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## New $\beta$ -Lactamase Inhibitors in the Clinic

Krisztina M. Papp-Wallace, PhD<sup>a,b</sup> and Robert A. Bonomo, MD<sup>a,b,c,d,e,\*</sup>

<sup>a</sup>Research Service, Louis Stokes Cleveland Department of Veterans Affairs Medical Center, 10701 East Boulevard, Cleveland, OH 44106, USA

<sup>b</sup>Department of Medicine, Case Western Reserve University, 10900 Euclid Avenue, Cleveland, OH 44106, USA

<sup>c</sup>Department of Biochemistry, Case Western Reserve University, 10900 Euclid Avenue, Cleveland, OH 44106, USA

<sup>d</sup>Department of Pharmacology, Case Western Reserve University, 10900 Euclid Avenue, Cleveland, OH 44106, USA

<sup>e</sup>Department of Molecular Biology and Microbiology, Case Western Reserve University, 10900 Euclid Avenue, Cleveland, OH 44106, USA

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## WHAT TRANSPIRED? THE FALL OF THE CURRENT CLINICALLY AVAILABLE $\beta$ -LACTAM- $\beta$ -LACTAMASE INHIBITOR COMBINATIONS

In gram-negative pathogens, the production of  $\beta$ -lactamases, which hydrolyze  $\beta$ -lactam antibiotics, is a foremost threat in modern medicine.<sup>1–3</sup>  $\beta$ -Lactam- $\beta$ -lactamase inhibitor combinations (ie, amoxicillin-clavulanic acid, ampicillin-sulbactam, cefoperazone-sulbactam, piperacillin-tazobactam, and ticarcillin-clavulanic acid) were first introduced into the clinic in the 1980s and 1990s (Fig. 1). The premise was simple: the  $\beta$ -lactamase inhibitor targeted the  $\beta$ -lactamase inactivating it, so that the partner  $\beta$ -lactam could inactivate the penicillin binding protein (PBP) target, eventually resulting in bacterial cell death. When introduced, these compounds were highly effective because they mimicked the  $\beta$ -lactam core. However, resistance to these inhibitors (ie, clavulanic acid, tazobactam, and sulbactam) is highly prevalent in the clinic because of 3 mechanisms. First, from the beginning, clavulanic acid, sulbactam, and tazobactam targeted only class A serine  $\beta$ -lactamases, thus 3 structurally and functionally distinct groups of  $\beta$ -lactamases: metallo- $\beta$ -lactamases (MBLs) of class B, AmpCs serine  $\beta$ -lactamases belonging to class C, and OXAs serine  $\beta$ -lactamases of class D, were resistant to inhibition. Second, variants of previously susceptible class A  $\beta$ -lactamases (eg, TEM-1 and SHV-1) evolved single amino acid substitutions (eg, S130G, K234R) that

\*Corresponding author. Research Service, Louis Stokes Cleveland Department of Veterans Affairs Medical Center, 10701 East Boulevard, Cleveland, OH 44106. robert.bonomo@va.gov.

resulted in these inhibitors failing to inactivate these enzymes.<sup>3</sup> These variant  $\beta$ -lactamases were considered inhibitor-resistant and are classically more resistant to clavulanic acid than the sulfone inhibitors, sulbactam and tazobactam. Last, new class A  $\beta$ -lactamases, such as KPC-2, evolved (circa 1996) with the ability to hydrolyze clavulanic acid, sulbactam, and tazobactam.<sup>4</sup> Thus, these first-generation  $\beta$ -lactam- $\beta$ -lactamase inhibitor combinations are unable to inactivate most  $\beta$ -lactamases expressed by multi-drug-resistant (MDR) clinical isolates today.

## MAJOR OBSTACLES IN $\beta$ -LACTAMASE INHIBITOR DEVELOPMENT

The development of novel inhibitors is an arduous task because the mechanisms by which  $\beta$ -lactamases are resistant to clavulanic acid, tazobactam and/or sulbactam, are different even within the same class of  $\beta$ -lactamase. The mechanistic characterization of the greater than 1600  $\beta$ -lactamases identified to date is critical to understanding how to evade their action (<http://www.lahey.org/Studies/>). In addition, most MDR Gram-negatives possess more than one of these  $\beta$ -lactamases. A large gap exists that needs to be filled by identifying novel  $\beta$ -lactamase inhibitors or modified  $\beta$ -lactams that can inhibit this ever-growing population of diverse enzymes. MBLs and OXA  $\beta$ -lactamases pose the most difficult challenge. MBLs possess a  $Zn^{2+}$ -mediated noncovalent mechanism, whereas the OXA class is extremely heterogeneous with over 500 different variants at the time this article was written. In addition, the  $\beta$ -lactam hydrolytic mechanism of OXAs is fundamentally different and not like the other serine-based mechanisms. As a result, a single “magic bullet”  $\beta$ -lactam- $\beta$ -lactamase inhibitor combination that targets all clinically important  $\beta$ -lactamases (eg, KPC-2, OXA-24/40, AmpC, and NDM-1) seems unlikely. This debated question is addressed and the novel  $\beta$ -lactam- $\beta$ -lactamase inhibitor combinations are discussed in this article.

## CHANGING THE $\beta$ -LACTAM PARTNER: CEFTOLOZANE-TAZOBACTAM

One technique to combat strains that carry multiple different classes of  $\beta$ -lactamases is to switch the  $\beta$ -lactam partner of a clinically available inhibitor. Cubist (now owned by Merck) used this approach with tazobactam when they paired tazobactam with the novel cephalosporin, ceftolozane (Fig. 2 and Table 1). Ceftolozane-tazobactam was approved in December 2014 by the US Food and Drug Administration (FDA) for the treatment of complicated intra-abdominal infections (cIAIs) and complicated urinary tract infections (cUTIs). Ceftolozane-tazobactam will also be tested in clinical trials for ventilator-associated pneumonia, cystic fibrosis patients, and for diabetic lower limb infections (Table 1). What makes this combination successful? Ceftolozane is more stable against the AmpC  $\beta$ -lactamase than the predecessor  $\beta$ -lactam partners of tazobactam. AmpC possesses a low catalytic efficiency ( $k_{cat}/K_m$ ) for ceftolozane.<sup>5</sup> Thus, ceftolozane inhibits PBPs and inhibitor-resistant TEMs and SHVs as well as AmpC (unlike tazobactam), allowing tazobactam to target class A serine  $\beta$ -lactamases (eg, TEM-1) and extended-spectrum beta-lactamases (ESBLs) (eg, CTX-M-15). In addition, ceftolozane works against some class D oxacillinases (eg, OXA-1).<sup>6</sup> Thus, the ceftolozane-tazobactam combination is able to target class A, C, and some class D  $\beta$ -lactamases; the major exception is carbapenemases. Ceftolozane or ceftolozane-tazobactam was demonstrated to be similarly effective or

superior to other  $\beta$ -lactams in a variety (eg, lung, urinary tract, burn wound, sepsis, and thigh) of animal infection models using *Pseudomonas aeruginosa*, *Escherichia coli*, or *Klebsiella pneumoniae*<sup>6</sup>; therefore, its utility in the clinic may be expanded for other infections. In addition, clinical trials are recruiting pediatric patients to test ceftolozane-tazobactam (Table 1).

Laboratory-mediated selection of resistance (minimum inhibitory concentration [MIC] range of 4–8 mg/L) to ceftolozane-tazobactam was slower to occur than that with ceftazidime, meropenem, and ciprofloxacin in *P aeruginosa* PAO1; resistance was attributed to global pleiotropic changes.<sup>7</sup> Higher levels of ceftolozane-tazobactam resistance (MIC range of 32–128 mg/L) were obtained only with a *mutS* strain of PAO1; in these resistant strains, mutations were identified in *bla*<sub>ampC</sub> and the *bla*<sub>ampC</sub> regulatory pathway. Since the writing of this article, additional amino acid substitutions in the AmpC from *Pseudomonas* were found to confer resistance to ceftolozane-tazobactam.<sup>8</sup>

## DIAZABICYCLOOCTANONES, THE “FUTURE” OF $\beta$ -LACTAMASE INHIBITOR MEDICINAL CHEMISTRY

DBOs are synthetic non- $\beta$ -lactam-based  $\beta$ -lactamase inhibitors that were discovered in the early 2000s. Over a period of 15 years, this class of compounds has expanded almost exponentially with most modifications occurring at the C2 side chain (Fig. 3). Most studies published to date indicate that DBOs possess class A and class C activity with minor class D activity. More recently, the DBOs, FPI-1465 and RG6080 (formerly, OP0595), showcased at the 2013 and 2014 Interscience Conference on Antimicrobial Agents and Chemotherapy (ICAAC) meetings, and WCK 5153 and WO2013/030735, claim activity against PBPs as well. Many pharmaceutical companies (eg, Actavis-Allergan [formerly Forest-Cerexa], AstraZeneca, Fedora Pharmaceuticals, Meiji Seika Pharmaceuticals, Merck [formerly Cubist], Naeja Pharmaceuticals, Roche, and Wockhardt) are developing DBO derivatives (see Fig. 3).

## AVIBACTAM, THE “PIONEER” DIAZABICYCLOOCTANONE IN THE CLINIC

In December 2014, avibactam was the first DBO to be approved by the FDA in combination with ceftazidime for the treatment of cUTIs and cIAIs; currently the combination is being tested in the pediatric population (see Fig. 3 and Table 1). Thus, the microbiological, pharmacologic, and biochemical characteristics of avibactam are the most known and are discussed in more detail than the other DBOs. Depending on the partner  $\beta$ -lactam (eg, ceftazidime, ceftaroline, aztreonam, cefepime, or imipenem),  $\beta$ -lactam-avibactam combinations have the potential to be highly effective against many MDR gram-negative pathogens, including Enterobacteriaceae and *P aeruginosa*, producing class A, B, C, and some D  $\beta$ -lactamases.<sup>1</sup> Avibactam has been studied primarily with 2 partner cephalosporins, ceftazidime and ceftaroline; these combinations target Gram-negatives expressing class A, C, and some D  $\beta$ -lactamases (see Fig. 3 and Table 1). However, partnership with aztreonam would expand the spectrum of activity to include class B MBLs, because aztreonam is not hydrolyzed by MBLs (see Fig. 5 and Table 1). Imipenem-avibactam and cefepime-avibactam or aztreonam-avibactam and ceftaroline-avibactam are effective against *E coli* and

*K pneumoniae* carrying *bla*<sub>OXA-48</sub>, respectively, but not *Acinetobacter baumannii* producing *bla*<sub>OXA5</sub>.<sup>9-12</sup> Avibactam restores susceptibility to ceftazidime when tested against clinical isolates of Enterobacteriaceae possessing porin and outer membrane permeability defects.<sup>13</sup>

Avibactam forms a stable carbamyl-adduct with serine  $\beta$ -lactamases that is reversible through recyclization of the 5-membered urea ring.<sup>14</sup> Thus far, decarbamylation-hydrolysis of avibactam was only observed after 24 hours with class A  $\beta$ -lactamase, KPC-2.<sup>15</sup> Formation of the carbamyl-enzyme complex, represented by the kinetic value,  $k_2/K$  and recyclization-decarbamylation to reform active avibactam, denoted by a  $k_{\text{off}}$  value differs for the various classes of serine  $\beta$ -lactamases. These kinetic parameters translate directly to efficacy within bacterial cells; thus, an ideal DBO is one that possesses a high  $k_2/K$  value and low  $k_{\text{off}}$  value. The tested class A and C  $\beta$ -lactamases (with the exception of the class A  $\beta$ -lactamase BlaC from *Mycobacterium tuberculosis*) acylate rapidly with high  $k_2/K$  values (range:  $10^4$  to  $10^6$   $\text{M}^{-1}\text{s}^{-1}$ ) and recyclize slowly with low  $k_{\text{off}}$  values (range:  $10^{-3}$  to  $10^{-4}$   $\text{s}^{-1}$ ).<sup>15,16</sup> Conversely, BlaC and class D  $\beta$ -lactamases are slow to acylate with low  $k_2/K$  values in the range of  $10^1$  to  $10^3$   $\text{M}^{-1}\text{s}^{-1}$ , but once acylated, recyclization is very slow with  $k_{\text{off}}$  values of  $10^{-5}$  to  $10^{-6}$   $\text{s}^{-1}$ . X-ray crystallography as well as molecular modeling revealed that avibactam adopts very similar active site conformations in class A, C, and D  $\beta$ -lactamases.<sup>16-20</sup>

Recent crystal structures and biochemical analysis with avibactam and OXA-24/40 and OXA-48 were conducted and provided some insights into why select class D  $\beta$ -lactamases are inhibited, whereas others are not.<sup>17,19</sup> OXA-24/40's  $k_2/K$  value is much lower than OXA-48's, whereas the  $k_{\text{off}}$  values are very similar. The crystal structures revealed that the binding pocket for avibactam in class D  $\beta$ -lactamases is more hydrophobic with fewer polar residues present, thus potentially affecting binding and acylation. In addition, OXA-24/40 possesses additional hydrophobic moieties (eg, hydrophobic bridge between Y112 and M223), whereas OXA-48 possesses polar residues (eg, T213, R214, and D101) that could aid in Michaelis-complex formation with avibactam. Slow recyclization was suggested to occur because of a decarboxylated Lys84/73.<sup>17,19</sup>

## RESISTANCE TO $\beta$ -LACTAM-AVIBACTAM COMBINATIONS

Selection via passaging of *E coli* producing *bla*<sub>CTX-M-15</sub> and *E cloacae* with de-repressed *bla*<sub>AmpC</sub> on ceftaroline-avibactam identified strains with elevated ceftaroline-avibactam MICs; however, most of the mutations were unstable.<sup>21</sup> *E coli* producing CTX-M-15 Lys237Gln variant conferred resistance with the cost of ESBL activity. Selected-resistant *E cloacae* possessed  $\Omega$  loop deletions within AmpC as well as porin loss. A similar approach was conducted using *P aeruginosa* with de-repressed *bla*<sub>AmpC</sub> with ceftazidime-avibactam.<sup>22</sup> Ceftazidime-avibactam-resistant variants possessed MICs from 64 to 256 mg/L as a result of deletions in the  $\Omega$  loop of the AmpC. Mechanistic analyses revealed that the AmpC variants were less susceptible to avibactam inactivation and possessed improved ceftazidime kinetics.

Resistance to ceftazidime-avibactam was observed in a panel of clinical isolates of *P aeruginosa*.<sup>23</sup> The mechanism of resistance was dissected by sequencing *bla*<sub>AmpC</sub>, the genes

in the *bla*<sub>AmpC</sub> regulon, *pbps*, and *oprD*, measuring efflux pump expression, and combination antibiotic therapy with ceftazidime-avibactam. Using these methods, membrane permeability and drug efflux were found to be the most important factor influencing ceftazidime-avibactam resistance in these isolates. Resistance to ceftazidime-avibactam was overcome with ceftazidime-avibactam/fosfomycin; this combination targets PBPs,  $\beta$ -lactamases, and MurA, and UDP-*N*-acetylglucosamine-3-enolpyruvyltransferase, which is involved in peptidoglycan synthesis.

In a panel of clinical isolates of *E coli* producing *bla*<sub>NDM-1</sub>, resistance to aztreonam-avibactam was detected.<sup>24</sup> Aztreonam was still able to inhibit the  $\beta$ -lactamases; however, a 4-amino-acid insertion was identified in PBP3.

Variant  $\beta$ -lactamases with single amino acid substitutions in residues that typically result in inhibitor resistance (ie, 69, 130, 234, 220/244, and 276) and ceftazidime-resistance (ie, 164, 167, 169, and 179) in clinical isolates were tested in SHV-1 and KPC-2  $\beta$ -lactamase isogenic *E coli* strain backgrounds with  $\beta$ -lactam-avibactam combinations.<sup>25–27</sup> The S130G, K234R, and R220M (KPC-2)/R244S (SHV-1) substitutions in the SHV-1 and KPC-2 backgrounds resulted in elevated MICs to ampicillin-avibactam when expressed in *E coli*.<sup>25,27</sup> The S130G variants of SHV-1 and KPC-2 were found to have severely compromised  $k_2/K$  values ( $\sim 1 \text{ M}^{-1}\text{s}^{-1}$ ), thus avibactam failed to inactivate these variants. S130 is an important residue for avibactam acylation. The resistance mechanisms of the K234R and R220M/R244S variants remain to be defined. The R164A, R164P, D179A, D179Q, and D179N substitutions in KPC-2 resulted in increased ceftazidime-avibactam MICs.<sup>26</sup> Loss of susceptibility to ceftazidime-avibactam is thought to be due to enhanced ceftazidime kinetics of the variants because avibactam was still able to inactivate the R164A and D179N variants. In another study, selection of Enterobacteriaceae producing blaKPC for resistance to ceftazidime-avibactam resulted in the identification of KPC variants with D179Y amino acid substitutions.<sup>28</sup> Resistance to a  $\beta$ -lactam- $\beta$ -lactamase inhibitor combination due to resistance to the partner  $\beta$ -lactam is a very intriguing observation. The choice of a  $\beta$ -lactam partner is critical, as described above with ceftolozane-tazobactam.

During the writing of this manuscript, the first clinical observation of ceftazidime-avibactam resistance was reported.<sup>29</sup> The resistance was observed in *Klebsiella pneumoniae* expressing blaKPC-3 and mechanism of resistance is unclear.

## DIAZABICYCLOOCTANONES, RELEBACTAM AND OP0595, ON THE HORIZON

Relebactam partnered with imipenem-cilastatin demonstrates a similar spectrum of activity as avibactam, thus lacking activity against MBLs and most OXAs (see Fig. 3 and Table 1).<sup>30–32</sup> RG6080 (formerly OP0595) not only is an inhibitor of class A and C  $\beta$ -lactamases but also inhibits PBP-2 of Enterobacteriaceae (see Fig. 3 and Table 1).<sup>33,34</sup> Thus, RG6080 is unique compared with avibactam and relebactam, because it does not need a  $\beta$ -lactam partner for antimicrobial activity. In addition, there is evidence that RG6080 acts to enhance the activity of  $\beta$ -lactams.<sup>35</sup>

## DIAZABICYCLOOCTANONES IN PRECLINICAL DEVELOPMENT

FPI-1465 when combined with aztreonam and ceftazidime possesses activity against Enterobacteriaceae containing ESBLs and class A, B, and D carbapenemases (see Fig. 3; Table 2).<sup>36,37</sup> In addition, FPI-1465 is also active against PBPs (ie, PBP2) from *E coli* and *P aeruginosa*.<sup>38</sup>

Wockhardt, Ltd has 3 DBOs in the pipeline: WCK 4234, WO2013/030735, and WCK 5153 (see Fig. 3 and Table 2).<sup>39</sup> The WCK 4234 combined with meropenem demonstrates activity against oxacillinase-producing strains of *A baumannii*.<sup>39</sup> WO2013/030735 and WCK 5153 possess antibacterial activity against *P aeruginosa* and *E coli*.

During the publication process of this article, Merck (formerly Cubist) published work with another DBO, CB-618.<sup>40</sup> CB-618 tested in combination with meropenem displays activity against clinical isolates of Enterobacteriaceae expressing the KPC-2, KPC-3, FOX-5, OXA-48, SHV-11, SHV-27, and/or TEM-1 beta-lactamases.

## BORONIC ACID $\beta$ -LACTAMASE INHIBITORS ARE MAKING GREAT STRIDES

In the late 1970s, boronic acids were recognized as inhibitors of serine  $\beta$ -lactamases in vitro.<sup>41</sup> Boron forms a reversible bond with the  $\beta$ -lactamase.<sup>42</sup> Boronic acids serve as competitive inhibitors and were not shown to be hydrolyzed by any  $\beta$ -lactamase to date. Historically, despite good affinities for many class A and C serine  $\beta$ -lactamases, boronic acids failed to make it into clinical development, but the tides are changing.

A novel cyclic boronic acid-based  $\beta$ -lactamase inhibitor, RPX7009, is in phase 3 clinical trials in combination with meropenem (RPX2014) under the name Carbavance; in addition, clinical trials in pediatric patients are in the works (Fig. 4; see Table 1).<sup>43</sup> Initially, development began with biapenem (RPX2003) (see Fig. 4 and Table 1). Biapenem-RPX7009 and meropenem-RPX7009 are most effective against Enterobacteriaceae producing class A carbapenemases and demonstrated against class A ESBLs and AmpC; impermeability had a negative impact on the activity of carbapenem-RPX7009 combinations.<sup>44,45</sup> In addition, biapenem-RPX7009 was not effective against Enterobacteriaceae expressing MBLs or OXA-48.<sup>44</sup>

RPX7009 did not potentiate the activity of carbapenems against nonfermenters *P aeruginosa* and *A baumannii*.<sup>45</sup> RPX7009 also did not increase the activity of biapenem against anaerobes.<sup>46</sup> A neutropenic lung model of infection in mice with *K pneumoniae* producing *bla*<sub>KPC-2</sub> revealed that RPX-7009 reduced colony forming units (CFUs) by 2 logs in combination with biapenem and meropenem compared with the carbapenem alone.<sup>43</sup>

## BORONIC ACIDS IN PRECLINICAL DEVELOPMENT

In addition, several boronic acid inhibitors in preclinical development showed promise because they reduced MICs or were successful in animal models of infection. Benzo(b)thiophene-2-boronic acids combined with ceftazidime are effective against Enterobacteriaceae and *P aeruginosa* (see Fig. 4 and Table 2).<sup>47</sup> S02030, a novel boronic

acid possessing thiophene and triazole carboxylate side chains demonstrates activity against Enterobacteriaceae carrying blaKPCs with a  $k_2/K$  value ( $1.2 \pm 0.2 \times 10(4) M(-1) s(-1)$ ) comparable to avibactam (Fig. 4 and Table 2).<sup>48</sup> TheraBor Pharmaceuticals and the Regents of the University of California developed and patented (patent WO2013/056079) several sulfonamide boronates (eg, CR161) that were shown to reduce ceftazidime MICs against Enterobacteriaceae and *P aeruginosa* (see Fig. 4 and Table 2).<sup>16,49</sup> Moreover, when mice were infected intraperitoneally with *E coli* overexpressing AmpC, after 120 hours, the mice treated with CR161 combined with cefotaxime possessed a 65% survival compared with cefotaxime alone at 15%. VenatoRx Pharmaceuticals also patented a series of cyclic boronic acids, 3,4-dihydro-2H-benzo[e][1,2]oxaborinine-8-carboxylic acids (US 8,912,169 B2), and a set of novel  $\alpha$ -aminoboronic acids (US 20100120715 A1) (see Fig. 4 and Table 2). Select cyclic boronates combined with either ceftazidime or meropenem demonstrated activity against *E cloacae*, *K pneumoniae* with bla<sub>KPC-3</sub>, *P aeruginosa* with bla<sub>VIM-2</sub>, and *K pneumoniae* with bla<sub>KPC-2</sub> and bla<sub>VIM-4</sub> and inhibited SHV-5, KPC-2, VIM-2, AmpC, and OXA-1 with concentration of inhibitor at which 50% inhibition of substrate hydrolysis is observed (IC<sub>50</sub>) values <1  $\mu$ M. Select  $\alpha$ -aminoboronic acids combined with ceftazidime possessed activity against *E coli* with bla<sub>SHV-5</sub>, *K pneumoniae* with bla<sub>CTX-M-15</sub>, *E cloacae* with bla<sub>P99</sub>, and *K pneumoniae* with bla<sub>KPC-2</sub> and inhibited SHV-5, CTX-M-15, P99, and KPC-2, with IC<sub>50</sub> values <0.1  $\mu$ M. Rempex, a subsidiary of The Medicines Company, also patented another group of cyclic boronic acids, 3,4-dihydro-2H-benzo[e][1,2]oxaborinine-8-carboxylic acids (see Fig. 4 and Table 2) (WO2014/107536 A1). Select compounds in this class have  $K_i$  values of less than 1  $\mu$ M against SHV-12, TEM-10, CTX-M-14, KPC-2, P99, CMY-2, OXA-48, VIM-1, and NDM-1, and when combined with carbapenems, demonstrated MICs less than 1 mg/L for Enterobacteriaceae producing bla<sub>NDM-1</sub>, bla<sub>VIM-1</sub>, and bla<sub>KPC-2</sub> and even *A baumannii* expressing bla<sub>NDM-1</sub>.

## NOVEL SULFONES AND CLAVAMS IN PRECLINICAL DEVELOPMENT

Allegra Therapeutics is developing a novel sulfone, AAI101, in combination with cefepime; cefepime-AAI101 demonstrated activity against some Enterobacteriaceae with ESBLs or carbapenemases (Fig. 5; see Table 2).<sup>50</sup> In a neutropenic thigh mouse infection model, the cefepime-AAI101 combination reduced bacterial CFUs by more than 0.5 log CFU for 12 of the 20 strains tested; cefepime alone only worked in 3 of 20 strains. Orchid Pharmaceuticals synthesized a series of sulfone derivatives with no R1 side chain and different R2 side chains, and some lowered meropenem and imipenem MICs from 32 to 64 mg/L to 1 to 2 mg/L against *K pneumoniae* with bla<sub>KPC-2</sub> (WO/2012/070071) when tested in combination (see Table 2). These sulfones were also tested in combination with imipenem or meropenem against other Enterobacteriaceae with bla<sub>KPC-2</sub> or bla<sub>KPC-3</sub>, and MICs decreased from 2 to 4 mg/L to 0.25 to 0.5 mg/L for some compounds. Another set of novel sulfones was developed by Dr John D. Buynak, and one of these compounds lowered meropenem and imipenem MICs for carbapenem-susceptible *A baumannii* producing bla<sub>OXA-24/40</sub> from 32 mg/L to 1 mg/L when tested in combination (see Table 2).<sup>51</sup> Nabriva Therapeutics created clavam spinoffs that possessed activity against *K pneumoniae* and *Citrobacter freundii* with MICs decreasing for ceftazidime from 26 mg/L to 0.2 mg/L and 3.2 mg/L, respectively, with a clavam derivative (see Table 2).

## PHOSPHONATES IN PRECLINICAL DEVELOPMENT

Like with the boronates, literature shows that phosphonates are good inhibitors of class A and C and even some B and D  $\beta$ -lactamases kinetically.<sup>52–54</sup> Mirati Therapeutics conducted preclinical studies on a phosphonate, MG96077, which is a novel broad-spectrum, non- $\beta$ -lactam  $\beta$ -lactamase inhibitor (Fig. 6; see Table 2).<sup>39,55</sup> Imipenem combined with MG96077 decreased greater than 90% of the MICs for imipenem-resistant *P aeruginosa* and *K pneumoniae* to 4 mg/L. In a mouse spleen infection model with imipenem-resistant *P aeruginosa*, imipenem-MG96077 caused a 4 to 6 log reduction in CFUs and increased mouse survival.

## MONOBACTAMS ARE PROMISING $\beta$ -LACTAMASE INHIBITORS OR EVADE $\beta$ -LACTAMASE ACTIVITY

Already available in the clinic, the monobactam, aztreonam, is a  $\beta$ -lactam that can also circumvent certain  $\beta$ -lactamases (Fig. 7). Aztreonam inhibits certain AmpC  $\beta$ -lactamases and binds poorly to MBLs.<sup>56–58</sup>

## MONOBACTAMS AND DERIVATIVES IN PRECLINICAL DEVELOPMENT

The monobactam scaffold is encouraging for future MBL inhibitors as well as inhibitors of AmpCs, and Basilea Pharmaceuticals, Merck, Pfizer, and Taiho Pharmaceutical Co are working on these agents.

For Basilea, expansion of the monobactam class has resulted in several new monobactams (eg, BAL30072 and BAL19764) and a novel class of bridged monobactams (eg, BAL29880) that serve as  $\beta$ -lactamase inhibitors. BAL30072 contains a 1,5-dihydroxy-4-pyridone group that is a siderophore moiety that allows for transport of the compound into the bacterial cell via the TonB iron transport system (see Fig. 7 and Table 2). BAL30072 possessed bactericidal activity against *Acinetobacter* spp, which include those with some *bla*<sub>OXA5</sub>, *P aeruginosa*, *Burkholderia* spp, Enterobacteriaceae with class A carbapenemases, and against strains that produced *bla*<sub>MBL5</sub>.<sup>59–62</sup> BAL30072 demonstrated efficacy in rat soft tissue infection by *A baumannii* as well as a mouse septicemia model when combined with meropenem against *Serratia marcescens* producing *bla*<sub>SME-1</sub> (a class A carbapenemase), *P aeruginosa*, and *A baumannii*.<sup>61,63</sup>

BAL30376 combines BAL19764, another siderophore monobactam, with the bridged monobactam BAL29880 and clavulanic acid (see Fig. 7 and Table 2).<sup>64,65</sup> BAL30376 was effective against most clinical Enterobacteriaceae isolates producing *bla*<sub>AmpC5</sub> and *bla*<sub>ESBL5</sub>, *P aeruginosa*, *A baumannii* producing *bla*<sub>OXA-23</sub>, or the southeast region clonal lineage and some *Stenotrophomonas maltophilia*. BAL30376 was not effective against Enterobacteriaceae expressing *bla*<sub>MBL5</sub>, *bla*<sub>KPC</sub>, or *bla*<sub>OXA-48</sub>, or those with impermeability, *P aeruginosa* or *B cepacia* complex isolated from cystic fibrosis patients, *P aeruginosa* expressing *bla*<sub>MBL5</sub>, and other *A baumannii*. In addition, in a mouse septicemia model of infection, BAL30376 demonstrated efficacy against selected isolates of *A baumannii*, *E cloacae*, and *P aeruginosa*.<sup>65</sup>



Merck is conducting preclinical testing with a bridged monobactam, MK-8712 (see Fig. 7 and Table 2).<sup>66</sup> MK-8712 combined with imipenem reduced MICs against *P aeruginosa* strain CL5701 from 32 mg/L to 4 mg/L. MK-8712 was also a potent inhibitor of AmpC of *P aeruginosa* with an IC<sub>50</sub> value of 1 μM. MK-8712 combined with imipenem-cilastatin resulted in a 4.6 log fold reduction in CFU in a mouse spleen model of infection compared with imipenem-cilastatin alone.<sup>67</sup>

Pfizer developed a series of siderophore monobactams, similar to BAL30072; however, the siderophore moiety is on the R2 side chain (see Fig. 7 and Table 2).<sup>68</sup> Several of these compounds worked in combination with aztreonam or meropenem to lower MIC values to the susceptible range against *P aeruginosa* and also possessed efficacy in a mouse pneumonia model of infection. The siderophore monobactams were modified to increase the length of the R2 side chain with linkers resulting in expansion of activity against *K pneumoniae*, *E coli*, and *A baumannii*, including *P aeruginosa* carrying a *bla*<sub>MBL</sub>; these compounds target the PBPs and evade MBL activity.<sup>69</sup>

Similar to Merck, Taiho Pharmaceutical Co is working on a siderophore monobactam, Syn2190, that inhibited AmpC β-lactamases and reduced ceftazidime MICs to the susceptible range against Enterobacteriaceae and *P aeruginosa* expressing *bla*<sub>AmpC</sub> (see Fig. 7 and Table 2).<sup>70</sup> In addition, Syn2190 combined with ceftazidime or ceftiprome demonstrated activity in mouse systemic and urinary tract infection models using *P aeruginosa*. However, Syn2190 induced expression of *bla*<sub>AmpC</sub>.

## A NEW SIDEROPHORE CEPHALOSPORIN IN CLINICAL DEVELOPMENT

S-649266, a novel catechol-substituted siderophore cephalosporin, is in phase II clinical trials (see Table 1 and Fig. 8).<sup>71</sup> This compound demonstrated potent in vitro activity against a diverse panel of gram-negative bacteria<sup>39</sup> S-649266 was not hydrolyzed by KPC-2, P99, or OXA-23; minimal hydrolysis with  $k_{cat}/K_m$  values in the 10<sup>3</sup>–10<sup>4</sup> M<sup>-1</sup>s<sup>-1</sup> for MBLs, IMP-1, VIM-2, L1, and CTX-M-15 was observed.<sup>72</sup> S-649266 demonstrated activity against *P aeruginosa*, *S maltophilia*, *K pneumoniae*, and *A baumannii* with MIC<sub>90</sub> values <2 mg/L.<sup>73</sup> Furthermore, against MDR strains of Enterobacteriaceae, *P aeruginosa*, and *A baumannii*, the MIC<sub>90</sub> values were less than 4 mg/L.<sup>73</sup> In a rat lung infection model, S-649266 was shown to have efficacy against *P aeruginosa* and *A baumannii*, including MDR strains.<sup>74</sup> In addition, several different mouse models of infections (ie, systemic, lung, urinary tract, and subcutaneous infection) using various Gram-negatives were used to assess the efficacy of S-649266; the compound was effective against *K pneumoniae* producing *bla*<sub>KPC-2</sub>.<sup>75</sup>

## NOVEL 3'-THIOBENZOYL CEPHALOSPORINS IN PRECLINICAL DEVELOPMENT

3'-Thiobenzoyl cephalosporins demonstrated inhibitory activity against class A, B, C, and D β-lactamases with IC<sub>50</sub> values in the range of 1.4 μM to 140 μM (Fig. 8 and see Table 2) (US 20120329770 A1). In addition, when combined with meropenem, selected compounds possessed activity against *P aeruginosa*, *S maltophilia*, and *Chryseobacterium meningosepticum*.

## NOVEL CARBAPENEMS IN PRECLINICAL DEVELOPMENT: FSI-1671 AND FSI-1686

Two novel carbapenems, FSI-1671 and FSI-1686, were shown to possess activity against MDR *A baumannii*, Enterobacteriaceae, and some *P aeruginosa* (Fig. 9; see Table 2).<sup>76</sup> Mice were intraperitoneally infected by carbapenem-resistant *A baumannii*, *K pneumoniae*, and *P aeruginosa* and treated with FSI-1671, FSI-1686, meropenem, doripenem, colistin, or tigecycline.<sup>77</sup> Both novel carbapenems demonstrated good potency (lower ED<sub>50</sub> [50% effective dose] values) against carbapenem-resistant *A baumannii*, *K pneumoniae*, and *P aeruginosa* compared with the other carbapenems tested. The  $\beta$ -lactamase inhibitory properties of these compounds were not assessed, but all other clinically available carbapenems are dual agents inhibiting PBPs and some  $\beta$ -lactamases.

## METALLO- $\beta$ -LACTAMASE-SPECIFIC INHIBITORS IN PRECLINICAL DEVELOPMENT: BISTHIAZOLIDINES AND ME1071

Four bisthiazolidine (BTZ) inhibitors were tested against VIM-2 and VIM-24 and possessed  $K_i$  values between 3.7 and 14  $\mu$ M (Fig. 10; see Table 2).<sup>78</sup> Most importantly, the BTZs restored imipenem susceptibility of clinical isolates, *P aeruginosa* producing *bla*<sub>VIM-2</sub> and *K pneumoniae* carrying *bla*<sub>VIM-24</sub>. In addition, BTZs demonstrate activity against NDM-1 with  $K_i$  values from 7 to 19  $\mu$ M and are effective against *A. baumannii*, *K. pneumoniae*, and *P. rettgeri* expressing *bla*<sub>NDM-1</sub> when combined with imipenem (ACS Infect. Dis., 2015, 1 (11), pp 544–54).

ME1071 is a maleic acid derivative, and when combined with ceftazidime, increased susceptibility to ceftazidime against *P aeruginosa* expressing *bla*<sub>IMP</sub> and *bla*<sub>VIM</sub>, although never for 100% of the isolates tested (see Fig. 10 and Table 2).<sup>79</sup> Biapenem-ME1071 combination decreased MICs for Enterobacteriaceae with *bla*<sub>IMP</sub> and *bla*<sub>VIM</sub>, but not *bla*<sub>NDM</sub>.<sup>80</sup> ME1071 in combination with biapenem was effective in a mouse model for ventilator-associated pneumonia by *P aeruginosa* producing *bla*<sub>MBL</sub>.<sup>81</sup>

## SUMMARY

When one thinks about designing drugs to treat infections caused by MDR bacteria, there are 2 approaches to keep in mind. One strategy involves designing “niche” drugs to target specific bacteria with certain resistance mechanisms. These “niche” agents would be highly useful especially for uncommon or hard-to-treat infections (eg, *S maltophilia* or MDR *A baumannii* or Gram-negatives producing *bla*<sub>MBL</sub>). However, using drugs that only target a specific pathogen will force clinicians to rely heavily on accurate and rapid molecular diagnostics. The second tactic is designing drugs that possess a very broad spectrum and can be given as empirical therapy. With this second scenario, “time is no longer the enemy,” and molecular diagnostics are not nearly as important. However, the risk of the bacteria evolving resistance to the novel agents is higher the more these agents are used in the clinic. If other older agents would work equally well, why accelerate the decline in utility of a novel agent?

Taking these ideas into consideration and looking back at this article, how do these approaches fit with the novel  $\beta$ -lactam- $\beta$ -lactamase inhibitor combinations described? Clavulanic acid and the sulfones seem to be classified between a “niche” and a broad-spectrum agent category because their spectrum is somewhat limited given the MDR organisms observed today. Moreover, because they were first introduced more than 30 years ago, further derivatization of these compounds has not resulted in any novel inhibitors reaching clinical development. There were slight gains such as increased penetration and the ability to target other classes of  $\beta$ -lactamase. However, based on this history, the novel inhibitors may be a better route.

The DBOs in the preclinical development target PBPs. Is this PBP activity enough to propel this class forward as a broad-spectrum agent? With that in mind, could a novel DBO be a potential “magic bullet” to target all Gram-negatives producing  $\beta$ -lactamases? Will  $\beta$ -lactamases ever evolve to hydrolyze DBOs? Probably, as KPC-2 hydrolyzes avibactam albeit at a very slow rate.

Boronates have historically possessed activity against class A and C  $\beta$ -lactamases. Now, boronates were expanded to inhibit class A and B carbapenemases and some class D  $\beta$ -lactamases with 3,4-dihydro-2H-benzo[e][1,2]oxaborinine-8-carboxylic acids showing the most promise. Can they be expanded to target OXA carbapenemases? Could a novel carbapenem-3,4-dihydro-2H-benzo[e][1,2]oxaborinine-8-carboxylic acid combination be a potential “magic bullet” to inhibit all Gram-negatives producing  $\beta$ -lactamases? Will  $\beta$ -lactamases evolve to hydrolyze them? Hydrolysis seems less likely because, to date, hydrolysis of boronic acids has not been documented. However,  $\beta$ -lactamases could still evolve to resist inhibition by boronates.

The other compounds discussed here possess more limited spectra compared with the DBOs and boronates that are in preclinical development. Some inhibitors would be categorized as “niche” agents, such as the BTZs and ME1071. Where does this leave us now, and what is the best path forward? Clinical agents need to be developed that treat the resistant pathogens that exist now. It should be kept in mind that other resistance mechanisms (eg, loss of porins, expression of efflux pumps, and other permeability barriers) are still going to be a challenge. Also, the evolution of resistance in both PBPs and  $\beta$ -lactamases to these novel  $\beta$ -lactams and  $\beta$ -lactamase inhibitor should be studied to understand their mechanisms of action in order to come up with strategies to circumvent resistance, when it eventually appears. The war against resistant pathogens will most likely not have a definitive end. However, luckily, several new agents are in the arsenal that momentarily will keep pace with the challenging pathogens in the clinic today.

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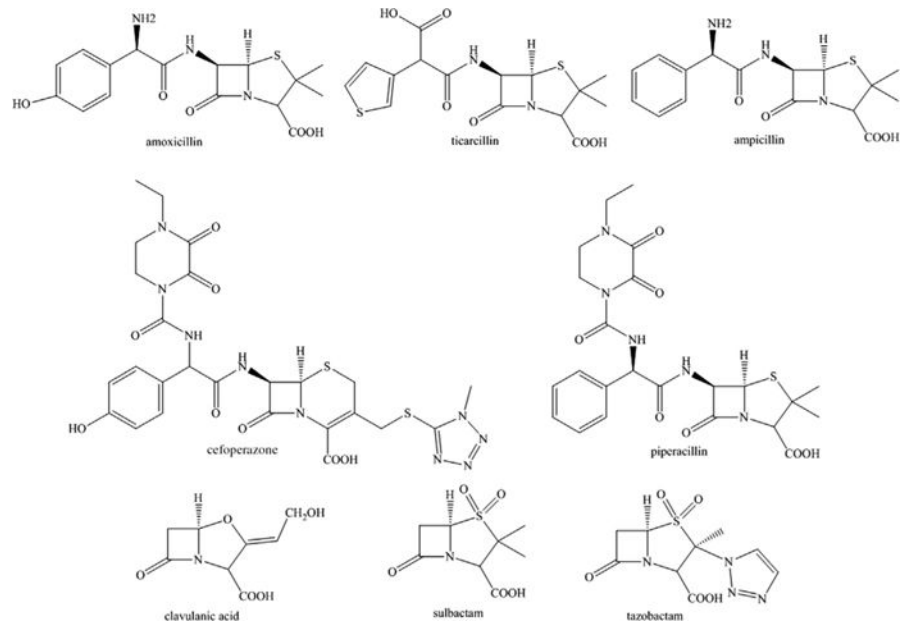
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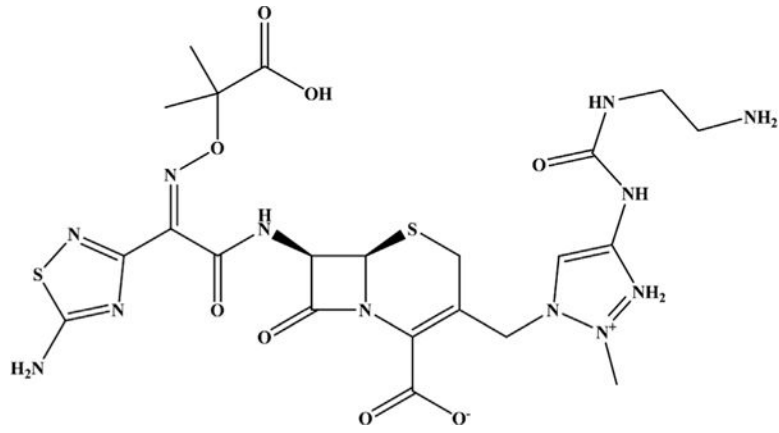


**KEY POINTS**

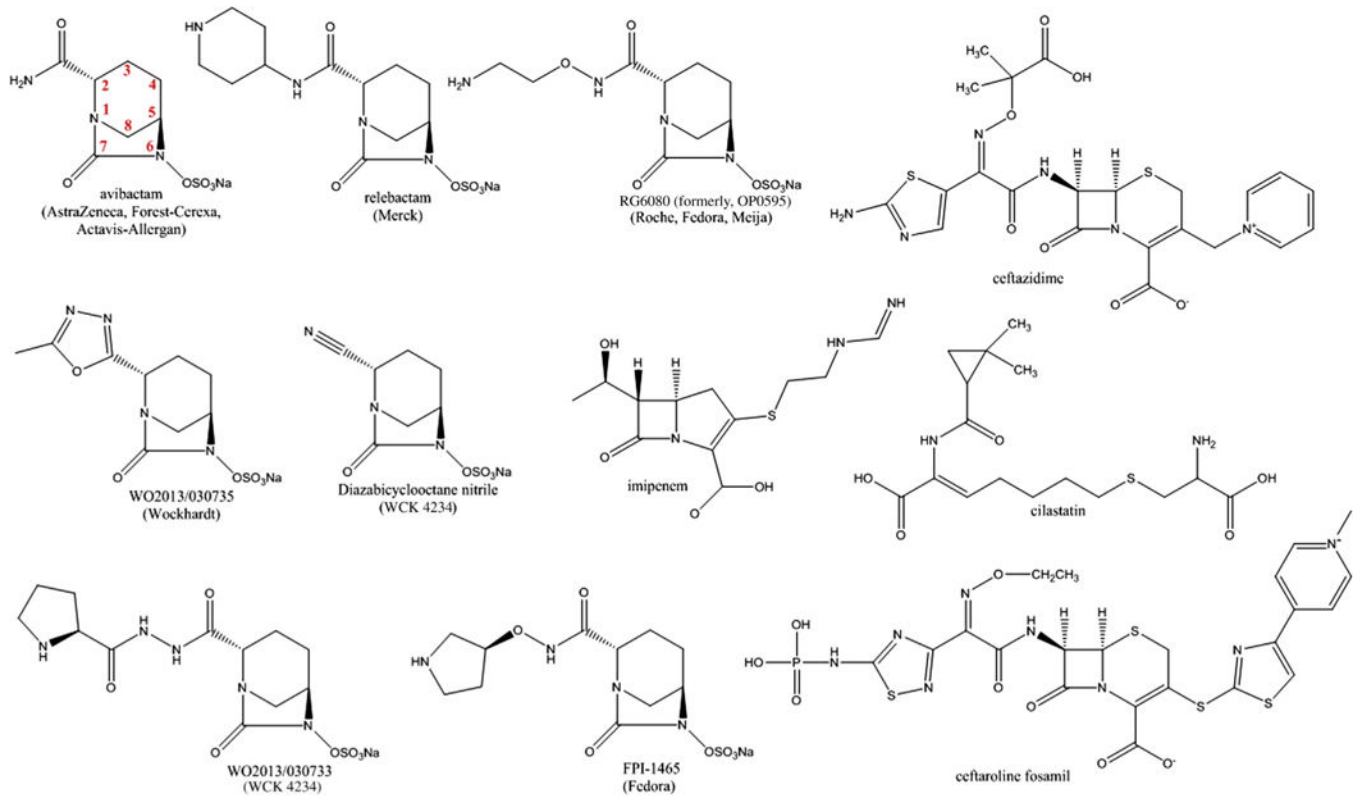
- Obstacles in the development of beta-lactamase inhibitors.
- Diazabicyclooctanones, an ever expanding novel beta-lactamase inhibitor class.
- Boronic acids, non-hydrolyzable beta-lactamase inhibitors.
- Beta-lactams as beta-lactamase inhibitors.



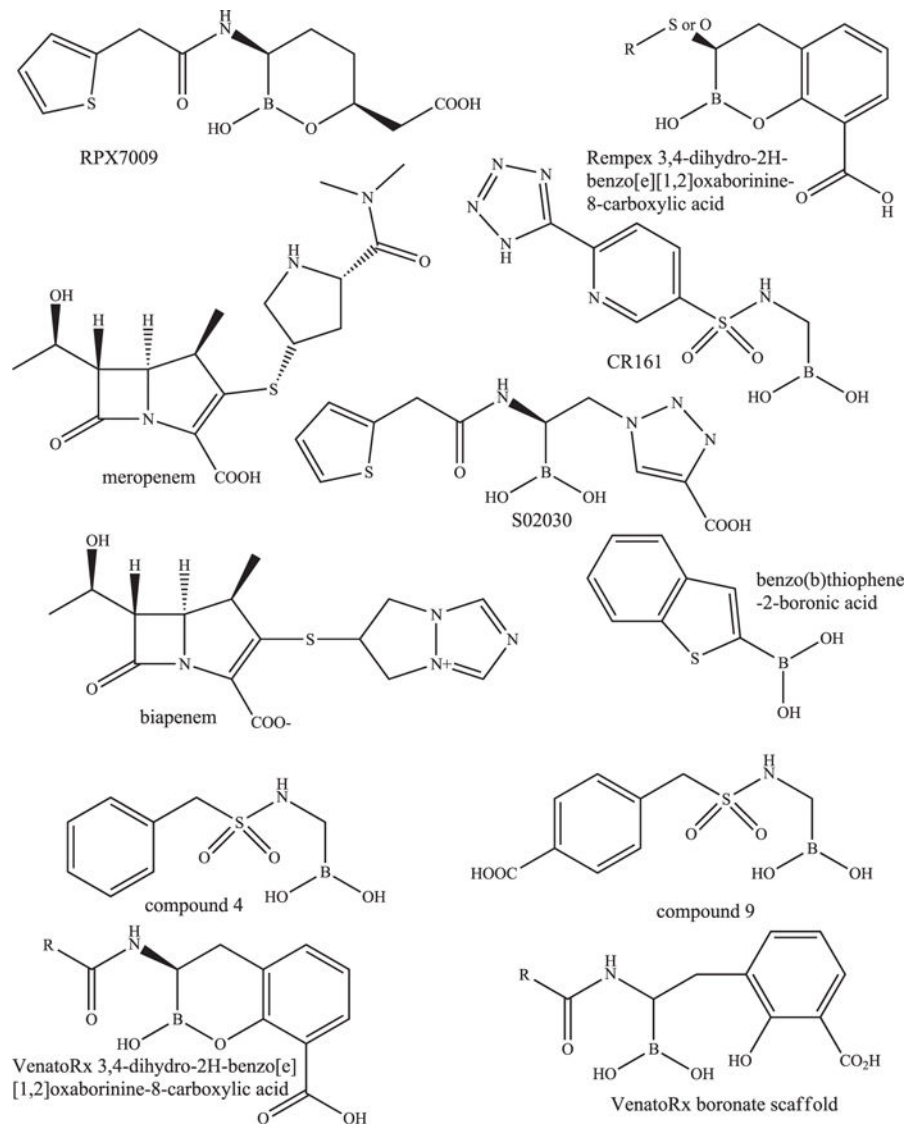
**Fig. 1.**  
 $\beta$ -Lactamase inhibitors of the past and their  $\beta$ -lactam partners.



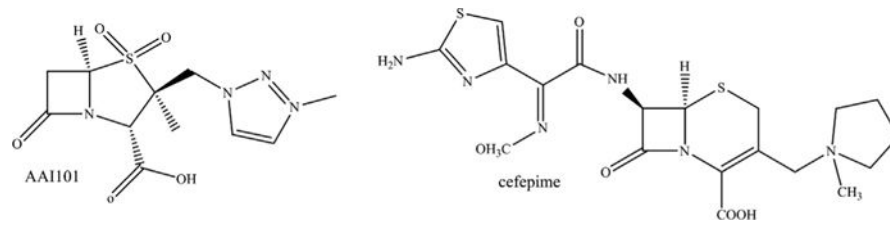
**Fig. 2.**  
Chemical structure of ceftolozane.



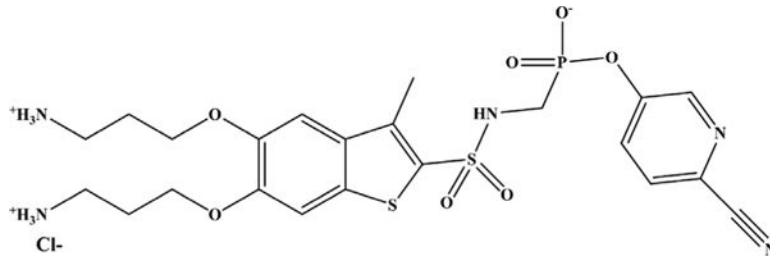
**Fig. 3.**  
DBOs and DBO  $\beta$ -lactam partners.



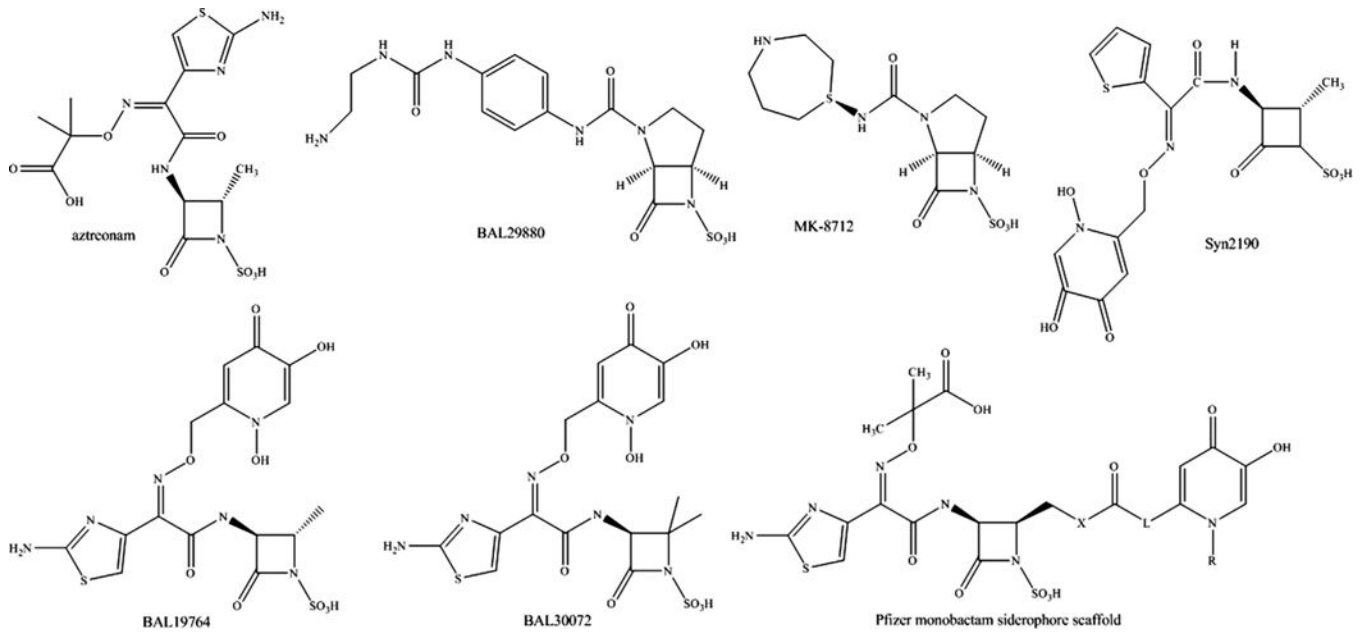
**Fig. 4.** Chemical structures of the carbavance (meropenem-RPX7009) combination, biapenem, and other boronates.



**Fig. 5.**  
Chemical structures of AAI101 and cefepime.

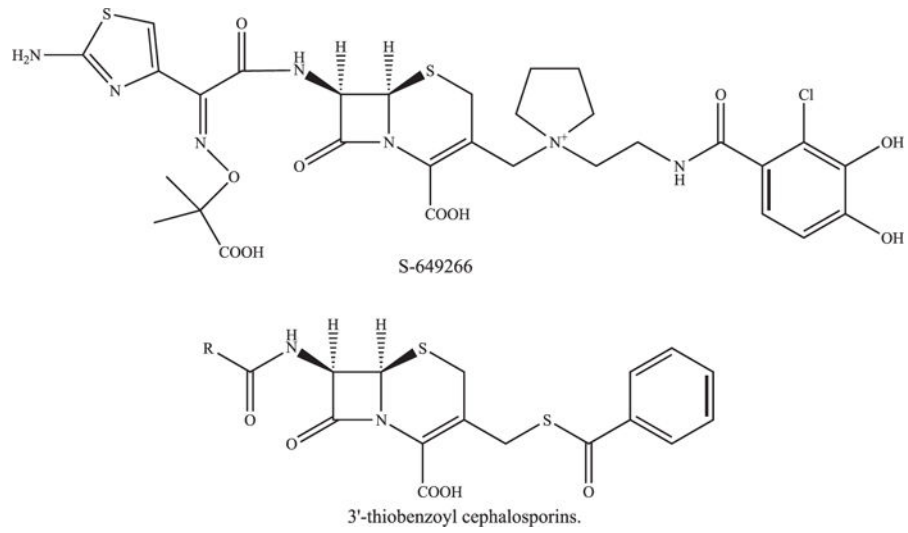


**Fig. 6.**  
Phosphonate, MG96077.

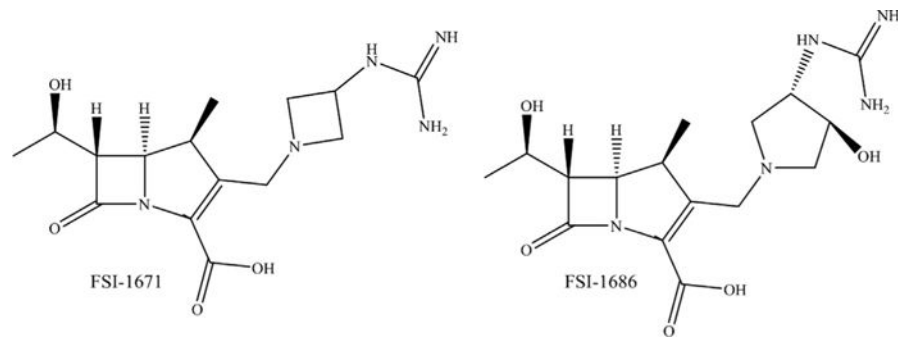


**Fig. 7.**  
Chemical structures of monobactams and bridged monobactams.

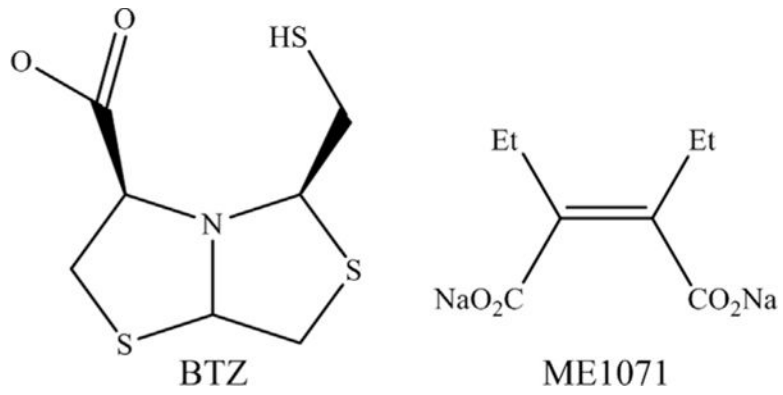




**Fig. 8.**  
Chemical structure of 3'-thiobenzoyl cephalosporins.



**Fig. 9.**  
Chemical structures of FSI-1671 and FSI-1686.



**Fig. 10.** Chemical structure of MBL-specific inhibitors the BTZ, (3R,5R,7aS)-5-(sulfanylmethyl) tetrahydro[1,3] thiazol[4,3-b][1,3]thiazole-3-carboxylic acid and ME1071.

**Table 1**New  $\beta$ -lactam- $\beta$ -lactamase inhibitor combinations in the clinic or in development

Combination	Company	Type of $\beta$ -Lactamase Inhibitor	Development Phase	US Clinical Trial Numbers (Status)
Ceftolozane-tazobactam	Merck/Cubist Pharmaceuticals	Sulfone	FDA approved (2014)	NCT01147640 (completed); NCT01853982 (terminated); NCT02266706, NCT02070757, and NCT02387372 (recruiting); NCT02508753 (completed); NCT02421120 (recruiting); NCT02620774 (not open yet)
Ceftazidime-avibactam	AstraZeneca Pharmaceuticals, Forest-Cerexa, Actavis-Allergan	DBO	FDA approved (2014)	NCT01395420, NCT01430910, NCT01290900, NCT01644643, NCT01291602, NCT00752219, NCT00690378, NCT01599806, NCT01595438, NCT01893346, NCT01499290, NCT01500239, NCT01920399, NCT01534247, and NCT01789528 (completed); NCT01726023 (completed); and NCT01808092 (completed); NCT02475733 and NCT02497781 (recruiting)
Ceftaroline-avibactam	AstraZeneca Pharmaceuticals, Forest-Cerexa, Actavis-Allergan	DBO	Phase 2	NCT01624246, NCT01281462, NCT01290900, and NCT01789528 (completed)
Aztreonam-avibactam	AstraZeneca Pharmaceuticals, Forest-Cerexa, Actavis-Allergan	DBO	Phase 1	NCT01689207 (completed); NCT02655419 (not open yet)
Imipenem-relebactam	Merck Sharp & Dohme Corporation	DBO	Phase 2	NCT01275170 and NCT01506271 (completed); and NCT01505634 (completed); NCT02452047 and NCT02493764 (recruiting)
RG6080 (formerly OP0595)	Meiji Seika Pharma Co, Ltd, Roche, and Fedora	DBO	Phase 1	NCT02134834 (completed)
Meropenem-RPX7009	Rempex Pharmaceuticals (The Medicines Company)	Boronate	Phase 3	NCT01897779, NCT02020434, and NCT02073812 (completed); and NTC02168946 and NCT02166476 (recruiting); NCT01751269 (completed); NCT02687906 (not open yet)
Biapenem-RPX7009	Rempex Pharmaceuticals (The Medicines Company)	Boronate	Phase 1	NCT01772836 (completed)
S-649266	Shionogi	Cephalosporin	Phase 2	NCT02321800 (recruiting); NCT02714595 (not open yet)

**Table 2**Promising new  $\beta$ -lactams or  $\beta$ -lactamase inhibitors in preclinical development

$\beta$ -Lactamase Inhibitor Name	Partner $\beta$ -Lactam	Company	Type of $\beta$ -Lactamase Inhibitor
FPI-1465	Aztreonam or ceftazidime	Fedora	DBO (also inhibits PBP activity)
WCK 4234	Meropenem	Wockhardt, Ltd	DBO
WO2013/030735	Not necessary?	Wockhardt, Ltd	DBO (also inhibits PBP activity)
WCK 5153	Not necessary?	Wockhardt, Ltd	DBO (also inhibits PBP activity)
Benzo(b)thiophene-2-boronic acid	Ceftazidime	Therabor and Regents of the University of California	Boronate
Sulfonamide boronates (CR161, compound 4, and compound 9)	Ceftazidime or cefotaxime	Therabor and Regents of the University of California	Boronate
S02030	Cefepime	Case Western Reserve University and Università degli Studi di Modena e Reggio Emilia	Boronate
3,4-dihydro-2H-benzo[e][1,2] oxaborinine-8-carboxylic acids	Ceftazidime or meropenem	VenatoRx Pharmaceuticals	Boronate
$\alpha$ -Aminoboronic acids	Ceftazidime	VenatoRx Pharmaceuticals	Boronate
3,4-Dihydro-2H-benzo[e][1,2] oxaborinine-8-carboxylic acids	Carbapenem	Rempex Pharmaceuticals (The Medicines Company)	Boronate
AA101	Cefepime	Allegra Therapeutics	Sulfone
Sulfone derivatives	Meropenem or imipenem	Orchid Pharmaceuticals	Sulfone
Sulfone derivatives	Meropenem or imipenem	Dr John D. Buynak (Southern Methodist University)	Sulfone
Clavam derivatives	Ceftazidime	Nabriva Therapeutics	Clavam
MG96077	Imipenem	Mirati Therapeutics	Phosphonate
BAL30072	Meropenem or no $\beta$ -lactam required	Basilea Pharmaceuticals	Siderophore monobactam
BAL30376 (BAL19764, BAL29880, & clavulanic acid)	No $\beta$ -lactam required	Basilea Pharmaceuticals	Siderophore monobactam, bridged monobactam, and a clavam
MK-8712	Imipenem	Merck Sharp & Dohme Corporation	Bridged monobactam
Siderophore monobactams	Aztreonam or meropenem	Pfizer	Siderophore monobactam
Syn2190	Ceftazidime	Taiho Pharmaceuticals Co	Siderophore monobactam
3'-Thiobenzoyl cephalosporins	Meropenem	University of Waterloo, Wilfrid Laurier University	$\beta$ -Lactam
FSI-1686 and FSI-1671	No $\beta$ -lactam required	FOB Synthesis Inc	$\beta$ -Lactam
BTZs	Imipenem	Universidad de la República, Montevideo, Uruguay	Bisthiazolidine
ME1071	Ceftazidime or biapenem	Meiji Seika Kaisha Ltd	Maleic acid derivative