

**Scheme 1.** Reagents and conditions: (a) (i) *p*-nitrobenzoic acid, DIAD, Ph<sub>3</sub>P, THF; (ii) NH<sub>3</sub>, MeOH (81% for two steps); (b) *p*MBnCl, NaH, DMF (86%); (c) (i) HCl, MeOH; (ii) *m*CPBA, CH<sub>2</sub>Cl<sub>2</sub>; (iii) Me<sub>2</sub>C(OMe)<sub>2</sub>, acetone, *p*TSA (80% for three steps); (d) adenine, NaH, 15-crown-6, DMF (**10**, 43%; **11**, 39%); (e) (i) TFA; (ii) 1 N HCl (83%); (f) (i) HC(OMe)<sub>2</sub>NMe<sub>2</sub>, DMF; (ii) MsCl, DMAP, CH<sub>2</sub>Cl<sub>2</sub>; (iii) NaOMe, THF/MeOH, reflux; (iv) MeOH, reflux (60% for four steps); (g) (i) 10% TFA, CH<sub>2</sub>Cl<sub>2</sub>; (ii) 1 N HCl (51% for two steps).

the  $\alpha$ -isomer **9**, which was desired for purine nucleophilic ring opening.<sup>11</sup> Since formation of the  $\beta$ -isomer could be rationalized by invoking *iso*-propylidene steric control embodied within the Henbest rule,<sup>11,12</sup> the following sequence to **9** as the only product was employed: deprotection of the *iso*-propylidene of **8** (to the 2,3-diol), epoxidation with *m*CPBA, and re-introduction of the *iso*-propylidene (step c, Scheme 1). Nucleophilic opening of **9** with adenine gave, after careful silica gel column chromatographic separation (MeOH/EtOAc, 1:30), **10** (43%) and **11** (39%). A similar result was achieved with sodium azide as nucleophilic source on a substrate analogous to **9**.<sup>11,13</sup>

The structure of **10** was established by comparing NMR spectra: (i) compound **10** with a similarly C-5' protected aristeromycin<sup>11</sup> and (ii) compound **12**, from deprotection of **10**, with a racemic analog.<sup>11</sup> Compound **11** was confirmed by using a proton–proton COSY NMR analysis, which will be detailed in the full paper on this research.

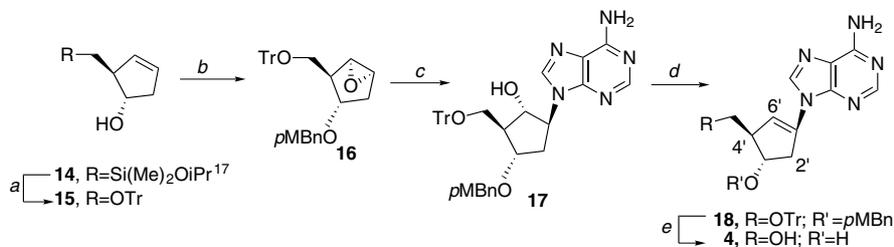
Introduction of a mesylate to C-6' of **10** was sought to provide the desired leaving group at this center. However, direct mesylation of **10** using methylsulfonyl chloride failed. In view of this, blocking the C-6 purine amine of **10** with *N,N*-dimethylformamide dimethyl acetal then allowed smooth mesylate formation. Refluxing this amino protected compound (not shown) with sodium methoxide in tetrahydrofuran followed by metha-

nol provided **13** (4 steps, 65% yield from **10**). Debenzylation of **13** with trifluoroacetic acid and, subsequent acidic deketalization completed the synthesis of **3**<sup>14</sup> (Scheme 1). Structural confirmation for **3** came from a proton NMR analysis<sup>15</sup>: (i) appearance of a peak at  $\delta$  6.60 is in the expected region<sup>16</sup> for H-6'; (ii) H-6' of isomeric neplanocin A has been reported to appear at  $\delta$  5.69<sup>15b</sup>; and (iii) presence of an H-4' peak for **3** ( $\delta$  2.79) while there was, of course, no corresponding peak for neplanocin A.<sup>15b</sup>

The synthesis of **4** began with a Tamao oxidation of **14**<sup>17</sup> (Scheme 2) followed by selective trityl protection of the resultant primary alcohol to give **15** in 65% yield. Epoxidation<sup>11</sup> of **15** followed by methoxybenzylation afforded **16**. Nucleophilic opening<sup>18</sup> of **16** with adenine resulted in the desired, single product **17**.<sup>18</sup> Following the same steps for converting **10**–**3** (steps f and g of Scheme 1), target **4**<sup>14b,19a</sup> was obtained from **17** (via **18**).

The regiochemistry of **4** was confirmed by analyzing its <sup>1</sup>H NMR spectrum,<sup>19</sup> which was substantially different from the isomeric 2'-deoxyneplanocin.<sup>19,20</sup> Particularly diagnostic was its H-6' absorption ( $\delta$  6.48), which is very similar to **3**, while the H-6' for 2'-deoxyneplanocin exists further upfield ( $\delta$  5.75).

The biological analysis of **3** and **4** is underway and will be presented in the full paper on this new class of nucleoside derivatives.



**Scheme 2.** Reagents and conditions: (a) (i) KF,  $\text{KHCO}_3$ ,  $\text{H}_2\text{O}_2$ , MeOH/THF (v/v, 50%), 75%; (ii) TrCl, pyridine, DMAP, 86%; (b) (i) *m*CPBA,  $\text{CH}_2\text{Cl}_2$ , 75%; (ii) *p*MBnCl, NaH, DMF, 94%; (c) adenine, NaH, 15-crown-6, DMF, 91%; (d) (i)  $\text{HC}(\text{OMe})_2\text{NMe}_2$ , DMF; (ii) MsCl, DMAP,  $\text{CH}_2\text{Cl}_2$ ; (iii) NaOMe, THF/MeOH, reflux; (iv) MeOH, reflux (65% for four steps); (e) (i) 10% TFA,  $\text{CH}_2\text{Cl}_2$ ; (ii) 1 N HCl (55% for two steps).

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### References and notes

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- (a) HRMS for compound **3**, 263.1014 (calcd 263.1018); (b) satisfactory microanalytical data was obtained for compounds **3** and **4**.
- (a) *NMR data for compound 3*:  $^1\text{H}$  (DMSO- $d_6$ )  $\delta$  8.31 (s, 1H), 8.20 (s, 1H), 7.35 (br s, 2H), 6.60 (d,  $J = 1.75$  Hz, 1H, H-6'), 5.30 (d,  $J = 4.25$  Hz, 1H), 4.95 (t,  $J = 6.50$  Hz, 1H), 4.83 (d,  $J = 7.0$  Hz, 1H), 4.14 (m, 1H), 3.85 (m, 1H), 3.67 (m, 1H), 3.42 (m, 1H), 2.79 (m, 1H, H-4').  $^{13}\text{C}$  (DMSO- $d_6$ )  $\delta$  161.6, 158.4, 154.7, 143.9, 143.3, 125.0, 124.5, 77.9, 76.4, 67.3, 57.8; (b) *NMR data for neplanocin*<sup>21</sup>:  $^1\text{H}$  (DMSO- $d_6$ )  $\delta$  8.11 (s, 1H), 8.05 (s, 1H), 7.20 (br s, 2H), 5.69 (s, 1H, H-6'), 5.33 (br d,  $J = 2.5$  Hz, 1H), 5.16–4.92 (br s, 3H), 4.42 (d,  $J = 5.4$  Hz, 1H), 4.30 (t,  $J = 5.4$  Hz, 1H), 4.11 (m, 2H).  $^{13}\text{C}$  (DMSO- $d_6$ )  $\delta$  156.0, 152.3, 150.1, 149.7, 139.5, 123.4, 119.2, 76.6, 72.2, 64.2, 58.6.
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- (a) *NMR data for compound 4*:  $^1\text{H}$  (DMSO- $d_6$ )  $\delta$  8.27 (s, 1H), 8.20 (s, 1H), 7.33 (br s, 2H), 6.48 (s instead of expected d, 1H, H-6'), 5.04 (d,  $J = 4.75$  Hz, 1H), 4.70 (t,  $J = 5.50$  Hz, 1H), 4.17 (m, 1H), 3.48–3.20 (m, 3H), 2.78–2.72 (m, 2H).  $^{13}\text{C}$  (DMSO- $d_6$ )  $\delta$  156.2, 153.1, 148.2, 148.1, 138.9, 133.1, 119.3, 116.7, 70.6, 62.5, 56.3, 41.1. Satisfactory microanalytical data was obtained for compound **4**; (b) *NMR data for 2'-deoxyneplanocin*<sup>20</sup>:  $^1\text{H}$  (DMSO- $d_6$ )  $\delta$  8.13 (s, 1H), 7.97 (s, 1H), 7.17 (br, 2H), 5.75 (d, 1H, H-6'), 5.64 (m, 1H), 5.06 (d, 1H), 4.85–5.0 (m, 2H), 4.15 (br, 2H), 2.2–2.4 (m, 2H).
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