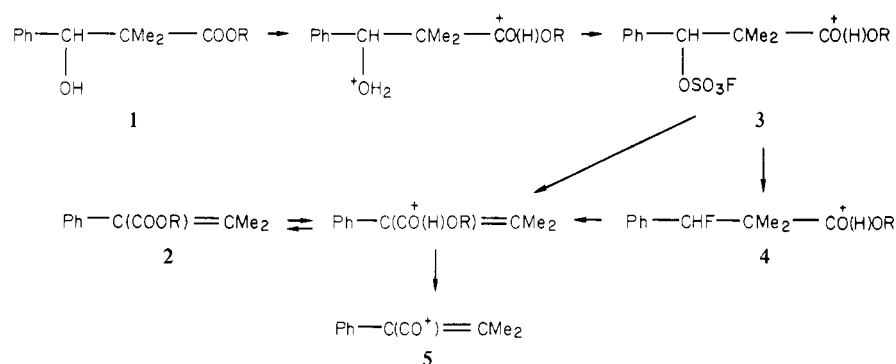


A 23 mM solution of Fp(*lin*-benzo-A) in 50 mM Tris-acetate buffer (pH 8.6) containing 13 mM MgCl<sub>2</sub> was treated with phosphodiesterase I from *C. adamanteus*<sup>53</sup> (ca. 0.05 unit in 15-μL total reaction volume).

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**Registry No. 3a**, 54193-15-6; **3b**, 80963-86-6; **5a**, 58-96-8; **5b**, 32456-54-5; **7**, 80963-87-7; **8**, 80963-89-9; **9**, 80963-91-3; **11**, 80963-92-4; **12**, 80963-93-5; **13**, 80963-94-6; **14a**, 61-19-8; **14b**, 67126-65-2; **15a**, 80963-95-7; **15b**, 80975-53-7; **16a**, 80963-97-9; **16b**, 80963-99-1; **18a**, 80964-01-8; **18b**, 80964-00-7; *N*-(2,3,5-tri-*O*-(*p*-nitrobenzoyl))-β-*D*-ribofuranosylformamide, 80964-02-9.

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Scheme 1<sup>a</sup>

<sup>a</sup> a, R = H; b, R = Me; c, R = Et.

of nearly pure, crystallized rearrangement product **2a** were isolated as the only carboxylic product: mp 149–150 °C (lit.<sup>6</sup> 151 °C); <sup>1</sup>H NMR (CDCl<sub>3</sub>) δ 7.26 (m, 5 H), 2.22 (s, 3 H), 1.70 (s, 3 H); <sup>13</sup>C NMR (CDCl<sub>3</sub>) 173.6 (C(1)), 150.7 (C(3)), 138.2–127.1 (Ph), 129.1 (C(2)), 24.4–22.8.

In order to establish the migration of the HOOC group, we prepared **1a** labeled with <sup>13</sup>C at C(3) by condensation of benzaldehyde-1-<sup>13</sup>C with isobutyric acid. When **1a**-3-<sup>13</sup>C was submitted to treatment with HSO<sub>3</sub>F/SO<sub>2</sub>ClF at 0 °C, the <sup>13</sup>C label appeared in the 2 position (δ 122.1) of protonated **2a** as well as in that of **3** (δ 94.2); it was equally visible in the <sup>1</sup>H spectra by coupling of <sup>13</sup>C with the protons of the methyl groups of protonated **2a** (δ 2.62, <sup>3</sup>J = 4.6 Hz, and δ 2.07, <sup>3</sup>J = 5.0 Hz) and of **3** (δ 2.85 and 2.47, <sup>3</sup>J = 6.0 Hz); similar values had been found for labeled **2c**.<sup>9</sup>

Definite proof for the HOOC group migration in the superacid comes from use of **1a** doubly <sup>13</sup>C labeled at C(1) (90% <sup>13</sup>C) and C(3) (69% <sup>13</sup>C).<sup>9</sup> In the rearranged (protonated) product **2a** the signals of the H<sub>2</sub>OOC<sup>+</sup> group at 180.2 ppm and of C(2) at 122.1 ppm (both increased by enrichment) are split into two doublets by direct <sup>13</sup>C,<sup>13</sup>C coupling (<sup>1</sup>J<sub>CC</sub> = 68.7 Hz). Furthermore, in quenching experiments starting with **1a**-<sup>13</sup>C<sub>2</sub>, the label appeared only in the positions 1 and 2 of (nonprotonated) **2a**, isolated and purified; <sup>13</sup>C NMR (acetone-*d*<sub>6</sub>) δ 170.4 (d, <sup>1</sup>J<sub>CC</sub> = 71.4, C(1)), 132.0 (d, <sup>1</sup>J<sub>CC</sub> = 70.5, C(2)). We conclude that the only reaction path available for the transformation **1a** → **2a** is a 1,2 shift of the HOOC group.<sup>17</sup>

In order to test whether the reaction is intermolecular (for instance via a decarbonylation–carbonylation process<sup>18</sup>), we

conducted a cross experiment using a 1:1 mixture of doubly labeled and unlabeled **1a**. There was no increase of monolabeled product in either the <sup>13</sup>C NMR spectra (judged by the amount of <sup>13</sup>C,<sup>13</sup>C-coupling between C(1) and C(2)) or in the mass spectra of isolated **2a**; this confirms the intramolecular character of the HOOC migration.

We attribute the preference for HOOC over Me migration to a difference in stability of the rearranged carbocations: if a Me group had been shifted, the positive charge would have appeared α to the carboxyl group, thus destabilizing this intermediate.

**Acknowledgment.** Financial support by the Swiss National Science Foundation is gratefully acknowledged.

**Registry No.** **1a**, 23985-59-3; **2a**, 4412-08-2; **4**, 81158-98-7.

**Supplementary Material Available:** All <sup>1</sup>H and <sup>13</sup>C NMR spectra mentioned in the text (19 pages). Ordering information is given on any current masthead page.

## Specific Peptide Sequences for Metal Ion Coordination. 1. Solid-Phase Synthesis of *cyclo*-(Gly-His)<sub>3</sub>

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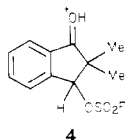
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Synthetic cyclic peptides<sup>1,2</sup> have been used to model various aspects of protein conformation and active sites.<sup>3–8</sup> The advantages of cyclic peptides<sup>3</sup> over linear peptides are the constrained geometry and the absence of free COO<sup>–</sup> and NH<sub>3</sub><sup>+</sup> terminals. Thus, a flexible polypeptide can be limited to a few desirable

(14) Other peaks observed in the reaction mixture correspond to **4** formed



from **1** (or from its fluorosulfate ester) by a Friedel–Crafts type cyclization and replacement of OH by OSO<sub>2</sub>F. The same peaks turned up upon treatment of authentic 3-hydroxy-2,2-dimethylindanone<sup>15</sup> with HSO<sub>3</sub>F/SO<sub>2</sub>ClF. This was the main product formed from **1a** with HSO<sub>3</sub>F in the absence of a solvent when the temperature was raised rapidly.

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(16) By following the reaction of **1a** with the superacid from –100 to +10 °C by <sup>1</sup>H and <sup>13</sup>C NMR spectroscopy we observed spectral changes analogous to those observed<sup>9</sup> with **1b** and **1c** and which we interpret as the consecutive transformation into the oxonium ion, the fluorosulfate, and (partially) the fluoride PhCHXCM<sub>2</sub>COOH<sub>2</sub><sup>+</sup>, X = OH<sub>2</sub><sup>+</sup>, OSO<sub>2</sub>F and F.

(17) It is not possible, however, to decide whether COOH migrates in its protonated or unprotonated form (though it might be felt that protonation would make the group too poor in electrons), nor can other transient species be excluded, e.g., a mixed anhydride (though their presence in the reaction mixture could not be detected).

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