

Systematic review of the effects of antimicrobial cycling on bacterial resistance rates within hospital settings

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Abstract

Aim Antimicrobial resistance is an evolving phenomenon with alarming public health consequences. Antibiotic cycling is a widely known antimicrobial stewardship initiative which encompasses periodical shifts in empirical treatment protocols with the aim to limit selective pressures on bacterial populations. Nonetheless, mathematical models have challenged its presumable efficacy by favouring a higher heterogeneity in antibiotic administration. We present a review of the evidence regarding the actual impact of antimicrobial cycling on bacterial resistance control within hospitals. **Methods** A systematic literature review was conducted using the PubMed/MedLine, Embase, CINAHL Plus and Global Health databases. **Results** A systematic search process retrieved a sole randomised study, and so we broadened inclusion criteria to encompass quasi-experimental designs. Fifteen studies formed our dataset including seven prospective trials and eight before-and-after studies. Nine studies evaluated cycling versus a control group and produced conflicting results whilst three studies compared cycling with antibiotic mixing, with none of the strategies appearing superior. The rest evaluated resistance dynamics of each of the on-cycle antibiotics with contradictory findings. Research protocols differed in parameters such as the cycle length, the choice of antibiotics, the opportunity to de-escalate to narrow-spectrum agents and the measurement of indicators of collateral damage. This limited our ability to evaluate the replicability of findings and the overall policy effects. **Conclusions** Dearth of robust designs and standardised protocols limits our ability to reach safe conclusions. Nonetheless, in view of the available data we find no reason to believe that cycling should be expected to improve antibiotic resistance rates within hospitals.

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Running head: Antimicrobial cycling and resistance control

Keywords: Antimicrobial stewardship, antibiotic cycling, antibiotic rotation, bacterial resistance

Abstract

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Methods

A systematic literature review was conducted using the PubMed/MedLine, Embase, CINAHL Plus and Global Health databases.

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A systematic search process retrieved a sole randomised study, and so we broadened inclusion criteria to encompass quasi-experimental designs. Fifteen studies formed our dataset including seven prospective trials and eight before-and-after studies. Nine studies evaluated cycling versus a control group and produced conflicting results whilst three studies compared cycling with antibiotic mixing, with none of the strategies appearing superior. The rest evaluated resistance dynamics of each of the on-cycle antibiotics with contradictory findings. Research protocols differed in parameters such as the cycle length, the choice of antibiotics, the opportunity to de-escalate to narrow-spectrum agents and the measurement of indicators of collateral damage. This limited our ability to evaluate the replicability of findings and the overall policy effects.

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Introduction

Evolving bacterial resistance to antimicrobial agents, one of the ten most critical public health threats according to the World Health Organization, demands immediate action[1]. Antimicrobial cycling or rotation is among the multitude of initiatives tried to streamline antibiotic prescribing, and fall within the umbrella term of antimicrobial stewardship. Cycling or rotation incurs scheduled shifts in empirical antibiotic treatment protocols, switching periodically between antimicrobial agents of similar spectrum. This practice is often adopted in high-risk settings such as Intensive Care Units and relies more or less on an intuitive perception that such scheduled rotations of antimicrobial agents could alter selective pressures on bacterial populations accordingly and thus stem the onset of resistant strains. The concept was probably further developed in the 90's when Gerding et al reported improvements in aminoglycoside resistance rates as a result of changes in the type of predominant aminoglycoside use[2][3].

However, mathematical models have challenged the strategy's presumable effectiveness by predicting that interventions which favoured a more heterogeneous antimicrobial use would be more successful in bacterial resistance control[4][5][6]. According to a 2006 systematic literature review very few studies met inclusion criteria and lack of rigorousness in study designs for those finally included was insufficient to draw safe inferences[7] A meta-analysis following almost ten years later suggested potential benefits by the application of the particular strategy without, however, performing an in-depth evaluation of the available studies[8].

We aim to provide an updated review and evaluation of the evidence with regard to the efficacy of antimicrobial cycling on bacterial resistance control within hospital settings. Our study is a composite element of a wider project with the objective to assess the impact of different antimicrobial stewardship initiatives on bacterial resistance rates which has led to the publication of two additional papers discussing the role of antimicrobial restrictions[9] and prospective audit with feedback[10].

Methods

A systematic literature review was conducted using the PubMed/MedLine, CINAHL Plus, Global Health and Embase databases. We sought to retrieve all studies of reasonable quality which assessed the impact of antimicrobial cycling strategies on bacterial resistance within clinical settings. We also recorded antimicrobial consumption and morbidity and/or mortality rates as secondary outcomes for a more thorough assessment of the observed results.

As this study was part of a wider project we designed a broad search algorithm on the basis of definitions provided by major organizations: Infectious Diseases Society of America (IDSA), Center for Disease Preven-

tion and Control (CDC)[11][12]. The search string covered three concepts, antimicrobial stewardship and its constituent strategies, antimicrobial resistance, and the hospital setting of the interventions:

1. (antimicrobial stewardship) OR (antibiotic stewardship) OR (audit “and” feedback) OR (restriction) OR (pre?authorization) OR (antibiotic combination*) OR (antimicrobial combination*) OR (antibiotic cycling) OR (antimicrobial cycling) OR (antibiotic rotation) OR (antimicrobial rotation) OR (antibiotic time?out*) OR (antimicrobial time?out*) OR (dose adjustment) OR (dose optimi#ation) OR (antibiotic mixing) OR (antimicrobial mixing) OR (antibiotic de?escalation) OR (antimicrobial de?escalation) OR (parenteral oral conversion) OR (intravenous oral conversion) OR (procalcitonin) OR (electronic alert*) OR (electronic system*) OR (computeri#ed alert*) OR (computeri#ed system*) OR (automat* stop order*)
2. Exp Drug Utilization
3. 1 OR 2
4. (antibiotic resistan*) OR (antimicrobial resistan*) OR (multi?drug resistan*) OR (bacterial resistan*) OR (bacterial susceptib*) OR (susceptib* phenotype*) OR (antibiotic susceptib*) OR (antimicrobial susceptib*)
5. 3 AND 4
6. (nosocomial OR hospital* OR in?patient OR intensive care OR ICU*)
7. 5 AND 6

8,922 papers covering the period to 1st April 2020 were screened for relevance. Randomised studies were scarce and for this reason we decided to broaden inclusion criteria by considering quasi-experimental designs. However, we excluded simple before-and-after studies which examined cohorts lasting less than one year, to minimise confounding due to seasonality and to facilitate comparability of results. We also excluded studies which combined changes in infection control practices or applied multidisciplinary interventions due to confounding and constraints on comparability. Studies which lacked historical or parallel cohorts for comparison were not included as interpretation is impossible without some kind of internal control or comparator. Data provided by grey literature such as congress papers and reports from governmental and non-governmental organizations were outside our scope due to lack of peer review. Finally, studies which did not apply suitable statistical methods to evaluate the significance of the reported results were also excluded.

A main distinction from prior meta-research on the topic is the fact that we considered changes in infection control as well as the application of additional antimicrobial stewardship interventions as important confounding factors which should not be overlooked; this led to the exclusion of several papers which other reviews have included.

Results

Study selection turned out to be challenging. This was partly due to the characteristics of the intervention studied which preclude blinding and limit options for even partial randomization. The search process retrieved a sole cluster randomised trial and for this reason we decided to include less robust study types and perform a sensitivity analysis in case of conflicting results.

Fifteen relevant studies formed our dataset including seven prospective trials and eight simple before-and-after studies (Table 1). Nine studies evaluated the effects of antibiotic cycling versus a control group[13][14][15][16][17][18][19][20][21]. Three papers compared antimicrobial cycling with antibiotic mixing[22][23][24], that is administering the scheduled antimicrobial agents on a successive patient basis. The last three assessed the resistance potential of each of the alternating on-cycle antibiotics, that is the variations in risk of antibiotic resistant infection and/or colonization during cycles of different predominant antibiotic use[25][26][27].

Fixed durations of each cycle ranged from one week to eight months. The rotating agents were piperacillin-tazobactam with cefepime in two cases[13][25], fluoroquinolones with beta lactams in three cases[18][26][27]. The rest rotated the aforementioned agents with carbapenems and aminoglycosides in varying combinations. In some protocols de-escalation to suitable narrow-spectrum agents was permitted but in others it was not,

with six teams proceeding to de-escalation in view of bacterial susceptibility results[16][17][19][23][24][27], five teams avoiding de-escalation to increase the on-cycle antimicrobial use[14][15][18][21][26] and four teams not clarifying their practices enough for their readers to be able to ascertain specifically what they did[13][20][22][25]. Four studies provided bacterial typing data to assist in the evaluation of cross-transmission dynamics[14][18][25][27]. Furthermore, methodologies differed as to whether surveillance cultures or cultures from clinically presumed infections, unit-wide or patient-specific, were recorded as indicators of resistance incidence.

Among those studies which compared an experimental with a control cohort there were seven simple before-and-after and two prospective trials. Seven of these provided data with regard to antimicrobial protocols in the control group[14][15][16][18][19][20][21] and two did not set out their standard practice[13][17]. Oddly, many studies fail to state any explicit goal of their chosen intervention, but the available information suggests that the institution of an antimicrobial rotation policy aimed to increase heterogeneity of antimicrobial administration in the intervention group by utilising more antimicrobial classes of similar spectrum in a scheduled fashion. The results, however, appear rather conflicting.

In particular, if one takes into account bacterial susceptibilities to the rotated agents which are apparently a more straightforward indicator of the policy's effectiveness four studies did not achieve any measurable success and five reported variable improvement (Table 1). The most noteworthy study in the group reporting negative findings is probably the trial conducted by Toltzis et al. Its main distinctive feature is the use of a contemporary control group, and its use of bacterial typing data facilitates interpretation of the available findings. The researchers observed no benefits even when only clonally discordant isolates were taken into account[14].

The group reporting positive findings encompassed two studies which observed an increase in *P. aeruginosa* susceptibility to one and two of the rotated agents respectively[17][18] and two studies which reported improvements in Extended-spectrum Beta Lactamase (ESBL) incidence ($p < 0.05$)[20][21]. One of the latter used a rather small sample while none of the aforementioned seemingly successful studies utilized bacterial typing. Thus, the possibility that the observed findings could be a result of horizontal transfer of bacterial clones due to breaks in infection control cannot be excluded as in the study conducted by Toltzis et al.

Nijssen et al reported lower colonization rates for ciprofloxacin-resistant isolates in the intervention group but no changes for cephalosporin-resistant isolates[18]. Highly homogeneous prescription of fluoroquinolones in the control arm, a radical reduction in ciprofloxacin administration in the intervention arm along with the main mechanism of fluoroquinolone resistance which incurs spontaneous chromosomal mutations favoured by increased selective pressures could perhaps explain the observed results, but no firm interpretation is possible.

Frequency of cycling did not appear to be associated with the possibility of positive or inconclusive outcomes as it varied widely in both groups. Furthermore, the fact that universal lack of randomization and blinding would potentially predispose to some degree of selection and information bias in favour of more positive outcomes, and while no specific biases were evident, this inevitable contextual bias should be taken into account.

Three studies assessed antimicrobial rotation compared to administering the agents on a successive patient basis to maximise antibiotic heterogeneity, a practice known as antibiotic mixing. Two of those, including one using the robust cluster-randomised cross-over design, observed no significant differences[23][24]. Jayashree et al reported lower resistance rates in both cycling and mixing periods compared to a three-month baseline period. The latter, however, was too short to be informative[24]. The third reported higher cefepime susceptibility rates for *P. aeruginosa* during cycling ($p = 0.01$) but no further improvements[22]. De-escalation as well as combination therapy were permitted in two instances[23][24], and their allowability was not clarified in the third[22]. None of the teams used typing data to assess cross-transmission dynamics.

As for the remaining studies, Ginn et al cycled piperacillin-tazobactam with cefepime and found that cefepime showed as a more important driver for the onset of bacterial resistance with the proportion of ad-

missions complicated by resistant infections during cefepime cycles being more than twice as high compared to piperacillin-tazobactam cycles ($p < 0.001$) [25]. Van Loon et al cycled levofloxacin with ceftazidime and piperacillin-tazobactam, concluding that levofloxacin use was associated with higher levofloxacin-resistance rates, but ceftazidime was seemingly not prone to the selection of ceftazidime-resistant strains [26]. Tsukayama et al rotated fluoroquinolones with piperacillin-tazobactam but did not find any significant correlations between the on-cycle antibiotic class and the probability of resistance onset. However, the authors report high use of off-cycle antibiotics which could potentially act as a confounding factor [27].

Finally, all but two studies provided some data regarding the on- and off-cycle antimicrobial consumption during the experimental period, while seven studies measured variable side effects as indicators of the policy's potential collateral damage including morbidity and/or mortality rates reported by six studies [15] [16] [19] [22] [23] [24]. None of these recorded worrying trends in intervention groups.

Discussion

Bacterial resistance to antimicrobial agents is an incessantly evolving phenomenon which threatens one of the greatest achievements of medical science, the effective treatment of infectious diseases. Overprescribing and suboptimal selection of antimicrobial agents are believed to have contributed to the acceleration of the selection of resistant strains. Thus antimicrobial stewardship has provoked the interest of the medical community as a multifaceted set of interventions which aim to optimise antimicrobial use and thus stem the onset of resistant bacterial strains.

Despite, however, the public health importance of this issue, there is a notable lack of standardised high-quality research on the field to provide definite answers as to which, if any, initiatives are effective or not. We have already examined antimicrobial restrictions and audit with feedback in two papers that were recently published [9] [10]. The absence of randomised models and the great heterogeneity in study protocols limited the ability to draw any firm conclusions on the aspects researched. It highlights the need for future high-quality, reproducible research. Standardisation in study design would increase the utility of clinical research in this field, as meta-synthesis of studies would be possible, providing greater statistical power to detect and map the effects of intervening to try to reduce resistance, and guide clinicians.

Examination of the available literature on the potential efficacy of antimicrobial cycling gives an overall impression of rather limited success. Research papers could be roughly divided to those which evaluated cycling versus a control group and produced conflicting results and those that compared cycling with mixing with none of the strategies appearing superior to the other. Lack of success becomes more evident if one takes into account the most rigorous studies conducted by Toltzis et al [14] as well as Van Duijn et al [23] both of which failed to record any favourable results comparing cycling with a control group and a mixing group respectively.

Fair interpretation of the relevant data must take into account some core limitations which could influence results in either way. One such limitation is the lack of standardization of antibiotic protocols across intervention and control groups of different studies, though a general tendency to increase heterogeneity of antibiotic administration in the experimental arms was observable. It is rational to assume that the relevant baseline practices would influence whether significant changes in antibiotic resistance patterns would be recorded post-intervention. A pertinent paradigm is probably provided by Nijssen et al who compared antibiotic rotation with a control group receiving fluoroquinolones in a highly homogeneous manner. Fluoroquinolone resistance rates were decreased in the rotation arm, a trend not seen for cephalosporins. It is well-known that the main mechanism of fluoroquinolone resistance comprises point mutations in chromosomal DNA which are obviously particularly prone to selective pressures. Radical reduction in fluoroquinolone administration along with the main relevant mechanism of resistance could provide a likely explanation for the observed results further supported in the clinical literature after the application of restrictive fluoroquinolone strategies [9].

We cannot exclude the possibility that the potential of success could be pathogen-specific and depending on the monitoring protocol it could be potentially missed; a pathogen-specific effect has indeed been suggested

by researchers in the past[8]. It is true that the majority of the available positive findings in our dataset relate to *P. aeruginosa* although we are not aware of any pathophysiological mechanism that would account for such a theory.

Failure of antibiotic cycling to produce clear benefits is consistent with the theoretical predictions generated by many mathematical models that challenge its intuitively presumed efficacy. On the basis of the aforementioned models, though, one would expect that antibiotic mixing would be more effective via maximising heterogeneous antimicrobial use. Neither assumption was confirmed in practice. Although there is high variability in research protocols and the overall quality of our data is far from satisfying to reach definite conclusions, we should bear in mind that the evolution of bacterial resistance is a complex process and the strategies tested may rely on an oversimplified model of how it may be manipulated. It is worth mentioning that antimicrobial agents of similar spectrum may possess totally different mechanisms of action, and thus may affect bacteria in different ways. In addition, infection control is a hard to standardise parameter which could influence relevant studies drastically.

At this point, it would be useful to discuss the third set of studies included in our review. The latter evaluated resistance dynamics of each of the on-cycle antibiotics during the application of antimicrobial cycling protocols. They provide little information as to the overall efficacy of cycling but could offer some ground for future research as to which agents are actually less prone to the selection of resistant strains. Ginn et al compared periods of predominant cefepime and piperacillin-tazobactam use and found that cefepime, a fourth-generation cephalosporin, was associated with higher overall resistance rates (including co- and cross-resistance). There is plenty of observational research which supports the notion that piperacillin-tazobactam is a less important driver of antibiotic resistance than broad-spectrum cephalosporins[9]. A rational explanation could lie on the fact that broad-spectrum cephalosporins are less effective than inhibitor-based beta-lactams in vitro against ESBLs, which are the among most widespread multidrug-resistant strains within nosocomial environments and could be theoretically preferentially selected under the pressure of inappropriate antibiotic treatment.

On the other hand, Van Loon et al concluded that the homogeneous use of ceftazidime, another fourth-generation cephalosporin, was not associated with an increase in the incidence of ceftazidime-resistant strains, while both piperacillin-tazobactam and levofloxacin use provoked resistance. The results of those studies are seemingly contradictory and could be confounded either by seasonality or breaks in infection control. Such discrepancies underline the importance of the use of contemporaneous controls as well as the need for bacterial typing data in future research to facilitate a more meaningful interpretation of the data. Bacterial typing becomes especially important in view of the fact that most studies to date have used the unit-wide incidence of resistant strains as the primary outcome indicator, but this is easily affected by changes in colonization pressure and/or breaks in infection control. An idea for future research would also be to differentiate colonization rates in patient groups within the same ward who have and have not participated in study protocols and use additional wards with similar baseline characteristics as comparison units.

Lack of standardization of research protocols was once again a crucial issue which limited our ability to evaluate with confidence the replicability of findings and reach safer conclusions. Research protocols differed in terms of the cycle length, the choice of empirical agents, the opportunity to de-escalate, the acquisition of typing data to assess cross-transmission dynamics, and the measurement of indicators of potential collateral damage induced by the established policies. Among the studies of our dataset it was only Van Duijn et al in 2018 who utilised a cluster-randomised cross-over design to compare cycling with mixing, which was a stronger study design than most. A more thorough evaluation would be possible only if the study included control groups and/or baseline data as well as bacterial typing to assess bacterial clonality. It is true that the conduct of research well-designed and rigorous to be of practical use to clinicians requires specialist expertise of multiple kinds, and is logistically difficult. Nevertheless, it is a worthwhile investment which should be co-ordinated by national or international public health agencies with the ultimate aim to safeguard the future value of antimicrobial agents.

Conclusion

Although we cannot exclude the possibility that yet unexplored cycling protocols could show benefits in the future we believe that the routine use of the currently tested options in current clinical practice should not be expected to improve bacterial resistance rates to any appreciable extent. We hope that this review will inspire a more standardised and rigorous approach in the future, as with some upgrading, this type of research could create an enormous contribution to the control of pathogenic bacteria worldwide.

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Figure 1: PRISMA chart depicting the study selection process

Table 1: Catalogue of the studies assessing the effects of antimicrobial cycling on bacterial resistance rates; A p value<0.05 was regarded as the statistical threshold of significance in all studies and is accordingly

recorded as such.

Authors	Study Design	Setting	Protocol	Outcomes	Indicator
Toltzis P et al 2002	Controlled trial	Neonatal ICU	Monthly cycling of gentamicin, piperacillin-tazobactam and ceftazidime for suspected infections due to Gram-negative pathogens versus standard practice in the control group (usually ampicillin and gentamicin for suspected infection at birth, vancomycin and gentamicin for hospital-acquired infection, ampicillin and cefotaxime for meningitis, and piperacillin-tazobactam for necrotizing enterocolitis) No de-escalation Typing to assess clonality of bacterial isolates	PRIMARY Similar incidence of colonization with resistant bacilli to any antibiotic Similar incidence of colonization with resistant bacilli to the rotated antibiotics (even when only data regarding clonally discordant isolates were considered) OTHER On-cycle antibiotic use 84.3% for the rotation team Predominant use of gentamicin in the control team Similar overall antibiotic use Similar length of stay	Unit-wide surveillance cultures

Authors	Study Design	Setting	Protocol	Outcomes	Indicator
Cadena J et al 2007	Before-and-after	Haematology-Oncology Unit	Cycling of piperacillin-tazobactam and cefepime for the empirical therapy of neutropenic fever every three months versus standard practice during a baseline period (not further clarified) Potential of de-escalation not clarified No typing of bacterial isolates to assess clonality	PRIMARY Inconclusive changes in relevant susceptibilities of Enterobacterales and <i>P. aeruginosa</i> Decrease in ampicillin-susceptible <i>Enterococcus</i> spp, erythromycin- and clindamycin-susceptible <i>S. aureus</i> OTHER Increase in cefepime and piperacillin-tazobactam consumption index from 0.003 to 0.88 Increase in cefepime use	Unit-wide clinically indicated cultures

Authors	Study Design	Setting	Protocol	Outcomes	Indicator
Bennett KM et al 2007	Before-and-after	Surgical ICU	Cycling of piperacillin-tazobactam, imipenem, ceftazidime and ciprofloxacin every month for the empirical treatment of suspected Gram-negative infections (Ciprofloxacin discarded later) versus standard practice during a baseline period (not further clarified) De-escalation permitted No typing of bacterial isolates to assess clonality	PRIMARY Increase in piperacillin-tazobactam and ceftazidime-susceptible <i>P. aeruginosa</i> proportions; No changes for the Medical ICU (Used as a comparison unit) Inconclusive changes for <i>E. coli</i> and <i>K. pneumoniae</i> in the Surgical ICU; Increase in piperacillin-tazobactam-resistant <i>E. coli</i> proportions and inconclusive changes for <i>K. pneumoniae</i> in the Medical ICU OTHER No information provided regarding secondary outcomes	Unit-wide clinically indicated cultures
Smith R et al 2008	Before-and-after	Surgical ICU	Cycling of vancomycin and linezolid for suspected Gram-positive infections every three months versus primary vancomycin use during a baseline period De-escalation permitted No typing of bacterial isolates to assess clonality	PRIMARY Decrease in MRSA incidence rates during cycling Similar VRE incidence rates OTHER Similar percentage of in-hospital deaths according to initial empirical therapy Similar incidence rates of <i>C. difficile</i> colitis	Unit-wide clinically indicated cultures

Authors	Study Design	Setting	Protocol	Outcomes	Indicator
Nijssen S et al 2009	Prospective comparative cross-over trial	2 ICUs (Medical ICU and Neurosurgery ICU)	Weekly cycling of ceftriaxone, amoxicillin-clavulanate and levofloxacin or ciprofloxacin as empirical treatment versus the homogeneous administration of ciprofloxacin or levofloxacin No de-escalation Typing of isolates to exclude clonal outbreaks	PRIMARY Higher colonization rates for ciprofloxacin-resistant isolates (including ciprofloxacin-resistant cephalosporin-resistant isolates) during the homogeneous period Similar colonization rates for cephalosporin-resistant Enterobacteriaceae OTHER Similar overall antibiotic use Higher ciprofloxacin use during the homogeneous period Lower third-generation cephalosporin use during the homogeneous period	Unit-wide surveillance cultures

Authors	Study Design	Setting	Protocol	Outcomes	Indicator
Raineri E et al 2010	Before-and-after	2 ICUs	Cycling of piperacillin-tazobactam, fluoroquinolones, carbapenems, cefepime/ceftazidime every three months for the empirical treatment of VAP versus standard practice in a baseline period (most commonly piperacillin-tazobactam or levofloxacin) No de-escalation No typing of bacterial isolates to assess clonality	PRIMARY Similar incidence of VAP due to antibiotic-resistant bacteria Decrease in cefepime- and aminoglycoside-resistant <i>P. aeruginosa</i> isolates Decrease in cefazolin-resistant <i>K. pneumoniae</i> and <i>E. coli</i> isolates No other conclusive changes OTHER On-cycle antibiotic use 83% in Unit 1 and 88% in Unit 2 Increase in carbapenem and extended-spectrum penicillin use Decrease in aminoglycoside, fluoroquinolone, 3GC and 4GC use Similar mortality rates	Respiratory cultures derived from Ventilator-associated Pneumonia cases

Authors	Study Design	Setting	Protocol	Outcomes	Indicator
Cumpston A et al 2012	Before-and-after	Blood and Marrow Transplantation Unit	Pre-cycling period: No prophylaxis for neutropenia;* Piperacillin-tazobactam for the empirical treatment of febrile neutropenia Period A: Cycling of imipenem, cefepime plus+ tobramycin and piperacillin-tazobactam plus tobramycin every eight months for the empirical treatment of febrile neutropenia; Levofloxacin as prophylaxis for neutropenia* Period B: Cycling of agents every three months; Addition of tobramycin in the imipenem arm; Levofloxacin as prophylaxis for neutropenia* *Addition of vancomycin at the discretion of the clinician De-escalation permitted No typing to assess clonality of bacterial isolates	PRIMARY Increase in quinolone-resistant Enterobacterales incidence rates Increase in VRE incidence rates No other conclusive changes in resistance patterns OTHER Decrease in vancomycin use Similar use of cefepime, piperacillin-tazobactam and imipenem across the four most recent years of cycling Decrease in incidence rate of <i>Klebsiella</i> and <i>E. coli</i> bacteremia and candidemia Similar morbidity and mortality incidence rates	Unit-wide blood cultures

Authors	Study Design	Setting	Protocol	Outcomes	Indicator
Chong Y et al 2013	Before-and-after	Haematology Unit	Monthly cycling of piperacillin-tazobactam, ciprofloxacin, meropenem and cefepime for the empirical treatment of neutropenic fever versus the homogeneous use of cefepime during a baseline period Potential of de-escalation not clarified No typing of bacterial isolates to assess clonality	PRIMARY Blood isolates: Decrease in cefepime-resistant isolate incidence from 6/13 (70% of those were ESBLs) to 0/14 (p=0.007); Decrease in ciprofloxacin-resistant isolate incidence Stool isolates: Decrease in ESBL and ciprofloxacin-resistant <i>E. coli</i> incidence OTHER Similar mortality rates 65.9% decrease in unit-wide cefepime-use	Blood and stool cultures from patients with neutropenic fever
Teranishi H et al 2017	Before-and-after	Paediatric Haematology Unit	Monthly cycling of piperacillin-tazobactam, meropenem and cefepime versus the homogeneous prescription of cefpirome as empirical treatment for neutropenic fever during a baseline period No de-escalation No typing of bacterial isolates to assess clonality	PRIMARY Blood isolates: Decrease in ESBL incidence from 5/15 to 0/15 isolates (p<0.05) Nasal and stool isolates: Decrease in ESBL incidence from 15/33 to 0/33 isolates (p<0.01) Similar MRSA and VRE incidence OTHER No information provided regarding secondary outcomes	Blood, nasal and stool cultures from patients with neutropenic fever

Authors	Study Design	Setting	Protocol	Outcomes	Indicator
Tsukayama D et al 2004	Comparative trial	ICU	Cycling of ciprofloxacin or levofloxacin plus clindamycin or metronidazole with piperacillin-tazobactam every four months as first-line empirical treatment De-escalation permitted Typing to assess clonality of bacterial isolates	PRIMARY No correlation between particular antibiotic class consumption and onset of resistance OTHER Off-cycle antibiotic use not drastically reduced	Unit-wide surveillance units
Van Loon H et al 2005	Comparative trial	ICU	Cycling of levofloxacin plus aminoglycoside with beta-lactam plus aminoglycoside (cefpime in one cycle and piperacillin-tazobactam in the other) every four months for suspected Gram-negative infections No de-escalation No typing of bacterial isolates to assess clonality	PRIMARY Colonization rates for Gram-negative bacteria resistant to levofloxacin higher in periods of exposure Colonization rates for Gram-negative bacteria resistant to cefpime similar between periods of exposure and non-exposure Colonization rates for Gram-negative bacteria resistant to piperacillin-tazobactam higher in periods of exposure OTHER On-cycle antibiotic use 88.5%-100%	Unit-wide surveillance cultures

Authors	Study Design	Setting	Protocol	Outcomes	Indicator
Ginn A et al 2012	Comparative trial	2 ICUs	Cycling of piperacillin- tazobactam and cefepime for the empirical therapy of sepsis every four months Potential of de-escalation not clarified Typing of isolates to exclude clonal outbreaks	PRIMARY Proportion of admissions complicated by antibiotic- resistant isolates higher in cefepime cycles Proportion of admissions complicated by MRSA higher in cefepime cycles OTHER Similar risk of admissions complicated by any infection On-cycle antibiotic use>60% of total use Off-cycle antibiotic use<15% of total use	Unit-wide clinically indicated cultures

Authors	Study Design	Setting	Protocol	Outcomes	Indicator
Martinez J et al 2006	Comparative cross-over trial	2 ICUs	1 st arm: Cycling of cefepime (or ceftazidime), ciprofloxacin, carbapenems, and piperacillin- tazobactam every month for suspected <i>Pseudomonas</i> infections 2 nd arm: Successive administration of these agents to consecutive patients Potential of de-escalation not clarified Combination therapy permitted No typing to assess clonality of bacterial isolates	PRIMARY Higher proportion of patients colonised with cefepime- resistant <i>P.</i> <i>aeruginosa</i> during mixing Inconclusively higher proportion of ceftazidime and carbapenem- resistant <i>P.</i> <i>aeruginosa</i> during mixing (p=. 0.06 and 0.07 respectively) No other significant differences with regard to other Gram-negatives species OTHER Higher mortality rates during cycling only for Unit 2 Similar mortality rates during cycling and mixing for Unit 1 Higher use of carbapenems and piperacillin- tazobactam and lower use of cephalosporins during mixing	Unit-wide surveillance cultures

Authors	Study Design	Setting	Protocol	Outcomes	Indicator
Van Duijn PJ et al 2018	Cluster randomised cross-over trial	Multi-centre ICU	Cycling of 3GC (or 4GC), carbapenems and piperacillin-tazobactam every six weeks versus mixing those agents (administering those successively to consecutive patients) for empirical treatment of suspected Gram-negative infections De-escalation permitted Combination therapy permitted No typing to assess clonality of bacterial isolates	PRIMARY Similar prevalence of antibiotic-resistant Gram-negative bacteria Similar incidence rate ratio of antibiotic-resistant Gram-negative bacteria adjusted for hand hygiene compliance, patient-sex and proportion of short-stay patients Similar prevalence of ESBLs, piperacillin-tazobactam- or carbapenem-resistant non-fermenters OTHER Similar mortality rates and similar length of stay during periods of mixing and cycling Similar overall use of antibiotics and similar use of study antibiotics between study periods Three times higher use of on-cycle antibiotics compared to off-cycle use	Unit-wide surveillance cultures

Authors	Study Design	Setting	Protocol	Outcomes	Indicator
Jayashree M et al 2020	Comparative trial	Paediatric ICU	Period 1: Mixing piperacillin-tazobactam, imipenem and cefepime (administering those successively to consecutive patients) for suspected Gram-negative infections Period 2: Cycling the aforementioned agents every month De-escalation permitted Combination therapy permitted No typing to assess clonality of bacterial isolates	PRIMARY Higher percentage of resistant isolates during baseline period than in mixing, cycling and washout periods Similar percentage of resistant isolates during mixing and cycling OTHER Similar mortality rates between periods Similar episodes of healthcare-associated infections during mixing and cycling but lower than baseline Similar overall use of antibiotics between all phases	Unit-wide surveillance cultures