

Supporting Information

A hydrazine-hydrazone adamantine compound that target MmpL3 shows antimycobacterial activity

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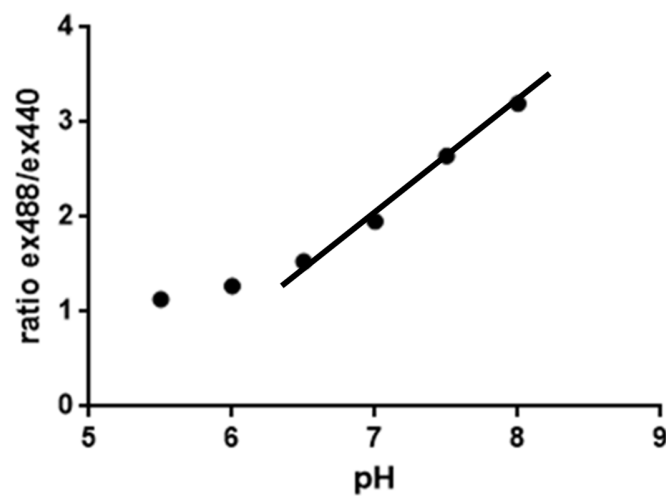


Figure S1. Intracellular calibration of the pH-sensitive dye BCECF-AN. Measurements of the fluorescence ratio (λ_{ex} 488 nm/ λ_{ex} 440 nm) of BCECF in *M. bovis* BCG were averaged (triplicates). pH calibration curve was sigmoidal with linear range in the physiological pH range of 6,5 - 8.

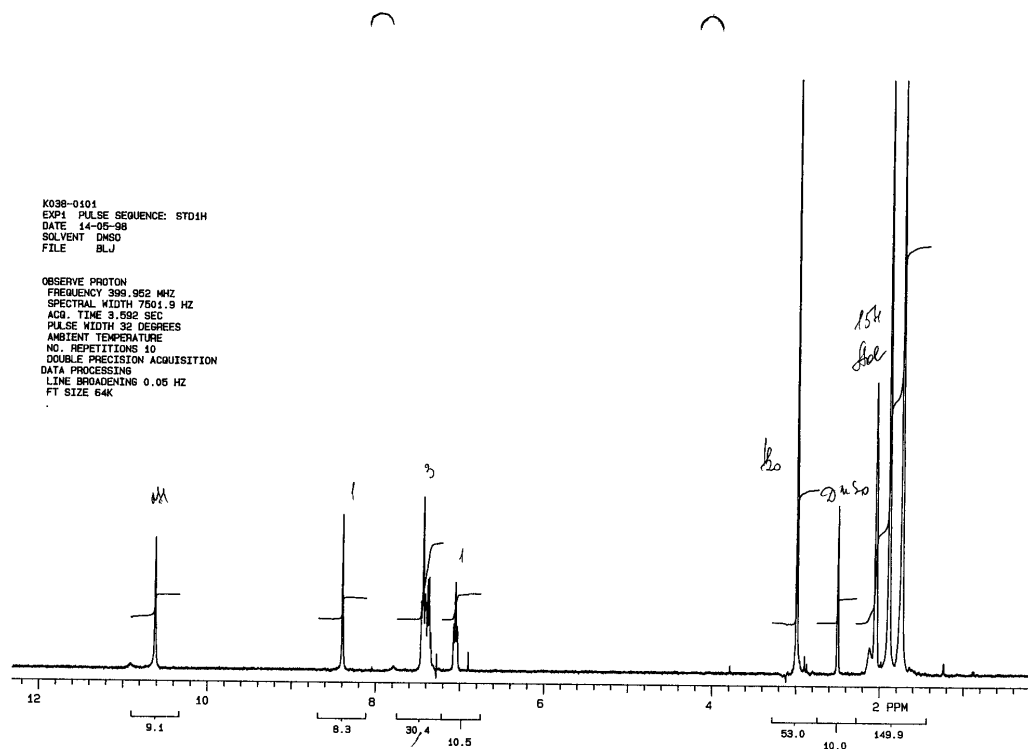


Figure S2. ^1H -NMR spectrum of compound 1 (400 MHz).

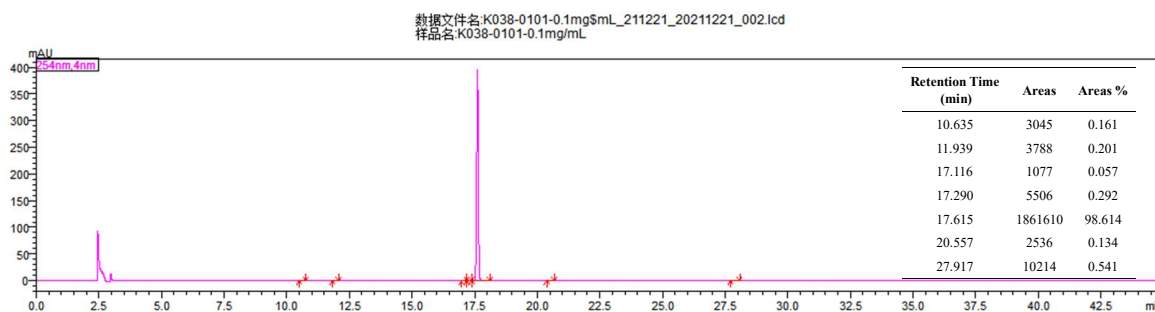


Figure S3. HPLC trace of compound 1. The purity of compound 1 is 98.6%. HPLC conditions: Column: Phenomenex Kinetex XB-C18, 5 μ m, 100 Å, 4.6*250 mm. Mobile A: acetonitrile, mobile B: 2 mM ammonium formate; gradient: 0-10 min with 10-60 % A, 10-30 min with 60-95 % A, 30-40 min with 95 % A, 40-40.1 min with 95-10 % A, 40.1-45 min with 10 % A. Flow rate: 1 mL/min; column temperature: 35°C; detector: 254 nm; injection volume: 10 μ L. Sample: 0.1 mg/mL in DMSO.

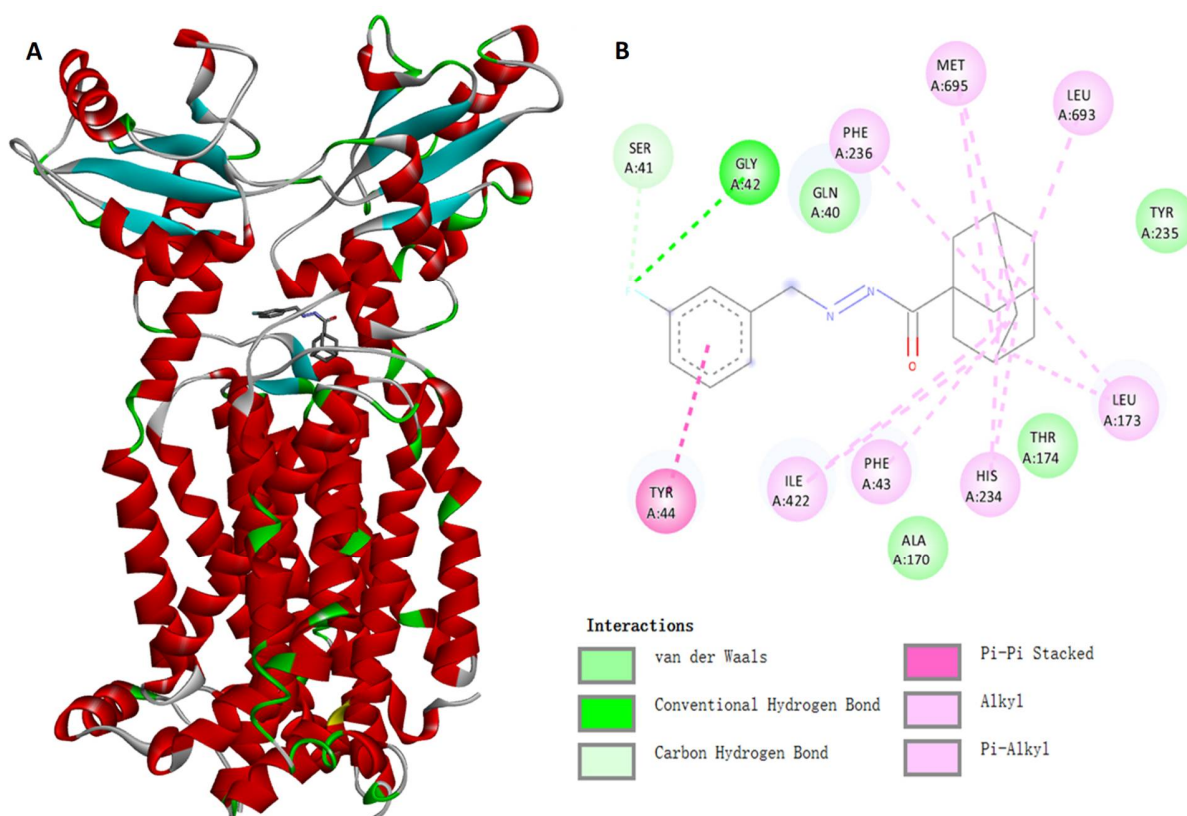


Figure S4. The docking site of compound 1 with *M. tuberculosis* MmpL3. A. Representation of an X-ray structure of the *M. tuberculosis* MmpL3 (PDB ID: 7NVH) with docked ligand compound 1 in the cavity of the protein. Compound 1 is depicted in stick representation in grey. B. 2D interaction diagram for the complex MmpL3-compound 1 drawn using Discovery Studio Visualizer.

Clinical strain	Mutation in <i>rpoB</i>	
	Base	Amino Acid
M9	1590C>T;1714A>G	Ser531Leu; Ile572Val
M20	1590C>T	Ser531Leu
M34	1598T>C	Leu533Pro

Table S1. Mutation in *rpoB* in clinical strains Rif-R used in the study.

Primers	5' to 3' Sequence
MSMEG_0250-F	ccttggagtttgtcgatc
MSMEG_0250-R	cggtaggcattcctcaaca

Table S2. Primers used in this study.

Compound	SMILE	Molecular Formula	Molecular Weight
Compound 1	<chem>Fc1cccc(c1)/C=N/NC(=O)C12CC3CC(C1)CC(C2)C3</chem>	C ₁₈ H ₂₁ FN ₂ O	300.4

Table S3. Molecular formula strings.