



Hapten-phosphoramidites based on 6-[(2*E*)-*N*-(hexyl)prop-2-enamidyl]-2'-deoxyuridine

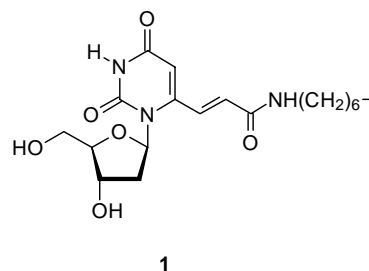
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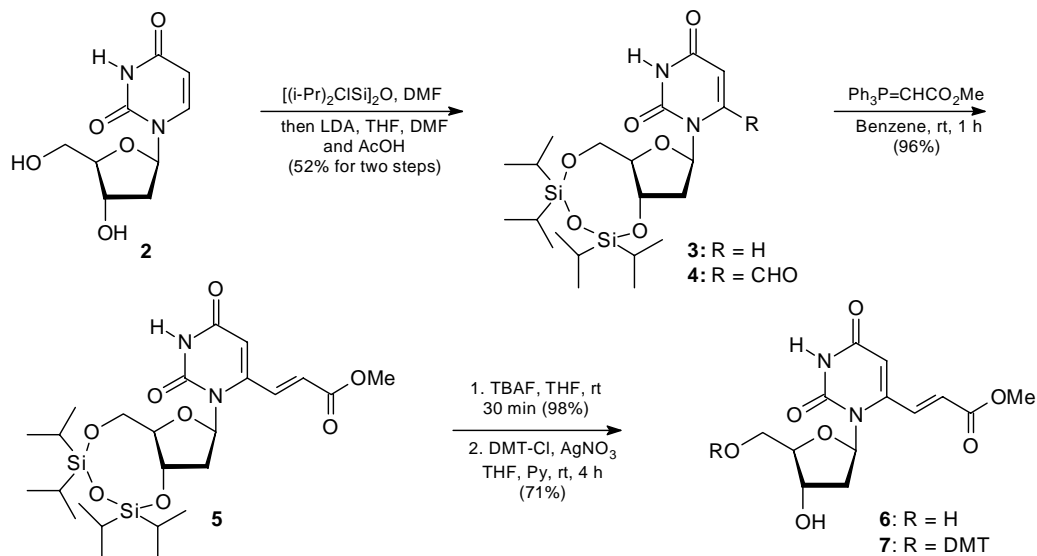
Abstract—Novel hapten-phosphoramidites **11a–c** were prepared from 2'-deoxyuridine (**2**) by functionalization at the 6-position and subsequent conjugation with adamantane, carbazole and dansyl reporter groups in good overall yield. © 2001 Elsevier Science Ltd. All rights reserved.

The inclusion of reporter groups in synthetic oligonucleotides is essential for the detection, quantification and identification of nucleic acid target sequences as a part of medical diagnoses or characterization of biotechnological processes.^{1–3} Such reporter molecules are a diverse group that include fluorescent⁴ and chemiluminescent species⁵ that can be directly measured, and other low molecular weight compounds that facilitate indirect detection. The latter category includes species with high affinity receptors (biotin/avidin)^{6–9} and an assortment of haptens (molecules that can be measured by immunoassays).¹⁰ The reporter groups are conveniently introduced by solid-phase phosphoramidite chemistry using both nucleoside^{6,9,11–14} and non-nucleoside scaffolds.^{2,15} The nucleoside scaffold approach first described by Ruth^{16,17} utilized an amino-containing linker at the 5-position of deoxyuridine to which the reporter group is attached. However, functionalization of deoxyuridine at the 6-position for this purpose has not been exploited. An explanation may lie in the report that 6-methyl-2'-deoxyuridine lowers the melt temperature (T_m) of some antisense oligonucleotides by varying degrees depending on the overall sequence, placement within the sequence and frequency of substitution.^{18,19} The authors believed that 6-Me-dU adopted a *syn* conformation which was unfavorable for Watson–Crick hydrogen bonding. In light of our success incorporating reporter groups into oligonucleotide probes using *non-nucleoside scaffolds*^{4,10} which also lack Watson–Crick hydrogen bonding, and using them in hybridization assays, we proceeded to prepare a series of hapten-phosphoramidites (**11a–c**) with a common scaffold, 6-[(2*E*)-*N*-(hexyl)prop-2-enamidyl]-2'-deoxyuridine (**1**).



Thus, 3'- and 5'-hydroxyl groups of 2'-deoxyuridine (**2**) (Scheme 1) were protected with 1,1,3,3-tetraisopropyl disioxane-1,3-diyl (TIPDS)^{20,21} group to give **3**, which was then deprotonated at the 6-position using lithium diisopropylamide and reacted with DMF in the presence of HMPA.¹³ The crude reaction mixture was treated with acetic acid at -78°C to give the aldehyde (**4**) in 52% yield after purification by silica-gel column chromatography (unreacted starting material **3**, could be recovered and recycled). The aldehyde **4** was then subjected to a Wittig reaction with methyl (triphenylphosphoranylidene)acetate in benzene to afford unsaturated ester **5** in almost quantitative yield as a *E*-isomer (>98%). The next step was to exchange the silyl protective group at the 5'-position of **5** to a 4,4'-dimethoxytrityl group (DMT), which is compatible for automated oligonucleotide synthesis. Thus, TIPDS protective group in **5** was hydrolyzed by treatment with tetra-*n*-butylammonium fluoride in THF and the crude product was purified by silica-gel column chromatography to afford diol (+)-**6** in excellent yield (98%). The 5'-hydroxyl group in **6** was protected as DMT ether by treatment with 4,4'-dimethoxytrityl chloride in the presence of silver nitrate and pyridine. The crude com-

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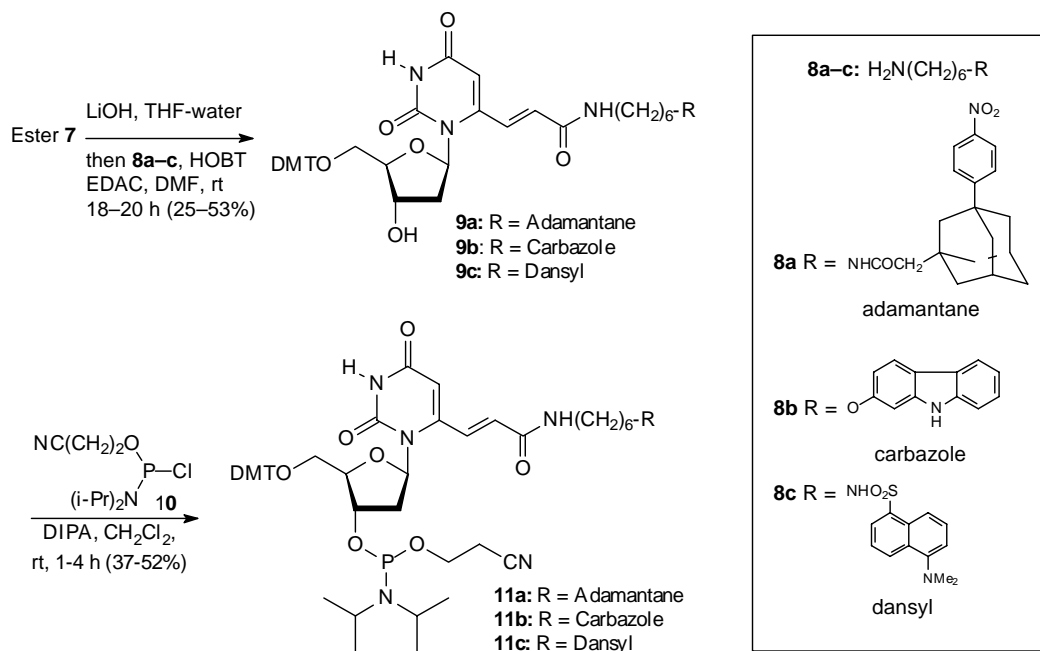


Scheme 1.

compound was purified by silica-gel column chromatography using EtOAc:Et₃N:MeOH in 97:2:1 ratio to afford ester **7** in 76% yield as a colorless solid (mp: 100–101°C). The ester **7** has the required protective group (DMT) at 5'-hydroxyl group and free 3'-hydroxy functionality for introduction of 2-(cyanoethyl)-*N,N*-diisopropylphosphoramidite, which are needed for incorporation of modified nucleotide building block into oligonucleotide via solid-phase synthesis.

The next step in synthesis of modified nucleotide building blocks **11a–c** was to conjugate ester **7** to the hapten reporter groups. Accordingly, adamantane, carbazole and dansyl reporter groups were coupled to a

C-6 linking arm containing a terminal amino functionality to provide **8a–c**.²² The ester **7** was then subjected to hydrolysis (Scheme 2) with lithium hydroxide in THF–water and the resulting crude acid was conjugated with amines **8a–c** using HOBt and EDAC in anhydrous DMF. The conjugates **9a–c** were purified by silica-gel column chromatography in 25–53% yield. Finally, **9a–c** were treated with (2-cyanoethyl)-*N,N*-diisopropylchlorophosphoramidite (**10**) in THF in the presence of 4.0 equiv. of diisopropylethylamine and the crude product was purified by preparative reversed-phase HPLC²³ to afford the phosphoramidites **11a–c** in 37–52% yield²⁴ as a mixture of diastereomers.



Scheme 2.

Adamantane, carbazole and dansyl haptens (**8a–c**) form highly antigenic protein conjugates that elicit selective, high affinity antibodies.^{25–27} The distinctive structure of the haptens ensures minimal cross-reactivity with potentially interfering substances when they are used in nucleic acid testing (NAT). The requisite hapten-phosphoramidites (**11a–c**) necessary for NAT were prepared on a 6-[(2*E*)-*N*-(6-hexyl)prop-2-enamidyl]-2'-deoxyuridine scaffold in good overall yield from 2'-deoxyuridine.

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22. The amine **8a** was prepared from 3-(4-nitrophenyl)adamantaneacetic¹⁰ and 2-amino-6-(*tert*-butoxycarbonyl)aminohexane in two steps (HOBT, Et₃N, EDAC in CH₂Cl₂ then 6.0 M aq HCl); amine **8b** was prepared from 2-hydroxy-carbazole and 6-(*p*-toluenesulfonyloxy)-*N*-(*tert*-butoxycarbonyl)aminohexane in two steps (anhydrous K₂CO₃, NaI, methyl ethyl ketone then 4.0 M aq HCl); amine **8c** was prepared from dansyl chloride and 2-amino-6-(*tert*-butoxycarbonyl)-aminohexane in two steps (Na₂CO₃ THF–water then 6.0 M aq HCl).
23. Preparative reversed-phase (RP) HPLC was carried out using Waters column (Symmetry, C18, 7.0 μm, 10×100 mm; MeCN:water/75:25, 45.0 mL/min at 225 nm). Analytical reversed phase (RP) HPLC was carried out using Waters column (Symmetry, C18, 7.0 μm, 10×100 mm).
24. Selected data for DNA Probes **11a–c**: Probe **11a**: Analytical RP HPLC: MeCN:water/76:24, 2.0 mL/min at 225 nm, *t*_R: 11.59 and 14.60 min, 96%; ¹H NMR (CD₃CN): 9.15 (br s, 1H), 8.14–8.10 (m, 2H), 7.56–7.52 (m, 2H), 7.45–7.40 (m, 2H), 7.34–7.15 (m, 8 H), 6.88–6.78 (m, 5H), 6.50–6.45 (m, 1H), 6.40 (br t, 1H), 6.14–6.06 (m, 1H), 5.69 (d, 1H, *J*=1.4 Hz), 4.56–4.36 (m, 1H), 4.08–3.96 (m, 1H), 3.75–3.62 (m, 8 H), 3.56–3.28 (m, 5H), 3.23–3.08 (m, 5H), 2.75–2.65 (m, 1H), 2.58 (t, 1H, *J*=6.0 Hz), 2.45 (t, 1H, *J*=6.0 Hz), 2.31–2.18 (m, 1H), 2.17–2.11 (m, 2H), 1.89–1.81 (m, 4H), 1.75 (br s, 1H), 1.69–1.62 (m, 6H), 1.48–1.38 (m, 4H), 1.36–1.27 (m, 4H), 1.14–1.05 (m, 9H), 0.92 (d, 3H, *J*=6.8 Hz); ³¹P NMR (CD₃CN): 148.63, 148.55; ESI–MS (*m/z*): 1197 (M+H)⁺, 1218 (M+Na)⁺. Probe **11b**: Analytical RP HPLC: MeCN:water/75:25, 2.0 mL/min at 225 nm, *t*_R: 14.47 and 18.14 min, 98%; ¹H NMR (CD₃CN): 9.28 (br s, 1H), 7.93 (d, 1H, *J*=7.7 Hz), 7.89 (d, 1H, *J*=8.5 Hz), 7.42–7.11 (m, 13H), 6.96 (d, 1H, *J*=2.2 Hz), 6.84–6.72 (m, 6H), 6.45–6.39 (m, 1H), 6.17–6.07 (m, 1H), 5.70 (d, 1H, *J*=2.5 Hz), 4.55–4.34 (m, 1H), 4.03 (t, 2H, *J*=6.5 Hz), 4.0–3.92 (m, 1H), 3.74–3.62 (m, 7H), 3.55–3.31 (m, 5H), 3.27–3.18 (m, 3H), 2.73–2.63 (m, 1H), 2.57 (t, 1H, *J*=6.3 Hz), 2.44 (t, 1H, *J*=6.0 Hz), 2.30–2.22 (m, 1H), 1.83–1.74 (m, 2H), 1.56–1.35 (m, 6H), 1.12–1.03 (m, 9H), 0.92 (d, 3H, *J*=6.6 Hz); ³¹P NMR (CD₃CN): 148.66, 148.59; ESI–MS (*m/z*): 1065 (M+H)⁺. Probe **11c**: Analytical RP HPLC: MeCN:water/72:28, 2.0 mL/min at 225 nm, *t*_R: 13.98 and 17.76 min, 96%; ¹H NMR (CD₃CN): 8.97 (brs, 1H), 8.51 (d, 1H, *J*=8.5 Hz), 8.27 (d, 1H, *J*=8.5 Hz), 8.18–8.15 (m, 1H), 7.62–7.56 (m, 2H), 7.45–7.40 (m, 2H), 7.34–7.16 (m, 9H), 6.84–6.77 (m, 4H), 6.59 (br q, 1H, *J*=5.5 Hz), 6.44–6.39 (m, 1H), 6.16–6.07 (m, 1H), 5.81 (t, 1H, *J*=6.0 Hz), 5.70–5.69 (m, 1H), 4.55–4.36 (m, 1H), 4.01–3.93 (m, 1H), 3.75–3.51 (m, 6H), 3.52–3.60 (m, 2H), 3.56–3.24 (m, 4H), 3.31–3.03 (m, 2H), 2.85–2.77 (m, 8H), 2.74–2.64 (m, 1H), 2.57 (t, 1H, *J*=6.0 Hz), 2.45 (t, 1H, *J*=6.0 Hz), 2.31–2.16 (m, 1H), 1.30–1.20 (m, 4H), 1.12–1.03 (m, 13H), 0.92 (d, 3H, *J*=6.6 Hz); ³¹P NMR (CD₃CN): 148.64, 148.55; ESI–MS (*m/z*): 1132 (M+H)⁺.
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