

The Synthesis and SAR of Rhodanines as Novel Class C β -Lactamase Inhibitors

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Received 27 March 2000; accepted 24 July 2000

Abstract— β -Lactam antibiotics such as the cephalosporins and penicillins have diminished clinical effectiveness due to the hydrolytic activity of diverse β -lactamases, especially those in molecular classes A and C. A structure–activity relationship (SAR) study of a high-throughput screening lead resulted in the discovery of a potent and selective non- β -lactam inhibitor of class C β -lactamases. © 2000 Elsevier Science Ltd. All rights reserved.

Bacterial resistance to β -lactam antibiotics can result from their overuse in treating mild infections and from patient failure in following a prescribed dosing regimen. The surviving bacteria often become resistant due to the β -lactam hydrolyzing enzymes known as β -lactamases.¹ To circumvent this resistance, bacterial infections have been treated by administering a β -lactamase inhibitor with the β -lactam antibiotic.

β -Lactamases are divided into four classes based on their molecular structure. Class C and class A β -lactamases, with an active site serine, represent the majority of all β -lactamases.² Commercially available inhibitors such as clavulanic acid (**1**), tazobactam (**2**, TZB), and sulbactam (**3**) inactivate most common class A β -lactamases, but are ineffective clinically against class C β -lactamases (Fig. 1).³ Chromosomally-encoded class C β -lactamases rapidly degrade cephalosporins, and they are generally not inhibited by the existing β -lactamase inhibitors. The most active broad-spectrum β -lactamase inhibitor that is effective against class A and C β -lactamases is BRL 42715 (**4**). However, this compound was not developed as a clinical agent.⁴

The class C cephalosporin-hydrolyzing β -lactamase class is an attractive target for drug discovery. As a dominant resistance mechanism in Gram-negative bacteria, it is expected to become more widespread in the absence of a commercially available inhibitor.

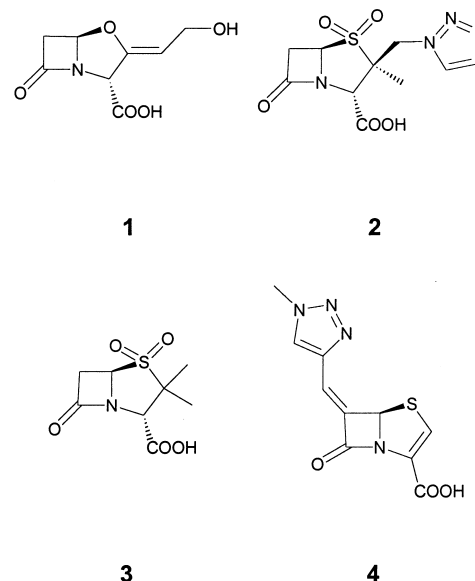


Figure 1.

High-throughput screening of our chemical library identified rhodanine **5** as a promising lead inhibitor candidate (Fig. 2). P99, isolated from *Enterobacter cloacae*, and TEM-1, isolated from *Escherichia coli*, were used in these studies as prototypical class C and class A β -lactamases.⁵ Rhodanine **5** is a non- β -lactam with an IC_{50} of 2.6 μ M against the class C enzyme P99 and an IC_{50} of 8.7 μ M against the class A enzyme TEM-1. This paper describes our efforts to discover a

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rhodanine β -lactamase inhibitor which is selective for P99 over TEM-1.

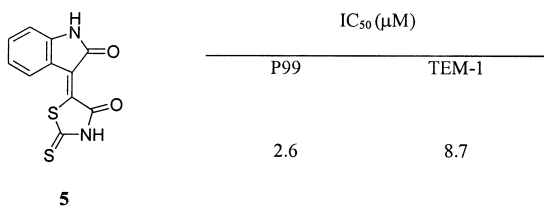


Figure 2.

The simplest modification of **5** was to excise the pyrrolidone ring and to construct analogues by condensation of substituted aromatic aldehydes with 2-thio-thiazolidi-4-none (Fig. 3). In general, synthesis of these alkylidene rhodanines was facile, and only the *Z* isomer was isolated.⁶

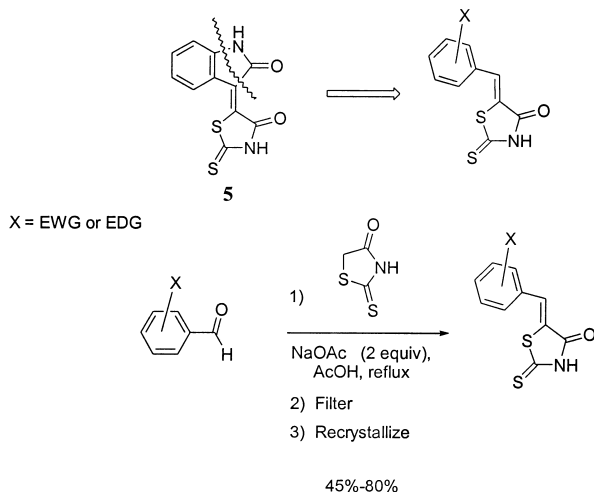


Figure 3.

A wide variety of rhodanines was prepared and tested as inhibitors of P99 and TEM-1 β -lactamases. Examining the activity of selected rhodanines, two general trends emerged. First, the most selective compounds for P99 lacked a substituent at the ring nitrogen (Table 1). Second, the analogues with a nitro group on the benzyldene phenyl ring had superior activity.

The relatively weak but selective activity of these analogues encouraged us to extend the sidechain with a two-carbon spacer in order to mimic the cephalosporin substrate of class C β -lactamases. The addition of the two-carbon spacer led to improved IC₅₀'s, while maintaining essentially the same trends observed for the simpler analogues (Table 2). Interestingly, the activity of the saturated analogue **19** was superior to unsaturated **17**, however the selectivity for P99 was diminished. Preparation of the *E* isomer at C3 of **17** was attempted, but was unsuccessful.

Finally, we decided to examine heterocyclic sidechains. Substitution of an electron withdrawing triazole for the double bond yielded a moderately active analogue, **20**, while the imidazole analogue **21** was inactive. Replacement of the phenyl group with a furan to give analogue **23** further increased the activity. The most active analogue in this study was the 5-nitrofuran analogue **24**. Rhodanine **24** was 200-fold more selective for P99 than TEM-1, as compared to our original rhodanine **5** which was only threefold more selective.

We next examined **24** for synergy with the β -lactam piperacillin in a six-organism panel using TZB as a positive control (Table 3). This panel includes bacterial strains producing class A enzymes and isogenic strains of *S. marcescens* and *P. aeruginosa* producing class C β -lactamases in an inducible or constitutive state. As demonstrated in this panel, piperacillin MICs were lowered against *S. marcescens* and *P. aeruginosa* strains

Table 1. Inhibitory activity of substituted rhodanines⁷ against P99 and TEM-1⁸

Compound	R ¹	R ²	IC ₅₀ (μ M)	
			P99	TEM-1
6	Phenyl	H	455	>700
7	Phenyl	Ethyl	240	432
8	Phenyl		>700	>700
9	Phenyl		246	138
10	Phenyl		435	>700
11		H	335	>700
12		H	>700	>700
13		H	290	>700
14		H	43	>700
15		H	500	>700
16		H	>700	190

Table 2. Inhibitory activity of substituted rhodanines against P99 and TEM-1

Compound	X	R ¹	IC ₅₀ (μM)	
			P99	TEM-1
6	S		455	>700
17	S		76	>700
18	S		4.2	100
19	S		9.7	32
20	S		83	>700
21	S		>700	>700
22	S		8	>100
23	S		62	100
24	S		0.45	100
25	O		>50	>50
2		TZB	7.7 ^a	0.06

^aAverage IC₅₀ of tazobactam in 10 experiments.**Table 4.** Antimicrobial activity of rhodanines **24** and **25**^a

Bacteria	Enzyme	MIC (μg/mL)	
		24	25
<i>E. coli</i> OC 4075	TEM-1	2	1
<i>E. cloacae</i> OC 4080	P99	2	0.5
<i>E. cloacae</i> OC 4092	E2 (ind)	32	4
<i>E. cloacae</i> OC 4081	E2 (const)	32	4
<i>P. aeruginosa</i> OC 4270	AmpC (ind)	32	32
<i>P. aeruginosa</i> OC 4290	AmpC (const)	32	32

^aSee Table 3 footnotes for abbreviations.

with inducible or constitutive production of class C cephalosporinases. Because of the inherent antibiotic activity of **24**, it was impossible to evaluate the synergistic potential at concentrations greater than its MIC.¹⁰

The nitrofuranyl moiety of **24** was suspected to be associated with the antibacterial activity since 2-nitrofurans, such as nitrofurantoin, are known antibacterials.¹² To test this hypothesis, thiazolidinedione **25** was prepared and tested in the biological assays (Table 4). Thiazolidinedione **25** had better MICs against *E. cloacae* isolates as compared to **24**, but **25** lacked inhibitory activity at the concentrations tested (IC₅₀ >50 μM). Therefore, it is likely that the nitrofuranyl moiety was not responsible for inhibition of P99 in **24**, but is responsible for the antibacterial activity of **24**.

In conclusion, the rhodanines represent a novel class of β-lactamase inhibitors that can be modified to show selectivity for class C β-lactamases. The rhodanines differ chemically from traditional β-lactamase inhibitors such as clavulanic acid and tazobactam, but do exhibit inhibitory activity against both class A and class C β-lactamases.

Table 3. Susceptibility testing of **24** in combination with piperacillin

Bacteria	Enzyme	Molecular class	MIC (μg/mL) ^a				
			Piperacillin	PIP/TZB ^b	24	PIP/24 (1) ^c	PIP/24 (4) ^c
<i>E. coli</i> OC 4075	TEM-1	A	>64	8	2	>64	NA ^d
<i>E. cloacae</i> OC 4080	P99	C	>64	64	2	>64	NA
<i>S. marcescens</i> OC 4101	AmpC (ind) ^e	C	8	4	8	2	2
<i>S. marcescens</i> OC 4294	AmpC (const) ^f	C	>64	32	8	32	8
<i>P. aeruginosa</i> OC 4270	AmpC (ind) ^e	C	8	8	32	4	4
<i>P. aeruginosa</i> OC 4290	AmpC (const) ^f	C	64	32	32	16	8

^aMICs were determined using standard methods according to NCCLS.¹¹^bPIP, piperacillin; TZB, tazobactam at 4 μg/mL.^cInhibitor concentration in μg/mL shown in parentheses.^dNA, not applicable.^eInducible β-lactamase production.^fConstitutive β-lactamase production.

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