# A TRAPPED DERIVATIVE IN STEROID BACKBONE REARRANGEMENT 

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. When treated with $\mathrm{BCl}_{3}$ at $-70^{\circ} \mathrm{C}$, compound 1 undergoes a backbone rearrangement to give compound 3 (1) via the ionic species 2 . A second derivative 4 , was also obtained (ratio 3/4: 2/1), fig. 1.

The structure of the latter compound, $\mathrm{C}_{21} \mathrm{H}_{35} \mathrm{ClO}$, investigated by $300 \mathrm{MHz}{ }^{1} \mathrm{H} \mathrm{NMR}$, indicates a partially rearranged steroid with ${ }^{21}{ }^{35}$ same primary alcohol function at C -4, like 3. A tertiary position for the chlorine atom was deduced from its chemical stability.

Up to now, the only partially rearranged compounds to be isolated are those arising from self-stabilization by proton elimination (2) to give olefinic products. Because of poor nucleophilicity, the anionic part of usual acidic reagents to be used (HF (3), $\mathrm{H}_{2} \mathrm{SO}_{4}$ (4), AcOH/TsOH (5), $\mathrm{CF}_{3} \mathrm{COOH}(6) .$. ) does not trap any cationic intermediary during the process. The use of $\mathrm{BCl}_{3}$ gives chlorine ions in the mixture through organoborane formation with the steroidal $\mathrm{CH}_{2} \mathrm{O}^{-}$molety. Its nucleophilicity is important anough to stop the rearrangement by trapping one of the cations 2. Based on actual know-


Fig. 1 ledge of the different mechanisms invoked for such rearrangement (7), the chlorine atom is expected to be located at C-8, C-9, C-10 or C-14 position for a half rearrangement, although, the $\mathrm{C}-17$ position, corresponding to a complete rearrangement, could not be firmly rouled out.

In fact, only one compound is isolated. To determine the position of the chlorine atom, an X-ray analysis of 4 was undertaken.

Compound 4, isolated by tlc, crystallizes from $\mathrm{CHCl}_{3}$ as large orthorhombic prisms, space group P2 2,2 , with $z=8$, $a=11.69 ; b=23.65 ; c=14.01 \AA$.

3942 reflections were recorded, in the $\theta / 2 \theta$ mode, on a four-circle computer controlled diffractometer (Cuka radiation, $\lambda=1.5418 \AA$, filtered by a graphite monochramator). The data set was corrected for Lorentz polarization but not for absorption.
The structure was solved by application of the multi-solution mathod ( 8 ) with some difficulty. The refinement of the atomic positional parameters and individual anisotropic thermal factors led to a final $R=5.8 \%$. All hydrogen atoms, located fram difference Fourier synthesis, were included in the final stage of the refinement, but their positions were not refined.

The two molecules of the asymmetric unit have the seme conformation. The QRTEP (10) drawing, fig. 2, of one of the molecules, indicates clearly the chlorine atom to be $\beta$ on the C-14 atom and a cis C/D ring junction,

As located, this chlorine atom has strong syn-axial interactions leading to a more strained situation in 4 than in the more flattaned compound 3. In the same way, the high value of the $4 / 3$ retio to be obtained in the final step is rather unexpected on the simple basis of a pure thermodynamic control of the reaction and the hypothesis of a slow Me-18 migration, which has been recently suggested (7) in amino-steroid backbone rearrangement, could be involved in our case to be the most important pathway of the cormpound 4 synthesis.


| ATOM | $x$ | molecture m" 1 |  |  |
| :---: | :---: | :---: | :---: | :---: |
|  |  | $\mathbf{Y}$ | 2 | "B" |
| C- 1 | 8157 | 1012 | -1958 | 4.8 |
| c- 2 | 8781 | 504 | -1536 | 5.6 |
| c- 3 | 8004 | 176 | -873 | 5.6 |
| C- 4 | 7472 | 517 | -78 | 4.1 |
| C- 5 | 6865 | 1063 | -499 | 3.5 |
| c- 6 | 6505 | 1477 | 311 | 4.5 |
| c-7 | 5953 | 2012 | -73 | 4.7 |
| c- 8 | 6774 | 2330 | -721 | 3. |
| C-9 | 7189 | 1951 | -1544 | 3.2 |
| $\mathrm{C}=10$ | 7692 | 1395 | -1163 | 3.4 |
| C-11 | 806 a | 2276 | -2145 | 4.2 |
| C-12 | 7606 | 2836 | -2513 | 4.1 |
| C=13 | 7154 | 3230 | -1736 | 4.0 |
| C-14 | 6323 | 2915 | -1048 | 4.0 |
| C-15 | 6245 | 3317 | -195 | 5.6 |
| C-16 | 7482 | 3534 | -73 | 6.6 |
| C-17 | 8071 | 3437 | -1041 | 4.8 |
| Me- 4 | 6592 | 137 | 471 | 5.3 |
| $\mathrm{Ma}-5$ | 5758 | 878 | -1049 | 5.9 |
| Me-18 | 6603 | 3752 | -2225 | 6.1 |
| Cl-14 | 4921 | 2615 | -1581 | 6. |


|  | molectins \% 2 : |  |  |  |
| :---: | :---: | :---: | :---: | :---: |
| A rOM | I | 1 | 2 | "B" |
| C- 1 | 4094 | 8626 | 3227 | 5.3 |
| C- 2 | 3850 | 9230 | 3016 | 6.9 |
| c- 3 | 3221 | 9528 | 3874 | 7.4 |
| E- 4 | 2140 | 9231 | 4202 | 5.7 |
| c- 5 | 2418 | 8577 | 4395 | 4.8 |
| c- 6 | 1327 | 8231 | 4618 | 5.7 |
| c- 7 | 1566 | 7602 | 4794 | 5.1 |
| c- 8 | 2163 | 7329 | 3952 | 4.1 |
| C-9 | 3225 | 7655 | 3618 | 4.2 |
| C-10 | 2999 | 8296 | 3505 | 4.4 |
| c-11 | 3612 | 7400 | 2685 | 5.3 |
| c-12 | 3871 | 6772 | 2750 | 6.1 |
| c-13 | 2922 | 6414 | 3199 | 5.1 |
| c-14 | 2395 | 6698 | 4084 | 4.0 |
| c-15 | 1289 | 6352 | 4239 | 3.8 |
| c-16 | 838 | 6257 | 3256 | 7.6 |
| c-17 | 1812 | 6367 | 2546 | 6.5 |
| me-4 | 1657 | 9539 | 5079 | 8.5 |
| Ms- 5 | 3212 | 8539 | 5310 | 6.6 |
| Mo-18 | 3389 | S814 | 3384 | 7.1 |
| c1-14 | 3322 | 6605 | 5113 | 5.7 |

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