Supplementary Material

Anti-virulence *C*-mannosides as antibiotic sparing, oral therapeutics for urinary tract infections

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1. Prophylactic model of acute UTI with mannosides and prodrugs of mannoside 23 dosed at 25 mg/kg PO, 30 minutes prior to infection.



Figure S1. Bladder titers of 23 (269), 23a (Prodrug 269), 23b (Prodrug 1233), and 23c (Prodrug 6123) following a 25 mg/kg PO dose in mice. Also shown on right is 23, 23a, 21R (240), and 25 (254).



2. Pharmacodynamics of 24 after oral dosing in chronic UTI model.

3. Rat pharmacokinetics of 21R, 23, and 25

The pharmacokinetic profile of FIM-4269 (23), FIM-5240 (21R), FIM-5254 (25), was evaluated after a single intravenous (IV) injection or oral (PO) gavage dose to male Sprague Dawley (SD) rats. The study was performed by Seventh Wave Laboratories LLC 743 Spirit 40 Park Drive, Suite 209, Chesterfield, MO 63005.

Eighteen male rats were divided into six groups (3 rats per group). On Day 1 of the study, rats were administered 23, 21R, and 25 as a single IV dose at 3 mg/kg via surgically implanted jugular catheters at a dose volume of 1 mL/kg (Groups 1-3) or a single PO dose of 10 mg/kg at a dose volume of 2 mL/kg (Groups 4-6). The dose volume was calculated using individual rat body weight taken prior to dosing on Day 1. There were no abnormal clinical signs observed in any rats during the study. Blood was collected from all rats, via the tail vein, into K₂EDTA tubes at the protocol stated time points; 5, 15, 30 minutes, 1, 2, 4, 8 and 24 hour post dose for Groups 1-3 (IV Groups) and at 15 and 30 minutes, and 1, 2, 4, 8 and 24 hr post dose for Groups 4-6 (PO Groups). Plasma was separated via refrigerated centrifugation and Bioanalysis and PK analysis was performed to evaluate the systemic exposure to 23, 21R, and 25. Using an internal standard, plasma and urine samples were analyzed for test article levels by LC-MS/MS with (+) ESI with Ion Spray detection using a Phenomenex Kinetex C18 (2.1 x 50 mm, 2.6 µm) reversed phase column with Mobile Phase, Aqueous: water containing 0.1% formic acid, Organic: acetonitrile containing 0.1% formic acid. The following MS/MS fragmentations [precursor mass/charge ratio (m/z)/product m/z] were found: compound 21R, 418.20/238.20 amu; compound 23, 404.20/242.20 amu; compound **25**, 414.20/252.20 amu.

Evidence of plasma exposure to FIM 4269, FIM-5240, FIM-5254, was observed in all FIM 4269, FIM-5240, FIM-5254-treated rats following a single IV and PO administration at 3 and 10 mg/kg, respectively. Exposure (as assessed by Cmax and AUCall) was the highest following IV administration for FIM-5240, followed by FIM-4269, and FIM-5254 with mean values of 3410 2440, 1590, 1360, and 337 ng/mL (Cmax) and 1440, 767, 521, 410, and 126 h*ng/mL (AUCall), respectively. FM-5240 had a moderate clearance in rat and the lowest of the three compounds at 34.9 mL/min/kg with the other compounds having markedly higher clearance values of 98.4 and 408 mL/min/kg for FIM-4269 and FIM-5254, respectively. All compounds had a short terminal half-life with values ranging from 0.28 to 0.49 hours following IV administration at 3 mg/kg.

Absorption (as assessed by Tmax) following PO administration at10 mg/kg was most rapid for FIM-5254, followed by FIM-5240, and FIM-4269 with mean Tmax values of 0.333, 0.667, and 1.42 hours, respectively. Exposure (as assessed by Cmax and AUCall) was the highest following PO administration for FIM-5240, followed by FIM-5254, and FIM-4269 with mean values of 18.6, 6.96, and 5.00 ng/mL (Cmax) and 93.7, 29.6, and 23.2 h*ng/mL (AUCall), respectively. The percent of the recovered dose in the urine was greatest for FIM-4269 followed by FIM5240, and FIM-5254 with mean values of 0.24, 0.19, and 0.13%, respectively. The low recovery of all five compounds in the urine over 24 hours suggests low renal clearance of unchanged parent compound following PO administration. Bioavailability for all compounds was low with the highest bioavailability observed for FIM-5254, followed by FIM-5240, and FIM-5240, and FIM-5254, it hughest bioavailability observed for FIM-5254, followed by FIM-5240, and FIM-4269 with the highest bioavailability observed for FIM-5254, followed by FIM-5240, and FIM-4269 with the highest bioavailability observed for FIM-5254, followed by FIM-5240, and FIM-4269 with the highest bioavailability observed for FIM-5254, followed by FIM-5240, and FIM-4269 with values of 7.05, 1.95, and 1.34, respectively.

⁷ mourn	Douto	Dose	Compound	Animal	C ₀	C _{max}	T _{max}	AUC _{all}	AUCINFobs	Clobs	Vss _{obs}	$HL_{\lambda z}$
nouh	Koute	(mg/kg)	Compound	ID	(ng/mL)	(ng/mL)	(h)	(h*ng/mL)	(h*ng/mL)	(mL/min/kg)	(L/kg)	(h)
				1M001	2040	1280	0.083	485	480	104	1.85	0.326
				1M002	3760	1980	0.083	622	619	80.8	0.965	0.301
			22	1M003	2920	1520	0.083	455	454	110	1.16	0.308
1	IV	3	23 (EIM 4260)	Ν	3	3	3	3	3	3	3	3
			(FINI-4209)	Mean	2910	1590	0.083	521	518	98.4	1.33	0.312
				SD	860	356	0.00	89.1	88.9	15.6	0.467	0.0129
				CV%	29.6	22.3	0.0	17.1	17.2	15.8	35.2	4.1
		3	21R (FIM-5240)	2M001	6520	4100	0.083	1480	1480	33.8	0.583	0.511
	IV			2M002	3520	2830	0.083	1470	1460	34.2	0.881	0.471
				2M003	4710	3310	0.083	1370	1370	36.6	0.759	0.482
2				Ν	3	3	3	3	3	3	3	3
				Mean	4920	3410	0.083	1440	1440	34.9	0.741	0.488
				SD	1510	641	0.00	60.9	60.6	1.51	0.150	0.0209
				CV%	30.7	18.8	0.0	4.2	4.2	4.3	20.2	4.3
				3M001	700	435	0.083	152	151	331	4.79	0.279
				3M002	382	269	0.083	108	107	467	8.37	0.278
		3	25	3M003	429	306	0.083	118	117	426	7.04	0.284
3	IV		25 (FIM-5254)	Ν	3	3	3	3	3	3	3	3
				Mean	504	337	0.083	126	125	408	6.73	0.280
				SD	172	87.1	0.00	23.0	23.0	69.7	1.81	0.0032
				CV%	34.1	25.9	0.0	18.3	18.4	17.1	26.9	1.1

Table S1. Individual Animal and Group Mean Pharmacokinetic Summary Data; IVAdministration

Figure S2. 23 (FIM-4269) Plasma Concentration versus Time—Individual Animal Comparison; Group 1



Figure S3. 21R (FIM-5240) Plasma Concentration versus Time—Individual Animal Comparison; Group 2



Figure S4. 25 (FIM-5254) Plasma Concentration versus Time—Individual Animal Comparison; Group 3



Table S2. Individual Animal and Group Mean Pharmacokinetic Summary Data; POAdministration

Group	Route	Dose	Compound	Animal	C_{max}	T _{max}	AUC _{all}	AUC _{INFobs}	$HL_{\lambda z}$	F
		(mg/kg)		ID	(ng/mL)	(h)	(h*ng/mL)	(h*ng/mL)	(h)	(fraction)
				6M001	4.41	2.00	34.6	NC	NC	NA
				6M002	6.53	2.00	22.1	NC	NC	NA
			22	6M003	4.05	0.250	12.8	16.0	2.85	NA
6	РО	10	(FIM-4269)	Ν	3	3	3	1	1	1
				Mean	5.00	1.42	23.2	16.0	2.85	0.0134
				SD	1.34	1.01	10.9	NA	NA	NA
				CV%	26.8	71.3	47.2	NA	NA	NA
			21R (FIM-5240)	7M001	23.8	0.500	83.6	75.0	3.56	NA
				7M002	15.5	1.00	117	103	3.22	NA
				7M003	16.4	0.500	80.9	74.0	3.81	NA
7	PO	10		Ν	3	3	3	3	3	1
				Mean	18.6	0.667	93.7	83.8	3.53	0.0195
				SD	4.55	0.289	19.8	16.2	0.298	NA
				CV%	24.5	43.3	21.2	19.3	8.4	NA

				8M001	6.41	0.250	23.3	NC	NC	NA
8	РО	10	25 (FIM-5254)	8M002	8.53	0.500	18.3	NC	NC	NA
				8M003	5.93	0.250	47.3	NC	NC	NA
				Ν	3	3	3	0	0	1
				Mean	6.96	0.333	29.6	NA	NA	0.0705
				SD	1.38	0.144	15.5	NA	NA	NA
				CV%	19.9	43.3	52.3	NA	NA	NA

Figure S5. 23 (FIM-4269) Plasma Concentration versus Time—Individual Animal Comparison; Group 6



Figure S6. 21R (FIM-5240) Plasma Concentration versus Time—Individual Animal Comparison; Group 7







All raw data, the Protocol, Protocol Amendments, supporting documents and any reports will be temporarily retained at Seventh Wave Laboratories LLC, 743 Spirit 40 Park Drive, Chesterfield, MO 63005 until transferred to the Sponsor or Sponsor's designee.

4. X-ray crystallization Studies of 23

A. Protein Crystallization. FimH lectin domain (FimH_L; residues 1-158) from E. coli strain J96 was expressed and purified as described previously and dialyzed into 10 mM MES pH 5.8, 50 mM NaCl. To generate crystals of FimH_L bound to 23, FimH_L was pre-incubated with 23 at final concentrations of 1.27 mM (approx. 21 mg/mL) and 1.7 μ M, respectively. 100 nL x 100 nL drops were set up using a TTP Labtech mosquito® crystal robot using 70 μ L mother liquor (0.2 M AmSO4, 20% PEG 3350.) Cuboidal crystals appeared 72 hours of incubation at 18 °C. After approximately two weeks, crystals were transferred to mother liquor supplemented with 20% v/v glycerol as cryoprotectant and flash frozen in liquid nitrogen prior to storage and data collection.

B. X-ray data collection and structure determination. Diffraction data were collected at 100 ° K. X-rays were generated by a Rigaku MicroMax 007 generator coupled to a Rayonix Marmux X-ray source (Evanston, IL.) Images were collected on a Mar345 image plate dector. Data were indexed and integrated in iMosflm and scaled by Scala. The space group was then determined by POINTLESS. PHases were obtained by molecular replacement using PHASER and a .pdb of a previously solved FimH_L structure (PDB ID 3MCY) stripped of all waters, additives, and ligands. The Fo-Fc map unambiguously indicated electron density corresponding to Mannoside 23 in two distinct binding conformations within the ligand binding pocket. Chemical restraints on 23 were generated by eLBOW and used in subsequent rounds of refinement with phenix.refine from the PHENIX package.

Data Collection	
Space Group	P 21 21 21
Cell dimensions	
<i>a, b, c</i> (Å)	48.97, 55.66, 60.89
α, β, γ (°)	90, 90, 90
Resolution	24.48 - 1.76
Completeness (%)	99.7 (24.48-1.76)
R_{merge} (%)	0.05
<i σ(i)=""></i>	4.23 (at 1.76)
Total/Unique Reflections	248304/17107

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Refinement	
R _{work} /R _{free}	0.1352 / 0.1718
No. atoms (non-H)	
Protein	1209
Ligand	58
Water	237
Wilson B-Factor (Å ²)	15.2
R.M.S. Deviations	
Bond Lengths (Å)	0.09
Bond Angles (°)	1.410

5. X-Ray Experimental, and Projection View of Compound 27.

Crystals of appropriate dimension were obtained from Et₂O/Hexanes. A crystal with approximate dimensions $0.52 \times 0.46 \times 0.37 \text{ mm}^3$ was mounted on a MiteGen cryoloop in random orientation. Preliminary examination and data collection were performed using a Bruker X8 Kappa Apex II Charge Coupled Device (CCD) Detector system single crystal X-Ray diffractometer equipped with an Oxford Cryostream LT device. All data were collected using graphite monochromated Mo K α radiation (λ = 0.71073 Å) from a fine focus sealed tube X-Ray source. Preliminary unit cell constants were determined with a set of 36 narrow frame scans. Typical data sets consist of combinations of ϖ and φ scan frames with scan width of 0.5° and counting time of 15 seconds/frame at a crystal to detector distance of 4.0 cm. The collected frames were integrated using an orientation matrix determined from the narrow frame scans. Apex II and SAINT software packagesⁱ were used for data collection and data integration. Analysis of the integrated data did not show any decay. Final cell constants were determined by global refinement of reflections harvested from the complete data set. Collected data were corrected for systematic errors using SADABSⁱⁱ based on the Laue symmetry using equivalent reflections.



Figure S8. Compound **27** Projection view with 50% probability ellipsoids. Structure solution and refinement were carried out using the SHELXTL- PLUS software packageⁱⁱⁱ. The structure was solved by direct methods and refined successfully in the space group, P1. Full matrix least-squares refinements was carried out by minimizing $\Sigma w (F_o^2 - F_c^2)^2$. The non-hydrogen atoms were refined anisotropically to convergence. All hydrogen atoms were treated using appropriate riding model (AFIX m3). Absolute structure was determined with the Flack x parameter refining to -0.011(1) for 6453 selected quotients using Parson's method^{iv} Complete listings of positional and isotropic displacement parameters for hydrogen atoms, anisotropic displacement parameters for the non-hydrogen atoms are listed below (**Tables S5-S9**). Table of calculated and observed structure factors are available in electronic format.

Identification code	j25315/lt/x8/Janetka-Mydock/Fimbrion			
Empirical formula	C ₂₄ H ₂₉ Br O ₁₁			
Formula weight	573.38			
Temperature	100(2) K			
Wavelength	0.71073 Å			
Crystal system	Triclinic			
Space group	P1			
Unit cell dimensions	a = 8.1051(4) Å	α= 82.568(3)°.		
	b = 8.9473(5) Å	β= 79.666(3)°.		
	c = 9.4597(5) Å	$\gamma = 88.134(2)^{\circ}$.		
Volume	669.18(6) Å ³			
Z	1			
Density (calculated)	1.423 Mg/m ³			
Absorption coefficient	1.592 mm ⁻¹			
F(000)	296			
Crystal size	0.519 x 0.463 x 0.365 mm ³			
Theta range for data collection	2.206 to 39.500°.			
Index ranges	-14≤h≤14, -16≤k≤16, -16≤l≤16			
Reflections collected	59865			
Independent reflections	15286 [R(int) = 0.022]			
Completeness to theta = 25.242°	100.0 %			
Absorption correction	Semi-empirical from equivalents			
Max. and min. transmission	0.6514 and 0.6044			
Refinement method	Full-matrix least-squares on F ²			
Data / restraints / parameters	15286 / 4 / 340			
Goodness-of-fit on F ²	1.054			
Final R indices [I>2sigma(I)]	R1 = 0.0240, wR2 = 0.0564			
R indices (all data)	R1 = 0.0269, wR2 = 0.0573			
Absolute structure parameter	-0.0108(10)			
Largest diff. peak and hole	0.868 and -0.255 e.Å ⁻³			

Table S4. Crystal data and structure refinement for jj25315 (27).

	Х	У	Z	U(eq)
Br(1)	360(1)	3108(1)	413(1)	22(1)
O(1)	3160(1)	4929(1)	5885(1)	14(1)
O(2)	4149(1)	7059(1)	7644(1)	15(1)
O(3)	5720(1)	9102(1)	7614(1)	25(1)
O(4)	7354(1)	5870(1)	7477(1)	16(1)
O(5)	9062(1)	7335(1)	5744(1)	24(1)
O(6)	6921(1)	2963(1)	6793(1)	18(1)
O(7)	6165(2)	2055(1)	9141(1)	30(1)
O(8)	2666(1)	2051(1)	5175(1)	21(1)
O(9)	2256(1)	-335(1)	6196(1)	21(1)
O(10)	4688(1)	8165(1)	3290(1)	17(1)
O(11)	7494(7)	7919(8)	2500(30)	39(2)
O(11')	7280(30)	7997(17)	2090(30)	42(3)
C(1)	3807(1)	6395(1)	5352(1)	13(1)
C(2)	5070(1)	6835(1)	6238(1)	13(1)
C(2")	3403(2)	8516(2)	9562(1)	26(1)
C(2')	4562(1)	8291(1)	8189(1)	17(1)
C(3)	6388(1)	5593(1)	6408(1)	14(1)
C(3")	9632(2)	6964(2)	8176(2)	26(1)
C(3')	8700(1)	6780(1)	6990(1)	17(1)
C(4)	5589(1)	4065(1)	6898(1)	13(1)
C(4")	8677(2)	1126(2)	7758(2)	28(1)
C(4')	7117(1)	2066(1)	8017(1)	18(1)
C(5)	4413(1)	3776(1)	5872(1)	14(1)
C(6)	3531(1)	2292(1)	6329(1)	17(1)
C(6")	1184(2)	512(2)	4014(2)	27(1)
C(6')	2071(1)	653(1)	5246(1)	17(1)
C(7)	4577(1)	6547(1)	3729(1)	13(1)
C(7")	6188(2)	10410(2)	2473(2)	32(1)
C(7')	6216(2)	8722(1)	2670(1)	25(1)
C(8)	3560(1)	5793(1)	2845(1)	13(1)

Table S5. Atomic coordinates (x 10⁴) and equivalent isotropic displacement parameters ($Å^2x$ 10³) for jj25315. U(eq) is defined as one third of the trace of the orthogonalized U^{ij} tensor.

C(9)	4344(1)	4819(1)	1877(1)	14(1)
C(10)	6212(1)	4568(2)	1575(1)	21(1)
C(11)	3356(1)	4040(1)	1154(1)	15(1)
C(12)	1636(1)	4259(1)	1374(1)	16(1)
C(13)	845(1)	5263(1)	2286(1)	18(1)
C(14)	1825(1)	6013(1)	3026(1)	16(1)

Br(1)-C(12)	1.8992(10)
O(1)-C(5)	1.4233(12)
O(1)-C(1)	1.4243(11)
O(2)-C(2')	1.3507(13)
O(2)-C(2)	1.4386(12)
O(3)-C(2')	1.2094(15)
O(4)-C(3')	1.3582(13)
O(4)-C(3)	1.4344(12)
O(5)-C(3')	1.2068(14)
O(6)-C(4')	1.3515(14)
O(6)-C(4)	1.4369(12)
O(7)-C(4')	1.1954(16)
O(8)-C(6')	1.3436(12)
O(8)-C(6)	1.4411(13)
O(9)-C(6')	1.2027(13)
O(10)-C(7')	1.3532(15)
O(10)-C(7)	1.4541(12)
O(11)-C(7')	1.239(7)
O(11')-C(7')	1.160(15)
C(1)-C(2)	1.5249(13)
C(1)-C(7)	1.5412(13)
C(1)-H(1)	1.0000
C(2)-C(3)	1.5300(14)
C(2)-H(2)	1.0000
C(2")-C(2')	1.4922(17)
C(2")-H(2"A)	0.9800
C(2")-H(2"B)	0.9800
C(2")-H(2"C)	0.9800
C(3)-C(4)	1.5126(13)
C(3)-H(3)	1.0000
C(3")-C(3')	1.4888(16)
C(3")-H(3"A)	0.9800
C(3")-H(3"B)	0.9800
C(3")-H(3"C)	0.9800

Table S6. Bond lengths [Å] and angles [°] for jj25315.

C(4)-C(5)	1.5263(14)
C(4)-H(4)	1.0000
C(4")-C(4')	1.4969(16)
C(4")-H(4"1)	0.9800
C(4")-H(4"2)	0.9800
C(4")-H(4"3)	0.9800
C(5)-C(6)	1.5044(13)
C(5)-H(5)	1.0000
C(6)-H(6A)	0.9900
C(6)-H(6B)	0.9900
C(6")-C(6')	1.4951(17)
C(6")-H(6"A)	0.9800
C(6")-H(6"B)	0.9800
C(6")-H(6"C)	0.9800
C(7)-C(8)	1.5047(13)
C(7)-H(7)	1.0000
C(7")-C(7')	1.4979(19)
C(7")-H(7"A)	0.9800
C(7")-H(7"B)	0.9800
C(7")-H(7"C)	0.9800
C(8)-C(14)	1.3965(14)
C(8)-C(9)	1.4010(14)
C(9)-C(11)	1.3968(14)
C(9)-C(10)	1.5049(15)
C(10)-H(10A)	0.9800
C(10)-H(10B)	0.9800
С(10)-Н(10С)	0.9800
C(11)-C(12)	1.3829(14)
С(11)-Н(11)	0.9500
C(12)-C(13)	1.3879(15)
C(13)-C(14)	1.3877(15)
С(13)-Н(13)	0.9500
C(14)-H(14)	0.9500
C(5)-O(1)-C(1)	113.75(7)
C(2')-O(2)-C(2)	115.78(8)

C(3')-O(4)-C(3)	115.71(8)
C(4')-O(6)-C(4)	117.58(8)
C(6')-O(8)-C(6)	115.27(8)
C(7')-O(10)-C(7)	117.15(9)
O(1)-C(1)-C(2)	111.17(7)
O(1)-C(1)-C(7)	112.46(7)
C(2)-C(1)-C(7)	110.78(8)
O(1)-C(1)-H(1)	107.4
C(2)-C(1)-H(1)	107.4
C(7)-C(1)-H(1)	107.4
O(2)-C(2)-C(1)	107.17(8)
O(2)-C(2)-C(3)	109.27(7)
C(1)-C(2)-C(3)	111.23(8)
O(2)-C(2)-H(2)	109.7
C(1)-C(2)-H(2)	109.7
C(3)-C(2)-H(2)	109.7
C(2')-C(2")-H(2"A)	109.5
C(2')-C(2")-H(2"B)	109.5
H(2"A)-C(2")-H(2"B)	109.5
C(2')-C(2")-H(2"C)	109.5
H(2"A)-C(2")-H(2"C)	109.5
H(2"B)-C(2")-H(2"C)	109.5
O(3)-C(2')-O(2)	123.43(10)
O(3)-C(2')-C(2")	125.18(11)
O(2)-C(2')-C(2")	111.38(10)
O(4)-C(3)-C(4)	107.00(8)
O(4)-C(3)-C(2)	110.83(8)
C(4)-C(3)-C(2)	111.64(8)
O(4)-C(3)-H(3)	109.1
C(4)-C(3)-H(3)	109.1
C(2)-C(3)-H(3)	109.1
C(3')-C(3")-H(3"A)	109.5
C(3')-C(3")-H(3"B)	109.5
H(3"A)-C(3")-H(3"B)	109.5
C(3')-C(3")-H(3"C)	109.5
H(3"A)-C(3")-H(3"C)	109.5

H(3"B)-C(3")-H(3"C)	109.5
O(5)-C(3')-O(4)	122.66(10)
O(5)-C(3')-C(3")	126.09(10)
O(4)-C(3')-C(3")	111.24(9)
O(6)-C(4)-C(3)	107.02(8)
O(6)-C(4)-C(5)	108.63(8)
C(3)-C(4)-C(5)	109.20(8)
O(6)-C(4)-H(4)	110.6
C(3)-C(4)-H(4)	110.6
C(5)-C(4)-H(4)	110.6
C(4')-C(4")-H(4"1)	109.5
C(4')-C(4")-H(4"2)	109.5
H(4"1)-C(4")-H(4"2)	109.5
C(4')-C(4")-H(4"3)	109.5
H(4"1)-C(4")-H(4"3)	109.5
H(4"2)-C(4")-H(4"3)	109.5
O(7)-C(4')-O(6)	123.62(10)
O(7)-C(4')-C(4")	125.84(11)
O(6)-C(4')-C(4")	110.54(11)
O(1)-C(5)-C(6)	107.48(8)
O(1)-C(5)-C(4)	109.11(8)
C(6)-C(5)-C(4)	111.88(8)
O(1)-C(5)-H(5)	109.4
C(6)-C(5)-H(5)	109.4
C(4)-C(5)-H(5)	109.4
O(8)-C(6)-C(5)	106.42(8)
O(8)-C(6)-H(6A)	110.4
C(5)-C(6)-H(6A)	110.4
O(8)-C(6)-H(6B)	110.4
C(5)-C(6)-H(6B)	110.4
H(6A)-C(6)-H(6B)	108.6
C(6')-C(6")-H(6"A)	109.5
C(6')-C(6")-H(6"B)	109.5
H(6"A)-C(6")-H(6"B)	109.5
C(6')-C(6")-H(6"C)	109.5
H(6"A)-C(6")-H(6"C)	109.5

H(6"B)-C(6")-H(6"C)	109.5
O(9)-C(6')-O(8)	122.70(10)
O(9)-C(6')-C(6")	125.81(10)
O(8)-C(6')-C(6")	111.49(9)
O(10)-C(7)-C(8)	111.95(8)
O(10)-C(7)-C(1)	104.30(7)
C(8)-C(7)-C(1)	113.19(8)
O(10)-C(7)-H(7)	109.1
C(8)-C(7)-H(7)	109.1
C(1)-C(7)-H(7)	109.1
C(7')-C(7")-H(7"A)	109.5
C(7')-C(7")-H(7"B)	109.5
H(7"A)-C(7")-H(7"B)	109.5
C(7')-C(7")-H(7"C)	109.5
H(7"A)-C(7")-H(7"C)	109.5
H(7"B)-C(7")-H(7"C)	109.5
O(11')-C(7')-O(10)	122.9(5)
O(11)-C(7')-O(10)	122.7(3)
O(11')-C(7')-C(7")	124.0(5)
O(11)-C(7')-C(7")	125.4(3)
O(10)-C(7')-C(7")	111.08(12)
C(14)-C(8)-C(9)	119.72(9)
C(14)-C(8)-C(7)	120.25(8)
C(9)-C(8)-C(7)	119.98(8)
C(11)-C(9)-C(8)	118.93(9)
C(11)-C(9)-C(10)	118.04(9)
C(8)-C(9)-C(10)	123.03(9)
C(9)-C(10)-H(10A)	109.5
C(9)-C(10)-H(10B)	109.5
H(10A)-C(10)-H(10B)	109.5
С(9)-С(10)-Н(10С)	109.5
H(10A)-C(10)-H(10C)	109.5
H(10B)-C(10)-H(10C)	109.5
C(12)-C(11)-C(9)	120.06(9)
C(12)-C(11)-H(11)	120.0
C(9)-C(11)-H(11)	120.0

C(11)-C(12)-C(13)	121.79(9)
C(11)-C(12)-Br(1)	117.87(8)
C(13)-C(12)-Br(1)	120.33(8)
C(14)-C(13)-C(12)	118.03(9)
С(14)-С(13)-Н(13)	121.0
С(12)-С(13)-Н(13)	121.0
C(13)-C(14)-C(8)	121.39(9)
C(13)-C(14)-H(14)	119.3
C(8)-C(14)-H(14)	119.3

Symmetry transformations used to generate equivalent atoms:

	U ¹¹	U ²²	U ³³	U ²³	U ¹³	U ¹²
Br(1)	20(1)	25(1)	24(1)	-9(1)	-8(1)	-3(1)
O(1)	15(1)	10(1)	15(1)	0(1)	-2(1)	-3(1)
O(2)	19(1)	13(1)	11(1)	-3(1)	-2(1)	-2(1)
O(3)	26(1)	20(1)	31(1)	-10(1)	-7(1)	-6(1)
O(4)	16(1)	17(1)	16(1)	0(1)	-4(1)	-5(1)
O(5)	23(1)	27(1)	20(1)	-1(1)	0(1)	-11(1)
O(6)	16(1)	16(1)	20(1)	-3(1)	1(1)	3(1)
O(7)	48(1)	26(1)	15(1)	-5(1)	-6(1)	19(1)
O(8)	30(1)	12(1)	22(1)	1(1)	-11(1)	-8(1)
O(9)	26(1)	12(1)	27(1)	1(1)	-9(1)	-4(1)
O(10)	21(1)	13(1)	15(1)	2(1)	-2(1)	-5(1)
O(11)	21(1)	27(1)	63(5)	0(2)	7(2)	-7(1)
O(11')	37(4)	30(2)	50(5)	-5(3)	21(3)	-13(3)
C(1)	16(1)	10(1)	11(1)	-1(1)	-2(1)	-3(1)
C(2)	16(1)	12(1)	12(1)	-1(1)	-2(1)	-3(1)
C(2")	36(1)	27(1)	17(1)	-10(1)	-5(1)	4(1)
C(2')	22(1)	16(1)	17(1)	-6(1)	-8(1)	1(1)
C(3)	15(1)	14(1)	14(1)	-1(1)	-2(1)	-3(1)
C(3")	26(1)	26(1)	29(1)	3(1)	-14(1)	-9(1)
C(3')	15(1)	16(1)	21(1)	-2(1)	-3(1)	-4(1)
C(4)	14(1)	11(1)	15(1)	-1(1)	-1(1)	0(1)
C(4")	21(1)	22(1)	48(1)	-14(1)	-14(1)	9(1)
C(4')	21(1)	15(1)	22(1)	-8(1)	-10(1)	5(1)
C(5)	16(1)	11(1)	15(1)	-2(1)	-2(1)	-2(1)
C(6)	22(1)	12(1)	18(1)	1(1)	-6(1)	-6(1)
C(6")	37(1)	21(1)	28(1)	-1(1)	-15(1)	-9(1)
C(6')	19(1)	12(1)	21(1)	-2(1)	-4(1)	-4(1)
C(7)	16(1)	11(1)	12(1)	-1(1)	-2(1)	-2(1)
C(7")	43(1)	20(1)	31(1)	4(1)	-2(1)	-14(1)
C(7')	28(1)	19(1)	25(1)	-1(1)	4(1)	-10(1)
C(8)	14(1)	13(1)	11(1)	-3(1)	-1(1)	0(1)
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Table S7. Anisotropic displacement parameters (Å²x 10³) for jj25315. The anisotropic displacement factor exponent takes the form: $-2\pi^2$ [h² a^{*2}U¹¹ + ... + 2 h k a^{*} b^{*} U¹²]

C(9)	13(1)	16(1)	13(1)	-4(1)	-1(1)	0(1)
C(10)	13(1)	28(1)	24(1)	-13(1)	-1(1)	1(1)
C(11)	15(1)	17(1)	15(1)	-6(1)	-2(1)	0(1)
C(12)	15(1)	18(1)	15(1)	-4(1)	-4(1)	-1(1)
C(13)	14(1)	23(1)	19(1)	-8(1)	-3(1)	1(1)
C(14)	14(1)	19(1)	16(1)	-7(1)	-2(1)	1(1)

	Х	у	Z	U(eq)
H(1)	2845	7124	5471	15
H(2)	5633	7793	5758	16
H(2"A)	2372	9016	9334	39
H(2"B)	3948	9145	10115	39
H(2"C)	3129	7536	10137	39
H(3)	7155	5562	5458	17
H(3"A)	8832	7119	9055	39
H(3"B)	10369	7838	7887	39
H(3"C)	10306	6056	8368	39
H(4)	4966	4013	7916	16
H(4"1)	8795	477	8652	43
H(4"2)	9653	1785	7457	43
H(4"3)	8606	500	6995	43
H(5)	5067	3788	4869	17
H(6A)	4351	1469	6480	21
H(6B)	2724	2326	7244	21
H(6"A)	492	1414	3847	41
H(6"B)	468	-379	4250	41
H(6"C)	2011	409	3139	41
H(7)	5734	6101	3611	16
H(7"A)	5174	10772	3067	48
H(7"B)	7181	10782	2771	48
H(7"C)	6191	10782	1451	48
H(10A)	6777	5542	1318	32
H(10B)	6504	3964	771	32
H(10C)	6572	4033	2441	32
H(11)	3865	3358	510	19
H(13)	-332	5431	2401	22
H(14)	1304	6691	3669	20

Table S8. Hydrogen coordinates ($x \ 10^4$) and isotropic displacement parameters (Å²x 10³) for jj25315.

Table S9. Torsion angles [°] for jj25315.

C(5)-O(1)-C(1)-C(2)	-58.95(10)
C(5)-O(1)-C(1)-C(7)	65.93(10)
C(2')-O(2)-C(2)-C(1)	-136.07(9)
C(2')-O(2)-C(2)-C(3)	103.29(9)
O(1)-C(1)-C(2)-O(2)	-69.46(9)
C(7)-C(1)-C(2)-O(2)	164.73(7)
O(1)-C(1)-C(2)-C(3)	49.92(10)
C(7)-C(1)-C(2)-C(3)	-75.90(9)
C(2)-O(2)-C(2')-O(3)	-6.07(15)
C(2)-O(2)-C(2')-C(2'')	173.29(9)
C(3')-O(4)-C(3)-C(4)	152.01(9)
C(3')-O(4)-C(3)-C(2)	-86.06(10)
O(2)-C(2)-C(3)-O(4)	-50.01(10)
C(1)-C(2)-C(3)-O(4)	-168.12(7)
O(2)-C(2)-C(3)-C(4)	69.17(10)
C(1)-C(2)-C(3)-C(4)	-48.95(10)
C(3)-O(4)-C(3')-O(5)	0.53(16)
C(3)-O(4)-C(3')-C(3")	-178.75(10)
C(4')-O(6)-C(4)-C(3)	120.84(9)
C(4')-O(6)-C(4)-C(5)	-121.38(9)
O(4)-C(3)-C(4)-O(6)	-67.85(10)
C(2)-C(3)-C(4)-O(6)	170.74(8)
O(4)-C(3)-C(4)-C(5)	174.75(8)
C(2)-C(3)-C(4)-C(5)	53.33(10)
C(4)-O(6)-C(4')-O(7)	6.56(17)
C(4)-O(6)-C(4')-C(4'')	-172.87(9)
C(1)-O(1)-C(5)-C(6)	-174.86(8)
C(1)-O(1)-C(5)-C(4)	63.64(10)
O(6)-C(4)-C(5)-O(1)	-175.67(7)
C(3)-C(4)-C(5)-O(1)	-59.28(10)
O(6)-C(4)-C(5)-C(6)	65.53(10)
C(3)-C(4)-C(5)-C(6)	-178.08(8)
C(6')-O(8)-C(6)-C(5)	167.55(9)
O(1)-C(5)-C(6)-O(8)	68.60(10)

C(4)-C(5)-C(6)-O(8)	-171.65(8)
C(6)-O(8)-C(6')-O(9)	-0.66(17)
C(6)-O(8)-C(6')-C(6")	179.55(11)
C(7')-O(10)-C(7)-C(8)	-113.23(11)
C(7')-O(10)-C(7)-C(1)	124.04(10)
O(1)-C(1)-C(7)-O(10)	163.23(8)
C(2)-C(1)-C(7)-O(10)	-71.68(9)
O(1)-C(1)-C(7)-C(8)	41.31(11)
C(2)-C(1)-C(7)-C(8)	166.40(8)
C(7)-O(10)-C(7')-O(11')	24(2)
C(7)-O(10)-C(7')-O(11)	-1.8(14)
C(7)-O(10)-C(7')-C(7")	-171.87(10)
O(10)-C(7)-C(8)-C(14)	-70.87(12)
C(1)-C(7)-C(8)-C(14)	46.67(12)
O(10)-C(7)-C(8)-C(9)	111.62(10)
C(1)-C(7)-C(8)-C(9)	-130.85(9)
C(14)-C(8)-C(9)-C(11)	-2.56(14)
C(7)-C(8)-C(9)-C(11)	174.97(9)
C(14)-C(8)-C(9)-C(10)	177.51(10)
C(7)-C(8)-C(9)-C(10)	-4.96(15)
C(8)-C(9)-C(11)-C(12)	1.40(15)
C(10)-C(9)-C(11)-C(12)	-178.67(10)
C(9)-C(11)-C(12)-C(13)	1.10(16)
C(9)-C(11)-C(12)-Br(1)	-177.71(8)
C(11)-C(12)-C(13)-C(14)	-2.36(16)
Br(1)-C(12)-C(13)-C(14)	176.42(8)
C(12)-C(13)-C(14)-C(8)	1.15(16)
C(9)-C(8)-C(14)-C(13)	1.30(16)
C(7)-C(8)-C(14)-C(13)	-176.22(10)

Symmetry transformations used to generate equivalent atoms:

D-HA	d(D-H)	d(HA)	d(DA)	<(DHA)
C(1)-H(1)O(9)#1	1.00	2.47	3.2835(13)	138.6
C(3")-H(3"B)O(9)#2	0.98	2.49	3.4074(15)	156.6
C(3")-H(3"C)Br(1)#3	0.98	3.07	3.8929(13)	142.9
C(6)-H(6A)O(3)#4	0.99	2.56	3.5102(15)	160.8
C(6")-H(6"B)O(5)#5	0.98	2.51	3.4382(16)	157.8
C(7")-H(7"C)O(7)#6	0.98	2.33	3.3045(18)	170.7
C(11)-H(11)O(7)#7	0.95	2.43	3.3197(14)	156.3

Table S10. Hydrogen bonds for jj25315 [Å and °].

Symmetry transformations used to generate equivalent atoms:

#1 x,y+1,z #2 x+1,y+1,z #3 x+1,y,z+1 #4 x,y-1,z

#5 x-1,y-1,z #6 x,y+1,z-1 #7 x,y,z-1

6. ¹H and ¹³C NMR Spectra of New Compounds



Compound 3



Compound 5

















S35

Compound 13











Compound 21R









S42







Compound 28R



Compound 29R



Compound 30



Compound 31R



Compound 32R



Compound 33



References.

- ⁱ Bruker Analytical X-Ray, Madison, WI, **2010** ⁱⁱ Bruker Analytical X-Ray, Madison, WI, **2010** ⁱⁱⁱ Sheldrick, G.M. (**2008**). *Acta Cryst.* A64,112-122 ^{iv} Parsons, S. & Flack, H. (**2004**). Acta Cryst., A, s61. 122