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Synthesis of Clustered Disaccharide Polyphosphate Analogues of Adenophostin A

Martin de Kort, A. Rob P.M. Valentijn, Gijs A. van der Marel and Jacques H. van Boom*

Leiden Institute of Chemistry, Gorlaeus Laboratories, Leiden University, P.O. Box 9502, 2300 RA Leiden, The Netherlands

Abstract. Three new potential ligands for the IP₃ receptor (*i.e.* compounds 5-7) were prepared by Sonogashira coupling of propargyl 2-O-acetyl-5-O-benzyl-3-O-(3,4-di-O-acetyl-2,6-di-O-benzyl- α -D-glucopyranosyl)- β -D-ribofuranoside (15) with iodobenzene, 1,2-diiodobenzene and 1,2,4,5-tetraiodobenzene, followed by deacetylation, phosphorylation and deprotection. © 1997 Elsevier Science Ltd.

The adenophostins A and B (1 and 2, Fig. 1), isolated from the fermentation broth of *Penicillium* brevicompactum, are full agonists¹ of the mammalian D-myo-inositol 1,4,5-trisphosphate receptor (IP₃R). Interestingly, the binding affinity of both ligands for the IP₃R and the Ca²⁺-mobilizing activity are 10-100 times higher in comparison with the natural ligand D-myo-inositol 1,4,5-trisphosphate (IP₃, **3**, Fig. 1).^{2, 3}

Earlier studies⁴ indicated that the IP₃R, a glycoprotein spanning the membrane of the endoplasmic reticulum, harbors four independent ligand binding sites in a fourfold symmetrical spatial arrangement. It has been proposed⁵ that the IP₃R forms a Ca²⁺-channel upon the sequential binding of IP₃ to the four subunits (*i.e.* cooperative opening). On the other hand, the possibility that the binding of a *single* IP₃ molecule suffices to release Ca²⁺ into the cytosol (*i.e.* non-cooperative opening) is not excluded.⁶

Figure 1



It occurred to us that a ligand, in which four IP_3 units are anchored via a spacer to the 1,2,4,5 positions of a central phenyl core moiety, would be a useful tool in solving the existing ambiguity concerning the precise mechanism of IP_3 -mediated Ca^{2+} -channel opening. However, it may be expected⁷ that the nature and orientation of the individual hydroxyl functions in *myo*-inositol will pose a formidable barrier in constructing a D-myo-inositol derivative suitable for coupling to the core unit. Recently, Jenkins *et al.*⁸ disclosed that the adenophostin A analogue 4 displayed Ca²⁺-mobilizing potency similar to IP₃. The latter finding implies that a molecule in which the anomeric methyl group of compound 4 is replaced by an appropriate spacer would be an acceptable substitute for the corresponding IP₃-spacer containing derivative. On the basis of these considerations, we here present a route of synthesis to the mono-, di- and tetravalent adenophostin A analogues 5, 6, and 7.

Target compounds 5-7 are composed of a central phenyl core, which is anchored *via* propyl spacers to one, two or four phosphorylated glucosyl α -1,3 ribose disaccharides. Retrosynthetic analysis reveals that the assembly of these mono-, di- and tetravalent molecules can be achieved by Sonogashira coupling⁹ of iodobenzene, 1,4-diiodobenzene or 1,2,4,5-tetraiodobenzene with the common building block propargyl 2-*O*-acetyl-5-*O*-benzyl-3-*O*-(3,4-di-*O*-acetyl-2,6-di-*O*-benzyl- α -D-glucopyranosyl)- β -D-ribofuranoside (**15**). The α -glucosidic linkage in key disaccharide **15** can in principle be introduced by condensing, as reported¹⁴ for the synthesis of adenophostin A, the ribose unit **11** (see Scheme 1) with ethyl 3,4,6-tri-*O*-acetyl-2-*O*-benzyl-1-thio- β -D-glucopyranoside. In addition, it was established that the latter glucosyl donor could be replaced by the more easily accessible thioglucoside **10** (see Scheme 1).

Scheme 1



Reagents and conditions: (i) butane-2,3-dione (1.1 eq), CH(OCH₃)₃, cat. CSA, CH₃OH, reflux, 1h, 78% (2 regioisomers); (ii) BnBr, NaH, DMF, 98%; (iii) NIS/cat. TfOH, Et₂O, 30 min., 83% (α:β = 1:0); (iv) a. TBAF (1M in THF)/1,4-dioxane, 1/4, v/v, 50°C, 8h; b. BnBr, NaH, DMF, 92%; (v) a. HOAc/H₂O/(HOCH₂)₂, 14/6/3, v/v/v, reflux, 1h; b. Ac₂O, pyr, 16h, 81%. (vi) C₃H₃OH (2 eq), TMSOTf, (CH₂Cl)₂, 30 min., 81%.

The requisite ethyl thioglucoside **10** was easily available by the following two-step procedure (see Scheme 1). Protection of known¹⁰ ethyl 1-thio- β -D-glucopyranoside (**8**) with 2,2,3,3-tetramethoxybutane,¹¹ prepared *in situ* by reaction of trimethyl orthoformate with butane-2,3-dione¹² in the presence of a catalytic amount of camphorsulfonic acid (CSA) gave, after purification¹³ by silica gel column chromatography, 3,4-butane diacetal (BDA) **9**. Benzylation of **9** with benzyl bromide (BnBr) and sodium hydride (NaH) proceeded smoothly to give the fully protected glucosyl donor **10**. Glycosylation of known¹⁴ ribose acceptor **11** with **10** in the presence of the promotor *N*-iodosuccinimide (NIS) and a catalytic amount of trifluoromethanesulfonic acid (TfOH) proceeded in a stereoselective fashion to give the α -linked disaccharide **12** in 83% yield. Removal of the 5'-*O*-*t*-butyldiphenylsilyl group in **12** with tetra-*n*-butylammonium fluoride (TBAF), followed by benzylation of the resulting primary hydroxyl function, yielded compound **13**. Removal of both the 3,4-

butane diacetal and isopropylidene groups in 13 under mild acid catalysed transacetalisation conditions¹⁴ proceeded smoothly without any concomitant cleavage of the interglycosidic bond. Acetylation of the free hydroxyl functions afforded the fully protected dimer 14 as a mixture of anomers. Glycosidation of 14 with propargyl alcohol under the agency of a catalytic amount of trimethylsilyl triflate (TMSOTf) gave building block 15 in 50% yield based on 11.



Reagents and conditions: (i) 5 mol% PdCl₂(PPh₃)₂, 10 mol% Cul, Et₃N/DMF, 1/20, v/v, 16h, 80% (ii) NaOCH₃, CH₃OH, 1h, 100%; (iii) a. **21**, 1*H*-tetrazole, (CH₂Cl)₂/CH₃CN, 3/1, v/v, 30 min.; b. *t*-BuOOH, 0°C, 1h, 80%; (iv) Pd/C, H₂ (1 atm.), NaOAc, 1,4-dioxane/*iso*-propanol/H₂O, 4/2/1, v/v/v, 16h.

At this stage, attention was focused on the assembly of the target compounds 5-7. Sonogashira coupling⁹ (see Scheme 2) of terminal acetylene 15 (1.25 mmol) with iodobenzene 16 (1.00 mmol) in DMF (5 mL) under the influence of PdCl₂(PPh₃)/CuI/Et₃N gave the phenyl acetylene derivative 17 and the cross coupling product 20 in a 2:1 ratio. Formation of the latter compound was prevented by adding a solution of 15 in DMF (2 mL) over a period of 1h to the iodobenzene/catalyst solution (3 mL). Zemplén deacetylation of 17 gave 18 which was phosphitylated with dibenzyl *N*,*N*-diisopropyl phosphoramidite¹⁵ (21) followed by *in situ* oxidation of the resulting phosphite triesters with *tert*-butyl hydroperoxide to afford fully benzylated trisphosphate 19 (80% over the three steps). Debenzylation and contemporary reduction of the acetylene moiety was effected by hydrogenolysis (1 atm. H₂) over Pd-C in a buffered (NaOAc) solution to give, after purification by HW-40 gel filtration, the monovalent derivative 5 (Na⁺-salt). In a similar way, di- and tetravalent derivatives 6 and 7 were readily available starting from 1,4-diiodobenzene and 1,2,4,5-tetraiodobenzene,¹⁶ respectively. The identity and homogeneity of target compounds 5-7 were fully



Figure 2. Part of the 600 MHz ¹H (A) and ³¹P-¹H COSY (B) NMR spectra of the tetravalent adenophostin A analogue 7.

ascertained by ¹H-, ¹³C- and ³¹P-NMR spectroscopy as well as ESI mass spectrometry. For example, the symmetrical substitution pattern of the central phenyl core with four disaccharide units in 7, as well as the

position of the individual phosphate functions, was firmly established by ¹H- and ³¹P-¹H COSY NMR spectroscopy (see Figure 2).

In conclusion, a straightforward and successful approach to clustered adenophostin A analogues has been presented. It is also of interest to note that coupling of the intermediate building block 14 with different terminal alkyn-1-ols allows adaptation of the spacer length. The latter possibility is in all likelyhood required for optimal binding of the clustered disaccharide to the IP_3R . The Ca^{2+} -releasing potential and mode of channel opening (*i.e.* cooperative or non-cooperative) by analogues 5-7 is currently under investigation.

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