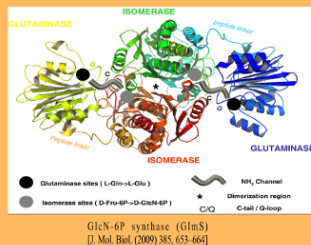
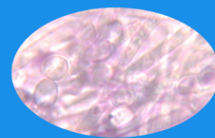


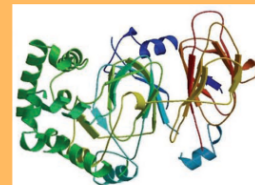
In search of new antifungal agents. Synthesis of benzyl and thiophenyl 2-deoxy-6-*O*-diethylphosphonato-2-hydroxyimino-D-hexopyranosides

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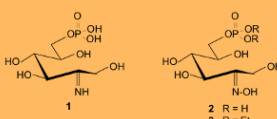
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Emerging challenge of systemic fungal infections, especially in immunocompromised patients, and a limited repertoire of effective antifungals stimulate a search for novel targets and drug candidates. Enzymes involved in biosynthesis of the fungal cell wall components are of a special interest in this respect. One of them is GlcN-6P synthase (GlmS), enzyme catalyzing the first committed step in chitin biosynthesis pathway, that is transformation of D-fructose-6-phosphate (Fru-6P) to D-glucosamine-6-phosphate (GlcN-6P). Another one is phosphomannose isomerase (PMI), that catalyses the reversible isomerization of D-mannose-6-phosphate (Man-6P) and D-fructose-6-phosphate (Fru-6P). PMI is reported to play a crucial role in the biosynthesis of many mannosylated structures, including the cell wall components of fungi. Both enzymes are proposed as the targets for antifungal chemotherapy and a search for their selective inhibitors has been continued. The reaction performed by GlmS is believed to proceed through the formation of intermediate 1, a Schiff base created between the keto group of the sugar and the ammonia generated from the glutamine amide function. Mechanism of the reaction catalyzed by PMI is similar to that catalyzed by GlmS.



Phosphomannose isomerase (PMI)
[J. Biol. Chem. (2004) Vol. 279, No. 2, p. 1491-1498]



In the course of search of antifungal agents, we plan to synthesize 2-deoxy-2-hydroxyimino-6-*O*-phosphono-D-glucitol (**2**) and its diethyl ester (**3**), and test them as inhibitors of mentioned above enzymes and for antifungal activity. Ethyl residues are incorporated into **2** to increase a lipophilicity of the molecule, which is supposed to be advantageous for better penetration of the derivative through the cytoplasmic cell membrane. Probably, diethyl ester will be metabolized to **2** inside a cell.

Here, the first steps of our synthesis are presented. This starts from the transformation of D-glucose (**4**), via 3,4,6-tri-*O*-acetyl-2-deoxy-2-nitroso- α -D-glucopyranosyl chloride (**5**), into benzyl (**6-8**) and thiophenyl 3,4,6-tri-*O*-acetyl-2-deoxy-2-hydroxyimino-D-hexopyranosides (**9-11**).

Some of the obtained glycosides (**6, 7, 10**) were *O*-deacetylated (**12-15**) and then 6-*O*-phosphorylated with diethylchlorophosphate to provide benzyl and thiophenyl 2-deoxy-6-*O*-diethylphosphonato-2-hydroxyimino-D-hexopyranosides (**16-18**).

Removing of aglycone from **16, 17** and **18**, respectively, followed by reduction will give us (2*Z*) and (2*E*)-2-deoxy-2-hydroxyimino-6-*O*-diethylphosphonato-D-glucitols (**3**).

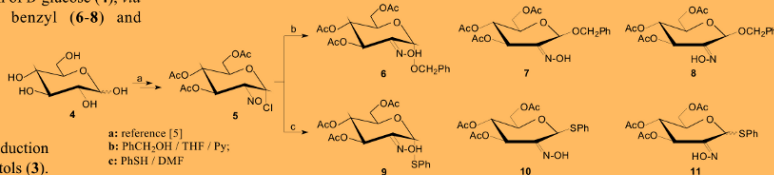


Table 1
Characteristic and Maltidof MS data for **6-18**.

No.	Configuration	Yield (%)	Form (mp [°C])	[α] _D ²⁵ (c, solvent)	M (g/mol)	MALDI/TOF MS
6	4 <i>Z</i> - α -D-arabinofuranose	26	symp	+81.5° (c=0.8, CHCl ₃)	409	432 (M ⁺ + 23)
7	4 <i>Z</i> - β -D-arabinofuranose	9	symp		409	432 (M ⁺ + 23)
8	4 <i>E</i> - β -D-arabinofuranose	14	symp	-8.4° (c=1, CHCl ₃)	409	432 (M ⁺ + 23)
9	4 <i>Z</i> - α -D-arabinofuranose	43	symp		411	412 (M ⁺ + 1)
10	4 <i>Z</i> - β -D-arabinofuranose	24	symp		411	412 (M ⁺ + 1)
11	4 <i>E</i> - α -D-arabinofuranose	1	symp		411	412 (M ⁺ + 1)
12	4 <i>E</i> - α -D-arabinofuranose	11	symp		282	305 (M ⁺ + 23)
13	4 <i>Z</i> - α -D-arabinofuranose	43	solid (144-146)	+103.0° (c=0.8, CH ₂ OH)	282	305 (M ⁺ + 23)
14	4 <i>E</i> - β -D-arabinofuranose	96	solid (146-147)	-23.0° (c=0.8, CH ₂ OH)	282	305 (M ⁺ + 23)
15	4 <i>Z</i> - α -D-arabinofuranose	53	solid		285	
16	4 <i>Z</i> - α -D-arabinofuranose	42	symp		419	442 (M ⁺ + 23)
17	4 <i>E</i> - β -D-arabinofuranose	14	symp		419	
18	4 <i>Z</i> - α -D-arabinofuranose	8	symp		421	

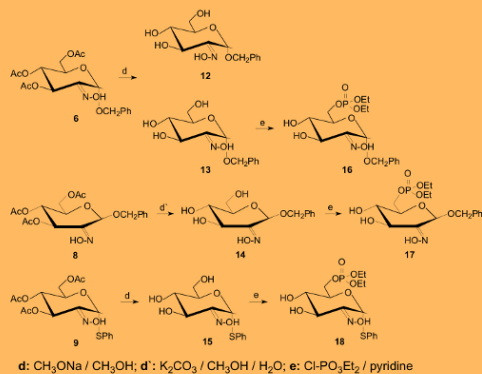


Table 2
Chemical shifts (ppm) of the protons in the ¹H NMR spectra of **6-18**.

No.	H-1	H-3	H-4	H-5	H-6'	H-6''	OAc	O-CH ₂ -Ph	N-OH	-CH ₂ CH ₂ -CH ₂	O-SPh
	H-A	H-B	-C ₆ H ₅								
6	6.09 (s) 1H	5.32 (d) 1H	5.21 (m) 4.19 (dd) 1H	4.02 (dd) 1H	4.29 (dd) 1H	2.09 (s) 3H 2.06 (s) 3H 2.04 (s) 3H	4.68 (d) 1H 4.75 (d) 1H 7.36 (m) 5H	8.22 (s) 1H	-	-	-
7	5.89 (s) 1H	5.54 (d) 1H	5.32 (m) 4.84 (s) 1H	4.44 (d) 2H	4.36 (dd) 1H	2.07 (s) 3H 2.08 (s) 3H 2.09 (s) 3H	4.70 (d) 1H 4.91 (d) 1H 7.34 (m) 5H	-	-	-	-
8	5.23 (s) 1H	6.12 (d) 1H	5.68 (dd) 3.97 (dd) 1H	4.25 (dd) 1H	4.36 (dd) 1H	2.02 (s) 3H 2.05 (s) 3H	4.68 (d) 1H 4.92 (d) 1H 7.37 (m) 5H	7.31 (s) 1H	-	-	-
9	6.55 (s) 1H	5.54 (d) 1H	5.97 (t) 4.63 (m) 1H	4.04 (dd) 1H	4.26 (dd) 1H	1.97 (s) 3H 2.07 (s) 3H 2.09 (s) 3H	-	-	12.11 (s) 1H	-	7.40 3H 7.58 2H
10	6.25 (s) 1H	5.33 (d) 1H	5.84 (dd) 3.81 (m) 1H	4.29 (dd) 1H	4.34 (dd) 1H	2.02 (s) 3H 2.07 (s) 3H 2.09 (s) 3H	-	-	12.19 (s) 1H	-	7.10 1H 7.40 2H 7.55 2H
11a	6.38 (s) 1H	5.96 (d) 1H	4.98 (s) 4.30 (m) 1H	4.20 (m) 1H	4.07 (m) 1H	-2.80 9H	-	-	12.00 (s) 1H	-	7.40 3H 7.55 2H
11b	5.96 (s) 1H	6.00 (d) 1H	5.23 (dd) 4.05 (m) 1H	4.20 (m) 2H	-2.80 9H	-	-	-	11.82 (s) 1H	-	7.40 3H 7.55 2H
12	6.13 (s) 1H	5.01 (d) 1H	-3.72 -3.84	-	-	-	4.62 (d) 1H 4.74 (d) 1H 7.32 (m) 5H	-	-	-	-
13	6.08 (s) 1H	4.54 (d) 1H	3.41 (t) 3.78 (dd) 1H	3.84 (dd) 1H	3.70 (dd) 1H	-	4.61 (d) 1H 4.78 (d) 1H 7.34 (m) 5H	-	-	-	-
14	5.12 (s) 1H	4.55 (d) 1H	4.19 (dd) 3.47 (m) 1H	3.85 (dd) 1H	3.73 (dd) 1H	-	4.61 (d) 1H 4.83 (d) 1H 7.32 (m) 5H	-	-	-	-
15	6.67 (s) 1H	4.32 (d) 1H	3.57 (t) 4.25 (dd) 1H	3.92 (dd) 1H	3.86 (dd) 1H	-	-	-	-	-	7.39 3H 7.65 2H
16	6.07 (s) 1H	4.51 (d) 1H	3.66 (t) 3.91 (m) 1H	4.16 1H	4.37 (dd) 1H	-	4.61 (d) 1H 4.72 (d) 1H 7.30 (m) 5H	-	4.11 (m) 4H 1.31 (s) 6H	-	-
17	5.14 (s) 1H	4.64 (d) 1H	4.43 (dd) 3.63 (m) 1H	4.22 (dd) 1H	4.35 (dd) 1H	-	4.57 (d) 1H 4.80 (d) 1H 7.26 (m) 5H	10.23 (s) 1H	4.09 (m) 4H 4.02 (m) 4H 1.17 (dd) 3H	1.28 (dd) 3H	-
18	6.65 (s) 1H	4.45 (d) 1H	3.74 (dd) 4.37 (dd) 1H	4.45 (m) 1H	4.28 (dd) 1H	-	-	-	4.08 (m) 4H 1.25 (s) 6H	1.25 (s) 6H 7.28 3H	-

Table 3
¹H-¹H and ¹H-³¹P coupling constants (Hz) in the ¹H NMR spectra of **6-18**.

No.	J _{1,2}	J _{2,3}	J _{3,4}	J _{4,5}	J _{5,6}	J _{6,7}	J _{6,8}	J _{6,9}	J _{6,10}	J _{6,11}	J _{6,12}
6	10.0	9.6	4.4	2.4	12.0	12.0	-	-	-	-	-
7	5.6	6.0	6.4	-	11.6	-	-	-	-	-	-
8	7.6	8.4	4.8	3.6	11.6	12.0	-	-	-	-	-
9	9.8	10.0	5.4	2.2	12.5	-	-	-	-	-	-
10	2.2	7.3	6.8	3.9	12.0	-	-	-	-	-	-
11a	4.4	8.3	*	*	*	-	-	-	-	-	-
11b	5.9	7.8	*	*	*	-	-	-	-	-	-
12	8.8	*	*	*	11.6	-	-	-	-	-	-
13	9.2	9.6	5.2	2.0	11.6	12.0	-	-	-	-	-
14	8.4	9.6	5.6	2.8	11.6	12.0	-	-	-	-	-
15	10.0	9.2	4.8	2.8	12.4	-	-	-	-	-	-
16	9.6	9.6	3.6	*	11.6	11.6	6.8	8.0	*	-	-
17	8.8	9.6	3.6	2.0	11.2	12.0	6.8	8.4	7.6	-	-
18	9.6	9.6	4.4	*	11.6	-	6.8	8.0	7.6	-	-

* not determined

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