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Synthesis of potential bisubstrate inhibitors of *Leishmania* elongating α -D-mannosyl phosphate transferase

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Abstract—Synthesis of a potential mechanism-based bisubstrate inhibitor 1 of the elongating α -D-mannosyl phosphate transferase in *Leishmania*, comprising a guanosine subunit bound to the synthetic acceptor substrate through the methylenebisphosphonate linker, as well as its analogues 2 and 3 has been successfully accomplished. © 2003 Elsevier Ltd. All rights reserved.

Sandfly-transmitted protozoan parasites of the Leishmania genus cause a variety of debilitating or fatal diseases throughout the tropical and subtropical regions of the globe. It has been shown that survival and infectivity of the parasite in the mammalian host and in the insect vector are both dependent on the lipophosphoglycan (LPG) molecules, which are ubiquitous at the parasite's cell surface.¹ The biosynthesis of the polymeric backbone portion of this glycoconjugate [-6)-β-D-Galp- $(1 \rightarrow 4)$ - α -D-Manp- $(1-PO_3^-)_n$ proceeds through the consecutive action of the elongating α -mannosyl phosphate transferase (eMPT) and β -galactosyltransferase enzyme activities with the former only being found in the Leishmania parasite,² thus endorsing the search for selective eMPT inhibitors as prospective anti-leishmaniasis drugs. As a part of our ongoing programme towards elucidation of the enzyme's mode of action by means of synthetic probes,³ we wish to disclose here preliminary results aimed at the preparation of potential mechanism-based eMPT bisubstrate inhibitors.²

Little is known so far about the structure of the eMPT active site. However, a putative transition state model (TS) for α -mannosyl phosphate transfer could be

devised (Fig. 1) taking into account that α -mannosyl phosphate is transferred to the growing end of the LPG chain (i.e., an acceptor substrate, **AS**, X = O, R = LPG) from the donor substrate GDP-mannose en block with retention of anomeric configuration.^{2,3} Whereas the synthetic phosphonodisaccharide, dec-9-enyl β -D-ga-lactopyranosyl-(1 \rightarrow 4)- α -D-mannopyranosyl methane-phosphonate was shown to be an effective exogenous acceptor substrate [**AS**, X = CH₂, R = (CH₂)₈CH=CH₂] for the eMPT,⁵ we assume the enzyme simultaneously hosts the acceptor and the donor substrates, while the entire bond breaking-bond forming process occurs in one step in the ternary complex formed.

Thus, the covalently bound transition state analogue 1 with a nonhydrolysable methylenebisphosphonate (MBP) bridge mimicking the pyrophosphate moiety of GDP-mannose while linking together the acceptor substrate and the guanosine residue was recognised as a potential eMPT bisubstrate-type inhibitor. Moreover, the simpler analogues 2 and 3, either lacking the nucleoside subunit (2) or bearing a 2-dimethylaminoethylphosphonate residue instead of MBP (3), both being, however, able to take part in the chelation phenomena with divalent metal ion(s) presumably present in the active site of the enzyme, were also considered as synthetic targets.

Our first goal was the preparation of the simplified analogue 3 (Scheme 1). Phosphonylation of the 6'-OH in

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Figure 1. Putative transition state model for the α -D-mannosyl phosphate transfer and structures of the potential bisubstrate-type inhibitors.



Scheme 1. Reagents and conditions: (i) DCM, 1,4-dioxane, rt, 30 min, then MeOH excess, 16 h, 85%; (ii) Et_3N , DCM, rt, 30 min, 81%; (iii) NaI, Me₂CO, 60 °C, 4 h, then Me₂NH, THF, rt, 16 h, 54%; (iv) Me₂NH, THF, rt, 16 h, 86%; (v) PhSH, Et_3N , DMF, rt, 16 h, then MeONa, MeOH, rt, 5 h, 90%.

the phosphonodisaccharide 4^6 with bis(benzotriazolyl) 2-bromoethylphosphonate 5^7 provided the mixed phosphonodiester 6^8 in 85% yield. However, further transformation into the required *N*,*N*-dimethylamino derivative **8** either by direct treatment of **6** with Me₂NH/ THF or through the intermediate iodide was not as effective as expected. Repeatedly, the yield of the required product did not exceed 54%. This product was obtained more effectively through the conjugate addition of the same reagent to the vinyl phosphonate **7** easily prepared from **6** by short treatment with Et₃N.⁷ Finally, deprotection provided the target compound **3** as the bis-ammonium salt after isolation by ion-exchange chromatography.⁹

Encouraged by the straightforward implementation of the first objective we turned our efforts to the preparation of the methylenebisphosphonate derivative **2** (Scheme 2). Direct introduction of the MBP group via the monochloridate reagent **13** (derived from methylenebisphosphonic acid trimethyl ester **10**¹⁰) in the presence of Et₃N and DMAP,¹¹ while being successful when applied to the simple galactose derivative **9** (65% isolated yield of MBP-carrying glycoside **11**), completely failed in the case of the phosphonodisaccharide **4**. These results were replicated in a parallel Mitsunobu esterification¹² of **9** and **4** with the triester **10**. In this case, a high yield of the phosphonylated monosaccharide derivative **11** (83%) again contrasted with the formation of only traces of the desired disaccharide product **16**. In agreement with our previous observations,⁶ the apparently diminished reactivity of the 6'-OH in compound **4** could be accounted for by the steric hindrance imposed by the benzoate protection at 3-OH of the mannose residue closely flanking the reactive site, as revealed by inspection of molecular models.

At this juncture we decided that the synthetic approach had to be diverted to the consecutive esterification–gly-cosylation concept developed by us previously.⁶ Thus, a suitable glycosyl donor already bearing the MBP moiety had to be prepared from **11** and reacted with the known mannosyl methanephosphonate acceptor 17^6 to give access to a suitable precursor of **2**.

All attempts to elaborate the 2-(trimethylsilyl)ethyl galactoside 11 (via the hemiacetal derivative) into the corresponding glycosyl trichloroacetimidate or bromide proved to be unsuccessful due to the instability of the MBP substructure towards either basic (DBU or Cs_2CO_3/DCM) or mild brominating agents [HBr/DCM or (COBr)₂/DMF/DCM]. The solution was eventually found when we tested the MBP-carrying thioglycoside 14 as a stable glycosyl donor. Mitsunobu esterification of the parent thiogalactoside 12 with the triester 10 provided 14 in 67% optimised yield along with ca. 20% of the *exo*-methylene by-product 15. Rewardingly, the MeOTf-promoted glycosylation of 17 with 14 proceeded



Scheme 2. Reagents and conditions: (i) (COCl)₂, DMF cat, DCM, 0 °C, 1 h; (ii) Et₃N, DMAP, DCM, 0 °C to rt, 6 h, 65% for 9+13, no reaction for 4+13; (iii) PPh₃, DIAD, THF, 60 °C, 1 h, 83% for 9+10, 67% for 12+10, traces for 4+10; (iv) MeOTf, MS 4Å, DCM, rt, 2.5 h, 72%; (v) TFA, DCM–H₂O, rt, 15 min, then PhSH, Et₃N, DMF, rt, 16 h, then MeONa, MeOH, rt, 5 h, 80%.



Scheme 3. Reagents and conditions: (i) PhSH, Et₃N, DMF, rt, 16 h, quant; (ii) PhCHN₂, *t*-BuOMe, HBF₄ cat, rt, 67%; (iii) KCN, DMF, 70 °C, 6 h, 88%; (iv) PPh₃, DIAD, THF, 60 °C, 4 h, 32% for **22+23**, 65% for **22+24**; (v) MeOTf, MS 4 Å, DCM, rt, 4 h, 50%.

with complete stereoselectivity to give the β -linked phosphonodisaccharide **18** ($J_{1',2'} = 8.0$ Hz; $\delta_{C1'} = 101.7-101.9$) as a mixture of four diastereomers at phosphorus in 72% yield. Finally, the stepwise removal of the protecting groups from **18** afforded the required compound **2** as the tris-ammonium salt after purification by ion-exchange chromatography.¹³

Going ahead with the construction of the guanosinecontaining compound 1 we first intended this to be accomplished through the stereoselective glycosylation of the mannosyl methanephosphonate acceptor **17** with the complex guanosine–MBP–thioglycoside donor **25** (Scheme 3) that could be prepared, in turn, via the Mitsunobu esterification of the monodeprotected MBPcarrying thioglycoside **19** with a suitable guanosine derivative **23**.^{14a} It is worthy to note, that the protective group pattern in the derivative **23** meant that the final deprotection could be performed in a nonhydrogenolytic way^{14b} thus keeping the dec-9-enyl moiety, important for the preparation of neoglycoconjugates,¹⁵ intact in the targeted product.

In the event, an attempted *regioselective* monodemethylation of the phosphonic dimethyl ester in 14 with either neat t-BuNH₂ or DABCO in THF only resulted in the formation of an inseparable 1:1 mixture of 19 and its regioisomer with the phosphonic monoacid proximal to the galactose residue.¹⁶ Obviously, the protecting groups at the MBP moiety had to be remodeled to make a chemoselective deprotection feasible. To fulfil this objective the starting compound 14 was treated with PhSH-Et₃N-DMF¹⁷ to produce the phosphonic diacid 20 almost quantitatively. This was exhaustively esterified¹⁸ with phenyl diazomethane in the presence of HBF₄ to provide the compound **21** with the selectively protected MBP group in 67% yield, which was smoothly converted into the required monoacid derivative 22 on treatment with KCN in DMF.19

The Mitsunobu esterification of 22 with the guanosine synthone 23 provided the targeted thiogalactoside-MBP-guanosine building block 25 albeit in a meagre 32% yield. Predictably, when 6-O-protected guanosine derivative 24^{20} was used as the hydroxylic component in the same reaction, the yield of the corresponding ester 26 was almost twice as high. The standard MeOTf promoted glycosylation of the mannosyl methanephosphonate acceptor 17 with the thioglycoside 26 did furnish some unstable material with a newly formed β -glycosidic bond ($\delta_{Cl'} = 101.2 - 101.4$). On the basis of the MALDI-TOF spectrum however, the structure of this product was tentatively assigned as the corresponding N-methylimidazolium salt 27 in accordance with the presence of a molecular ion with m/z 2037.72, exactly one methyl group heavier than required.

Despite seemingly being unproductive, the above approach already encompassed all the synthetic essentials needed for the successful preparation of the target compound 1 (vide infra). Specifically, with the established protocol of chemoselective monodeprotection of the MBP moiety in hand the introduction of the guanosine block could be advantageously switched further to the end of the synthetic sequence thus avoiding the presence of the incompatible heterocyclic base at the glycosylation step. This required, however, that the methyl protecting group for the phosphonodiester function in the mannosyl methanephosphonate acceptor 17 had to be changed to a benzyl one in order to be stable during the MBP monodeprotection.

Fortunately, preparation of the required mannosyl methanephosphonate acceptor 31 proved to be quite straightforward (Scheme 4). Selective monodeprotection of the starting mixed phosphonodiester 28⁶ with PhSH– Et₃N (nearly quantitative yield) followed by the Mitsunobu-type esterification with benzyl alcohol (92%) yield) and removal of the *p*-methoxybenzyl (MBn) protecting group (90%) provided the desired product seamlessly. Next, the standard MeOTf promoted glycosylation of the acceptor 31 with the MBP-carrying thioglycoside 21 was again completely β -selective giving the phosphonodisaccharide MBP derivative 32 $(\delta_{C1'} = 101.6 - 101.8)$ in an acceptable yield of 47% along with some 20% of compound 33 presumably arising from transesterification of one of the phosphonodiester groups present in the molecule with MeOTf. Selective saponification of the single methyl phosphonic ester in 32 with KCN in DMF proceeded without incident (75% yield), as did the crucial Mitsunobu esterification of 34 thus prepared with the guanosine derivative 23, which produced the required compound 35 in 50% yield.



Scheme 4. Reagents and conditions: (i) PhSH, Et₃N, DMF, rt, 16 h, quant; (ii) BnOH, PPh₃, DIAD, THF, rt, 1 h, 92%; (iii) DDQ, DCM–H₂O, rt, 2 h, 90%; (iv) MeOTf, MS, 4Å, DCM, rt, 4 h, 47%; (v) KCN, DMF, 70 °C, 6 h, 75%; (vi) PPh₃, DIAD, THF, rt, 6 h, 50%; (vii) TFA, DCM–H₂O, rt, 15 min, then PhSH, Et₃N, DMF, rt, 16 h, then *t*-BuOK, *t*-BuOH–THF–H₂O, rt, 16 h, 80%.

Finally, sequential deprotection²¹ of **35** performed without isolation of any intermediates followed by ionexchange chromatographic purification delivered the phosphonodisaccharide–MBP–guanosine hybrid 1^{22} as the tris-ammonium salt in 80% yield.

In conclusion, we report here the synthesis of a potential mechanism-based bisubstrate inhibitor of the elongating α -mannosyl phosphate transferase in *Leishmania* designed with the emphasis on the incorporation of a guanosine moiety linked to the acceptor substrate through the methylenebisphosphonate bridge mimicking the important guanosine-pyrophosphate motif present in the natural substrate donor GDP-mannose, as well as its simplified analogues. The results of a biological evaluation of these compounds will be disclosed elsewhere in due course.

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References and notes

- (a) Turco, S. J.; Spath, G. F.; Beverley, S. M. Trends Parasitol. 2001, 17, 223–226; (b) Ilg, T. Parasitol. Today 2000, 16, 489–497.
- (a) Carver, M. A.; Turco, S. J. J. Biol. Chem. 1991, 266, 10974–10981;
 (b) Carver, M. A.; Turco, S. J. Arch. Biochem. Biophys. 1992, 295, 309–317.
- Routier, F. H.; Higson, A. P.; Ivanova, I. A.; Ross, A. J.; Tsvetkov, Y. E.; Yashunsky, D. V.; Bates, P. A.; Nikolaev, A. V.; Ferguson, M. A. J. *Biochemistry* 2000, 39, 8017–8025.
- For the definition of the bisubstrate inhibition concept: Palcic, M.; Heerze, L.; Srivastava, O. P.; Hindsgaul, O. J. Biol. Chem. 1989, 264, 17174–17181; For recent advances in this area: Hinou, H.; Sun, X.-L.; Ito, Y. Tetrahedron Lett. 2002, 43, 9147–9150, and references cited therein.
- Borodkin, V. S.; Ferguson, M. A. J.; Nikolaev, A. V. *Tetrahedron Lett.* 2001, 42, 5305–5308.
- Borodkin, V. S.; Milne, F. C.; Ferguson, M. A. J.; Nikolaev, A. V. *Tetrahedron Lett.* 2002, 43, 7821–7825.
- Van der Klein, P. M. A.; Dreef, C. E.; van der Marel, G. A.; van Boom, J. H. *Tetrahedron Lett.* **1989**, *30*, 5473– 5476.
- 8. All new compounds showed satisfactory analytical and spectral data.
- 9. Compound 3: ¹H NMR (300 MHz, D₂O): δ 1.16–1.31 (10H, m, 5×CH₂), 1.51 (2H, quintet, J 6.5, -OCH₂CH₂-), 1.85–2.07 (6H, m, H-1*a, -1*b, -CH₂CH₂NMe₂ and -CH₂CH=CH₂), 2.76 and 2.77 (6H, 2×s, -NMe₂), 3.23 (2H, m, -CH₂NMe₂), 3.44 (1H, dd, J_{2',3'} 9.8, H-2'), 3.58 (1H, dd, J_{3',4'} 3.5, H-3'), 3.60 (1H, m, H-5), 3.70–4.00 (11H, m, H-2, -3, -4, -4', -5', -6a, -6b, -6'a, -6'b and -OCH₂CH₂-), 4.13 (1H, m, H-1), 4.35 (1H, d, J_{1',2'} 7.8, H-1'), 4.86 and 4.94 (2H, 2×br d, J 10.4, J 17.2, -CH₂CH=CH₂), 5.81 (1H, ddt, J_{H,CH2} 6.5, -CH₂CH=CH₂); ¹³C NMR (75 MHz, D₂O): δ 22.0 (d, J_{C,P} 133.7, -CH₂CH₂CH₂), 33.1 (-CH₂CH=CH₂), 42.3 (-NMe₂), 53.1 (-CH₂CH=CH₂), 60.3 (C-6), 63.1 (d, J_{C,P} 4.5, C-6'), 64.7

(d, $J_{C,P}$ 4.5, $-OCH_2CH_2-$), 68.1 (C-4'), 69.1 (C-3), 70.7 (C-2'), 70.8 (d, $J_{C,P}$ 9.0, C-2), 72.3 (C-3'), 72.5 (C-5), 73.1 (C-1), 74.1 (d, $J_{C,P}$ 8.5, C-5'), 77.7 (C-4), 103.2 (C-1'), 114.0 ($-CH=CH_2$), 140.2 ($-CH=CH_2$); ³¹P NMR (121 MHz, D₂O): δ 20.6 (P'), 22.3 (P); ES-MS (–) data: m/z 692.28 (100%, [M–H][–]) (expected m/z 692.29; $C_{27}H_{53}NO_{15}P_2$ requires M, 693.29); $[\alpha]_D^{22}$ +15.2° (*c* 0.5, MeOH).

- 10. Compound 10 was prepared from tetramethyl methylenebisphosphonate by saponification with neat t-BuNH₂ at rt for 28 h, followed by evaporation of the amine, treatment of the residue with Dowex-50 (H⁺) in MeOH and concentration. The triester 10 (which contained up to 10% of the symmetric dimethyl ester) was used without additional purification.
- 11. Malachowski, W. P.; Coward, J. K. J. Org. Chem. 1994, 59, 7616–7624.
- Campbell, D. A.; Bermack, J. C. J. Org. Chem. 1994, 59, 658–660.
- 13. Compound **2**: ¹H NMR (300 MHz, D_2O): δ 1.20–1.35 $(10H, m, 5 \times CH_2), 1.54 (2H, quintet, J 6.6, -OCH_2CH_2-),$ 1.89-2.02 (4H, m, H-1*a, -1*b and -CH₂CH=CH₂), 2.10 (2H, br t, J_{H,P} 19.5, -PCH₂P-), 3.46 (1H, dd, J_{2',3'} 10.0, H-2'), 3.50 (3H, d, J_{H,P} 10.3, OMe), 3.62 (1H, dd, J_{3',4'} 3.5, H-3'), 3.65 (1H, m, H-5), 3.74-3.98 (11H, m, H-2, -3, -4, -4', -5', -6a, -6b, -6'a, -6'b and -OCH₂CH₂-), 4.16 (1H, m, H-1), 4.38 (1H, d, J_{1',2'} 7.8, H-1'), 4.89 and 4.97 (2H, 2×br d, J 10.2, J 17.3, -CH₂CH=CH₂), 5.85 (1H, ddt, J_{H,CH2}) 6.7, $-CH_2CH=CH_2$); ¹³C NMR (75 MHz, D₂O): δ 25.2(t, $J_{C,P}$ 125.9, $-PCH_2P-$), 26.8 (d, $J_{C,P}$ 133.2, C-1*), 25.0, 28.2, 28.3, 28.4, 28.6 (5×CH₂), 30.2 (d, $J_{C,P}$ 6.0, $-OCH_2CH_2-$), 33.2 (-CH₂CH=CH₂), 51.6 (d, J_{C,P} 3.5, -OMe), 60.4 (C-6), 62.6 (br, C-6'), 64.9 (d, J_{C,P} 5.6, -OCH₂CH₂-), 68.0 (C-4'), 69.3 (C-3), 70.9 (d, J_{C,P} 8.0, C-2), 71.0 (C-2'), 72.4 (C-3'), 72.9 (C-5), 73.5 (C-1), 73.8 (br, C-5'), 77.6 (C-4), 103.3 (C-1'), 114.0 (-CH=CH₂), 140.6 (-CH=CH₂); ³¹P NMR (121 MHz, D_2O): δ 18.8 (br, P' + P"), 22.5 (P); ES-MS (-) data: m/z 364.33 (100%, $[M-2H]^{2-}$) (expected m/z 364.10), 729.05 (40%, [M-H]⁻) (expected m/z 729.21; $C_{25}H_{49}O_{18}P_3$ requires M, 730.21); $[\alpha]_D^{22}$ +11.2° (c 0.52, MeOH-H₂O, 1:1).
- (a) Turner, J. J.; Filipov, D. V.; Overhand, M.; van der Marel, G. A.; van Boom, J. A. *Tetrahedron Lett.* 2001, 42, 5763–5767; (b) Vincent, S. P.; Mioskowski, C.; Lebeau, L. *Nucleos. Nucleot.* 1999, 18, 2127–2139.
- 15. Routier, F. H.; Nikolaev, A. V.; Ferguson, M. A. J. *Glycoconjugate J.* **2000**, *16*, 773–780.
- For the regioselective monodeprotection of the nucleosidemethylenebisphosphonate tribenzyl ester derivative: Ikeda, H.; Abushanab, E.; Marquez, V. E. *Bioorg. Med. Chem. Lett.* **1999**, *9*, 3069–3074.
- 17. Daub, G. W.; Van Tamelen, E. E. J. Am. Chem. Soc. 1977, 99, 3526–3528.
- (a) Creary, X. Org. Synth. 1985, 64, 207–216; (b) Liotta, L.
 J.; Ganem, B. Tetrahedron Lett. 1989, 30, 4759–4762.
- Saady, M.; Lebeau, L.; Mioskowski, C. *Helv. Chim. Acta* 1995, 78, 670–678.
- Compound 24 was synthesised following the preparation of 23 (Ref. 14a), by treatment of the common 6'-O-TBDPS precursor with Ph₂NCOCl–DIPEA in pyridine (rt, 16h, 50%) prior to desilylation with TBAF–AcOH in THF (rt, 1 h, 90%).
- Simultaneous de-O-acylation and cleavage of the N,Ndimethylformamidine N-protecting group was performed with t-BuOK in t-BuOH–THF–H₂O as recommended in Ref. 14b.
- Compound 1: ¹H NMR (300 MHz, D₂O): δ 1.15–1.31 (10H, m, 5×CH₂), 1.51 (2H, quintet, J 6.8, –OCH₂CH₂–), 1.87–1.98 (4H, m, H-1*a, -1*b and –CH₂CH=CH₂),

2.11 (2H, br t, $J_{\text{H,P}}$ 19.9, -PCH₂P–), 3.42 (2H, m, H-2' and -3'), 3.65 (1H, dt, $J_{4,5}$ 8.7, $J_{5,6a} = J_{5,6b} = 4.1$, H-5), 3.65–3.95 [11H, includes d (at δ 3.84), $J_{3',4'}$ 2.6, H-4' and m, H-2, -3, -4, -5', -6a, -6b, -6'a, -6'b and -OCH₂CH₂–], 4.07 (2H, m, H-5''a and -5''b), 4.15 (1H, m, H-1), 4.24 (1H, m, H-4''), 4.27 (1H, d, $J_{1',2'}$ 7.7, H-1'), 4.42 (1H, dd, $J_{2'',3''}$ 5.1, $J_{3'',4''}$ 3.6, H-3''), 4.70 (1H, dd, H-2''), 4.86 and 4.94 (2H, 2×br d, J 10.2, J 17.3, -CH₂CH=CH₂), 5.81 (1H, ddt, $J_{\text{H,CH2}}$ 6.7, -CH₂CH=CH₂), 5.84 (1H, d, $J_{1'',2''}$ 5.9, H-1''), 8.09 (1H, br, H-8'''); ¹³C NMR (75 MHz, D₂O): δ 26.9 (d, $J_{\text{C,P}}$ 133.1, C-1*), 25.0, 28.2, 28.3, 28.5, 28.6 (5×CH₂), 30.2 (d, $J_{\text{C,P}}$ 5.8, -OCH₂CH₂–), 33.2 (-CH₂CH=CH₂), 60.5 (C-6), 62.8 (br, C-6'), 63.9 (br, C-5''), 64.9 (d, $J_{\text{C,P}}$ 5.6,

–OCH₂CH₂–), 68.0 (C-4′), 69.3 (C-3), 70.5 (C-3″), 70.9 (2C, C-2′ and d, $J_{C,P}$ 8.5, C-2), 72.4 (C-3′), 72.9 (C-5), 73.6 (br, C-1), 73.8 (C-2″), 73.9 (br, C-5′), 77.8 (C-4), 84.0 (d, $J_{C,P}$ 6.1, C-4″), 87.0 (C-1″), 103.3 (C-1′), 114.0 (–CH=CH₂), 116.3 (br, C-5″), 137.8 (br, C-8″″), 140.5 (–CH=CH₂), 151.9 (br, C-4″′), 154.0 (C-2″″), 158.9 (C-6″″); ³¹P NMR (121 MHz, D₂O): δ 17.2, 17.5 (2×br, P′ and P″), 22.3 (P); ES-MS (–) data: m/z 489.67 (50%, [M–2H]^{2–}) (expected m/z 489.63), 500.69 (50%, [M–3H+Na]^{2–}) (expected m/z 500.62), 508.65 (100%, [M–3H+K]^{2–}) (expected m/z 508.61; C₃₄H₅₈N₅O₂₂P₃ requires M, 981.28); [α]²²_D –2.5° (*c* 0.5, MeOH–H₂O, 1:1).