

Syntheses and studies of quinolone-cephalosporins as potential anti-tuberculosis agents

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Abstract—The syntheses and anti-tuberculosis activity of quinolone-cephalosporin conjugates (**1** and **2**) are described. Both showed broad-spectrum antibacterial activity and significant anti-TB activity. The carbamate-linked quinolone-cephem **2** showed better antimycobacterial activity, including anti-TB activity, than the direct amine-linked quinolone-cephem **1**, while quinolone-cephem **1** was slightly more effective against some Gram-negative bacterial strains.
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Tuberculosis (TB) is one of the leading causes of death and suffering worldwide. The increasing microbial resistance, toxicity, and side effects of currently used anti-tuberculosis drugs also emphasize the urgent need for new, safer, and more effective anti-tuberculosis agents.^{1–5} Quinolonyl- β -lactams are a class of multifunctional β -lactams and some examples showed high potency and broad antibacterial activity against a range of clinically significant bacterial pathogens, especially against newly emerging strains of quinolone- and β -lactam-resistant bacteria, for example, methicillin-resistant ciprofloxacin-resistant (MRCR) *Staphylococcus aureus*, penicillin-resistant *Streptococcus pneumoniae*, and vancomycin-resistant *Enterococcus faecium*.⁶

However, no anti-TB activity data of quinolonyl- β -lactams were reported. Based on our previous work on the synthesis of a carbacephalosporin-quinolone conjugate,⁷ we describe here the syntheses and anti-TB activity of direct amine-linked quinolonyl-cephem **1** and carbamate-linked quinolonyl-cephem **2** (see Fig. 1).

The synthesis of the direct amine-linked quinolone-cephalosporin **1** is shown in Scheme 1. 3-Chloromethyl-7-phenylacetyl amino cephalosporanic acid *p*-methoxybenzyl ester (GCLE) was converted to 3-iodomethyl cephalosporin **3** in 92% yield according to the literature procedure.⁸ The key intermediate **4** was obtained in 30% yield by coupling compound **3** with ciprofloxacin that had been pretreated with *N*-methyl-*N*-(trimethylsilyl) trifluoro acetamide (MSTFA).⁹ Subsequent deprotection with TFA in the presence of anisole provided the desired product **1** in 70% yield.

The synthesis of carbamate-linked quinolone-cephalosporin compound **2** is shown in Scheme 2. This synthesis began with a controlled hydrolysis to remove the acetyl group of commercially available 7-amino cephalosporanic acid (7-ACA). Reaction with phenylacetyl chloride

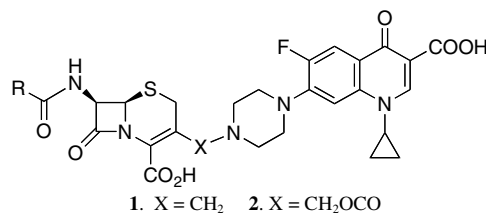
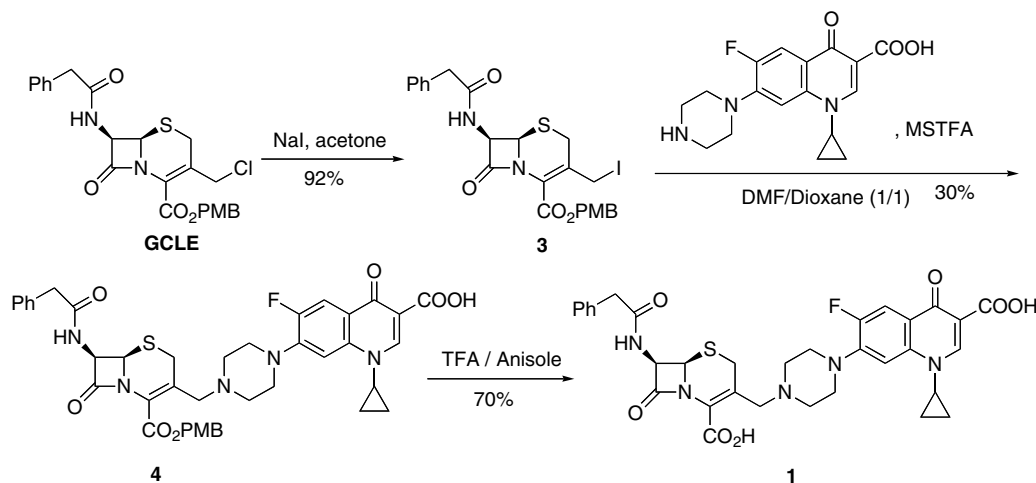


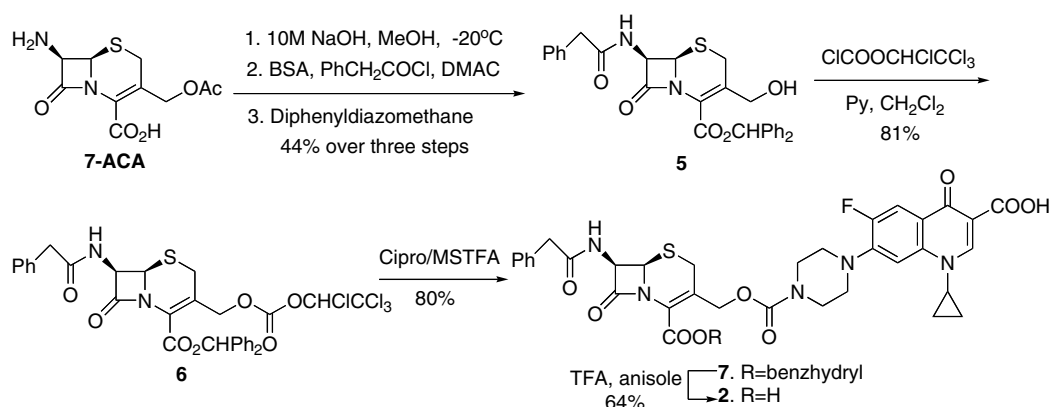
Figure 1. QLAs: quinolone-cephalosporin conjugates.

Keywords: Anti-tuberculosis; Antibacterial agents; Quinolonyl- β -lactams.

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Scheme 1. Procedure for synthesis of amine-linked quinolone-cephalosporin conjugate.



Scheme 2. Procedure for synthesis of carbamate-linked quinolone-cephalosporin conjugate.

followed by treatment with diphenyldiazomethane provided **5** in 44% yield for three steps.¹⁰ Subsequent coupling with tetrachloroethyl chloroformate gave intermediate carbonate **6** in 81% yield. Direct reaction with ciprofloxacin under basic aqueous conditions followed by deprotection gave the desired product, **2**, in 51% yield for two steps.¹¹

Quinolone-cephalosporin compounds **1**, **2**, **4**, **7**, as well as GCLE, ciprofloxacin, mixtures, and precursors **3** and **5**, were tested for their antibacterial activities against various strains of Gram-positive and Gram-negative bacteria, several strains of mycobacteria, as well as against *Mycobacterium tuberculosis* (Table 1).^{12–17} Compound **2** was found to be comparable to or by 1 to 2 dilution steps more active than compound **1** for all fast growing mycobacteria, and also for *M. tuberculosis* H37Rv. Interestingly, the very hydrophobic benzhydryl ester, compound **7**, was the most active compound against enterobacteria (*Escherichia coli* and *Enterobacter cloacae*), against staphylococci and enterococci as well as fast growing mycobacteria. On the other hand, **7** and ciprofloxacin were less effective against *Micrococcus luteus*, while conjugates **1**, **2**, and **4** were highly active

toward *M. luteus*. For the Gram-negative bacteria and for the enterococcus strain used here, ciprofloxacin alone was still more active than the corresponding β -lactam conjugates **1**, **4**, and **7**. As noted, the hydrophobic protecting group of **7** increased activity against staphylococci and enterococci relative to compounds **1**, **2**, and **4** that are more hydrophobic. Generally, compounds **1** and **2** with hydrophilic properties (or without hydrophobic protecting groups) were found to be more active than conjugate **4** with a hydrophobic (PMB) protecting group. Activity of all compounds against *Pseudomonas aeruginosa* did not depend on outer membrane permeability since essentially the same MIC values were obtained for the wild-type strain and for the mutant strain with an impaired outer membrane barrier. Activity of all compounds (except for the inactive precursor β -lactams **3** and **5**) against *E. coli* was increased for the permeability mutant DC2 in contrast to the wild-type strain DC0.

Compounds **1**, **2**, and **7** also showed good anti-TB activity, but starting GCLE, **4**, and conjugate precursors **3** and **5** displayed weak to no activity. Carbamate-based conjugate **2** was active against *M. tuberculosis* H37Rv

Table 1. Antibacterial activity of quinolone-cephalosporin compounds

Species/strains	MIC (μM)							
	1	4	2	7	Cipro ^a /GCLE tested separately	3	5	GCLE + Cipro
<i>Escherichia coli</i> DC0 wild type	3.12	25	3.12	1.56	0.4/200	200	>200	3.12
<i>Escherichia coli</i> DC2 permeability mutant	0.8	6.25	0.8	0.2	<0.1/NT ^c	200	200	NT
<i>Enterobacter cloacae</i> P99	0.1	0.4	0.2	<0.1	<0.1/NT	100	200	NT
<i>Micrococcus luteus</i> ATCC 10240	0.4	3.12	0.2	12.5	25/50	100	25	50
<i>Pseudomonas aeruginosa</i> KW 799/WT wild type	1.56	12.5	3.12	1.56	0.8/100	100	100	3.12
<i>Pseudomonas aeruginosa</i> KW 799/61 permeability mutant	1.56	12.5	3.12	1.56	0.4/50	100	100	1.56
<i>Mycobacterium vaccae</i> IMET 10670	3.12	3.12	1.56	1.56	0.8/100	100	200	6.25
<i>Mycobacterium smegmatis</i> SG987	3.12	12.5	3.12	1.56	0.8/NT	>200	200	NT
<i>Mycobacterium aurum</i> SB66	0.4	0.8	0.2	0.2	0.1/NT	100	50	NT
<i>Mycobacterium aurum</i> DSM 43436	0.4	1.56	0.2	0.2	0.1/NT	50	25	NT
<i>Mycobacterium fortuitum</i> B	3.12	12.5	0.8	0.8	0.2/100	200	200	1.56
<i>Staphylococcus aureus</i> SG 511	1.56	1.56	0.4	0.2	0.8/1.56	6.25	3.12	1.56
<i>Staphylococcus aureus</i> 134/93 MRSA	200	>200	200	100	50/NT	>200	>200	NT
<i>Enterococcus faecalis</i> 1528 VRE	12.5	50	6.25	3.12	1.56/NT	200	200	NT
<i>Mycobacterium tuberculosis</i> H37Rv MABA ^b /GAS ^c	3.8	62.3	1.6	8.0	1.8/31.5	>128	>128	1.8
Vero cells IC ₅₀ (μM) ^d	>128	13.1	>128	>128	NT/NT	NT	NT	NT

^a Ciprofloxacin from BioChemika.^b MABA, microplate alamar blue assay.¹⁵^c GAS, glycerol-alanine-salts.¹⁶^d Concentration resulting in 50% inhibition of the growth of VERO cells.¹⁷^e NT, not tested.

with an MIC of 1.6 μM . Interestingly, its benzhydryl ester **7** also was active with an MIC of 8.0 μM . The amine-linked quinolone-cephalosporin compound **1** was slightly less active than carbamate **2**. Compounds **1**, **2**, and **7** were found not to be cytotoxic against Vero cells, thus, making them worthy of consideration as potential anti-tuberculosis agents.

In summary, we have synthesized direct amine-linked quinolone-cephem **1** and carbamate-linked quinolone-cephem **2**, and evaluated them and their corresponding esters **4** and **7** for antibacterial activity. Of all conjugates, **2** had superior antimycobacterial/anti-TB activity.

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Supplementary data

Supplementary data associated with this article can be found, in the online version, at [doi:10.1016/j.bmcl.2006.08.045](https://doi.org/10.1016/j.bmcl.2006.08.045).

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