Electronic Supplementary Material (ESI) for Organic & Biomolecular Chemistry. This journal is © The Royal Society of Chemistry 2017

Supporting Information

for

Zwitterionic pyrrolidine-phosphonates: Inhibitors of the glycoside hydrolase-like phosphorylase *Streptomyces coelicolor* GlgEI-V279S

Sri Kumar Veleti†‡, Cecile Petit†, Jared J. Lindenberger‡, Donald R. Ronning*

and Steven J. Sucheck*

Department of Chemistry and Biochemistry and School of Green Chemistry and Engineering,

The University of Toledo, 2801 W. Bancroft Street, Toledo, Ohio 43606, USA.

Steve.Sucheck@UToledo.edu

Table of	Contents

	Spectra
Table S1. Data collection and refinement statistics for <i>Sco</i> GlgEI V279S-5	S-3
and ScoGlgEI V279S-6 complex	
Dibenzyl vinylphosphonate (10)	S-4, 5, 6
Dibenzyl allylphosphonate (14)	S-7, 8, 9
Dibenzyl ((3-hydroxy-2,5-bis(hydroxymethyl)-(3-O-a-D-	S-10, 11, 12
glucopyranosyl)pyrrolidin-1-yl)methyl)phosphonate (16)	
((3-hydroxy-2,5-bis(hydroxymethyl)-(3-O-α-D-glucopyranosyl)pyrrolidin-	S-13, 14, 15
1-yl)methyl)phosphonic acid (5)	
Dibenzyl ((3-hydroxy-2,5-bis(hydroxymethyl)-(3-O-a-D-	S-16, 17, 18
glucopyranosyl)pyrrolidin-1-yl)ethyl)phosphonate (17)	
((3-Hydroxy-2,5-bis(hydroxymethyl)-(3-O-a-D-glucopyranosyl)pyrrolidin-	S-19, 20, 21
1-yl)ethyl)phosphonic acid (6)	

Table S1. Data collection and refinement statistics. The necessary data were obtained from one crystal. Values in parentheses are for the highest shells.

	Sco GlgEI-V279S-5 complex	<i>Sco</i> GlgEI-V279S-6 complex		
Data collection				
Wavelength (Å)				
Space group	P 41	P 4 ₁ 2 ₁ 2		
Cell dimensions				
a, b, c (Å)	113.3, 113.3 314.3	113.9, 113.9, 317.3		
α, β, γ	90 °, 90 °, 90 °	90 °, 90 °, 90 °		
Resolution (Å)	50.0 - 3.20	50.0 - 2.45		
R merge	12.4 (61.7)	17.2 (33.0)		
Ι/σΙ	11.5 (2.4)	14.8 (2.8)		
Completeness (%)	99.6 (99.8)	99.1 (95.1)		
Redundancy	4.3 (4.3)	12.7 (13.3		
Refinement				
Resolution (Å)	39.06 - 3.21	45.88 - 2.45		
No. unique reflections	64,431	76,413 (7,224)		
$R_{\text{work}}/R_{\text{free}}$ (%)	20.8/24.0	19.2/23.7		
No. atoms				
Protein	20,461	10,724		
Ligand/ion	108	64		
Water	57	359		
<i>B</i> -factors ($Å^2$)				
Protein	68.0	52.6		
Ligand/ion	70.5	66.0		
Water	50.8	47.1		
R.m.s deviations				
Bond lengths (Å)	0.005	0.008		
Bond angles (°)	1.047	1.18		
Ramachandran				
Favored (%)	97.98	97.38		
Outliers (%)	0.12	0.00		

¹H NMR of Dibenzyl vinylphosphonate (10)



¹³C NMR of Dibenzyl vinylphosphonate (10)



³¹P NMR of Dibenzyl vinylphosphonate (10)







³¹P NMR of Dibenzyl allylphosphonate (14)





¹H NMR of dibenzyl ((3-hydroxy-2,5-bis(hydroxymethyl)-(3-*O*-α-D-glucopyranosyl)pyrrolidin-1-yl)methyl)phosphonate (16):



¹³C NMR of dibenzyl ((3-hydroxy-2,5-bis(hydroxymethyl)-(3-*O*-α-D-glucopyranosyl)pyrrolidin-1-yl)methyl)phosphonate (16):

200 190 180 170 160 150 140 130 120 110 100 90 80 70 60 50 40 30 20 10 0 -10 ppm





¹H NMR of ((3-hydroxy-2,5-bis(hydroxymethyl)-(3-*O*-α-D-glucopyranosyl)pyrrolidin-1-yl)methyl)phosphonic acid (5)





³¹P NMR of ((3-hydroxy-2,5-bis(hydroxymethyl)-(3-*O*-α-D-glucopyranosyl)pyrrolidin-1-yl)methyl)phosphonic acid (5)





¹H NMR of dibenzyl ((3-hydroxy-2,5-bis(hydroxymethyl)-(3-O-α-D-glucopyranosyl)pyrrolidin-1-yl)ethyl)phosphonate (17)

¹³C NMR of dibenzyl ((3-hydroxy-2,5-bis(hydroxymethyl)-(3-*O*-α-D-glucopyranosyl)pyrrolidin-1-yl)ethyl)phosphonate (17)







¹H NMR of ((3-hydroxy-2,5-bis(hydroxymethyl)-(3-*O*-α-D-glucopyranosyl)pyrrolidin-1-yl)ethyl)phosphonic acid (6)

¹³C NMR of ((3-hydroxy-2,5-bis(hydroxymethyl)-(3-*O*-α-D-glucopyranosyl)pyrrolidin-1-yl)ethyl)phosphonic acid (6)



³¹P NMR of ((3-hydroxy-2,5-bis(hydroxymethyl)-(3-*O*-α-D-glucopyranosyl)pyrrolidin-1-yl)ethyl)phosphonic acid (6)

