

Cite this: *RSC Chem. Biol.*, 2021, 2, 368

Brief survey on organometalated antibacterial drugs and metal-based materials with antibacterial activity

Przemysław Biegański, ^a Łukasz Szczupak, ^a Manuel Arruebo ^{bcd} and Konrad Kowalski *^a

Rising bacterial antibiotic resistance is a global threat. To deal with it, new antibacterial agents and antiseptic materials need to be developed. One alternative in this quest is the organometallic derivatization of well-established antibacterial drugs and also the fabrication of advanced metal-based materials having antibacterial properties. Metal-based agents and materials often show new modes of antimicrobial action which enable them to overcome drug resistance in pathogenic bacterial strains. This review summarizes recent (2017–2020) progress in the field of organometallic-derived antibacterial drugs and metal-based materials having antibacterial activity. Specifically, it covers organometallic derivatives of antibacterial drugs including β -lactams, ciprofloxacin, isoniazid, trimethoprim, sulfadoxine, sulfamethoxazole, and ethambutol as well as non-antibacterial drugs like metformin, phenformin and aspirin. Recent advances and reported clinical trials in the use of metal-based nanomaterials as antibiofouling coatings on medical devices, as photocatalytic agents in indoor air pollutant control, and also as photodynamic/photothermal antimicrobial agents are also summarized.

Received 26th November 2020,
Accepted 4th January 2021

DOI: 10.1039/d0cb00218f

rsc.li/rsc-chembio

^a Department of Organic Chemistry, Faculty of Chemistry, University of Łódź, Tamka 12, 91-403 Łódź, Poland. E-mail: kondor15@wp.pl, konrad.kowalski@chemia.uni.lodz.pl; Tel: +48-42-635-5759

^b Instituto de Nanociencia y Materiales de Aragón (INMA), CSIC-Universidad de Zaragoza, Zaragoza 50009, Spain

^c Department of Chemical Engineering, University of Zaragoza, Campus Río Ebro - Edificio I + D, C/Poeta Mariano Esquillor S/N, 50018 Zaragoza, Spain

^d Networking Research Center on Bioengineering, Biomaterials and Nanomedicine, CIBER-BBN, 28029 Madrid, Spain

1. Introduction

Bacteria are unicellular organisms usually a few micrometers in length which together with archaea (formerly archaebacteria) belong to the prokaryote domain of life. High-resolution microscopy and isotopic analyses of some Archaean (4–2.5 Ga) rocks show that ancestors of modern bacteria existed on earth between 3.47 and 2.7 Ga,^{1–3} whereas studies on 1.88 Ga old



Przemysław Biegański

Przemysław Biegański obtained his MSc degree in chemistry in 2020 from the Łódź University of Technology (Poland). He then started his PhD at the University of Łódź under the supervision of Prof. Konrad Kowalski. His PhD project is focused on the chemistry of organometallic bioconjugates and organometallic nucleoside analogues.



Łukasz Szczupak

Łukasz Szczupak obtained his MSc degree in chemistry in 2012 from the University of Łódź (Poland). In 2017, he received his PhD degree in chemical sciences working under the supervision of Prof. Konrad Kowalski. In 2017, he came to Mabion biotechnological company and worked on monoclonal antibody therapy. In 2018, he rejoined Prof. K. Kowalski's group at the University of Łódź. His main current interests are in developing new metal-based antimicrobial and anticancer agents.



stromatolites showed Fe₂O₃-mineralized microfossils of bacterial cells.⁴ Bacteria inhabit all ecological niches including extreme environments like hydrothermal vents,⁵ hot springs,⁶ the deep sub-seafloor,⁷ sub-polar snowpacks,⁸ the Antarctic desert⁹ and even nuclear waste.¹⁰ The human body also represents an ecological niche which harbors more than 100 trillion bacteria and other microorganisms of microbiota.¹¹ Especially human gut microbiota can influence our physiology and even the risk of non-infectious disease development in organisms.¹¹ This is exemplified by recent studies which show links between the presence of *Ruminococcus flavefaciens* and hypertension as well as between *Clostridium* and platelet counts.¹² Other studies have revealed links between colon-cancer development and production of colibactin, which is a genotoxic metabolite produced by *Escherichia coli*.¹³

Interaction between humans and infective bacterial pathogens is as old as the history of mankind,¹⁴ although the modern era of antibacterial agents began in 1928 with the discovery of penicillin by Alexander Fleming.¹⁵ Penicillin was introduced in hospitals in the 1940s but its effectiveness was soon questioned by the appearance of penicillin-resistant *Staphylococcus aureus* strains.¹⁶ A similar scenario occurred in the case of streptomycin, discovered by Waksman,¹⁷ as the first streptomycin-resistant *Mycobacterium tuberculosis* was reported just a few years after its discovery.¹⁸ Likewise, the first methicillin-resistant *S. aureus* (MRSA) appeared rapidly after the introduction of this antibiotic in 1959.^{19,20} While time goes by, the progressive appearance of new drug-resistant strains including the emergence and spread of multidrug resistant (MDR) ESKAPE pathogens has continued.^{21–26} According to the WHO, today MDR bacteria are one of the key problems for health-care systems.²⁷ Thus, severe infection caused by some drug-resistant Gram-negative strains of *Pseudomonas aeruginosa*, *Acinetobacter baumannii*, *Escherichia coli*, *Klebsiella pneumoniae*, *Mycobacterium tuberculosis* or *Enterobacter spp.* may,

despite the best medical care, produce a lethal outcome.^{25,28–36} Also infections caused by some Gram-positive *S. aureus* (MRSA), vancomycin-resistant enterococci (VRE) and penicillin-resistant *Streptococcus pneumoniae* (PRSP) pathogenic strains can cause life-threatening problems during treatment.^{25,37–40}

Where does drug-resistance arise and what are the key mechanisms of resistance in bacteria? To answer the first question one has to look back into the long evolutionary history of bacteria. During billions of years of evolution, bacteria developed a number of genetically driven biochemical processes and biomolecules which enabled them to withstand the presence of different competitor microorganisms invading or sharing their environment. In return, the attacked competitors have protected themselves with their own set of molecular mechanisms, which ultimately results in a never ending arms race. Drug-resistance in bacteria originates from this evolutionary molecular competition. Uncontrolled use of antibacterial drugs in medicine, veterinary practice and agriculture increased the selection pressure on bacteria and enabled drug-resistant strains to develop, survive, and spread.²⁶ Drug resistance is a dynamic phenomenon mediated by mobile genetic vectors like R plasmids (containing transposons), bacteriophages or naked DNA following conjugation, transduction, and transformation processes, respectively.⁴¹ Transposons are able to move from plasmid to plasmid and back and forth to the bacterial chromosome. Drug resistance can spread within the same bacterial strain as well as horizontally between different bacteria.

To answer the second question – there are many specific mechanisms of resistance in bacteria.²² All of them, however, can be classified into three different general mechanistic groups (Fig. 1).

The first mechanism pertains to the bacterial development of enzymes which either degrade or alter antibacterial drugs to make them inactive. The second pertains to the biosynthesis of



Manuel Arruebo

Manuel Arruebo is a Full Professor in Chemical Engineering at the University of Zaragoza, Spain. He is the co-author of more than 150 scientific publications, including 4 book chapters. A number of those papers have been published in journals having high-impact factors including Nature Nanotechnology, Nature Catalysis, PNAS, Nano Today, Trends in Biotechnology, Biomaterials, Chemistry of Materials, Small, and Hepatology, among others, having in 2020

an h-index of 38 (WoS) and 41 (Google scholar). He was awarded in 2014 an ERC Consolidator Grant (selected from 3673 applicants throughout Europe), which helped him to consolidate his own nanobiotechnology research group.



Konrad Kowalski

Konrad Kowalski received his PhD degree in chemical sciences in 2003 at the University of Łódź (Poland). Afterwards, he worked as a postdoctoral fellow at Imperial College London (UK) with Prof. Nicholas J. Long and at the University of Regensburg (Germany) with Prof. Rainer F. Winter. In 2011, he obtained his DSc (habilitation) degree and became Associate Professor at the University of Łódź. In 2020, he was promoted to Full Professor

of Chemistry. His current scientific interests include biology-oriented chemistry of organometallic compounds, xeno-nucleic acids and luminescent probes for bioimaging applications.



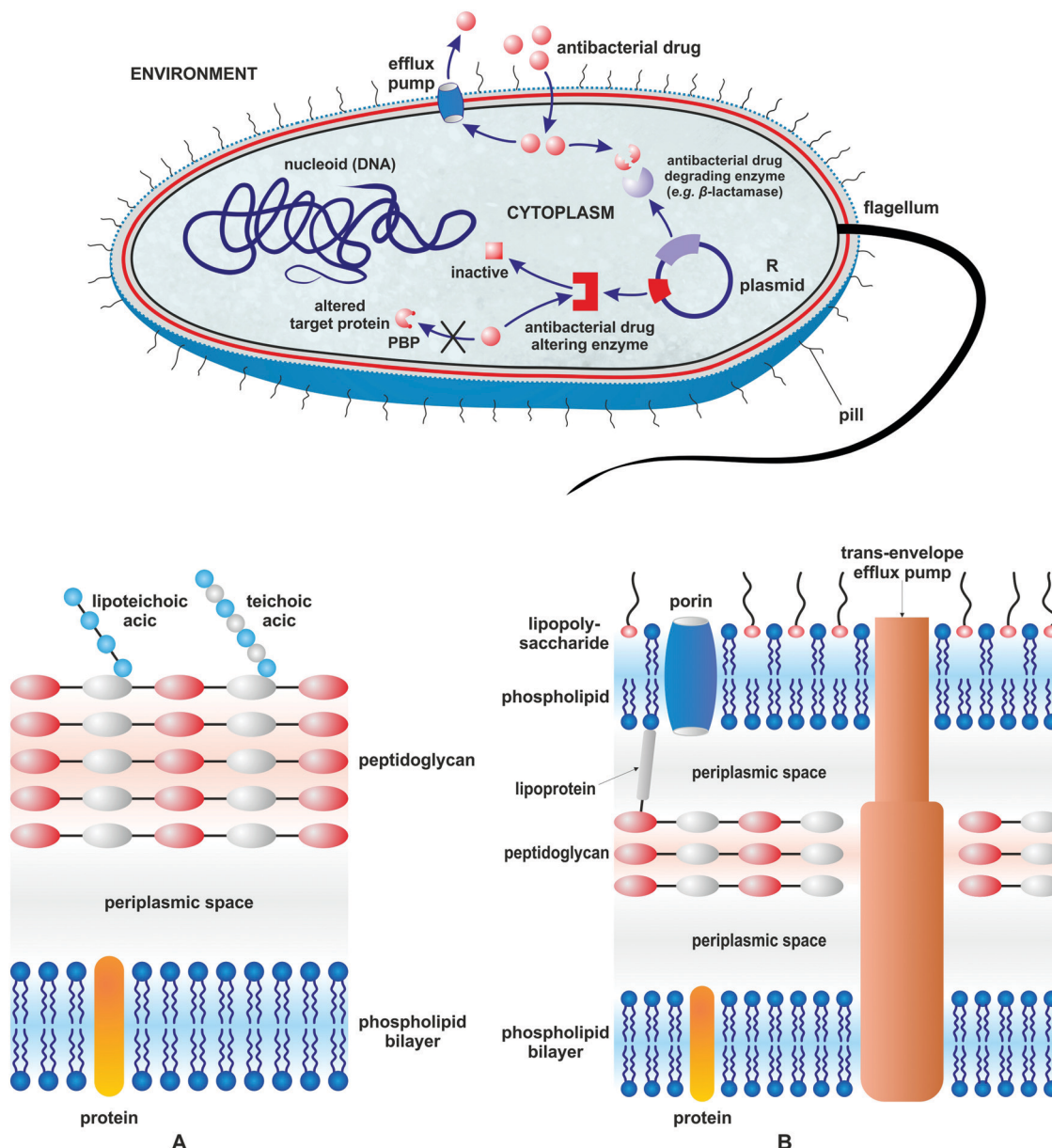


Fig. 1 Different mechanisms of drug resistance in bacteria and a schematic representation of the Gram-positive (A) and Gram-negative (B) bacterial cell envelope.

structurally modified proteins which help them to be protected from inhibition by antibacterial agents (target modification). The last mechanism relates to the modification of the bacterial cell wall to impair antibacterial drug influx and/or development of protein pump systems to enable drug efflux. The most effective way to eliminate drug resistant bacteria from a given ecological niche is to outnumber their population by non-pathogenic (non-resistant) strains in the absence of selecting factors. Obviously this approach cannot be widely applied to cure patients, especially those with acute and fast progressing infections. Thus, antibacterial drugs are a second and clinically most relevant option. They are over 200 marketed antibacterial drugs which belong to over 11 large classes.^{42,43} Historically, antibacterial agents diverge into those isolated from natural

source antibiotics and those of synthetic origin. They either eliminate bacteria (bactericidal effect) or inhibit their growth (bacteriostatic effect). Antibacterial drugs interfere with different targets and biochemical processes in the bacterial cell. Table 1 summarizes the major classes of antibacterial drugs, their mechanisms of action, and the year of their introduction.

Over the last decades the number of new antibacterial drugs has decreased.⁴³ In 2000–2020 only twenty seven antibacterial drugs have been approved for clinical use.⁴⁴ In parallel, bacteria have developed resistance to some of the “last resort drugs” like quinupristin/dalfopristin dyad, linezolid and daptomycin.^{45,46} In the face of vanishing effectiveness of all antibacterial classes, the need for new compounds is immense. In that respect, natural products are still at the forefront of research as exemplified by the

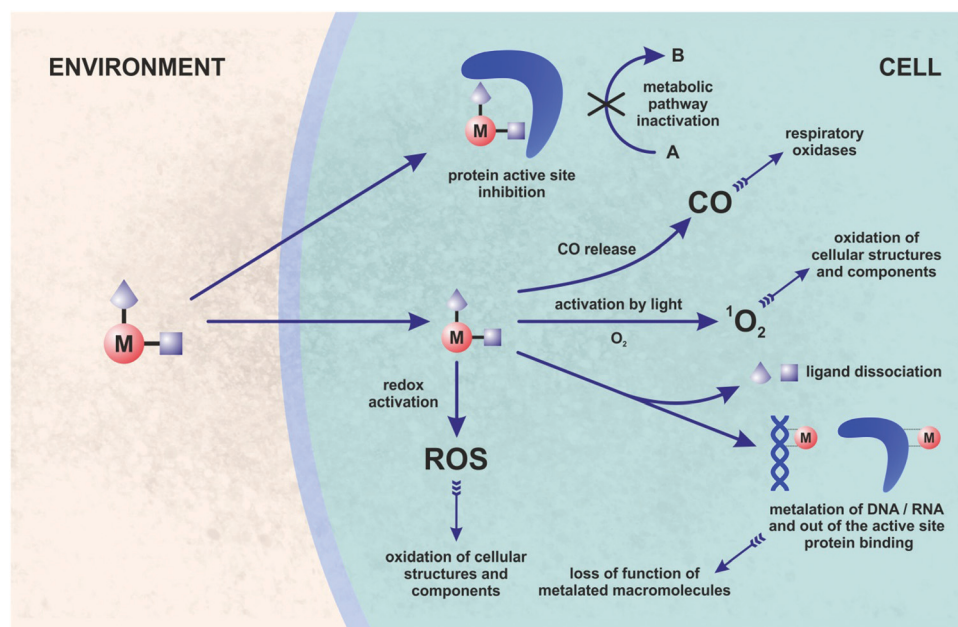


Table 1 Major classes of antibacterial drugs, and their representative examples, mechanisms of action and year of introduction

Antibacterial drug class (year of introduction, representative drug(s))	Mechanism of action
Sulfonamides (1935, prontosil) and trimethoprim (1960s)	Inhibitors of folic acid synthesis (inhibitors of dihydropteroate synthase, dihydrofolate reductase)
Quinolones (1960s) (ciprofloxacin, ozenoxacin)	Inhibitors of DNA replication (inhibitors of topoisomerase IV, gyrase)
Rifamycin (1957, does not belong to the large antibiotic families)	Inhibitor of RNA synthesis
β -Lactams (1940s) (penicillins, cephalosporins, carbapenems, monobactams) (selected examples: ampicillin, cefiderocol)	Inhibitors of cell wall synthesis and remodeling (PBP inhibitors)
Glycopeptides (1959) (vancomycin, oritavancin), lipopeptides (2003, daptomycin) and other cell wall inhibitors which do not belong to the large antibiotic families: cycloserine (1955), isoniazid (1952)	Inhibitors of cell wall synthesis
Phenylpropanoids (1949, chloramphenicol), aminoglycosides (1950, tobramycin), macrolides (1952, erythromycin), polyketides (1949, tetracycline), oxazolidinones (2000, linezolid), streptogramins (1962, quinupristin/dalfopristin), lincosamides (1960s, clindamycin)	Inhibitors of protein synthesis
Nitroimidazoles (1960, metronidazole)	Cellular damage of DNA and proteins

isolation of darobactin,⁴⁷ teixobactin,⁴⁸ numerous components present in plants⁴⁹ and other sources of antibacterially active compounds.^{43,50} Problems with culturing some antibiotic-producing microorganisms⁵¹ and limited industrial-scale synthetic accessibility to complex antibacterial natural products are two drawbacks of this approach. Therefore, culture-independent methods capable of identifying and delivering first-in class synthetic antibacterial compounds and derivatization of well-established antibacterial drugs are two attractive alternatives. Other complementary strategies comprise photodynamic (PDT)^{52–54} and photothermal (PTT) antibacterial therapies in which light is used to produce cytotoxic reactive oxygen species (ROS) and thermal damage, respectively.⁵⁵ Also, photoactivated carbon monoxide-releasing molecules (PhotoCORMs) are an emerging class of antimicrobials with promising activity against antibiotic resistant bacteria.^{56–58}

One of the alternative approaches to combating bacterial pathogens is based on inorganic^{59–62} and organometallic compounds.^{63–78} The latter are defined by the presence of at least one metal-carbon (M-C) bond. The biological activity of organometallic compounds has been a subject of many studies, especially in the field of anticancer therapy.^{79–83} Organometallic compounds are attractive candidates for medical applications as their mechanisms of action are often multi-modal, and thus not commonly accessible with purely organic pharmacophores. This feature increases the chances for organometallic compounds to overcome drug-resistance in bacteria. Fig. 2 shows the multi-modal mechanisms of action for a hypothesized bioorganometallic compound in a bacterial cell. They comprise: (a) direct protein inhibition, (b) activation by light followed by singlet oxygen generation or CO release, (c) ligand(s) dissociation and macromolecule metalation, and (d) redox activation and ROS/oxidative stress generation.

**Fig. 2** General view on possible mechanisms of action of organometallic compounds in bacteria.

Antibiotics affecting more than one cellular or biomolecular target have been recognized as highly attractive pharmacological options. Fourteen from 27 recently introduced antibacterial drugs have demonstrated synergistic or multi-target mechanisms of action.⁴⁴ Of note is also the fact that the analysis of the mechanisms of action for well-established antibacterial drugs revealed their ability to generate oxidative stress in bacteria, which increases/complements their target-specific modes of action.^{84–86}

Applications of gold in therapy have long roots in mankind's history.⁸⁷ However, it was not until Robert Koch's observation of the bacteriostatic activity of $K[Au(CN)_2]$ followed by the introduction of salvarsan by Paul Ehrlich that metals entered into the antibacterial field following the scientific method.^{88,89} Furthermore, the antiproliferative activity of cisplatin was first observed on *E. coli* strains.⁹⁰ The next breakthrough was the discovery of ferrocene.^{91–94} This sandwich, redox active, iron-containing complex rapidly gained importance within organometallic,⁹⁵ bioorganometallic^{96–98} and anti-infective organometallic chemistry.⁶⁶ Beyond ferrocene, a number of other organometallic compounds have been studied as antibacterial agents.^{63–68} Strategies for the development of antibacterially active organometallic compounds are diverse. A major group relies on the derivatization of well-established antibacterial drugs and natural products,^{63,97} whereas others focus on the synthesis of entirely new species.⁹⁹ This short review focuses mainly on organometallic-antibacterial drugs and on metal-based materials with antibacterial activity. It is not exhaustive; we just focus on the most important advancements and, in respect of organometallic compounds, we cover only the recent literature from 2017 to 2020. In respect of antibacterially active materials, a wider time frame was applied, providing a general view on the field. Readers broadly

interested in the rapidly burgeoning field of organometallic antibacterial agents are encouraged to read preceding reviews.^{63–68}

2. Organometallic derivatives of antibacterial drugs

2.1. Derivatives of β -lactams

β -lactam antibiotics have been at the forefront of antibacterial therapy since the 1940s. Their target-specific mechanism of action relies on penicillin binding protein (PBP) inhibition. PBPs are enzymes that catalyze the last steps of bacterial cell wall synthesis.^{100,101} β -lactams inhibit PBPs by irreversible acylation of a serine in a catalytic site.^{102,103} One of the major resistance mechanisms developed by bacteria against β -lactams involves the production of β -lactam hydrolyzing enzymes – β -lactamases.¹⁰⁴ β -lactamases of class A, C and D all share a common Ser residue in an active site, whereas class B are Zn^{2+} -dependent enzymes.¹⁰⁵ The first ferrocenyl conjugates of β -lactams were obtained in the 1970s.¹⁰⁶ In 2017, Kowalski, Chen and co-workers first reported ferrocenyl (Fc) and ruthenocenyl (Rc) 7-aminodesacetoxycephalosporanic acid (7-ADCA) derivatives 1–6 (Fig. 3).¹⁰⁷

The inhibitory activity of these metallocenyl- β -lactams was examined against DD-carboxypeptidase 64-575 from *Saccharopolyspora erythraea* 64-575, CTX-M-14 class A β -lactamase and *Bacillus cereus* 569/H9 class B metallo- β -lactamase. DD-Carboxypeptidase 64-575, similar to PBPs, shows affinity to a number of β -lactam antibiotics and therefore serves as a good model for studies on PBP inhibition. The active site of CTX-M-14 β -lactamase also shares catalytic features with those of PBPs.^{108,109} In comparison with penicillin G, compounds 1–6 have shown enhanced inhibitory activities, substantiating the

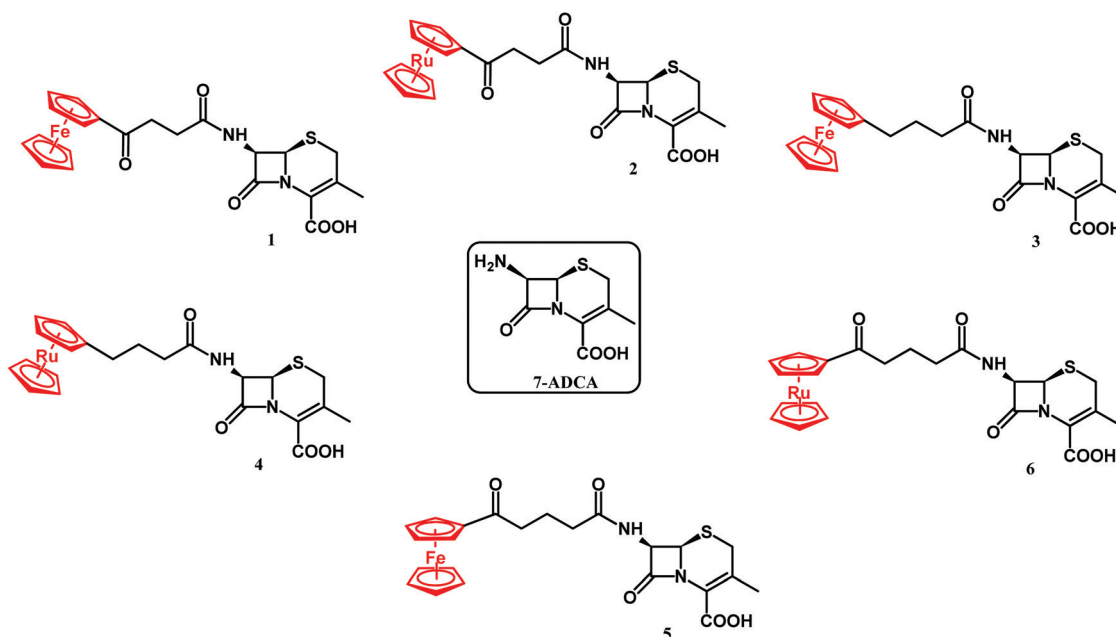


Fig. 3 Structures of 7-ADCA and compounds 1–6.



role of the metallocenyl entity in protein binding. The highest inhibition found was against DD-carboxypeptidase 64-575 with an inhibitory concentration (IC_{50}) value at the nanomolar level. Remarkably, ruthenocenyl derivatives were better inhibitors than their ferrocenyl congeners. The most active Rc derivative **4** showed an IC_{50} value of 27 nM against DD-carboxypeptidase 64-575 and 7 μ M against CTX-M-14 β -lactamase, whereas the inhibitory properties of compound **2** and **6** were characterized by IC_{50} values of 52 nM (DD-carboxypeptidase 64-575) and 44 μ M (CTX-M-14 β -lactamase), and 81 nM (DD-carboxypeptidase 64-575) and 65 μ M (569/H9 metallo- β -lactamase), respectively. Metallocenyl- β -lactams **1–6** showed lower activity than penicillin G and ampicillin in cell-based tests against model Gram-positive strains of methicillin-sensitive *Staphylococcus aureus* (MSSA), MRSA, vancomycin-intermediate *S. aureus* (VISA) and *S. epidermidis*. This feature can be explained by an impaired uptake of bacterial cells. On the other hand, the high inhibitory activities of **1–6** justified further studies aiming to understand the molecular basis of the interaction between the enzyme and the organometallic β -lactams.¹⁰⁷ Indeed, X-ray crystallographic analysis of the CTX-M E166A mutant with ruthenocenyl derivative **2** proved the formation of a covalent

acyl-enzyme complex between the β -lactam ring carbon of **2** and the Ser70O γ of the protein. Fig. 4 shows compound **2** bound to the CTX-M E166A enzyme.

XRD analysis revealed also an intact compound **2** captured at the crystal-packing interface.¹⁰⁷ Finding a non-hydrolyzed **2** in the crystal was unique as metallocenyl β -lactams eluded small-molecule crystallographic characterization. Recently, Kowalski, Chen and co-workers reported on ferrocenyl and ruthenocenyl 7-aminocephalosporanic acid (7-ACA) derivatives **7** and **8** (Fig. 5).¹¹⁰

The antibacterial activity of the two conjugates was assayed against reference Gram-negative bacterial strains (*Escherichia coli* ATCC 25922, *E. coli* NCTC 8196, *Proteus vulgaris* ATCC 49990, and *Pseudomonas aeruginosa* NCTC 6749), Gram-positive bacterial strains (*S. aureus* ATCC 6538, *S. aureus* ATCC 29213, *S. aureus* ATCC 25923, *S. epidermidis* ATCC 12228, and *Enterococcus faecalis* ATCC 29212), two methicillin-resistant *S. aureus* bacterial strains, and twelve clinical isolates of *S. aureus*, including two MRSA isolates. No antibacterial activity against Gram-negative strains and significant activity against Gram-positive Staphylococci was observed. Ruthenocenyl compound **8** showed MIC values 2–8 times lower than that of the ferrocenyl derivative **7**. Compared with the activity of metallocenyl-7-ADCA compounds **1–6** against the ATCC 29213 strain, the 7-ACA compounds **7** and **8** had reduced the MIC values by 64 \times and 16 \times for the Fc and Rc conjugates, respectively. Compounds **7** and **8** showed particularly high antibacterial activity against clinical strains of *S. aureus* isolated from the naso-pharynx and from ulcers/furuncles. In all these cases, the activity of ruthenocenyl compound **8** (MIC value 0.85 μ M) was higher than that of **7** (MIC value 15 μ M) and ampicillin. In the case of drug-resistant bone isolates of *S. aureus*, the ruthenocenyl derivative **8** again showed higher activity than its Fc counterpart and ampicillin. Furthermore, conjugate **8** showed two-fold higher antibacterial activity than ampicillin did against the Meca positive strain of *S. aureus* EDCC 5443. A lack of toxicity toward mammalian cells is the essential feature of any new antibacterial drug candidate. To address this issue, the cytotoxicity of **7** and **8** was tested *in vitro* against mouse murine fibroblast L929 and human cervical epithelioid carcinoma HeLa cells. Both tested compounds showed negligible toxicity with IC_{50} higher than 440 μ M. The antibacterial activity studies on **7** and **8** were further augmented by the identification of the complex

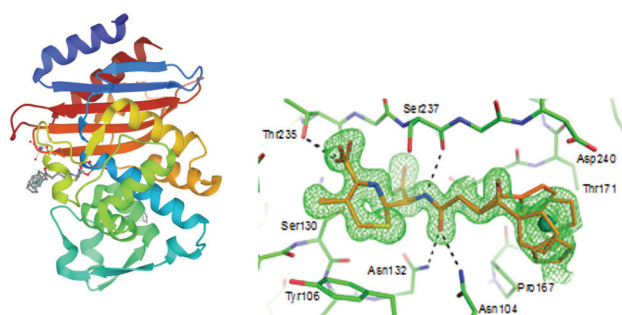


Fig. 4 Left: Structure of the acyl-enzyme complex of CTX-M E166A with compound **2** (compound **2** depicted in grey; adopts two alternative conformations). Image from the RCSB PDB (rcsb.org) of PDB ID 5UJO (E. M. Lewandowski, Ł. Szczupak, S. Wong, J. Skiba, A. Guśpiel, J. Solecka, V. Vrček, K. Kowalski and Y. Chen, *Organometallics*, 2017, **36**, 1673–1676). Right: Enlarged view of the acyl-enzyme complex nested in the protein binding site. The protein and compound are shown in green and gold, respectively. The unbiased $F_o - F_c$ density map is shown in green at 2σ . The figure was reproduced from ref. 107 with permission from the American Chemical Society.

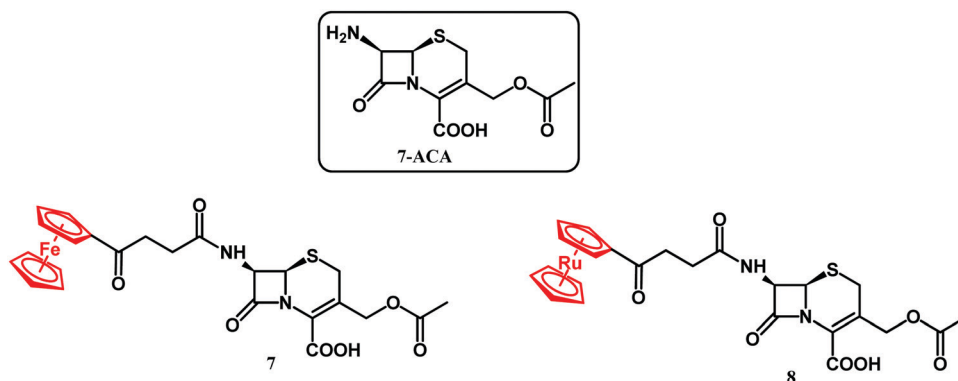


Fig. 5 Structures of 7-ACA and compounds **7** and **8**.



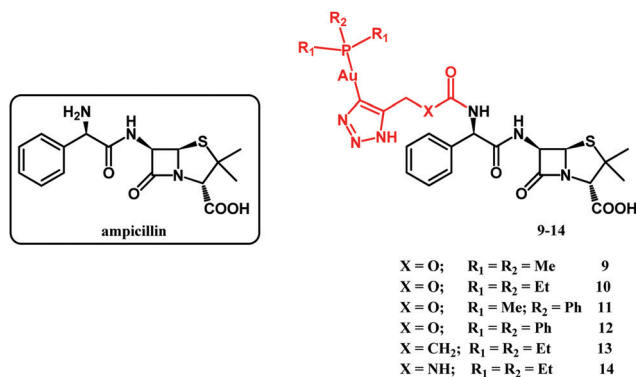


Fig. 6 Structures of ampicillin and compounds 9–14.

crystal structure of compound **8** with CTX-M-14 β -lactamase and by scanning electron microscopy (SEM) micrographs.¹¹⁰ In conclusion, the higher antibacterial activity of the ruthenocenyl β -lactams compared to the ferrocenyl counterparts can be ascribed to favorable interactions of the Rc entity with amino acid residues in the active site of PBP enzymes.

In 2020, Mislin and co-workers reported on Au(I)-ampicillin complexes **9–14** (Fig. 6).¹¹¹ The antibacterial activity of compounds **9–14** was tested against Gram-positive *S. aureus*, *S. epidermidis*, *E. faecalis* and *E. faecium* as well as against Gram-negative *E. coli* strains. It turned out that the size of the gold-coordinated phosphine has a key role in the activity, whereas the nature of the X linker has not. Accordingly, the bulkiest

triphenylphosphine derivative **12** was less active than the rest of that series as well as ampicillin. On the contrary, less sterically hindered triethylphosphine compounds **10**, **13** and **14** all showed improved (or comparable) activity compared to ampicillin against Gram-positive strains. The most potent against *Staphylococcus* species and several *Enterococcus* strains compound **10** was tested to evaluate its toxicity in healthy human hepatocytes. It did not affect eukaryotic cell viability at concentrations up to 10 μ M. Above that concentration the compound was cytotoxic. Despite this drawback, compound **10** remained as a good starting point for the design of new gold-based β -lactam antibacterial agents.

2.2. Derivatives of ciprofloxacin

Quinolones are by far the most successful synthetic antibacterial drugs on a global scale. The mechanism of antibacterial activity of quinolones involves inhibition of type II bacterial DNA topoisomerases; namely gyrase (the primary target in G-negative bacteria) and topoisomerase IV (the primary target in G-positive bacteria).^{112,113} Both of these proteins are crucial for bacterial physiology and are broadly distributed. Topoisomerase inhibition by quinolones collides with bacterial replication forks and transcription complexes, which ultimately leads to SOS system induction (SOS after international telegraph (or optical) distress signal “SOS” in the Morse alphabet). When DNA strand breaks surpass the SOS response, the bacterial cell dies. Kowalski, Stączek and co-workers reported on six organometallic ciprofloxacin derivatives **9–14** (Fig. 7).¹¹⁴

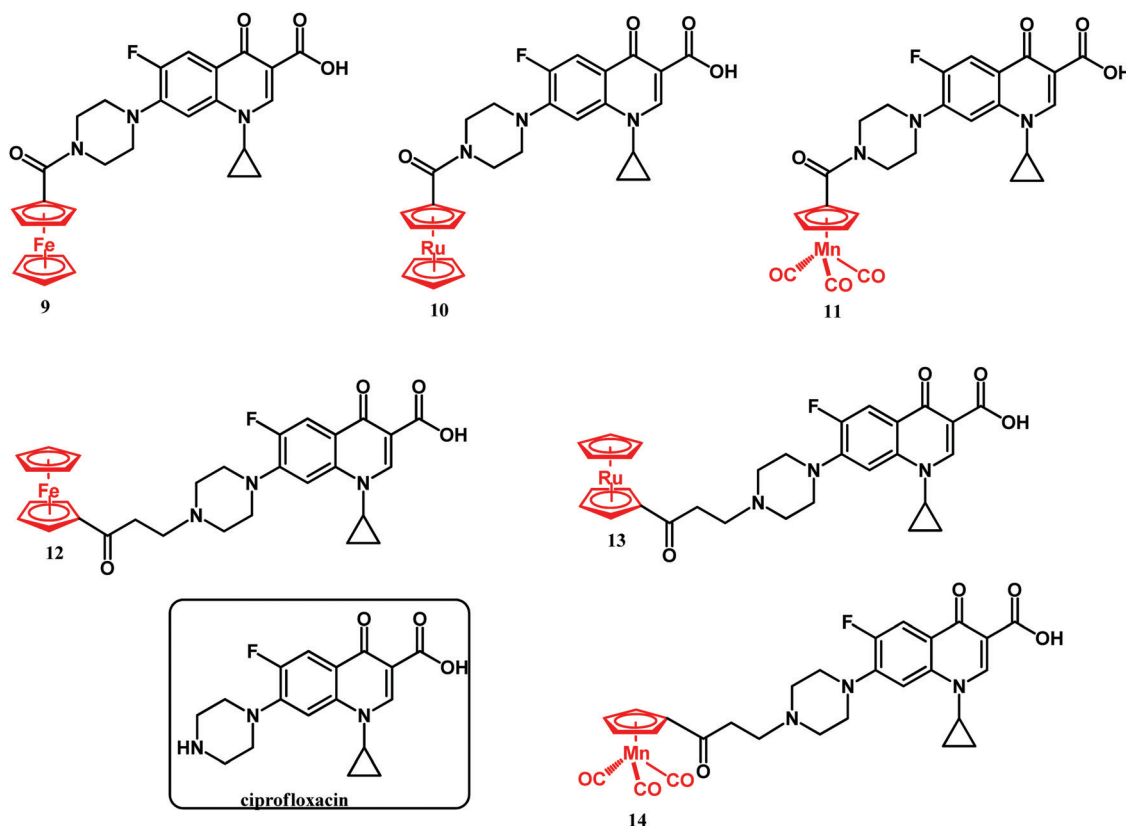


Fig. 7 Structures of ciprofloxacin and compounds 9–14.



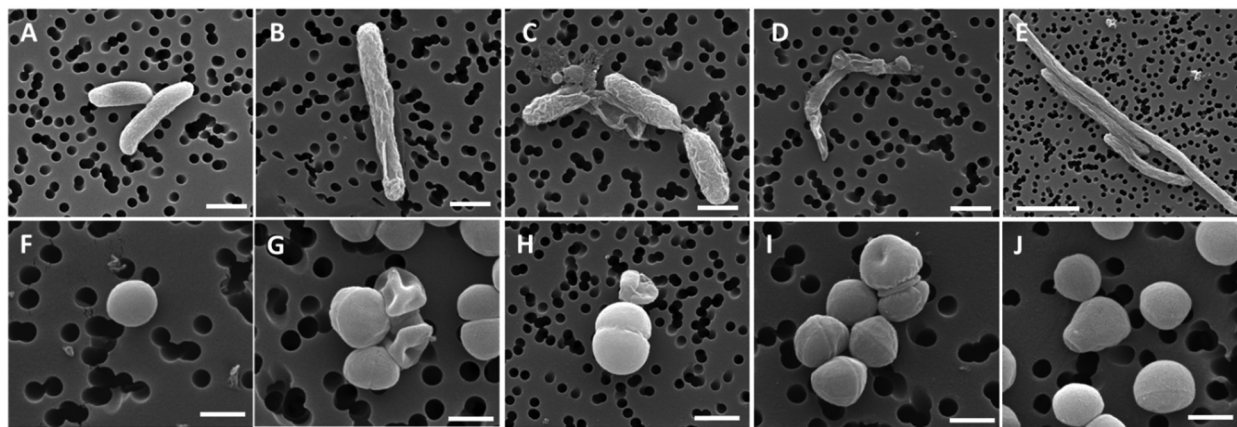


Fig. 8 SEM micrographs of the bacterial cell morphology after exposure to compounds **12–14** and ciprofloxacin. Top panels, *E. coli* cells; bottom panels, *S. aureus* cells. (A and F) control, non-treated samples; and bacteria treated with compounds **12** (B and G), **13** (C and H), **14** (D and I), and ciprofloxacin (E and J). Scale bars: 500 nm in F, G, I, and J; 1 μm in A–D and H; and 3 μm in E. Reproduced from ref. 114 with permission from the Royal Society of Chemistry.

Antibacterial activity studies of **9–14** showed that conjugation of the organometallic moiety to the ciprofloxacin scaffold enhances the bactericidal effect. Accordingly, the *N*-alkyl derivatives **12–14** were the most active compounds, although derivatives **9–11** also showed significant activity. Compounds **13** and **14** were substantially more active against the Gram-negative *E. coli* ATCC 25922 strain than ciprofloxacin. Their MIC values were 0.0006 and 0.0001 μM , respectively, while the MIC of ciprofloxacin against the same strain was 0.01 μM . Thus, in the case of the cymantrenyl derivative **14**, the MIC value was 100 times lower than that of ciprofloxacin. Furthermore, compound **14** was more active than ciprofloxacin against the *S. aureus* ATCC 6538 and *K. pneumoniae* ATCC 13883 strains with MICs of 0.4 and 0.001 μM , respectively. For the latter, the MIC value of **14** was 50 times lower than that of ciprofloxacin. Cymantrenyl derivative **11** overcame drug-resistance in two clinical bone isolates of *S. aureus* (MRSA) strains. Another feature of organometallic ciprofloxacin derivatives was their ability to eradicate pathogenic bacteria in the stationary growth phase. In this regard, the most active was cymantrene conjugate **14**. It decreased the viability of the *E. coli* ATCC strain from 3.3×10^8 (untreated control) to 2.2×10^6 CFU mL^{-1} . The remarkable

antibacterial activity of compounds **9–14** stems from their dual mechanism of action. With the exception of **9**, all compounds inhibited the introduction of supercoils by *E. coli* gyrase and the decatenation process by *S. aureus* topoisomerase IV. This is the first mechanism of action and it originated from the ciprofloxacin portion of the conjugates. The second mechanism of action pertains to the ability of oxidative stress induction in bacterial cells and stems from the organometallic portion of those conjugates. The synergistic effect between the two mechanisms enables them to overcome drug resistance in bacteria as well as to eliminate them even in the stationary phase of growth. Fig. 8 shows the structural changes in the bacterial morphology upon treatment with the organometallic ciprofloxacin derivatives **12–14**.

In 2018, Pokharia and co-workers reported on triorganotin(IV) ciprofloxacin complexes **15** and **16** (Fig. 9).¹¹⁵ Recently, a closely related diphenyltin(IV) derivative **17** has been reported by Hadjikakou and co-workers (Fig. 9).¹¹⁶

Compounds **15** and **16** showed better antibacterial activity than ciprofloxacin against Gram-positive (*S. aureus* and *E. faecalis*), and Gram-negative (*K. pneumoniae*, *E. coli*, *P. aeruginosa* and *P. mirabilis*) strains. Likewise, diphenyltin complex **17** was more

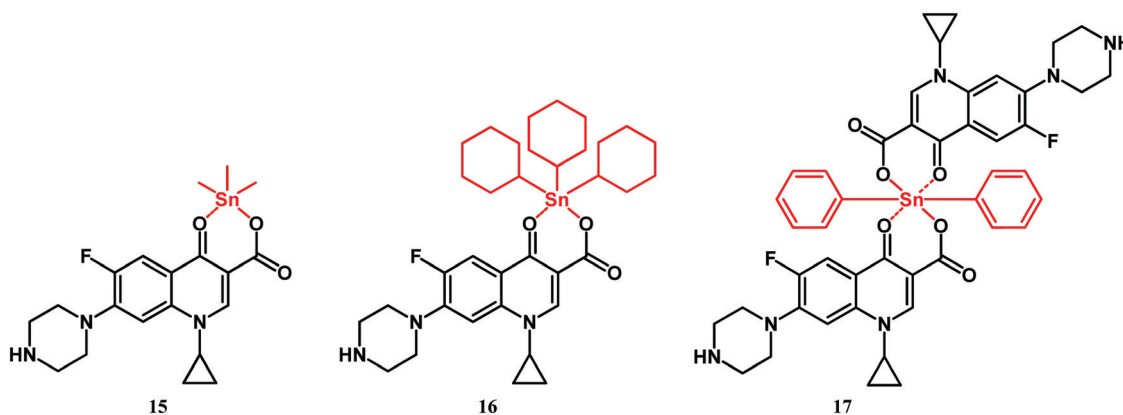


Fig. 9 Structures of compounds **15–17**.



active than its constituents (ciprofloxacin and diphenyl dichloride) against *S. aureus* and *S. epidermidis* as well as against *E. coli* and *P. aeruginosa* strains. The MICs of **17** for these strains were in the nanomolar concentration range. Furthermore, complex **17** eradicated the biofilm of *P. aeruginosa* and *S. aureus* more effectively than ciprofloxacin did.

2.3. Derivatives of isoniazid, pyrazinamide and ethambutol

Rifampicin, isoniazid, pyrazinamide and ethambutol are four drugs used in treatment of the initial phase of tuberculosis.¹¹⁷ Isoniazid is a prodrug which upon transformation to the isonicotinoyl radical by the *M. tuberculosis* KatG enzyme and further reaction with NAD(P) inhibits mycobacterial cell wall formation.^{41,117} Smith and co-workers reported on ferrocenyl isoniazid (**18–20**), ferrocenyl pyrazinoic acid hydrazide (**21–23**) and half-sandwich derivatives **24–26** (Fig. 10).¹¹⁸

The activity of compounds **18–26** has been examined against the *M. tuberculosis* H37Rv strain in glycerol-based GAST-Fe and in glucose-based Middlebrook 7H9-ADC growth media. The compounds showed interesting growth medium-dependent *M. tuberculosis* eradication properties. Their antituberculosis activity in the glycerol-based GAST-Fe medium was noticeably higher than that in the glucose-based Middlebrook medium. The most active compound was the isoniazid derivative **18**. The MIC₉₀ value of 0.39 μM was comparable to that of the parental isoniazid. Incorporation of the second metal center gave no improvement in their activity, as the MICs of binuclear compounds **24–26** were in the 0.416 to 0.968 μM range. The authors

hypothesized that the mechanisms of action of compound **18** might involve the disruption of the glycerol metabolism, resulting in the accumulation of toxic products in *M. tuberculosis* H37Rv cells.

In 2019, an interesting report on the cationic pentamethyl-cyclopentadienyl-Ir(III) ethambutol complex **27** (Fig. 11) was published by Merola and co-workers.¹¹⁹

This complex showed no activity against *M. tuberculosis*, which could be due to its ligand-dissociation inertness together with its positive charge, which impairs cellular uptake. On the other hand, **27** showed activity against *S. aureus* MSSA (MIC = 35 $\mu\text{g mL}^{-1}$) and MRSA (MIC = 40 $\mu\text{g mL}^{-1}$). Studies aiming to understand the mechanisms of antistaphylococci action are underway.

2.4. Derivatives of sulfonamides and trimethoprim

Sulfonamides are antimetabolites (antifolates) with bacteriostatic activity. Their mechanism of action stems from competitive binding of *para*-aminobenzoic acid (PABA) to dihydropteroate synthase (DHPS).^{41,117} DHPS plays a key role in dihydrofolic acid (DHF) biosynthesis. Similar to sulfonamides, trimethoprim also interferes with the folate biosynthesis pathway. It acts as an inhibitor of dihydrofolate reductase (DHFR), an enzyme which transforms DHF into tetrahydrofolic acid (THF). Foliates are cofactors for nucleobase biosynthesis in bacteria. Thus, depletion of their cellular availability prevents DNA synthesis and impairs bacterial cell division and growth. In 2017, Arancibia and co-workers reported on ferrocenyl and cyrhetrenyl (Cyr)

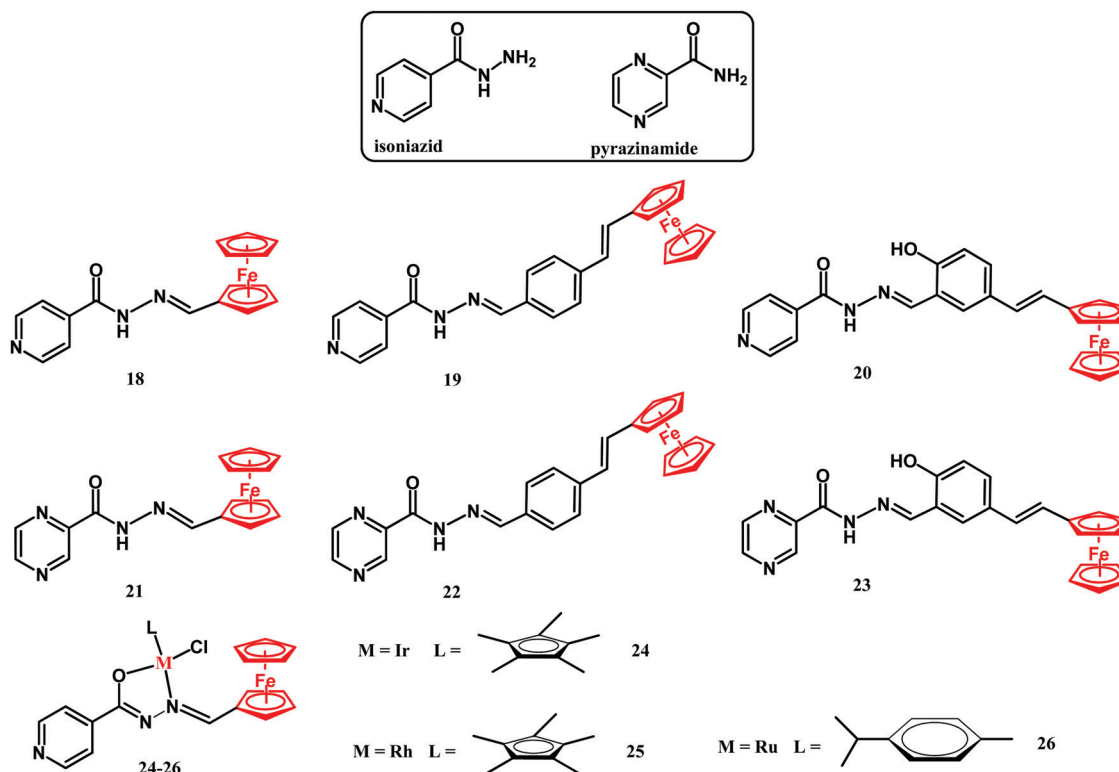


Fig. 10 Structures of isoniazid, pyrazinamide and compounds **18–26**.



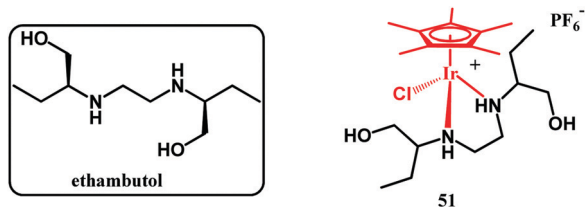


Fig. 11 Structures of ethambutol and compound 27.

derivatives 28–33 (Fig. 12), all having a sulfonamide structural core and variable substituents in the phenyl *para* position.¹²⁰

Compounds 28–33 were tested for antitubercular activity against the *M. tuberculosis* mc²6230 model strain. Isoniazid was used as reference drug and had MIC₉₉ = 0.4 μM. The antibacterial activity of Cyr derivatives 28–30 was higher than that of their ferrocenyl congeners 31–33 and *ca.* 465 times lower than isoniazid. The better antibacterial activity of the Cyr derivatives over the Fc compounds was explained by the electron-withdrawing *vs.* electron-donating properties of the former *vs.* the latter moiety.

In 2018, Sadler and co-workers reported on half-sandwich Ru(II), Rh(III) and Ir(III) complexes 34–47 (Fig. 13) containing sulfadoxine derived ligands.¹²¹ Sulfadoxine is a drug administered together with pyrimethamine in antimalarial therapy. They both block folate biosynthesis in the *Plasmodium falciparum* parasite by targeting DHPS and DHFR.

In vitro biological activity studies of 34–47 have been primarily directed toward the screening of their inhibitory activity against the *P. falciparum* 3D7 (chloroquine-sensitive), Dd2 (chloroquine-sensitive) and NG54 Late Stage Gametocyte (LSG) strains as well as against *Trichomonas vaginalis* parasite strain G3. Furthermore, the antibacterial potential of compounds 34–47 has been tested against the laboratory *M. tuberculosis* H37Rv strain. Ruthenium complexes 34 and 41 showed no antibacterial activity under the experimental conditions applied (MIC₅₀ = 100 μM). The antituberculosis activity of pyridylimino-sulfadoxine rhodium complexes 35–37 was superior to the activity of their iridium counterparts 38–40. Interestingly, the reverse trend in activity has been observed for quinolylimino-sulfadoxine derivatives 42–47.

In this respect, the most active compounds were iridium complexes 46 and 47 (both with MIC₅₀ = 3.13 μM), whereas their rhodium counterparts 43 and 44 showed MIC₅₀ = 6.25 and 50 μM, respectively. Compounds 35, 36, 43, 44, 46 and 47 exhibited antimycobacterial activity significantly higher than that of the parental sulfadoxine drug. On the other hand, none of the fourteen assayed complexes rivaled rifampicin and isoniazid antimycobacterial drugs in activity. The structure–activity relationship (SAR) for antimycobacterial activity is not straightforward and has not been provided in those studies. Compounds 34–47 showed no or limited activity against the *T. vaginalis* G3 strain. In respect of the antiplasmodial activity, the rhodium complexes were, in general, more active than the iridium compounds, whereas the ruthenium derivatives showed no activity. An interesting observation was that compounds 35–40 and 42–47 were active against sexual last stage gametocytes (LSG), whereas sulfadoxine, pyrimethamine and chloroquine were inactive. This feature clearly shows a beneficial role of the metal in the sulfadoxine activity improvement. SAR studies have shown that the quinolylimino-sulfadoxine compounds are more potent than their pyridyl-sulfadoxine analogues. Other activity-control factors are the size/lipophilicity of the Cp ligand and the rate of the chloride to water ligand exchange reaction.

A recent account on organometallic sulfonamide derivatives was published in 2019 by Metzler-Nolte and co-workers.¹²² They obtained sulfamethoxazole Ru(II) and Re(I) complexes 48–53 (shown in Fig. 14) and they tested them against four *S. aureus* strains (DSM 20231, methicillin-resistant ATCC 43300, BAA 976 and BAA977), as well as against *Acinetobacter baumannii* and *Pseudomonas aeruginosa* strains. Sulfamethoxazole is a DHPS inhibitor and is clinically used in combination with DHF inhibitor trimethoprim in a formulation known as cotrimoxazole (sulfamethoxazole:trimethoprim, 20:1 formulation) to treat a wide range of bacterial infections.^{123,124}

Sulfamethoxazole alone was not active against the tested bacterial strains up to a concentration of 512 μg mL⁻¹. On the contrary, trimethoprim alone was highly active against all strains except for Gram-negative *A. baumannii* and *P. aeruginosa*. For the majority of the strains, the cotrimoxazole activity was more potent

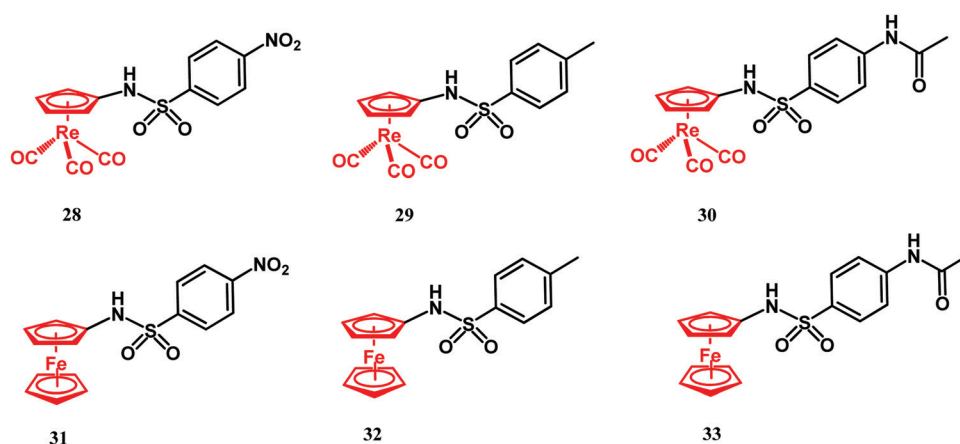


Fig. 12 Structures of compounds 28–33.



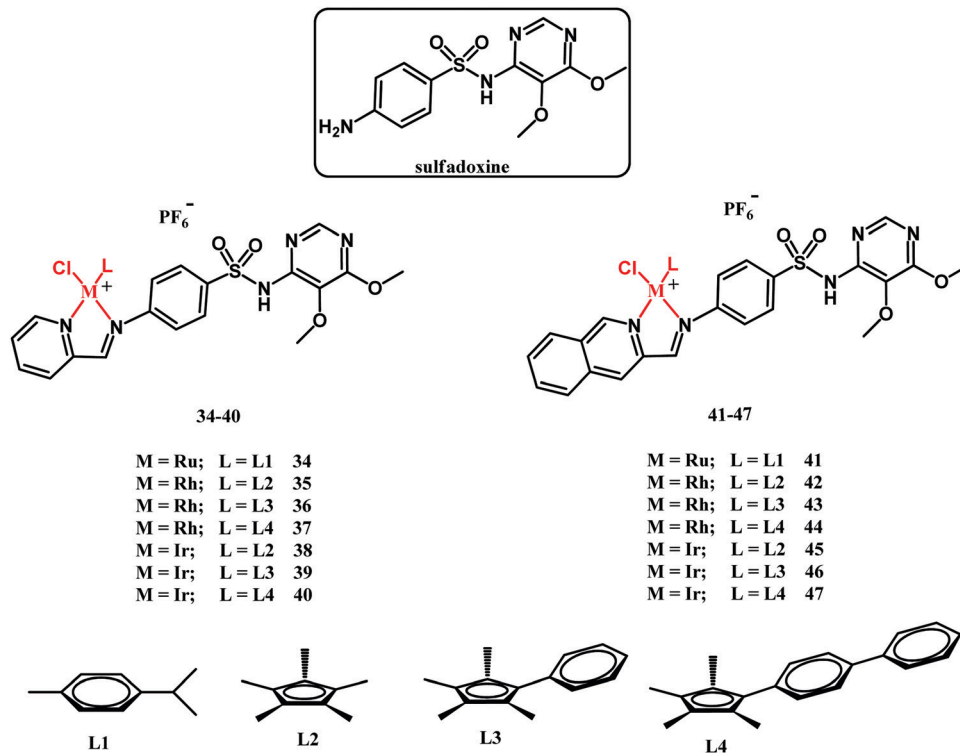


Fig. 13 Structures of sulfadoxine and compounds 34–47.

on a molar basis than trimethoprim alone. Organometallic complexes 50, 51 and 52 showed medium activity against the *S. aureus* strains with MIC = 190–750 μM . This activity was lower than that of trimethoprim alone (MIC = 6.9–14.0 μM). Noticeably, an

improvement in the activity was observed when a combination of complexes 50 and 51 with trimethoprim was tested. Like in the case of cotrimoxazole, a molar ratio of 20 : 1 (Re complex:trimethoprim) was formulated. Both Re-trimethoprim formulations

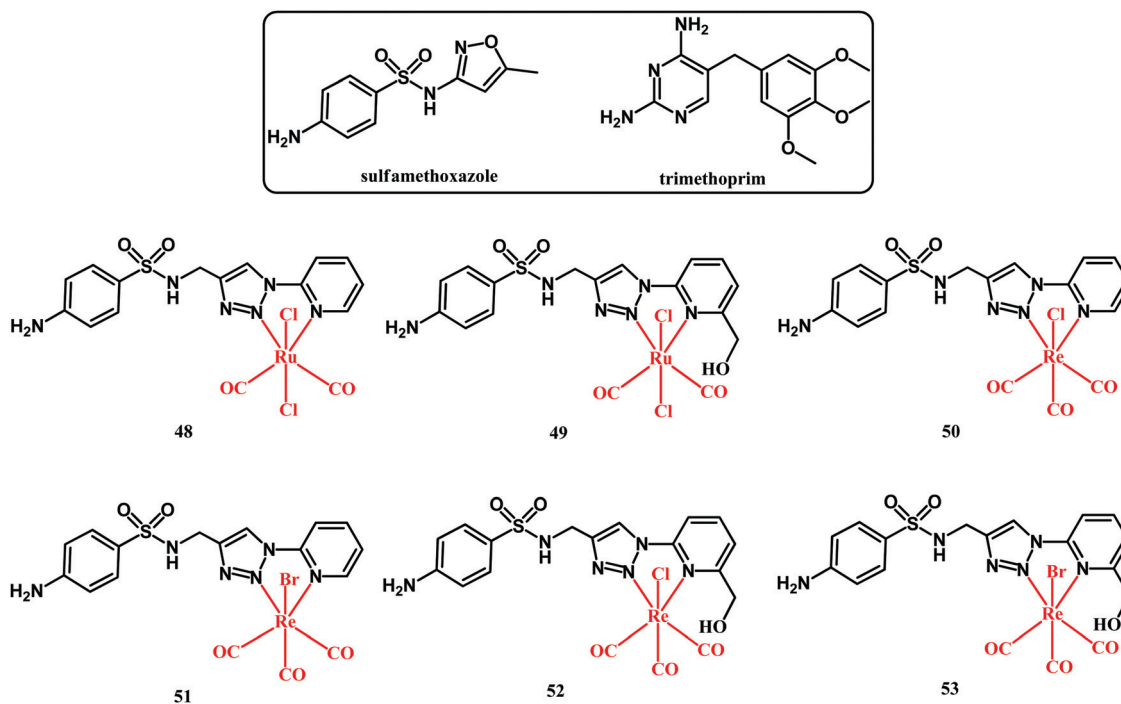


Fig. 14 Structures of sulfamethoxazole, trimethoprim and compounds 48–53.



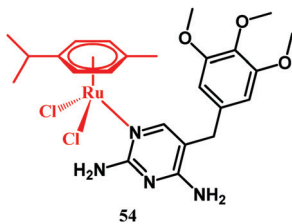


Fig. 15 Structure of compound 54.

showed similar activity toward all four *S. aureus* strains including the MRSA ATCC 43300 strain. In the latter case, cotrimoxazole had a MIC of 960:48 μM , whereas the MICs for the 50:trimethoprim and 51:trimethoprim formulations were 98:4.9 and 46:2.3 μM , respectively. This is an indication of a significant anti-MRSA activity increase. Although the mechanism behind such an activity increase was not given, the obtained results pinpoint the advantages offered by combined organometallic-organics for antibacterial therapy.

Recently Das, Ganeshpandian and co-workers reported on half-sandwich $\text{Ru}(\eta^6\text{-}p\text{-cymene})$ -trimethoprim compound 54 (Fig. 15).¹²⁵ The complex has been tested alone as well as after being loaded into polydiacetylene(PDA)-based liposomes against healthy human HEK-293, lung adenocarcinoma A549, breast carcinoma MCF-7 and liver carcinoma HepG2 cells. Compound 54 showed cytotoxicity toward healthy HEK-293 cells. Yet, this undesirable effect was diminished by the liposomal formulation of the drug. Neither 54 alone nor when encapsulated in liposomes showed activity against A549 and MCF-7 cells. On the contrary, compound 54 showed cytotoxic activity against liver carcinoma HepG2 cells. This activity was potentiated by the encapsulation of 54 into liposomal micelles. Insight into the action of compound 54 revealed that the Ru complex shows proapoptotic induction in HepG2 cells together with the ability of DNA scission. The antibacterial activity of 54 alone and when loading 54 into a liposomal carrier has been tested with the agar disk diffusion method against *S. aureus* and *P. aeruginosa* strains. The selection of these two strains was

rationalized due to their prevalence as one of the most common pathogens which are responsible for nosocomial infections in patients with cancer.¹²⁶ Rather unexpectedly, ruthenium trimethoprim complex 54 showed no improvement in antibacterial activity over the trimethoprim drug alone. The zone of inhibition for trimethoprim was 26 and 29 mm against the *S. aureus* and *P. aeruginosa* strains, respectively, at a 100 μM concentration. At the same high concentration, the zone of inhibition for 54 was 23 and 20 mm, respectively. Likewise, almost the same diameters of inhibition were obtained for 54 and trimethoprim at the lowest concentrations tested. For completeness, the liposome loaded 54 showed no antibacterial activity even at a 300 μM concentration. This can likely be explained by the hindered uptake of the liposome through the bacterial envelope. Although the mechanism of antibacterial activity of 54 was not examined, it can be predicted that the trimethoprim portion of the complex acts as a folate synthesis inhibitor, whereas the role of the $\text{Ru}(\eta^6\text{-}p\text{-cymene})$ portion could be more speculative and may pertain to DNA/protein metalation.

3. Derivatives of non-antibacterial drugs (metformin, phenformin and aspirin)

Biguanides are a group of nitrogen-rich organic compounds with established pharmacological relevance. Belonging to this class are metformin and phenformin, two drugs used for diabetes treatment. In 2018, Sadler and co-workers reported on the synthesis and antimicrobial activity studies of a series of biguanide $\text{Ir}(\text{II})$ complexes.¹²⁷ Within the studied series of compounds, metformin and phenformin complexes 55–57 and 58 (Fig. 16) were evaluated, respectively. The complexes were assayed against a wide panel of G-negative bacteria including *E. coli*, *K. pneumoniae*, *P. aeruginosa*, and *A. baumannii*, and G-positive bacteria *B. subtilis*, *S. pyogenes*, *E. faecalis*, *S. epidermidis* and *S. aureus* (MSSA and MRSA), as well as

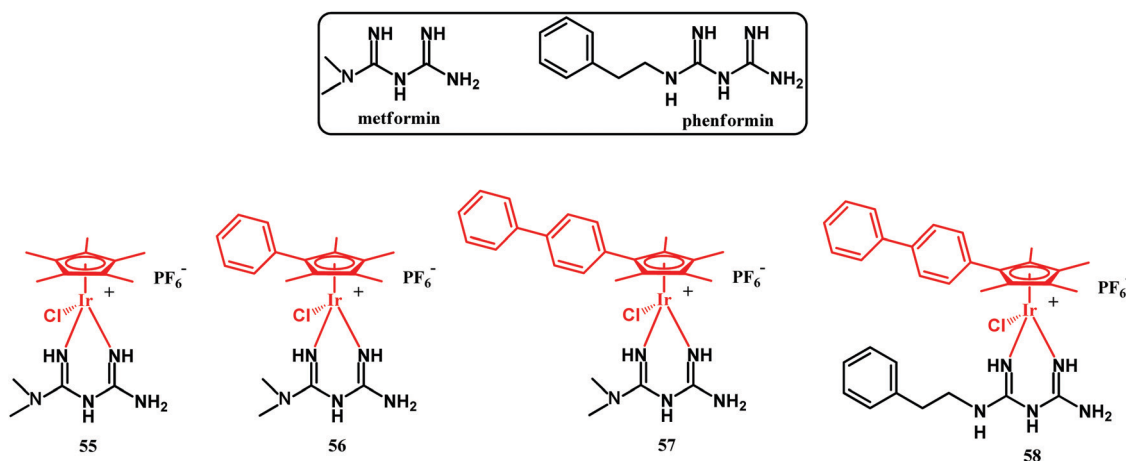


Fig. 16 Structures of metformin, phenformin and compounds 55–58.



against the pathogenic fungi *Candida albicans* and *Cryptococcus neoformans*.

The more hydrophilic complexes **55** and **56** were weakly active against the tested bacterial strains, with MICs over 54.3 μM . The more hydrophobic biphenyl derivative **57** showed instead superior antibacterial activity, with MICs in the range 3.2 to 12.6 μM against Gram-positive bacteria and 25 μM against an *E. coli* strain. The better activity of **57** can be explained by the increased uptake of the compound into bacterial cells. Likewise, even the more hydrophobic phenformin complex **58** showed superior antibacterial activity against Gram-positive bacteria with MICs in the range 0.17 to 2.7 μM . It also showed increased activity against the Gram-negative pathogens *E. coli* (MIC = 5.4 μM), *K. pneumoniae* (MIC = 21.6 μM), *P. aeruginosa* (MIC = 43.2 μM) and *A. baumannii* (MIC = 5.4 μM). Combined fluorescence microscopy and transmission electron microscopy (TEM) studies showed that the Ir(III) biguanide complexes do not disrupt the bacterial cell wall. Instead, Ir(III) biguanides can act as delivery systems to bacterial cells, where they dissociate to release free biguanidine ligands and reactive Ir(III) species. The latter can react with thiol-containing biomolecules, e.g., L-cysteine, to form dimers or other byproducts.¹²⁷ On the other hand, the released biguanide can bind cellular metal ions and inhibit vital metabolic pathways in bacteria, which ultimately leads to the experimentally observed bactericidal effect. Importantly, as studied by Sadler, Ir(III) biguanide complexes showed overall low toxicity toward mammalian cells as well as low hemolytic activity. The Ir(III) biguanide complexes stand as an example of repurposing essentially non-antibacterial drugs into highly active antibacterial agents.

Another recent example of an organometallic derivative of a non-antibacterial drug which was examined for antibacterial activity is the aspirin Re(I) phenanthroline complex **59** (Fig. 17).¹²⁸

It has been reported by Kowalski and co-workers, with the main aim to study its optical properties, its accumulation in living mammalian cells by confocal microscopy and its anti-cancer activity. In connection with these major studies, the antibacterial activity of **59** was screened against *S. aureus* ATCC 6538 and ATCC 29213 and *E. coli* NCTC 8196 model strains.

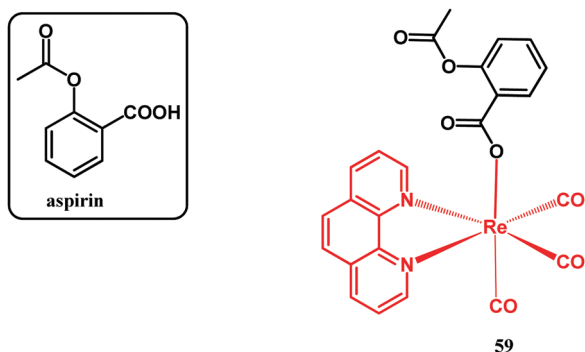


Fig. 17 Structure of aspirin and compound **59**.

The examined complex showed activity with a MIC of 50.8 μM against all strains tested. This activity was better than that of nitrofurantoin and weaker than that of ampicillin, which were used as references in the course of the assays.

4. Metal-containing antibacterially active materials

Metals have been used as antimicrobial materials since antiquity, mainly for water disinfection, food preservation, crop protection, and health care. Their toxicity depends on several parameters including their physicochemical form (*i.e.*, metal speciation), either in their elemental or oxidized states, their affinity to bind DNA, and their ability to disrupt membrane function, to bind donor ligands (including extra- and intracellular proteins and enzymes), to impair nutrient uptake, to alter signal transduction, and to induce oxidative stress, among others.¹²⁹ In their elemental (zerovalent) form, metals can form nanoparticles under a controlled environment which confines crystal nucleation and growth. Those nanoparticles due to their large area per volume ratio are prone to rapid oxidation in physiological media and consequently their released ions are chemisorbed on the surface of the metal nanoparticles, the nanoparticle acting as a reservoir for the sustained release of oxidized species.¹³⁰ Once in solution, their oxidation state can also change upon contact with cells. In this regard, Balfourier and co-workers demonstrated that gold nanoparticles are rapidly endocytosed inside endosomes and biodegraded due to the acidic environment and the participation of NADPH oxidase as catalyst for the production of the highly oxidizing superoxide free radicals.¹³¹ Subsequently, metal-binding proteins (*i.e.*, metallothioneins) are able to biomineralize dissolved species to form recrystallized nanoparticles.

The oxidation state of metal nanoparticles is key to determine their antimicrobial action. In this regard, Xiu and co-workers demonstrated that under anaerobic incubation conditions the antimicrobial action of silver nanoparticles against *E. coli* strain K12 (ATCC 25404) was impaired due to the lack of ionic silver.¹³² Direct contact also plays a key role in the antimicrobial action. We demonstrated that silver and gold-loaded chitosan films were efficient in the elimination of *Staphylococcus aureus* ATCC 6538 and 9213 strains, whereas, under the same conditions, the exudates released from those films were unable to reduce bacterial contamination.^{133,134}

The antimicrobial action also depends on the availability of the cytotoxic released ions. In 19th-century Europe, silver nitrate was used as eyedrops in newborns for the prevention of ophthalmia, reducing its incidence by a factor of 20–30.¹³⁵ In this case, ionic silver was immediately available in aqueous solution, but, later on, sustained release systems have been developed using different materials as ion hosts to reduce the large cytotoxicity associated with uncontrolled release. Inorganic (*e.g.*, zeolites, clays, *etc.*), inorganic–organic hybrid (MOFs, POMs, *etc.*) and organic (metal exchange resins) hosts are used to provide sustained release of antimicrobial ionic



metals. Different application sectors have incorporated those materials to take advantage of their long-term antimicrobial action.

As we mentioned before, antimicrobial metals are commonly used in water disinfection, food preservation, crop protection, and health care. It is important to point out that not only the metal itself is responsible for its antimicrobial action but also its surface chemistry including ionic metallic species chemisorbed on the surface and also the presence of different ligands used during surface functionalization. In this section of the review we will focus only on health care applications including antimicrobial medical devices, surfaces and indoor air pollutant control with special emphasis on those studies which have been assessed in clinical settings.

4.1. Antibiofouling medical surfaces and indoor air pollutant control

Medical surfaces can be coated with metal containing paints as a prophylactic measure to prevent biofouling. Zinc oxide and titanium oxide are the most frequent semiconductor metal oxides used as nanofillers in paints to prevent healthcare associated nosocomial infections in healthcare facilities. Those nanofillers are added to modify the rheological properties of the paints, promoting fast drying and avoiding paint drips, and also to take advantage of their photocatalytic action against pathogenic microorganisms. Recent advances focus on the development of highly active visible-light induced photocatalytic coatings based on those materials instead of the traditional UV-activated ones. In this regard, Krumdieck and co-workers developed intricate titanium dioxide nanostructures containing carbon and prepared by chemical vapor deposition on stainless steel doors and faucet handles, rendering a 100 times larger surface area and enhanced antimicrobial action compared to conventional titania coatings and powders.¹³⁶ However, Hu and co-workers have recently shown that non harmful bacteria can become harmful (*i.e.*, spore forming bacteria) when using antimicrobial paints and they state that we should be judicious in the use of antimicrobial products.¹³⁷ Most antimicrobial paints are validated using model Gram + (*e.g.*, *S. aureus*) and Gram – bacteria (*e.g.*, *E. coli*), but it is important to recall that a large amount of known bacteria can remain in a viable but nonculturable state. In this regard, Robben and co-workers demonstrated that commercially available household cleaners in combination with inorganic salts can induce a viable but non-culturable state in five human pathogens.¹³⁸

Besides Ti and Zn, several other metals have been reported as antimicrobials on medical wards. Copper nanoparticles and copper-zeolite nanocomposites have been used as nanofillers within polymeric paints to render antimicrobial coatings on plastic waiting room chairs and on metal hospital IV pools, respectively, reducing the total viable microorganisms present, regardless of the microorganism tested.¹³⁹ A randomized control trial between 2010 and 2011 in the ICUs of 3 hospitals was carried out by distributing patients admitted in those units in rooms with or without copper alloy surfaces, and the rates of incident acquired infections and/or colonization with MRSA or

vancomycin-resistant *Enterococcus* (VRE) in each type of room were compared.¹⁴⁰ Patients cared for in rooms having copper alloy surfaces (*i.e.*, bed rails, overbed tables, IV poles, and arms of the visitor's chair) had a significantly lower rate of incident infection and/or colonization than patients treated in standard rooms. A prospective cohort study involving 621 patients hospitalized in a medical intensive care unit having titania-based photocatalysts coated on high touch surfaces and walls also revealed that the MRSA acquisition rate was significantly reduced.¹⁴¹

Commercial formulations of silver including AgION Technology's AgION™ (*i.e.*, silver/zinc ions contained in an LTA type zeolite) have been demonstrated as efficient antimicrobial coatings.¹⁴² Silver exchanged Y-type zeolites have not only been used on surfaces but also as filters to remove airborne pathogens (bacteria and fungi) in medical facilities.¹⁴³ Titania-coated cordierite foams have also been used to photo-catalytically degrade gaseous acetaldehyde, which is associated with sick building syndrome, and airborne or droplet-based infectious pathogens: *E. coli*, *P. aeruginosa*, *L. pneumophila*, *K. pneumoniae*, and MRSA.¹⁴⁴

4.2. Antimicrobial medical devices and metal-based treatments

The extensive medicinal use of metals was displaced with the discovery of antibiotics, although there are still several medical devices in use which incorporate metals for the treatment of pathogenic microorganisms.

Silver-based catheters (*e.g.*, AcryMed's SilvaGard™) are widely used in clinical settings. A 12-month randomized cross-over trial with 27 878 patients compared rates of nosocomial catheter-associated urinary tract infection in patients with silver-coated and uncoated catheters, revealing a decrease in the risk of infection by 21% among study wards randomized to silver-coated catheters and by 32% among patients in whom silver-coated catheters were used on the wards.¹⁴⁵ A meta-analysis of 117 reports and eight trials with a total of 2355 patients satisfying the inclusion criteria revealed that silver alloy catheters are significantly more effective in preventing urinary tract infections than silver oxide catheters are.¹⁴⁶ Silver-containing catheters are recommended for short term use only and their benefit for patients with long-term catheters remains unclear.

Endotracheal tubes (ETTs) used in mechanically ventilated patients can be colonized by biofilm forming bacteria and thus contribute to the development of ventilator-associated pneumonia. Thorarinsdottir and co-workers reported that compared to uncoated PVC-based ETTs, the use of noble-metal-coated (containing silver, gold, and palladium (Bactiguard® AB, Sweden)) PVC-based ETTs was associated with reduced high-grade biofilm formation, although no significant difference was observed between silicon-coated ETTs and noble-metal-coated ETTs.¹⁴⁷ ETTs containing silver sulfadiazine in the interior of their lumen showed the lack of bacterial biofilm in 23 patients intubated in intensive care units following a phase I–II randomized clinical trial.¹⁴⁸



Silver coated tumor endoprostheses have been associated with a lower rate of early periprosthetic infection in 85 patients and also debridement with antibiotic treatment and retention of the implant appeared to be more successful with those silver-coated implants.¹⁴⁹ Implantcast Ltd's Mutars[®] (*i.e.*, silver coated bone replacements) showed a reduced rate of infection in 51 sarcoma patients receiving those megaprotheses compared to those receiving uncoated titanium controls.¹⁵⁰

Silver sulfadiazine, as a topical antiseptic used in partial and full thickness burns to prevent infection, promotes regeneration and reduces inflammation. Several common formulations include silver, copper and bismuth in their formulation and are commercialized as topical antiseptics (*e.g.*, Silvadene[®], Xeroform[®], *etc.*).¹⁵¹ Chronic wounds are difficult to treat and frequently polymicrobial populations are colonizing those and tissue debridement and antibiotic therapy fail in their management. Antimicrobial wound dressings have been designed to release antimicrobial compounds and some of them include metals in their formulation. To date, more studies are needed to corroborate that antimicrobial-releasing dressings are more clinically and cost efficient than conventional wound dressings in the management of chronic infected wounds.¹⁵² For instance, after the analysis of 12 randomized controlled trials reporting 13 comparisons, O'Meara and co-workers concluded that current evidence does not support the use of silver-based products in the management of venous leg ulcers.¹⁵³

Metal ions are also used in the treatment of eye infections and in periodontal and peri-implant diseases. Eye drops, mouthwashes, dentifrices, dental implants and delivery devices have incorporated metallic ions (*i.e.*, silver, copper, zinc, *etc.*) to prevent infection, but special attention should be paid to their potential toxicological effects.¹⁵⁴

Not only are metals incorporated into devices but also they can be applied systemically. In this regard, a phase 1 clinical trial in individuals with cystic fibrosis and chronic *P. aeruginosa* airway infections intravenously treated with gallium (as a disruptor of bacterial iron metabolism) revealed improved lung function inhibiting *P. aeruginosa* growth.¹⁵⁵ The use of iron chelators (such as gallium) and other metals as siderophore synthesis inhibitors opens new avenues in the management of pathogenic bacteria.

4.3. Photodynamic and photothermal antimicrobial therapy based on metal nanoparticles

PDT uses light and tissue oxygen to generate ROS with the aid of organic photosensitizers. Those organic molecules (*e.g.*, porphyrins, chlorophylls and dyes) are prone to photobleaching and, consequently, metal nanoparticles have been introduced in this field to avoid such a limitation. Upconversion nanoparticles (*i.e.*, rare-earth based lanthanide- or actinide-doped transition metals) and large band-gap semiconductor nanoparticles (*e.g.*, zinc oxide, titanium oxide, copper oxide, quantum dots, *etc.*) have been used in the photoinactivation of pathogenic bacteria usually colonizing infected topical wounds due to the inherent limitation in the light penetration depth achievable. For instance, rapid sterilization and accelerated

wound healing have been reported by using Zn²⁺ and graphene oxide in the management of bacteria-infected wounds using preclinical models.¹⁵⁶ Mao and co-workers recently reported the use of MOFs for the *in vivo* labelling of bacteria and simultaneous photodynamic treatment and therapy guided observation by using fluorescence imaging.¹⁵⁷ PEGylated W₁₈O₄₉ nanosheets have been used for the multi-modal imaging of the gastrointestinal tract and imaging-guided photothermal (PTT) sterilization *in vivo* using externally applied near infrared light.¹⁵⁸ The combination of both PDT and PTT is also possible by combining photothermal MoS₂ films and an organic photosensitizer (IR780) assisted by glutathione oxidation accelerated by NIR light, rendering synergistic and rapid killing of *Staphylococcus aureus* biofilms *in vivo*.¹⁵⁹

Despite all those successful approaches, future research should be focused on the synthesis of metal nanoparticles having reduced toxicity and fast biodegradability in order to compete with conventional organic photosensitizers.

6. Conclusion and outlook

This review primarily focuses on organometallic drug derivatives and metal-containing materials having antibacterial activity. In addition, this antibacterial action is also reported for non-antibacterial drugs like metformin, phenformin and aspirin. The data discussed herein show that the combination of an organometallic moiety with an organic pharmacophore (drug) in many cases results in derivatives which are able to circumvent drug resistance in bacteria. This activity is achieved by the bimodal (or multimodal) mode of action of organometallic–drug conjugates. In such cases, the chances of developing antibiotic resistance are reduced because simultaneous bacterial mutations against several mechanisms of action are improbable. On the other hand, the impaired uptake of some organometallic–drug derivatives and their potential toxicity against mammalian cells hinder the full exploration of their antibacterial activity in the clinic. This review also shows that metals and metal nanoparticles can benefit traumatology, orthopedic surgery, wound management, and ocular, periodontal and respiratory diseases thanks to their prophylactic and bactericidal use; however, despite all recent advances and even having several metal-based products on the market, more multicenter prospective clinical trials are needed to validate the large amount of scientific literature encompassing metal based antimicrobial materials.

Conflicts of interest

There are no conflicts to declare.

Acknowledgements

Support from University of Łódź, Poland is gratefully acknowledged.



References

- 1 N. J. Planavsky, D. Asael, A. Hofmann, C. T. Reinhard, S. V. Lalonde, A. Knudsen, X. Wang, F. O. Ossa, E. Pecoits, A. J. B. Smith, N. J. Beukes, A. Bekker, T. M. Johnson, K. O. Konhauser, T. W. Lyons and O. J. Rouxel, *Nat. Geosci.*, 2014, **7**, 283–286.
- 2 C. Thomazo, M. Ader and P. Philippot, *Geobiology*, 2011, **9**, 107–120.
- 3 K. Lepot, *Earth-Sci. Rev.*, 2020, **209**, 103296.
- 4 K. Lepot, A. Addad, A. H. Knoll, J. Wang, D. Troadec, A. Béché and E. J. Javaux, *Nat. Commun.*, 2017, **8**, 14890–14900.
- 5 H. W. Jannasch and M. J. Mottl, *Science*, 1985, **229**, 717–725.
- 6 E. Spieck, M. Spohn, K. Wendt, E. Bock, J. Shively, J. Frank, D. Indenbirken, M. Alawi, S. Lücker and J. Hüpeden, *ISME J.*, 2020, **14**, 364–379.
- 7 A. Schippers, L. N. Neretin, J. Kallmeyer, T. G. Ferdelman, B. A. Cragg, R. J. Parkes and B. B. Jørgensen, *Nature*, 2005, **433**, 861–864.
- 8 K. R. Redeker, J. P. J. Chong, A. Aguion, A. Hodson and D. A. Pearce, *J. R. Soc., Interface*, 2017, **14**, 20170729.
- 9 D. A. Cowan, *Proc. Natl. Acad. Sci. U. S. A.*, 2009, **106**, 19749–19750.
- 10 J. K. Fredrickson, J. M. Zachara, D. L. Balkwill, D. Kennedy, S. W. Li, H. M. Kostandarithes, M. J. Daly, M. F. Romine and F. J. Brockman, *Appl. Environ. Microbiol.*, 2004, **70**, 4230–4241.
- 11 R. Conrad and A. V. Vlassov, *Med. Sci. Rev.*, 2015, **2**, 92–103.
- 12 H. G. Groot, Y. J. van de Vegte, N. Verweij, E. Lipsic, J. C. Karper and P. van der Harst, *Sci. Rep.*, 2020, **10**, 14771–14779.
- 13 C. Pleguezuelos-Manazo, J. Puschhof, A. R. Huber, A. van Hoeck, H. M. Wood, J. Nomburg, C. Gurjao, F. Manders, G. Dalmaso, P. B. Stege, F. L. Paganelli, M. H. Geurts, J. Beumer, T. Mizutani, Y. Miao, R. van der Linden and S. van der Elst, Genomics England Research Consortium, K. C. Garcia, J. Top, R. J. L. Willems, M. Giannakis, R. Bonnet, P. Quirke, M. Meyerson, E. Cuppen, R. van Boxtel and H. Clevers, *Nature*, 2020, **580**, 269–273.
- 14 M. A. Spyrou, K. I. Bos, A. Herbig and J. Krause, *Nat. Rev. Genetics*, 2019, **20**, 323–340.
- 15 A. Fleming, *Rev. Infect. Dis.*, 1929, **2**, 129–139.
- 16 M. Barber, *Lancet*, 1948, **2**, 641–644.
- 17 A. Schatz, E. Bugie and S. A. Waksman, *Exp. Biol. Med.*, 1944, **55**, 66–69.
- 18 J. Cotton and D. A. Mitchison, *Br. Med. J.*, 1948, **2**, 1009–1015.
- 19 M. P. Jevons, *Br. Med. J.*, 1961, **1**, 124–125.
- 20 M. T. Parker and J. H. Hewitt, *Lancet*, 1970, **295**, 800–804.
- 21 L. Elwell, M. Roberts, L. Mayer and S. Falkow, *Antimicrob. Agents Chemother.*, 1977, **11**, 528–533.
- 22 J. M. Blair, M. A. Webber, A. J. Baylay, D. O. Ogbolu and L. J. Piddock, *Nat. Rev. Microbiol.*, 2015, **13**, 42–51.
- 23 B. Marshall, M. Roberts, A. Smith and S. B. Levy, *J. Infect. Dis.*, 1984, **149**, 1028–1029.
- 24 M. D. Iseman, *N. Eng. J. Med.*, 1993, **329**, 784–791.
- 25 D. M. P. De Oliveira, B. M. Forde, T. J. Kidd, P. N. A. Harris, M. A. Schembri, S. A. Beatson, D. L. Paterson and M. J. Walker, *Clin. Microbiol. Rev.*, 2020, **33**, e00181.
- 26 S. B. Levy and B. Marshall, *Nat. Med.*, 2004, **10**, 122–129.
- 27 WHO. WHO's global report on antimicrobial resistance 2019, <https://www.who.int/antimicrobial-resistance/interagency-coordination-group/final-report/en/>.
- 28 P. Wiggins, *Exp. Opin. Invest. Drugs*, 2004, **13**, 889–902.
- 29 D. J. Hoban, D. J. Biedenbach, A. H. Mutnick and R. N. Jones, *Microbiol. Infect. Dis.*, 2003, **45**, 279–285.
- 30 D. M. Morens, G. K. Folkers and A. S. Fauci, *Nature*, 2004, **430**, 242–249.
- 31 S. Kumari and V. J. Ram, *Drugs Today*, 2004, **40**, 487–500.
- 32 D. A. Mitchison, *Am. J. Respir. Crit. Care Med.*, 2005, **171**, 699–706.
- 33 S. B. Levy, *Sci. Am.*, 1998, **278**, 46–53.
- 34 K. Bush, *Clin. Infect. Dis.*, 2001, **32**, 1085–1089.
- 35 D. L. Paterso, W.-C. Ko, A. Von Gottberg, S. Mohapatra, J. M. Casellas, H. Goossens, L. Mulazimoglu, G. Trenholme, K. P. Klugman, R. A. Bonomo, L. B. Rice, M. M. Wagener, J. G. McCormack and V. L. Yu, *Ann. Intern. Med.*, 2004, **140**, 26–32.
- 36 P. A. Bradford, *Clin. Microbiol. Rev.*, 2001, **14**, 933–951.
- 37 J. F. Barrett, *Exp. Opin. Ther. Targets*, 2004, **8**, 515–519.
- 38 C. D. Salgado, B. M. Farr and D. P. Calfee, *Clin. Infect. Dis.*, 2003, **36**, 131–139.
- 39 C. G. Whitney, M. M. Farley, J. Hadler, L. H. Harrison, C. Lexau, A. Reingold, L. Lefkowitz, P. R. Cieslak, M. Cetron, E. R. Zell, J. H. Jorgensen and A. Schuchat, *N. Engl. J. Med.*, 2000, **343**, 1917–1924.
- 40 K. A. Gordon, D. J. Bidenbach and R. N. Jones, *Diagn. Microbiol. Infect. Dis.*, 2003, **46**, 285–289.
- 41 R. J. Anderson, P. W. Groundwater, A. Todd and A. J. Worsley, *Antibacterial Agents. Chemistry, Mode of Action, Mechanisms of Resistance and Clinical Applications*, Wiley and Sons, Ltd, Chichester, West Sussex, UK, 2012.
- 42 S. Levy, *The Antibiotic Paradox: How Misuse of Antibiotics Destroys their Curative Powers*, Perseus, Cambridge, 2002.
- 43 F. von Nussbaum, M. Brands, B. Hinzen, S. Weigand and D. Häbich, *Angew. Chem., Int. Ed.*, 2006, **45**, 5072–5129.
- 44 C. Wetzel, M. Lonneman and C. Wu, *Eur. J. Med. Chem.*, 2021, **209**, 112931.
- 45 T. T. Tran, D. Panesso, N. N. Mishra, E. Mileykovskaya, Z. Guan, J. M. Munita, J. Reyes, L. Diaz, G. M. Weinstock, B. E. Murray, Y. Shamoo, W. Dowhan, A. S. Bayer and C. A. Arias, *mBio-Am. Soc. Microbiol.*, 2013, **4**, e00281–13.
- 46 V. G. Meka and H. S. Gold, *Clin. Infect. Dis.*, 2004, **39**, 1010–1015.
- 47 Y. Imai, K. J. Meyer, A. Iinishi, Q. Favre-Godal, R. Green, S. Manuse, M. Caboni, M. Mori, S. Niles, M. Ghiglieri, C. Honrao, X. Ma, J. J. Guo, A. Makriyannis, L. Linares-Otoya, N. Böhringer, Z. G. Wuisan, H. Kaur, R. Wu, A. Mateus, A. Typas, M. M. Savitski, J. L. Espinoza, A. O'Rourke, K. E. Nelson, S. Hiller, N. Noinaj,



- T. F. Schäberle, A. D'Onofrio and K. Lewis, *Nature*, 2019, **576**, 459–464.
- 48 L. L. Ling, T. Schneider, A. J. Peoples, A. L. Spoering, I. Engels, B. P. Conlon, A. Mueller, T. F. Schäberle, D. E. Hughes, S. Epstein, M. Jones, L. Lazarides, V. A. Steadman, D. R. Cohen, C. R. Felix, K. A. Fetterman, W. P. Millett, A. G. Nitti, A. M. Zullo, C. Chen and K. Lewis, *Nature*, 2015, **517**, 455–459.
- 49 S. Gibbons, *Nat. Prod. Rep.*, 2013, **30**, 988–1027.
- 50 K. M. G. O'Connell, J. T. Hodgkinson, H. F. Sore, M. Welch, G. P. C. Salmond and D. R. Spring, *Angew. Chem., Int. Ed.*, 2013, **52**, 10706–10733.
- 51 J. T. Staley and A. Konopka, *Annu. Rev. Microbiol.*, 1985, **39**, 321–346.
- 52 F. Cieplik, D. Deng, W. Crielaard, W. Buchalla, E. Hellwig, A. Al-Ahmad and T. Maisch, *Crit. Rev. Microbiol.*, 2018, **44**, 571–589.
- 53 N. A. Smith, P. Zhang, S. E. Greenough, M. D. Horbury, G. J. Clarkson, D. McFeely, A. Habtemariam, L. Salassa, V. G. Stavros, C. G. Dowson and P. J. Sadler, *Chem. Sci.*, 2017, **8**, 395–404.
- 54 E. Sauvageot, M. Elie, S. Gaillard, R. Daniellou, P. Fechter, I. J. Schalk, V. Gasser, J.-L. Renaud and G. L. A. Mislin, *Metallomics*, 2017, **9**, 1820–1827.
- 55 S. Wu, A. Li, X. Zhao, C. Zhang, B. Yu, N. Zhao and F.-J. Xu, *ACS Appl. Mater. Interfaces*, 2019, **11**, 17177–17183.
- 56 M. Tinajero-Trejo, N. Rana, C. Nagel, H. E. Jesse, T. W. Smith, L. K. Wareham, M. Hippler, U. Schatzschneider and R. K. Poole, *Antioxid. Redox Signaling*, 2016, **24**, 765–780.
- 57 J. Betts, C. Nagel, U. Schatzschneider, R. Poole and R. M. la Ragione, *PLoS One*, 2017, **12**, e0186359.
- 58 N. Rana, H. Jesse, M. Tinajero-Trejo, J. Butler, M. L. von und zur Mühlen, C. Nagel, U. Schatzschneider and R. K. Poole, *Microbiology*, 2017, **163**, 1477–1489.
- 59 A. Regiel-Futyra, J. M. Dąbrowski, O. Mazuryk, K. Śpiwak, A. Kyzioł, B. Pucelik, M. Brindell and G. Stochel, *Coord. Chem. Rev.*, 2017, **351**, 76–117.
- 60 A. Frei, *Antibiotics*, 2020, **9**, 90.
- 61 M. B. Harbut, C. Vilchère, X. Luo, M. E. Hensler, H. Guo, B. Yang, A. K. Chatterjee, V. Nizet, W. R. Jacobs Jr., P. G. Schultz and F. Wang, *Proc. Natl. Acad. Sci. U. S. A.*, 2015, **112**, 4453–4458.
- 62 F. Li, J. G. Collins and F. R. Keene, *Chem. Soc. Rev.*, 2015, **44**, 2529–2542.
- 63 M. Patra, G. Gasser and N. Metzler-Nolte, *Dalton Trans.*, 2012, **41**, 6350–6358.
- 64 M. A. Sierra, L. Casarrubios and M. C. de la Torre, *Chem. – Eur. J.*, 2019, **25**, 7232–7242.
- 65 U. Schatzschneider, Antimicrobial activity of organometal compounds: Past, present, and future prospects, in *Advances in Bioorganometallic Chemistry*, ed. T. Hirao and T. Moriuchi, Elsevier, Amsterdam, 2018, ISBN 978-0-12-814197-7.
- 66 B. S. Ludwig, J. D. G. Correia and F. E. Kühn, *Coord. Chem. Rev.*, 2019, **396**, 22–48.
- 67 A. Frei, J. Zuegg, A. G. Elliott, M. Baker, S. Braese, C. Brown, F. Chen, C. G. Dowson, G. Dujardin, N. Jung, A. P. King, A. M. Mansour, M. Massi, J. Moat, H. A. Mohamed, A. K. Renfrew, P. J. Rutledge, P. J. Sadler, M. H. Todd, C. E. Willans, J. J. Wilson, M. A. Cooper and M. A. T. Blaskovich, *Chem. Sci.*, 2020, **11**, 2627–2639.
- 68 B. Albada and N. Metzler-Nolte, *Acc. Chem. Res.*, 2017, **50**, 2510–2518.
- 69 P. Güntzel, C. Nagel, J. Weigelt, J. W. Betts, C. A. Patrick, H. M. Southam, R. M. La Ragione, R. K. Poole and U. Schatzschneider, *Metallomics*, 2019, **11**, 2033–2042.
- 70 Z. Ude, I. Romero-Canelón, B. Twamley, D. F. Hughes, P. J. Sadler and C. J. Marmion, *J. Inorg. Biochem.*, 2016, **160**, 210–217.
- 71 C. Schmidt, L. Albrecht, S. Balaupramaniam, R. Misgeld, B. Karge, M. Brönstrup, A. Prokop, K. Baumann, S. Reichl and I. Ott, *Metallomics*, 2019, **11**, 533–545.
- 72 C. Schmidt, B. Karge, R. Misgeld, A. Prokop, R. Franke, M. Brönstrup and I. Ott, *Chem. – Eur. J.*, 2017, **23**, 1869–1880.
- 73 F. Chen, J. Moat, D. McFeely, G. Clarkson, I. J. Hands-Portman, J. P. Furner-Pardoe, F. Harrison, C. G. Dowson and P. J. Sadler, *J. Med. Chem.*, 2018, **61**, 7330–7344.
- 74 Q. Laurent, L. K. Batchelor and P. J. Dyson, *Organometallics*, 2018, **37**, 915–923.
- 75 A. Frei, M. Amado, M. A. Cooper and M. A. T. Blaskovich, *Chem. – Eur. J.*, 2020, **26**, 2852–2858.
- 76 A. Mansour and K. Radacki, *Dalton Trans.*, 2020, **49**, 4491–4501.
- 77 A. Gupta, P. Prasad, S. Gupta and P. K. Sasmal, *ACS Appl. Mater. Interfaces*, 2020, **12**, 35967–35976.
- 78 Y. Zheng, W. Liu, Y. Chen, H. Jiang, H. Yan, I. Kosenko, L. Chekulaeva, I. Sivaev, V. Bregadze and X. Wang, *Organometallics*, 2017, **36**, 3484–3490.
- 79 G. Jaouen, A. Vessières and S. Top, *Chem. Soc. Rev.*, 2015, **44**, 8802–8817.
- 80 G. Gasser, I. Ott and N. Metzler-Nolte, *J. Med. Chem.*, 2011, **54**, 3–25.
- 81 E. J. Anthony, E. M. Bolitho, H. E. Bridgewater, O. W. L. Carter, J. M. Donnelly, C. Imberti, E. C. Lant, F. Lermyte, R. J. Needham, M. Palau, P. J. Sadler, H. Shi, F.-X. Wang, W.-Y. Zhang and Z. Zhang, *Chem. Sci.*, 2020, **11**, 12888–12917.
- 82 G. Palermo, A. Magistrato, T. Riedel, T. von Erlach, C. A. Davey, P. J. Dyson and U. Rothlisberger, *ChemMedChem*, 2016, **11**, 1199–1210.
- 83 E. Alessio, *Eur. J. Inorg. Chem.*, 2017, 1549–1560.
- 84 S. R. Martínez, A. M. Durantini, M. C. Becerra and G. Cosa, *ACS Infect. Dis.*, 2020, **6**, 2468–2477.
- 85 P. Belenky, J. D. Ye, C. B. M. Porter, N. R. Cohen, M. A. Lobritz, T. Ferrante, S. Jain, B. J. Korry, E. G. Schwarz, G. C. Walker and J. J. Collins, *Cell Rep.*, 2015, **13**, 968–980.
- 86 J. J. Foti, B. Devadoss, J. A. Winkler, J. J. Collins and G. C. Walker, *Science*, 2012, **336**, 315–319.
- 87 R. Rubbiani, B. Wahrig and I. Ott, *J. Biol. Inorg. Chem.*, 2014, **19**, 961–965.



- 88 R. Koch, *Dtsch. Med. Wochenstr.*, 1890, **16**, 756–757.
- 89 N. C. Lloyd, H. W. Morgan, B. K. Nicholson and R. S. Ronimus, *Angew. Chem., Int. Ed.*, 2005, **44**, 941–944.
- 90 B. Rosenberg, L. van Camp and T. Krigas, *Nature*, 1965, **205**, 698–699.
- 91 T. J. Kealy and P. L. Pauson, *Nature*, 1951, **168**, 1039–1040.
- 92 S. A. Miller, J. A. Tebboth and J. F. Tremaine, *J. Chem. Soc.*, 1952, 632–635.
- 93 G. Wilkinson, M. Rosenblum, M. C. Whiting and R. B. Woodward, *J. Am. Chem. Soc.*, 1952, **74**, 2125–2126.
- 94 E. O. Fischer and W. Pfab, *Z. Naturforsch., B: J. Chem. Sci.*, 1952, **7**, 377–379.
- 95 *Ferrocenes: Ligands, Materials, and Biomolecules*, ed. P. Štěpnička, Wiley-VCH, Chichester, 2008.
- 96 M. Patra and G. Gasser, *Nat. Rev. Chem.*, 2017, **1**, 0066.
- 97 K. Kowalski, *Coord. Chem. Rev.*, 2018, **366**, 91–108.
- 98 D. R. van Staveren and N. Metzler-Nolte, *Chem. Rev.*, 2020, **104**, 5931–5986.
- 99 M. Wenzel, M. Patra, C. H. R. Senges, I. Ott, J. J. Stepanek, A. Pinto, P. Prochnow, C. Vuong, S. Langklotz, N. Metzler-Nolte and J. E. Bandow, *ACS Chem. Biol.*, 2013, **8**, 1442–1450.
- 100 P. Macheboeuf, C. Contreras-Martel, V. Job, O. Dideberg and A. Dessen, *FEMS Microbiol. Rev.*, 2006, **30**, 673–691.
- 101 E. Sauvage, F. Kerff, M. Terrak, J. A. Ayala and P. Charlier, *FEMS Microbiol. Rev.*, 2008, **32**, 234–258.
- 102 L. I. Llarrull, S. A. Testero, J. F. Fisher and S. Mobashery, *Curr. Opin. Microbiol.*, 2010, **13**, 551–557.
- 103 Z. Yao, D. Kahne and R. Kishony, *Mol. Cell*, 2012, **48**, 705–712.
- 104 K. Bush and J. F. Fisher, *Annu. Rev. Microbiol.*, 2011, **65**, 455–478.
- 105 D. M. Livermore, *Clin. Microbiol. Rev.*, 1995, **8**, 557–584.
- 106 E. I. Edwards, R. Epton and G. Marr, *J. Organomet. Chem.*, 1976, **107**, 351–357.
- 107 E. M. Lewandowski, Ł. Szczupak, S. Wong, J. Skiba, A. Guśpiel, J. Solecka, V. Vrček, K. Kowalski and Y. Chen, *Organometallics*, 2017, **36**, 1673–1676.
- 108 J. D. Smith, M. Kumarasiri, W. Zhang, D. Heseck, M. Lee, M. Toth, S. Vakulenko, J. F. Fisher, S. Mobashery and Y. Chen, *Antimicrob. Agents Chemother.*, 2013, **57**, 3137–3146.
- 109 C. J. Adamski, A. M. Cardenas, N. G. Brown, L. B. Horton, B. Sankaran, B. V. V. Prasad, H. F. Gilbert and T. Palzkill, *Biochemistry*, 2015, **54**, 447–457.
- 110 E. M. Lewandowski, Ł. Szczupak, A. Kowalczyk, G. Mendoza, M. Arruebo, L. M. C. Jacobs, P. Stączek, Y. Chen and K. Kowalski, *ChemBioChem*, 2020, **21**, 2187–2195.
- 111 M. Michaut, A. Steffen, J.-M. Contreras, C. Morice, A. Paulen, I. J. Schalk, P. Plésiat and G. L. A. Mislin, *Bioorg. Med. Chem. Lett.*, 2020, **30**, 127098.
- 112 L. A. Mitscher, *Chem. Rev.*, 2005, **105**, 559–592.
- 113 D. J. Dwyer, M. Kohanski, B. Hayete and J. J. Collins, *Mol. Syst. Biol.*, 2007, **3**, 91.
- 114 Ł. Szczupak, A. Kowalczyk, D. Trzybiński, K. Woźniak, G. Mendoza, M. Arruebo, D. Steverding, P. Stączek and K. Kowalski, *Dalton Trans.*, 2020, **49**, 1403–1415.
- 115 R. Joshi, S. K. Yadav, H. Mishra, N. Pandey, R. Tilak and S. Pokharia, *Heteroatom Chem.*, 2018, **29**, 21433–21446.
- 116 M. P. Chrysouli, C. N. Banti, N. Kourkoumelis, E. E. Moushi, A. J. Tasiopoulos, A. Douvalis, C. Papachristodoulou, A. G. Hatzidimitriou, T. Bakas and S. K. Hadjikakou, *Dalton Trans.*, 2020, **49**, 11522–11535.
- 117 R. W. Lacey, *J. Antimicrob. Chemother.*, 1979, **5**(suppl. B), 75–83.
- 118 T. Stringer, R. Seldon, N. Liu, D. F. Warner, C. Tam, L. W. Cheng, K. M. Land, P. J. Smith, K. Chibale and G. S. Smith, *Dalton Trans.*, 2017, **46**, 9875–9885.
- 119 C. M. DuChane, G. W. Karpin, M. Ehrich, J. O. Falkinham III and J. S. Merola, *Med. Chem. Commun.*, 2019, **10**, 1391–1398.
- 120 C. Quintana, G. Silva, A. H. Klahn, V. Artigas, M. Fuentealba, C. Biot, I. Halloum, L. Kremer, N. Novoa and R. Arancibia, *Polyhedron*, 2017, **134**, 166–172.
- 121 P. Chellan, V. M. Avery, S. Duffy, J. A. Triccas, G. Nagalingam, C. Tam, L. W. Cheng, J. Liu, K. M. Land, G. J. Clarkson, I. Romero-Canelón and P. J. Sadler, *Chem. – Eur. J.*, 2018, **24**, 10078–10090.
- 122 R. G. Miller, M. Vázquez-Hernández, P. Prochnow, J. E. Bandow and N. Metzler-Nolte, *Inorg. Chem.*, 2019, **58**, 9404–9413.
- 123 R. Raz, B. Chazan, Y. Kennes, R. Colodner, E. Rottensterich, M. Dan, I. Lavi and W. Stamm, *Clin. Infect. Dis.*, 2002, **34**, 1165–1169.
- 124 M. A. Foltzer and R. E. Reese, *Med. Clin. North Am.*, 1987, **71**, 1177–1194.
- 125 D. Gopalakrishnan, C. Sumithaa, A. M. Kumar, N. S. P. Bhuvanesh, S. Ghorai, P. Das and M. Ganeshpandian, *New J. Chem.*, 2020, **44**, 20047–20059.
- 126 M. Kamboj and K. A. Sepkowitz, *Lancet Oncol.*, 2009, **10**, 589–597.
- 127 F. Chen, J. Moat, D. McFeely, G. Clarkson, I. J. Hands-Portman, J. P. Furner-Pardoe, F. Harrison, C. G. Dowson and P. J. Sadler, *J. Med. Chem.*, 2018, **61**, 7330–7344.
- 128 J. Skiba, A. Kowalczyk, P. Stączek, T. Bernaś, D. Trzybiński, K. Woźniak, U. Schatzschneider, R. Czerwieńec and K. Kowalski, *New J. Chem.*, 2019, **43**, 573–583.
- 129 J. Lemire, J. Harrison and R. Turner, *Nat. Rev. Microbiol.*, 2013, **11**, 371–384.
- 130 A. Henglein, *Chem. Mater.*, 1998, **10**, 444–450.
- 131 A. Balfourier, N. Luciani, G. Wang, G. Lelong, O. Ersen, A. Khelfa, D. Alloyeau, F. Gazeau and F. Carn, *Proc. Natl. Acad. Sci. U. S. A.*, 2020, **117**, 103–113.
- 132 Z. Xiu, Q. Zhang, H. L. Puppala, V. L. Colvin and P. J. J. Alvarez, *Nano Lett.*, 2012, **12**, 4271–4275.
- 133 A. Regiel, S. Irusta, A. Kyzioł, M. Arruebo and J. Santamaria, *Nanotechnology*, 2012, **24**, 015101.
- 134 A. Regiel-Futyr, M. Kus-Liśkiewicz, V. Sebastian, S. Irusta, M. Arruebo, G. Stochel and A. Kyzioł, *ACS Appl. Mater. Interfaces*, 2015, **7**, 1087–1099.
- 135 C. S. F. Crede, Die verhütung der augenentzündung der neugeborenen, *Arch. Gynakol.*, 1881, **17**, 50–53.
- 136 S. P. Krumdieck, R. Boichot, R. Gorthy, J. G. Land, S. Lay, A. J. Gardecka, M. I. J. Polson, A. Wasa, J. E. Aitken,



- J. A. Heinemann, G. Renou, G. Berthome, F. Charlot, T. Encinas, M. Braccini and C. M. Bishop, *Sci. Rep.*, 2019, **9**, 1883.
- 137 J. Hu, S. B. Maamar, A. J. Glawe, N. Gottel, J. A. Gilbert and E. M. Hartmann, *Indoor Air*, 2019, **29**, 551–562.
- 138 C. Robben, S. Fister, A. K. Witte, D. Schoder, P. Rossmannith and P. Mester, *Sci. Rep.*, 2018, **8**, 15132.
- 139 H. Palza, M. Nuñez, R. Bastías and K. Delgado, *Int. J. Antimicrob. Agents*, 2018, **51**, 912–917.
- 140 C. D. Salgado, K. A. Sepkowitz, J. F. John, J. R. Cantey, H. H. Attaway, K. D. Freeman, P. A. Sharpe, H. T. Michels and M. G. Schmidt, *Infect. Control Hosp. Epidemiol.*, 2013, **34**, 479–486.
- 141 M. H. Kim, S. G. Lee, K. S. Kim, Y. J. Heo, J. E. Oh and S. J. Jeong, *BMC Infect. Dis.*, 2018, **18**, 610.
- 142 M. M. Cowan, K. Z. Abshire, S. L. Houk and S. M. Evans, *J. Ind. Microbiol. Biotechnol.*, 2003, **30**, 102–106.
- 143 J. H. Shen, Y. S. Wang, J. P. Lin, S. H. Wu and J. J. Horng, *J. Air Waste Manag. Assoc.*, 2013, **64**, 13–18.
- 144 Y. Yao, T. Ochiai, H. Ishiguro, R. Nakano and Y. Kubota, *Appl. Catal., B*, 2011, **106**, 592–599.
- 145 T. B. Karchmer, E. T. Giannetta, C. A. Muto, B. A. Strain and B. M. Farr, *Arch. Intern. Med.*, 2000, **160**, 3294–3298.
- 146 S. Saint, J. G. Elmore, S. D. Sullivan, S. S. Emerson and T. D. Koepsell, *Am. J. Med.*, 1998, **105**, 236–241.
- 147 H. R. Thorarinsdottir, T. Kander, A. Holmberg, S. Petronis and B. Klarin, *Crit. Care*, 2020, **24**, 382.
- 148 L. Berra, T. Kolobow, P. Laquerriere, B. Pitts, S. Bramati, J. Pohlmann, C. Marelli, M. Panzeri, P. Brambillasca, F. Villa, A. Baccarelli, S. Bouthors, H. T. Stelfox, L. M. Bigatello, J. Moss and A. Pesenti, *Intensive Care Med.*, 2008, **34**, 1030–1037.
- 149 H. Wafa, R. J. Grimer, K. Reddy, L. Jeys, A. Abudu, S. R. Carter and R. M. Tillman, *Bone Joint J.*, 2015, **97-B**, 252–257.
- 150 J. Hardes, C. von Eiff, A. Streitbuerger, M. Balke, T. Budny, M. P. Henrichs, G. Hauschild and H. Ahrens, *J. Surg. Oncol.*, 2010, **101**, 389–395.
- 151 J. Cambiaso-Daniel, S. Boukavalas, G. H. Bitz, L. K. Branski, D. N. Herndon and D. M. Culnan, *Ann. Plast. Surg.*, 2018, **1**.
- 152 G. Norman, J. Christie, Z. Liu, M. J. Westby, J. M. Jefferies, T. Hudson, J. Edwards, D. P. Mohapatra, I. A. Hassan and J. C. Dumville, *Cochrane Database Syst. Rev.*, 2017, **7**, CD011821.
- 153 S. O'Meara, D. Al-Kurdi, Y. Ologun, L. G. Ovington, M. Martyn-St James and R. Richardson, *Cochrane Database Syst. Rev.*, 2014, **10**, CD003557.
- 154 R. P. Allaker, *J. Dent. Res.*, 2010, **89**, 1175–1186.
- 155 C. H. Goss, Y. Kaneko, L. Khuu, G. D. Anderson, S. Ravishankar, M. L. Aitken, N. Lechtzin, G. Zhou, D. M. Czyz, K. McLean, O. Olakanmi, H. A. Shuman, M. Teresi, E. Wilhelm, E. Caldwell, S. J. Salipante, D. B. Hornick, R. J. Siehnel, L. Becker, B. E. Britigan and P. K. Singh, *Sci. Transl. Med.*, 2018, **10**, eaat7520.
- 156 Y. Li, X. Liu, L. Tan, Z. Cui, X. Yang, Y. Zheng, K. W. K. Yeung, P. K. Chu and S. Wu, *Adv. Funct. Mater.*, 2018, **28**, 1800299.
- 157 D. Mao, F. Hu, Kenry, S. Ji, W. Wu, D. Ding, D. Kong and B. Liu, *Adv. Mater.*, 2018, **30**, 1706831.
- 158 Z. Liu, J. Liu, R. Wang, Y. Du, J. Ren and X. Qu, *Biomaterials*, 2015, **56**, 206–218.
- 159 M. Li, L. Li, K. Su, X. Liu, T. Zhang, Y. Liang, D. Jing, X. Yang, D. Zheng, Z. Cui, Z. Li, S. Zhu, K. W. K. Yeung, Y. Zheng, X. Wang and S. Wu, *Adv. Sci.*, 2019, **6**, 1900599.

