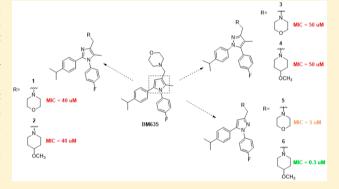
Novel Pyrazole-Containing Compounds Active against Mycobacterium tuberculosis

Giovanna Poce,*,†© Sara Consalvi,† Giulia Venditti,† Salvatore Alfonso,‡ Nicoletta Desideri,† Raquel Fernandez-Menendez,‡ Robert H. Bates,‡ Lluis Ballell,‡ David Barros Aguirre,‡© Joaquin Rullas,‡ Alessandro De Logu,§ Michelle Gardner, Thomas R. Ioerger, Eric J. Rubin, and Mariangela Biava*,†

Supporting Information

ABSTRACT: In this study, a series of 49 five-membered heterocyclic compounds containing either a pyridine- or a pyrrole-type nitrogen were synthesized and tested against Mycobacterium tuberculosis. Among them, only the 1,3,5trisubstituted pyrazoles 5-49 exhibited minimum inhibitory concentration values in the low micromolar range, and some also exhibited an improved physicochemical profile without cytotoxic effects. Three pyrazoles were subjected to an animal tuberculosis efficacy model, and compound 6 induced a statistically significant difference in lung bacterial counts compared with untreated mice. Moreover, to determine the target of this series, resistors were generated, and whole genome sequencing revealed mutations in the mmpL3 gene.



KEYWORDS: Tuberculosis, pyrazoles, MmpL3, drug discovery, antimycobacterials

uberculosis (TB) is the leading cause of death worldwide from a single infectious agent and ranks above HIV/ AIDS. An estimated 1.3 million people died of TB and ~10 million fell ill with TB in 2017. Luckily, TB mortality has fallen 47% since 1990, with an estimated 53 million saved lives between 2000 and 2017. Despite that, the emergence and spread of multi- and extensively drug-resistant (MDR and XDR) strains have made the control of the disease more difficult. Therefore, there is a high priority on the development of new anti-TB drugs with bactericidal mechanisms different from those of the presently available agents.²

Our interest in developing new antitubercular agents reflects our work on developing a class of 1,5-diphenyl pyrroles. An initial phenotypic screening of an in-house chemical library identified BM212 (Figure 1) as a "soft hit", and further medicinal chemistry efforts provided an improved hit (BM635, Chart 1) active against both replicating and nonreplicating bacilli and efficacious in a murine model of TB infection.⁴ These pyrroles act by inhibiting the Mycobacterium membrane protein Large 3 (MmpL3).6,7 MmpL3 belongs to the resistance, nodulation, and cell division protein superfamily and is likely involved in trehalose monomycolate flipping across the inner membrane of M. tuberculosis (Mtb). We have

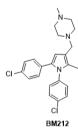


Figure 1. Chemical structure of BM212.

previously demonstrated that BM212 binds MmpL3 directly and inhibits its activity.8

After the discovery of BM635, we followed diverse strategies for improving BM635's drug-like profile (low water solubility and high lipophilicity), such as preparing different pharmaceutical salts⁹ and making modifications around the central pyrrole core. ¹⁰

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Department of Chemistry and Technologies of Drug, Sapienza University of Rome, Piazzale A. Moro 5, 00185 Rome, Italy

[‡]Diseases of the Developing World, GlaxoSmithKline, Calle Severo Ochoa 2, 28760 Tres Cantos, Madrid, Spain

[§]Department of Life and Environmental Sciences, University of Cagliari, via Ospedale 72, 09124 Cagliari, Italy

Department of Immunology and Infectious Diseases, Harvard T.H. Chan School of Public Health, 677 Huntington Avenue, Boston, Massachusetts 02115, United States

¹Department of Computer Science, Texas A&M University, 3112 TAMU, College Station, Texas 77843, United States

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Chart 1. Chemical Structures of BM635 and Compounds 1-6

To further develop this class of compounds, and with the aim of improving both the solubility and the polarity of the BM series, we have designed new compounds containing a second nitrogen atom on the heterocyclic moiety that could easily lead to the formation of hydrogen bonds. Therefore, the central core has been replaced with two five-membered heterocycles containing either a pyridine- or a pyrrole-type nitrogen: ^{11–15} imidazole and pyrazole. Herein we report the synthesis and the full biological and pharmacological profiling of such novel five-membered heterocyclic MmpL3 inhibitors.

We first investigated which five-membered ring could replace the central pyrrole core of **BM635** by preparing compounds 1-6 (Chart 1), but only the 1,3,5-trisubstituted pyrazoles 5 and 6 showed good activity against *Mtb*, with 6 providing a minimum inhibitory concentration (MIC) of 0.3 μ M (Table 1).

Thus we selected the 1,3,5-trisubstituted pyrazole as a suitable central core and prepared analogues 7–49 (Tables 2 and 3) decorated with the best substituents resulting from the previous works. We explored which amine could replace the morpholine by introducing different six-membered cyclic amines at position 3. To date, compounds 13, 23, 40, and 47 incorporate a silicon amine, which was shown, in a previous work by Ramesh and coworkers, to be responsible for a remarkable increase in activity. ¹⁶ Then, we examined positions 1 and 5 by introducing substituents that in the pyrrole series produced the best activities (compounds 14–49, Tables 2 and 3). ^{4,10,17}

Compounds were evaluated in vitro to determine the minimum concentration required to inhibit Mtb growth in the culture as well as the compound cytotoxicity by measuring the concentration of the compound resulting in 50% inhibition (CC50) of HepG2 cells. Moreover, compounds with MICs < 1 μ M were evaluated to determine the drug metabolism and

pharmacokinetics (DMPK) profile (Chrom log $D_{\rm pH7.4}$ values, artificial membrane permeability, percentage of binding to human serum albumin (HSA), kinetic solubility, solubility in the biorelevant medium to simulate fasted states in vivo (FaSSIF), and human ether-a-go-go-related gene (hERG)) binding, and those providing the best profile were selected to determine their efficacy in vivo. Finally, having modified the main chemical motif of the parent compounds, we verified if the new pyrazoles act by inhibiting MmpL3 by generating spontaneous resistant mutants and whole genome sequencing.

Structure-activity relationship (SAR) studies were performed on a series of 1,3,5-trisubstituted pyrazoles. Compounds 1-49 were prepared as reported in the Supporting Information. Antituberculosis assays revealed that within the different substitutions at position five, only the 4-isopropylphenyl (compounds 6, 7, 11-17, 20, 22-24, and 26-29) and the trifluoromethylphenyl (compounds 36, 37, 39-47, and 49) ones were well tolerated, confirming a trend already seen for the pyrrole series (Tables 2 and 3). Compounds 6, 12– 14, 16, 22, 23, 27, 36, 40, and 47 showed remarkable activities against Mtb (0.00925 to 0.6 μ M, Tables 1-3) with 6, 12, 14, 22, 27, and 36 providing low cytotoxicities (3.6-64 μ M, Tables 1-3), whereas, a different trend was observed for the substituent at position one (compounds 24–26, 48, and 49): The isopropyl chain, which provided one of the best pyrrole compounds, did not work well for the pyrazoles (MIC = 1.75, 50, 1, 23, and 2.5 μ M, respectively). In general, the activity of this series of pyrazoles seemed to be more influenced by the cyclic amine at position three. All of the compounds incorporating a silicon amine (13, 23, 40, and 47) showed a remarkable activity increase (0.00925, 0.04, 0.02, and 0.07 μ M, respectively), but unfortunately they exhibited high cytotoxicity together with high levels of hERG activity, low water solubility, as well as low membrane permeability (Table 4). Both the high toxicity and the DMPK may be explained by the high lipophilicity displayed by silicon compounds. In view of the limited safety and poor DMPK, silicon-incorporating compounds were not further progressed. Instead, compounds bearing either the 4-methoxypiperidine or the pyridinepiperazine at position three (6, 12, 16, 22, and 36) showed the best results. In general, these compounds showed good activities and low cytotoxicities combined with good DMPKs and low hERG activities (Table 4). Surprisingly, when moved to the pyrazoles, the morpholine ring that did work well for the previous pyrrole compounds^{4,10} did provide only two active compounds: 14 and 27.

We sought to establish physicochemical, DMPK, and hERG profiles for selected compounds. Compounds showing MICs <1 μ M were profiled for physicochemical properties, in vitro DMPK properties, and hERG activities, as shown in Table 4. Overall, the introduction of a free pyrrole-type nitrogen led to both increased water solubility and reduced binding to HSA, as

Table 1. Activity against Mtb H37RV (MIC) and Cytotoxicity on HepG2 Cells (CC50) of Compounds 1-6, BM635, and Isoniazid

	1	2	3	4	5	6	BM635	isoniazid
$MIC (\mu M)^a$	40	40	50	50	5	0.3	0.12	1.8
CC50 $(\mu M)^b$	50.1	nd ^c	>100	nd ^c	50.1	31.6	40	
PI $(CC50/MIC)^d$	1.25		>2		10	105	333	

^aMinimum inhibitory concentration required to inhibit *Mtb* growth. ^bConcentration of compound resulting in 50% inhibition of HepG2 cell line. ^cNot determined. ^dProtection Index (CC50/MIC).

Table 2. Activity against Mtb H37RV (MIC) and Cytotoxicity on HepG2 Cells (CC50) of Compounds 7-29

		R	,		Comp.	R	MIC ^α (μM)	CC50 ^b (μM)	PI ^d (CC50/MIC) 40
						\$-N_N_O	1	40	40
F					21	ξ−N_NH	20	nd ^c	-
-	,OCH₃	(μM)	(μM)	(CC50/MIC)	22	₹-N N-{ }	0.5	40	80
7	ξ−N	2.4	32		23	}_N_	0.04	12.6	315
8	ξ− N → OH	10	20	2		\$ 14	0.04 R	12.0	
9	OH OH	20	25.1	1.25			N	N	
				0.6		\rightarrow		_	
10	{-N_N-	20	12.6		Comp.	R	MIC ^α (μM)	CC50 ^b (μΜ)	PI ^d (CC50/MIC)
11	\$ N N	2	80	40	24	ξ−NO	1.75	79.4	45
12	}-N_N-_N_	0.5	64	128	25	ξ−N →OCH ₃	50	nd ^c	-
13	ξ-N Si	0.00925	12.6	1362	26	Ş-N OH	1	50	50
	\ <u></u>	R	N			\ \	O N R) N	
Comp.	R	MIC ^a (μM)	CC50 ^b (μM)	PI ^d (CC50/MIC)	Comp.	R	MIC ^a (μM)	СС50 ^ь (µМ)	PI ^d (CC50/MIC)
14	§−N _O	0.4	63.1	158	27	\	0.4	50	125
15	₹-N_S	1.3	40	31	28	₹— О	2	>100	>50
16	ξ-N → OCH ₃	0.6	12.6	21			H ₃ CO		
17	QCH ₃	2.5	20	8		Y	N.R	1	
18	ξ− N —ОН	42	63	1.5	Comp.	R	MICα (μM)	CC50 ^b (μΜ)	PI ^d (CC50/MIC)
19	}−NN−	21	10	0.5	29	₹— <u></u>	2.5	50	20

^aMinimum inhibitory concentration required to inhibit Mtb growth. ^bConcentration of compound resulting in 50% inhibition of HepG2 cells. ^cNot determined. ^dProtection Index (CC50/MIC).

Table 3. Activity against Mtb H37RV (MIC) and Cytotoxicity on HepG2 Cells (CC50) of Compounds 30-49

		R'N						F ₃ C	N		
Comp	R	R'	MIC ^α (μM	СС50 ^ь (µМ)	PI ^d (CC50 /MIC)	Comp.	R	R'	ΜΙC ^α (μΜ)	СС50 ^ь (µМ)	PI ^d (CC50 /MIC)
30	-СНЗ	ξ−NO	2.5	50	20	41		§−N_O	2.5	50	20
31	-СНЗ	och3	50	>100	>2	42	* —	§−N S	1.2	63.1	53
32	N Zho	§−N_O	50	>100	>2	43	\ 	och3	1.2	31.6	26
33	N The	Serie N	50	>100	>2	44	\\	ξ−N	2.5	50	20
	OCH ₃					45		NNO	2	>100	>50
	F ₃ C					46	X —	see N	1	50	50
Comp.	R		MIC ^α (μM)	СС50 ^ь (µМ)	PI ^d (CC50 /MIC)	47	\	₹-N Si	0.07	12.6	180
34	× 0		20	100	5	48	***	ξ−N_O	23	>100	>4
35	§-N_s		4.7	79.4	17	49	~~~	ser' N	2.5	79.4	32
36	~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~		0.3	31.6	105	BM63		OCH ₃			333
37	OCH ₃		2.5	40	16	5	-	-	0.12	40	
	2,3		2.3	40	>25	Isonia zid	-	-	1.8	-	-
38	o v v v v v v v v v v v v v v v v v v v		4	>100	723						
39	road N		2	32	16						
40	§−N Si		0.02	16	800						

 $[^]a$ Minimum inhibitory concentration required to inhibit Mtb growth. b Concentration of compound resulting in 50% inhibition of HepG2 cells. c Protection Index (CC50/MIC).

Table 4. DMPK and Safety Profile of Compounds 6, 12, 13, 14, 22, 23, 27, 36, 40, 47 and BM635

comp	Chrom log $D_{\rm pH7.4}^{a}$	membrane permeability $\mathrm{cm/s}^b$	%HSA ^c	KS $(\mu M)^d$	FaSSIF $(\mu g/mL)^e$	hERG IC50 $(\mu M)^f$
6	7.31	4.4×10^{-5}	96.42	152	nd^g	6.31
12	8.76	$<0.3 \times 10^{-5}$	98.45	6	nd^g	25.1
13	8.82	4-Far	97.49	41	nd^g	2
14	8.29	3.5×10^{-5}	97.11	78.5	100	>50
22	9.83	$<0.03 \times 10^{-5}$	98.41	15	nd^g	>50
23	nd^g	4-Far	97.35	46	nd^g	8
27	7.37	6.1×10^{-5}	96.72	305	139	32
36	6.67	1.7×10^{-5}	96.47	258	nd^g	3.2
40	8.13	4-Far	96.65	53	nd^g	1.26
47	8.49	4-Far	96.32	69	nd^g	2.5
BM635	8.1	2.4×10^{-5}	98.37	<1	5	10

^aLipophilicity expressed as Chrom log $D_{pH7.4}$. ^bArtificial membrane permeability. ^cPercentage of binding to human serum albumin (HSA). ^dKinetic solubility. ^eSolubility in the biorelevant medium to simulate the fasted states in vivo (FaSSIF). ^fHuman ether-a-go-go-related gene (hERG) binding. ^gNot determined.

for compounds 6, 14, 27, and 36 (Table 4), compared with the previous pyrroles. ¹⁰ The ability of compounds to cross biological membranes was assessed by an artificial membrane permeability assay, which is based on permeation through planar polycarbonate-filter-supported lipid bilayers. Most of the tested compounds proved to be highly permeable, with values in the 10⁻⁵ range, except for silicon-containing compounds 13, 23, 40, and 47 (Table 4). Finally, whereas compounds 6, 13, 23, 36, 40, and 47 were found to be low micromolar inhibitors of the potassium channel, 12, 14, 22, and 27 had a clean profile in the hERG activity (Table 4).

Because the pyrazoles and pyrroles have different SARs, they could possibly have different targets. To determine this, we isolated resistant mutants to two of the compounds, **6** (at $3 \times$ MIC) and **22** (at $3 \times$ and $5 \times$ MIC), with a resistance frequency of 10^{-7} and performed whole genome sequencing on DNA from isolated colonies (Table 5). We found no mutations in a colony grown on the higher concentration of **22**. At the lower compound concentrations, we found multiple mutations;

Table 5. MICs and Amino Acid Substitutions Found in Mtb Mutants Isolated as Resistant to Compounds 6 and 22

comp	MIC (μM)	gene	mutation
6	12.5	mmpL3	V240A
6	12.5	ppsB	L1476S
22	20	mmpL3	P239H
22	25	lprM	V86V
22	20	fadD26	+g in aa 336

however, both had mutations that altered the coding sequence of MmpL3. For 6, we found a substitution of alanine for valine at position 240 (along with a missense mutation in ppsB). For 22, we found a substitution at the adjacent position, 239, of histidine for proline (along with a missense mutation in lprM and an insertion mutation in fadD26). These mutations both lie in the fourth transmembrane helix, where other resistance mutations to other MmpL3-targeting compounds are known to exist. ¹⁹

Finally, to evaluate the potential of these compounds, we selected **6**, **14**, and **27** to be tested in a fast standardized 9 day acute assay intended to measure the efficacy of drugs against *M. tuberculosis* growing in the lungs of immunocompetent mice. ¹⁸ Compounds **6**, **14**, and **27** where chosen because of their low cytotoxicities (31.60, 63.10, and 40 μ M) but

especially because of the high water solubility (152, 78.50, and 305.50 μ M), which is expected to play a crucial role in the in vivo efficacy. Compounds were administered at a dose of 200 mg/kg per day for 4 days starting on day 5 after infection. Surprisingly, only 6 induced a statistically significant difference in lung bacterial counts compared with untreated mice (Table 6).

Despite the good permeability and good water solubility, 14 and 27 proved to be inactive. Because poor metabolic stability may account for the lack of in vivo efficacy, we tested 14 and 27 on mouse and human liver microsomes to determine the

Table 6. Therapeutic Efficacy of 6, 14, and 27 and Moxifloxacin^a

comp	dose (mg/kg)	administration	difference in log CFU/lungs	P^2
6	200	once per day	1.5	p < 0.05
14	200	once per day	0.4	p < 0.05
27	200	once per day	0.3	p < 0.05
moxifloxacin	30	once per day	3.5	p < 0.05

^aDifferences in the lung microorganism burden (log 10 CFUs/lungs) with respect to untreated controls (day 9 after infection).

intrinsic clearance (Clint) and the metabolic stability (%LBF). On the basis of clearance in mouse microsomes (7.98 and 7.04 mL/min/g) and applying a well-stirred model, an in vivo blood clearance of \sim 75% for both compounds is predicted (Table 7). This would explain the lack of efficacy of those compounds related to an expected high first-pass liver effect and a quick elimination process.

In summary, following up on our medicinal chemistry campaign based on the implementation of a class of potent MmpL3 inhibitors, we sought to replace the central pyrrole core and prepared 49 five-membered heterocyclic compounds.

Table 7. Intrinsic Clearance (Clint) in Mouse and Human Microsomes of Compounds 14 and 27

	Clint (m	L/min/g)	%LBF ^a		
comp	mouse	human	mouse	human	
14	7.98	3.4	76.3	82.2	
27	7.04	2.8	74	79.2	

^aLiver blood flow, predicted in vivo blood clearance.

The introduction of a free pyrrole-type nitrogen gave several compounds that proved to be active against Mtb with a low level of cytotoxicity but, above all, with improved water solubility and reduced hERG activity—two of the major concerns in TB drug development. Moreover, one of the tested pyrazoles showed efficacy in a murine model of acute TB infection. This, combined with the results of whole genome sequencing highlighting MmpL3 as the possible target, confirms the high potential of the MmpL3 inhibitor class for development in TB drug discovery.

■ ASSOCIATED CONTENT

S Supporting Information

The Supporting Information is available free of charge on the ACS Publications website at DOI: 10.1021/acsmedchemlett.9b00204.

Detailed synthetic procedure, spectroscopic data, and full biological characterizations of all new compounds (PDF)

AUTHOR INFORMATION

Corresponding Authors

*E-mail: giovanna.poce@uniroma1.it. *E-mail: mariangela.biava@uniroma1.it.

ORCID ®

Giovanna Poce: 0000-0001-7466-8861 David Barros Aguirre: 0000-0002-4099-0438

Author Contributions

All authors have given approval to the final version of the manuscript.

Notes

The authors declare no competing financial interest.

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ABBREVIATIONS

CFUs, colony forming units; DMPK, drug metabolism and pharmacokinetics; FaSSIF, fasted simulated fluid; hERG, human ether-a-go-go-related gene; HSA, human serum albumin; LBF, liver blood flow; MDR, multi-drug-resistant; MIC, minimum inhibitory concentration; MmpL3, *Mycobacte-rium* membrane protein large 3; Mtb, M. tuberculosis; SAR, structure—activity relationship; TB, tuberculosis; WHO, World Health Organization; XDR, extensively drug resistant.

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