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## Context

*Mycobacterium tuberculosis* (*M. tb*) is responsible for 10.6 million infections and 1.6 million deaths worldwide in 2020.<sup>[1]</sup> However, in Europe and North America, the incidence of **non-tuberculous mycobacteria (NTM)** infections exceeds that of *M. tb*.<sup>[2]</sup> Among NTM with **pulmonary pathogenicity**, *M. abscessus* (*M. abs*) and *M. avium complex* (MAC) are the most common.<sup>[2]</sup> As for tuberculosis cure, NTM treatments require a **combination of antibiotics** which can induce **many side effects** for a **long period** (Figure 1).<sup>[2,3]</sup>

There is an urgent need to develop **new antimycobacterial drugs** that are safer, more specific to NTM and, if possible, with a new mechanism of action to limit cross-resistance.

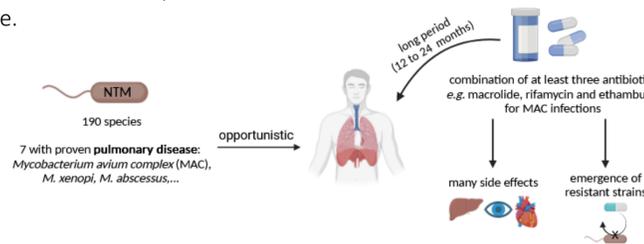


Figure 1: NTM as emerging virulent pathogens responsible for difficult-to-treat lung diseases.

## Antimycobacterial quinolines

Quinoline-based pharmacophore is found in **bedaquiline (BQ)** and **mefloquine (MQ)** (Figure 2). They are **active against NTM** and target **ATP synthase** but not only in the case of MQ.<sup>[4,5,6]</sup> However, BQ and MQ can induce liver and central nervous system disorders respectively.<sup>[7,8]</sup> Thus, we are interested in developing new **amino-alcohol-quinolines (AAQ)** as MQ analogs to establish new structure-activity relationships (SAR) and to reduce MQ toxicity.

*In vitro* evaluations of previous series have identified the **hit A**. It is **effective on MAC** (MIC = 4 µg/mL) and has a **better selectivity index (SI)** than MQ (SI = 2.9 vs 0.4).<sup>[5]</sup> To increase their SI, **new AAQ with piperazine core** grafted with aryl or alkyl groups have been designed.

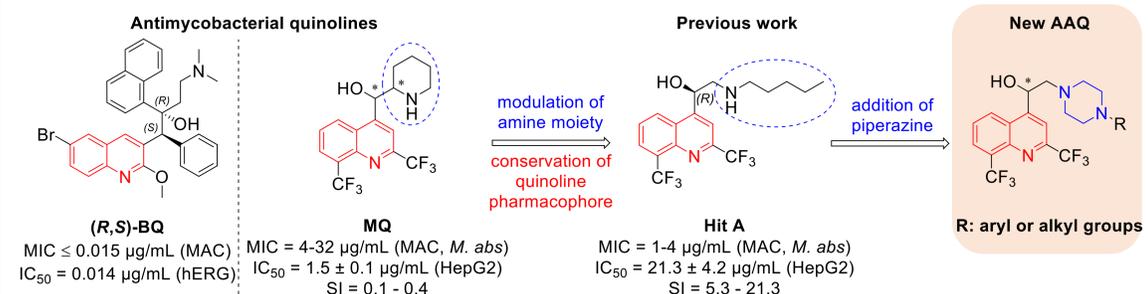
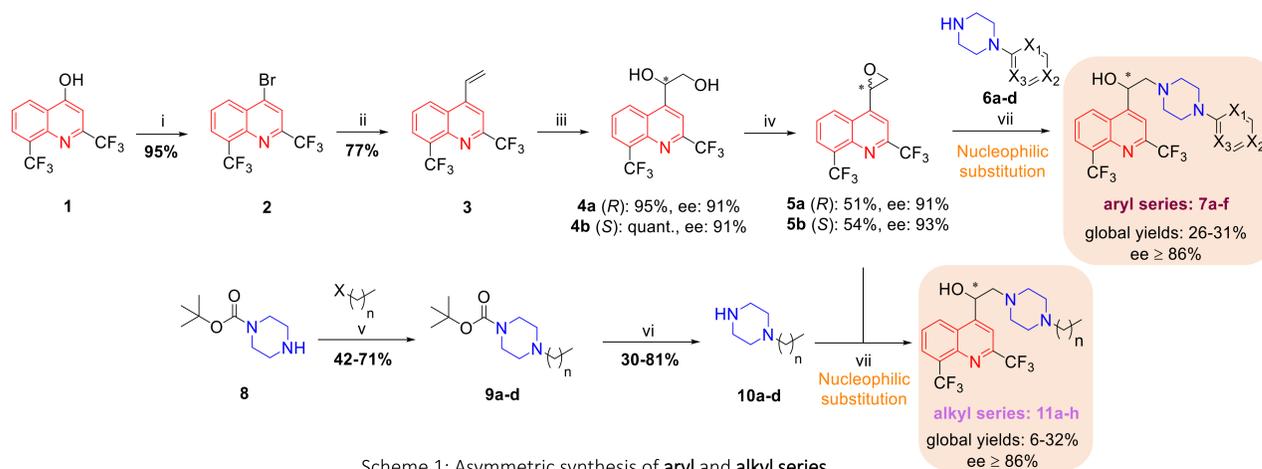


Figure 2: Known antimycobacterial quinolines and development of new amino-alcohol-quinolines.

## Synthesis

To synthesise **new aryl** and **alkyl series**, two intermediates were required: i) the **epoxydes 5a/b** obtained from 4-hydroxyquinoline **1** in four steps and, ii) the **amines 6a-d** and **10a-d** (Scheme 1). Aryl-piperazines **6a-d** were commercially available, while alkyl-piperazines **10a-d** were obtained in two-step synthesis from *tert*-butyl piperazine-1-carboxylate **8**. Both entities were conjugated by nucleophilic substitution to afford **three enantiomer pairs 7a-f** and **four enantiomer pairs 11a-h** in 6 to 31 % of global yields and an enantiomeric excess (ee) greater than 86 % (Table 1).



Scheme 1: Asymmetric synthesis of aryl and alkyl series.

i: POBr<sub>3</sub>, 150 °C; ii: potassium vinyltrifluoroborate, Cs<sub>2</sub>CO<sub>3</sub>, PdCl<sub>2</sub>(dppf), DCM, THF/H<sub>2</sub>O 9/1, 70 °C; iii: AD-mix α or β, K<sub>2</sub>[(OsO<sub>2</sub>(OH))<sub>2</sub>], tBuOH/H<sub>2</sub>O 1/1; iv: 1) MeC(OMe)<sub>3</sub>, PTSA, H<sub>2</sub>O, DCM, 2) TMSBr, DCM, 3) K<sub>2</sub>CO<sub>3</sub>, MeOH; v: DIPEA, MeCN; vi: 1) TFA, DCM, 2) NaOH, DCM; vii: 130 °C, 150 W, EtOH.

Table 1: Global yields and ee of aryl and alkyl series.

Compound	X <sub>1,2,3</sub>	n	Global yield (%)	ee (%)
7a (R)	X <sub>1,3</sub> = CH, X <sub>2</sub> = C-Me	-	30	94
7b (S)		-	31	92
7c (R)	X <sub>1</sub> = N, X <sub>2,3</sub> = CH	-	28	87
7d (S)		-	26	86
7e (R)	X <sub>1,3</sub> = N, X <sub>2</sub> = CH	-	31	85
7f (S)		-	26	86
11a (R)	-	3	14	89
11b (S)	-	3	18	91
11c (R)	-	4	6	86
11d (S)	-	4	32	87
11e (R)	-	5	10	86
11f (S)	-	5	17	88
11g (R)	-	6	15	85
11h (S)	-	6	13	85

## Physicochemical and pharmacokinetic properties

*In silico* physicochemical (PC) and pharmacokinetic (PK) profiles of **aryl** and **alkyl compounds 7a-f** and **11a-h** were performed using Qikprop software (Figure 3). **Lipinski's** and **Weber's rules** were chosen for the PC data. Solubility, metabolic reactions, cardiac toxicity and permeability of Caco-2 cells parameters were selected to reflect **A.D.M.E.T properties**.

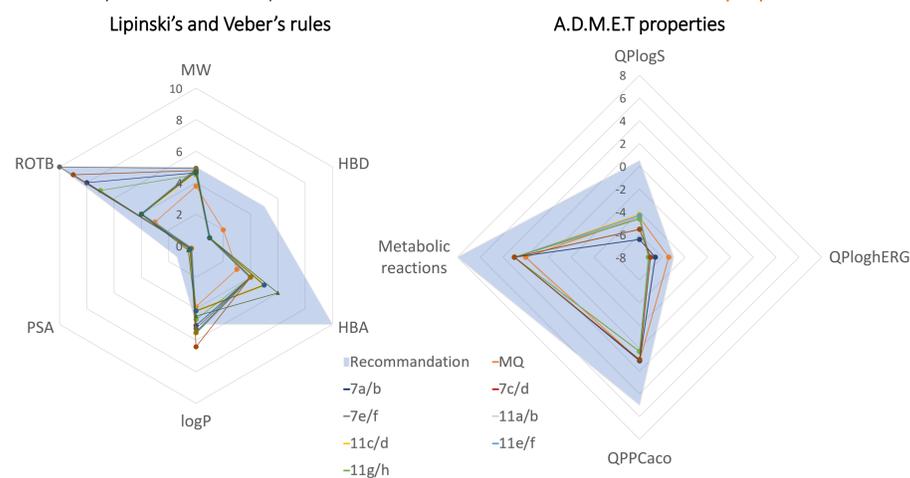


Figure 3: Radar chart of *in silico* physicochemical and pharmacokinetic properties of aryl and alkyl series.

→ Aryl and alkyl series comply with Lipinski's and Weber's rules  
→ Aryl and alkyl series could be potential drug candidates

## Conclusion and perspectives

**Fourteen new AAQ** in 6 to 32 % of global yields and 86 to 94 % of ee were obtained by a **short asymmetric synthesis** (5 or 7 steps).

While aryl series **7** was inactive *in vitro* against *M. abs S* and *R*, alkyl AAQ **11** displayed **anti-NTM activity** from 4 to 64 µg/mL.

SI values for alkyl AAQ **11** highlighted **eight compounds AAQ 11** safer than MQ (calculated according their cytotoxicity on HepG2). The enantiomer pair **11g/11h** is the **hit** of new series.

In the future, *in vitro* **antimycobacterial activity** on slowly growing rate NTM such as MAC or *M. xenopi*, and **cytotoxicity** on THP-1 cell lines for alkyl AAQ will be evaluated. In order to explore **new SAR**, further modulations focusing on the MQ core are planned. A few compounds with the highest SI will be selected for ***in vivo* evaluation on mouse model**.

## Biological results

The **efficacy** of final compounds **7a-f** and **11a-h** was tested on *M. abs smooth* (*S*) and *rough* (*R*) phenotypes. While **aryl compounds were inactive**, **alkyl series displayed moderate activity** (Table 2). **AAQ 11g/h** with a long alkyl chain (n = 6) were more efficient than **11a-d** (n = 3 or 4). No difference in activity was observed between *S* and *R* enantiomers of the same pair. **Cytotoxicity assays** on HepG2 cell lines revealed that **new series 7 and 11 are less toxic than MQ** (IC<sub>50</sub> = 26.2-96.3 vs 1.5 µg/mL). **SI of alkyl AAQ** are equal or **better** than those of MQ (SI = 0.4-5.0 vs 0.1-0.4).

Table 2: *In vitro* antimycobacterial activity and cytotoxicity of aryl and alkyl series 7 and 11.

Compound	X <sub>1,2,3</sub>	n	MIC (µg/mL) <sup>a</sup>		IC <sub>50</sub> (µg/mL) <sup>b</sup>	SI <sup>c</sup>	
			<i>M. abs S</i>	<i>M. abs R</i>		<i>M. abs S</i>	<i>M. abs R</i>
7a (R)	X <sub>1,3</sub> = CH, X <sub>2</sub> = C-Me	-	ND <sup>d</sup>	ND <sup>d</sup>	92.3 ± 4.3	ND	ND
7b (S)		-	ND <sup>d</sup>	ND <sup>d</sup>	51.1 ± 0.1	ND	ND
7c (R)	X <sub>1</sub> = N, X <sub>2,3</sub> = CH	-	ND <sup>d</sup>	ND <sup>d</sup>	48.6 ± 2.5	ND	ND
7d (S)		-	> 128	> 128	39.9 ± 3.5	< 0.2	< 0.2
7e (R)	X <sub>1,3</sub> = N, X <sub>2</sub> = CH	-	ND <sup>d</sup>	ND <sup>d</sup>	50.6 ± 0.7	ND	ND
7f (S)		-	> 128	> 128	95.7 ± 1.0	< 0.1	< 0.1
11a (R)	-	3	64	32	58.6 ± 4.6	0.8	0.8
11b (S)	-	3	64	32-64	60.1 ± 6.3	0.4	0.4-0.8
11c (R)	-	4	32	32-64	70.6 ± 6.5	1.0	1.0-2.0
11d (S)	-	4	32	32	NA <sup>e</sup>	0.6	0.6
11e (R)	-	5	16	8-16	96.3 ± 1.0	2.3	1.2-2.3
11f (S)	-	5	8-16	16	26.2 ± 0.7	0.8	0.8
11g (R)	-	6	8	4	41.0 ± 3.7	5.0	1.3-5.0
11h (S)	-	6	4-16	4	33.5 ± 5.8	4.1	2.1-4.1
MQ	-	-	32	16-32	1.5 ± 0.1	0.1	0.1
Clarithromycin	-	-	≥ 32	≥ 32	ND <sup>d</sup>	ND <sup>d</sup>	ND <sup>d</sup>
Ciprofloxacin	-	-	4	4	ND <sup>d</sup>	ND <sup>d</sup>	ND <sup>d</sup>
Amikacin	-	-	≤ 2	≤ 2	ND <sup>d</sup>	ND <sup>d</sup>	ND <sup>d</sup>

<sup>a</sup> MIC *M. abs S* and *R* DSM 44196, MIC were determined with technical and biological duplicates,

<sup>b</sup> IC<sub>50</sub> results are expressed as mean ± SD of duplicate experiments, <sup>c</sup> SI = IC<sub>50</sub>/MIC on *M. abs S* or *R* strains,

<sup>d</sup> ND not determined, <sup>e</sup> NA not applicable.

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