499]	Food, Animal	Surveillance	Surveillance & Risk Assessment
2019, Bae K [500]	Human	Diagnostics	Research; Optimisation of Antimicrobial Use
2019, Bazan EL [501]	Human	Therapeutics	Research
2019, Bharadwaj S [502]	Human	Intervention	Prevention & Control of Infection
2019, Chan JC [503]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2019, Chan YY [504]	Human	Knowledge, Attitudes, Practices	Education
2019, Chen H [505]	Human, Environment	Transmission	Research
2019, Chen WK [506]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2019, Chew KL [507]	Human	Transmission	Surveillance & Risk Assessment
2019, Chew KL [508]	Human	Diagnostics	Research; Optimisation of Antimicrobial Use
2019, Chiu JKH [509]	Human	Diagnostics	Research
2019, Choudhury S [510]	Human	Diagnostics	Research; Optimisation of Antimicrobial Use
2019, Chua AQ [511]	Human	Public Health and Epidemiology	Education; Surveillance & Risk Assessment; Research; Prevention & Control of Infection; Optimisation of Antimicrobial Use
2019, Dupont C [512]	Human	Therapeutics	Research
2019, Ero R [513]	Human	Therapeutics	Research
2019, Fong J [514]	Human	Intervention	Research; Prevention & Control of Infection
2019, Guo S [515]	Food	Surveillance	Surveillance & Risk Assessment
2019, Guo S [516]	Food	Microbiology	Surveillance & Risk Assessment; Research
2019, Guo S [517]	Food	Microbiology	Surveillance & Risk Assessment; Research
2019, Ho HJ [518]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2019, Ho PL [519]	Human	Therapeutics	Research
2019, Htun HL [520]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2019, Keerthisinghe TP [521]	Environment	Intervention	Research
2019, Ko KKK [522]	Human	Surveillance	Surveillance & Risk Assessment
2019, Kyaw BM [21]	Human	Intervention	Education; Research; Prevention & Control of Infection
2019, La M-V [523]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment

Article ID	Sector(s)	Research Domain	Core Strategy(ies)
2019, Lee SY [524]	Human	Diagnostics	Surveillance & Risk Assessment; Research
2019, Legido-Quigley H [525]	Human	Public Health and Epidemiology	Education; Surveillance & Risk Assessment; Research; Prevention & Control of Infection; Optimisation of Antimicrobial Use
2019, Leung CM [526]	Human	Intervention	Research; Prevention & Control of Infection
2019, Li D [527]	Food	Intervention	Research; Prevention & Control of Infection
2019, Li M [528]	Human	Therapeutics	Research
2019, Liu H [529]	Environment	Microbiology	Research
2019, Loke HY [530]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2019, Loo LW [531]	Human	Intervention	Optimisation of Antimicrobial Use
2019, Marimuthu K [532]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2019, Martinez-Vega R [9]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2019, Mayandi V [533]	Human	Therapeutics	Research
2019, Mo Y [534]	Human	Knowledge, Attitudes, Practices	Education
2019, Mo Y [535]	Human	Social and Economic Impacts	Education
2019, Ng C [536]	Environment	Intervention	Surveillance & Risk Assessment; Research
2019, Ngo T-M [537]	Human	Diagnostics	Research; Optimisation of Antimicrobial Use
2019, Obuobi, S [538]	Human	Therapeutics	Research
2019, Octavia S [539]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2019, Pada, SMK [15]	Human	Intervention	Prevention & Control of Infection
2019, Pang X [540]	Food	Intervention	Research
2019, Quek WM [541]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2019, Ryanputra D [542]	Human	Therapeutics	Research
2019, Selcuk A [543]	Human	Surveillance	Surveillance & Risk Assessment
2019, Singh SR [544]	Human	Public Health and Epidemiology	Education; Surveillance & Risk Assessment; Research; Prevention & Control of Infection; Optimisation of Antimicrobial Use
2019, Sorayah R [545]	Human	Therapeutics	Research
2019, Stewardson AJ [546]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2019, Su CT-T [547]	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2019, Tan GSE [548]	Human	Transmission	Surveillance & Risk Assessment; Research
2019, Tan YE [549]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2019, Tay MYF [550]	Food, Human	Surveillance	Surveillance & Risk Assessment
2019, Teo JQM [551]	Human	Therapeutics	Research
2019, Teo JQM [552]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2019, Teo SW [553]	Human	Therapeutics	Research
2019, Tong JX [554]	Human	Therapeutics	Research
2019, Tran NH [555]	Environment	Surveillance	Surveillance & Risk Assessment
2019, Vasoo S [556]	Human	Diagnostics	Research; Optimisation of Antimicrobial Use
2019, Yang C [557]	Human	Therapeutics	Research
2019, Yi X [558]	Environment	Surveillance	Surveillance & Risk Assessment
2019, Yi X [559]	Environment	Surveillance	Surveillance & Risk Assessment
2019, Yuan Y [560]	Human	Therapeutics	Research
2019, Yuan Y [561]	Human	Therapeutics	Research
2019, Zhang EX [562]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2019, Zhang J [563]	Food, Environment	Intervention	Research
2019, Zhang K [564]	Human	Therapeutics	Research
2019, Zhang L [565]	Environment	Intervention	Research
2019, Zheng S-W [566]	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2019, Zhou C [567]	Human	Intervention	Research; Prevention & Control of Infection

Table 2. List of AMR research or review articles published between 2009 and 2019 where Singapore either contributed samples or data, or had researchers involved as part of the research team led by their overseas counterparts.

Articles were assigned to the most relevant research domain, sector(s) and their contribution to the NSAP core strategy(ies).

Article ID	Lead or collaborating country	Sector(s)	Research Domain	Core Strategy(ies)
Contributing samples to multinational studies				
2009, Bouchillon SK [568]	USA	Human	Surveillance	Surveillance & Risk Assessment
2009, Chuang C-H [569]	Taiwan	Human	Surveillance	Surveillance & Risk Assessment
2009, Hawser SP [570]	Switzerland	Human	Surveillance	Surveillance & Risk Assessment
2009, Hurt AC [571]	Australia	Human	Surveillance	Surveillance & Risk Assessment
2009, Ko W-C [572]	Taiwan	Human	Surveillance	Surveillance & Risk Assessment
2009, Lee H-Y [573]	Taiwan South Korea	Human	Surveillance	Surveillance & Risk Assessment
2009, Mendes RE [574]	USA	Human	Surveillance	Surveillance & Risk Assessment
2009, Yau W [575]	Australia	Human	Surveillance	Surveillance & Risk Assessment
2010, Christiansen KJ [576]	UK	Human	Surveillance	Surveillance & Risk Assessment; Research
2010, Farrell DJ [577]	USA	Human	Surveillance	Surveillance & Risk Assessment
2010, Higgins PG [578]	Germany	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2010, Hsueh P-R [579]	Taiwan	Human	Surveillance	Surveillance & Risk Assessment
2011, Chen Y-H [580]	Taiwan	Human	Surveillance	Surveillance & Risk Assessment
2011, Chung DR [581]	South Korea	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2011, Hurt AC [582]	Australia	Human	Surveillance	Surveillance & Risk Assessment
2011, Hurt AC [583]	Australia	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2011, Lee MY [584]	South Korea	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2011, Roberts JA [585]	USA	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2011, Wang H [586]	China Taiwan	Human	Surveillance	Surveillance & Risk Assessment
2012, Bouchillon SK [587]	USA	Human	Surveillance	Surveillance & Risk Assessment
2012, Kiratisin P [588]	Thailand	Human	Surveillance	Surveillance & Risk Assessment
2012, Lin Y-T [589]	Taiwan	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2012, Lu P-L [590]	Taiwan	Human	Surveillance	Surveillance & Risk Assessment
2012, Namdari H [591]	USA	Human	Surveillance	Surveillance & Risk Assessment
2012, Yang Y [592]	China (Hong Kong SAR)	Environment	Surveillance	Surveillance & Risk Assessment
2013, Holden MTG [593]	UK Germany Ireland	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2013, Kim DH [594]	South Korea	Human	Transmission	Surveillance & Risk Assessment
2013, Leang S-K [595]	Australia	Human	Surveillance	Surveillance & Risk Assessment
2013, Mendes RE [596]	USA	Human	Surveillance	Surveillance & Risk Assessment
2013, Sader HS [597]	USA	Human	Surveillance	Surveillance & Risk Assessment
2013, Sheng W-H [598]	Taiwan	Human	Surveillance	Surveillance & Risk Assessment
2014, Ginn AN [599]	Australia	Human	Diagnostics	Surveillance & Risk Assessment; Research
2015, Holt KE [600]	Australia UK	Human, Animal	Public Health and Epidemiology	Surveillance & Risk Assessment
2015, Pfaller MA [601]	USA	Human	Surveillance	Surveillance & Risk Assessment
2016, Jean S-S [602]	Taiwan	Human	Surveillance	Surveillance & Risk Assessment
2016, Tan TY [603]	Singapore	Human	Surveillance	Surveillance & Risk Assessment
2016, Torumkuney D [604]	UK	Human	Surveillance	Surveillance & Risk Assessment
2017, Blackwell GA [605]	Australia	Human	Transmission	Surveillance & Risk Assessment; Research
2017, Blackwell GA [606]	Australia	Human	Microbiology	Surveillance & Risk Assessment; Research
2017, Chang Y-T [607]	Taiwan	Human	Surveillance	Surveillance & Risk Assessment

Article ID	Lead or collaborating country	Sector(s)	Research Domain	Core Strategy(ies)
2017, Cheong HS [608]	South Korea	Human	Surveillance	Surveillance & Risk Assessment
2017, Jean S-S [609]	Taiwan	Human	Surveillance	Surveillance & Risk Assessment
2017, Karlowsky JA [610]	US	Human	Surveillance	Surveillance & Risk Assessment
2017, Ma L [611]	China (Hong Kong SAR)	Environment	Surveillance	Surveillance & Risk Assessment
2018, Harris PNA [612]	Australia	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2018, Harris PNA [613]	Australia	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2018, Khor WC [614]	Malaysia	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2018, Mendis SM [615]	Singapore USA	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2018, Pfaller MA [616]	USA	Human	Surveillance	Surveillance & Risk Assessment; Optimisation of Antimicrobial Use
2018, Versporten A [617]	Belgium	Human	Surveillance	Surveillance & Risk Assessment
2018, Vilaichone RK [618]	Thailand	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2018, Yarbrough ML [619]	USA	Human	Diagnostics	Research; Prevention & Control of Infection
2019, Blot S [620]	Belgium	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2019, Chen SL [621]	Singapore	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2019, George CRR [622]	Australia	Human	Surveillance	Surveillance & Risk Assessment
2019, Hendriksen RS [623]	Denmark	Environment	Surveillance	Surveillance & Risk Assessment
2019, Hsia Y [624]	UK	Human	Surveillance	Surveillance & Risk Assessment; Optimisation of Antimicrobial Use
2019, Hsia Y [625]	UK	Human	Surveillance	Surveillance & Risk Assessment; Optimisation of Antimicrobial Use
2019, Hu YJ [626]	China (Hong Kong SAR)	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2019, Luo Y [627]	China USA	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2019, Ma L [628]	China (Hong Kong SAR)	Environment	Surveillance	Surveillance & Risk Assessment
2019, Papadopoulos A [629]	Greece	Human	Public Health and Epidemiology	Surveillance & Risk Assessment

International Research Collaborations on AMR, 2009-2019

2009, Cervantes S [630]	USA	Human	Diagnostics	Research
2009, Tam VH [631]	USA	Human	Therapeutics	Research
2009, Valvatne H [632]	Norway	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2011, Dhorda M [633]	France	Human	Transmission	Surveillance & Risk Assessment; Research
2011, Leitsch D [634]	Australia	Human	Therapeutics	Research
2011, Massi MN [635]	Indonesia	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2011, Siswantoro H [636]	Australia	Human	Therapeutics	Research
2012, Apisarnthanarak A [637]	Thailand	Human, Environment	Public Health and Epidemiology	Surveillance & Risk Assessment
2012, Brunner R [638]	Switzerland	Human	Therapeutics	Research
2012, Cervantes S [639]	USA	Human	Diagnostics	Research
2012, Dunn LA [640]	Australia	Human	Therapeutics	Research
2012, Guiton PS [641]	USA	Human	Intervention	Research; Prevention & Control of Infection
2012, Ho KKK [642]	Australia	Human	Intervention	Research; Prevention & Control of Infection
2012, Köser CU [643]	UK	Human	Diagnostics	Research; Prevention & Control of Infection
2012, Lee M [644]	South Korea	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2012, Shields RK [645]	USA	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2012, Zhang L [646]	China	Human	Microbiology	Research

Article ID	Lead or collaborating country	Sector(s)	Research Domain	Core Strategy(ies)
2013, Barraud N [647]	Australia	Human	Therapeutics	Research
2013, Bowers DR [648]	Singapore USA	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2013, Chiang W-C [649]	Denmark	Human	Therapeutics	Research
2013, Ciesielczuk H [650]	UK	Human	Diagnostics	Research
2013, Jakobsen TH [651]	Denmark	Human	Microbiology	Research
2013, Kelesidis T [652]	USA	Human	Surveillance	Surveillance & Risk Assessment
2013, Liu Y [653]	Denmark	Human	Therapeutics	Research
2013, Richmond GE [654]	UK	Human	Diagnostics	Research
2013, Wang H [655]	Denmark	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2013, Yepuri NR [656]	Australia	Human	Intervention	Research; Prevention & Control of Infection
2013, Zhang Y [657]	USA	Animal, Environment	Surveillance	Surveillance & Risk Assessment
2013, Zhang Y [658]	USA	Animal, Environment	Surveillance	Surveillance & Risk Assessment
2014, Apisarnthanarak A [659]	Thailand	Human	Intervention	Surveillance & Risk Assessment; Prevention & Control of Infection
2014, Butler J [660]	Australia	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2014, Duan S [661]	USA	Human	Microbiology	Research
2014, Karunakaran R [662]	Malaysia	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2014, Kelesidis T [663]	USA	Human, Animal	Transmission	Surveillance & Risk Assessment; Research
2014, Lai C-C [664]	Taiwan	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2014, Landelle C [665]	Switzerland	Human, Environment	Intervention	Prevention & Control of Infection
2014, Stryjewski ME [666]	Argentina	Human	Therapeutics	Research
2014, Veiga MI [667]	Portugal	Human	Microbiology	Surveillance & Risk Assessment
2015, Aamodt H [668]	Norway	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2015, Baranovich T [669]	USA	Human	Therapeutics	Research
2015, Farrukee R [670]	Australia	Human	Therapeutics	Research
2015, Malmquist NA [671]	France	Human	Therapeutics	Research
2015, Matsunaga S [672]	Japan	Human	Diagnostics	Research
2015, Nguyen D [673]	Australia	Human	Therapeutics	Research
2015, Regmi SM [674]	Thailand	Human	Surveillance	Surveillance & Risk Assessment
2015, Rhee S-Y [675]	USA	Human	Diagnostics	Research; Optimisation of Antimicrobial Use
2015, Soetaert K [676]	Belgium	Human	Therapeutics	Research
2015, Zowawi HM [677]	Australia	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2016, Arribas JR [678]	Spain	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2016, Auburn S [679]	Australia	Human	Diagnostics	Research
2016, Coker OO [680]	Thailand	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2016, Cunningham SA [681]	USA	Human	Diagnostics	Research; Optimisation of Antimicrobial Use
2016, Grigg MJ [682]	Australia	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2016, Hanafi A [683]	Malaysia	Human	Microbiology	Research
2016, Harris RC [684]	UK	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2016, Mac Aogáin M [685]	Ireland	Human	Surveillance	Surveillance & Risk Assessment
2016, Nguyen T-K [686]	Australia	Human	Therapeutics	Research
2016, Nilsson M [687]	Denmark	Human	Microbiology	Research
2016, Phyo AP [688]	Thailand	Human	Therapeutics	Research
2016, Richmond GE [689]	UK	Human	Microbiology	Research
2016, Stenvang M [690]	Denmark	Human	Therapeutics	Research

Article ID	Lead or collaborating country	Sector(s)	Research Domain	Core Strategy(ies)
2017, Atarashi K [691]	Japan	Human	Microbiology	Research
2017, Basilico N [692]	Italy	Human	Therapeutics	Research
2017, Bazaka K [693]	Australia	Human	Therapeutics	Research
2017, Belousoff MJ [694]	Australia	Human	Therapeutics	Research
2017, Bhuyan GS [695]	Bangladesh	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2017, Chang MJ [696]	Republic of Korea (South Korea)	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2017, Chen F [697]	China	Human	Therapeutics	Research
2017, Cunningham SA [698]	USA	Human	Diagnostics	Research; Optimisation of Antimicrobial Use
2017, Gowrisankar G [699]	India	Environment	Microbiology	Research
2017, Gunawan C [700]	Australia	Human	Intervention	Research; Prevention & Control of Infection
2017, Howlin RP [701]	UK	Human	Therapeutics	Research
2017, Hutchison C [702]	China	Human	Social and Economic Impacts	Research
2017, Kathirvel S [703]	India	Human	Knowledge, Attitudes, Practices	Education
2017, Lamothe F [704]	USA	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2017, Landier J [705]	Thailand	Human	Therapeutics	Research
2017, Liao J-H [706]	Taiwan	Human	Microbiology	Research
2017, Thai VC [707]	Vietnam	Human	Therapeutics	Research
2018, Ahmed W [708]	Australia	Environment	Surveillance	Surveillance & Risk Assessment
2018, Antonoplis A [709]	USA	Human	Therapeutics	Research
2018, Beattie RE [710]	USA	Environment	Surveillance	Surveillance & Risk Assessment
2018, Chen C [711]	USA	Human	Therapeutics	Research
2018, Allix-Béguec C [712]	UK	Human	Diagnostics	Research; Optimisation of Antimicrobial Use
2018, Dunn DT [713]	UK	Human	Diagnostics	Research
2018, Fang T [714]	China	Environment	Surveillance	Surveillance & Risk Assessment
2018, Germond A [715]	Japan	Human	Microbiology	Research
2018, Grigg MJ [716]	Australia	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2018, Hoppe A [717]	UK	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2018, Jiang Y [718]	China	Environment	Surveillance	Surveillance & Risk Assessment
2018, Kano R [719]	Japan	Animal	Therapeutics	Research
2018, Malkawi R [720]	UK	Human	Therapeutics	Research
2018, Mather AE [721]	Vietnam	Human	Transmission	Surveillance & Risk Assessment; Research
2018, Merchant S [722]	USA	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2018, Nordström R [723]	Sweden	Human	Therapeutics	Research
2018, Oonsivilai M [724]	Cambodia	Human	Diagnostics	Research; Optimisation of Antimicrobial Use
2018, Ravensdale JT [725]	Australia	Human	Surveillance	Surveillance & Risk Assessment
2018, Stockdale AJ [726]	UK	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2018, Subedi D [727]	Australia	Human	Microbiology	Research
2018, Subedi D [728]	Australia	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2018, Subedi D [729]	Australia	Human	Transmission	Surveillance & Risk Assessment; Research
2018, Tzou PL [730]	USA	Human	Diagnostics	Surveillance & Risk Assessment; Research; Optimisation of Antimicrobial Use
2018, Vente A [731]	Germany	Human	Therapeutics	Research; Optimisation of Antimicrobial Use
2018, Zhou C [732]	China	Human	Intervention	Research; Prevention & Control of Infection
2019, Brunton LA [733]	UK	Environment	Public Health and Epidemiology	Surveillance & Risk Assessment
2019, Capci A [734]	Germany	Human	Therapeutics	Research
2019, Chen Y [735]	China	Environment	Surveillance	Surveillance & Risk Assessment

Article ID	Lead or collaborating country	Sector(s)	Research Domain	Core Strategy(ies)
2019, Chen Y [736]	China	Environment	Surveillance	Surveillance & Risk Assessment
2019, Faksri K [737]	Thailand	Human	Diagnostics	Research
2019, González A [738]	Spain	Human	Therapeutics	Research
2019, González A [739]	Spain	Human	Therapeutics	Research
2019, Jabbar A [740]	Pakistan	Human	Public Health and Epidemiology	Surveillance & Risk Assessment
2019, Juhas M [741]	Switzerland	Human	Therapeutics	Research
2019, Li H [742]	Denmark	Food	Transmission	Surveillance & Risk Assessment; Research
2019, Li H [743]	Denmark	Food	Surveillance	Surveillance & Risk Assessment
2019, Limmathurotsakul D [744]	Thailand	Human, Animal, Food	Knowledge, Attitudes, Practices	Education
2019, Long S [745]	China	Human	Diagnostics	Research
2019, Nilsson M [746]	Denmark	Human	Therapeutics	Research
2019, Nilsson M [747]	Denmark	Human	Therapeutics	Research
2019, Pei M [748]	China	Environment	Intervention	Research
2019, Penesyan A [749]	Australia	Human	Therapeutics	Research
2019, Phelan JE [750]	UK Philippines	Human	Microbiology	Research
2019, Ram M R [751]	Malaysia	Human	Therapeutics	Research
2019, Safi H [752]	USA	Human	Microbiology	Research
2019, Sosibo SC [753]	South Africa	Human	Therapeutics	Research
2019, Subedi D [754]	Australia	Human	Microbiology	Research
2019, Thompson JA [755]	UK	Human	Therapeutics	Research
2019, Yang DL [756]	China	Human	Therapeutics	Research
2019, Zhang N [757]	China	Environment	Surveillance	Surveillance & Risk Assessment

Annex C. AMR Research from Singapore, 2009-2019 ([Section IV\(A\)](#))

Figure 1. AMR Research – Annual Publication Trend 2009-2019.

Annual Publication Trend, 2009-2019

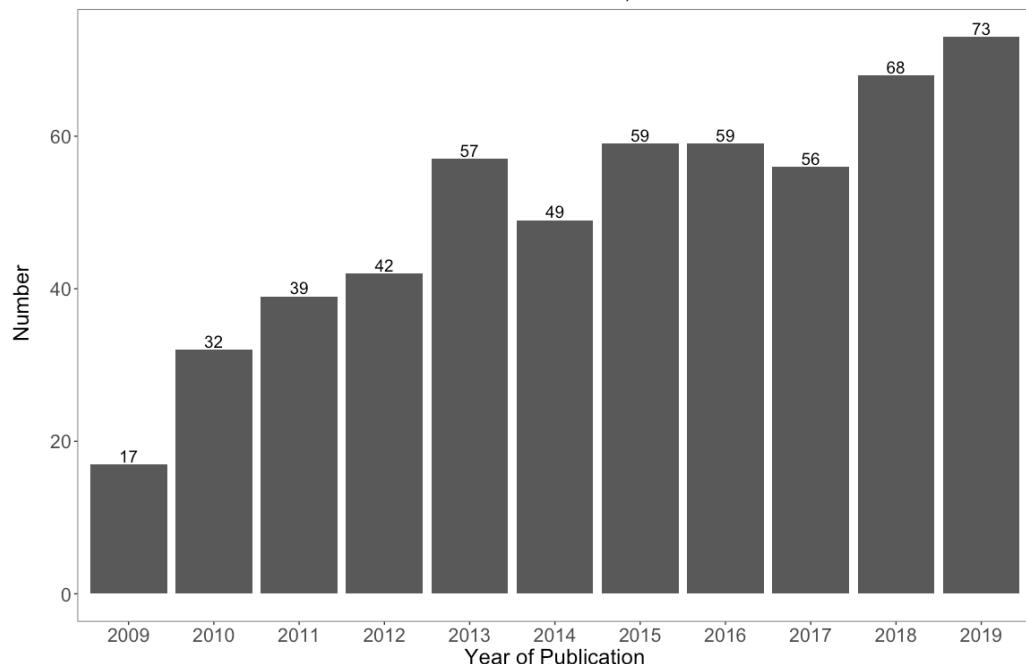
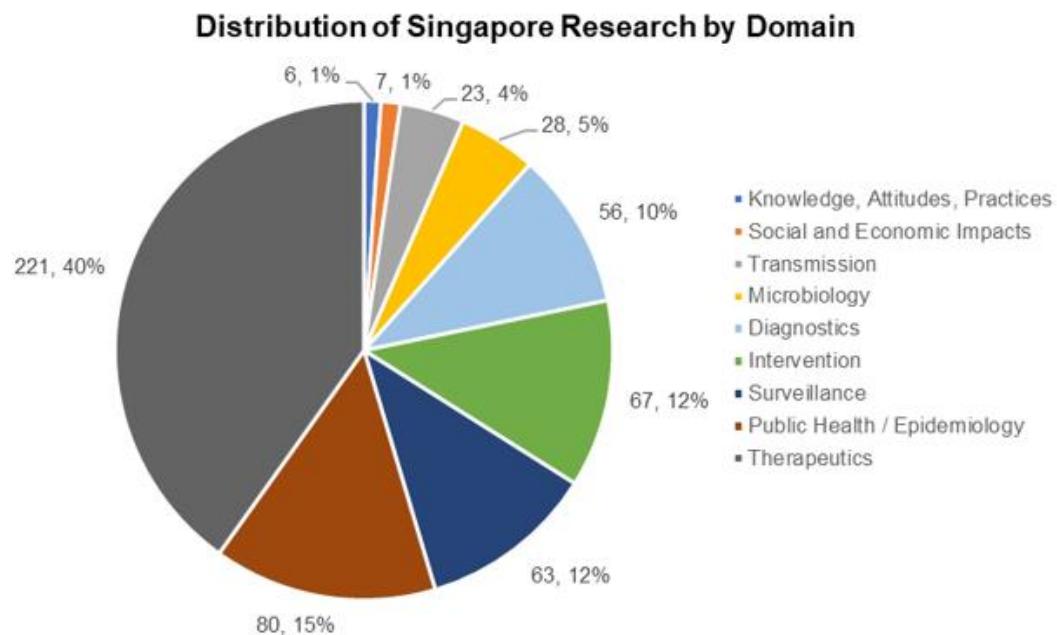


Figure 2. Distribution of AMR research by research domain.

(A) Overall distribution of AMR research by research domain.



(B) Annual trend of publication by research domain, 2009-2019.

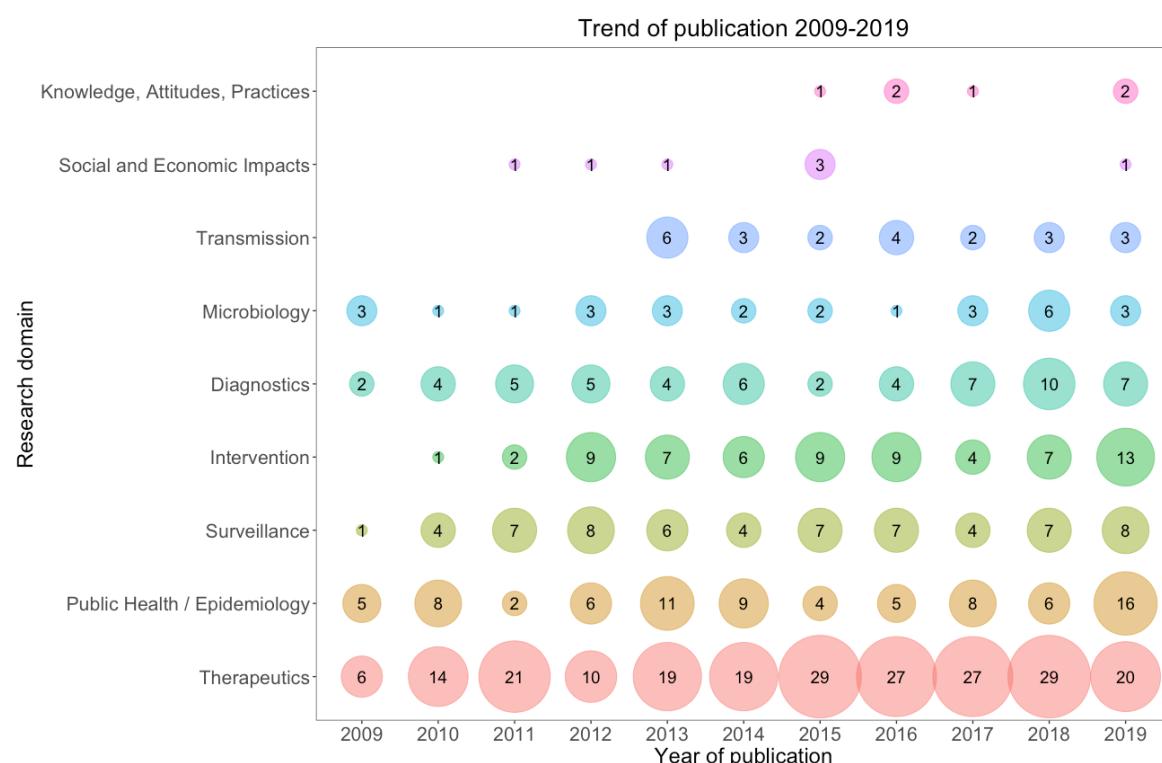
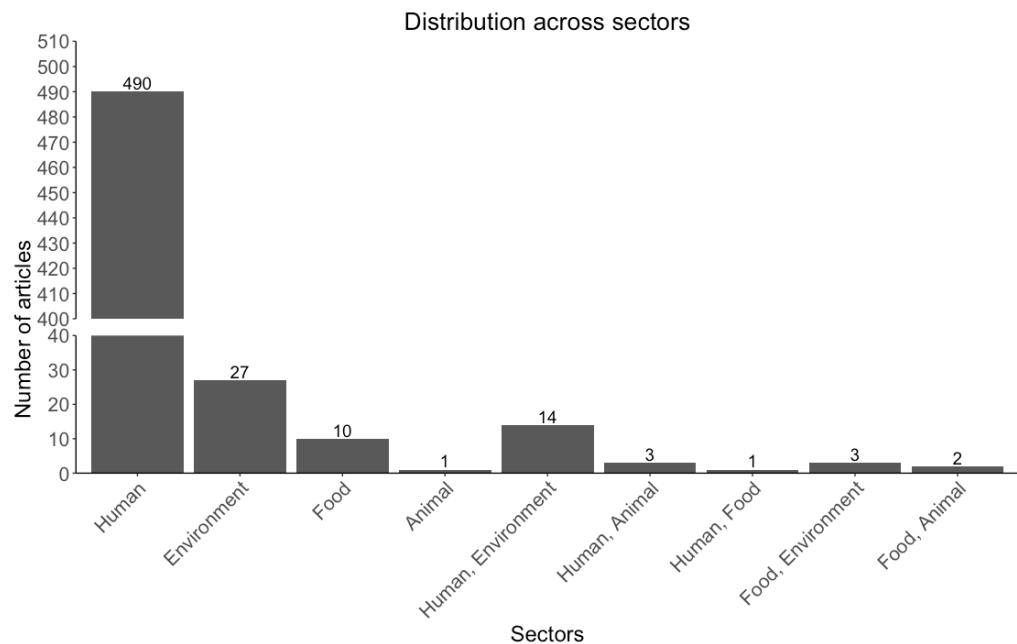


Figure 3. AMR Publication from One Health sectors.

(A) Total publication across One Health sectors.



(B) Annual publication trend from One Health sectors.

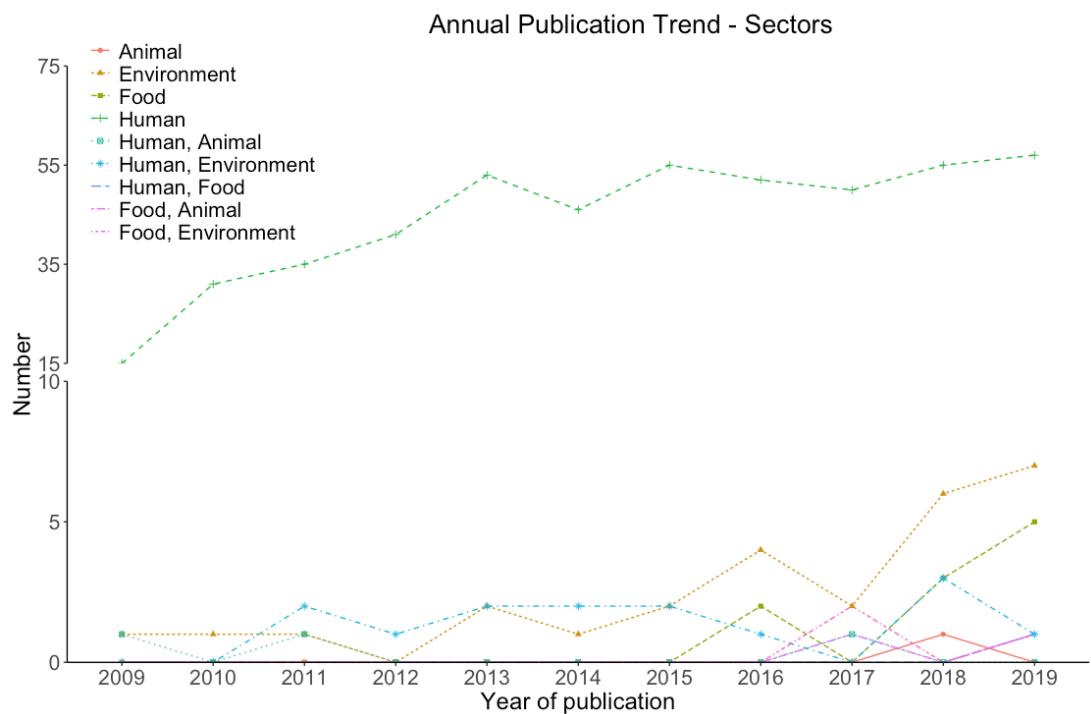


Figure 4. Microorganisms studied between 2009 to 2019.

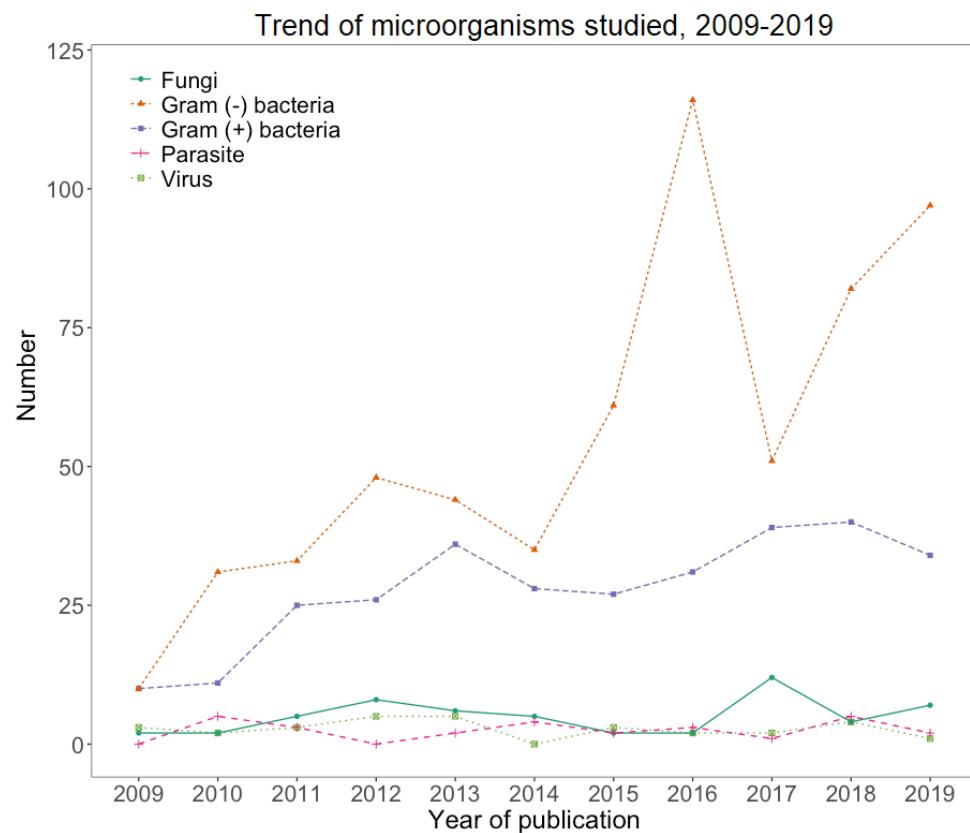


Figure 5. Institutions and their areas of research.

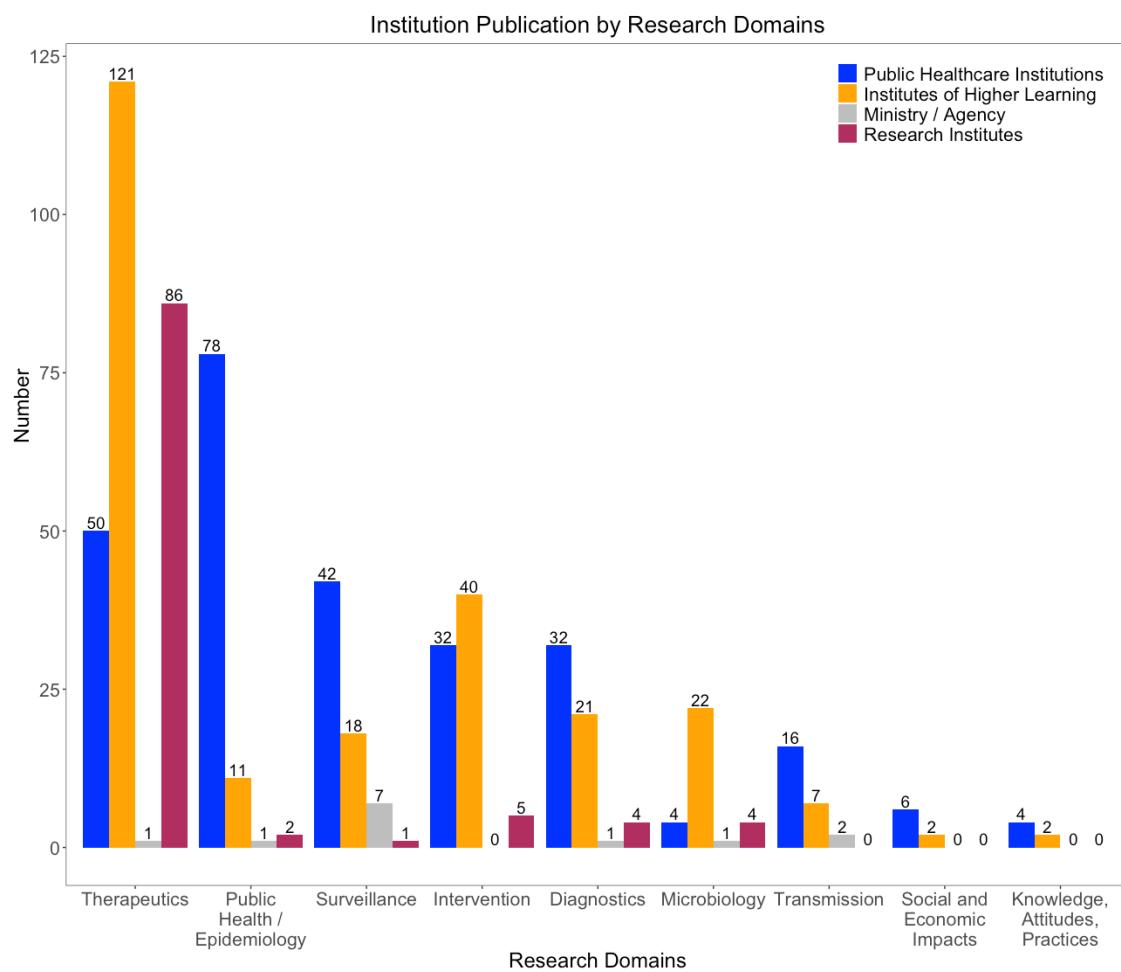


Table 1. Number of research articles in the Therapeutics domain that corresponded to WHO's priority pathogens' list for research and development of novel therapeutics^{vi}.

AMR Pathogen	Number of articles
Priority 1: Critical	
<i>Acinetobacter baumannii</i> , carbapenem-resistant	11
<i>Pseudomonas aeruginosa</i> , carbapenem-resistant	43
Enterobacterales, carbapenem-resistant, ESBL-producing	40
Priority 2: High	
<i>Enterococcus faecium</i> , vancomycin-resistant	9
<i>Staphylococcus aureus</i> , methicillin-resistant, vancomycin-intermediate and resistant	59
<i>Helicobacter pylori</i> , clarithromycin-resistant	2
<i>Campylobacter</i> spp., fluoroquinolone-resistant	0
<i>Salmonellae</i> , fluoroquinolone-resistant	2
<i>Neisseria gonorrhoeae</i> , cephalosporin-resistant, fluoroquinolone-resistant	0
Priority 3: Medium	
<i>Streptococcus pneumoniae</i> , penicillin-non-susceptible	1
<i>Haemophilus influenzae</i> , ampicillin-resistant	0
<i>Shigella</i> spp., fluoroquinolone-resistant	0

^{vi} [WHO publishes list of bacteria for which new antibiotics are urgently needed](#)

Table 2. Volume of AMR research output by institutions, 2009-2019.

Singapore Institutions (by groups)	Number of articles
Institutes of Higher Learning	245
Duke-NUS Medical School	11
Nanyang Technological University	85
National University of Singapore	140
Ngee Ann Polytechnic	2
Republic Polytechnic	2
Singapore Institute of Technology	2
Singapore Polytechnic	1
Singapore University of Technology & Design	2
Ministry / Agency	13
Research institutes	102
Public Healthcare Institutions	260
Healthcare Cluster: National Healthcare Group	75
Khoo Teck Puat Hospital	1
National Skin Centre	2
Tan Tock Seng Hospital / National Centre for Infectious Diseases	72
Healthcare Cluster: National University Health System	83
National University Hospital	80
Ng Teng Fong General Hospital / Jurong Community Hospital	3
Healthcare Cluster: Singapore Health Services (SingHealth)	102
Bedok Polyclinic	1
Changi General Hospital	17
Eastern Health Alliance	1
KK Women's and Children's Hospital	4
National Cancer Centre Singapore	5
Singapore General Hospital	74

Table 3. Sources of funding and the number of research articles funded each, 2009-2019.

Source of funding	Number of articles
Ministry/Agency	344
Healthcare cluster grants	44
IHL funding	52
Public Healthcare Institutions Internal Funding	18
Others	2

Annex D. Summary of AMR research from Singapore in each research domain
(Section IV(A))

1. Knowledge, Attitudes, Practices. Studies in this domain were only published from 2015 and conducted in the human sector. Amongst healthcare professionals, factors associated with antibiotic prescription and acceptance of computer-generated recommendations were studied. In non-healthcare professionals, the overall knowledge of antibiotic use was poor, and factors associated with higher antibiotic use were also identified, such as ethnicity, housing type, lower education and higher social economic status. Such studies in different population groups could provide insights for the development of appropriate or effective education interventions for the respective audiences, to improve their awareness and response toward AMR.
2. Social and Economic Impacts. AMR infections resulted in higher median medical costs to hospitals and patients, arising from an increased length of stay and post-discharge care costs. AMR infections also caused higher mortality rates in patients. There were a few studies that compared the cost-effectiveness of AMR interventions, such as screen and isolate programmes and recommendations by hospitals' antimicrobial stewardship programme (ASP). In terms of the social and psychological impact of AMR, one study found that isolation programmes led to emotional trauma, depression and anxiety in patients
3. Transmission. Another key area of research was to identify sources for the transfer or spread of AMR microorganisms and antibiotic resistance genes (ARGs). Studies identified from this review covered environmental sources, nosocomial, vector-borne or zoonotic transmission as transmission pathways for microorganisms. For MRSA and tuberculosis, transmission often occurred at venues with prolonged close contact, such as gaming centres, prisons or schools. International travel could also contribute to transmission or introduction of new AMR strains. At the molecular level, phage transduction, plasmid transconjugation or acquisition, and the presence of mobile genetic elements were identified as possible contributors to increased transmission. Insights from these studies provided avenues for the development of appropriate intervention strategies to control the spread of AMR.
4. Diagnostics. To enhance detection of AMR, research on diagnostics identified from this review had concentrated on developing more robust tools that would improve the turnaround time, accuracy and reliability of existing methods. Diagnostic tools were also developed to measure the presence of antimicrobials or residues from environmental samples. In general, diagnostic tools could be further segregated into assay-based methods, genetic approaches, imaging or spectrometry. In addition to developing detection technologies, studies that evaluated predictive scoring methods for patient cohorts and sample collection methods to increase sensitivity of AMR detection were also identified.
5. Intervention. Research to mitigate the transmission or development of AMR was primarily done in the human, environment and food sectors. Infection

prevention and control (IPC) strategies – including active surveillance and decontamination, control bundles, use of gloves – and ASP recommendations reduced the nosocomial spread of AMR. Bioengineering methods were also developed to disrupt growth of AMR microorganisms, using biomaterials and photodynamic inactivation that could be applied to catheters or building surfaces. Specific to the environment sector, the removal of AMR pathogens and genes from waste treatment was studied and optimised, while in the food sector, methods to prevent biofilm growth through ultra-violet (UV) treatment or novel compounds were studied. Unfortunately, this landscape review identified only one study each targeted at patients and healthcare professionals to improve antibiotic prescribing.

6. *Public Health and Epidemiology*. Overall, studies in this domain aimed to identify the impact of AMR on the health of a population as a whole, as well as expand surveillance efforts through identifying potential risk factors of AMR in target populations, treatment outcomes and adverse outcomes when treating patients for AMR infections. Research identified in this domain was also diverse, covering the spectrum of developing risk evaluation tools, modelling microbial population response, longitudinal studies to map the changes in drug susceptibilities of microbes, identifying associations between resistance factors and resistance rates, correlating AMR with genetic polymorphisms, and changes in minimum inhibition concentration values. Semi-structured interviews were also used to elicit expert opinion in the social, political, cultural and behavioural spheres.
7. *Surveillance*. Surveillance studies reported the prevalence of AMR microorganisms from the One Health sectors, the abundance of resistance genes in the environment, the genetic relatedness of resistance genes in the human sector, the susceptibility profiles of microorganisms, and antimicrobial usage in healthcare settings. Clinicians also reported emerging cases of resistant microorganisms identified from hospitals, demonstrating that passive surveillance is ongoing and an essential component of healthcare professional duties. Studies that described the presence of AMR microorganisms or antimicrobial content from the environment were also identified.
8. *Therapeutics*. Articles assigned to this domain comprised of laboratory-based basic research, bioinformatics research, clinical research, and reviews. Laboratory-based basic research was diverse, and comprised the identification and development of novel compounds or from drug libraries that had promising activities against AMR microorganisms or biofilms, or repurposing existing approved drugs for treatment of AMR infections. Some studies elucidated the mechanisms of action of therapeutics, or ways the microorganisms developed resistance. Bioinformatics approaches were also used to model mechanisms of resistance *in silico*, or used to model pharmacokinetics-pharmacodynamics responses using retrospective clinical data. In terms of clinical research, retrospective clinical data were analysed to understand appropriateness of antimicrobial therapy, or where possible, identify when culture-guided step-down therapies or the use of effective narrow spectrum or empirical therapeutics based on *in vitro* combination testing could be recommended. However, majority of these studies often employed retrospective analyses and

were limited by sample size and patient demographics. Lastly, reviews on therapeutics summarised the current status of research on therapeutics or treatments against AMR microorganisms.

9. *Microbiology*. Articles assigned to this domain were research that generated further insights on AMR and generally investigated a particular aspect of the microorganism to gain such understanding. These included analysing the dynamics of biofilm formation in a co-culture of two bacteria species, correlating genetics to resistance phenotypes, studying the stages of AMR development in bacteria, and studying novel responses or novel mutations in resistant microorganisms, which could provide avenues for the identification of novel targets for therapeutics development.

Annex E. Involvement in Multinational Research Efforts to Address AMR ([Section IV\(B\)](#))

Figure 1. Multinational AMR research – annual publication trend 2009-2019.

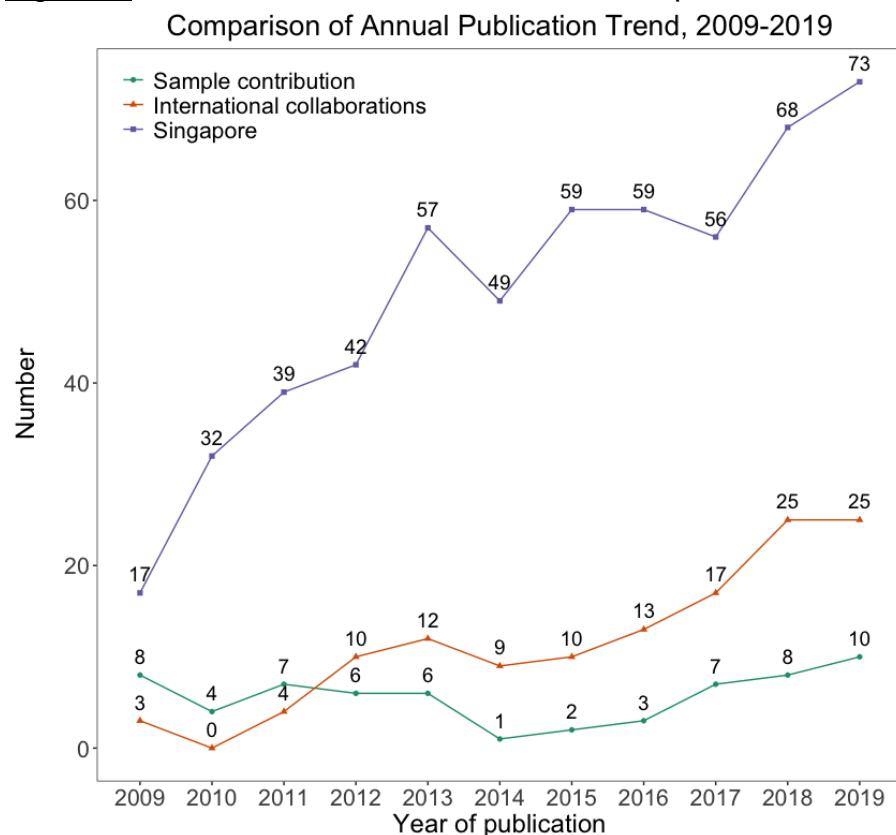


Figure 2. Domain distribution of collaborative AMR research.

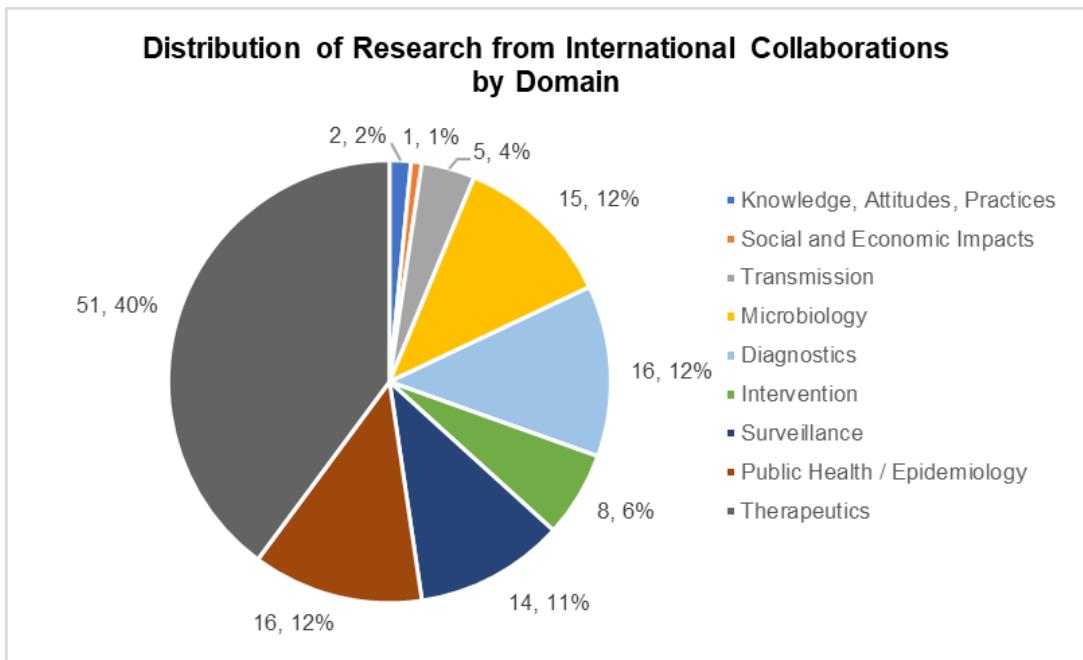
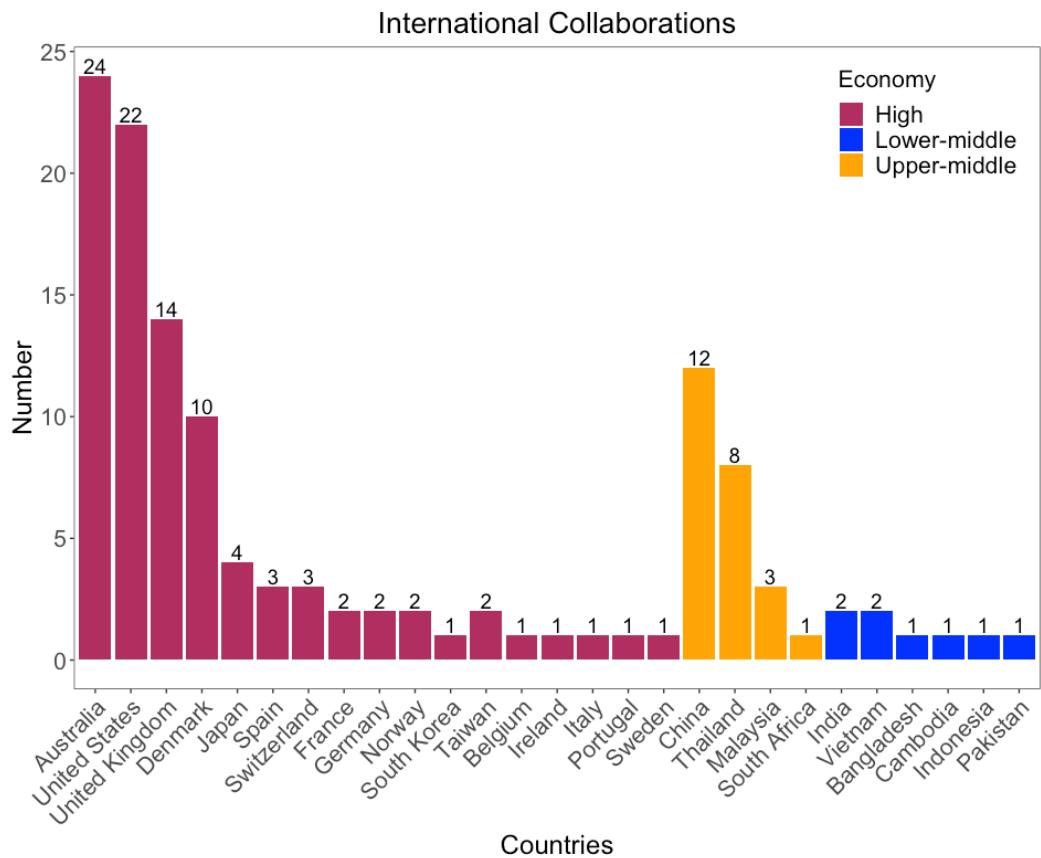


Figure 3. Countries that had AMR research collaborations with Singapore, grouped according to the World Bank country classification by income level^{vii}.



^{vii} [World Bank Country and Lending Groups](#).

Annex F. Summary of the types of AMR research that Singapore contributed samples or data to ([Section IV\(B\)](#))

Domain	Sector	Types of studies	Reference
Surveillance	Human	Most studies reported the resistance trends of microorganisms to antimicrobials. Some of these studies reported surveillance data from regional studies such as the Study for Monitoring Antimicrobial Resistance Trends (SMART), the COMPACT study, the Survey of Antibiotic Resistance (SOAR), the SENTRY antifungal surveillance programme, the Tigecycline Evaluation and Surveillance Trial (TEST), and the Community-Acquired Respiratory Tract Infection Pathogen Surveillance (CARTIPS) study.	[568, 569, 572, 573, 575-577, 579, 580, 586-588, 590, 591, 596, 597, 601-604, 607, 610, 616, 622]
		A few studies reported the trends in antimicrobial consumption in adults and children.	[617, 624, 625]
		Emergence and distribution of ESBL, AmpC beta-lactamases and carbapenemases were reported.	[570, 598, 608, 609]
		The emergence and spread of carbapenemase genes was reported in one study from the SENTRY Surveillance Programme.	[574]
		Studies on the emergence and spread of influenza A(H1N1) towards antivirals. All studies identified focused after the 2009 pandemic when there was increased use of antivirals.	[571, 582, 595]
	Environment	Drinking water samples from households or point of use were collected to investigate the antibiotic resistome.	[611, 628]
		AMR gene abundance from urban sewage or sewage treatment plants were studied to identify variations and diversity.	[592, 623]
Public Health and Epidemiology	Human	Studies in the human sector were diverse and consisted of investigations into the genetic linkages with resistance phenotypes.	[584, 612]
		Elucidating resistance profile with pathogen characteristics.	[615, 621]
		Correlating resistance profile with the type of antimicrobial treatment received.	[581, 629]
		Population characteristics for the acquisition or colonisation by resistant microorganisms.	[589, 614, 620, 626, 627]
		The pattern and spread of resistant <i>Helicobacter pylori</i> , carbapenem-resistance <i>Acinetobacter baumannii</i> and MRSA were studied.	[578, 593, 618]
		One study on resistant influenza H1N1 isolated from patients receiving antiviral therapy.	[583]
	Human-animal	<i>Klebsiella pneumoniae</i> from human and animal sources were analysed to determine its diversity and population structure, as well as their virulence and AMR.	[600]
Diagnostics	Human	One study that identified gene targets that could be used to predict resistance to 3GC and aminoglycosides in <i>Klebsiella pneumoniae</i> and <i>Escherichia coli</i> . Another study evaluated the performance of the Xpert MRSA NxG assay in detecting MRSA directly from nasal swabs.	[599, 619]

Therapeutics	Human	Clinical trial to investigate the activity of piperacillin-tazobactam and meropenem in patients with bacteraemia due to ceftriaxone-resistant <i>E. coli</i> or <i>K. pneumoniae</i> . Pharmacodynamic simulation of carbapenem infusions to determine the dosing regimens required to achieve effective and optimal cumulative fraction of response against resistant bacteria.	[585, 613]
Transmission	Human	Genetic determinants of spread were investigated in <i>Acinetobacter baumannii</i> .	[594, 605]
Microbiology	Human	Presence of an <i>armA</i> gene (a 16S methyltransferase shown to confer resistance to several aminoglycosides) in carbapenem-resistant <i>Acinetobacter baumannii</i> was analysed.	[606]

Annex G. Mapping research relevant to NSAP core strategies and opportunities for future research. ([Section VI\(A\)](#))

	Education	Surveillance & risk assessment	Prevention & control of infection	Optimisation of antimicrobial use (AMU)
Research	Existing / Identified			
Recommendations				
	<p>Knowledge, Attitude, Practices</p> <ul style="list-style-type: none"> Studies provided insights on the public's knowledge on antimicrobial utilisation, and healthcare professionals' antibiotic prescribing behaviour <p>Intervention</p> <ul style="list-style-type: none"> Two studies evaluated existing education for the public and healthcare professionals, both showing mixed or lack of effectiveness 	<p>Surveillance; Public Health/ Epidemiology</p> <ul style="list-style-type: none"> Studies provided information on the prevalence and incidence of AMR/AMU, risk factors related to AMR <p>Transmission</p> <ul style="list-style-type: none"> Studies could contribute information or provide updated information on the spread of resistant pathogens when conducting risk assessments 	<p>Intervention</p> <ul style="list-style-type: none"> Institutional IPC^a programmes evaluated for their effectiveness to prevent and control the spread of infection, sustainability of some of the initiatives Engineering innovations to minimise transmission, such as coatings or modification of surfaces that prevent infection or growth by AMR pathogens and biofilms 	<p>Intervention</p> <ul style="list-style-type: none"> Studies evaluated the impact of ASP[#] recommendations and CDSS recommendations <p>Therapeutics</p> <ul style="list-style-type: none"> Clinical trials to identify effective therapeutics <p>Diagnostics</p> <ul style="list-style-type: none"> Developing improved diagnostic tools for rapid identification of resistance profile for suitable treatment
	<p>Knowledge, Attitudes, Practices</p> <ul style="list-style-type: none"> Encourage studies from the animal, environment and food sectors <p>Intervention</p> <ul style="list-style-type: none"> Encourage studies on development of education interventions and evaluate their effectiveness 	<p>Surveillance</p> <ul style="list-style-type: none"> Encourage cross-sector surveillance studies <p>Public Health/ Epidemiology</p> <ul style="list-style-type: none"> Correlate AMR and AMU <p>Diagnostics</p> <ul style="list-style-type: none"> Development for cross-sector applications to aid in cross-sector surveillance efforts 	<p>Intervention</p> <ul style="list-style-type: none"> Exploratory studies on vaccines to reduce infection by AMR Encourage translation of engineering innovations to practical use in relevant sectors and measure efficacy in real world settings 	<p>Multiple research domains</p> <ul style="list-style-type: none"> Similar research conducted in the human sector could be applied in the animal sector, e.g. antimicrobial prescription, methods to optimise usage

^aIPC – Infection, prevention & control; [#]ASP – antimicrobial stewardship programme

(X) References

1. O'Neill J. Tackling Drug-Resistant Infections Globally: Final Report and Recommendations. London, UK: 2016.
2. Hutchings MI, Truman AW, Wilkinson B. Antibiotics: past, present and future. *Curr Opin Microbiol.* 2019 Oct;51:72-80. PMID: 31733401. doi: 10.1016/j.mib.2019.10.008.
3. World Health Organization. Global action plan on antimicrobial resistance. Geneva: World Health Organization; 2015 2015. ISBN: 9789241509763
9789246509768 (Arabic)
9789245509769 (Chinese).
4. National Strategic Action Plan On Antimicrobial Resistance. Singapore: 2017.
5. Dejene H, Birhanu R, Tarekegn ZS. Knowledge, attitude and practices of residents toward antimicrobial usage and resistance in Gondar, Northwest Ethiopia. *One Health Outlook.* 2022 May 18;4(1):10. PMID: 35581659. doi: 10.1186/s42522-022-00066-x.
6. ul Haq N, Hassali MA, Shafie AA, Saleem F, Farooqui M, Aljadhey H. A cross sectional assessment of knowledge, attitude and practice towards Hepatitis B among healthy population of Quetta, Pakistan. *BMC Public Health.* 2012 2012/08/23;12(1):692. doi: 10.1186/1471-2458-12-692.
7. Boerlin P, Reid-Smith RJ. Antimicrobial resistance: its emergence and transmission. *Anim Health Res Rev.* 2008 Dec;9(2):115-26. PMID: 19102787. doi: 10.1017/S146625230800159X.
8. Burnham CD, Leeds J, Nordmann P, O'Grady J, Patel J. Diagnosing antimicrobial resistance. *Nat Rev Microbiol.* 2017 Oct 12;15(11):697-703. PMID: 29021600. doi: 10.1038/nrmicro.2017.103.
9. Martinez-Vega R, Jaunekaitė E, Thoon KC, Chua HY, Chua AH, Khong WX, et al. Risk factor profiles and clinical outcomes for children and adults with pneumococcal infections in Singapore: A need to expand vaccination policy? *PLoS One.* 2019;14(10):e0220951. PMID: 31618204. doi: 10.1371/journal.pone.0220951.
10. Vasoo S, Singh K, Chow C, Lin RTP, Hsu L-Y, Tambyah PA. Pneumococcal carriage and resistance in children attending day care centers in Singapore in an early era of PCV-7 uptake. *J Infect.* 2010 Jun;60(6):507-9. PMID: 20226209. doi: 10.1016/j.jinf.2010.03.004.
11. Vasoo S, Singh K, Hsu L-Y, Chiew YF, Chow C, Lin RTP, et al. Increasing antibiotic resistance in *Streptococcus pneumoniae* colonizing children attending day-care centres in Singapore. *Respirology.* 2011 Nov;16(8):1241-8. PMID: 21848708. doi: 10.1111/j.1440-1843.2011.02036.x.
12. Chow ALP, Hon PY, Tin G, Zhang W, Poh BF, Ang BSP. Intranasal octenidine and universal antiseptic bathing reduce methicillin-resistant *Staphylococcus aureus* (MRSA) prevalence in extended care facilities. *Epidemiol Infect.* 2018 Dec;146(16):2036-41. PMID: 30176951. doi: 10.1017/s0950268818002522.
13. Ho HJ, Poh BF, Choudhury S, Krishnan PU, Ang BSP, Chow ALP. Alcohol handrubbing and chlorhexidine handwashing are equally effective in removing methicillin-resistant *Staphylococcus aureus* from health care workers' hands: A randomized controlled trial. *Am J Infect Control.* 2015 Nov;43(11):1246-8. PMID: 26190381. doi: 10.1016/j.ajic.2015.06.005.
14. Ling ML, How KB. Impact of a hospital-wide hand hygiene promotion strategy on healthcare-associated infections. *Antimicrob Resist Infect Control.* 2012 Mar 23;1(1):13. PMID: 22958911. doi: 10.1186/2047-2994-1-13.
15. Pada SMK, Chee PL, Rathnam S, Ng KS, Alenton LS, Poh L, et al. Effectiveness of a Ward level target accountability strategy for hand hygiene. *Antimicrob Resist Infect Control.* 2019;8:177. PMID: 31788234. doi: 10.1186/s13756-019-0641-0.
16. Balm MN, Salmon S, Jureen R, Teo C, Mahdi R, Seetoh T, et al. Bad design, bad practices, bad bugs: frustrations in controlling an outbreak of *Elizabethkingia meningoseptica* in intensive care units. *J Hosp Infect.* 2013 Oct;85(2):134-40. PMID: 23958153. doi: 10.1016/j.jhin.2013.05.012.
17. Ng DHL, Marimuthu K, Lee JJ, Khong WX, Ng OT, Zhang W, et al. Environmental colonization and onward clonal transmission of carbapenem-resistant *Acinetobacter baumannii* (CRAB) in a medical intensive care unit: the case for environmental hygiene. *Antimicrob Resist Infect Control.* 2018;7:51. PMID: 29644052. doi: 10.1186/s13756-018-0343-z.
18. Tan TY, Tan JSM, Tay H, Chua GH, Ng LSY, Syahidah N. Multidrug-resistant organisms in a routine ward environment: differential propensity for environmental dissemination and implications for infection control. *J Med Microbiol.* 2013 May;62(Pt 5):766-72. PMID: 23393110. doi: 10.1099/jmm.0.052860-0.
19. Ling ML, How KB. *Pseudomonas aeruginosa* outbreak linked to sink drainage design. *Healthcare infection.* 2013 2013/12/01/;18(4):143-6. doi: <https://doi.org/10.1071/HI13015>.
20. Lee MHM, Pan DST, Huang JH, Chen MI-C, Chong JWC, Goh EH, et al. Results from a Patient-Based Health Education Intervention in Reducing Antibiotic Use for Acute Upper Respiratory Tract

- Infections in the Private Sector Primary Care Setting in Singapore. *Antimicrob Agents Chemother*. 2017 May;61(5). PMID: 28193663. doi: 10.1128/aac.02257-16.
21. Kyaw BM, Tudor Car L, van Galen LS, van Agtmael MA, Costelloe CE, Ajuebor O, et al. Health Professions Digital Education on Antibiotic Management: Systematic Review and Meta-Analysis by the Digital Health Education Collaboration. *J Med Internet Res*. 2019 Sep 12;21(9):e14984. PMID: 31516125. doi: 10.2196/14984.
 22. Huttner B, Saam M, Moja L, Mah K, Sprenger M, Harbarth S, et al. How to improve antibiotic awareness campaigns: findings of a WHO global survey. *BMJ Global Health*. 2019;4(3):e001239. doi: 10.1136/bmigh-2018-001239.
 23. Chow ALP, Win M-K, Wong CS, Leo YS. Universal methicillin-resistant *Staphylococcus aureus* (MRSA) screening: comparison of anatomic screening sites for patients with high and low prevalence of MRSA carriage. *Infect Control Hosp Epidemiol*. 2012 Mar;33(3):315-7. PMID: 22314076. doi: 10.1086/664042.
 24. Win M-K, Soliman TAA, Lee LK, Wong CS, Chow ALP, Ang BSP, et al. Review of a two-year methicillin-resistant *Staphylococcus aureus* screening program and cost-effectiveness analysis in Singapore. *BMC Infect Dis*. 2015 Sep 29;15:391. PMID: 26419926. doi: 10.1186/s12879-015-1131-5.
 25. Tudor Car L, Poon S, Kyaw BM, Cook DA, Ward V, Atun R, et al. Digital Education for Health Professionals: An Evidence Map, Conceptual Framework, and Research Agenda. *J Med Internet Res*. 2022 Mar 17;24(3):e31977. PMID: 35297767. doi: 10.2196/31977.
 26. Uddin TM, Chakraborty AJ, Khusro A, Zidan BMRM, Mitra S, Emran TB, et al. Antibiotic resistance in microbes: History, mechanisms, therapeutic strategies and future prospects. *Journal of Infection and Public Health*. 2021 2021/12/01;14(12):1750-66. doi: <https://doi.org/10.1016/j.jiph.2021.10.020>.
 27. World Health Organization. A systematic approach for undertaking a research priority-setting exercise: guidance for WHO staff. Geneva: World Health Organization; 2020 2020. ISBN: 9789240009622 (electronic version)
9789240009639 (print version).
 28. World Health Organization. Invitation to participate in a survey on research questions for the development of a One Health Priority Research Agenda on Antimicrobial Resistance. World Health Organization; 2021; Available from: <https://www.who.int/news-room/articles-detail/invitation-to-participate-in-a-survey-on-research-questions-for-the-development-of-a-one-health-priority-research-agenda-on-antimicrobial-resistance>.
 29. Humeniuk C, Arlet G, Gautier V, Grimont P, Labia R, Philippon A. Beta-lactamases of Kluyvera ascorbata, probable progenitors of some plasmid-encoded CTX-M types. *Antimicrob Agents Chemother*. 2002 Sep;46(9):3045-9. PMID: 12183268. doi: 10.1128/AAC.46.9.3045-3049.2002.
 30. Kumarasamy KK, Toleman MA, Walsh TR, Bagaria J, Butt F, Balakrishnan R, et al. Emergence of a new antibiotic resistance mechanism in India, Pakistan, and the UK: a molecular, biological, and epidemiological study. *Lancet Infect Dis*. 2010 Sep;10(9):597-602. PMID: 20705517. doi: 10.1016/S1473-3099(10)70143-2.
 31. Rossolini GM, D'Andrea MM, Mugnaioli C. The spread of CTX-M-type extended-spectrum beta-lactamases. *Clin Microbiol Infect*. 2008 Jan;14 Suppl 1:33-41. PMID: 18154526. doi: 10.1111/j.1469-0691.2007.01867.x.
 32. Chan KS, Ling ML, Hsu L-Y, Tan AL. Methicillin-resistant *Staphylococcus aureus* throat colonization among healthcare workers during an outbreak in Singapore General Hospital. *Infect Control Hosp Epidemiol*. 2009 Jan;30(1):95-7. PMID: 19046049. doi: 10.1086/593123.
 33. Chuwa EW, Wong C, Tan Y, Hong G. MRSA breast abscesses in postpartum women. *Asian J Surg*. 2009 Jan;32(1):55-8. PMID: 19321404. doi: 10.1016/s1015-9584(09)60010-5.
 34. Deepak RN, Koh TH, Chan KS. Plasmid-mediated quinolone resistance determinants in urinary isolates of *Escherichia coli* and *Klebsiella pneumoniae* in a large Singapore hospital. *Ann Acad Med Singap*. 2009 Dec;38(12):1070-3. PMID: 20052442.
 35. Donaldson AD, Razak L, Liang LJ, Fisher DA, Tambyah PA. Carbapenems and subsequent multiresistant bloodstream infection: does treatment duration matter? *Int J Antimicrob Agents*. 2009 Sep;34(3):246-51. PMID: 19628129. doi: 10.1016/j.ijantimicag.2009.04.007.
 36. Fan C, Lee PK, Ng WJ, Alvarez-Cohen L, Brodie EL, Andersen GL, et al. Influence of trace erythromycin and erythromycin-H₂O on carbon and nutrients removal and on resistance selection in sequencing batch reactors (SBRs). *Appl Microbiol Biotechnol*. 2009 Nov;85(1):185-95. PMID: 19727707. doi: 10.1007/s00253-009-2201-7.
 37. Fong RKC, Low J, Koh TH, Kurup A. Clinical features and treatment outcomes of vancomycin-intermediate *Staphylococcus aureus* (VISA) and heteroresistant vancomycin-intermediate

- Staphylococcus aureus (hVISA) in a tertiary care institution in Singapore. *Eur J Clin Microbiol Infect Dis.* 2009 Aug;28(8):983-7. PMID: 19387707. doi: 10.1007/s10096-009-0741-5.
38. Ho YM, Sun Y-J, Wong S-Y, Lee AS. Contribution of dfrA and inhA mutations to the detection of isoniazid-resistant *Mycobacterium tuberculosis* isolates. *Antimicrob Agents Chemother.* 2009 Sep;53(9):4010-2. PMID: 19581462. doi: 10.1128/aac.00433-09.
39. Koh TH, Low BS, Leo N, Hsu L-Y, Lin RTP, Krishnan PU, et al. Molecular epidemiology of vancomycin-resistant enterococci in Singapore. *Pathology.* 2009;41(7):676-80. PMID: 19670074. doi: 10.3109/00313020903073047.
40. Lee C, Sun Y-J, Barkham TMS, Leo YS. Primary drug resistance and transmission analysis of HIV-1 in acute and recent drug-naïve seroconverters in Singapore. *HIV Med.* 2009 Jul;10(6):370-7. PMID: 19490177. doi: 10.1111/j.1468-1293.2009.00698.x.
41. Lim T-P, Tan TY, Lee WHL, Sasikala S, Tan TT, Hsu L-Y, et al. In vitro activity of various combinations of antimicrobials against carbapenem-resistant *Acinetobacter* species in Singapore. *J Antibiot (Tokyo).* 2009 Dec;62(12):675-9. PMID: 19876075. doi: 10.1038/ja.2009.99.
42. Liu L, Xu K, Wang H, Tan JP, Fan W, Venkatraman SS, et al. Self-assembled cationic peptide nanoparticles as an efficient antimicrobial agent. *Nat Nanotechnol.* 2009 Jul;4(7):457-63. PMID: 19581900. doi: 10.1038/nnano.2009.153.
43. Luan J, Yuan J, Li X, Jin S, Yu L, Liao M, et al. Multiplex Detection of 60 Hepatitis B Virus Variants by MALDI-TOF Mass Spectrometry. *CLINICAL CHEMISTRY.* 2009 AUG;55(8):1503-9. PMID: WOS:000268557500009. doi: 10.1373/clinchem.2009.124859.
44. Prabakaran M, Prabhu N, Fang H, Qian H, Ho H-T, Jia Q, et al. Combination therapy using chimeric monoclonal antibodies protects mice from lethal H5N1 infection and prevents formation of escape mutants. *PLoS One.* 2009 May 22;4(5):e5672. PMID: 19478856. doi: 10.1371/journal.pone.0005672.
45. Tan TY, Ng LSY, He J, Koh TH, Hsu L-Y. Evaluation of screening methods to detect plasmid-mediated AmpC in *Escherichia coli*, *Klebsiella pneumoniae*, and *Proteus mirabilis*. *Antimicrob Agents Chemother.* 2009 Jan;53(1):146-9. PMID: 18955528. doi: 10.1128/aac.00862-08.
46. Teo JWP, Ng KY, Lin RTP. Detection and genetic characterisation of qnrB in hospital isolates of *Klebsiella pneumoniae* in Singapore. *Int J Antimicrob Agents.* 2009 Feb;33(2):177-80. PMID: 18993034. doi: 10.1016/j.ijantimicag.2008.08.019.
47. Tin S, Sakharkar KR, Lim CSD, Sakharkar MK. Activity of Chitosans in combination with antibiotics in *Pseudomonas aeruginosa*. *Int J Biol Sci.* 2009;5(2):153-60. PMID: 19173037. doi: 10.7150/ijbs.5.153.
48. Wang S-Q, Du Q-S, Huang R-B, Zhang D-W, Chou K-C. Insights from investigating the interaction of oseltamivir (Tamiflu) with neuraminidase of the 2009 H1N1 swine flu virus. *BIOCHEMICAL AND BIOPHYSICAL RESEARCH COMMUNICATIONS.* 2009 AUG 28;386(3):432-6. PMID: WOS:000268262100004. doi: 10.1016/j.bbrc.2009.06.016.
49. Cheow WS, Chang MW, Hadinoto K. Antibacterial efficacy of inhalable antibiotic-encapsulated biodegradable polymeric nanoparticles against *E. coli* biofilm cells. *Journal of Biomedical Nanotechnology.* 2010;6(4):391-403. doi: 10.1166/jbn.2010.1116.
50. Cheow WS, Chang MW, Hadinoto K. Antibacterial Efficacy of Inhalable Levofloxacin-Loaded Polymeric Nanoparticles Against *E. coli* Biofilm Cells: The Effect of Antibiotic Release Profile. *PHARMACEUTICAL RESEARCH.* 2010 AUG;27(8):1597-609. PMID: WOS:000279506100011. doi: 10.1007/s11095-010-0142-6.
51. Ch'ng J-H, Kotturi S, Chong A-L, Lear MJ, Tan KSW. A programmed cell death pathway in the malaria parasite *Plasmodium falciparum* has general features of mammalian apoptosis but is mediated by clan CA cysteine proteases. *Cell Death Dis.* 2010;1(2):e26. PMID: 21364634. doi: 10.1038/cddis.2010.2.
52. Ding C, He J. Effect of antibiotics in the environment on microbial populations. *Appl Microbiol Biotechnol.* 2010 Jul;87(3):925-41. PMID: 20508933. doi: 10.1007/s00253-010-2649-5.
53. Donaldson AD, Tang WY, Tan AL, Barkham TMS. *Neisseria meningitidis* with reduced susceptibility to quinolones in Singapore. *J Antimicrob Chemother.* 2010 Feb;65(2):362-4. PMID: 20007332. doi: 10.1093/jac/dkp437.
54. Gan LSH, Loh JP. Rapid identification of chloroquine and atovaquone drug resistance in *Plasmodium falciparum* using high-resolution melt polymerase chain reaction. *Malar J.* 2010 May 21;9:134. PMID: 20487570. doi: 10.1186/1475-2875-9-134.
55. Ho J, Tambyah PA, Paterson DL. Multiresistant Gram-negative infections: a global perspective. *Curr Opin Infect Dis.* 2010 Dec;23(6):546-53. PMID: 20802331. doi: 10.1097/QCO.0b013e32833f0d3e.

56. Hsu L-Y, Tan TY, Tam VH, Kwa AL-H, Fisher DA, Koh TH. Surveillance and correlation of antibiotic prescription and resistance of Gram-negative bacteria in Singaporean hospitals. *Antimicrob Agents Chemother*. 2010 Mar;54(3):1173-8. PMID: 20065055. doi: 10.1128/aac.01076-09.
57. Hsu L-Y, Leong ML-N, Balm MN, Chan DSG, Huggan P, Tan TY, et al. Six cases of daptomycin-non-susceptible *Staphylococcus aureus* bacteraemia in Singapore. *J Med Microbiol*. 2010 Dec;59(Pt 12):1509-13. PMID: 20705729. doi: 10.1099/jmm.0.022533-0.
58. Husain N, Tkaczuk KL, Tulsidas SR, Kaminska KH, Cubrilo S, Sivaraman J, et al. Structural basis for the methylation of G1405 in 16S rRNA by aminoglycoside resistance methyltransferase Sgm from an antibiotic producer: a diversity of active sites in m7G methyltransferases. *Nucleic Acids Res*. 2010 Jul;38(12):4120-32. PMID: 20194115. doi: 10.1093/nar/gkq122.
59. Inoue M, Barkham TMS, Leo YS, Chan K-P, Chow ALP, Wong CW, et al. Emergence of oseltamivir-resistant pandemic (H1N1) 2009 virus within 48 hours. *Emerg Infect Dis*. 2010 Oct;16(10):1633-6. PMID: 20875299. doi: 10.3201/eid1610.100688.
60. Jayaraman P, Sakharkar MK, Lim CSD, Tang TH, Sakharkar KR. Activity and interactions of antibiotic and phytochemical combinations against *Pseudomonas aeruginosa* in vitro. *Int J Biol Sci*. 2010 Sep 21;6(6):556-68. PMID: 20941374. doi: 10.7150/ijbs.6.556.
61. Koh TH, Khoo CT, Tan TT, Mohamed Arshad MAB, Ang LP, Lau LJ, et al. Multilocus sequence types of carbapenem-resistant *Pseudomonas aeruginosa* in Singapore carrying metallo-beta-lactamase genes, including the novel bla(IMP-26) gene. *J Clin Microbiol*. 2010 Jul;48(7):2563-4. PMID: 20463166. doi: 10.1128/jcm.01905-09.
62. Koh TH, Khoo CT, Wijaya L, Leong HN, Lo YL, Lim LC, et al. Global spread of New Delhi metallo-β-lactamase 1. *Lancet Infect Dis*. 2010 Dec;10(12):828. PMID: 21109168. doi: 10.1016/s1473-3099(10)70274-7.
63. Kurup A, Chlebicka N, Tan KY, Chen EX, Oon LL-E, Tan AL, et al. Active surveillance testing and decontamination strategies in intensive care units to reduce methicillin-resistant *Staphylococcus aureus* infections. *Am J Infect Control*. 2010 Jun;38(5):361-7. PMID: 20189267. doi: 10.1016/j.ajic.2009.09.018.
64. Lim LG, Aung MO, Seet BL, Tan C, Dan YY, Lee YM, et al. Alanine aminotransferase is an inadequate surrogate marker for detecting lamivudine resistance. *World J Gastroenterol*. 2010 Oct 7;16(37):4691-6. PMID: 20872970. doi: 10.3748/wjg.v16.i37.4691.
65. Lim PL, Mok YJ, Lye DCB, Leo YS. Imported chloroquine-resistant *Plasmodium vivax* in Singapore: case report and literature review. *J Travel Med*. 2010 Jan-Feb;17(1):69-71. PMID: 20074103. doi: 10.1111/j.1708-8305.2009.00376.x.
66. Liu R, Teo W, Tan S, Feng H, Padmanabhan P, Xing B. Metallic nanoparticles bioassay for *Enterobacter cloacae* P99 beta-lactamase activity and inhibitor screening. *ANALYST*. 2010;135(5):1031-6. PMID: WOS:000277037500026. doi: 10.1039/b926909f.
67. Ng ES-T, Liew YX, Koh LP, Hsu L-Y. Fluoroquinolone prophylaxis against febrile neutropenia in areas with high fluoroquinolone resistance--an Asian perspective. *J Formos Med Assoc*. 2010 Sep;109(9):624-31. PMID: 20863989. doi: 10.1016/s0929-6646(10)60102-7.
68. Nzila A, Rottmann M, Chitnumsub P, Kiara SM, Kamchonwongpaisan S, Maneeruttanarungroj C, et al. Preclinical Evaluation of the Antifolate QN254, 5-Chloro-N' 6'-(2,5-Dimethoxy-Benzyl)-Quinazoline-2,4,6-Triamine, as an Antimalarial Drug Candidate. *ANTIMICROBIAL AGENTS AND CHEMOTHERAPY*. 2010 JUN;54(6):2603-10. PMID: WOS:000277756000041. doi: 10.1128/AAC.01526-09.
69. Ong DC, Yam W-C, Siu GK, Lee AS. Rapid detection of rifampicin- and isoniazid-resistant *Mycobacterium tuberculosis* by high-resolution melting analysis. *J Clin Microbiol*. 2010 Apr;48(4):1047-54. PMID: 20164280. doi: 10.1128/jcm.02036-09.
70. Rottmann M, McNamara C, Yeung BK, Lee MC, Zou B, Russell B, et al. Spiroindolones, a potent compound class for the treatment of malaria. *Science*. 2010 Sep 3;329(5996):1175-80. PMID: 20813948. doi: 10.1126/science.1193225.
71. Samy RP, Gopalakrishnakone P, Bow H, Puspharaj PN, Chow VT. Identification and characterization of a phospholipase A2 from the venom of the Saw-scaled viper: Novel bactericidal and membrane damaging activities. *Biochimie*. 2010 Dec;92(12):1854-66. PMID: 20723574. doi: 10.1016/j.biochi.2010.07.012.
72. Soe WM, Lin RTP, Lim CSD, Sakharkar KR, Sakharkar MK. In vitro drug interactions of gallates with antibiotics in *Staphylococcus Aureus*. *Front Biosci (Elite Ed)*. 2010 Jan 1;2(2):668-72. PMID: 20036910. doi: 10.2741/e126.
73. Soe WM, Giridharan M, Lin RTP, Sakharkar KM, Sakharkar RK. Effect of Combinations of Antibiotics and Gallates on Biofilm Formation in *Staphylococcus aureus*. *Letters in Drug Design & Discovery*. 2010;7(3):160-4. doi: <http://dx.doi.org/10.2174/157018010790596687>.

74. Sun Y-J, Luo J-T, Wong S-Y, Lee AS. Analysis of rpsL and rrs mutations in Beijing and non-Beijing streptomycin-resistant *Mycobacterium tuberculosis* isolates from Singapore. *Clin Microbiol Infect.* 2010 Mar;16(3):287-9. PMID: 19519851. doi: 10.1111/j.1469-0691.2009.02800.x.
75. Tan TY, Ng LSY, He J, Hsu L-Y. CTX-M and ampC beta-lactamases contributing to increased prevalence of ceftriaxone-resistant *Escherichia coli* in Changi General Hospital, Singapore. *Diagn Microbiol Infect Dis.* 2010 Feb;66(2):210-3. PMID: 19800752. doi: 10.1016/j.diagmicrobio.2009.08.019.
76. Tan TY, Tan AL, Tee NWS, Ng LSY, Chee CW. The increased role of non-albicans species in candidaemia: results from a 3-year surveillance study. *Mycoses.* 2010 Nov;53(6):515-21. PMID: 19619263. doi: 10.1111/j.1439-0507.2009.01746.x.
77. Wijaya L, Hsu L-Y. Community-Associated Methicillin-Resistant *Staphylococcus aureus* Skin and Soft Tissue Infections. *Proceedings of Singapore Healthcare.* 2010 09/01;19:212-9. doi: 10.1177/201010581001900307.
78. Wong L, Liu G. Protein interactome analysis for countering pathogen drug resistance. *Journal of Computer Science and Technology.* 2010;25(1):124-30. doi: 10.1007/s11390-010-9310-8.
79. Zhou C, Qi X, Li P, Chen WN, Mouad L, Chang MW, et al. High potency and broad-spectrum antimicrobial peptides synthesized via ring-opening polymerization of alpha-aminoacid-N-carboxyanhydrides. *Biomacromolecules.* 2010 Jan 11;11(1):60-7. PMID: 19957992. doi: 10.1021/bm900896h.
80. Bahadin J, Teo S, Mathew S. Aetiology of community-acquired urinary tract infection and antimicrobial susceptibility patterns of uropathogens isolated. *Singapore Med J.* 2011 Jun;52(6):415-20. PMID: 21731993.
81. Chan HE, Poon LM, Chan S, Teo JWP. The perils of medical tourism: NDM-1-positive *Escherichia coli* causing febrile neutropenia in a medical tourist. *Singapore Med J.* 2011 Apr;52(4):299-302. PMID: 21552793.
82. Chantratita N, Rholl DA, Sim B, Wuthiekanun V, Limmathurotsakul D, Amornchai P, et al. Antimicrobial resistance to ceftazidime involving loss of penicillin-binding protein 3 in *Burkholderia pseudomallei*. *PROCEEDINGS OF THE NATIONAL ACADEMY OF SCIENCES OF THE UNITED STATES OF AMERICA.* 2011 OCT 11;108(41):17165-70. PMID: WOS:000295973800058. doi: 10.1073/pnas.1111020108.
83. Cheow WS, Chang MW, Hadinoto K. The roles of lipid in anti-biofilm efficacy of lipid-polymer hybrid nanoparticles encapsulating antibiotics. *Colloids and Surfaces A: Physicochemical and Engineering Aspects.* 2011;389(1-3):158-65. doi: 10.1016/j.colsurfa.2011.08.035.
84. Ch'ng J-H, Liew K, Goh A-P, Sidhartha E, Tan KSW. Drug-induced permeabilization of parasite's digestive vacuole is a key trigger of programmed cell death in *Plasmodium falciparum*. *Cell Death Dis.* 2011 Oct 13;2(10):e216. PMID: 21993392. doi: 10.1038/cddis.2011.97.
85. Fan C, He J. Proliferation of antibiotic resistance genes in microbial consortia of sequencing batch reactors (SBRs) upon exposure to trace erythromycin or erythromycin-H₂O. *Water Res.* 2011 May;45(10):3098-106. PMID: 21482429. doi: 10.1016/j.watres.2011.03.025.
86. Hsu L-Y, Tan TY, Koh TH, Kwa AL-H, Krishnan PU, Tee NWS, et al. Decline in *Clostridium difficile*-associated disease rates in Singapore public hospitals, 2006 to 2008. *BMC Res Notes.* 2011 Mar 23;4:77. PMID: 21429188. doi: 10.1186/1756-0500-4-77.
87. Hussain N, Obranic S, Koscienski L, Seetharaman J, Babic F, Sivaraman J, et al. Structural basis for the methylation of A1408 in 16S rRNA by a panaminoglycoside resistance methyltransferase NpmA from a clinical isolate and analysis of the NpmA interactions with the 30S ribosomal subunit. *Nucleic Acids Res.* 2011 Mar;39(5):1903-18. PMID: 21062819. doi: 10.1093/nar/gkq1033.
88. Kyaw BM, Arora S, Win KN, Lim CSD. Prevention of emergence of fusidic acid and rifampicin resistance in *Staphylococcus aureus* using phytochemicals. *African Journal of Microbiology Research.* 2011 10/16;5:3684-92. doi: 10.5897/AJMR11.471.
89. Kyaw BM, Arora S, Lim CSD. Anti-pseudomonal and anti-biofilm activity of tannic acid in combination with antibiotics. *International Journal of Integrative Biology.* 2011;11(3):110-6.
90. Lee CY, Kam Y-W, Fric J, Malleret B, Koh EG, Prakash C, et al. Chikungunya virus neutralization antigens and direct cell-to-cell transmission are revealed by human antibody-escape mutants. *PLoS Pathog.* 2011 Dec;7(12):e1002390. PMID: 22144891. doi: 10.1371/journal.ppat.1002390.
91. Lee HK, Lee CK, Loh TP, Tang JW-T, Tambyah PA, Koay ES-C. High-resolution melting approach to efficient identification and quantification of H275Y mutant influenza H1N1/2009 virus in mixed-virus-population samples. *J Clin Microbiol.* 2011 Oct;49(10):3555-9. PMID: 21865430. doi: 10.1128/jcm.01087-11.
92. Leung GY, Li H, Toh Q-Y, Ng AM-Y, Sum RJ, Chen DY-K, et al. Total Synthesis and Biological Evaluation of the Fab-Inhibitory Antibiotic Platencin and Analogues Thereof. *EUROPEAN JOURNAL*

- OF ORGANIC CHEMISTRY. 2011 JAN;2011(1):183-96. PMID: WOS:000286774900021. doi: 10.1002/ejoc.201001281.
93. Liew YX, Lee WHL, Kwa AL-H, Lye DCB, Yeo C-L, Hsu L-Y. Inappropriate carbapenem use in Singapore public hospitals: opportunities for antimicrobial stewardship. *Int J Antimicrob Agents*. 2011 Jan;37(1):87-8. PMID: 21074375. doi: 10.1016/j.ijantimicag.2010.09.008.
94. Liew YX, Chlebicki MP, Lee WHL, Hsu L-Y, Kwa AL-H. Use of procalcitonin (PCT) to guide discontinuation of antibiotic use in an unspecified sepsis is an antimicrobial stewardship program (ASP). *Eur J Clin Microbiol Infect Dis*. 2011 Jul;30(7):853-5. PMID: 21279532. doi: 10.1007/s10096-011-1165-6.
95. Liew YX, Krishnan PU, Yeo C-L, Tan TY, Lee S-Y, Lim W-P, et al. Surveillance of broad-spectrum antibiotic prescription in Singaporean hospitals: a 5-year longitudinal study. *PLoS One*. 2011;6(12):e28751. PMID: 22174889. doi: 10.1371/journal.pone.0028751.
96. Lim PL, Ling ML, Lee H, Koh TH, Tan AL, Kuijper E, et al. Isolation of the first three cases of Clostridium difficile polymerase chain reaction ribotype 027 in Singapore. *Singapore Med J*. 2011 May;52(5):361-4. PMID: 21633771.
97. Lim T-P, Tan TY, Lee WHL, Sasikala S, Teo JQM, Hsu L-Y, et al. Effective antibiotics in combination against extreme drug-resistant *Pseudomonas aeruginosa* with decreased susceptibility to polymyxin B. *PLoS One*. 2011;6(12):e28177. PMID: 22162759. doi: 10.1371/journal.pone.0028177.
98. Lim T-P, Tan TY, Lee WHL, Sasikala S, Tan TT, Hsu L-Y, et al. In-vitro activity of polymyxin B, rifampicin, tigecycline alone and in combination against carbapenem-resistant *Acinetobacter baumannii* in Singapore. *PLoS One*. 2011 Apr 21;6(4):e18485. PMID: 21533030. doi: 10.1371/journal.pone.0018485.
99. Lingegowda PB, Tan CK, Tan AL, Tan BH. Selection of resistant fungi in liver transplant recipients during use of newer antifungal agents -- a report of two cases. *Ann Acad Med Singap*. 2011 Jun;40(6):287-90. PMID: 21779617.
100. Mirza H, Teo JDW, Upcroft JA, Tan KSW. A rapid, high-throughput viability assay for *Blastocystis* spp. reveals metronidazole resistance and extensive subtype-dependent variations in drug susceptibilities. *Antimicrob Agents Chemother*. 2011 Feb;55(2):637-48. PMID: 21098237. doi: 10.1128/aac.00900-10.
101. Mok S, Imwong M, MacKinnon MJ, Sim J, Ramadoss R, Yi P, et al. Artemisinin resistance in *Plasmodium falciparum* is associated with an altered temporal pattern of transcription. *BMC Genomics*. 2011 Aug 3;12:391. PMID: 21810278. doi: 10.1186/1471-2164-12-391.
102. My NH, Van DU, Hirao H, Morokuma K. Computational studies of bacterial resistance to β -lactam antibiotics: Mechanism of covalent inhibition of the penicillin-binding protein 2a (PBP2a). *Journal of Chemical Information and Modeling*. 2011;51(12):3226-34. doi: 10.1021/ci2004175.
103. Nederberg F, Zhang Y, Tan JP, Xu K, Wang H, Yang C, et al. Biodegradable nanostructures with selective lysis of microbial membranes. *Nature Chemistry*. 2011;3(5):409-14. doi: 10.1038/nchem.1012.
104. Ng LSY, Teh W, Ng S, Eng LC, Tan TY. Bacterial contamination of hands and the environment in a microbiology laboratory. *J Hosp Infect*. 2011 Jul;78(3):231-3. PMID: 21481970. doi: 10.1016/j.jhin.2011.01.025.
105. Ong DC, Koh TH, Syahidah N, Krishnan PU, Tan TY. Rapid detection of the blaNDM-1 gene by real-time PCR. *J Antimicrob Chemother*. 2011 Jul;66(7):1647-9. PMID: 21565805. doi: 10.1093/jac/dkr184.
106. Pada SMK, Ding Y, Ling ML, Earnest A, Lee T-E, Yong H-C, et al. Economic and clinical impact of nosocomial meticillin-resistant *Staphylococcus aureus* infections in Singapore: a matched case-control study. *J Hosp Infect*. 2011 May;78(1):36-40. PMID: 21269733. doi: 10.1016/j.jhin.2010.10.016.
107. Phua CK, Chee CBE, Chua AP-G, Gan SH, Ahmed AD, Wang Y-T. Managing a case of extensively drug-resistant (XDR) pulmonary tuberculosis in Singapore. *Ann Acad Med Singap*. 2011 Mar;40(3):132-5. PMID: 21603731.
108. Saeidi N, Wong CK, Lo T-M, Nguyen HX, Ling H, Leong SSJ, et al. Engineering microbes to sense and eradicate *Pseudomonas aeruginosa*, a human pathogen. *Molecular Systems Biology*. 2011;7. doi: 10.1038/msb.2011.55.
109. Samy RP, Stiles BG, Gopalakrishnakone P, Chow VT. Antimicrobial proteins from snake venoms: direct bacterial damage and activation of innate immunity against *Staphylococcus aureus* skin infection. *Curr Med Chem*. 2011;18(33):5104-13. PMID: 22050758. doi: 10.2174/092986711797636108.
110. Sim SH, Liu Y, Tan J, Thong TW, Wang D, Ooi EE, et al. Antimicrobial activity of cathelicidin peptides against *Burkholderia pseudomallei*, the causative agent of melioidosis. *International Journal of Antimicrobial Agents*. 2011;38(3):270-1. doi: 10.1016/j.ijantimicag.2011.06.002.

111. Soe WM, Myint NL, Lim CSD, Sakharkar MK, Sakharkar KR. Ethyl Gallate as a Combination Drug Can Overcome Resistance in MRSA. *LETTERS IN DRUG DESIGN & DISCOVERY*. 2011 JAN;8(1):65-8. PMID: WOS:000285781700008.
112. Suhaila M, Tang JW-T, Lee HK, Lin C, Tambyah PA, Yap HK, et al. Mixtures of oseltamivir-sensitive and -resistant pandemic influenza A/H1N1/2009 viruses in immunocompromised hospitalized children. *Pediatr Infect Dis J*. 2011 Jul;30(7):625-7. PMID: 21206398. doi: 10.1097/INF.0b013e31820929ab.
113. Tan TY, Lim T-P, Lee WHL, Sasikala S, Hsu L-Y, Kwa AL-H. In vitro antibiotic synergy in extensively drug-resistant *Acinetobacter baumannii*: the effect of testing by time-kill, checkerboard, and Etest methods. *Antimicrob Agents Chemother*. 2011 Jan;55(1):436-8. PMID: 20956606. doi: 10.1128/aac.00850-10.
114. Teo BW, Low SJ-X, Ding Y, Koh TH, Hsu L-Y. High prevalence of mupirocin-resistant staphylococci in a dialysis unit where mupirocin and chlorhexidine are routinely used for prevention of catheter-related infections. *J Med Microbiol*. 2011 Jun;60(Pt 6):865-7. PMID: 21317195. doi: 10.1099/jmm.0.024539-0.
115. Teo BW, Sau PY, Xu H, Ma V, Vathsala A, Lee EJ. Microbiology of tunnelled catheter-related infections in a multi-ethnic South-east Asian patient population. *Nephron Clin Pract*. 2011;118(2):c86-92. PMID: 21150216. doi: 10.1159/000319866.
116. Teo JWP, Krishnan PU, Jureen R, Lin RTP. Detection of an unusual van genotype in a vancomycin-resistant *Enterococcus faecium* hospital isolate. *J Clin Microbiol*. 2011 Dec;49(12):4297-8. PMID: 21998432. doi: 10.1128/jcm.05524-11.
117. Xing B, Jiang T, Bi W, Yang Y, Li L, Ma M, et al. Multifunctional divalent vancomycin: the fluorescent imaging and photodynamic antimicrobial properties for drug resistant bacteria. *Chem Commun (Camb)*. 2011 Feb 7;47(5):1601-3. PMID: 21109901. doi: 10.1039/c0cc04434b.
118. Bai Y, Liu S, Lakshminarayanan R, Sarawathi P, Tang C, Ho D, et al. Progressive structuring of a branched antimicrobial peptide on the path to the inner membrane target. *J Biol Chem*. 2012 Aug 3;287(32):26606-17. PMID: 22700968. doi: 10.1074/jbc.M112.363259.
119. Balm MN, Ngan G, Jureen R, Lin RTP, Teo JWP. Molecular characterization of newly emerged blaKPC-2-producing *Klebsiella pneumoniae* in Singapore. *J Clin Microbiol*. 2012 Feb;50(2):475-6. PMID: 22116160. doi: 10.1128/jcm.05914-11.
120. Cai Y, Chan JP, Fisher DA, Koh TH, Krishnan PU, Kwa AL-H, et al. Vancomycin-resistant Enterococci in Singaporean hospitals: 5-year results of a multi-centre surveillance programme. *Ann Acad Med Singap*. 2012 Feb;41(2):77-81. PMID: 22498854.
121. Chen C, Gunawan P, Lou XWD, Xu R. Silver Nanoparticles Deposited Layered Double Hydroxide Nanoporous Coatings with Excellent Antimicrobial Activities. *ADVANCED FUNCTIONAL MATERIALS*. 2012 FEB 22;22(4):780-7. PMID: WOS:000300447500013. doi: 10.1002/adfm.201102333.
122. Chen T, Wang R, Xu L, Kang E-T, Neoh KG. Carboxymethyl Chitosan-Functionalized Magnetic Nanoparticles for Disruption of Biofilms of *Staphylococcus aureus* and *Escherichia coli*. *INDUSTRIAL & ENGINEERING CHEMISTRY RESEARCH*. 2012 OCT 10;51(40):13164-72. PMID: WOS:000309624500020. doi: 10.1021/ie301522w.
123. Chen Y-T, Lin A-C, Siu LK, Koh TH. Sequence of closely related plasmids encoding bla(NDM-1) in two unrelated *Klebsiella pneumoniae* isolates in Singapore. *PLoS One*. 2012;7(11):e48737. PMID: 23139815. doi: 10.1371/journal.pone.0048737.
124. Cheong CSJ, Tan LML, Ngo RYS. Clinical audit of the microbiology of otorrhoea referred to a tertiary hospital in Singapore. *Singapore Med J*. 2012 Apr;53(4):244-8. PMID: 22511046.
125. Chew KK, Ng KY, Khong WX, Kaur P, Yap JK, Chua A, et al. Clinical evaluation of an in-house human immunodeficiency virus (HIV) genotyping assay for the detection of drug resistance mutations in HIV-1 infected patients in Singapore. *Ann Acad Med Singap*. 2012 Dec;41(12):553-8. PMID: 23303111.
126. Chien JMF, Koh TH, Chan KS, Chuah THC, Tan TT. Successful treatment of NDM-1 *Klebsiella pneumoniae* bacteraemia in a neutropenic patient. *Scand J Infect Dis*. 2012 Apr;44(4):312-4. PMID: 22126430. doi: 10.3109/00365548.2011.633549.
127. Choudhury S, Krishnan PU, Ang BSP. Prevalence of high-level mupirocin resistance among meticillin-resistant *Staphylococcus aureus* isolates in a tertiary care hospital in Singapore. *J Hosp Infect*. 2012 Sep;82(1):56-7. PMID: 22854355. doi: 10.1016/j.jhin.2012.07.002.
128. Choudhury S, Chan KS, Koh TH. Use of meropenem Adatabs dissolved in MacConkey agar for screening NDM-1 positive Enterobacteriaceae in faecal surveillance cultures. *Pathology*. 2012 Aug;44(5):465-8. PMID: 22772339. doi: 10.1097/PAT.0b013e3283557697.

129. Chow ALP, Arah OA, Chan S-P, Poh BF, Krishnan PU, Ng W-K, et al. Alcohol handrubbing and chlorhexidine handwashing protocols for routine hospital practice: a randomized clinical trial of protocol efficacy and time effectiveness. *Am J Infect Control.* 2012 Nov;40(9):800-5. PMID: 22325731. doi: 10.1016/j.ajic.2011.10.005.
130. Cui Y, Kim SH, Kim H, Yeom J, Ko K, Park W, et al. AFM probing the mechanism of synergistic effects of the green tea polyphenol (-)-epigallocatechin-3-gallate (EGCG) with cefotaxime against extended-spectrum beta-lactamase (ESBL)-producing *Escherichia coli*. *PLoS ONE.* 2012;7(11):e48880. doi: <http://dx.doi.org/10.1371/journal.pone.0048880>.
131. Du H, Lo T-M, Sitompul J, Chang MW. Systems-level analysis of *Escherichia coli* response to silver nanoparticles: the roles of anaerobic respiration in microbial resistance. *Biochem Biophys Res Commun.* 2012 Aug 10;424(4):657-62. PMID: 22771582. doi: 10.1016/j.bbrc.2012.06.134.
132. Fukushima K, Tan JP, Korevaar PA, Nelson A, Maune H, Coady DJ, et al. Broad-spectrum antimicrobial supramolecular assemblies with distinctive size and shape. *ACS Nano.* 2012;6(10):9191-9. doi: 10.1021/nn3035217.
133. Han N, Liu X-W, Mu Y. Exploring the mechanism of zanamivir resistance in a neuraminidase mutant: a molecular dynamics study. *PLoS One.* 2012;7(9):e44057. PMID: 22970161. doi: 10.1371/journal.pone.0044057.
134. Huang Y, Wiradharma N, Xu K, Ji Z, Bi S, Li L, et al. Cationic amphiphilic alpha-helical peptides for the treatment of carbapenem-resistant *Acinetobacter baumannii* infection. *Biomaterials.* 2012;33(34):8841-7. doi: 10.1016/j.biomaterials.2012.08.026.
135. Kanagarajan V, Gopalakrishnan M. Pyrimidino piperazinyl acetamides: innovative class of hybrid acetamide drugs as potent antimicrobial and antimycobacterial agents. *PHARMACEUTICAL CHEMISTRY JOURNAL.* 2012 APR;46(1):26-34. PMID: WOS:000304169300007. doi: 10.1007/s11094-012-0729-9.
136. Koh TH, Cao DY, Chan KS, Wijaya L, Low SB, Lam MS, et al. bla(OXA-181)-positive *Klebsiella pneumoniae*, Singapore. *Emerg Infect Dis.* 2012 Sep;18(9):1524-5. PMID: 22931561. doi: 10.3201/eid1809.111727.
137. Koh TH, Tan TT, Khoo CT, Ng LSY, Tan TY, Hsu L-Y, et al. *Acinetobacter calcoaceticus-Acinetobacter baumannii* complex species in clinical specimens in Singapore. *Epidemiol Infect.* 2012 Mar;140(3):535-8. PMID: 21733253. doi: 10.1017/s0950268811001129.
138. Kyaw BM, Arora S, Lim CSD. Bactericidal antibiotic-phytochemical combinations against methicillin resistant *Staphylococcus aureus*. *Braz J Microbiol.* 2012 Jul;43(3):938-45. PMID: 24031910. doi: 10.1590/s1517-838220120003000013.
139. Win M-K, Lee LK, Wong CS, Chow ALP, Ang BSP, Leo YS. Prevalence of and risk factors for MRSA colonization in HIV-positive outpatients in Singapore. *AIDS Res Ther.* 2012 Nov 6;9(1):33. PMID: 23126233. doi: 10.1186/1742-6405-9-33.
140. Lee AS, Ong DC, Wong JC, Siu GK, Yam W-C. High-resolution melting analysis for the rapid detection of fluoroquinolone and streptomycin resistance in *Mycobacterium tuberculosis*. *PLoS One.* 2012;7(2):e31934. PMID: 22363772. doi: 10.1371/journal.pone.0031934.
141. Lee CK, Lee HK, Loh TP, Sethi SK, Koay ES-C, Tang JW-T. An in-house HIV genotyping assay for the detection of drug resistance mutations in Southeast Asian patients infected with HIV-1. *J Med Virol.* 2012 Mar;84(3):394-401. PMID: 22246824. doi: 10.1002/jmv.23202.
142. Liew YX, Lee WHL, Cai Y, Tang SSL, Ong RWQ, Lim CLL, et al. Utility and safety of procalcitonin in an antimicrobial stewardship program (ASP) in patients with malignancies. *Eur J Clin Microbiol Infect Dis.* 2012 Nov;31(11):3041-6. PMID: 22678350. doi: 10.1007/s10096-012-1662-2.
143. Liew YX, Lee WHL, Loh JCZ, Cai Y, Tang SSL, Lim CLL, et al. Impact of an antimicrobial stewardship programme on patient safety in Singapore General Hospital. *Int J Antimicrob Agents.* 2012 Jul;40(1):55-60. PMID: 22591837. doi: 10.1016/j.ijantimicag.2012.03.004.
144. Lye DCB, Earnest A, Ling ML, Lee T-E, Yong H-C, Fisher DA, et al. The impact of multidrug resistance in healthcare-associated and nosocomial Gram-negative bacteraemia on mortality and length of stay: cohort study. *Clin Microbiol Infect.* 2012 May;18(5):502-8. PMID: 21851482. doi: 10.1111/j.1469-0691.2011.03606.x.
145. Ng ES-T, Earnest A, Lye DCB, Ling ML, Ding Y, Hsu L-Y. The excess financial burden of multidrug resistance in severe gram-negative infections in Singaporean hospitals. *Ann Acad Med Singap.* 2012 May;41(5):189-93. PMID: 22760715.
146. Poon LM, Jin J, Chee YL, Ding Y, Lee YM, Chng WJ, et al. Risk factors for adverse outcomes and multidrug-resistant Gram-negative bacteraemia in haematology patients with febrile neutropenia in a Singaporean university hospital. *Singapore Med J.* 2012 Nov;53(11):720-5. PMID: 23192498.

147. Sarathy JP, Dartois V, Lee EJD. The role of transport mechanisms in *Mycobacterium Tuberculosis* drug resistance and tolerance. *Pharmaceuticals.* 2012;5(11):1210-35. doi: 10.3390/ph5111210.
148. Shao Q, Xing B. Enzyme responsive luminescent ruthenium(II) cephalosporin probe for intracellular imaging and photoinactivation of antibiotics resistant bacteria. *Chem Commun (Camb).* 2012 Feb 7;48(12):1739-41. PMID: 22057185. doi: 10.1039/c1cc16165b.
149. Teo JQM, Cai Y, Tang SSL, Lee WHL, Tan TY, Tan TT, et al. Risk factors, molecular epidemiology and outcomes of ertapenem-resistant, carbapenem-susceptible Enterobacteriaceae: a case-case-control study. *PLoS One.* 2012;7(3):e34254. PMID: 22461908. doi: 10.1371/journal.pone.0034254.
150. Teo JQM, Kwa AL-H, Loh J, Chlebicki MP, Lee WHL. The effect of a whole-system approach in an antimicrobial stewardship programme at the Singapore General Hospital. *Eur J Clin Microbiol Infect Dis.* 2012 Jun;31(6):947-55. PMID: 21904857. doi: 10.1007/s10096-011-1391-y.
151. Teo JWP, Ngan G, Balm MN, Jureen R, Krishnan PU, Lin RTP. Molecular characterization of NDM-1 producing Enterobacteriaceae isolates in Singapore hospitals. *Western Pac Surveill Response J.* 2012 Jan;3(1):19-24. PMID: 23908903. doi: 10.5365/wpsar.2011.2.4.010.
152. Vasoo S, Singh K, Chow C, Parthasarathy P, Lin RTP, Hsu L-Y, et al. Health care-associated methicillin-resistant *Staphylococcus aureus* colonization in children attending day care centers in Singapore. *Pediatr Infect Dis J.* 2012 Feb;31(2):213-4. PMID: 22252212. doi: 10.1097/INF.0b013e318243e209.
153. Venkatachalam I, Teo JWP, Balm MN, Fisher DA, Jureen R, Lin RTP. *Klebsiella pneumoniae* Carbapenemase-producing enterobacteria in hospital, Singapore. *Emerg Infect Dis.* 2012 Aug;18(8):1381-3. PMID: 22840461. doi: 10.3201/eid1808.110893.
154. Verrall AJ, Llorin R, Tam VH, Lye DCB, Sulaiman Z, Zhong L, et al. Efficacy of continuous infusion of vancomycin for the outpatient treatment of methicillin-resistant *Staphylococcus aureus* infections. *J Antimicrob Chemother.* 2012 Dec;67(12):2970-3. PMID: 22915464. doi: 10.1093/jac/dks328.
155. Wozniak M, Tiuryn J, Wong L. An approach to identifying drug resistance associated mutations in bacterial strains. *BMC GENOMICS.* 2012 DEC 13;13. PMID: WOS:000317183100003. doi: 10.1186/1471-2164-13-S7-S23.
156. Yeo C-L, Chan DSG, Earnest A, Wu T-S, Yeoh S-F, Lim R, et al. Prospective audit and feedback on antibiotic prescription in an adult hematology-oncology unit in Singapore. *Eur J Clin Microbiol Infect Dis.* 2012 Apr;31(4):583-90. PMID: 21845470. doi: 10.1007/s10096-011-1351-6.
157. Zou G, Kukkaro P, Lok S-M, Ng JK, Tan GK, Hanson BJ, et al. Resistance analysis of an antibody that selectively inhibits dengue virus serotype-1. *Antiviral Res.* 2012 Sep;95(3):216-23. PMID: 22771779. doi: 10.1016/j.antiviral.2012.06.010.
158. Ang TL, Wang L, Ang D, Chiam P, Fock KM, Teo EK. Is there still a role for empiric first-line triple therapy using proton pump inhibitor, amoxicillin and clarithromycin for *Helicobacter pylori* infection in Singapore? Results of a time trend analysis. *J Dig Dis.* 2013 Feb;14(2):100-4. PMID: 23253473. doi: 10.1111/1751-2980.12024.
159. Balm MN, La M-V, Krishnan PU, Jureen R, Lin RTP, Teo JWP. Emergence of *Klebsiella pneumoniae* co-producing NDM-type and OXA-181 carbapenemases. *Clin Microbiol Infect.* 2013 Sep;19(9):E421-3. PMID: 23668475. doi: 10.1111/1469-0691.12247.
160. Balm MN, Lover AA, Salmon S, Tambyah PA, Fisher DA. Progression from new methicillin-resistant *Staphylococcus aureus* colonisation to infection: an observational study in a hospital cohort. *BMC Infect Dis.* 2013 Oct 22;13:491. PMID: 24148135. doi: 10.1186/1471-2334-13-491.
161. Balm MN, Ngan G, Jureen R, Lin RTP, Teo JWP. OXA-181-producing *Klebsiella pneumoniae* establishing in Singapore. *BMC Infect Dis.* 2013 Feb 1;13:58. PMID: 23374756. doi: 10.1186/1471-2334-13-58.
162. Chee CBE, Hsu L-Y, Sng L-H, Leo YS, Cutter J, Wang Y-T. MDR TB transmission, Singapore. *Emerg Infect Dis.* 2013 Jul;19(7):1151-2. PMID: 23763791. doi: 10.3201/eid1907.120372.
163. Chin W, Yang C, Ng VWL, Huang Y, Cheng J, Tong YW, et al. Biodegradable Broad-Spectrum Antimicrobial Polycarbonates: Investigating the Role of Chemical Structure on Activity and Selectivity. *MACROMOLECULES.* 2013 NOV 26;46(22):8797-807. PMID: WOS:000327752900006. doi: 10.1021/ma4019685.
164. Ch'ng J-H, Mok S, Bozdech Z, Lear MJ, Boudhar A, Russell B, et al. A whole cell pathway screen reveals seven novel chemosensitizers to combat chloroquine resistant malaria. *Sci Rep.* 2013;3:1734. PMID: 23615863. doi: 10.1038/srep01734.

165. Chong C-W, Ormston VE, Tan AB-H. Epidemiology of hand infection--a comparative study between year 2000 and 2009. *Hand Surg.* 2013;18(3):307-12. PMID: 24156570. doi: 10.1142/s0218810413500317.
166. Chow WL, Soe TA, Lim WW, Lim JFY, Kurup A, Ling ML, et al. Efficacy of titanium dioxide compounds in preventing environmental contamination by meticillin resistant *Staphylococcus aureus* (MRSA). *International Journal of Infection Control.* 2013 09/06;9. doi: 10.3396/IJIC.v9i3.022.13.
167. Chua SL, Tan SY-Y, Rybtke MT, Chen Y, Rice SA, Kjelleberg S, et al. Bis-(3'-5')-cyclic dimeric GMP regulates antimicrobial peptide resistance in *Pseudomonas aeruginosa*. *Antimicrob Agents Chemother.* 2013 May;57(5):2066-75. PMID: 23403434. doi: 10.1128/aac.02499-12.
168. Fisher DA, Tambyah PA, Lin RTP, Jureen R, Cook AR, Lim A, et al. Sustained meticillin-resistant *Staphylococcus aureus* control in a hyper-endemic tertiary acute care hospital with infrastructure challenges in Singapore. *J Hosp Infect.* 2013 Oct;85(2):141-8. PMID: 24011440. doi: 10.1016/j.jhin.2013.07.005.
169. Fukushima K, Liu S, Wu H, Engler AC, Coady DJ, Maune H, et al. Supramolecular high-aspect ratio assemblies with strong antifungal activity. *Nature Communications.* 2013;4. doi: 10.1038/ncomms3861.
170. Goh S, Hussain H, Chang BJ, Emmett W, Riley TV, Mullany P. Phage phi C2 Mediates Transduction of Tn6215, Encoding Erythromycin Resistance, between *Clostridium difficile* Strains. *MBIO.* 2013 NOV-DEC;4(6):e00840-13. PMID: WOS:000329174500027. doi: 10.1128/mBio.00840-13.
171. Grant D, Koh TH, Tan YE, Hsu L-Y, Kurup A, Donahue SK, et al. An Outbreak of Community Associated Methicillin Resistant *Staphylococcus aureus* Subtype USA300 at an International School in Singapore. *Ann Acad Med Singap.* 2013 Nov;42(11):575-8. PMID: 24356653.
172. Heng YK, Tan KT, Sen P, Chow ALP, Leo YS, Lye DCB, et al. *Staphylococcus aureus* and topical fusidic acid use: results of a clinical audit on antimicrobial resistance. *Int J Dermatol.* 2013 Jul;52(7):876-81. PMID: 23432159. doi: 10.1111/j.1365-4632.2012.05747.x.
173. Hon PY, Chan KS, Holden MT, Harris SR, Tan TY, Zu Y-B, et al. Arginine catabolic mobile element in methicillin-resistant *Staphylococcus aureus* (MRSA) clonal group ST239-MRSA-III isolates in Singapore: implications for PCR-based screening tests. *Antimicrob Agents Chemother.* 2013 Mar;57(3):1563-4. PMID: 23318798. doi: 10.1128/aac.02518-12.
174. Koh J-J, Qiu S, Zou H, Lakshminarayanan R, Li J, Zhou X, et al. Rapid bactericidal action of alpha-mangostin against MRSA as an outcome of membrane targeting. *Biochim Biophys Acta.* 2013 Feb;1828(2):834-44. PMID: 22982495. doi: 10.1016/j.bbamem.2012.09.004.
175. Koh TH, Cao DY, Quek YS, Bacon A, Hsu L-Y, Ooi EE. Acquired carbapenemases in Enterobactericeae in Singapore, 1996-2012. *Pathology.* 2013 Oct;45(6):600-3. PMID: 24018814. doi: 10.1097/PAT.0b013e3283650b1e.
176. Lee GH, Aung MO, Dan YY, Lee YM, Mak B, Low HC, et al. Do different lamivudine-resistant hepatitis B genotypes carry the same risk of entecavir resistance? *J Med Virol.* 2013 Jan;85(1):26-33. PMID: 23023992. doi: 10.1002/jmv.23392.
177. Lee GH, Inoue M, Toh JK-P, Chong RH-H, Aung MO, Koay ES-C, et al. Two-step evolution of the hepatitis B drug-resistant mutations in a patient who developed primary entecavir resistance. *Liver Int.* 2013 Apr;33(4):642-6. PMID: 23346997. doi: 10.1111/liv.12104.
178. Lee LK, Win M-K, Veeraraghavan MA, Wong CS, Chow ALP, Leo YS. Short communication: risk factors for methicillin-resistant *Staphylococcus aureus* colonization among HIV patients at hospital admission. *AIDS Res Hum Retroviruses.* 2013 May;29(5):796-8. PMID: 23517521. doi: 10.1089/aid.2012.0074.
179. Li Y, Fukushima K, Coady DJ, Engler AC, Liu S, Huang Y, et al. Broad-spectrum antimicrobial and biofilm-disrupting hydrogels: stereocomplex-driven supramolecular assemblies. *Angew Chem Int Ed Engl.* 2013 Jan 7;52(2):674-8. PMID: 23161823. doi: 10.1002/anie.201206053.
180. Liew YX, Tan TT, Lee WHL, Ng J-L, Chia D-Q, Wong GC, et al. Risk factors for extreme-drug resistant *Pseudomonas aeruginosa* infections in patients with hematologic malignancies. *Am J Infect Control.* 2013 Feb;41(2):140-4. PMID: 22795726. doi: 10.1016/j.ajic.2012.02.025.
181. Lim CLL, Lee WHL, Lee AL-C, Liew LT-T, Nah SC, Wan CN, et al. Evaluation of Ertapenem use with impact assessment on extended-spectrum beta-lactamases (ESBL) production and gram-negative resistance in Singapore General Hospital (SGH). *BMC Infect Dis.* 2013 Nov 6;13:523. PMID: 24195651. doi: 10.1186/1471-2334-13-523.
182. Lim S-G, Lee GH, Lim K, Tan PS. Resistance is no Longer a Problem with Entecavir and Tenofovir. *Current Hepatitis Reports.* 2013 06/01;12. doi: 10.1007/s11901-013-0168-4.
183. Marimuthu K, Ng TM, Teng CB, Lim T-P, Koh TH, Tan TY, et al. Risk factors and treatment outcome of ertapenem non-susceptible enterobacteriaceae bacteraemia. *J Infect.* 2013 Mar;66(3):294-6. PMID: 23201151. doi: 10.1016/j.jinf.2012.11.010.

184. Molton JS, Tambyah PA, Ang BSP, Ling ML, Fisher DA. The global spread of healthcare-associated multidrug-resistant bacteria: a perspective from Asia. *Clin Infect Dis.* 2013 May;56(9):1310-8. PMID: 23334810. doi: 10.1093/cid/cit020.
185. Ng VWL, Ke X-Y, Lee AL, Yang YY, Hedrick JL. Synergistic co-delivery of membrane-disrupting polymers with commercial antibiotics against highly opportunistic bacteria. *Advanced Materials.* 2013;25(46):6730-6. doi: 10.1002/adma.201302952.
186. Ong ZY, Gao S, Yang YY. Short Synthetic beta-Sheet Forming Peptide Amphiphiles as Broad Spectrum Antimicrobials with Antibiofilm and Endotoxin Neutralizing Capabilities. *ADVANCED FUNCTIONAL MATERIALS.* 2013 AUG 7;23(29):3682-92. PMID: WOS:000327490500011. doi: 10.1002/adfm.201202850.
187. Qiu G, Zeng P, Duan L, Xiao S, Song Y. Combination of upflow anaerobic sludge blanket (UASB) and membrane bioreactor (MBR) for berberine reduction from wastewater and the effects of berberine on bacterial community dynamics. *Journal of Hazardous Materials.* 2013;246-247:34-43. doi: 10.1016/j.jhazmat.2012.12.010.
188. Russell B, Malleret B, Suwanarusk R, Anthony C, Kanlaya S, Lau Y, et al. Field-based flow cytometry for ex vivo characterization of *Plasmodium vivax* and *P. falciparum* antimalarial sensitivity. *Antimicrob Agents Chemother.* 2013 Oct;57(10):5170-4. PMID: 23877705. doi: 10.1128/aac.00682-13.
189. Samy PR, Gopalakrishnakone P, Satyanarayana-Jois SD, Stiles BG, Chow VT. Snake Venom Proteins and Peptides as Novel Antibiotics Against Microbial Infections. *Current Proteomics.* 2013;10(1):10-28. doi: <http://dx.doi.org/10.2174/157016461131001003>.
190. Samy RP, Manikandan J, Qahtani MA. Evaluation of aromatic plants and compounds used to fight multidrug resistant infections. *Evid Based Complement Alternat Med.* 2013;2013:525613. PMID: 24223059. doi: 10.1155/2013/525613.
191. Sarathy JP, Dartois V, Dick T, Gengenbacher M. Reduced drug uptake in phenotypically resistant nutrient-starved nonreplicating *Mycobacterium tuberculosis*. *Antimicrob Agents Chemother.* 2013 Apr;57(4):1648-53. PMID: 23335744. doi: 10.1128/aac.02202-12.
192. Sarathy JP, Lee EJD, Dartois V. Polyamines inhibit porin-mediated fluoroquinolone uptake in mycobacteria. *PLoS One.* 2013;8(6):e65806. PMID: 23755283. doi: 10.1371/journal.pone.0065806.
193. Seah J, Lye DCB, Ng TM, Krishnan PU, Choudhury S, Teng CB. Vancomycin monotherapy vs. combination therapy for the treatment of persistent methicillin-resistant *Staphylococcus aureus* bacteremia. *Virulence.* 2013 Nov 15;4(8):734-9. PMID: 24165210. doi: 10.4161/viru.26909.
194. Shao Q, Zheng Y, Dong X, Tang K, Yan X, Xing B. A covalent reporter of β -lactamase activity for fluorescent imaging and rapid screening of antibiotic-resistant bacteria. *Chemistry.* 2013 Aug 12;19(33):10903-10. PMID: 23836436. doi: 10.1002/chem.201301654.
195. Sim JHC, Ng LSY, Eng LC, Chan KS, Tan TY. Characterisation of significant Gram positive bacilli from soft tissue infections. *Pathology.* 2013 Feb;45(2):167-71. PMID: 23277175. doi: 10.1097/PAT.0b013e32835c9863.
196. Soon MML, Madigan E, Jones KR, Salata RA. An exploration of the psychologic impact of contact isolation on patients in Singapore. *Am J Infect Control.* 2013 Oct;41(10):e111-3. PMID: 23663862. doi: 10.1016/j.ajic.2013.01.037.
197. Tan PS, Aung MO, Dan YY, Lee YM, Lim K, Low HC, et al. Liver disease progression and virological response: entecavir rescue still possible in the setting of rtM204I lamivudine-resistant mutation. *Gut.* 2013 May;62(5):801-2. PMID: 23144055. doi: 10.1136/gutjnl-2012-303647.
198. Tan SY-Y, Chua SL, Liu Y, Høiby N, Andersen LP, Givskov M, et al. Comparative genomic analysis of rapid evolution of an extreme-drug-resistant *Acinetobacter baumannii* clone. *Genome Biol Evol.* 2013;5(5):807-18. PMID: 23538992. doi: 10.1093/gbe/evt047.
199. Teo JQM, Tan TY, Hon PY, Lee WHL, Koh TH, Krishnan PU, et al. ST22 and ST239 MRSA duopoly in Singaporean hospitals: 2006-2010. *Epidemiol Infect.* 2013 Jan;141(1):153-7. PMID: 22394568. doi: 10.1017/s0950268812000337.
200. Teo JWP, La M-V, Krishnan PU, Ang BSP, Jureen R, Lin RTP. *Enterobacter cloacae* producing an uncommon class A carbapenemase, IMI-1, from Singapore. *J Med Microbiol.* 2013 Jul;62(Pt 7):1086-8. PMID: 23558141. doi: 10.1099/jmm.0.053363-0.
201. Teo JWP, Kurup A, Lin RTP, Koh TH. Emergence of clinical *Klebsiella pneumoniae* producing OXA-232 carbapenemase in Singapore. *New Microbes New Infect.* 2013 Oct;1(1):13-5. PMID: 25356318. doi: 10.1002/2052-2975.4.
202. Vasudevan A, Li J, Mukhopadhyay A, Goh EYY, Chuang L, Tambyah PA. Inappropriate empirical antimicrobial therapy for multidrug-resistant organisms in critically ill patients with pneumonia is not an independent risk factor for mortality: Results of a prospective observational study of 758 patients. *J Glob Antimicrob Resist.* 2013 Sep;1(3):123-30. PMID: 27873622. doi: 10.1016/j.jgar.2013.03.008.

203. Vasudevan A, Mukhopadhyay A, Goh EYY, Li J, Tambyah PA. Risk factors for infection/colonization caused by resistant Gram negative bacilli in critically ill patients (an observational study of 1633 critically ill patients). *Prev Med*. 2013;57 Suppl:S70-3. PMID: 23246839. doi: 10.1016/j.ypmed.2012.12.003.
204. Verrall AJ, Merchant R, Dillon J, Ying D, Fisher DA. Impact of nursing home residence on hospital epidemiology of meticillin-resistant *Staphylococcus aureus*: a perspective from Asia. *J Hosp Infect*. 2013 Mar;83(3):250-2. PMID: 23374286. doi: 10.1016/j.jhin.2012.11.021.
205. Win M-K, Yung C-F, Poh BF, Krishnan PU, Seet S-K, Leo YS, et al. Evaluation of universal methicillin-resistant *Staphylococcus aureus* screening using nasal polymerase chain reaction compared with nasal, axilla, and groin and throat and perianal cultures in a hospital setting. *Infect Control Hosp Epidemiol*. 2013 Dec;34(12):1335-7. PMID: 24225629. doi: 10.1086/673989.
206. Xia E, Khong WX, Marimuthu K, Xu W, Ong RT-H, Tan EL, et al. Draft Genome Sequence of a Multidrug-Resistant New Delhi Metallo- β -Lactamase-1 (NDM-1)-Producing *Escherichia coli* Isolate Obtained in Singapore. *Genome Announc*. 2013 Dec 19;1(6). PMID: 24356827. doi: 10.1128/genomeA.01020-13.
207. Xiong P, Hu J. Inactivation/reactivation of antibiotic-resistant bacteria by a novel UVA/LED/TiO₂ system. *Water Res*. 2013 Sep 1;47(13):4547-55. PMID: 23764604. doi: 10.1016/j.watres.2013.04.056.
208. Yang Y, Chua KL. Assessment of the effect of efflux pump inhibitors on *in vitro* antimicrobial susceptibility of multidrug-resistant *Acinetobacter baumannii*. *Int J Antimicrob Agents*. 2013 Sep;42(3):283-4. PMID: 23837925. doi: 10.1016/j.ijantimicag.2013.05.011.
209. Yokokawa F, Wang G, Chan WL, Ang SH, Wong J, Ma I, et al. Discovery of tetrahydropyrazolopyrimidine carboxamide derivatives as potent and orally active antitubercular agents. *ACS Med Chem Lett*. 2013 May 9;4(5):451-5. PMID: 24900693. doi: 10.1021/ml400071a.
210. Yuan X, Setyawati MI, Tan AS, Ong CN, Leong DT, Xie J. Highly luminescent silver nanoclusters with tunable emissions: cyclic reduction-decomposition synthesis and antimicrobial properties. *NPG ASIA MATERIALS*. 2013 FEB;5. PMID: WOS:000315349500005. doi: 10.1038/am.2013.3.
211. Zou H, Koh J-J, Li J, Qiu S, Aung TT, Lin H, et al. Design and synthesis of amphiphilic xanthone-based, membrane-targeting antimicrobials with improved membrane selectivity. *J Med Chem*. 2013 Mar 28;56(6):2359-73. PMID: 23441632. doi: 10.1021/jm301683j.
212. Bandyopadhyay S, Loh RJ, Lim B, Sanjeev R, Woon YX, Chong KY, et al. Solution structures and model membrane interactions of Ctriporin, an anti-methicillin-resistant *Staphylococcus aureus* Peptide from Scorpion Venom. *Biopolymers*. 2014 Dec;101(12):1143-53. PMID: 24947608. doi: 10.1002/bip.22519.
213. Bayen S, Yi X, Segovia E, Zhou Z, Kelly BC. Analysis of selected antibiotics in surface freshwater and seawater using direct injection in liquid chromatography electrospray ionization tandem mass spectrometry. *J Chromatogr A*. 2014 Apr 18;1338:38-43. PMID: 24642398. doi: 10.1016/j.chroma.2014.02.034.
214. Castillo CF, Ling MH. Resistant traits in digital organisms do not revert preselection status despite extended deselection: implications to microbial antibiotics resistance. *Biomed Res Int*. 2014;2014:648389. PMID: 24977157. doi: 10.1155/2014/648389.
215. Chen HH, Hon PY, Hsu L-Y. Ceftaroline--An Anti-MRSA Cephalosporin and Its Implications for Singapore. *Ann Acad Med Singap*. 2014 Mar;43(3):177-86. PMID: 24714713.
216. Ch'ng J-H, Lee YQ, Gun S, Chia WN, Chang Z-W, Wong L-K, et al. Validation of a chloroquine-induced cell death mechanism for clinical use against malaria. *Cell Death Dis*. 2014 Jun 26;5(6):e1305. PMID: 24967967. doi: 10.1038/cddis.2014.265.
217. Chow JY, Yang Y, Tay SB, Yew WS, Chua KL. Disruption of biofilm formation by the human pathogen *Acinetobacter baumannii* using engineered quorum-quenching lactonases. *Antimicrob Agents Chemother*. 2014;58(3):1802-5. PMID: 24379199. doi: 10.1128/aac.02410-13.
218. Coady DJ, Ong ZY, Lee PS, Venkataraman S, Chin W, Engler AC, et al. Enhancement of Cationic Antimicrobial Materials via Cholesterol Incorporation. *ADVANCED HEALTHCARE MATERIALS*. 2014 JUN;3(6):882-9. PMID: WOS:000337692900013. doi: 10.1002/adhm.201300554.
219. Deng C-L, Yeo H, Ye H-Q, Liu S-Q, Shang B-D, Gong P, et al. Inhibition of enterovirus 71 by adenosine analog NITD008. *J Virol*. 2014 Oct;88(20):11915-23. PMID: 25100827. doi: 10.1128/jvi.01207-14.
220. Deng Y, Lim A, Lee J, Chen S, An S, Dong Y-H, et al. Diffusible signal factor (DSF) quorum sensing signal and structurally related molecules enhance the antimicrobial efficacy of antibiotics against some bacterial pathogens. *BMC Microbiol*. 2014 Feb 27;14:51. PMID: 24575808. doi: 10.1186/1471-2180-14-51.

221. Gopal P, Dick T. Reactive dirty fragments: implications for tuberculosis drug discovery. *Curr Opin Microbiol.* 2014 Oct;21:7-12. PMID: 25078318. doi: 10.1016/j.mib.2014.06.015.
222. Hon PY, Koh TH, Tan TY, Krishnan PU, Leong JW-Y, Jureen R, et al. Changing molecular epidemiology and high rates of mupirocin resistance among meticillin-resistant *Staphylococcus aureus* in Singaporean hospitals. *J Glob Antimicrob Resist.* 2014 Mar;2(1):53-5. PMID: 27873639. doi: 10.1016/j.jgar.2013.10.002.
223. Jauneikaite E, Jefferies JMC, Churton NWV, Lin RTP, Hibberd ML, Clarke SC. Genetic diversity of *Streptococcus pneumoniae* causing meningitis and sepsis in Singapore during the first year of PCV7 implementation. *Emerging Microbes and Infections.* 2014;3. doi: 10.1038/emi.2014.37.
224. Khara JS, Wang Y, Ke X-Y, Liu S, Newton SM, Langford PR, et al. Anti-mycobacterial activities of synthetic cationic α -helical peptides and their synergism with rifampicin. *Biomaterials.* 2014 Feb;35(6):2032-8. PMID: 24314557. doi: 10.1016/j.biomaterials.2013.11.035.
225. Koh TH, Cao DY, Tee NWS, Teo JWP. *Escherichia coli* with bla(IMP-8) in Singapore. *Antimicrob Agents Chemother.* 2014;58(1):617. PMID: 24145544. doi: 10.1128/aac.01754-13.
226. La M-V, Jureen R, Lin RTP, Teo JWP. Unusual detection of an *Acinetobacter* class D carbapenemase gene, blaOXA-23, in a clinical *Escherichia coli* isolate. *J Clin Microbiol.* 2014 Oct;52(10):3822-3. PMID: 25031438. doi: 10.1128/jcm.01566-14.
227. Lescar J, Meyer I, Akshita K, Srinivasaraghavan K, Verma CS, Palous M, et al. *Aspergillus fumigatus* harbouring the sole Y121F mutation shows decreased susceptibility to voriconazole but maintained susceptibility to itraconazole and posaconazole. *J Antimicrob Chemother.* 2014 Dec;69(12):3244-7. PMID: 25125676. doi: 10.1093/jac/dku316.
228. Li X, Li P, Saravanan R, Basu A, Mishra B, Lim SH, et al. Antimicrobial functionalization of silicone surfaces with engineered short peptides having broad spectrum antimicrobial and salt-resistant properties. *Acta Biomater.* 2014 Jan;10(1):258-66. PMID: 24056098. doi: 10.1016/j.actbio.2013.09.009.
229. Liu S, Venkataraman S, Ong ZY, Chan JM, Yang C, Yang YY, et al. Overcoming multidrug resistance in microbials using nanostructures self-assembled from cationic bent-core oligomers. *Small.* 2014 Oct 29;10(20):4130-5. PMID: 24975599. doi: 10.1002/smll.201303921.
230. Loh CC, Suwanarusk R, Lee YQ, Chan KW, Choy K-Y, Rénia L, et al. Characterization of the commercially-available fluorescent chloroquine-BODIPY conjugate, LynxTag-CQGREEN, as a marker for chloroquine resistance and uptake in a 96-well plate assay. *PLoS One.* 2014;9(10):e110800. PMID: 25343249. doi: 10.1371/journal.pone.0110800.
231. Mani V, Wang S, Inci F, De Libero G, Singhal A, Demirci U. Emerging technologies for monitoring drug-resistant tuberculosis at the point-of-care. *Adv Drug Deliv Rev.* 2014 Nov 30;78:105-17. PMID: 24882226. doi: 10.1016/j.addr.2014.05.015.
232. Marimuthu K, Teo JWP, Poh BF, Ong JHC, Kum JQ, Lye DCB, et al. First report of emergence of OXA-48 carbapenemase-producing Enterobacteriaceae in Singapore: proactive or reactive infection control strategy? *Am J Infect Control.* 2014 May;42(5):577-8. PMID: 24773801. doi: 10.1016/j.ajic.2014.01.021.
233. Marimuthu K, Pittet D, Harbarth S. The effect of improved hand hygiene on nosocomial MRSA control. *Antimicrob Resist Infect Control.* 2014;3:34. PMID: 25937922. doi: 10.1186/2047-2994-3-34.
234. Marimuthu K, Harbarth S. Screening for methicillin-resistant *Staphylococcus aureus* . . . all doors closed? *Current Opinion in Infectious Diseases.* 2014;27(4):356-62. doi: 10.1097/QCO.0000000000000081.
235. Mok S, Liang K-Y, Lim E-H, Huang X, Zhu L, Preiser PR, et al. Structural polymorphism in the promoter of pfmrp2 confers *Plasmodium falciparum* tolerance to quinoline drugs. *Mol Microbiol.* 2014 Mar;91(5):918-34. PMID: 24372851. doi: 10.1111/mmi.12505.
236. Ng TM, Liew YX, Teng CB, Ling LM, Ang BSP, Lye DCB. An interactive, point-of-care, computerised antibiotic prescription decision support system and quality of antibiotic prescription in the management of complicated urinary tract infection. *J Glob Antimicrob Resist.* 2014 Jun;2(2):127-8. PMID: 27873592. doi: 10.1016/j.jgar.2014.03.001.
237. Ng TM, Teng CB, Lye DCB, Apisarnthanarak A. A multicenter case-case control study for risk factors and outcomes of extensively drug-resistant *Acinetobacter baumannii* bacteremia. *Infect Control Hosp Epidemiol.* 2014 Jan;35(1):49-55. PMID: 24334798. doi: 10.1086/674387.
238. Ng VWL, Chan JM, Sardon H, Ono RJ, García JM, Yang YY, et al. Antimicrobial hydrogels: A new weapon in the arsenal against multidrug-resistant infections. *ADVANCED DRUG DELIVERY REVIEWS.* 2014 NOV 30;78:46-62. PMID: WOS:000358460400005. doi: 10.1016/j.addr.2014.10.028.
239. Ong ZY, Cheng J, Huang Y, Xu K, Ji Z, Fan W, et al. Effect of stereochemistry, chain length and sequence pattern on antimicrobial properties of short synthetic β -sheet forming peptide amphiphiles. *Biomaterials.* 2014 Jan;35(4):1315-25. PMID: 24211081. doi: 10.1016/j.biomaterials.2013.10.053.

240. Samy RP, Manikandan J, Sethi G, Franco OL, Okonkwo JC, Stiles BG, et al. Snake venom proteins: Development into antimicrobial and wound healing agents. *Mini-Reviews in Organic Chemistry*. 2014;11(1):4-14. doi: 10.2174/1570193X1101140402100131.
241. Seah XVF, Ong YLR, Tan SW, Krishnaswamy G, Chong CY, Tan NWH, et al. Impact of an antimicrobial stewardship program on the use of carbapenems in a tertiary women's and children's hospital, Singapore. *Pharmacotherapy*. 2014 Nov;34(11):1141-50. PMID: 25283969. doi: 10.1002/phar.1490.
242. Shekar S, Yeo ZX, Wong JC, Chan MKL, Ong DC, Tongyoo P, et al. Detecting novel genetic variants associated with isoniazid-resistant *Mycobacterium tuberculosis*. *PLoS One*. 2014;9(7):e102383. PMID: 25025225. doi: 10.1371/journal.pone.0102383.
243. Singhal A, Liu J, Kumar P, Gan SH, Leow MK-S, Paleja B, et al. Metformin as adjunct antituberculosis therapy. *Sci Transl Med*. 2014 Nov 19;6(263):263ra159. PMID: 25411472. doi: 10.1126/scitranslmed.3009885.
244. Song M, Ang TL. Second and third line treatment options for *Helicobacter pylori* eradication. *World J Gastroenterol*. 2014 Feb 14;20(6):1517-28. PMID: 24587627. doi: 10.3748/wjg.v20.i6.1517.
245. Tan MW, Lye DCB, Ng TM, Nikolaou M, Tam VH. Mathematical model to quantify the effects of risk factors on carbapenem-resistant *Acinetobacter baumannii*. *Antimicrob Agents Chemother*. 2014 Sep;58(9):5239-44. PMID: 24957824. doi: 10.1128/aac.02791-14.
246. Tan TT, Ling ML, Tan BH, Koh TH. Multidrug resistant, blaVVB positive *Pseudomonas aeruginosa* causing high mortality among haematology patients. *Pathology*. 2014 Dec;46(7):650-2. PMID: 25393259. doi: 10.1097/pat.0000000000000177.
247. Tan YE, Ng LSY, Tan TY. Evaluation of *Enterococcus faecalis* clinical isolates with 'penicillin-resistant, ampicillin-susceptible' phenotype as reported by Vitek-2 Compact system. *Pathology*. 2014 Oct;46(6):544-50. PMID: 25158809. doi: 10.1097/pat.0000000000000146.
248. Tang SSL, Apisarnthanarak A, Hsu L-Y. Mechanisms of β-lactam antimicrobial resistance and epidemiology of major community- and healthcare-associated multidrug-resistant bacteria. *Adv Drug Deliv Rev*. 2014 Nov 30;78:3-13. PMID: 25134490. doi: 10.1016/j.addr.2014.08.003.
249. Teo JWP, Tan SY-Y, Liu Y, Tay M, Ding Y, Li Y, et al. Comparative genomic analysis of malaria mosquito vector-associated novel pathogen *Elizabethkingia anophelis*. *Genome Biol Evol*. 2014 May 6(5):1158-65. PMID: 24803570. doi: 10.1093/gbe/evu094.
250. Teo JWP, Tan P, La M-V, Krishnan PU, Tee NWS, Koh TH, et al. Surveillance trends of carbapenem-resistant Enterobacteriaceae from Singapore, 2010-2013. *J Glob Antimicrob Resist*. 2014 Jun;2(2):99-102. PMID: 27873598. doi: 10.1016/j.jgar.2013.11.003.
251. Vasudevan A, Mukhopadhyay A, Li J, Goh EYY, Tambyah PA. A prediction tool for nosocomial multi-drug Resistant Gram-Negative Bacilli infections in critically ill patients - prospective observational study. *BMC Infect Dis*. 2014 Nov 25;14:615. PMID: 25420613. doi: 10.1186/s12879-014-0615-z.
252. Venkatachalam I, Hsu L-Y, Fisher DA, Lye DCB, Ling ML, Tambyah PA, et al. Multidrug-resistant gram-negative bloodstream infections among residents of long-term care facilities. *Infect Control Hosp Epidemiol*. 2014 May;35(5):519-26. PMID: 24709720. doi: 10.1086/675823.
253. Wang Y, Ke X-Y, Khara JS, Bahety P, Liu S, Seow SV, et al. Synthetic modifications of the immunomodulating peptide thymopentin to confer anti-mycobacterial activity. *Biomaterials*. 2014 Mar;35(9):3102-9. PMID: 24411680. doi: 10.1016/j.biomaterials.2013.12.049.
254. Wen H, Wang K, Liu Y, Tay M, Lauro FM, Huang H, et al. Population dynamics of an *Acinetobacter baumannii* clonal complex during colonization of patients. *Journal of Clinical Microbiology*. 2014;52(9):3200-8. doi: 10.1128/JCM.00921-14.
255. Wozniak M, Tiuryn J, Wong L. GWAMAR: Genome-wide assessment of mutations associated with drug resistance in bacteria. *BMC GENOMICS*. 2014 DEC 12;15. PMID: WOS:000346166900010. doi: 10.1186/1471-2164-15-S10-S10.
256. Wu Z, Mirza H, Tan KSW. Intra-subtype variation in enteroadhesion accounts for differences in epithelial barrier disruption and is associated with metronidazole resistance in *Blastocystis* subtype-7. *PLoS Negl Trop Dis*. 2014 May;8(5):e2885. PMID: 24851944. doi: 10.1371/journal.pntd.0002885.
257. Yeoh LY, Tan FLG, Willis GC, Ooi ST. Methicillin-resistant *Staphylococcus aureus* carriage in hospitalized chronic hemodialysis patients and its predisposing factors. *Hemodial Int*. 2014 Jan;18(1):142-7. PMID: 23763574. doi: 10.1111/hdi.12061.
258. Young BE, Lye DCB, Krishnan PU, Chan S-P, Leo YS. A prospective observational study of the prevalence and risk factors for colonization by antibiotic resistant bacteria in patients at admission to hospital in Singapore. *BMC Infect Dis*. 2014 Jun 2;14:298. PMID: 24889720. doi: 10.1186/1471-2334-14-298.

259. Zhou YP, Wilder-Smith A, Hsu L-Y. The role of international travel in the spread of methicillin-resistant *Staphylococcus aureus*. *J Travel Med.* 2014 Jul-Aug;21(4):272-81. PMID: 24894491. doi: 10.1111/jtm.12133.
260. Zhou YP, Chua NGS, Chlebicki MP, Lee WHL, Kwa AL-H. Nonattenuated Polymyxin B Used for the Treatment of Extreme-Drug Resistant *Acinetobacter baumannii*-Related Infections in Patients with Preexisting End Stage Renal Failure. *Case Rep Infect Dis.* 2014;2014:573279. PMID: 24995135. doi: 10.1155/2014/573279.
261. Anusha S, Sinha A, Babu Rajeev C, Chu TT, Mathai J, Huang X, et al. Synthesis, characterization and in vitro evaluation of novel enantiomerically-pure sulphonamide antimalarials. *Org Biomol Chem.* 2015 Nov 21;13(43):10681-90. PMID: 26347024. doi: 10.1039/c5ob01479d.
262. Cai Y, Leck H, Lim T-P, Teo JQM, Lee WHL, Hsu L-Y, et al. Using an Adenosine Triphosphate Bioluminescent Assay to Determine Effective Antibiotic Combinations against Carbapenem-Resistant Gram Negative Bacteria within 24 Hours. *PLoS One.* 2015;10(10):e0140446. PMID: 26460891. doi: 10.1371/journal.pone.0140446.
263. Chee CBE, Gan SH, Ong RT-H, Sng L-H, Wong CW, Cutter J, et al. Multidrug-resistant tuberculosis outbreak in gaming centers, Singapore, 2012. *Emerg Infect Dis.* 2015 Jan;21(1):179-80. PMID: 25531726. doi: 10.3201/eid2101.141159.
264. Cheng J, Chin W, Dong H, Xu L, Zhong G, Huang Y, et al. Biodegradable Antimicrobial Polycarbonates with In Vivo Efficacy against Multidrug-Resistant MRSA Systemic Infection. *Adv Healthc Mater.* 2015 Oct;4(14):2128-36. PMID: 26331284. doi: 10.1002/adhm.201500471.
265. Cherng BPZ, Tan TT, Tan BH. Resistant cytomegalovirus infection in renal transplant recipients. *PROCEEDINGS OF SINGAPORE HEALTHCARE.* 2015 DEC;24(4):243-8. PMID: WOS:000218309300008. doi: 10.1177/2010105815611811.
266. Ch'ng J-H, Ursing J, Tan KSW. Microbial hara-kiri: Exploiting lysosomal cell death in malaria parasites. *Microb Cell.* 2015 Jan 12;2(2):57-8. PMID: 28357276. doi: 10.15698/mic2015.02.186.
267. Choudhury S, Leong JW-Y, Krishnan PU. In vitro susceptibilities of clinical isolates of carbapenemase-producing Enterobacteriaceae to fosfomycin and tigecycline. *Clin Microbiol Infect.* 2015 Oct;21(10):e75-6. PMID: 26086572. doi: 10.1016/j.cmi.2015.06.005.
268. Chow ALP, Lye DCB, Arah OA. Psychosocial determinants of physicians' acceptance of recommendations by antibiotic computerised decision support systems: A mixed methods study. *Int J Antimicrob Agents.* 2015 Mar;45(3):295-304. PMID: 25434998. doi: 10.1016/j.ijantimicag.2014.10.009.
269. Chow ALP, Lye DCB, Arah OA. Mortality Benefits of Antibiotic Computerised Decision Support System: Modifying Effects of Age. *Sci Rep.* 2015 Nov 30;5:17346. PMID: 26617195. doi: 10.1038/srep17346.
270. Chua AP-G, Hoo GS-R, Chee CBE, Wang Y-T. First use of bedaquiline in a patient with XDR-TB in Singapore. *BMJ Case Rep.* 2015 Sep 23;2015. PMID: 26400590. doi: 10.1136/bcr-2015-210961.
271. Chua NGS, Zhou YP, Tan TT, Lingegowda PB, Lee WHL, Lim T-P, et al. Polymyxin B with dual carbapenem combination therapy against carbapenemase-producing *Klebsiella pneumoniae*. *J Infect.* 2015 Mar;70(3):309-11. PMID: 25305144. doi: 10.1016/j.jinf.2014.10.001.
272. Chung SJ, Lim T-P, Kwa AL-H. Intravenous ceftazidime 200 mg administered every 8 h is safe and adequate for meticillin-resistant *Staphylococcus aureus* bloodstream infections in end-stage renal failure patients on haemodialysis: a case study. *Int J Antimicrob Agents.* 2015 Dec;46(6):720-1. PMID: 26607339. doi: 10.1016/j.ijantimicag.2015.09.012.
273. Duan R, Lazim R, Zhang D-W. Understanding the basis of I50V-induced affinity decrease in HIV-1 protease via molecular dynamics simulations using polarized force field. *J Comput Chem.* 2015 Sep 30;36(25):1885-92. PMID: 26198456. doi: 10.1002/jcc.24020.
274. Feng G, Yuan Y, Fang H, Zhang R, Xing B, Zhang G, et al. A light-up probe with aggregation-induced emission characteristics (AIE) for selective imaging, naked-eye detection and photodynamic killing of Gram-positive bacteria. *Chem Commun (Camb).* 2015 Aug 11;51(62):12490-3. PMID: 26149530. doi: 10.1039/c5cc03807c.
275. Gopal P, Dick T. The new tuberculosis drug Perchlozone® shows cross-resistance with thiacetazone. *Int J Antimicrob Agents.* 2015 Apr;45(4):430-3. PMID: 25704063. doi: 10.1016/j.ijantimicag.2014.12.026.
276. Harris PN, Mo Y, Jureen R, Chew J, Ali J, Paynter S, et al. Comparable outcomes for β -lactam/ β -lactamase inhibitor combinations and carbapenems in definitive treatment of bloodstream infections caused by cefotaxime-resistant *Escherichia coli* or *Klebsiella pneumoniae*. *Antimicrob Resist Infect Control.* 2015;4:14. PMID: 25932324. doi: 10.1186/s13756-015-0055-6.
277. Harris PN, Tambyah PA, Paterson DL. β -lactam and β -lactamase inhibitor combinations in the treatment of extended-spectrum β -lactamase producing Enterobacteriaceae: time for a reappraisal in

- the era of few antibiotic options? *Lancet Infect Dis.* 2015 Apr;15(4):475-85. PMID: 25716293. doi: 10.1016/s1473-3099(14)70950-8.
278. Harris PN, Le BD, Tambyah PA, Hsu L-Y, Pada SMK, Archuleta S, et al. Antiseptic Body Washes for Reducing the Transmission of Methicillin-Resistant Staphylococcus aureus: A Cluster Crossover Study. *Open Forum Infect Dis.* 2015 Apr;2(2):ofv051. PMID: 26125031. doi: 10.1093/ofid/ofv051.
279. Haver HL, Chua A, Ghode P, Lakshminarayana SB, Singhal A, Mathema B, et al. Mutations in genes for the F420 biosynthetic pathway and a nitroreductase enzyme are the primary resistance determinants in spontaneous in vitro-selected PA-824-resistant mutants of *Mycobacterium tuberculosis*. *Antimicrob Agents Chemother.* 2015 Sep;59(9):5316-23. PMID: 26100695. doi: 10.1128/aac.00308-15.
280. Hsu L-Y, Harris SR, Chlebowicz MA, Lindsay JA, Koh TH, Krishnan PU, et al. Evolutionary dynamics of methicillin-resistant *Staphylococcus aureus* within a healthcare system. *Genome Biol.* 2015 Apr 23;16(1):81. PMID: 25903077. doi: 10.1186/s13059-015-0643-z.
281. Koh J-J, Lin H, Caroline V, Chew YS, Pang LM, Aung TT, et al. N-Lipidated Peptide Dimers: Effective Antibacterial Agents against Gram-Negative Pathogens through Lipopolysaccharide Permeabilization. *J Med Chem.* 2015 Aug 27;58(16):6533-48. PMID: 26214729. doi: 10.1021/acs.jmedchem.5b00628.
282. Koh J-J, Lin S, Aung TT, Lim FH, Zou H, Bai Y, et al. Amino acid modified xanthone derivatives: novel, highly promising membrane-active antimicrobials for multidrug-resistant Gram-positive bacterial infections. *J Med Chem.* 2015 Jan 22;58(2):739-52. PMID: 25474410. doi: 10.1021/jm501285x.
283. Koh TH, Ko KKK, Jureen R, Deepak RN, Tee NWS, Tan TY, et al. High counts of carbapenemase-producing Enterobacteriaceae in hospital sewage. *Infect Control Hosp Epidemiol.* 2015 May;36(5):619-21. PMID: 25732673. doi: 10.1017/ice.2015.44.
284. Kumar A, Ting YP. Presence of *Pseudomonas aeruginosa* influences biofilm formation and surface protein expression of *Staphylococcus aureus*. *Environ Microbiol.* 2015 Nov;17(11):4459-68. PMID: 25925222. doi: 10.1111/1462-2920.12890.
285. Lau QY, Tan YYF, Goh VCY, Lee DJQ, Ng FM, Ong EHQ, et al. An FDA-Drug Library Screen for Compounds with Bioactivities against Meticillin-Resistant *Staphylococcus aureus* (MRSA). *Antibiotics (Basel).* 2015 Oct 9;4(4):424-34. PMID: 27025633. doi: 10.3390/antibiotics4040424.
286. Lau QY, Choo XY, Lim ZX, Kong XN, Ng FM, Ang MJ, et al. A Head-to-Head Comparison of the Antimicrobial Activities of 30 Ultra-Short Antimicrobial Peptides Against *Staphylococcus aureus*, *Pseudomonas aeruginosa* and *Candida albicans*. *International Journal of Peptide Research and Therapeutics.* 2015 2015/03/01;21(1):21-8. doi: 10.1007/s10989-014-9440-x.
287. Lau QY, Ng FM, Cheong JWD, Yap YYA, Tan YYF, Jureen R, et al. Discovery of an ultra-short linear antibacterial tetrapeptide with anti-MRSA activity from a structure-activity relationship study. *Eur J Med Chem.* 2015 Nov 13;105:138-44. PMID: 26489599. doi: 10.1016/j.ejmech.2015.10.015.
288. Lee HK, Loh TP, Hurt AC, Oon LL-E, Tang JW-T, Koay ES-C. Molecular surveillance of antiviral drug resistance of influenza A/H3N2 virus in Singapore, 2009-2013. *PLoS One.* 2015;10(1):e0117822. PMID: 25635767. doi: 10.1371/journal.pone.0117822.
289. Lew KY, Ng TM, Tan MW, Tan SH, Lew EL, Ling LM, et al. Safety and clinical outcomes of carbapenem de-escalation as part of an antimicrobial stewardship programme in an ESBL-endemic setting. *J Antimicrob Chemother.* 2015 Apr;70(4):1219-25. PMID: 25473028. doi: 10.1093/jac/dku479.
290. Li J, Liu S, Koh J-J, Zou H, Lakshminarayanan R, Bai Y, et al. A novel fragment based strategy for membrane active antimicrobials against MRSA. *Biochim Biophys Acta.* 2015 Apr;1848(4):1023-31. PMID: 25582665. doi: 10.1016/j.bbamem.2015.01.001.
291. Liew YX, Lee WHL, Tay D, Tang SSL, Chua NGS, Zhou YP, et al. Prospective audit and feedback in antimicrobial stewardship: is there value in early reviewing within 48 h of antibiotic prescription? *Int J Antimicrob Agents.* 2015 Feb;45(2):168-73. PMID: 25511192. doi: 10.1016/j.ijantimicag.2014.10.018.
292. Liew YX, Lee WHL, Kwa AL-H, Chlebicki MP. Cost effectiveness of an antimicrobial stewardship programme. *Int J Antimicrob Agents.* 2015 Nov;46(5):594-5. PMID: 26493229. doi: 10.1016/j.ijantimicag.2015.08.008.
293. Lim K, Chua RRY, Ho B, Tambyah PA, Hadinoto K, Leong SSJ. Development of a catheter functionalized by a polydopamine peptide coating with antimicrobial and antibiofilm properties. *Acta Biomater.* 2015 Mar;15:127-38. PMID: 25541344. doi: 10.1016/j.actbio.2014.12.015.
294. Lim T-P, Ong RT-H, Hon PY, Hawkey J, Holt KE, Koh TH, et al. Multiple Genetic Mutations Associated with Polymyxin Resistance in *Acinetobacter baumannii*. *Antimicrob Agents Chemother.* 2015 Dec;59(12):7899-902. PMID: 26438500. doi: 10.1128/aac.01884-15.

295. Lim T-P, Cai Y, Hong Y, Chan ECY, Sasikala S, Teo JQM, et al. In vitro pharmacodynamics of various antibiotics in combination against extensively drug-resistant Klebsiella pneumoniae. *Antimicrob Agents Chemother*. 2015 May;59(5):2515-24. PMID: 25691628. doi: 10.1128/aac.03639-14.
296. Ling ML, Tee YM, Tan SG, Amin IM, How KB, Tan KY, et al. Risk factors for acquisition of carbapenem resistant Enterobacteriaceae in an acute tertiary care hospital in Singapore. *Antimicrob Resist Infect Control*. 2015;4:26. PMID: 26106476. doi: 10.1186/s13756-015-0066-3.
297. Loo LW, Liew YX, Lee WHL, Chlebicki MP, Kwa AL-H. Impact of Antimicrobial Stewardship Program (ASP) on Outcomes in Patients with Acute Bacterial Skin and Skin Structure Infections (ABSSSIs) in an Acute-Tertiary Care Hospital. *Infect Dis Ther*. 2015 Sep;4(Suppl 1):15-25. PMID: 26362296. doi: 10.1007/s40121-015-0085-7.
298. Lu S, Liu F, Xing B, Yeow EKL. Nontoxic colloidal particles impede antibiotic resistance of swarming bacteria by disrupting collective motion and speed. *Phys Rev E Stat Nonlin Soft Matter Phys*. 2015 Dec;92(6):062706. PMID: 26764726. doi: 10.1103/PhysRevE.92.062706.
299. Manjunatha UH, Rao SP, Kondreddi RR, Noble CG, Camacho LR, Tan BH, et al. Direct inhibitors of InhA are active against Mycobacterium tuberculosis. *Sci Transl Med*. 2015 Jan 7;7(269):269ra3. PMID: 25568071. doi: 10.1126/scitranslmed.3010597.
300. Ng C, Le T-H, Goh SG, Liang L, Kim Y, Rose JB, et al. A Comparison of Microbial Water Quality and Diversity for Ballast and Tropical Harbor Waters. *PLoS One*. 2015;10(11):e0143123. PMID: 26575481. doi: 10.1371/journal.pone.0143123.
301. Ng LSY, Kwang LL, Rao S, Tan TY. Anaerobic bacteraemia revisited: species and susceptibilities. *Ann Acad Med Singap*. 2015 Jan;44(1):13-8. PMID: 25703492.
302. Ng PS, Manjunatha UH, Rao SP, Camacho LR, Ma NL, Herve M, et al. Structure activity relationships of 4-hydroxy-2-pyridones: A novel class of antituberculosis agents. *Eur J Med Chem*. 2015 Dec 1;106:144-56. PMID: 26544629. doi: 10.1016/j.ejmech.2015.10.008.
303. Ng TM, Teng CB, Lew EL, Ling LM, Ang BSP, Lye DCB. Potential for Cefazolin as De-escalation Therapy for Klebsiella Pneumoniae Bacteraemia. *Ann Acad Med Singap*. 2015 Dec;44(12):571-4. PMID: 27090077.
304. Paton NI, Stöhr W, Arenas-Pinto A, Fisher M, Williams I, Johnson M, et al. Protease inhibitor monotherapy for long-term management of HIV infection: a randomised, controlled, open-label, non-inferiority trial. *Lancet HIV*. 2015 Oct;2(10):e417-26. PMID: 26423649. doi: 10.1016/s2352-3018(15)00176-9.
305. Phoon YW, Chan YYC, Koh TH. Isolation of multidrug-resistant Salmonella in Singapore. *Singapore Med J*. 2015 Aug;56(8):e142-4. PMID: 26311915. doi: 10.11622/smedj.2015129.
306. Samy RP, Thwin MM, Stiles BG, Satyanarayana-Jois SD, Chinnathambi A, Zayed M, et al. Novel phospholipase A2 inhibitors from python serum are potent peptide antibiotics. *Biochimie*. 2015 Apr;111:30-44. PMID: 25583073. doi: 10.1016/j.biochi.2015.01.003.
307. Samy RP, Stiles BG, Chinnathambi A, Zayed M, Alharbi SA, Franco OL, et al. Viperatoxin-II: A novel viper venom protein as an effective bactericidal agent. *FEBS Open Bio*. 2015;5:928-41. PMID: 26793432. doi: 10.1016/j.fob.2015.10.004.
308. Seneviratne CJ, Fong PH, Wong SS, Lee VH. Antifungal susceptibility and phenotypic characterization of oral isolates of a black fungus from a nasopharyngeal carcinoma patient under radiotherapy. *BMC Oral Health*. 2015 Mar 20;15:39. PMID: 25887752. doi: 10.1186/s12903-015-0023-9.
309. Tan S, Gan C, Li R, Ye Y, Zhang S, Wu X, et al. A novel chemosynthetic peptide with β -sheet motif efficiently kills Klebsiella pneumoniae in a mouse model. *International Journal of Nanomedicine*. 2015;10:1045-59. doi: 10.2147/IJN.S73303.
310. Tang SSL, Lye DCB, Jureen R, Sng L-H, Hsu L-Y. Rapidly growing mycobacteria in Singapore, 2006-2011. *Clin Microbiol Infect*. 2015 Mar;21(3):236-41. PMID: 25658536. doi: 10.1016/j.cmi.2014.10.018.
311. Teng CB, Ng TM, Tan MW, Tan SH, Tay M, Lim SF, et al. Safety and effectiveness of improving carbapenem use via prospective review and feedback in a multidisciplinary antimicrobial stewardship programme. *Ann Acad Med Singap*. 2015 Jan;44(1):19-25. PMID: 25703493.
312. Teo JWP, La M-V, Jureen R, Lin RTP. Emergence of a New Delhi metallo- β -lactamase-1-producing *Pseudomonas aeruginosa* in Singapore. *Emerg Microbes Infect*. 2015 Nov 25;4(11):e72. PMID: 26632659. doi: 10.1038/emi.2015.72.
313. Vasoo S, Barreto JN, Tosh PK. Emerging issues in gram-negative bacterial resistance: an update for the practicing clinician. *Mayo Clin Proc*. 2015 Mar;90(3):395-403. PMID: 25744116. doi: 10.1016/j.mayocp.2014.12.002.
314. Vasoo S, Cunningham SA, Cole NC, Kohner PC, Menon SR, Krause KM, et al. In Vitro Activities of Ceftazidime-Avibactam, Aztreonam-Avibactam, and a Panel of Older and Contemporary

- Antimicrobial Agents against Carbapenemase-Producing Gram-Negative Bacilli. *Antimicrob Agents Chemother.* 2015 Dec;59(12):7842-6. PMID: 26392487. doi: 10.1128/aac.02019-15.
315. Vasudevan A, Memon BI, Mukhopadhyay A, Li J, Tambyah PA. The costs of nosocomial resistant gram negative intensive care unit infections among patients with the systemic inflammatory response syndrome- a propensity matched case control study. *Antimicrob Resist Infect Control.* 2015;4(1):3. PMID: 25653851. doi: 10.1186/s13756-015-0045-8.
316. Yi X, Bayen S, Kelly BC, Li X, Zhou Z. Improved detection of multiple environmental antibiotics through an optimized sample extraction strategy in liquid chromatography-mass spectrometry analysis. *Analytical and Bioanalytical Chemistry.* 2015;407(30):9071-83. doi: 10.1007/s00216-015-9074-7.
317. Yoon BK, Jackman JA, Kim MC, Cho N-J. Spectrum of Membrane Morphological Responses to Antibacterial Fatty Acids and Related Surfactants. *Langmuir.* 2015 Sep 22;31(37):10223-32. PMID: 26325618. doi: 10.1021/acs.langmuir.5b02088.
318. Ang TL, Fock KM, Ang D, Kwek ABE, Teo EK, Dhamodaran S. The Changing Profile of *Helicobacter pylori* Antibiotic Resistance in Singapore: A 15-Year Study. *Helicobacter.* 2016 Aug;21(4):261-5. PMID: 26774006. doi: 10.1111/hel.12291.
319. Ariyasu S, Too PC, Mu J, Goh CC, Ding Y, Tnay YL, et al. Glycopeptide antibiotic analogs for selective inactivation and two-photon imaging of vancomycin-resistant strains. *Chem Commun (Camb).* 2016 Mar 28;52(25):4667-70. PMID: 26953360. doi: 10.1039/c5cc10230h.
320. Arora S, Kyaw BM, Lim CSD. Combination antibiotic-phytochemical effects on resistance adaptation in *Staphylococcus aureus*. *African Journal of Microbiology Research.* 2016 12/14;10:1973-82. doi: 10.5897/AJMR11.1650.
321. Ashajyothi C, Harish KH, Dubey N, Chandrakanth RK. Antibiofilm activity of biogenic copper and zinc oxide nanoparticles-antimicrobials collegiate against multiple drug resistant bacteria: a nanoscale approach. *JOURNAL OF NANOSTRUCTURE IN CHEMISTRY.* 2016 DEC;6(4):329-41. PMID: WOS:000388101400005. doi: 10.1007/s40097-016-0205-2.
322. Aung KT, Lo JACY, Chau ML, Kang JSL, Yap HM, Gutiérrez RA, et al. MICROBIOLOGICAL SAFETY ASSESSMENT AND RISK MITIGATION OF INDIAN ROJAK (DEEP FRIED READY-TO-EAT FOOD) IN SINGAPORE. *Southeast Asian J Trop Med Public Health.* 2016 Nov;47(6):1231-45. PMID: 29634190.
323. Aung TT, Yam JKH, Lin S, Salleh SM, Givskov M, Liu S, et al. Biofilms of Pathogenic Nontuberculous Mycobacteria Targeted by New Therapeutic Approaches. *Antimicrob Agents Chemother.* 2016 Jan;60(1):24-35. PMID: 26459903. doi: 10.1128/aac.01509-15.
324. Boudhar A, Ng XW, Loh CY, Chia WN, Tan ZM, Nosten FH, et al. Overcoming Chloroquine Resistance in Malaria: Design, Synthesis, and Structure-Activity Relationships of Novel Hybrid Compounds. *Antimicrob Agents Chemother.* 2016 May;60(5):3076-89. PMID: 26953199. doi: 10.1128/aac.02476-15.
325. Boudhar A, Ng XW, Loh CY, Chia WN, Tan ZM, Nosten FH, et al. Overcoming chloroquine resistance in malaria: Design, synthesis and structure-activity relationships of novel chemoreversal agents. *Eur J Med Chem.* 2016 Aug 25;119:231-49. PMID: 27173385. doi: 10.1016/j.ejmech.2016.04.058.
326. Cai B, Cai Y, Liew YX, Chua NGS, Teo JQM, Lim T-P, et al. Clinical Efficacy of Polymyxin Monotherapy versus Nonvalidated Polymyxin Combination Therapy versus Validated Polymyxin Combination Therapy in Extensively Drug-Resistant Gram-Negative Bacillus Infections. *Antimicrob Agents Chemother.* 2016 Jul;60(7):4013-22. PMID: 27090177. doi: 10.1128/aac.03064-15.
327. Cai Y, Shek PY, Teo I, Tang SSL, Lee WHL, Liew YX, et al. A multidisciplinary antimicrobial stewardship programme safely decreases the duration of broad-spectrum antibiotic prescription in Singaporean adult renal patients. *Int J Antimicrob Agents.* 2016 Jan;47(1):91-6. PMID: 26706421. doi: 10.1016/j.ijantimicag.2015.10.021.
328. Cai Y, Chua NGS, Lim T-P, Teo JQM, Lee WHL, Kurup A, et al. From Bench-Top to Bedside: A Prospective In Vitro Antibiotic Combination Testing (iACT) Service to Guide the Selection of Rationally Optimized Antimicrobial Combinations against Extensively Drug Resistant (XDR) Gram Negative Bacteria (GNB). *PLoS One.* 2016;11(7):e0158740. PMID: 27441603. doi: 10.1371/journal.pone.0158740.
329. Cai Y, Lim T-P, Teo JQM, Sasikala S, Lee WHL, Hong Y, et al. In Vitro Activity of Polymyxin B in Combination with Various Antibiotics against Extensively Drug-Resistant *Enterobacter cloacae* with Decreased Susceptibility to Polymyxin B. *Antimicrob Agents Chemother.* 2016 Sep;60(9):5238-46. PMID: 27324776. doi: 10.1128/aac.00270-16.
330. Cao Y, Nguyen GKT, Chuah S, Tam JP, Liu C-F. Butelase-Mediated Ligation as an Efficient Bioconjugation Method for the Synthesis of Peptide Dendrimers. *Bioconjug Chem.* 2016 Nov 16;27(11):2592-6. PMID: 27723303. doi: 10.1021/acs.bioconjchem.6b00538.

331. Chen YT, Ahmad Murad K, Ng LSY, Seah JTH, Park J-J, Tan TY. In Vitro Efficacy of Six Alternative Antibiotics against Multidrug Resistant Escherichia Coli and Klebsiella Pneumoniae from Urinary Tract Infections. *Ann Acad Med Singap.* 2016 Jun;45(6):245-50. PMID: 27412057.
332. Chen Y-T, Siu LK, Tsai Y-K, Lin F-M, Koh TH, Chen J-H. A Common Flanking Region in Promiscuous Plasmids Encoding blaNDM-1 in Klebsiella pneumoniae Isolated in Singapore. *Microb Drug Resist.* 2016 Mar;22(2):109-14. PMID: 26308279. doi: 10.1089/mdr.2015.0132.
333. Chia G, Kum JQ, Marimuthu K, Poh BF, Ang BSP. Identifying Patients at High Risk for Carbapenem-Resistant Enterobacteriaceae at Admission: Nurse-Led or Doctor-Led? *Infect Control Hosp Epidemiol.* 2016 Feb;37(2):238-9. PMID: 26691540. doi: 10.1017/ice.2015.288.
334. Chow ALP, Win NN, Ng PY, Lee WHL, Win M-K. Vancomycin-resistant enterococci with reduced daptomycin susceptibility in Singapore: prevalence and associated factors. *Epidemiol Infect.* 2016 Sep;144(12):2540-5. PMID: 27174845. doi: 10.1017/s0950268816000923.
335. Chow ALP, Ang A, Chow CZ, Ng TM, Teng CB, Ling LM, et al. Implementation hurdles of an interactive, integrated, point-of-care computerised decision support system for hospital antibiotic prescription. *Int J Antimicrob Agents.* 2016 Feb;47(2):132-9. PMID: 26774157. doi: 10.1016/j.ijantimicag.2015.12.006.
336. Chua SL, Yam JKH, Hao P, Adav SS, Salido MM, Liu Y, et al. Selective labelling and eradication of antibiotic-tolerant bacterial populations in *Pseudomonas aeruginosa* biofilms. *Nat Commun.* 2016 Feb 19;7:10750. PMID: 26892159. doi: 10.1038/ncomms10750.
337. Fisher DA, Pang L, Salmon S, Lin RTP, Teo C, Tambyah PA, et al. A Successful Vancomycin-Resistant Enterococci Reduction Bundle at a Singapore Hospital. *Infect Control Hosp Epidemiol.* 2016 Jan;37(1):107-9. PMID: 26486456. doi: 10.1017/ice.2015.251.
338. Ghode P, Ramachandran S, Bifani P, Sivaraman J. Structure and mapping of spontaneous mutational sites of PyrR from *Mycobacterium tuberculosis*. *Biochem Biophys Res Commun.* 2016 Mar 18;471(4):409-15. PMID: 26902118. doi: 10.1016/j.bbrc.2016.02.071.
339. Gopal P, Yee M, Sarathy J, Low JL, Sarathy JP, Kaya F, et al. Pyrazinamide Resistance Is Caused by Two Distinct Mechanisms: Prevention of Coenzyme A Depletion and Loss of Virulence Factor Synthesis. *ACS Infect Dis.* 2016 Sep 9;2(9):616-26. PMID: 27759369. doi: 10.1021/acsinfecdis.6b00070.
340. Hemu X, Qiu Y, Nguyen GKT, Tam JP. Total Synthesis of Circular Bacteriocins by Butelase 1. *J Am Chem Soc.* 2016 Jun 8;138(22):6968-71. PMID: 27206099. doi: 10.1021/jacs.6b04310.
341. Ho HJ, Toh CY, Ang BSP, Krishnan PU, Lin RTP, La M-V, et al. Outbreak of New Delhi metallo-β-lactamase-1-producing *Enterobacter cloacae* in an acute care hospital general ward in Singapore. *Am J Infect Control.* 2016 Feb;44(2):177-82. PMID: 26454748. doi: 10.1016/j.ajic.2015.08.028.
342. Husain N, Tulsian NK, Chien WL, Suresh S, Anand GS, Sivaraman J. Ligand-mediated changes in conformational dynamics of NpmA: implications for ribosomal interactions. *Sci Rep.* 2016 Nov 15;6:37061. PMID: 27845431. doi: 10.1038/srep37061.
343. Isenman H, Michaels J, Fisher DA. Global variances in infection control practices for vancomycin resistant *Enterococcus* - results of an electronic survey. *Antimicrob Resist Infect Control.* 2016;5:41. PMID: 27822366. doi: 10.1186/s13756-016-0140-5.
344. Isenman H, Fisher DA. Advances in prevention and treatment of vancomycin-resistant *Enterococcus* infection. *Curr Opin Infect Dis.* 2016 Dec;29(6):577-82. PMID: 27584589. doi: 10.1097/qco.0000000000000311.
345. Khara JS, Priestman M, Uhía I, Hamilton MS, Krishnan N, Wang Y, et al. Unnatural amino acid analogues of membrane-active helical peptides with anti-mycobacterial activity and improved stability. *J Antimicrob Chemother.* 2016 Aug;71(8):2181-91. PMID: 27118774. doi: 10.1093/jac/dkw107.
346. Khong WX, Marimuthu K, Teo JWP, Ding Y, Xia E, Lee JJ, et al. Tracking inter-institutional spread of NDM and identification of a novel NDM-positive plasmid, pSg1-NDM, using next-generation sequencing approaches. *J Antimicrob Chemother.* 2016 Nov;71(11):3081-9. PMID: 27494913. doi: 10.1093/jac/dkw277.
347. Khong WX, Xia E, Marimuthu K, Xu W, Teo Y-Y, Tan EL, et al. Local transmission and global dissemination of New Delhi Metallo-Beta-Lactamase (NDM): a whole genome analysis. *BMC Genomics.* 2016 Jun 13;17:452. PMID: 27297071. doi: 10.1186/s12864-016-2740-0.
348. Koh J-J, Zou H, Lin S, Lin H, Soh RT, Lim FH, et al. Nonpeptidic Amphiphilic Xanthone Derivatives: Structure-Activity Relationship and Membrane-Targeting Properties. *Journal of Medicinal Chemistry.* 2016;59(1):171-93. doi: 10.1021/acs.jmedchem.5b01500.
349. Kong YL, Ker KJ, Tan WD, Tey HL. Colonization and acquisition of methicillin-resistant *Staphylococcus aureus* and secondary bacterial infections among dermatological in-patients. *Journal of the European Academy of Dermatology and Venereology.* 2016;30(11):e111-e3.

350. Lakshminarayanan R, Tan WX, Aung TT, Goh ETL, Muruganantham N, Li J, et al. Branched Peptide, B2088, Disrupts the Supramolecular Organization of Lipopolysaccharides and Sensitizes the Gram-negative Bacteria. *Sci Rep.* 2016 May 13;6:25905. PMID: 27174567. doi: 10.1038/srep25905.
351. Le T-H, Ng C, Chen H, Yi X, Koh TH, Barkham TMS, et al. Occurrences and Characterization of Antibiotic-Resistant Bacteria and Genetic Determinants of Hospital Wastewater in a Tropical Country. *Antimicrob Agents Chemother.* 2016 Dec;60(12):7449-56. PMID: 27736769. doi: 10.1128/aac.01556-16.
352. Lee SH, Teo JWP, Heng D, Ng WK, Zhao Y, Tan RBH. Tailored Antibiotic Combination Powders for Inhaled Rotational Antibiotic Therapy. *J Pharm Sci.* 2016 Apr;105(4):1501-12. PMID: 27019964. doi: 10.1016/j.xphs.2016.02.007.
353. Li M, Neoh KG, Xu L, Yuan L, Leong DT, Kang E-T, et al. Sugar-Grafted Cyclodextrin Nanocarrier as a "Trojan Horse" for Potentiating Antibiotic Activity. *Pharm Res.* 2016 May;33(5):1161-74. PMID: 26792570. doi: 10.1007/s11095-016-1861-0.
354. Lim EJZ, Ho SX, Cao DYH, Lau QC, Koh TH, Hsu L-Y. Extended-Spectrum Beta-Lactamase-Producing Enterobacteriaceae in Retail Chicken Meat in Singapore. *Ann Acad Med Singap.* 2016 Dec;45(12):557-9. PMID: 28062884.
355. Lim MY-X, LaMonte G, Lee MCS, Reimer C, Tan BH, Corey V, et al. UDP-galactose and acetyl-CoA transporters as Plasmodium multidrug resistance genes. *Nat Microbiol.* 2016 Sep 19;1:16166. PMID: 27642791. doi: 10.1038/nmicrobiol.2016.166.
356. Lim WS, Phang KKS, Tan AH-M, Li SF-Y, Ow DS-W. Small Colony Variants and Single Nucleotide Variations in Pf1 Region of PB1 Phage-Resistant *Pseudomonas aeruginosa*. *Front Microbiol.* 2016;7:282. PMID: 27014207. doi: 10.3389/fmicb.2016.00282.
357. Low A, Ng C, He J. Identification of antibiotic resistant bacteria community and a GeoChip based study of resistome in urban watersheds. *Water Res.* 2016 Dec 1;106:330-8. PMID: 27750121. doi: 10.1016/j.watres.2016.09.032.
358. Mandakhalikar KD, Chua RRY, Tambyah PA. New Technologies for Prevention of Catheter Associated Urinary Tract Infection. *Current Treatment Options in Infectious Diseases.* 2016 2016/03/01;8(1):24-41. doi: 10.1007/s40506-016-0069-5.
359. Marimuthu K, Ng OT, Khong WX, Xia E, Teo Y-Y, Ong RT-H, et al. Reactive Infection Control Strategy for Control of New Delhi Metallo- β -Lactamase (NDM)-Producing Enterobacteriaceae Analyzed Using Whole-Genome Sequencing: Hits and Misses. *Infect Control Hosp Epidemiol.* 2016 Aug;37(8):987-90. PMID: 27143092. doi: 10.1017/ice.2016.92.
360. Ng TM, Khong WX, Harris PNA, De PP, Chow ALP, Tambyah PA, et al. Empiric Piperacillin-Tazobactam versus Carbapenems in the Treatment of Bacteraemia Due to Extended-Spectrum Beta-Lactamase-Producing Enterobacteriaceae. *PLoS One.* 2016;11(4):e0153696. PMID: 27104951. doi: 10.1371/journal.pone.0153696.
361. Pan DST, Huang JH, Lee MHM, Yu Y, Chen MI-C, Goh EH, et al. Knowledge, attitudes and practices towards antibiotic use in upper respiratory tract infections among patients seeking primary health care in Singapore. *BMC Fam Pract.* 2016 Nov 3;17(1):148. PMID: 27809770. doi: 10.1186/s12875-016-0547-3.
362. Seneviratne CJ, Rajan S, Wong SSW, Tsang DNC, Lai CKC, Samaranayake LP, et al. Antifungal Susceptibility in Serum and Virulence Determinants of *Candida* Bloodstream Isolates from Hong Kong. *FRONTIERS IN MICROBIOLOGY.* 2016 FEB 26;7. PMID: WOS:000370872000001. doi: 10.3389/fmicb.2016.00216.
363. Song CT, Hwee J, Song C, Tan BK, Chong SJ. Burns infection profile of Singapore: prevalence of multidrug-resistant *Acinetobacter baumannii* and the role of blood cultures. *Burns Trauma.* 2016;4:13. PMID: 27574683. doi: 10.1186/s41038-016-0038-8.
364. Su CT-T, Ling W-L, Lua W-H, Haw Y-X, Gan SK-E. Structural analyses of 2015-updated drug-resistant mutations in HIV-1 protease: an implication of protease inhibitor cross-resistance. *BMC Bioinformatics.* 2016 Dec 22;17(Suppl 19):500. PMID: 28155724. doi: 10.1186/s12859-016-1372-3.
365. Teng CP, Zhou T, Ye E, Liu S, Koh LD, Low M, et al. Effective Targeted Photothermal Ablation of Multidrug Resistant Bacteria and Their Biofilms with NIR-Absorbing Gold Nanocrosses. *Adv Healthc Mater.* 2016 Aug;5(16):2122-30. PMID: 27336752. doi: 10.1002/adhm.201600346.
366. Teo JQM, Cai Y, Lim T-P, Tan TT, Kwa AL-H. Carbapenem Resistance in Gram-Negative Bacteria: The Not-So-Little Problem in the Little Red Dot. *Microorganisms.* 2016 Feb 16;4(1). PMID: 27681907. doi: 10.3390/microorganisms4010013.
367. Teo JWP, Chew KL, Lin RTP. Transmissible colistin resistance encoded by mcr-1 detected in clinical Enterobacteriaceae isolates in Singapore. *Emerg Microbes Infect.* 2016 Aug 17;5(8):e87. PMID: 27530747. doi: 10.1038/emi.2016.85.

368. Teo JWP, La M-V, Lin RTP. Development and evaluation of a multiplex real-time PCR for the detection of IMP, VIM, and OXA-23 carbapenemase gene families on the BD MAX open system. *Diagn Microbiol Infect Dis.* 2016 Dec;86(4):358-61. PMID: 27645609. doi: 10.1016/j.diagmicrobio.2016.08.019.
369. Tran NH, Chen H, Do TV, Reinhard M, Ngo HH, He Y, et al. Simultaneous analysis of multiple classes of antimicrobials in environmental water samples using SPE coupled with UHPLC-ESI-MS/MS and isotope dilution. *Talanta.* 2016 Oct 1;159:163-73. PMID: 27474294. doi: 10.1016/j.talanta.2016.06.006.
370. Tran NH, Chen H, Reinhard M, Mao F, Gin KY-H. Occurrence and removal of multiple classes of antibiotics and antimicrobial agents in biological wastewater treatment processes. *Water Res.* 2016 Nov 1;104:461-72. PMID: 27585426. doi: 10.1016/j.watres.2016.08.040.
371. Truong T, Zeng G, Lin Q, Lim TK, Cao T, Chan FY, et al. Comparative Ploidy Proteomics of *Candida albicans* Biofilms Unraveled the Role of the AHP1 Gene in the Biofilm Persistence Against Amphotericin B. *Mol Cell Proteomics.* 2016 Nov;15(11):3488-500. PMID: 27644984. doi: 10.1074/mcp.M116.061523.
372. Tun ZM, Moorthy M, Linster M, Su YCF, Coker RJ, Ooi EE, et al. Patterns of medication use and factors associated with antibiotic use among adult fever patients at Singapore primary care clinics. *Antimicrob Resist Infect Control.* 2016;5:47. PMID: 27904748. doi: 10.1186/s13756-016-0146-z.
373. Wee KB, Lee RTC, Lin J, Pramono ZAD, Maurer-Stroh S. Discovery of Influenza A Virus Sequence Pairs and Their Combinations for Simultaneous Heterosubtypic Targeting that Hedge against Antiviral Resistance. *PLoS Comput Biol.* 2016 Jan;12(1):e1004663. PMID: 26771381. doi: 10.1371/journal.pcbi.1004663.
374. Wong EHH, Khin MM, Ravikumar V, Si Z, Rice SA, Chan-Park MB. Modulating Antimicrobial Activity and Mammalian Cell Biocompatibility with Glucosamine-Functionalized Star Polymers. *Biomacromolecules.* 2016 Mar 14;17(3):1170-8. PMID: 26859230. doi: 10.1021/acs.biomac.5b01766.
375. Wong JGX, Chen MI-C, Win M-K, Ng PY, Chow ALP. Length of stay an important mediator of hospital-acquired methicillin-resistant *Staphylococcus aureus*. *Epidemiol Infect.* 2016 Apr;144(6):1248-56. PMID: 26538070. doi: 10.1017/s0950268815002733.
376. Teo JQM, Ong RT-H, Xia E, Koh TH, Khor C-C, Lee SJ-Y, et al. mcr-1 in Multidrug-Resistant blaKPC-2-Producing Clinical Enterobacteriaceae Isolates in Singapore. *Antimicrob Agents Chemother.* 2016 Oct;60(10):6435-7. PMID: 27503652. doi: 10.1128/aac.00804-16.
377. Zhang J, Zhang L, Loh K-C, Dai Y, Tong YW. Enhanced anaerobic digestion of food waste by adding activated carbon: Fate of bacterial pathogens and antibiotic resistance genes. *Biochemical Engineering Journal.* 2017 2017/12/15/;128:19-25. doi: <https://doi.org/10.1016/j.bej.2017.09.004>.
378. Ang MLT, Zainul Rahim SZ, de Sessions PF, Lin W, Koh V, Pethe K, et al. EthA/R-Independent Killing of *Mycobacterium tuberculosis* by Ethionamide. *Front Microbiol.* 2017;8:710. PMID: 28487681. doi: 10.3389/fmicb.2017.00710.
379. Aung KT, Hsu L-Y, Koh TH, Hapuarachchi HC, Chau ML, Gutiérrez RA, et al. Prevalence of methicillin-resistant *Staphylococcus aureus* (MRSA) in retail food in Singapore. *Antimicrob Resist Infect Control.* 2017;6:94. PMID: 28904790. doi: 10.1186/s13756-017-0255-3.
380. Aw J, Widjaja F, Ding Y, Mu J, Yang L, Xing B. Enzyme-responsive reporter molecules for selective localization and fluorescence imaging of pathogenic biofilms. *Chem Commun (Camb).* 2017 Mar 16;53(23):3330-3. PMID: 28067350. doi: 10.1039/c6cc09296a.
381. Aziz DB, Low JL, Wu M-L, Gengenbacher M, Teo JWP, Dartois V, et al. Rifabutin Is Active against *Mycobacterium abscessus* Complex. *Antimicrob Agents Chemother.* 2017 Jun;61(6):e00155-17. PMID: 28396540. doi: 10.1128/aac.00155-17.
382. Cai Y, Venkatachalam I, Tee NWS, Tan TY, Kurup A, Wong S-Y, et al. Prevalence of Healthcare-Associated Infections and Antimicrobial Use Among Adult Inpatients in Singapore Acute-Care Hospitals: Results From the First National Point Prevalence Survey. *Clin Infect Dis.* 2017 May 15;64(suppl_2):S61-s7. PMID: 28475790. doi: 10.1093/cid/cix103.
383. Cai Y, Lim T-P, Teo JQM, Sasikala S, Chan ECY, Hong Y, et al. Evaluating Polymyxin B-Based Combinations against Carbapenem-Resistant *Escherichia coli* in Time-Kill Studies and in a Hollow-Fiber Infection Model. *Antimicrob Agents Chemother.* 2017 Jan;61(1). PMID: 27795375. doi: 10.1128/aac.01509-16.
384. Chee CBE, KhinMar K-W, Sng L-H, Jureen R, Cutter J, Lee VJM, et al. The shorter multidrug-resistant tuberculosis treatment regimen in Singapore: are patients from South-East Asia eligible? *Eur Respir J.* 2017 Aug;50(2). PMID: 28798092. doi: 10.1183/13993003.00753-2017.
385. Chen S, Larsson M, Robinson RC, Chen SL. Direct and convenient measurement of plasmid stability in lab and clinical isolates of *E. coli*. *Sci Rep.* 2017 Jul 6;7(1):4788. PMID: 28684862. doi: 10.1038/s41598-017-05219-x.

386. Chew KL, Cheng JWS, Jureen R, Lin RTP, Teo JWP. ERG11 mutations are associated with high-level azole resistance in clinical *Candida tropicalis* isolates, a Singapore study. Mycoscience. 2017 2017/03/01;/58(2):111-5. doi: <https://doi.org/10.1016/j.myc.2016.11.001>.
387. Chew KL, Cheng JWS, Hudaa Osman N, Lin RTP, Teo JWP. Predominance of clarithromycin-susceptible *Mycobacterium massiliense* subspecies: Characterization of the *Mycobacterium abscessus* complex at a tertiary acute care hospital. J Med Microbiol. 2017 Oct;66(10):1443-7. PMID: 28874233. doi: 10.1099/jmm.0.000576.
388. Chew KL, Lin RTP, Teo JWP. *Klebsiella pneumoniae* in Singapore: Hypervirulent Infections and the Carbapenemase Threat. Front Cell Infect Microbiol. 2017;7:515. PMID: 29312894. doi: 10.3389/fcimb.2017.00515.
389. Chew KL, La M-V, Lin RTP, Teo JWP. Colistin and Polymyxin B Susceptibility Testing for Carbapenem-Resistant and mcr-Positive Enterobacteriaceae: Comparison of Sensititre, MicroScan, Vitek 2, and Etest with Broth Microdilution. J Clin Microbiol. 2017 Sep;55(9):2609-16. PMID: 28592552. doi: 10.1128/jcm.00268-17.
390. Chin JSF, Sinha S, Nalaparaju A, Yam JKH, Qin Z, Ma L, et al. *Pseudomonas aeruginosa* Psl Exopolysaccharide Interacts with the Antimicrobial Peptide LG21. WATER. 2017 SEP;9(9). PMID: WOS:000411567200054. doi: 10.3390/w9090681.
391. Chong SM, Wong WK, Lee WY, Tan ZB, Tay YH, Teo XH, et al. Streptococcus agalactiae outbreaks in cultured golden pomfret, *Trachinotus blochii* (Lacépède), in Singapore. J Fish Dis. 2017 Jul;40(7):971-4. PMID: 28000924. doi: 10.1111/jfd.12570.
392. Chow ALP, Lim VW, Khan A, Pettigrew KA, Lye DCB, Kanagasabai K, et al. MRSA Transmission Dynamics Among Interconnected Acute, Intermediate-Term, and Long-Term Healthcare Facilities in Singapore. Clin Infect Dis. 2017 May 15;64(suppl_2):S76-s81. PMID: 28475785. doi: 10.1093/cid/cix072.
393. Gopal P, Nartep W, Ragunathan P, Sarathy JP, Kaya F, Yee M, et al. Pyrazinoic Acid Inhibits Mycobacterial Coenzyme A Biosynthesis by Binding to Aspartate Decarboxylase PanD. ACS Infect Dis. 2017 Nov 10;3(11):807-19. PMID: 28991455. doi: 10.1021/acsinfecdis.7b00079.
394. Gopal P, Tasneen R, Yee M, Lanoix J-P, Sarathy JP, Rasic G, et al. In Vivo-Selected Pyrazinoic Acid-Resistant *Mycobacterium tuberculosis* Strains Harbor Missense Mutations in the Aspartate Decarboxylase PanD and the Unfoldase ClpC1. ACS Infect Dis. 2017 Jul 14;3(7):492-501. PMID: 28271875. doi: 10.1021/acsinfecdis.7b00017.
395. Hou Z, Shankar YV, Liu Y, Ding F, Subramanion JL, Ravikumar V, et al. Nanoparticles of Short Cationic Peptidopolysaccharide Self-Assembled by Hydrogen Bonding with Antibacterial Effect against Multidrug-Resistant Bacteria. ACS Appl Mater Interfaces. 2017 Nov 8;9(44):38288-303. PMID: 29028315. doi: 10.1021/acsami.7b12120.
396. Hsu L-Y, Apisarnthanarak A, Khan E, Suwantarat N, Ghafur A, Tambyah PA. Carbapenem-Resistant *Acinetobacter baumannii* and Enterobacteriaceae in South and Southeast Asia. Clin Microbiol Rev. 2017 Jan;30(1):1-22. PMID: 27795305. doi: 10.1128/CMR.masthead.30-1.
397. Hsu L-Y, Holden MTG, Koh TH, Pettigrew KA, Cao DYH, Hon PY, et al. ST3268: a geographically widespread primate MRSA clone. J Antimicrob Chemother. 2017 Aug 1;72(8):2401-3. PMID: 28444294. doi: 10.1093/jac/dkx120.
398. Khara JS, Obuobi S, Wang Y, Hamilton MS, Robertson BD, Newton SM, et al. Disruption of drug-resistant biofilms using de novo designed short α -helical antimicrobial peptides with idealized facial amphiphilicity. Acta Biomater. 2017 Jul 15;57:103-14. PMID: 28457962. doi: 10.1016/j.actbio.2017.04.032.
399. Kityo CM, Thompson JA, Nankya I, Hoppe A, Ndashimye E, Warambwa C, et al. HIV Drug Resistance Mutations in Non-B Subtypes After Prolonged Virological Failure on NNRTI-Based First-Line Regimens in Sub-Saharan Africa. J Acquir Immune Defic Syndr. 2017 Jun 1;75(2):e45-e54. PMID: 28129253. doi: 10.1097/qai.0000000000001285.
400. Lee TH, Wong JGX, Lye DCB, Chen MI-C, Loh VWK, Leo YS, et al. Medical and psychosocial factors associated with antibiotic prescribing in primary care: survey questionnaire and factor analysis. Br J Gen Pract. 2017 Mar;67(656):e168-e177. PMID: 28093423. doi: 10.3399/bjgp17X688885.
401. Li PQ, Piper A, Schmueser I, Mount AR, Corrigan DK. Impedimetric measurement of DNA-DNA hybridisation using microelectrodes with different radii for detection of methicillin resistant *Staphylococcus aureus* (MRSA). Analyst. 2017 May 30;142(11):1946-52. PMID: 28492640. doi: 10.1039/c7an00436b.
402. Lin S, Koh J-J, Aung TT, Lim FH, Li J, Zou H, et al. Symmetrically Substituted Xanthone Amphiphiles Combat Gram-Positive Bacterial Resistance with Enhanced Membrane Selectivity. J Med Chem. 2017 Feb 23;60(4):1362-78. PMID: 28122182. doi: 10.1021/acs.jmedchem.6b01403.

403. Lin S, Koh J-J, Aung TT, Sin WWL, Lim FH, Wang L, et al. Semisynthetic Flavone-Derived Antimicrobials with Therapeutic Potential against Methicillin-Resistant *Staphylococcus aureus* (MRSA). *J Med Chem.* 2017 Jul 27;60(14):6152-65. PMID: 28636355. doi: 10.1021/acs.jmedchem.7b00380.
404. Lin S, Sin WWL, Koh J-J, Lim FH, Wang L, Cao D, et al. Semisynthesis and Biological Evaluation of Xanthone Amphiphilics as Selective, Highly Potent Antifungal Agents to Combat Fungal Resistance. *J Med Chem.* 2017 Dec 28;60(24):10135-50. PMID: 29155590. doi: 10.1021/acs.jmedchem.7b01348.
405. Liu S, Ono RJ, Wu H, Teo JY, Liang ZC, Xu K, et al. Highly potent antimicrobial polyionenes with rapid killing kinetics, skin biocompatibility and in vivo bactericidal activity. *Biomaterials.* 2017 May;127:36-48. PMID: 28279920. doi: 10.1016/j.biomaterials.2017.02.027.
406. Loo LH, Soong HY, Maiwald M, Tee NWS. Assessment of Genotypic Macrolide Resistance among *Mycoplasma pneumoniae* Infections in Children in Singapore. *Ann Acad Med Singap.* 2017 Jul;46(7):290-2. PMID: 28821894.
407. Marimuthu K, Venkatachalam I, Khong WX, Koh TH, Cherng BPZ, La M-V, et al. Clinical and Molecular Epidemiology of Carbapenem-Resistant Enterobacteriaceae Among Adult Inpatients in Singapore. *Clin Infect Dis.* 2017 May 15;64(suppl_2):S68-s75. PMID: 28475792. doi: 10.1093/cid/cix113.
408. Ng C, Tay M, Tan B, Le T-H, Haller L, Chen H, et al. Characterization of Metagenomes in Urban Aquatic Compartments Reveals High Prevalence of Clinically Relevant Antibiotic Resistance Genes in Wastewaters. *Front Microbiol.* 2017;8:2200. PMID: 29201017. doi: 10.3389/fmicb.2017.02200.
409. Ng SMS, Yap YYA, Cheong JWD, Ng FM, Lau QY, Barkham TMS, et al. Antifungal peptides: a potential new class of antifungals for treating vulvovaginal candidiasis caused by fluconazole-resistant *Candida albicans*. *J Pept Sci.* 2017 Mar;23(3):215-21. PMID: 28105725. doi: 10.1002/psc.2970.
410. Ng SMS, Ching HSV, Xu G, Ng FM, Ong EHQ, Lau QY, et al. Screening for a Potent Antibacterial Peptide to Treat Mupirocin-Resistant MRSA Skin Infections. *INTERNATIONAL JOURNAL OF PEPTIDE RESEARCH AND THERAPEUTICS.* 2017 DEC;23(4):481-91. PMID: WOS:000415163300007. doi: 10.1007/s10989-017-9580-x.
411. Ng SMS, Teo SW, Yong YE, Ng FM, Lau QY, Jureen R, et al. Preliminary investigations into developing all-D Omiganan for treating Mupirocin-resistant MRSA skin infections. *Chem Biol Drug Des.* 2017 Dec;90(6):1155-60. PMID: 28581672. doi: 10.1111/cbdd.13035.
412. Paton NI, Kityo CM, Thompson JA, Nankya I, Bagenda L, Hoppe A, et al. Nucleoside reverse-transcriptase inhibitor cross-resistance and outcomes from second-line antiretroviral therapy in the public health approach: an observational analysis within the randomised, open-label, EARNEST trial. *Lancet HIV.* 2017 Aug;4(8):e341-e8. PMID: 28495562. doi: 10.1016/s2352-3018(17)30065-6.
413. Peng J, Cao J, Ng FM, Hill J. *Pseudomonas aeruginosa* develops Ciprofloxacin resistance from low to high level with distinctive proteome changes. *J Proteomics.* 2017 Jan 30;152:75-87. PMID: 27771372. doi: 10.1016/j.jprot.2016.10.005.
414. Pu Y, Du Y, Khin MM, Ravikumar V, Rice SA, Duan H, et al. Using Diphenylphosphoryl Azide (DPPA) for the Facile Synthesis of Biodegradable Antiseptic Random Copolypeptides. *Macromol Rapid Commun.* 2017 Apr;38(7). PMID: 28169482. doi: 10.1002/marc.201600601.
415. Rashid R, Cazenave-Gassiot A, Gao IH, Nair ZJ, Kumar JK, Gao L, et al. Comprehensive analysis of phospholipids and glycolipids in the opportunistic pathogen *Enterococcus faecalis*. *PLoS One.* 2017;12(4):e0175886. PMID: 28423018. doi: 10.1371/journal.pone.0175886.
416. Samy PR, Stiles BG, Franco OL, Sethi G, Lim LHK. Animal venoms as antimicrobial agents. *Biochem Pharmacol.* 2017 Jun 15;134:127-38. PMID: 28288817. doi: 10.1016/j.bcp.2017.03.005.
417. Seneviratne CJ, Suriyanarayanan T, Swarup S, Chia KHB, Nagarajan N, Zhang C. Transcriptomics Analysis Reveals Putative Genes Involved in Biofilm Formation and Biofilm-associated Drug Resistance of *Enterococcus faecalis*. *J Endod.* 2017 Jun;43(6):949-55. PMID: 28457636. doi: 10.1016/j.joen.2017.01.020.
418. Tan JH, Vidaillac C, Yam JKH, Chua SL, Givskov M, Yang L. In Vitro and In Vivo Efficacy of an LpxC Inhibitor, CHIR-090, Alone or Combined with Colistin against *Pseudomonas aeruginosa* Biofilm. *Antimicrob Agents Chemother.* 2017 Jul;61(7). PMID: 28461320. doi: 10.1128/aac.02223-16.
419. Tan JPK, Coady DJ, Sardon H, Yuen A, Gao S, Lim SW, et al. Broad Spectrum Macromolecular Antimicrobials with Biofilm Disruption Capability and In Vivo Efficacy. *Adv Healthc Mater.* 2017 Aug;6(16). PMID: 28504348. doi: 10.1002/adhm.201601420.
420. Tan TY, Ng LSY, Kwang LL, Rao S, Eng LC. Clinical characteristics and antimicrobial susceptibilities of anaerobic bacteremia in an acute care hospital. *Anaerobe.* 2017 Feb;43:69-74. PMID: 27890724. doi: 10.1016/j.anaerobe.2016.11.009.

421. Tan TY, Jiang B, Ng LSY. Faster and economical screening for vancomycin-resistant enterococci by sequential use of chromogenic agar and real-time polymerase chain reaction. *J Microbiol Immunol Infect.* 2017 Aug;50(4):448-53. PMID: 26442675. doi: 10.1016/j.jmii.2015.08.003.
422. Teo JQM, Candra SR, Lee SJ-Y, Chia SY-H, Leck H, Tan AL, et al. Candidemia in a major regional tertiary referral hospital - epidemiology, practice patterns and outcomes. *Antimicrob Resist Infect Control.* 2017;6:27. PMID: 28293420. doi: 10.1186/s13756-017-0184-1.
423. Venkatesh M, Barathi VA, Goh ETL, Anggara R, Fazil MHUT, Ng AJY, et al. Antimicrobial Activity and Cell Selectivity of Synthetic and Biosynthetic Cationic Polymers. *Antimicrob Agents Chemother.* 2017 Oct;61(10). PMID: 28784676. doi: 10.1128/aac.00469-17.
424. Wang A, Zhou K, Liu Y, Yang L, Zhang Q, Guan J, et al. A potential role of transposon IS431 in the loss of *mecA* gene. *Sci Rep.* 2017 Jan 25;7:41237. PMID: 28120911. doi: 10.1038/srep41237.
425. Wang J, Xu C, Lun Z-R, Meshnick SR. Unpacking 'Artemisinin Resistance'. *Trends Pharmacol Sci.* 2017 Jun;38(6):506-11. PMID: 28473165. doi: 10.1016/j.tips.2017.03.007.
426. Wang K, Chen Y-q, Salido MM, Kohli GS, Kong J-I, Liang H, et al. The rapid in vivo evolution of *Pseudomonas aeruginosa* in ventilator-associated pneumonia patients leads to attenuated virulence. *Open Biol.* 2017 Sep;7(9). PMID: 28878043. doi: 10.1098/rsob.170029.
427. Yee M, Gopal P, Dick T. Missense Mutations in the Unfoldase ClpC1 of the Caseinolytic Protease Complex Are Associated with Pyrazinamide Resistance in *Mycobacterium tuberculosis*. *Antimicrob Agents Chemother.* 2017 Feb;61(2). PMID: 27872068. doi: 10.1128/aac.02342-16.
428. Yew PYM, Chee PL, Owh C, Zhang K, Liow SS, Loh XJ. Quarterized Short Polyethylenimine Shows Good Activity against Drug-Resistant Bacteria. *Macromolecular Materials and Engineering.* 2017;302(9). doi: 10.1002/mame.201700186.
429. Yi X, Tran NH, Yin T, He Y, Gin KY-H. Removal of selected PPCPs, EDCs, and antibiotic resistance genes in landfill leachate by a full-scale constructed wetlands system. *Water Res.* 2017 Sep 15;121:46-60. PMID: 28511040. doi: 10.1016/j.watres.2017.05.008.
430. Zhong G, Cheng J, Liang ZC, Xu L, Lou W, Bao C, et al. Short Synthetic β -Sheet Antimicrobial Peptides for the Treatment of Multidrug-Resistant *Pseudomonas aeruginosa* Burn Wound Infections. *Adv Healthc Mater.* 2017 Apr;6(7). PMID: 28135045. doi: 10.1002/adhm.201601134.
431. Zhou C, Wu Y, Thappeta KRV, Subramanian JL, Pranantyo D, Kang E-T, et al. In Vivo Anti-Biofilm and Anti-Bacterial Non-Leachable Coating Thermally Polymerized on Cylindrical Catheter. *ACS Appl Mater Interfaces.* 2017 Oct 18;9(41):36269-80. PMID: 28945343. doi: 10.1021/acsami.7b07053.
432. Aziz DB, Teo JWP, Dartois V, Dick T. Teicoplanin - Tigecycline Combination Shows Synergy Against *Mycobacterium abscessus*. *Front Microbiol.* 2018;9:932. PMID: 29867841. doi: 10.3389/fmicb.2018.00932.
433. Baek J-S, Tan CH, Ng NKJ, Yeo YP, Rice SA, Loo SCJ. A programmable lipid-polymer hybrid nanoparticle system for localized, sustained antibiotic delivery to Gram-positive and Gram-negative bacterial biofilms. *Nanoscale Horiz.* 2018 May 1;3(3):305-11. PMID: 32254078. doi: 10.1039/c7nh00167c.
434. Balne PK, Harini S, Dhand C, Dwivedi N, Chalasani MLS, Verma NK, et al. Surface characteristics and antimicrobial properties of modified catheter surfaces by polypyrogallol and metal ions. *Mater Sci Eng C Mater Biol Appl.* 2018 Sep 1;90:673-84. PMID: 29853139. doi: 10.1016/j.msec.2018.04.095.
435. Budigi Y, Ong EZ, Robinson LN, Ong LC, Rowley KJ, Winnett A, et al. Neutralization of antibody-enhanced dengue infection by VIS513, a pan serotype reactive monoclonal antibody targeting domain III of the dengue E protein. *PLoS Negl Trop Dis.* 2018 Feb;12(2):e0006209. PMID: 29425203. doi: 10.1371/journal.pntd.0006209.
436. Cai Q, Hu J. Effect of UVA/LED/TiO₂) photocatalysis treated sulfamethoxazole and trimethoprim containing wastewater on antibiotic resistance development in sequencing batch reactors. *Water Res.* 2018 Sep 1;140:251-60. PMID: 29723814. doi: 10.1016/j.watres.2018.04.053.
437. Cai Y, Seah CL, Leck H, Lim T-P, Teo JQM, Lee WHL, et al. Rapid Antibiotic Combination Testing for Carbapenem-Resistant Gram-Negative Bacteria within Six Hours Using ATP Bioluminescence. *Antimicrob Agents Chemother.* 2018 Sep;62(9). PMID: 29967021. doi: 10.1128/aac.00183-18.
438. Chan HL, Lyu L, Aw J, Zhang W, Li J, Yang H-h, et al. Unique Fluorescent Imaging Probe for Bacterial Surface Localization and Resistant Enzyme Imaging. *ACS Chem Biol.* 2018 Jul 20;13(7):1890-6. PMID: 29595947. doi: 10.1021/acscchembio.8b00172.
439. Chan LY, Teo JDW, Tan KSW, Sou K, Kwan WL, Lee C-LK. Near Infrared Fluorophore-Tagged Chloroquine in *Plasmodium falciparum* Diagnostic Imaging. *Molecules.* 2018 Oct 14;23(10). PMID: 30322183. doi: 10.3390/molecules23102635.

440. Chan MKL, Koo SH, Quek Q, Pang WS, Jiang B, Ng LSY, et al. Development of a real-time assay to determine the frequency of qac genes in methicillin resistant *Staphylococcus aureus*. *J Microbiol Methods*. 2018 Oct;153:133-8. PMID: 30267717. doi: 10.1016/j.mimet.2018.09.017.
441. Chew KL, Cheng B, Lin RTP, Teo JWP. Elizabethkingia anophelis Is the Dominant Elizabethkingia Species Found in Blood Cultures in Singapore. *J Clin Microbiol*. 2018 Mar;56(3). PMID: 29237782. doi: 10.1128/jcm.01445-17.
442. Chew KL, Tay MKL, Cheng B, Lin RTP, Octavia S, Teo JWP. Aztreonam-Avibactam Combination Restores Susceptibility of Aztreonam in Dual-Carbapenemase-Producing Enterobacteriaceae. *Antimicrob Agents Chemother*. 2018 Aug;62(8). PMID: 29760136. doi: 10.1128/aac.00414-18.
443. Chiang RZ-H, Gan SK-E, Su CT-T. A computational study for rational HIV-1 non-nucleoside reverse transcriptase inhibitor selection and the discovery of novel allosteric pockets for inhibitor design. *Biosci Rep*. 2018 Apr 27;38(2). PMID: 29437904. doi: 10.1042/bsr20171113.
444. Chiew CJ, Ho HJ, Win M-K, Tan A, Lim J-W, Ang BSP, et al. Persistence of meticillin-resistant *Staphylococcus aureus* carriage in re-admitted patients. *J Hosp Infect*. 2018 Nov;100(3):350-4. PMID: 29649554. doi: 10.1016/j.jhin.2018.04.001.
445. Chilambi GS, Gao IH, Yoon BK, Park S, Kawakami LM, Ravikumar V, et al. Membrane adaptation limitations in *Enterococcus faecalis* underlie sensitivity and the inability to develop significant resistance to conjugated oligoelectrolytes. *RSC ADVANCES*. 2018;8(19):10284-93. PMID: WOS:000428572600015. doi: 10.1039/c7ra11823f.
446. Chin W, Zhong G, Pu Q, Yang C, Lou W, de Sessions PF, et al. A macromolecular approach to eradicate multidrug resistant bacterial infections while mitigating drug resistance onset. *Nat Commun*. 2018 Mar 2;9(1):917. PMID: 29500445. doi: 10.1038/s41467-018-03325-6.
447. Chuang L, Ratnayake L. Overcoming challenges of treating extensively drug-resistant *Acinetobacter baumannii* bacteraemic urinary tract infection. *Int J Antimicrob Agents*. 2018 Oct;52(4):521-2. PMID: 30081140. doi: 10.1016/j.ijantimicag.2018.07.016.
448. Ding Y, Teo JWP, Drautz-Moses DI, Schuster SC, Givskov M, Yang L. Acquisition of resistance to carbapenem and macrolide-mediated quorum sensing inhibition by *Pseudomonas aeruginosa* via ICE(Tn4371) 6385. *Commun Biol*. 2018;1:57. PMID: 30271939. doi: 10.1038/s42003-018-0064-0.
449. Ding Y, Zwe YH, Chin JSF, Kohli GS, Drautz-Moses DI, Givskov M, et al. Characterization of a novel multidrug resistance plasmid pSGB23 isolated from *Salmonella enterica* subspecies enterica serovar Saintpaul. *Gut Pathog*. 2018;10:20. PMID: 29881467. doi: 10.1186/s13099-018-0249-6.
450. Gao J, Wang H, Li Z, Wong AH-H, Wang Y-Z, Guo Y, et al. *Candida albicans* gains azole resistance by altering sphingolipid composition. *Nat Commun*. 2018 Oct 29;9(1):4495. PMID: 30374049. doi: 10.1038/s41467-018-06944-1.
451. Haller L, Chen H, Ng C, Le T-H, Koh TH, Barkham TMS, et al. Occurrence and characteristics of extended-spectrum β-lactamase- and carbapenemase-producing bacteria from hospital effluents in Singapore. *Sci Total Environ*. 2018 Feb 15;615:1119-25. PMID: 29751417. doi: 10.1016/j.scitotenv.2017.09.217.
452. Hartantyo SHP, Chau ML, Fillon L, Mohamad Ariff AZB, Kang JSL, Aung KT, et al. Sick pets as potential reservoirs of antibiotic-resistant bacteria in Singapore. *Antimicrob Resist Infect Control*. 2018;7:106. PMID: 30186596. doi: 10.1186/s13756-018-0399-9.
453. Heng ST, Chen SL, Wong JGX, Lye DCB, Ng TM. No association between resistance mutations, empiric antibiotic, and mortality in ceftriaxone-resistant *Escherichia coli* and *Klebsiella pneumoniae* bacteremia. *Sci Rep*. 2018 Aug 24;8(1):12785. PMID: 30143706. doi: 10.1038/s41598-018-31081-6.
454. Ho ZJM, Chee CBE, Ong RT-H, Sng L-H, Peh WLJ, Cook AR, et al. Investigation of a cluster of multi-drug resistant tuberculosis in a high-rise apartment block in Singapore. *Int J Infect Dis*. 2018 Feb;67:46-51. PMID: 29253709. doi: 10.1016/j.ijid.2017.12.010.
455. Htun HL, Win M-K, de Sessions PF, Low L, Hibberd ML, Chow ALP, et al. Methicillin-resistant *Staphylococcus aureus* colonisation: epidemiological and molecular characteristics in an acute-care tertiary hospital in Singapore. *Epidemiol Infect*. 2018 Oct;146(14):1785-92. PMID: 30019657. doi: 10.1017/s0950268818001966.
456. Hu B, Owh C, Chee PL, Leow WR, Liu X, Wu Y-L, et al. Supramolecular hydrogels for antimicrobial therapy. *Chem Soc Rev*. 2018 Sep 17;47(18):6917-29. PMID: 29697128. doi: 10.1039/c8cs00128f.
457. Kalimuddin S, Chan YFZ, Phillips R, Ong SP, Archuleta S, Lye DCB, et al. A randomized phase 2B trial of vancomycin versus daptomycin for the treatment of methicillin-resistant *Staphylococcus aureus* bacteremia due to isolates with high vancomycin minimum inhibitory concentrations - results of

- a prematurely terminated study. *Trials*. 2018 Jun 1;19(1):305. PMID: 29859132. doi: 10.1186/s13063-018-2702-8.
458. Koh J-J, Lin S, Sin WWL, Ng ZH, Jung DY, Beuerman RW, et al. Design and synthesis of oligolipidated arginyl peptide (OLAP) dimers with enhanced physicochemical activity, peptide stability and their antimicrobial actions against MRSA infections. *Amino Acids*. 2018 Oct;50(10):1329-45. PMID: 30066172. doi: 10.1007/s00726-018-2607-6.
459. Koh TH, Abdul Rahman NB, Teo JWP, La M-V, Periaswamy B, Chen SL. Putative Integrative Mobile Elements That Exploit the Xer Recombination Machinery Carrying bla(IMI)-Type Carbapenemase Genes in *Enterobacter cloacae* Complex Isolates in Singapore. *Antimicrob Agents Chemother*. 2018 Jan;62(1). PMID: 29038281. doi: 10.1128/aac.01542-17.
460. Lakshminarayanan R, Ye E, Young DJ, Li Z, Loh XJ. Recent Advances in the Development of Antimicrobial Nanoparticles for Combating Resistant Pathogens. *Adv Healthc Mater*. 2018 Jul;7(13):e1701400. PMID: 29717819. doi: 10.1002/adhm.201701400.
461. Lau QY, Li J, Sani M-A, Sinha S, Li Y, Ng FM, et al. Elucidating the bactericidal mechanism of action of the linear antimicrobial tetrapeptide BRBR-NH(2). *Biochim Biophys Acta Biomembr*. 2018 Aug;1860(8):1517-27. PMID: 29758185. doi: 10.1016/j.bbamem.2018.05.004.
462. Le T-H, Ng C, Tran NH, Chen H, Gin KY-H. Removal of antibiotic residues, antibiotic resistant bacteria and antibiotic resistance genes in municipal wastewater by membrane bioreactor systems. *Water Res*. 2018 Nov 15;145:498-508. PMID: 30193193. doi: 10.1016/j.watres.2018.08.060.
463. Lee GH, Inoue M, Chong RH-H, Toh JK-P, Wee SY, Loh K-S, et al. Pyrosequencing method for sensitive detection of HBV drug resistance mutations. *J Med Virol*. 2018 Jun;90(6):1071-9. PMID: 29488627. doi: 10.1002/jmv.25066.
464. Li J, Zhang K, Ruan L, Chin JSF, Wickramasinghe N, Liu H, et al. Block Copolymer Nanoparticles Remove Biofilms of Drug-Resistant Gram-Positive Bacteria by Nanoscale Bacterial Debridement. *Nano Lett*. 2018 Jul 11;18(7):4180-7. PMID: 29902011. doi: 10.1021/acs.nanolett.8b01000.
465. Lim CLL, Chua AQ, Teo JQM, Cai Y, Lee WHL, Kwa AL-H. Importance of control groups when delineating antibiotic use as a risk factor for carbapenem resistance, extreme-drug resistance, and pan-drug resistance in *Acinetobacter baumannii* and *Pseudomonas aeruginosa*: A systematic review and meta-analysis. *Int J Infect Dis*. 2018 Nov;76:48-57. PMID: 29870795. doi: 10.1016/j.ijid.2018.05.017.
466. Lim T-P, Wang R, Poh GQ, Koh TH, Tan TY, Lee WHL, et al. Integrated pharmacokinetic-pharmacodynamic modeling to evaluate empiric carbapenem therapy in bloodstream infections. *Infect Drug Resist*. 2018;11:1591-6. PMID: 30310294. doi: 10.2147/idr.S168191.
467. Lim YH, Wong FT, Yeo WL, Ching KC, Lim YW, Heng E, et al. Auroramycin: A Potent Antibiotic from *Streptomyces roseosporus* by CRISPR-Cas9 Activation. *Chembiochem*. 2018 May 25. PMID: 29799651. doi: 10.1002/cbic.201800266.
468. Lou W, Venkataraman S, Zhong G, Ding B, Tan JPK, Xu L, et al. Antimicrobial polymers as therapeutics for treatment of multidrug-resistant *Klebsiella pneumoniae* lung infection. *Acta Biomater*. 2018 Sep 15;78:78-88. PMID: 30031912. doi: 10.1016/j.actbio.2018.07.038.
469. Mendez AR, Tan TY, Low HY, Otto KH, Tan H, Khoo X. Micro-textured films for reducing microbial colonization in a clinical setting. *J Hosp Infect*. 2018 Jan;98(1):83-9. PMID: 28797757. doi: 10.1016/j.jhin.2017.08.001.
470. Ng C, Chen H, Goh SG, Haller L, Wu Z, Charles FR, et al. Microbial water quality and the detection of multidrug resistant *E. coli* and antibiotic resistance genes in aquaculture sites of Singapore. *Mar Pollut Bull*. 2018 Oct;135:475-80. PMID: 30301062. doi: 10.1016/j.marpolbul.2018.07.055.
471. Ng C, Goh SG, Saeidi N, Gerhard WA, Gunsch CK, Gin KY-H. Occurrence of *Vibrio* species, beta-lactam resistant *Vibrio* species, and indicator bacteria in ballast and port waters of a tropical harbor. *Sci Total Environ*. 2018 Jan 1;610-611:651-6. PMID: 28822933. doi: 10.1016/j.scitotenv.2017.08.099.
472. Ng C, Gu X, Goh SG, Chen H, Haller L, Tan B, et al. Draft Genome Sequences of Four Multidrug-Resistant *Pseudomonas aeruginosa* Isolates from Hospital Wastewater in Singapore. *Microbiol Resour Announc*. 2018 Nov;7(19). PMID: 30533798. doi: 10.1128/mra.01193-18.
473. Ng SMS, Sioson JSP, Yap JM, Ng FM, Ching HSV, Teo JWP, et al. Repurposing Zidovudine in combination with Tigecycline for treating carbapenem-resistant Enterobacteriaceae infections. *Eur J Clin Microbiol Infect Dis*. 2018 Jan;37(1):141-8. PMID: 29019016. doi: 10.1007/s10096-017-3114-5.
474. Ng SMS, Yap JM, Lau QY, Ng FM, Ong EHQ, Barkham TMS, et al. Structure-activity relationship studies of ultra-short peptides with potent activities against fluconazole-resistant *Candida albicans*. *Eur J Med Chem*. 2018 Apr 25;150:479-90. PMID: 29549835. doi: 10.1016/j.ejmech.2018.03.027.

475. Obuobi S, Wang Y, Khara JS, Riegger A, Kuan SL, Ee PLR. Antimicrobial and Anti-Biofilm Activities of Surface Engineered Polycationic Albumin Nanoparticles with Reduced Hemolytic Activity. *Macromol Biosci.* 2018 Oct;18(10):e1800196. PMID: 30066983. doi: 10.1002/mabi.201800196.
476. Ong CH, Ratnayake L, Ang MLT, Lin RTP, Chan DSG. Diagnostic Accuracy of BD Phoenix CPO Detect for Carbapenemase Production in 190 Enterobacteriaceae Isolates. *J Clin Microbiol.* 2018 Dec;56(12). PMID: 30209186. doi: 10.1128/jcm.01043-18.
477. Parmar A, Lakshminarayanan R, Iyer A, Mayandi V, Goh ETL, Lloyd DG, et al. Design and Syntheses of Highly Potent Teixobactin Analogues against *Staphylococcus aureus*, Methicillin-Resistant *Staphylococcus aureus* (MRSA), and Vancomycin-Resistant Enterococci (VRE) in Vitro and in Vivo. *J Med Chem.* 2018 Mar 8;61(5):2009-17. PMID: 29363971. doi: 10.1021/acs.jmedchem.7b01634.
478. Rocamora FM, Zhu L, Liang K-Y, Dondorp AM, Miotto O, Mok S, et al. Oxidative stress and protein damage responses mediate artemisinin resistance in malaria parasites. *PLoS Pathog.* 2018 Mar;14(3):e1006930. PMID: 29538461. doi: 10.1371/journal.ppat.1006930.
479. Selcuk A, Teng CB, Chan SY, Yap KZ. Antimicrobial use and drug-drug interactions among nursing home residents in Singapore: a multicentre prevalence study. *Int J Clin Pharm.* 2018 Oct;40(5):1044-50. PMID: 30054787. doi: 10.1007/s11096-018-0683-z.
480. Sinha S, Harioudh MK, Dewangan RP, Ng WJ, Ghosh JK, Bhattacharjya S. Cell-Selective Pore Forming Antimicrobial Peptides of the Prodomain of Human Furin: A Conserved Aromatic/Cationic Sequence Mapping, Membrane Disruption, and Atomic-Resolution Structure and Dynamics. *ACS Omega.* 2018 Nov 30;3(11):14650-64. PMID: 30555984. doi: 10.1021/acsomega.8b01876.
481. Su CT-T, Kwoh C-K, Verma CS, Gan SK-E. Modeling the full length HIV-1 Gag polyprotein reveals the role of its p6 subunit in viral maturation and the effect of non-cleavage site mutations in protease drug resistance. *J Biomol Struct Dyn.* 2018 Dec;36(16):4366-77. PMID: 29237328. doi: 10.1080/07391102.2017.1417160.
482. Su W, Kumar V, Ding Y, Ero R, Serra A, Lee BST, et al. Ribosome protection by antibiotic resistance ATP-binding cassette protein. *Proc Natl Acad Sci U S A.* 2018 May 15;115(20):5157-62. PMID: 29712846. doi: 10.1073/pnas.1803313115.
483. Tan BH, De Guzman MRT, Donato LKS, Kalimuddin S, Lee WHL, Tan AL, et al. Impact of an alternating first-line antibiotics strategy in febrile neutropenia. *PLoS One.* 2018;13(11):e0208039. PMID: 30485357. doi: 10.1371/journal.pone.0208039.
484. Tan D, Htun HL, Koh J, Kanagasabai K, Lim J-W, Hon PY, et al. Comparative Epidemiology of Vancomycin-Resistant Enterococci Colonization in an Acute-Care Hospital and Its Affiliated Intermediate- and Long-Term Care Facilities in Singapore. *Antimicrob Agents Chemother.* 2018 Dec;62(12). PMID: 30224534. doi: 10.1128/aac.01507-18.
485. Tan YE, Tan AL. Arrival of *Candida auris* Fungus in Singapore: Report of the First 3 Cases. *Ann Acad Med Singap.* 2018 Jul;47(7):260-2. PMID: 30120434.
486. Tang YW, Cheng B, Yeoh SF, Lin RTP, Teo JWP. Tedizolid Activity Against Clinical *Mycobacterium abscessus* Complex Isolates-An in vitro Characterization Study. *Front Microbiol.* 2018;9:2095. PMID: 30245674. doi: 10.3389/fmicb.2018.02095.
487. Teo JWP, Marimuthu K, Venkatachalam I, Ng OT, Lin RTP, Octavia S. mcr-3 and mcr-4 Variants in Carbapenemase-Producing Clinical Enterobacteriaceae Do Not Confer Phenotypic Polymyxin Resistance. *J Clin Microbiol.* 2018 Mar;56(3). PMID: 29237785. doi: 10.1128/jcm.01562-17.
488. Teo JWP, Octavia S, Cheng JWS, Lin RTP. Evaluation of an in-house developed multiplex real-time PCR for the detection of IMP, OXA-23, GES carbapenemases and the transmissible colistin-resistant mcr gene on the BD MAX™ open system. *Diagn Microbiol Infect Dis.* 2018 Jan;90(1):67-9. PMID: 29132936. doi: 10.1016/j.diagmicrobio.2017.09.010.
489. Tong JX, Chandramohanadas R, Tan KSW. High-Content Screening of the Medicines for Malaria Venture Pathogen Box for *Plasmodium falciparum* Digestive Vacuole-Disrupting Molecules Reveals Valuable Starting Points for Drug Discovery. *Antimicrob Agents Chemother.* 2018 Mar;62(3). PMID: 29311064. doi: 10.1128/aac.02031-17.
490. Xu HV, Zheng XT, Wang C, Zhao Y, Tan YN. Bioinspired Antimicrobial Nanodots with Amphiphilic and Zwitterionic-like Characteristics for Combating Multidrug-Resistant Bacteria and Biofilm Removal. *ACS Applied Nano Materials.* 2018;1(5):2062-8. doi: 10.1021/acsanm.8b00465.
491. Yang SS, Long V, Liau MM, Lee SH, Toh M, Teo JWP, et al. A profile of *Propionibacterium acnes* resistance and sensitivity at a tertiary dermatological centre in Singapore. *Br J Dermatol.* 2018 Jul;179(1):200-1. PMID: 29368337. doi: 10.1111/bjd.16380.
492. Yason JA, Koh KARP, Tan KSW. Viability Screen of LOPAC(1280) Reveals Phosphorylation Inhibitor Auranofin as a Potent Inhibitor of *Blastocystis* Subtype 1, 4, and 7 Isolates. *Antimicrob Agents Chemother.* 2018 Aug;62(8). PMID: 29866860. doi: 10.1128/aac.00208-18.

493. Yeo CK, Vikhe YS, Li P, Guo Z, Greenberg P, Duan H, et al. Hydrogel Effects Rapid Biofilm Debridement with ex situ Contact-Kill to Eliminate Multidrug Resistant Bacteria in vivo. *ACS Appl Mater Interfaces*. 2018 Jun 20;10(24):20356-67. PMID: 29806938. doi: 10.1021/acsami.8b06262.
494. Yuan W, Yuk H-G. Antimicrobial efficacy of *Syzygium antisepticum* plant extract against *Staphylococcus aureus* and methicillin-resistant *S. aureus* and its application potential with cooked chicken. *Food Microbiol*. 2018 Jun;72:176-84. PMID: 29407395. doi: 10.1016/j.fm.2017.12.002.
495. Zhang Z, Han X, Wang Z, Yang Z, Zhang W, Li J, et al. A live bacteria SERS platform for the: In situ monitoring of nitric oxide release from a single MRSA. *Chemical Communications*. 2018;54(51):7022-5. doi: 10.1039/c8cc02855a.
496. Zhu L, Tripathi J, Rocamora FM, Miotto O, van der Pluijm R, Voss TS, et al. The origins of malaria artemisinin resistance defined by a genetic and transcriptomic background. *Nat Commun*. 2018 Dec 4;9(1):5158. PMID: 30514877. doi: 10.1038/s41467-018-07588-x.
497. Zwe YH, Tang VCY, Aung KT, Gutiérrez RA, Ng LC, Yuk H-G. Prevalence, sequence types, antibiotic resistance and, *gyrA* mutations of *Salmonella* isolated from retail fresh chicken meat in Singapore. *Food Control*. 2018 2018/08/01;90:233-40. doi: <https://doi.org/10.1016/j.foodcont.2018.03.004>.
498. Arfan G, Ong CYF, Ng SMS, Lau QY, Ng FM, Ong EHQ, et al. Designing an ultra-short antibacterial peptide with potent activity against Mupirocin-resistant MRSA. *Chem Biol Drug Des*. 2019 Jan;93(1):4-11. PMID: 30103288. doi: 10.1111/cbdd.13377.
499. Aung KT, Chen H-J, Chau ML, Yap G, Lim XF, Humaidi M, et al. *Salmonella* in Retail Food and Wild Birds in Singapore-Prevalence, Antimicrobial Resistance, and Sequence Types. *Int J Environ Res Public Health*. 2019 Oct 31;16(21). PMID: 31683716. doi: 10.3390/ijerph16214235.
500. Bae K, Zheng WS-h, Ma Y, Huang Z. Real-Time Monitoring of Pharmacokinetics of Antibiotics in Biofilms with Raman-Tagged Hyperspectral Stimulated Raman Scattering Microscopy. *Theranostics*. 2019;9(5):1348-57. PMID: 30867835. doi: 10.7150/thno.32043.
501. Bazan EL, Ruan L, Zhou C. Improving the antimicrobial efficacy against resistant *Staphylococcus aureus* by a combined use of conjugated oligoelectrolytes. *PLoS One*. 2019;14(11):e0224816. PMID: 31730663. doi: 10.1371/journal.pone.0224816.
502. Bharadwaj S, Ho SK, Khong KC, Seet A, Yeo KC, Chan XY, et al. Eliminating MRSA transmission in a tertiary neonatal unit-A quality improvement initiative. *Am J Infect Control*. 2019 Nov;47(11):1329-35. PMID: 31326262. doi: 10.1016/j.ajic.2019.06.001.
503. Chan JC, Chong CY, Thoon KC, Tee NWS, Maiwald M, Lam JCM, et al. Invasive paediatric *Elizabethkingia meningoseptica* infections are best treated with a combination of piperacillin/tazobactam and trimethoprim/sulfamethoxazole or fluoroquinolone. *J Med Microbiol*. 2019 Aug;68(8):1167-72. PMID: 31199227. doi: 10.1099/jmm.0.001021.
504. Chan YY, Bin Ibrahim MA, Wong CM, Ooi CK, Chow ALP. Determinants of antibiotic prescribing for upper respiratory tract infections in an emergency department with good primary care access: a qualitative analysis. *EPIDEMIOLOGY AND INFECTION*. 2019;147. PMID: WOS:000475912200042. doi: 10.1017/S095026881800331X.
505. Chen H, Gu X, Ng C, Haller L, Charles FR, Gin KY-H. Draft Genome Sequences of a Ceftazidime-Resistant *Acinetobacter baumannii* Donor and a Conjugal *Escherichia coli* Recipient with Acquired Resistance. *Microbiol Resour Announc*. 2019 Mar 28;8(13):e00024-19. PMID: 30923237. doi: 10.1128/mra.00024-19.
506. Chen WK, Yang Y, Tan BH. Increased mortality among carbapenemase-producing carbapenem-resistant Enterobacteriaceae carriers who developed clinical isolates of another genotype. *Open Forum Infectious Diseases*. 2019;6(2). doi: 10.1093/ofid/ofz006.
507. Chew KL, Octavia S, Ng OT, Marimuthu K, Venkatachalam I, Cheng B, et al. Challenge of drug resistance in *Pseudomonas aeruginosa*: clonal spread of NDM-1-positive ST-308 within a tertiary hospital. *J Antimicrob Chemother*. 2019 Aug 1;74(8):2220-4. PMID: 31081022. doi: 10.1093/jac/dkz169.
508. Chew KL, Octavia S, Lin RTP, Yan GZ, Teo JWP. Delay in effective therapy in anidulafungin-resistant *Candida tropicalis* fungaemia: Potential for rapid prediction of antifungal resistance with whole-genome-sequencing. *J Glob Antimicrob Resist*. 2019 Mar;16:105-7. PMID: 30583013. doi: 10.1016/j.jgar.2018.12.010.
509. Chiu JKH, Ong RT-H. ARGDT: a validation and integration toolkit for Antimicrobial Resistance Gene Databases. *BIOINFORMATICS*. 2019 JUL 15;35(14):2466-74. PMID: WOS:000477703600086. doi: 10.1093/bioinformatics/bty987.
510. Choudhury S, Lee KM, Ng ECX, Lee SJ, Vasoo S, Wickramasinghe SS, et al. Evaluation of in vitro susceptibility of Gram-positive pathogens from a tertiary care hospital in Singapore to a novel oxazolidinone, tedizolid, by a gradient diffusion method and broth microdilution. *J Clin Pathol*. 2019 Feb;72(2):181-4. PMID: 30446512. doi: 10.1136/jclinpath-2018-205473.

511. Chua AQ, Kwa AL-H, Tan TY, Legido-Quigley H, Hsu L-Y. Ten-year narrative review on antimicrobial resistance in Singapore. *Singapore Med J*. 2019 Aug;60(8):387-96. PMID: 31482178. doi: 10.11622/smedj.2019088.
512. Dupont C, Chen Y, Xu Z, Roquet-Banères F, Blaise M, Witt A-K, et al. A piperidinol-containing molecule is active against *Mycobacterium tuberculosis* by inhibiting the mycolic acid flippase activity of MmpL3. *J Biol Chem*. 2019 Nov 15;294(46):17512-23. PMID: 31562241. doi: 10.1074/jbc.RA119.010135.
513. Ero R, Kumar V, Su W, Gao Y-G. Ribosome protection by ABC-F proteins-Molecular mechanism and potential drug design. *Protein Sci*. 2019 Apr;28(4):684-93. PMID: 30746819. doi: 10.1002/pro.3589.
514. Fong J, Mortensen KT, Nørskov A, Qvortrup K, Yang L, Tan CH, et al. Itaconimides as Novel Quorum Sensing Inhibitors of *Pseudomonas aeruginosa*. *Front Cell Infect Microbiol*. 2019;8:443. PMID: 30666301. doi: 10.3389/fcimb.2018.00443.
515. Guo S, Tay MYF, Aung KT, Seow KLG, Ng LC, Purbojati RW, et al. Phenotypic and genotypic characterization of antimicrobial resistant *Escherichia coli* isolated from ready-to-eat food in Singapore using disk diffusion, broth microdilution and whole genome sequencing methods. *Food Control*. 2019 2019/05/01;99:89-97. doi: <https://doi.org/10.1016/j.foodcont.2018.12.043>.
516. Guo S, Tay MYF, Aung KT, Seow KLG, Zhong Y, Ng LC, et al. Conjugative IncX1 Plasmid Harboring Colistin Resistance Gene mcr-5.1 in *Escherichia coli* Isolated from Chicken Rice Retailed in Singapore. *Antimicrob Agents Chemother*. 2019 Nov;63(11). PMID: 31481439. doi: 10.1128/aac.01043-19.
517. Guo S, Aung KT, Tay MYF, Seow KLG, Ng LC, Schlundt J. Extended-spectrum β-lactamase-producing *Proteus mirabilis* with multidrug resistance isolated from raw chicken in Singapore: Genotypic and phenotypic analysis. *J Glob Antimicrob Resist*. 2019 Dec;19:252-4. PMID: 31639544. doi: 10.1016/j.jgar.2019.10.013.
518. Ho HJ, Tan MX, Chen MI-C, Tan TY, Koo SH, Koong AYL, et al. Interaction between Antibiotic Resistance, Resistance Genes, and Treatment Response for Urinary Tract Infections in Primary Care. *J Clin Microbiol*. 2019 Sep;57(9). PMID: 31243084. doi: 10.1128/jcm.00143-19.
519. Ho PL, Ong HK, Teo JWP, Ow DS-W, Chao S-H. HEXIM1 Peptide Exhibits Antimicrobial Activity Against Antibiotic Resistant Bacteria Through Guidance of Cell Penetrating Peptide. *Front Microbiol*. 2019;10:203. PMID: 30800117. doi: 10.3389/fmicb.2019.00203.
520. Htun HL, Hon PY, Holden MTG, Ang BSP, Chow ALP. Chlorhexidine and octenidine use, carriage of qac genes, and reduced antiseptic susceptibility in methicillin-resistant *Staphylococcus aureus* isolates from a healthcare network. *Clin Microbiol Infect*. 2019 Sep;25(9):1154.e1-e7. PMID: 30625411. doi: 10.1016/j.cmi.2018.12.036.
521. Keerthisinghe TP, Nguyen LN, Kwon EE, Oh S. Antiseptic chlorhexidine in activated sludge: Biosorption, antimicrobial susceptibility, and alteration of community structure. *J Environ Manage*. 2019 May 1;237:629-35. PMID: 30851591. doi: 10.1016/j.jenvman.2019.02.043.
522. Ko KKK, Chio MTW, Goh SS, Tan AL, Koh TH, Abdul Rahman NB. First Case of Ceftriaxone-Resistant Multidrug-Resistant *Neisseria gonorrhoeae* in Singapore. *Antimicrob Agents Chemother*. 2019 May;63(5). PMID: 30858209. doi: 10.1128/aac.02624-18.
523. La M-V, Lee B, Hong BZM, Yah JY, Koo SH, Jiang B, et al. Prevalence and antibiotic susceptibility of colistin-resistance gene (mcr-1) positive Enterobacteriaceae in stool specimens of patients attending a tertiary care hospital in Singapore. *Int J Infect Dis*. 2019 Aug;85:124-6. PMID: 31153981. doi: 10.1016/j.ijid.2019.05.029.
524. Lee SY, Octavia S, Chew KL. Detection of OXA-carbapenemase-producing Enterobacteriaceae with chromID CARBA SMART screening plate. *Pathology*. 2019 Jan;51(1):108-10. PMID: 30454880. doi: 10.1016/j.pathol.2018.08.017.
525. Legido-Quigley H, Khan MS, Durrance-Bagale A, Hanefeld J. Something Borrowed, Something New: A Governance and Social Construction Framework to Investigate Power Relations and Responses of Diverse Stakeholders to Policies Addressing Antimicrobial Resistance. *Antibiotics (Basel)*. 2019 Dec 24;8(1). PMID: 30586853. doi: 10.3390/antibiotics8010003.
526. Leung CM, Dhand C, Dwivedi N, Xiao A, Ong ST, Chalasani MLS, et al. Combating Microbial Contamination with Robust Polymeric Nanofibers: Elemental Effect on the Mussel-Inspired Cross-Linking of Electrospun Gelatin. *ACS APPLIED BIO MATERIALS*. 2019 FEB 18;2(2):807-23. PMID: WOS:000616368100022. doi: 10.1021/acsabm.8b00666.
527. Li D, Wong CH, Seet MF, Kuan N. Isolation, Characterization, and Inactivation of *Stenotrophomonas maltophilia* From Leafy Green Vegetables and Urban Agriculture Systems. *Front Microbiol*. 2019;10:2718. PMID: 31849874. doi: 10.3389/fmicb.2019.02718.

528. Li M, Kang E-T, Chua KL, Neoh KG. Sugar-powered nanoantimicrobials for combating bacterial biofilms. *BIOMATERIALS SCIENCE*. 2019 JUL 1;7(7):2961-74. PMID: WOS:000474601400024. doi: 10.1039/c9bm00471h.
529. Liu H, Sun H, Zhang M, Liu Y. Dynamics of microbial community and tetracycline resistance genes in biological nutrient removal process. *J Environ Manage*. 2019 May 15;238:84-91. PMID: 30849601. doi: 10.1016/j.jenvman.2019.02.123.
530. Loke HY, Win M-K, Chen MI-C, Lim J-W, Ang BSP, Chow ALP. Length of stay and odds of MRSA acquisition: a dose-response relationship? *Epidemiol Infect*. 2019 Jan;147:e223. PMID: 31364542. doi: 10.1017/s0950268819001110.
531. Loo LW, Liew YX, Lee WHL, Lee LW, Chlebicki MP, Kwa AL-H. Discontinuation of antibiotic therapy within 24 hours of treatment initiation for patients with no clinical evidence of bacterial infection: a 5-year safety and outcome study from Singapore General Hospital Antimicrobial Stewardship Program. *Int J Antimicrob Agents*. 2019 May;53(5):606-11. PMID: 30639630. doi: 10.1016/j.ijantimicag.2019.01.008.
532. Marimuthu K, Ng OT, Cherng BPZ, Fong RKC, Pada SMK, De PP, et al. Antecedent Carbapenem Exposure as a Risk Factor for Non-Carbapenemase-Producing Carbapenem-Resistant Enterobacteriaceae and Carbapenemase-Producing Enterobacteriaceae. *Antimicrob Agents Chemother*. 2019 Oct;63(10). PMID: 31383670. doi: 10.1128/aac.00845-19.
533. Mayandi V, Sridhar S, Fazil MHUT, Goh ETL, Htoon HM, Orive G, et al. Protective Action of Linear Polyethylenimine against *Staphylococcus aureus* Colonization and Exaggerated Inflammation in Vitro and in Vivo. *ACS Infect Dis*. 2019 Aug 9;5(8):1411-22. PMID: 31099239. doi: 10.1021/acsinfecdis.9b00102.
534. Mo Y, Seah I, Lye PSP, Kee XLJ, Wong KYM, Ko KKK, et al. Relating knowledge, attitude and practice of antibiotic use to extended-spectrum beta-lactamase-producing Enterobacteriaceae carriage: results of a cross-sectional community survey. *BMJ Open*. 2019 Mar 5;9(3):e023859. PMID: 30842108. doi: 10.1136/bmjopen-2018-023859.
535. Mo Y, Low I, Tambyah SK, Tambyah PA. The socio-economic impact of multidrug-resistant nosocomial infections: a qualitative study. *J Hosp Infect*. 2019 Aug;102(4):454-60. PMID: 30171886. doi: 10.1016/j.jhin.2018.08.013.
536. Ng C, Tan B, Jiang X-T, Gu X, Chen H-J, Schmitz BW, et al. Metagenomic and Resistome Analysis of a Full-Scale Municipal Wastewater Treatment Plant in Singapore Containing Membrane Bioreactors. *Front Microbiol*. 2019;10:172. PMID: 30833934. doi: 10.3389/fmicb.2019.00172.
537. Ngo T-M, Teo Y-Y. Genomic prediction of tuberculosis drug-resistance: benchmarking existing databases and prediction algorithms. *BMC Bioinformatics*. 2019 Feb 8;20(1):68. PMID: 30736750. doi: 10.1186/s12859-019-2658-z.
538. Obuobi S, Tay HK-L, Tram NDT, Selvarajan V, Khara JS, Wang Y, et al. Facile and efficient encapsulation of antimicrobial peptides via crosslinked DNA nanostructures and their application in wound therapy. *J Control Release*. 2019 Nov 10;313:120-30. PMID: 31629042. doi: 10.1016/j.jconrel.2019.10.013.
539. Octavia S, Marimuthu K, Venkatachalam I, Ng OT, Xu W, Sridatta PSR, et al. *Klebsiella pneumoniae* and *Klebsiella quasipneumoniae* define the population structure of blaKPC-2Klebsiella: a 5 year retrospective genomic study in Singapore. *J Antimicrob Chemother*. 2019 Nov 1;74(11):3205-10. PMID: 31504571. doi: 10.1093/jac/dkz332.
540. Pang X, Wong CH, Chung H-J, Yuk H-G. Biofilm formation of *Listeria monocytogenes* and its resistance to quaternary ammonium compounds in a simulated salmon processing environment. *Food Control*. 2019 2019/04/01/;98:200-8. doi: <https://doi.org/10.1016/j.foodcont.2018.11.029>.
541. Quek WM, Teng CB, Tan YZ, Chong K, Lye DCB, Ng TM. Outcomes of Fosfomycin Use in Ceftriaxone-Resistant Enterobacteriaceae Urinary Tract Infection in the Elderly. *Int J Antimicrob Agents*. 2019 Feb;53(2):195-6. PMID: 30722963. doi: 10.1016/j.ijantimicag.2018.10.008.
542. Ryanputra D, Wang D, Lee MB, Teo BW, Tok PL. Peritoneal Dialysis-Related Peritonitis from Carbapenemase-Producing *Klebsiella pneumoniae* with OXA-48 Type Gene. *Perit Dial Int*. 2019 Jan-Feb;39(1):97-8. PMID: 30692237. doi: 10.3747/pdi.2018.00234.
543. Selcuk A, Yap KZ, Wong CL, Yang JX, Yong PC, Chan SY, et al. A Point Prevalence Study of Antimicrobial Use and Practice Among Nursing Homes in Singapore. *Drugs Aging*. 2019 Jun;36(6):559-70. PMID: 30972716. doi: 10.1007/s40266-019-00651-2.
544. Singh SR, Chua AQ, Tan ST, Tam CC, Hsu L-Y, Legido-Quigley H. Combating Antimicrobial Resistance in Singapore: A Qualitative Study Exploring the Policy Context, Challenges, Facilitators, and Proposed Strategies. *Antibiotics (Basel)*. 2019 Oct 29;8(4). PMID: 31671826. doi: 10.3390/antibiotics8040201.

545. Sorayah R, Manimekalai MSS, Shin SJ, Koh W-J, Grüber G, Pethe K. Naturally-Occurring Polymorphisms in QcrB Are Responsible for Resistance to Telacebec in *Mycobacterium abscessus*. *ACS Infect Dis.* 2019 Dec 13;5(12):2055-60. PMID: 31599569. doi: 10.1021/acsinfecdis.9b00322.
546. Stewardson AJ, Marimuthu K, Sengupta S, Allignol A, El-Bouseary M, Carvalho MJ, et al. Effect of carbapenem resistance on outcomes of bloodstream infection caused by Enterobacteriaceae in low-income and middle-income countries (PANORAMA): a multinational prospective cohort study. *Lancet Infect Dis.* 2019 Jun;19(6):601-10. PMID: 31047852. doi: 10.1016/s1473-3099(18)30792-8.
547. Su CT-T, Koh DW-S, Gan SK-E. Reviewing HIV-1 Gag Mutations in Protease Inhibitors Resistance: Insights for Possible Novel Gag Inhibitor Designs. *Molecules.* 2019 Sep 6;24(18). PMID: 31489889. doi: 10.3390/molecules24183243.
548. Tan GSE, Octavia S, Teo JWP, Ang MLT, La M-V, Lin RTP, et al. In vivo development of amoxicillin and ceftriaxone resistance in *Salmonella enterica* serovar Typhi. *Clin Microbiol Infect.* 2019 Aug;25(8):1045-7. PMID: 30914348. doi: 10.1016/j.cmi.2019.03.015.
549. Tan YE, Teo JQM, Abdul Rahman NB, Ng OT, Marimuthu K, Tan AL, et al. *Candida auris* in Singapore: Genomic epidemiology, antifungal drug resistance, and identification using the updated 8.01 VITEK® system. *Int J Antimicrob Agents.* 2019 Dec;54(6):709-15. PMID: 31600556. doi: 10.1016/j.ijantimicag.2019.09.016.
550. Tay MYF, Pathirage S, Chandrasekaran L, Wickramasuriya U, Sadeepanie N, Waidyaratna KDK, et al. Whole-Genome Sequencing Analysis of Nontyphoidal *Salmonella enterica* of Chicken Meat and Human Origin Under Surveillance in Sri Lanka. *Foodborne Pathog Dis.* 2019 Jul;16(7):531-7. PMID: 31099590. doi: 10.1089/fpd.2018.2604.
551. Teo JQM, Lee SJ-Y, Tan AL, Lim RS-M, Cai Y, Lim T-P, et al. Molecular mechanisms of azole resistance in *Candida* bloodstream isolates. *BMC Infect Dis.* 2019 Jan 17;19(1):63. PMID: 30654757. doi: 10.1186/s12879-019-3672-5.
552. Teo JQM, Chang CW-T, Leck H, Tang C-Y, Lee SJ-Y, Cai Y, et al. Risk factors and outcomes associated with the isolation of polymyxin B and carbapenem-resistant Enterobacteriaceae spp.: A case-control study. *Int J Antimicrob Agents.* 2019 May;53(5):657-62. PMID: 30880229. doi: 10.1016/j.ijantimicag.2019.03.011.
553. Teo SW, Yong YE, Ng SMS, Ng FM, Teo JWP, Jureen R, et al. Identification of an Ultra-Short Peptide with Potent *Pseudomonas aeruginosa* Activity for Development as a Topical Antibacterial Agent. *INTERNATIONAL JOURNAL OF PEPTIDE RESEARCH AND THERAPEUTICS.* 2019 MAR;25(1):359-66. PMID: WOS:000457490500036. doi: 10.1007/s10989-018-9678-9.
554. Tong JX, Ang SEL, Tan EHN, Tan KSW. Viability Screen of LOPAC(1280) Reveals Tyrosine Kinase Inhibitor Tyrphostin A9 as a Novel Partner Drug for Artesunate Combinations To Target the Plasmodium falciparum Ring Stage. *Antimicrob Agents Chemother.* 2019 Apr;63(4). PMID: 30718250. doi: 10.1128/aac.02389-18.
555. Tran NH, Hoang L, Nghiem LD, Nguyen NMH, Ngo HH, Guo W, et al. Occurrence and risk assessment of multiple classes of antibiotics in urban canals and lakes in Hanoi, Vietnam. *Sci Total Environ.* 2019 Nov 20;692:157-74. PMID: 31344569. doi: 10.1016/j.scitotenv.2019.07.092.
556. Vasoo S, Hon PY, Hsu J-P, Koh TH, De PP. A pilot study on the analytic sensitivity and accuracy of the Check-Points Check Direct CPE, Cepheid Xpert Carba-R, and ChromID CARBA SMART chromogenic agar for detecting carbapenemase-producing Enterobacteriaceae. *J Glob Antimicrob Resist.* 2019 Sep;18:235-7. PMID: 31202978. doi: 10.1016/j.jgar.2019.06.003.
557. Yang C, Lou W, Zhong G, Lee ALZ, Leong J, Chin W, et al. Degradable antimicrobial polycarbonates with unexpected activity and selectivity for treating multidrug-resistant *Klebsiella pneumoniae* lung infection in mice. *Acta Biomater.* 2019 Aug;94:268-80. PMID: 31129359. doi: 10.1016/j.actbio.2019.05.057.
558. Yi X, Lin C, Ong EJL, Wang M, Li B, Zhou Z. Expression of resistance genes instead of gene abundance are correlated with trace levels of antibiotics in urban surface waters. *Environ Pollut.* 2019 Jul;250:437-46. PMID: 31026690. doi: 10.1016/j.envpol.2019.04.035.
559. Yi X, Wang M, Zhou Z. The potential impact of naturally produced antibiotics, environmental factors, and anthropogenic pressure on the occurrence of erm genes in urban soils. *Environ Pollut.* 2019 Feb;245:282-9. PMID: 30445415. doi: 10.1016/j.envpol.2018.11.009.
560. Yuan Y, Zhou F, Su H, Zhang Y. Structural design of microbicidal cationic oligomers and their synergistic interaction with azoles against *Candida albicans*. *Sci Rep.* 2019 Aug 15;9(1):11885. PMID: 31417167. doi: 10.1038/s41598-019-48322-x.
561. Yuan Y, Liang S, Li J, Zhang S, Zhang Y. Copolymers with both soft and rigid cationic rings as highly selective antimicrobials to combat antibiotic resistant microbes and biofilms. *J Mater Chem B.* 2019 Sep 25;7(37):5620-5. PMID: 31508648. doi: 10.1039/c9tb01264h.

562. Zhang EX, Chavatte J-M, See CXY, Tow C, Wong JY, Khan K, et al. Assessment of the risk posed to Singapore by the emergence of artemisinin-resistant malaria in the Greater Mekong Subregion. *Western Pac Surveill Response J.* 2019 Apr-Jun;10(2):6-13. PMID: 31720049. doi: 10.5365/wpsar.2018.9.2.011.
563. Zhang J, Chua QW, Mao F, Zhang L, He Y, Tong YW, et al. Effects of activated carbon on anaerobic digestion – Methanogenic metabolism, mechanisms of antibiotics and antibiotic resistance genes removal. *Bioresource Technology Reports.* 2019;5:113-20. doi: 10.1016/j.biteb.2019.01.002.
564. Zhang K, Du Y, Si Z, Liu Y, Turvey ME, Raju C, et al. Enantiomeric glycosylated cationic block co-beta-peptides eradicate *Staphylococcus aureus* biofilms and antibiotic-tolerant persisters. *Nat Commun.* 2019 Oct 21;10(1):4792. PMID: 31636263. doi: 10.1038/s41467-019-12702-8.
565. Zhang L, Loh K-C, Zhang J. Jointly reducing antibiotic resistance genes and improving methane yield in anaerobic digestion of chicken manure by feedstock microwave pretreatment and activated carbon supplementation. *Chemical Engineering Journal.* 2019 2019/09/15/;372:815-24. doi: <https://doi.org/10.1016/j.cej.2019.04.207>.
566. Zheng S-W. Disseminated nocardiosis due to *< i>Nocardia otitidiscavarium</i>*: A case report and literature review. *Asian Pacific Journal of Tropical Medicine.* 2019 April 1, 2019;12(4):185-94. doi: 10.4103/1995-7645.257120.
567. Zhou C, Song H, Zhang F, Liu J, Li J, Liu B, et al. A facile method to fabricate an antimicrobial coating based on poly(1-vinyl-3-allylimidazolium iodide) (PAVI) and poly(ethylene glycol) dimethyl acrylate (PEGDMA). *Polymer Bulletin.* 2019;76(10):5433-49. doi: 10.1007/s00289-018-2637-y.
568. Bouchillon SK, Iredell JR, Barkham TMS, Lee K, Dowzicky MJ. Comparative in vitro activity of tigecycline and other antimicrobials against Gram-negative and Gram-positive organisms collected from the Asia-Pacific Rim as part of the Tigecycline Evaluation and Surveillance Trial (TEST). *Int J Antimicrob Agents.* 2009 Feb;33(2):130-6. PMID: 18995992. doi: 10.1016/j.ijantimicag.2008.07.031.
569. Chuang C-H, Su L-H, Perera J, Carlos CC, Tan BH, Kumarasinghe G, et al. Surveillance of antimicrobial resistance of *Salmonella enterica* serotype Typhi in seven Asian countries. *Epidemiol Infect.* 2009 Feb;137(2):266-9. PMID: 18474127. doi: 10.1017/s0950268808000745.
570. Hawser SP, Bouchillon SK, Hoban DJ, Badal RE, Hsueh P-R, Paterson DL. Emergence of high levels of extended-spectrum-beta-lactamase-producing gram-negative bacilli in the Asia-Pacific region: data from the Study for Monitoring Antimicrobial Resistance Trends (SMART) program, 2007. *Antimicrob Agents Chemother.* 2009 Aug;53(8):3280-4. PMID: 19506060. doi: 10.1128/aac.00426-09.
571. Hurt AC, Ernest J, Deng Y-M, Iannello P, Besselaar TG, Birch C, et al. Emergence and spread of oseltamivir-resistant A(H1N1) influenza viruses in Oceania, South East Asia and South Africa. *Antiviral Res.* 2009 Jul;83(1):90-3. PMID: 19501261. doi: 10.1016/j.antiviral.2009.03.003.
572. Ko W-C, Hsueh P-R. Increasing extended-spectrum beta-lactamase production and quinolone resistance among Gram-negative bacilli causing intra-abdominal infections in the Asia/Pacific region: data from the Smart Study 2002-2006. *J Infect.* 2009 Aug;59(2):95-103. PMID: 19564044. doi: 10.1016/j.jinf.2009.06.003.
573. Lee H-Y, Su L-H, Tsai M-H, Kim S-W, Chang H-H, Jung S-I, et al. High rate of reduced susceptibility to ciprofloxacin and ceftriaxone among nontyphoid *Salmonella* clinical isolates in Asia. *Antimicrob Agents Chemother.* 2009 Jun;53(6):2696-9. PMID: 19332677. doi: 10.1128/aac.01297-08.
574. Mendes RE, Bell JM, Turnidge JD, Castanheira M, Jones RN. Emergence and widespread dissemination of OXA-23, -24/40 and -58 carbapenemases among *Acinetobacter* spp. in Asia-Pacific nations: report from the SENTRY Surveillance Program. *J Antimicrob Chemother.* 2009 Jan;63(1):55-9. PMID: 18957398. doi: 10.1093/jac/dkn434.
575. Yau W, Owen RJ, Poudyal A, Bell JM, Turnidge JD, Yu HH, et al. Colistin hetero-resistance in multidrug-resistant *Acinetobacter baumannii* clinical isolates from the Western Pacific region in the SENTRY antimicrobial surveillance programme. *J Infect.* 2009 Feb;58(2):138-44. PMID: 19058855. doi: 10.1016/j.jinf.2008.11.002.
576. Christiansen K, Ip M, Ker H, Mendoza M, Hsu L-Y, Kiratisin P, et al. In vitro activity of doripenem and other carbapenems against contemporary Gram-negative pathogens isolated from hospitalised patients in the Asia-Pacific region: results of the COMPACT Asia-Pacific Study. *Int J Antimicrob Agents.* 2010 Dec;36(6):501-6. PMID: 20869212. doi: 10.1016/j.ijantimicag.2010.08.002.
577. Farrell DJ, Turnidge JD, Bell JM, Sader HS, Jones RN. The in vitro evaluation of tigecycline tested against pathogens isolated in eight countries in the Asia-Western Pacific region (2008). *J Infect.* 2010 Jun;60(6):440-51. PMID: 20361999. doi: 10.1016/j.jinf.2010.03.024.
578. Higgins PG, Dammhayn C, Hackel MA, Seifert H. Global spread of carbapenem-resistant *Acinetobacter baumannii*. *J Antimicrob Chemother.* 2010 Feb;65(2):233-8. PMID: 19996144. doi: 10.1093/jac/dkp428.

579. Hsueh P-R, Badal RE, Hawser SP, Hoban DJ, Bouchillon SK, Ni Y, et al. Epidemiology and antimicrobial susceptibility profiles of aerobic and facultative Gram-negative bacilli isolated from patients with intra-abdominal infections in the Asia-Pacific region: 2008 results from SMART (Study for Monitoring Antimicrobial Resistance Trends). *Int J Antimicrob Agents*. 2010 Nov;36(5):408-14. PMID: 20728316. doi: 10.1016/j.ijantimicag.2010.07.002.
580. Chen Y-H, Hsueh P-R, Badal RE, Hawser SP, Hoban DJ, Bouchillon SK, et al. Antimicrobial susceptibility profiles of aerobic and facultative Gram-negative bacilli isolated from patients with intra-abdominal infections in the Asia-Pacific region according to currently established susceptibility interpretive criteria. *J Infect*. 2011 Apr;62(4):280-91. PMID: 21382411. doi: 10.1016/j.jinf.2011.02.009.
581. Chung DR, Song J-H, Kim SH, Thamlikitkul V, Huang S-G, Wang HC-H, et al. High prevalence of multidrug-resistant nonfermenters in hospital-acquired pneumonia in Asia. *Am J Respir Crit Care Med*. 2011 Dec 15;184(12):1409-17. PMID: 21920919. doi: 10.1164/rccm.201102-0349OC.
582. Hurt AC, Lee RTC, Leang S-K, Lin C, Deng Y-M, Phuah S, et al. Increased detection in Australia and Singapore of a novel influenza A(H1N1)2009 variant with reduced oseltamivir and zanamivir sensitivity due to a S247N neuraminidase mutation. *Euro Surveill*. 2011 Jun 9;16(23). PMID: 21679678.
583. Hurt AC, Deng Y-M, Ernest J, Caldwell N, Leang L, Iannello P, et al. Oseltamivir-resistant influenza viruses circulating during the first year of the influenza A(H1N1) 2009 pandemic in the Asia-Pacific region, March 2009 to March 2010. *Euro Surveill*. 2011 Jan 20;16(3). PMID: 21262183.
584. Lee MY, Kang C-I, Chung DR, Peck KR, Ko KS, Song J-H. High prevalence of CTX-M-15-producing Klebsiella pneumoniae isolates in Asian countries: diverse clones and clonal dissemination. *Int J Antimicrob Agents*. 2011 Aug;38(2):160-3. PMID: 21605960. doi: 10.1016/j.ijantimicag.2011.03.020.
585. Roberts JA, Kwa AL-H, Montakantikul P, Gomersall C, Kuti JL, Nicolau DP. Pharmacodynamic profiling of intravenous antibiotics against prevalent Gram-negative organisms across the globe: the PASSPORT Program-Asia-Pacific Region. *Int J Antimicrob Agents*. 2011 Mar;37(3):225-9. PMID: 21168997. doi: 10.1016/j.ijantimicag.2010.10.024.
586. Wang HC-H, Chen M, Xu Y, Sun H, Yang Q, Hu Y, et al. Antimicrobial susceptibility of bacterial pathogens associated with community-acquired respiratory tract infections in Asia: Report from the Community-Acquired Respiratory Tract Infection Pathogen Surveillance (CARTIPS) study, 2009-2010. *International Journal of Antimicrobial Agents*. 2011;38(5):376-83. doi: 10.1016/j.ijantimicag.2011.06.015.
587. Bouchillon SK, Hoban DJ, Badal RE, Hawser SP. Fluoroquinolone resistance among gram-negative urinary tract pathogens: global smart program results, 2009-2010. *Open Microbiol J*. 2012;6:74-8. PMID: 23002406. doi: 10.2174/187428580120610074.
588. Kiratisin P, Chongthaleong A, Tan TY, Lagamayo E, Roberts S, Garcia J, et al. Comparative in vitro activity of carbapenems against major Gram-negative pathogens: results of Asia-Pacific surveillance from the COMPACT II study. *Int J Antimicrob Agents*. 2012 Apr;39(4):311-6. PMID: 22386743. doi: 10.1016/j.ijantimicag.2012.01.002.
589. Lin Y-T, Siu LK, Lin J-C, Chen T-L, Tseng C-P, Yeh K-M, et al. Seroepidemiology of Klebsiella pneumoniae colonizing the intestinal tract of healthy Chinese and overseas Chinese adults in Asian countries. *BMC Microbiol*. 2012 Jan 19;12:13. PMID: 22260182. doi: 10.1186/1471-2180-12-13.
590. Lu P-L, Liu Y-C, Toh H-S, Lee Y-L, Liu Y-M, Ho C-M, et al. Epidemiology and antimicrobial susceptibility profiles of Gram-negative bacteria causing urinary tract infections in the Asia-Pacific region: 2009-2010 results from the Study for Monitoring Antimicrobial Resistance Trends (SMART). *Int J Antimicrob Agents*. 2012 Jun;40 Suppl:S37-43. PMID: 22749057. doi: 10.1016/s0924-8579(12)70008-0.
591. Namdari H, Tan TY, Dowzicky MJ. Activity of tigecycline and comparators against skin and skin structure pathogens: global results of the Tigecycline Evaluation and Surveillance Trial, 2004-2009. *INTERNATIONAL JOURNAL OF INFECTIOUS DISEASES*. 2012 JAN;16(1):E60-E6. PMID: WOS:000299013200011. doi: 10.1016/j.ijid.2011.09.021.
592. Yang Y, Zhang T, Zhang X-X, Liang D-W, Zhang M, Gao D-W, et al. Quantification and characterization of β -lactam resistance genes in 15 sewage treatment plants from East Asia and North America. *Appl Microbiol Biotechnol*. 2012 Sep;95(5):1351-8. PMID: 22202968. doi: 10.1007/s00253-011-3810-5.
593. Holden MT, Hsu L-Y, Kurt K, Weinert LA, Mather AE, Harris SR, et al. A genomic portrait of the emergence, evolution, and global spread of a methicillin-resistant *Staphylococcus aureus* pandemic. *Genome Res*. 2013 Apr;23(4):653-64. PMID: 23299977. doi: 10.1101/gr.147710.112.
594. Kim DH, Choi J-Y, Kim HW, Kim SH, Chung DR, Peck KR, et al. Spread of carbapenem-resistant *Acinetobacter baumannii* global clone 2 in Asia and AbaR-type resistance islands. *Antimicrob Agents Chemother*. 2013 Nov;57(11):5239-46. PMID: 23939892. doi: 10.1128/aac.00633-13.

595. Leang S-K, Deng Y-M, Shaw R, Caldwell N, Iannello P, Komadina N, et al. Influenza antiviral resistance in the Asia-Pacific region during 2011. *Antiviral Res.* 2013 Feb;97(2):206-10. PMID: 23274624. doi: 10.1016/j.antiviral.2012.12.016.
596. Mendes RE, Mendoza M, Singh KKB, Castanheira M, Bell JM, Turnidge JD, et al. Regional resistance surveillance program results for 12 Asia-Pacific nations (2011). *Antimicrob Agents Chemother.* 2013 Nov;57(11):5721-6. PMID: 23959306. doi: 10.1128/aac.01121-13.
597. Sader HS, Flamm RK, Jones RN. Antimicrobial activity of ceftaroline and comparator agents tested against bacterial isolates causing skin and soft tissue infections and community-acquired respiratory tract infections isolated from the Asia-Pacific region and South Africa (2010). *Diagn Microbiol Infect Dis.* 2013 May;76(1):61-8. PMID: 23535208. doi: 10.1016/j.diagmicrobio.2013.01.005.
598. Sheng W-H, Badal RE, Hsueh P-R. Distribution of extended-spectrum β -lactamases, AmpC β -lactamases, and carbapenemases among Enterobacteriaceae isolates causing intra-abdominal infections in the Asia-Pacific region: results of the study for Monitoring Antimicrobial Resistance Trends (SMART). *Antimicrob Agents Chemother.* 2013 Jul;57(7):2981-8. PMID: 23587958. doi: 10.1128/aac.00971-12.
599. Ginn AN, Wiklund AM, Zong Z, Lin RTP, Teo JWP, Tambyah PA, et al. Prediction of major antibiotic resistance in Escherichia coli and Klebsiella pneumoniae in Singapore, USA and China using a limited set of gene targets. *Int J Antimicrob Agents.* 2014 Jun;43(6):563-5. PMID: 24721234. doi: 10.1016/j.ijantimicag.2014.02.010.
600. Holt KE, Wertheim H, Zadoks RN, Baker S, Whitehouse CA, Dance D, et al. Genomic analysis of diversity, population structure, virulence, and antimicrobial resistance in Klebsiella pneumoniae, an urgent threat to public health. *Proc Natl Acad Sci U S A.* 2015 Jul 7;112(27):E3574-81. PMID: 26100894. doi: 10.1073/pnas.1501049112.
601. Pfaller MA, Messer SA, Jones RN, Castanheira M. Antifungal susceptibilities of Candida, Cryptococcus neoformans and Aspergillus fumigatus from the Asia and Western Pacific region: data from the SENTRY antifungal surveillance program (2010-2012). *J Antibiot (Tokyo).* 2015 Sep;68(9):556-61. PMID: 25899126. doi: 10.1038/ja.2015.29.
602. Jean S-S, Coombs G, Ling T, Balaji V, Rodrigues C, Mikamo H, et al. Epidemiology and antimicrobial susceptibility profiles of pathogens causing urinary tract infections in the Asia-Pacific region: Results from the Study for Monitoring Antimicrobial Resistance Trends (SMART), 2010-2013. *Int J Antimicrob Agents.* 2016 Apr;47(4):328-34. PMID: 27005459. doi: 10.1016/j.ijantimicag.2016.01.008.
603. Tan TY, Hsu L-Y, Alejandria MM, Chaiwarith R, Chinniah T, Chayakulkeeree M, et al. Antifungal susceptibility of invasive Candida bloodstream isolates from the Asia-Pacific region. *Med Mycol.* 2016 Jul 1;54(5):471-7. PMID: 26868904. doi: 10.1093/mmy/myv114.
604. Torumkuney D, Chaiwarith R, Reechaipichitkul W, Malatham K, Chareonphaibul V, Rodrigues C, et al. Results from the Survey of Antibiotic Resistance (SOAR) 2012-14 in Thailand, India, South Korea and Singapore. *J Antimicrob Chemother.* 2016 May;71 Suppl 1(Suppl 1):i3-19. PMID: 27048580. doi: 10.1093/jac/dkw073.
605. Blackwell GA, Hall RM. The tet39 Determinant and the msrE-mphE Genes in Acinetobacter Plasmids Are Each Part of Discrete Modules Flanked by Inversely Oriented pdif (XerC-XerD) Sites. *Antimicrob Agents Chemother.* 2017 Aug;61(8). PMID: 28533235. doi: 10.1128/aac.00780-17.
606. Blackwell GA, Holt KE, Bentley SD, Hsu L-Y, Hall RM. Variants of AbGRI3 carrying the armA gene in extensively antibiotic-resistant Acinetobacter baumannii from Singapore. *J Antimicrob Chemother.* 2017 Apr 1;72(4):1031-9. PMID: 28073968. doi: 10.1093/jac/dkw542.
607. Chang Y-T, Coombs G, Ling T, Balaji V, Rodrigues C, Mikamo H, et al. Epidemiology and trends in the antibiotic susceptibilities of Gram-negative bacilli isolated from patients with intra-abdominal infections in the Asia-Pacific region, 2010-2013. *Int J Antimicrob Agents.* 2017 Jun;49(6):734-9. PMID: 28435019. doi: 10.1016/j.ijantimicag.2017.01.030.
608. Cheong HS, Chung DR, Park M, Kim SH, Ko KS, Ha YE, et al. Emergence of an extended-spectrum β -lactamase-producing serotype K1 Klebsiella pneumoniae ST23 strain from Asian countries. *Epidemiol Infect.* 2017 Apr;145(5):990-4. PMID: 28031071. doi: 10.1017/s0950268816003113.
609. Jean S-S, Hsueh P-R. Distribution of ESBLs, AmpC β -lactamases and carbapenemases among Enterobacteriaceae isolates causing intra-abdominal and urinary tract infections in the Asia-Pacific region during 2008-14: results from the Study for Monitoring Antimicrobial Resistance Trends (SMART). *J Antimicrob Chemother.* 2017 Jan;72(1):166-71. PMID: 27703058. doi: 10.1093/jac/dkw398.
610. Karlowsky JA, Hoban DJ, Hackel MA, Lob SH, Sahm DF. Antimicrobial susceptibility of Gram-negative ESKAPE pathogens isolated from hospitalized patients with intra-abdominal and urinary tract infections in Asia-Pacific countries: SMART 2013-2015. *J Med Microbiol.* 2017 Jan;66(1):61-9. PMID: 28051952. doi: 10.1099/jmm.0.000421.

611. Ma L, Li B, Jiang X-T, Wang Y-L, Xia Y, Li A-D, et al. Catalogue of antibiotic resistome and host-tracking in drinking water deciphered by a large scale survey. *Microbiome*. 2017 Nov 28;5(1):154. PMID: 29179769. doi: 10.1186/s40168-017-0369-0.
612. Harris PNA, Ben Zakour NL, Roberts LW, Wailan AM, Zowawi HM, Tambyah PA, et al. Whole genome analysis of cephalosporin-resistant *Escherichia coli* from bloodstream infections in Australia, New Zealand and Singapore: high prevalence of CMY-2 producers and ST131 carrying blaCTX-M-15 and blaCTX-M-27. *J Antimicrob Chemother*. 2018 Mar 1;73(3):634-42. PMID: 29253152. doi: 10.1093/jac/dkx466.
613. Harris PNA, Tambyah PA, Lye DCB, Mo Y, Lee TH, Yilmaz M, et al. Effect of Piperacillin-Tazobactam vs Meropenem on 30-Day Mortality for Patients With *E. coli* or *Klebsiella pneumoniae* Bloodstream Infection and Ceftriaxone Resistance: A Randomized Clinical Trial. *Jama*. 2018 Sep 11;320(10):984-94. PMID: 30208454. doi: 10.1001/jama.2018.12163.
614. Khor WC, Puah SM, Koh TH, Tan JAMA, Puthucheary SD, Chua KH. Comparison of Clinical Isolates of *Aeromonas* from Singapore and Malaysia with Regard to Molecular Identification, Virulence, and Antimicrobial Profiles. *Microb Drug Resist*. 2018 May;24(4):469-78. PMID: 29461928. doi: 10.1089/mdr.2017.0083.
615. Mendis SM, Vasoo S, Johnston BD, Porter SB, Cunningham SA, Menon SR, et al. Clinical and Molecular Correlates of *Escherichia coli* Bloodstream Infection from Two Geographically Diverse Centers in Rochester, Minnesota, and Singapore. *Antimicrob Agents Chemother*. 2018 Oct;62(10). PMID: 30082285. doi: 10.1128/aac.00937-18.
616. Pfaller MA, Shortridge D, Sader HS, Castanheira M, Flamm RK. Ceftolozane/tazobactam activity against drug-resistant Enterobacteriaceae and *Pseudomonas aeruginosa* causing healthcare-associated infections in the Asia-Pacific region (minus China, Australia and New Zealand): report from an Antimicrobial Surveillance Programme (2013-2015). *Int J Antimicrob Agents*. 2018 Feb;51(2):181-9. PMID: 28993143. doi: 10.1016/j.ijantimicag.2017.09.016.
617. Versporten A, Zarb P, Caniaux I, Gros M-F, Drapier N, Miller M, et al. Antimicrobial consumption and resistance in adult hospital inpatients in 53 countries: results of an internet-based global point prevalence survey. *Lancet Glob Health*. 2018 Jun;6(6):e619-e29. PMID: 29681513. doi: 10.1016/s2214-109x(18)30186-4.
618. Vilaichone RK, Quach DT, Yamaoka Y, Sugano K, Mahachai V. Prevalence and Pattern of Antibiotic Resistant Strains of *Helicobacter pylori* Infection in ASEAN. *Asian Pac J Cancer Prev*. 2018 May 26;19(5):1411-3. PMID: 29802708. doi: 10.22034/apjcp.2018.19.5.1411.
619. Yarbrough ML, Warren DK, Allen K, Burkholder D, Daum R, Donskey C, et al. Multicenter Evaluation of the Xpert MRSA NxG Assay for Detection of Methicillin-Resistant *Staphylococcus aureus* in Nasal Swabs. *J Clin Microbiol*. 2018 Jan;56(1). PMID: 29118165. doi: 10.1128/jcm.01381-17.
620. Blot S, Antonelli M, Arvaniti K, Blot K, Creagh-Brown B, de Lange D, et al. Epidemiology of intra-abdominal infection and sepsis in critically ill patients: "AbSeS", a multinational observational cohort study and ESICM Trials Group Project. *INTENSIVE CARE MEDICINE*. 2019 DEC;45(12):1703-17. PMID: WOS:000493268200001. doi: 10.1007/s00134-019-05819-3.
621. Chen SL, Ding Y, Apisarnthanarak A, Kalimuddin S, Archuleta S, Omar SFS, et al. The higher prevalence of extended spectrum beta-lactamases among *Escherichia coli* ST131 in Southeast Asia is driven by expansion of a single, locally prevalent subclone. *Sci Rep*. 2019 Sep 13;9(1):13245. PMID: 31519972. doi: 10.1038/s41598-019-49467-5.
622. George CRR, Enriquez RP, Gatus BJ, Whiley DM, Lo Y-R, Ishikawa N, et al. Systematic review and survey of *Neisseria gonorrhoeae* ceftriaxone and azithromycin susceptibility data in the Asia Pacific, 2011 to 2016. *PLoS One*. 2019;14(4):e0213312. PMID: 30943199. doi: 10.1371/journal.pone.0213312.
623. Hendriksen RS, Munk P, Njage P, van Bunnik B, McNally L, Lukjancenko O, et al. Global monitoring of antimicrobial resistance based on metagenomics analyses of urban sewage. *Nature Communications*. 2019;10(1):1124. doi: 10.1038/s41467-019-08853-3.
624. Hsia Y, Lee BR, Versporten A, Yang Y, Bielicki JA, Jackson C, et al. Use of the WHO Access, Watch, and Reserve classification to define patterns of hospital antibiotic use (AWaRe): an analysis of paediatric survey data from 56 countries. *Lancet Glob Health*. 2019 Jul;7(7):e861-e71. PMID: 31200888. doi: 10.1016/s2214-109x(19)30071-3.
625. Hsia Y, Sharland M, Jackson C, Wong ICK, Magrini N, Bielicki JA. Consumption of oral antibiotic formulations for young children according to the WHO Access, Watch, Reserve (AWaRe) antibiotic groups: an analysis of sales data from 70 middle-income and high-income countries. *Lancet Infect Dis*. 2019 Jan;19(1):67-75. PMID: 30522834. doi: 10.1016/s1473-3099(18)30547-4.
626. Hu YJ, Ogyu A, Cowling BJ, Fukuda K, Pang HH. Available evidence of antibiotic resistance from extended-spectrum β-lactamase-producing Enterobacteriaceae in paediatric patients in 20 countries: a systematic review and meta-analysis. *Bulletin of the World Health Organization*.

- 2019;97(7):486-501B. PMID: 137322024. Language: English. Entry Date: 20190708. Revision Date: 20190709. Publication Type: Article. doi: 10.2471/BLT.18.225698.
627. Luo Y, Cheong E, Bian Q, Collins DA, Ye J, Shin JH, et al. Different molecular characteristics and antimicrobial resistance profiles of *Clostridium difficile* in the Asia-Pacific region. *Emerg Microbes Infect*. 2019;8(1):1553-62. PMID: 31662120. doi: 10.1080/22221751.2019.1682472.
628. Ma L, Li B, Zhang T. New insights into antibiotic resistome in drinking water and management perspectives: A metagenomic based study of small-sized microbes. *Water Res*. 2019 Apr 1;152:191-201. PMID: 30669041. doi: 10.1016/j.watres.2018.12.069.
629. Papadopoulos A, Ribera A, Mavrogenis AF, Rodriguez-Pardo D, Bonnet E, Salles MJ, et al. Multidrug-resistant and extensively drug-resistant Gram-negative prosthetic joint infections: Role of surgery and impact of colistin administration. *Int J Antimicrob Agents*. 2019 Mar;53(3):294-301. PMID: 30395988. doi: 10.1016/j.ijantimicag.2018.10.018.
630. Cervantes S, Prudhomme J, Carter D, Gopi KG, Li Q, Chang Y-T, et al. High-content live cell imaging with RNA probes: advancements in high-throughput antimalarial drug discovery. *BMC CELL BIOLOGY*. 2009 JUN 10;10. PMID: WOS:000267598200001. doi: 10.1186/1471-2121-10-45.
631. Tam VH, Ledesma KR, Schilling AN, Lim T-P, Yuan Z, Ghose R, et al. In vivo dynamics of carbapenem-resistant *Pseudomonas aeruginosa* selection after suboptimal dosing. *DIAGNOSTIC MICROBIOLOGY AND INFECTIOUS DISEASE*. 2009 AUG;64(4):427-33. PMID: WOS:000269337900011. doi: 10.1016/j.diagmicrobio.2009.03.031.
632. Valvatne H, Syre H, Kross M, Stavrum R, Ti T, Phyus S, et al. Isoniazid and rifampicin resistance-associated mutations in *Mycobacterium tuberculosis* isolates from Yangon, Myanmar: implications for rapid molecular testing. *JOURNAL OF ANTIMICROBIAL CHEMOTHERAPY*. 2009 OCT;64(4):694-701. PMID: WOS:000270592400005. doi: 10.1093/jac/dkp292.
633. Dhorda M, Nyehangane D, Rénia L, Piola P, Guirin PJ, Snounou G. Transmission of *Plasmodium vivax* in South-Western Uganda: Report of Three Cases in Pregnant Women. *PLOS ONE*. 2011 MAY 13;6(5). PMID: WOS:000290558500031. doi: 10.1371/journal.pone.0019801.
634. Leitsch D, Burgess AG, Dunn LA, Krauer KG, Tan K, Duchene M, et al. Pyruvate:ferredoxin oxidoreductase and thioredoxin reductase are involved in 5-nitroimidazole activation while flavin metabolism is linked to 5-nitroimidazole resistance in *Giardia lamblia*. *J Antimicrob Chemother*. 2011 Aug;66(8):1756-65. PMID: 21602576. doi: 10.1093/jac/dkr192.
635. Massi MN, Wahyuni S, Halik H, (NITD) A, Yusuf I, Leong F, et al. Drug resistance among tuberculosis patients attending diagnostic and treatment centres in Makassar, Indonesia. *International Journal of Tuberculosis and Lung Disease*. 2011;15(4):489-95. doi: 10.5588/ijtld.09.0730.
636. Siswantoro H, Russell B, Ratcliff A, Prasetyorini B, Chalfein F, Marfurt J, et al. In Vivo and In Vitro Efficacy of Chloroquine against *Plasmodium malariae* and *P. ovale* in Papua, Indonesia. *ANTIMICROBIAL AGENTS AND CHEMOTHERAPY*. 2011 JAN;55(1):197-202. PMID: WOS:000285577400024. doi: 10.1128/AAC.01122-10.
637. Apisarnthanarak A, Hsu L-Y, Warren DK. Termination of an Extreme-Drug Resistant-*Acinetobacter baumannii* Outbreak in a Hospital After Flooding: Lessons Learned. *Clin Infect Dis*. 2012 Dec;55(11):1589-90. PMID: 22918995. doi: 10.1093/cid/cis726.
638. Brunner R, Aissaoui H, Boss C, Bozdech Z, Brun R, Corminboeuf O, et al. Identification of a New Chemical Class of Antimalarials. *JOURNAL OF INFECTIOUS DISEASES*. 2012 SEP 1;206(5):735-43. PMID: WOS:000307501800014. doi: 10.1093/infdis/jis418.
639. Cervantes S, Stout PE, Prudhomme J, Engel S, Bruton M, Cervantes M, et al. High content live cell imaging for the discovery of new antimalarial marine natural products. *BMC INFECTIOUS DISEASES*. 2012 JAN 3;12. PMID: WOS:000300066900001. doi: 10.1186/1471-2334-12-1.
640. Dunn LA, Tan KSW, Vanelle P, Juspin T, Crozet M, Terme T, et al. Development of metronidazole-resistant lines of *Blastocystis* sp. *Parasitol Res*. 2012 Jul;111(1):441-50. PMID: 22362365. doi: 10.1007/s00436-012-2860-7.
641. Guiton PS, Cusumano CK, Kline KA, Dodson KW, Han Z, Janetka JW, et al. Combinatorial small-molecule therapy prevents uropathogenic *Escherichia coli* catheter-associated urinary tract infections in mice. *Antimicrobial Agents and Chemotherapy*. 2012;56(9):4738-45. doi: 10.1128/AAC.00447-12.
642. Ho KKK, Cole N, Chen R, Willcox MD, Rice SA, Kumar N. Immobilization of antibacterial dihydropyrrol-2-ones on functional polymer supports to prevent bacterial infections in vivo. *Antimicrobial Agents and Chemotherapy*. 2012;56(2):1138-41. doi: 10.1128/AAC.05814-11.
643. Köser CU, Holden MT, Ellington MJ, Cartwright EJ, Brown NM, Ogilvy-Stuart AL, et al. Rapid whole-genome sequencing for investigation of a neonatal MRSA outbreak. *N Engl J Med*. 2012 Jun 14;366(24):2267-75. PMID: 22693998. doi: 10.1056/NEJMoa1109910.

644. Lee M, Lee J, Carroll MW, Choi H, Min S, Song T, et al. Linezolid for Treatment of Chronic Extensively Drug-Resistant Tuberculosis. *NEW ENGLAND JOURNAL OF MEDICINE*. 2012 OCT 18;367(16):1508-18. PMID: WOS:000309904500008. doi: 10.1056/NEJMoa1201964.
645. Shields RK, Press EG, Kwa AL-H, Cheng S, Du C, Clancy CJ, et al. The presence of an FKS mutation rather than MIC is an independent risk factor for failure of echinocandin therapy among patients with invasive candidiasis due to *Candida glabrata*. *Antimicrob Agents Chemother*. 2012 Sep;56(9):4862-9. PMID: 22751546. doi: 10.1128/aac.00027-12.
646. Zhang L, Chiang W-C, Gao Q, Givskov M, Tolker-Nielsen T, Yang L, et al. The catabolite repression control protein Crc plays a role in the development of antimicrobial-tolerant subpopulations in *Pseudomonas aeruginosa* biofilms. *Microbiology (United Kingdom)*. 2012;158(12):3014-9. doi: 10.1099/mic.0.061192-0.
647. Barraud N, Buson A, Jarolimek W, Rice SA. Mannitol enhances antibiotic sensitivity of persister bacteria in *Pseudomonas aeruginosa* biofilms. *PLoS One*. 2013;8(12):e84220. PMID: 24349568. doi: 10.1371/journal.pone.0084220.
648. Bowers DR, Liew YX, Lye DCB, Kwa AL-H, Hsu L-Y, Tam VH. Outcomes of Appropriate Empiric Combination versus Monotherapy for *Pseudomonas aeruginosa* Bacteremia. *ANTIMICROBIAL AGENTS AND CHEMOTHERAPY*. 2013 MAR;57(3):1270-4. PMID: WOS:000314968100022. doi: 10.1128/AAC.02235-12.
649. Chiang W-C, Nilsson M, Jensen PØ, Høiby N, Nielsen TE, Givskov M, et al. Extracellular DNA shields against aminoglycosides in *Pseudomonas aeruginosa* biofilms. *Antimicrobial Agents and Chemotherapy*. 2013;57(5):2352-61. doi: 10.1128/AAC.00001-13.
650. Ciesielczuk H, Hornsey M, Choi V, Woodford N, Wareham D. Development and evaluation of a multiplex PCR for eight plasmid-mediated quinolone-resistance determinants. *J Med Microbiol*. 2013 Dec;62(Pt 12):1823-7. PMID: 24000223. doi: 10.1099/jmm.0.064428-0.
651. Jakobsen TH, Hansen MA, Jensen PØ, Hansen L, Riber L, Cockburn A, et al. Complete Genome Sequence of the Cystic Fibrosis Pathogen *Achromobacter xylosoxidans* NH44784-1996 Complies with Important Pathogenic Phenotypes. *PLoS ONE*. 2013;8(7):e68484. doi: 10.1371/journal.pone.0068484.
652. Kelesidis T, Humphries R, Chow ALP, Tsiodras S, Uslan DZ. Emergence of daptomycin-non-susceptible enterococci urinary tract isolates. *J Med Microbiol*. 2013 Jul;62(Pt 7):1103-5. PMID: 23598376. doi: 10.1099/jmm.0.056630-0.
653. Liu Y, Knapp KM, Yang L, Molin S, Franzén H, Folkesson A. High in vitro antimicrobial activity of β-peptoid-peptide hybrid oligomers against planktonic and biofilm cultures of *Staphylococcus epidermidis*. *International Journal of Antimicrobial Agents*. 2013;41(1):20-7. doi: 10.1016/j.ijantimicag.2012.09.014.
654. Richmond GE, Chua KL, Piddock LJ. Efflux in *Acinetobacter baumannii* can be determined by measuring accumulation of H33342 (bis-benzamide). *J Antimicrob Chemother*. 2013 Jul;68(7):1594-600. PMID: 23467176. doi: 10.1093/jac/dkt052.
655. Wang H, Ciofu O, Yang L, Wu H, Song Z, Oliver A, et al. High beta-Lactamase Levels Change the Pharmacodynamics of beta-Lactam Antibiotics in *Pseudomonas aeruginosa* Biofilms. *ANTIMICROBIAL AGENTS AND CHEMOTHERAPY*. 2013 JAN;57(1):196-204. PMID: WOS:000312958400024. doi: 10.1128/AAC.01393-12.
656. Yepuri NR, Barraud N, Mohammadi NS, Kardak BG, Kjelleberg S, Rice SA, et al. Synthesis of cephalosporin-3'-diazeniumdiolates: Biofilm dispersing NO-donor prodrugs activated by β-lactamase. *Chemical Communications*. 2013;49(42):4791-3. doi: 10.1039/c3cc40869h.
657. Zhang Y, Parker DB, Snow DD, Zhou Z, Li X. Intracellular and extracellular antimicrobial resistance genes in the sludge of livestock waste management structures. *Environmental Science and Technology*. 2013;47(18):10206-13. doi: 10.1021/es401964s.
658. Zhang Y, Zhang C, Parker DB, Snow DD, Zhou Z, Li X. Occurrence of antimicrobials and antimicrobial resistance genes in beef cattle storage ponds and swine treatment lagoons. *Science of the Total Environment*. 2013;463-464:631-8. doi: 10.1016/j.scitotenv.2013.06.016.
659. Apisarnthanarak A, Hsu L-Y, Lim T-P, Mundy LM. Increase in chlorhexidine minimal inhibitory concentration of *Acinetobacter baumannii* clinical isolates after implementation of advanced source control. *Infection Control and Hospital Epidemiology*. 2014;35(1):98-9. doi: 10.1086/674404.
660. Butler J, Hooper KA, Petrie S, Lee RTC, Maurer-Stroh S, Reh L, et al. Estimating the fitness advantage conferred by permissive neuraminidase mutations in recent oseltamivir-resistant A(H1N1)pdm09 influenza viruses. *PLoS Pathog*. 2014 Apr;10(4):e1004065. PMID: 24699865. doi: 10.1371/journal.ppat.1004065.

661. Duan S, Govorkova EA, Bahl J, Zaraket H, Baranovich T, Seiler P, et al. Epistatic interactions between neuraminidase mutations facilitated the emergence of the oseltamivir-resistant H1N1 influenza viruses. *Nat Commun.* 2014 Oct 9;5:5029. PMID: 25297528. doi: 10.1038/ncomms6029.
662. Karunakaran R, Tay ST, Rahim FF, Lim BB, Puthucheary SD. Molecular Analysis of Ciprofloxacin Resistance among Non-Typhoidal *Salmonella* with Reduced Susceptibility to Ciprofloxacin Isolated from Patients at a Tertiary Care Hospital in Kuala Lumpur, Malaysia. *JAPANESE JOURNAL OF INFECTIOUS DISEASES.* 2014 MAY;67(3):157-62. PMID: WOS:000336821700002. doi: 10.7883/yoken.67.157.
663. Kelesidis T, Chow ALP. Proximity to animal or crop operations may be associated with de novo daptomycin-non-susceptible *Enterococcus* infection. *EPIDEMIOLOGY AND INFECTION.* 2014 JAN;142(1):221-4. PMID: WOS:000332523900027. doi: 10.1017/S0950268813000885.
664. Lai C-c, Lee K, Xiao Y, Ahmad N, Veeraraghavan B, Thamlikitkul V, et al. High burden of antimicrobial drug resistance in Asia. *J Glob Antimicrob Resist.* 2014 Sep;2(3):141-7. PMID: 27873720. doi: 10.1016/j.jgar.2014.02.007.
665. Landelle C, Marimuthu K, Harbarth S. Infection control measures to decrease the burden of antimicrobial resistance in the critical care setting. *Curr Opin Crit Care.* 2014 Oct;20(5):499-506. PMID: 25032821. doi: 10.1097/mcc.0000000000000126.
666. Stryjewski ME, Lentnek A, O'Riordan W, Pullman J, Tambyah PA, Miró JM, et al. A randomized Phase 2 trial of telavancin versus standard therapy in patients with uncomplicated *Staphylococcus aureus* bacteremia: the ASSURE study. *BMC INFECTIOUS DISEASES.* 2014 MAY 23;14. PMID: WOS:000336931300001. doi: 10.1186/1471-2334-14-289.
667. Veiga MI, Osório NS, Ferreira PE, Franzén O, Dahlstrom S, Lum JK, et al. Complex polymorphisms in the *Plasmodium falciparum* multidrug resistance protein 2 gene and its contribution to antimalarial response. *Antimicrob Agents Chemother.* 2014 Dec;58(12):7390-7. PMID: 25267670. doi: 10.1128/aac.03337-14.
668. Aamodt H, Mohn SC, Maselle S, Manji KP, Willems R, Jureen R, et al. Genetic relatedness and risk factor analysis of ampicillin-resistant and high-level gentamicin-resistant enterococci causing bloodstream infections in Tanzanian children. *BMC Infect Dis.* 2015 Feb 28;15:107. PMID: 25884316. doi: 10.1186/s12879-015-0845-8.
669. Baranovich T, Bahl J, Marathe BM, Culhane M, Stigger-Rosser E, Darnell D, et al. Influenza A viruses of swine circulating in the United States during 2009-2014 are susceptible to neuraminidase inhibitors but show lineage-dependent resistance to adamantanes. *Antiviral Res.* 2015 May;117:10-9. PMID: 25701593. doi: 10.1016/j.antiviral.2015.02.004.
670. Farrukee R, Leang S-K, Butler J, Lee RTC, Maurer-Stroh S, Tilmanis D, et al. Influenza viruses with B/Yamagata- and B/Victoria-like neuraminidases are differentially affected by mutations that alter antiviral susceptibility. *J Antimicrob Chemother.* 2015 Jul;70(7):2004-12. PMID: 25786478. doi: 10.1093/jac/dkv065.
671. Malmquist NA, Sundriyal S, Caron J, Chen P, Witkowski B, Menard D, et al. Histone methyltransferase inhibitors are orally bioavailable, fast-acting molecules with activity against different species causing malaria in humans. *Antimicrob Agents Chemother.* 2015 Feb;59(2):950-9. PMID: 25421480. doi: 10.1128/aac.04419-14.
672. Matsugana S, Masaoka T, Sawasaki T, Morishita R, Iwantani Y, Tatsumi M, et al. A cell-free enzymatic activity assay for the evaluation of HIV-1 drug resistance to protease inhibitors. *Front Microbiol.* 2015;6:1220. PMID: 26583013. doi: 10.3389/fmicb.2015.01220.
673. Nguyen D, Nguyen T-K, Rice SA, Boyer C. CO-Releasing Polymers Exert Antimicrobial Activity. *Biomacromolecules.* 2015 Sep 14;16(9):2776-86. PMID: 26200390. doi: 10.1021/acs.biomac.5b00716.
674. Regmi SM, Chaiprasert A, Coker OO, Disratthakit A, Pramananan T, Suriyaphol P, et al. Draft Genome Sequence of an Extensively Drug-Resistant *Mycobacterium tuberculosis* Manu-Ancestor Spoligo-International Type 523 Isolate from Thailand. *Genome Announc.* 2015 Feb 19;3(1):e01589-14. PMID: 25700413. doi: 10.1128/genomeA.01589-14.
675. Rhee S-Y, Jordan MR, Raizes E, Chua A, Parkin N, Kantor R, et al. HIV-1 Drug Resistance Mutations: Potential Applications for Point-of-Care Genotypic Resistance Testing. *PLoS One.* 2015;10(12):e0145772. PMID: 26717411. doi: 10.1371/journal.pone.0145772.
676. Soetaert K, Rens C, Wang X-M, De Bruyn J, Lanéelle M-A, Laval F, et al. Increased Vancomycin Susceptibility in Mycobacteria: a New Approach To Identify Synergistic Activity against Multidrug-Resistant Mycobacteria. *Antimicrob Agents Chemother.* 2015 Aug;59(8):5057-60. PMID: 26033733. doi: 10.1128/aac.04856-14.
677. Zowawi HM, Harris PNA, Roberts MJ, Tambyah PA, Schembri MA, Pezzani MD, et al. The emerging threat of multidrug-resistant Gram-negative bacteria in urology. *Nat Rev Urol.* 2015 Oct;12(10):570-84. PMID: 26334085. doi: 10.1038/nrurol.2015.199.

678. Arribas JR, Girard P-M, Paton NI, Winston A, Marcelin A-G, Elbirt D, et al. Efficacy of protease inhibitor monotherapy vs. triple therapy: meta-analysis of data from 2303 patients in 13 randomized trials. *HIV Med.* 2016 May;17(5):358-67. PMID: 26709605. doi: 10.1111/hiv.12348.
679. Auburn S, Serre D, Pearson RD, Amato R, Sriprawat K, To S, et al. Genomic Analysis Reveals a Common Breakpoint in Amplifications of the *Plasmodium vivax* Multidrug Resistance 1 Locus in Thailand. *J Infect Dis.* 2016 Oct 15;214(8):1235-42. PMID: 27456706. doi: 10.1093/infdis/jiw323.
680. Coker OO, Chaiprasert A, Ngamphiw C, Tongsima S, Regmi SM, Clark TG, et al. Genetic signatures of *Mycobacterium tuberculosis* Nonthaburi genotype revealed by whole genome analysis of isolates from tuberculous meningitis patients in Thailand. *PeerJ.* 2016;4:e1905. PMID: 27114869. doi: 10.7717/peerj.1905.
681. Cunningham SA, Vasoo S, Patel R. Evaluation of the Check-Points Check MDR CT103 and CT103 XL Microarray Kits by Use of Preparatory Rapid Cell Lysis. *J Clin Microbiol.* 2016 May;54(5):1368-71. PMID: 26888905. doi: 10.1128/jcm.03302-15.
682. Grigg MJ, William T, Menon J, Barber BE, Wilkes CS, Rajahram GS, et al. Efficacy of Artesunate-mefloquine for Chloroquine-resistant *Plasmodium vivax* Malaria in Malaysia: An Open-label, Randomized, Controlled Trial. *Clin Infect Dis.* 2016 Jun 1;62(11):1403-11. PMID: 27107287. doi: 10.1093/cid/ciw121.
683. Hanafi A, Lee WC, Loke MF, Teh X, Shaari A, Dinarvand M, et al. Molecular and Proteomic Analysis of Levofloxacin and Metronidazole Resistant *Helicobacter pylori*. *Front Microbiol.* 2016;7:2015. PMID: 28018334. doi: 10.3389/fmicb.2016.02015.
684. Harris RC, Khan MS, Martin LJ, Allen V, Moore DAJ, Fielding K, et al. The effect of surgery on the outcome of treatment for multidrug-resistant tuberculosis: a systematic review and meta-analysis. *BMC Infect Dis.* 2016 Jun 10;16:262. PMID: 27283524. doi: 10.1186/s12879-016-1585-0.
685. Mac Aogáin M, Mjavlovic H, Moloney G, Chotirmall SH, Rogers TR, Smith SGJ. Identification of a novel sequence type of *Escherichia coli* as the causative agent of pyelonephritis and bloodstream infection. *JMM Case Rep.* 2016 Oct;3(5):e005061. PMID: 28348784. doi: 10.1099/jmmcr.0.005061.
686. Nguyen T-K, Selvanayagam R, Ho KKK, Chen R, Kutty SK, Rice SA, et al. Co-delivery of nitric oxide and antibiotic using polymeric nanoparticles. *Chem Sci.* 2016 Feb 1;7(2):1016-27. PMID: 28808526. doi: 10.1039/c5sc02769a.
687. Nilsson M, Rybtke MT, Givskov M, Høiby N, Twetman S, Tolker-Nielsen T. The dlt genes play a role in antimicrobial tolerance of *Streptococcus mutans* biofilms. *INTERNATIONAL JOURNAL OF ANTIMICROBIAL AGENTS.* 2016 SEP;48(3):298-304. PMID: WOS:000386022000010. doi: 10.1016/j.ijantimicag.2016.06.019.
688. Phy AP, Ashley EA, Anderson TJC, Bozdech Z, Carrara VI, Sriprawat K, et al. Declining Efficacy of Artemisinin Combination Therapy Against *P. falciparum* Malaria on the Thai-Myanmar Border (2003-2013): The Role of Parasite Genetic Factors. *Clin Infect Dis.* 2016 Sep 15;63(6):784-91. PMID: 27313266. doi: 10.1093/cid/ciw388.
689. Richmond GE, Evans LP, Anderson MJ, Wand ME, Bonney LC, Ivens A, et al. The *Acinetobacter baumannii* Two-Component System AdeRS Regulates Genes Required for Multidrug Efflux, Biofilm Formation, and Virulence in a Strain-Specific Manner. *mBio.* 2016 Apr 19;7(2):e00430-16. PMID: 27094331. doi: 10.1128/mBio.00430-16.
690. Stenvang M, Dueholm MS, Vad BS, Seviour T, Zeng G, Geifman-Shochat S, et al. Epigallocatechin Gallate Remodels Overexpressed Functional Amyloids in *Pseudomonas aeruginosa* and Increases Biofilm Susceptibility to Antibiotic Treatment. *J Biol Chem.* 2016 Dec 16;291(51):26540-53. PMID: 27784787. doi: 10.1074/jbc.M116.739953.
691. Atarashi K, Suda W, Luo C, Kawaguchi T, Motoo I, Narushima S, et al. Ectopic colonization of oral bacteria in the intestine drives T(H)1 cell induction and inflammation. *Science.* 2017 Oct 20;358(6361):359-65. PMID: 29051379. doi: 10.1126/science.aan4526.
692. Basilico N, Parapini S, Sparatore A, Romeo S, Misiano P, Vivas L, et al. In Vivo and In Vitro Activities and ADME-Tox Profile of a Quinolizidine-Modified 4-Aminoquinoline: A Potent Anti-*P. falciparum* and Anti-*P. vivax* Blood-Stage Antimalarial. *Molecules.* 2017 Dec 1;22(12):2102. PMID: 29194347. doi: 10.3390/molecules22122102.
693. Bazaka K, Bazaka O, Levchenko I, Xu S, Ivanova EP, Keidar M, et al. Plasma-potentiated small molecules—possible alternative to antibiotics? *Nano Futures.* 2017;1(2). doi: 10.1088/2399-1984/aa80d3.
694. Belousoff MJ, Eyal Z, Radjainia M, Ahmed T, Bamert RS, Matzov D, et al. Structural Basis for Linezolid Binding Site Rearrangement in the *Staphylococcus aureus* Ribosome. *mBio.* 2017 May 9;8(3). PMID: 28487427. doi: 10.1128/mBio.00395-17.

695. Bhuyan GS, Hossain MA, Sarker SK, Rahat A, Islam MT, Haque TN, et al. Bacterial and viral pathogen spectra of acute respiratory infections in under-5 children in hospital settings in Dhaka city. *PLoS One*. 2017;12(3):e0174488. PMID: 28346512. doi: 10.1371/journal.pone.0174488.
696. Chang MJ, Jin B, Chae J-w, Yun H-y, Kim ES, Lee YJ, et al. Population pharmacokinetics of moxifloxacin, cycloserine, p-aminosalicylic acid and kanamycin for the treatment of multi-drug-resistant tuberculosis. *Int J Antimicrob Agents*. 2017 Jun;49(6):677-87. PMID: 28408267. doi: 10.1016/j.ijantimicag.2017.01.024.
697. Chen F, Chen G, Liu Y, Jin Y, Cheng Z, Liu Y, et al. *Pseudomonas aeruginosa* oligoribonuclease contributes to tolerance to ciprofloxacin by regulating pyocin biosynthesis. *Antimicrobial Agents and Chemotherapy*. 2017;61(3). doi: 10.1128/AAC.02256-16.
698. Cunningham SA, Limbago B, Traczewski M, Anderson K, Hackel MA, Hindler J, et al. Multicenter Performance Assessment of Carba NP Test. *J Clin Microbiol*. 2017 Jun;55(6):1954-60. PMID: 28404676. doi: 10.1128/jcm.00244-17.
699. Gowrisankar G, Chelliah R, Ramakrishnan SR, Elumalai V, Dhanamadhavan S, Brindha K, et al. Chemical, microbial and antibiotic susceptibility analyses of groundwater after a major flood event in Chennai. *Sci Data*. 2017 Oct 10;4:170135. PMID: 28994821. doi: 10.1038/sdata.2017.135.
700. Gunawan C, Marquis CP, Amal R, Sotiriou GA, Rice SA, Harry EJ. Widespread and Indiscriminate Nanosilver Use: Genuine Potential for Microbial Resistance. *ACS NANO*. 2017 APR;11(4):3438-45. PMID: WOS:000400233200004. doi: 10.1021/acsnano.7b01166.
701. Howlin RP, Cathie K, Hall-Stoodley L, Cornelius V, Duignan C, Allan RN, et al. Low-Dose Nitric Oxide as Targeted Anti-biofilm Adjunctive Therapy to Treat Chronic *Pseudomonas aeruginosa* Infection in Cystic Fibrosis. *Molecular Therapy*. 2017;25(9):2104-16. doi: 10.1016/j.ymthe.2017.06.021.
702. Hutchison C, Khan MS, Yoong J, Lin X, Coker RJ. Financial barriers and coping strategies: a qualitative study of accessing multidrug-resistant tuberculosis and tuberculosis care in Yunnan, China. *BMC Public Health*. 2017 Feb 22;17(1):221. PMID: 28222724. doi: 10.1186/s12889-017-4089-y.
703. Kathirvel S, Tripathy JP, Tun ZM, Patro BK, Singh T, Bhalla A, et al. Physicians' compliance with the National Drug Policy on Malaria in a tertiary teaching hospital, India, from 2010 to 2015: a mixed method study. *Trans R Soc Trop Med Hyg*. 2017 Feb 1;111(2):62-70. PMID: 28460016. doi: 10.1093/trstmh/trx020.
704. Lamoth F, Chung SJ, Damonti L, Alexander BD. Changing Epidemiology of Invasive Mold Infections in Patients Receiving Azole Prophylaxis. *Clin Infect Dis*. 2017 Jun 1;64(11):1619-21. PMID: 28199491. doi: 10.1093/cid/cix130.
705. Landier J, Kajeechiwa L, Thwin MM, Parker DM, Chaumeau V, Wiladphaingern J, et al. Safety and effectiveness of mass drug administration to accelerate elimination of artemisinin-resistant falciparum malaria: A pilot trial in four villages of Eastern Myanmar. *Wellcome Open Res*. 2017;2:81. PMID: 29062913. doi: 10.12688/wellcomeopenres.12240.1.
706. Liao J-H, Tsai C-H, Patel SG, Yang J-T, Tu I-F, Lo Cicero M, et al. Acetylome of *Acinetobacter baumannii* SK17 Reveals a Highly-Conserved Modification of Histone-Like Protein HU. *Front Mol Biosci*. 2017;4:77. PMID: 29230394. doi: 10.3389/fmolb.2017.00077.
707. Thai VC, Lim TK, Le KPU, Lin Q, Nguyen TTH. iTRAQ-based proteome analysis of fluoroquinolone-resistant *Staphylococcus aureus*. *J Glob Antimicrob Resist*. 2017 Mar;8:82-9. PMID: 28039103. doi: 10.1016/j.jgar.2016.11.003.
708. Ahmed W, Zhang Q, Lobos A, Senkbeil J, Sadowsky MJ, Harwood VJ, et al. Precipitation influences pathogenic bacteria and antibiotic resistance gene abundance in storm drain outfalls in coastal sub-tropical waters. *Environ Int*. 2018 Jul;116:308-18. PMID: 29754026. doi: 10.1016/j.envint.2018.04.005.
709. Antonoplis A, Zang X, Huttner MA, Chong KKL, Lee YB, Co JY, et al. A Dual-Function Antibiotic-Transporter Conjugate Exhibits Superior Activity in Sterilizing MRSA Biofilms and Killing Persister Cells. *J Am Chem Soc*. 2018 Nov 28;140(47):16140-51. PMID: 30388366. doi: 10.1021/jacs.8b08711.
710. Beattie RE, Walsh M, Cruz MC, McAliley LR, Dodgen L, Zheng WU-e, et al. Agricultural contamination impacts antibiotic resistance gene abundances in river bed sediment temporally. *FEMS Microbiol Ecol*. 2018 Sep 1;94(9). PMID: 30010841. doi: 10.1093/femsec/fiy131.
711. Chen C, Gardete S, Jansen RS, Shetty A, Dick T, Rhee KY, et al. Verapamil Targets Membrane Energetics in *Mycobacterium tuberculosis*. *Antimicrob Agents Chemother*. 2018 May;62(5). PMID: 29463541. doi: 10.1128/aac.02107-17.
712. Allix-Béguec C, Arandjelovic I, Bi L, Beckert P, Bonnet M, Bradley P, et al. Prediction of Susceptibility to First-Line Tuberculosis Drugs by DNA Sequencing. *NEW ENGLAND JOURNAL OF MEDICINE*. 2018 OCT 11;379(15):1403-15. PMID: WOS:000446923400005. doi: 10.1056/NEJMoa1800474.

713. Dunn DT, Stöhr W, Arenas-Pinto A, Tostevin A, Mbisa JL, Paton NI. Next generation sequencing of HIV-1 protease in the PIVOT trial of protease inhibitor monotherapy. *J Clin Virol.* 2018 Apr;101:63-5. PMID: 29428459. doi: 10.1016/j.jcv.2018.02.003.
714. Fang T, Wang HC-e, Cui Q, Rogers M, Dong P. Diversity of potential antibiotic-resistant bacterial pathogens and the effect of suspended particles on the spread of antibiotic resistance in urban recreational water. *Water Res.* 2018 Nov 15;145:541-51. PMID: 30199799. doi: 10.1016/j.watres.2018.08.042.
715. Germond A, Ichimura T, Horinouchi T, Fujita H, Furusawa C, Watanabe TM. Raman spectral signature reflects transcriptomic features of antibiotic resistance in *Escherichia coli*. *Commun Biol.* 2018;1:85. PMID: 30271966. doi: 10.1038/s42003-018-0093-8.
716. Grigg MJ, William T, Piera KA, Rajahram GS, Jelip J, Aziz A, et al. Plasmodium falciparum artemisinin resistance monitoring in Sabah, Malaysia: in vivo therapeutic efficacy and kelch13 molecular marker surveillance. *Malar J.* 2018 Dec 10;17(1):463. PMID: 30526613. doi: 10.1186/s12936-018-2593-x.
717. Hoppe A, Giuliano M, Lugemwa A, Thompson JA, Floridia M, Walker AS, et al. HIV-1 viral load and resistance in genital secretions in patients taking protease-inhibitor-based second-line therapy in Africa. *Antivir Ther.* 2018;23(2):191-5. PMID: 29021409. doi: 10.3851/imp3200.
718. Jiang Y, Xu C, Wu X, Chen Y, Han W, Gin KY-H, et al. Occurrence, seasonal variation and risk assessment of antibiotics in Qingcaosha reservoir. *Water (Switzerland).* 2018;10(2). doi: 10.3390/w10020115.
719. Kano R, Hsiao Y-H, Han HS, Chen C, Hasegawa A, Kamata H. Resistance Mechanism in a Terbinafine-Resistant Strain of *Microsporum canis*. *Mycopathologia.* 2018 Jun;183(3):623-7. PMID: 29340910. doi: 10.1007/s11046-018-0242-0.
720. Malkawi R, Iyer A, Parmar A, Lloyd DG, Goh ETL, Taylor EJ, et al. Cysteines and Disulfide-Bridged Macroyclic Mimics of Teixobactin Analogues and Their Antibacterial Activity Evaluation against Methicillin-Resistant *Staphylococcus aureus* (MRSA). *Pharmaceutics.* 2018 Oct 11;10(4):183. PMID: 30314324. doi: 10.3390/pharmaceutics10040183.
721. Mather AE, Phuong TLT, Gao Y, Clare S, Mukhopadhyay S, Goulding DA, et al. New Variant of Multidrug-Resistant *Salmonella enterica* Serovar Typhimurium Associated with Invasive Disease in Immunocompromised Patients in Vietnam. *mBio.* 2018 Sep 4;9(5):e01056-18. PMID: 30181247. doi: 10.1128/mBio.01056-18.
722. Merchant S, Proudfoot EM, Quadri HN, McElroy HJ, Wright WR, Gupta A, et al. Risk factors for *Pseudomonas aeruginosa* infections in Asia-Pacific and consequences of inappropriate initial antimicrobial therapy: A systematic literature review and meta-analysis. *J Glob Antimicrob Resist.* 2018 Sep;14:33-44. PMID: 29454906. doi: 10.1016/j.jgar.2018.02.005.
723. Nordström R, Nyström L, Andrén OCJ, Malkoch M, Umerska A, Davoudi M, et al. Membrane interactions of microgels as carriers of antimicrobial peptides. *J Colloid Interface Sci.* 2018 Mar 1;513:141-50. PMID: 29145017. doi: 10.1016/j.jcis.2017.11.014.
724. Oonsivilai M, Mo Y, Luangasanatip N, Lubell Y, Miliya T, Tan P, et al. Using machine learning to guide targeted and locally-tailored empiric antibiotic prescribing in a children's hospital in Cambodia. *Wellcome Open Res.* 2018;3:131. PMID: 30756093. doi: 10.12688/wellcomeopenres.14847.1.
725. Ravensdale JT, Xian DTW, Wei CM, Lv Q, Wen X, Guo J, et al. PCR screening of antimicrobial resistance genes in faecal samples from Australian and Chinese children. *J Glob Antimicrob Resist.* 2018 Sep;14:178-81. PMID: 29614373. doi: 10.1016/j.jgar.2018.03.003.
726. Stockdale AJ, Saunders MJ, Boyd MA, Bonnett LJ, Johnston V, Wandeler G, et al. Effectiveness of Protease Inhibitor/Nucleos(t)ide Reverse Transcriptase Inhibitor-Based Second-line Antiretroviral Therapy for the Treatment of Human Immunodeficiency Virus Type 1 Infection in Sub-Saharan Africa: A Systematic Review and Meta-analysis. *Clin Infect Dis.* 2018 Jun 1;66(12):1846-57. PMID: 29272346. doi: 10.1093/cid/cix1108.
727. Subedi D, Vijay AK, Kohli GS, Rice SA, Willcox MDP. Association between possession of ExoU and antibiotic resistance in *Pseudomonas aeruginosa*. *PLoS One.* 2018;13(9):e0204936. PMID: 30265709. doi: 10.1371/journal.pone.0204936.
728. Subedi D, Vijay AK, Kohli GS, Rice SA, Willcox MDP. Comparative genomics of clinical strains of *Pseudomonas aeruginosa* strains isolated from different geographic sites. *Sci Rep.* 2018 Oct 23;8(1):15668. PMID: 30353070. doi: 10.1038/s41598-018-34020-7.
729. Subedi D, Vijay AK, Kohli GS, Rice SA, Willcox MDP. Nucleotide sequence analysis of NPS-1 β-lactamase and a novel integron (In1427)-carrying transposon in an MDR *Pseudomonas aeruginosa* keratitis strain. *J Antimicrob Chemother.* 2018 Jun 1;73(6):1724-6. PMID: 29547943. doi: 10.1093/jac/dky073.

730. Tzou PL, Ariyaratne P, Varghese V, Lee C, Rakhmanaliev E, Villy C, et al. Comparison of an In Vitro Diagnostic Next-Generation Sequencing Assay with Sanger Sequencing for HIV-1 Genotypic Resistance Testing. *J Clin Microbiol*. 2018 Jun;56(6). PMID: 29618499. doi: 10.1128/jcm.00105-18.
731. Vente A, Bentley C, Lückermann M, Tambyah PA, Dalhoff A. Early Clinical Assessment of the Antimicrobial Activity of Finafloxacin Compared to Ciprofloxacin in Subsets of Microbiologically Characterized Isolates. *Antimicrob Agents Chemother*. 2018 Apr;62(4). PMID: 29339393. doi: 10.1128/aac.02325-17.
732. Zhou C, Song H, Loh CJL, She J, Deng L, Liu B. Grafting antibiofilm polymer hydrogel film onto catheter by SARA SI-ATRP. *J Biomater Sci Polym Ed*. 2018 Dec;29(17):2106-23. PMID: 30141743. doi: 10.1080/09205063.2018.1507268.
733. Brunton LA, Desbois AP, Garza M, Wieland B, Mohan CV, Hässler B, et al. Identifying hotspots for antibiotic resistance emergence and selection, and elucidating pathways to human exposure: Application of a systems-thinking approach to aquaculture systems. *Sci Total Environ*. 2019 Oct 15;687:1344-56. PMID: 31412468. doi: 10.1016/j.scitotenv.2019.06.134.
734. Çapçı A, Lorion MM, Wang HD, Simon N, Leidenberger M, Silva MCB, et al. Artemisinin-(Iso)quinoline Hybrids by C-H Activation and Click Chemistry: Combating Multidrug-Resistant Malaria. *Angew Chem Int Ed Engl*. 2019 Sep 9;58(37):13066-79. PMID: 31290221. doi: 10.1002/anie.201907224.
735. Chen Y, Li P, Huang Y, Yu K, Chen H-J, Cui K, et al. Environmental media exert a bottleneck in driving the dynamics of antibiotic resistance genes in modern aquatic environment. *Water Res*. 2019 Oct 1;162:127-38. PMID: 31260828. doi: 10.1016/j.watres.2019.06.047.
736. Chen Y, Su J-Q, Zhang J, Li P, Chen H, Zhang B, et al. High-throughput profiling of antibiotic resistance gene dynamic in a drinking water river-reservoir system. *Water Res*. 2019 Feb 1;149:179-89. PMID: 30447523. doi: 10.1016/j.watres.2018.11.007.
737. Faksri K, Kaewprasert O, Ong RT-H, Suriyaphol P, Prammananan T, Teo Y-Y, et al. Comparisons of whole-genome sequencing and phenotypic drug susceptibility testing for *Mycobacterium tuberculosis* causing MDR-TB and XDR-TB in Thailand. *Int J Antimicrob Agents*. 2019 Aug;54(2):109-16. PMID: 30981926. doi: 10.1016/j.ijantimicag.2019.04.004.
738. González A, Casado J, Chueca E, Salillas S, Velázquez-Campoy A, Angarica VE, et al. Repurposing Dihydropyridines for Treatment of *Helicobacter pylori* Infection. *Pharmaceutics*. 2019 Dec 15;11(12). PMID: 31847484. doi: 10.3390/pharmaceutics11120681.
739. González A, Salillas S, Velázquez-Campoy A, Angarica VE, Fillat MF, Sancho J, et al. Identifying potential novel drugs against *Helicobacter pylori* by targeting the essential response regulator HsrA. *Sci Rep*. 2019 Aug 5;9(1):11294. PMID: 31383920. doi: 10.1038/s41598-019-47746-9.
740. Jabbar A, Phelan JE, de Sessions PF, Khan TA, Rahman H, Khan SN, et al. Whole genome sequencing of drug resistant *Mycobacterium tuberculosis* isolates from a high burden tuberculosis region of North West Pakistan. *Sci Rep*. 2019 Oct 18;9(1):14996. PMID: 31628383. doi: 10.1038/s41598-019-51562-6.
741. Juhas M, Widlake E, Teo JWP, Huseby DL, Tyrrell JM, Polikanov YS, et al. In vitro activity of apramycin against multidrug-, carbapenem- and aminoglycoside-resistant Enterobacteriaceae and *Acinetobacter baumannii*. *J Antimicrob Chemother*. 2019 Apr 1;74(4):944-52. PMID: 30629184. doi: 10.1093/jac/dky546.
742. Li H, Andersen PS, Stegger M, Sieber RN, Ingmer H, Staubrand N, et al. Antimicrobial Resistance and Virulence Gene Profiles of Methicillin-Resistant and -Susceptible *Staphylococcus aureus* From Food Products in Denmark. *Front Microbiol*. 2019;10:2681. PMID: 31920996. doi: 10.3389/fmicb.2019.02681.
743. Li H, Stegger M, Dalsgaard A, Leisner JJ. Bacterial content and characterization of antibiotic resistant *Staphylococcus aureus* in Danish sushi products and association with food inspector rankings. *Int J Food Microbiol*. 2019 Sep 16;305:108244. PMID: 31202150. doi: 10.1016/j.ijfoodmicro.2019.108244.
744. Limmathurotsakul D, Sandoe JAT, Barrett DC, Corley M, Hsu L-Y, Mendelson M, et al. 'Antibiotic footprint' as a communication tool to aid reduction of antibiotic consumption. *J Antimicrob Chemother*. 2019 Aug 1;74(8):2122-7. PMID: 31074489. doi: 10.1093/jac/dkz185.
745. Long S, Miao L, Li R, Deng F, Qiao Q, Liu X, et al. Rapid identification of bacteria by membrane-responsive aggregation of a pyrene derivative. *ACS Sensors*. 2019;4(2):281-5. doi: 10.1021/acssensors.8b01466.
746. Nilsson M, Givskov M, Twetman S, Tolker-Nielsen T. Inactivation of the *pgmA* Gene in *Streptococcus mutans* Significantly Decreases Biofilm-Associated Antimicrobial Tolerance. *Microorganisms*. 2019 Sep 3;7(9). PMID: 31484288. doi: 10.3390/microorganisms7090310.

747. Nilsson M, Jakobsen TH, Givskov M, Twetman S, Tolker-Nielsen T. Oxidative stress response plays a role in antibiotic tolerance of *Streptococcus mutans* biofilms. *Microbiology (Reading)*. 2019 Mar;165(3):334-42. PMID: 30663959. doi: 10.1099/mic.0.000773.
748. Pei M, Zhang B, He Y, Su J-Q, Gin KY-H, Lev O, et al. State of the art of tertiary treatment technologies for controlling antibiotic resistance in wastewater treatment plants. *Environ Int*. 2019 Oct;131:105026. PMID: 31351383. doi: 10.1016/j.envint.2019.105026.
749. Penesyan A, Nagy SS, Kjelleberg S, Gillings MR, Paulsen IT. Rapid microevolution of biofilm cells in response to antibiotics. *NPJ Biofilms Microbiomes*. 2019;5(1):34. PMID: 31728201. doi: 10.1038/s41522-019-0108-3.
750. Phelan JE, Lim DR, Mitarai S, de Sessions PF, Tujan MAA, Reyes LT, et al. *Mycobacterium tuberculosis* whole genome sequencing provides insights into the Manila strain and drug-resistance mutations in the Philippines. *Sci Rep*. 2019 Jun 26;9(1):9305. PMID: 31243306. doi: 10.1038/s41598-019-45566-5.
751. Ram M R, Teh X, Rajakumar T, Goh KL, Leow AHR, Poh BH, et al. Polymorphisms in the host CYP2C19 gene and antibiotic-resistance attributes of *Helicobacter pylori* isolates influence the outcome of triple therapy. *J Antimicrob Chemother*. 2019 Jan 1;74(1):11-6. PMID: 30403784. doi: 10.1093/jac/dky401.
752. Safi H, Gopal P, Lingaraju S, Ma S, Levine C, Dartois V, et al. Phase variation in *Mycobacterium tuberculosis* glpK produces transiently heritable drug tolerance. *Proc Natl Acad Sci U S A*. 2019 Sep 24;116(39):19665-74. PMID: 31488707. doi: 10.1073/pnas.1907631116.
753. Sosibo SC, Somboro AM, Amoako DG, Sekyere JO, Bester LA, Ngila JC, et al. Impact of Pyridyl Moieties on the Inhibitory Properties of Prominent Acyclic Metal Chelators Against Metallo- β -Lactamase-Producing Enterobacteriaceae: Investigating the Molecular Basis of Acyclic Metal Chelators' Activity. *Microb Drug Resist*. 2019 Apr;25(3):439-49. PMID: 30741600. doi: 10.1089/mdr.2018.0272.
754. Subedi D, Kohli GS, Vijay AK, Willcox MDP, Rice SA. Accessory genome of the multi-drug resistant ocular isolate of *Pseudomonas aeruginosa* PA34. *PLoS One*. 2019;14(4):e0215038. PMID: 30986237. doi: 10.1371/journal.pone.0215038.
755. Thompson JA, Kityo CM, Dunn DT, Hoppe A, Ndashimye E, Hakim J, et al. Evolution of Protease Inhibitor Resistance in Human Immunodeficiency Virus Type 1 Infected Patients Failing Protease Inhibitor Monotherapy as Second-line Therapy in Low-income Countries: An Observational Analysis Within the EARNEST Randomized Trial. *Clin Infect Dis*. 2019 Mar 19;68(7):1184-92. PMID: 30060027. doi: 10.1093/cid/ciy589.
756. Yang DL, Hu YL, Yin ZX, Zeng G, Li D, Zhang YQ, et al. Cis-2-dodecenoic Acid Mediates Its Synergistic Effect with Triazoles by Interfering with Efflux Pumps in Fluconazole-resistant *Candida albicans*. *Biomed Environ Sci*. 2019 Mar;32(3):199-209. PMID: 30987694. doi: 10.3967/bes2019.027.
757. Zhang N, Liu X, Liu R, Zhang T, Li M, Zhang Z, et al. Influence of reclaimed water discharge on the dissemination and relationships of sulfonamide, sulfonamide resistance genes along the Chaobai River, Beijing. *FRONTIERS OF ENVIRONMENTAL SCIENCE & ENGINEERING*. 2019 FEB;13(1). PMID: WOS:000454562300001. doi: 10.1007/s11783-019-1099-2.