Supporting Information

Anti-Biofilm Agents against *Pseudomonas aeruginosa*: a Structure-Activity Relationship Study of *C*-Glycosidic LecB Inhibitors

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6.6 6.4 6.2 6.0 5.8 5.6 5.4 5.2 5.0 4.8 4.6 4.4 4.2 4.0 3.8 3.6 3.4 3.2 3.0 2.8 2.6 2.4 2.2 2.0 1.8 1.6 1.4 1.2 1.0 0.8 0.(f1 (ppm)



















7.2 7.0 6.8 6.6 6.4 6.2 6.0 5.8 5.6 5.4 5.2 5.0 4.8 4.6 4.4 4.2 4.0 3.8 3.6 3.4 3.2 3.0 2.8 2.6 2.4 2.2 2.0 1.8 1.6 1.4 1.2 1.0 0.8 f1 (ppm)







S20









7.4 7.2 7.0 6.8 6.6 6.4 6.2 6.0 5.8 5.6 5.4 5.2 5.0 4.8 4.6 4.4 4.2 4.0 3.8 3.6 3.4 3.2 3.0 2.8 2.6 2.4 2.2 2.0 1.8 1.6 1.4 1.2 1.0 f1 (ppm)





$^1\text{H},\,^{13}\text{C}$ and ^{19}F spectra of new compounds









$^1\text{H},\,^{13}\text{C}$ and ^{19}F spectra of new compounds









S32



	LecB рао1 64.3 µМ	LecB рао1 60 µМ	LecB _{РАО1} 60 µМ
	22 1.0 mM	22 750 μΜ	22 600 μΜ
	TBS supplemented with 1 mM CaCl ₂	TBS supplemented with 1 mM CaCl ₂	TBS supplemented with 1 mM CaCl ₂
	Time (min) 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	Image: space of the s	Tine (mi) 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
Ka [1/M]	9.07E+05	7.14E+05	9.15E+05
$\triangle H$ [cal/mol]	-10980	-11420	-11520
$\triangle s$ [cal/molK]	-9.53	-11.5	-11.4
Kd [M]	1.1E-06	1.4E-06	1.09E-06
Ν	1.02	0.96	0.974
$\triangle G$ [cal/mol]	-8140	-7990	-8120

Table S1: ITC measurements with 22



	LecB _{PA14} in complex with 22			
Data collection				
Beamline			ID23-2	
Wavelength	0.87260			
Space group			P61	
Unit cell dimensions		49.88 4	9.88 288.75 90.00 90.00	90.00
Resolution (Å)			37 07-1 45 (1 47-1 45)	
Nb reflections			808656	
Nb uniques reflections			71656	
R _{merge}			0.057 (0.308)	
R_{pim}			0.026 (0.141)	
Mean $I / \sigma I$			27.1 (7.6)	
Completeness (%)			100.00 (100.00)	
Redundancy			11.3 (11.0)	
CC1/2			100.0 (98.0)	
Refinement				
Resolution (Å)			37.07-1.45	
No. reflections/No. free reflections			67895/3598	
Rweet / Roma			12 4 / 14 3	
\mathbf{R} m s Bond lengths (Å)			0.016	
K.m.s bond lenguis (A)			0.010	
Rmsd Bond angles (°)	2.043			
Rmsd Chiral (°)	0 103			
No. atoms (Chain)	А	В	С	D
Protein	862	883	853	868
Sugar	22	22	22	22
Calcium	2	2	2	2
Waters	150	175	166	169
B-factors (Å ²)				
Protein	10.5	10.0	10.4	9.6
Sugar	15.3	10.3	12.2	10.8
Calcium	8.6	7.4	8.3	7.0
Waters	24.0	25.8	24.7	24.6
Ramachandran				
Allowed	99.8			
Favored	97.1			
Outliers	1			
PDB Code			5MAZ	

Table S2: Data collection and refinement statistics for $LecB_{PA14}$ in complex with 22

name	start-stop [m/z]
naproxen	230-232
glipizide	444.5-447
procaine	236-237.5
procainamide	235.5-236.6
propoxycaine	294.4-295.8
17	320-321.8
22	349.8-351
23	335.8-337
27	389-390.4
29	421-423.1

Table S3: m/z search window for plasma stability assay

Table S4: Mass transitions of compound
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compound	Q1 mass	Q3 mass	Time [msec]	DP [volts]	CE [volts]	CXP [volts]
6	357.966	183.0	30	-140	-34	-19
	357.966	197.9	30	-140	-36	-23
7	321.888	146.8	30	-140	-30	-15
	321.888	230.0	30	-140	-28	-21
17	320.91	238.2	30	-30	-18	-13
	320.91	82.0	30	-30	-20	-9
	320.91	40.0	30	-30	-66	-17
22	349.98	174.6	30	-90	-30	-7
	349.98	110.9	30	-90	-38	-11
23	335.899	175.9	30	-115	-32	-17
	335.899	160.9	30	-115	-30	-15
29	421.93	182.9	30	-135	-40	-19
	421.93	198.8	30	-135	-34	-17
3	374.003	198.0	30	-135	-34	-19
	374.003	182.9	30	-135	-32	-17
45	337.881	161.8	30	-130	-34	-17
	337.881	146.8	30	-130	-30	-15
Naproxene	231.106	185.1	50	80	19	10
	231.106	170.2	50	80	33	12
Caffeine	195.024	138.0	30	80	25	14
	195.024	110.0	30	80	31	18



Figure S1: Rationale for the extension of the thiophene moiety in 7 to target an additional patch or a subpocket on the surface of LecB



Figure S2: Purity of key compounds 17, 22, 23, 27, 29 by HPLC-UV