## Mechanism of Nucleophilic Solvolysis of 1,2-Dibromo-1,2-dihydrocyclobuta[b]pyridine; X-Ray Crystal Structure Analysis of 1,2-Dihydro-1-methoxycyclobuta[b]pyridine Methiodide

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*trans*-1,2-Dibromo-1,2-dihydrocyclobuta[*b*]pyridine **1a** undergoes nucleophilic methanolysis with retention of configuration *via* 2-bromocyclobuta[*b*]pyridine **2**; the crystal structure of 1,2-dihydro-1-methoxycyclobuta[*b*]pyridine methiodide is described.

1,2-Dibromo-1,2-dihydrocyclobutabenzene is resistant to nucleophilic substitution, as a consequence of the strained ring imposing a relatively high degree of s-character on the carbon-bromine bonds. In contrast, we have found that *trans*-1,2-dibromo-1,2-dihydrocyclobuta[b]pyridine 1a<sup>1</sup> reacts with sodium methoxide in methanol to give predominantly (80%) the *trans*-1-substitution product 5a,<sup>†</sup> the <sup>1</sup>H NMR spectrum showing singlets for the cyclobutene protons at  $\delta$ 



Fig. 1 The structure of 1,2-dihydro-1-methoxycyclobuta[b]pyridine methiodide. Selected bond distances (Å) and angles (°): C(1)-C(2) 1.599(12), C(1)-C(6a) 1.649(13), C(1)-O(7) 1.061(13), C(2)-C(2a) 1.528(11), C(2a)-N(3) 1.324(10), C(2a)-C(6a) 1.421(10), N(3)-C(4) 1.355(10), N(3)-C(9) 1.482(9), C(4)-C(5) 1.381(11), C(5)-C(6) 1.436(14), C(6)-C(6a) 1.429(11), O(7)-C(8) 1.449(11), C(6a)-C(1)-C(2) 85.6(7), O(7)-C(1)-C(2) 129.3(9), O(7)-C(1)-C(6a) 136.0(9), C(2a)-C(2)-C(1) 84.0(6), N(3)-C(2a)-C(2) 139.6(6), C(6a)-C(2a)-C(2) 96.9(6), C(6a)-C(2a)-N(3) 122.5(7), C(4)-N(3)-C(2a) 116.7(6), C(9)-N(3)-C(2a) 120.0(6), C(9)-N(3)-C(4) 122.6(7), C(5)-C(4)-118.8(7), C(6a)-C(5) 109.4(7), C(2a)-C(6a)-C(1) 85.6(6), C(6)-C(5a)-C(1) 130.6(8), C(6)-C(5a)-C(2a) 113.4(8), C(8)-O(7)-C(1) 128.7(8).

4.96 (1-H) and 5.27 (2-H).<sup>‡</sup> This was accompanied by traces (ca. 5% each) of the corresponding cis-isomer 5b, characterised by doublets in the <sup>1</sup>H NMR spectrum for the cyclobutene protons, and the quinoline derivative 4. A similar ratio of products 4 and 5a, b was obtained under the same conditions from the cis-dibromide 1b. Preferential displacement of the 1-bromo-substituent in 1, expected from the relative electron densities at C(1) and C(2), was confirmed by catalytic hydrogenation of 5a to 1,2-dihydro-1-methoxycyclobuta[b]pyridine, followed by an X-ray crystal structure analysis§ of the corresponding methiodide salt (Fig. 1). The internal angles of the pyridine ring differ significantly from 120°, with the largest deviation for C(5)-C(6)-C(6a) of 109.4(7)°. This contrasts with 1,8-diazabiphenylene<sup>2</sup> and 1,2-dihydrocyclobuta[b]quinoline,<sup>3</sup> in which the most acute angles in the six-membered rings (at ca. 112°) are those subtended at the heteroatom; the analogous angle C(4)-N(3)-C(2a) in Fig. 1 is considerably larger, at 116.7(6)°.

The unexpected observation that substitution of **1a** proceeds with retention of the *trans*-configuration at C(1)–C(2) leads us to propose an elimination–addition pathway involving 2-bromocyclobuta[b]pyridine **2** (Scheme 1). While well-documented for aromatic and vinylic substitutions, this type of mechanism is unusual in aliphatic systems,<sup>4</sup> the more so in this case owing to the predicted high energy of the cyclobutadienoid intermediate **2**.<sup>5</sup> However, this pathway is consistent with the formation of the byproduct **4**, one of four possible 'angular' dimeric products of **2**. The structure **4** was established by comparison with substituted benzocyclobutene dimerisations<sup>6</sup> and by <sup>1</sup>H NMR spectroscopy, 7-H showing an NOE on irradiation of 6-H. The dimer **4** is the major product (88%) in Et<sub>2</sub>O solvent along with **5a** (6%); the *cis*-isomer **5b** could not be detected.

Further support for this mechanism was obtained from an analogous solvolysis of **1a** in CH<sub>3</sub>OD, which resulted in deuterium incorporation at C(2) in the product. Neither **1a** nor **5a** underwent H–D exchange under identical conditions (as determined by NMR monitoring). The elimination step is therefore essentially irreversible,  $E_{1cb}$  deprotonation at C(2) being facilitated by the  $\beta$ -nitrogen. The addition step appears to proceed *via* reprotonation of **2** followed by *anti*-attack on the resulting bromonium ion **3**; the alternative sequence of nucleophilic addition of methoxide anion to **2** followed by protonation should be less stereospecific.

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## Footnotes

† New compounds gave satisfactory microanalytical and mass spectroscopic data.

 dd, 2-H), 3.58 (3H, s, OMe), 3.66 (1H, dd, 2'-H), 4.98 (1H, dd, 1-H), 7.13 (1H, dd, 5-H), 7.49 (1H, dd, 6-H), 8.48 (1H, dd, 4-H).

§ Crystal data for C<sub>9</sub>H<sub>12</sub>NOI, M = 277.02, monoclinic, space group 539.07 Å<sup>3</sup>, F(000) = 268,  $\mu(Mo-K\alpha) = 27.24$  cm<sup>-1</sup>, Z = 2,  $D_c = 1.71$ g cm<sup>-3</sup>. Data were collected on a Philips PW 1100 diffractometer in the  $\theta$  range 3–26° with a scan width of 0.90°. Equivalent reflections were merged to give 986 absorption-corrected data with weights of  $l/\sigma(l) > 3.0$ . R = 0.0352,  $R_w = 0.0388$  with weights of  $w = 1/\sigma^2 F_o$ assigned to the individual reflections. Atomic coordinates, bond distances and angles, and thermal parameters have been deposited at the Cambridge Crystallographic Data Centre. See Information for Authors, Issue No. 1.

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