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Conformation of 17-chloro-16-formylandrosta-5,16-dien-3 β -yl acetate and 17-chloroandrosta-5,16-dien-3 β -yl acetate

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In the title compounds, $C_{22}H_{29}CIO_3$, (I), and $C_{21}H_{29}CIO_2$, (II), respectively, the B rings adopt a half-chair conformation and the D rings adopt an envelope conformation. A twist of the steroid skeleton of both compounds is observed. There is a positional disorder of the acetoxy group of (II), with the terminal atoms disordered over two positions with near equal occupancy. Quantum-mechanical *ab initio* calculations using a molecular orbital Hartree–Fock method were performed for the isolated molecules, thus allowing the distinction within the structural features of these two androstane derivatives of which characteristics are intrinsic to the molecules and which are due to packing effects. The skeletal twisting was found to be innate to the molecules, while the acetoxy disorder is due to packing effects.

Comment

Treatment of 17-oxoandrost-5-en-3 β -yl acetate with POCl₃ and dimethylformamide (Vilsmeier reagent) afforded 17-chloro-16-formylandrosta-5,16-dien-3 β -yl acetate, (I), as the major and 17-chloroandrosta-5,16-dien-3 β -yl acetate, (II), as the minor reaction product (Sciaky & Pallini, 1964; Marson, 1992; Siddiqui *et al.*, 1995). Compound (I) was found to be an important precursor for the synthesis of steroidal inhibitors of cytochrome P450 17 α -hydroxylase-C17,20-lyase as potential agents for prostate cancer treatment (Njar *et al.*, 1998; Handratta *et al.*, 2005; Moreira *et al.*, 2007). We report here the molecular structures of (I) and (II), determined by single-crystal X-ray analysis, and compare them with those of the free molecules given by quantum-mechanical *ab initio* calculations.

ORTEPII (Johnson, 1976) plots of (I) and (II) are shown in Figs. 1 and 2, respectively. The two compounds are very

similar: both have acetoxy and chloro substituents at the 3 and 17 positions, and compound (I) has an extra formyl group at the 16 position. In both compounds, the A/B junction is quasi-trans and the remaining rings are trans-fused. The acetoxy

AcO (II)
$$CI$$
 AcO AcO AcO AcO AcO AcO AcO AcO AcO AcO

substituents at C3 are in equatorial positions, with angles to the normal (Cremer & Pople, 1975) of ring A of 68.43 (15) and 68.4 (2)°, respectively, for (I) and (II). In compound (II), the substituent is disordered, with atoms O2 and C31 and methyl group C32 occupying two positions with near-equal occupancy. Rings A and C have slightly flattened chair conformations, as shown by the mean values of their torsion angles [52.97 (12)–55.33 (14)°]. The unsaturated ring B adopts a half-chair conformation, with puckering parameters (Cremer & Pople, 1975) Q = 0.474 (3) Å, $\theta = 52.0$ (4)° and $\varphi = 210.0$ (4)° for (I), and Q = 0.491 (3) Å, $\theta = 51.5$ (3)° and $\varphi = 211.0$ (5)° for (II).

The five-membered ring D features a double bond at C16—C17 and shows a conformation that can be described as an envelope on C14, with P = 11.3 (3)° and $\tau = 36.0$ (2)° [P = 12.0 (4)° and $\tau = 35.7$ (2)° for (II)]. The value of the bowing angle, the angle between the least-squares plane of ring A and

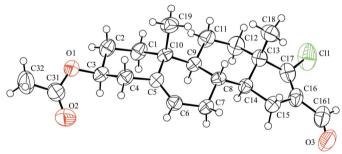
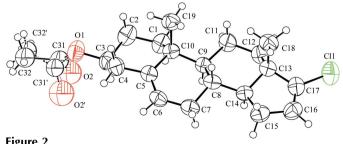


Figure 1

The molecular structure of compound (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as small spheres of arbitrary radii.



The molecular structure of compound (II), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as small spheres of arbitrary radii. The minor component of the disordered acetoxy group is shown with broken bonds.

organic compounds

the least-squares plane that includes the atoms of rings B, C and D, is 23.43 (9)° for (I) and 21.35 (4)° for (II). The corresponding distances between terminal atoms C3 and C16 are 8.921 (5) and 8.867 (5) Å, and the values for the pseudotorsion angle C19—C10···C13—C18 are 12.9 (5)° for (I) and 9.4 (5)° for (II).

We have performed quantum mechanical calculations of the equilibrium geometry of the free molecule using GAMESS (Schmidt et al., 1993) under conditions described previously (Ramos Silva et al., 2008). By comparing the calculated and experimental structural parameters it is possible to infer whether the molecular conformation is intrinsic to the free steroid molecule or is due to intermolecular interactions. The calculations reproduce the twist of the molecules as measured by the pseudo-torsion angle C19-C10···C13-C18, giving a slightly decreased value of 10.2° for (I) and a slightly increased value of 9.9° for (II). Such twists are therefore intrinsic to the molecule and not a packing effect, as confirmed by previous studies (Ramos Silva et al., 2008; Pinto et al., 2008; Paixão et al., 2004). For compound (I), the geometry of the isolated molecule that corresponds to the energy minimum is quite similar to that observed in the solid state. Larger differences are found for the head and tail groups; the calculated C3-O1-C31-O2 torsion angle is 0.6° [observed 1.5 (4)°] and the calculated C15-C16-C161-O3 torsion angle is -1.7° [observed $3.2 (5)^{\circ}$].

There are no hydrogen bonds joining the molecules of (I). They pack in such a way that the terminal acetoxy/formyl groups come close, with the shortest intermolecular contact being $O3 \cdots O1^i$ of 3.440 (3) Å [symmetry code: (i) $\frac{3}{2} - x$, 1 - y, $-\frac{1}{2} + z$]. Without the formyl substituent, the molecules of (II) pack in a different way. They assemble head-to-tail, with the shortest intermolecular contact being $C14 \cdots O2^{ni}$ of 3.311 (9) Å [symmetry code: (ii) -x + 1, $-\frac{1}{2} + y$, $\frac{1}{2} - z$].

Experimental

Details of the synthesis of the title compound have been reported previously (Siddiqui *et al.*, 1995; Njar *et al.*, 1998; Moreira *et al.*, 2007). Compounds (I) and (II) were each crystallized from a solution in a mixture of ethyl acetate and n-hexane (1:1 v/v) by slow evaporation.

Compound (I)

Crystal data

 $\begin{array}{lll} \text{C}_{22}\text{H}_{29}\text{CIO}_3 & V = 1997.24 \ (11) \ \text{Å}^3 \\ M_r = 376.90 & Z = 4 \\ \text{Orthorhombic, } P2_12_12_1 & \text{Mo } K\alpha \ \text{radiation} \\ a = 6.0689 \ (2) \ \text{Å} & \mu = 0.21 \ \text{mm}^{-1} \\ b = 13.2879 \ (4) \ \text{Å} & T = 293 \ (2) \ \text{K} \\ c = 24.7664 \ (8) \ \text{Å} & 0.42 \times 0.17 \times 0.09 \ \text{mm} \end{array}$

Data collection

 $\begin{array}{l} {\rm Bruker\ APEX\ CCD\ area-detector} \\ {\rm diffractometer} \\ {\rm Absorption\ correction:\ multi-scan} \\ {\rm (\it SADABS;\ Sheldrick,\ 2000)} \\ {\rm \it T_{min}=0.863,\ \it T_{max}=0.98} \end{array}$

38455 measured reflections 3757 independent reflections 2563 reflections with $I > 2\sigma(I)$ $R_{\rm int} = 0.084$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.044$ $wR(F^2) = 0.105$ S = 1.03 3757 reflections 238 parameters H-atom parameters constrained $\Delta \rho_{\rm max} = 0.14 \ {\rm e \ \AA^{-3}}$ $\Delta \rho_{\rm min} = -0.23 \ {\rm e \ \AA^{-3}}$ Absolute structure: Flack (1983), with 1551 Friedel pairs Flack parameter: 0.02 (10)

Compound (II)

Crystal data

 $\begin{array}{lll} {\rm C}_{21}{\rm H}_{29}{\rm CIO}_2 & V = 1913.58 \ (13) \ {\rm Å}^3 \\ M_r = 348.89 & Z = 4 \\ {\rm Orthorhombic}, P2_12_12_1 & {\rm Mo} \ K\alpha \ {\rm radiation} \\ a = 6.0271 \ (2) \ {\rm Å} & \mu = 0.21 \ {\rm mm}^{-1} \\ b = 15.4064 \ (6) \ {\rm Å} & T = 293 \ (2) \ {\rm K} \\ c = 20.6081 \ (9) \ {\rm Å} & 0.36 \times 0.13 \times 0.09 \ {\rm mm} \end{array}$

Data collection

Bruker APEX CCD area-detector diffractometer Absorption correction: multi-scan (SADABS; Sheldrick, 2000) $T_{\min} = 0.861$, $T_{\max} = 0.98$

35093 measured reflections 3615 independent reflections 2487 reflections with $I > 2\sigma(I)$ $R_{\rm int} = 0.076$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.063$ $wR(F^2) = 0.165$ S = 1.263615 reflections 219 parameters H-atom parameters constrained $\Delta \rho_{\rm max} = 0.42 \ {\rm e \ \mathring{A}^{-3}}$ $\Delta \rho_{\rm min} = -0.25 \ {\rm e \ \mathring{A}^{-3}}$ Absolute structure: Flack (1983), with 1496 Friedel pairs Flack parameter: -0.14 (13)

All H atoms were refined as riding on their parent atoms, with C— H = 0.93–0.98 Å and $U_{\rm iso}({\rm H})$ = 1.2 $U_{\rm eq}({\rm C})$ or 1.5 $U_{\rm eq}({\rm methyl~C})$. Atoms O2, C31 and C32 (and their riding H atoms) are disordered over two alternative conformations. The final refined occupancy of the major components was 0.504 (9). The disordered atoms were refined isotropically. The absolute configuration known from the synthesis route was confirmed from the X-ray data.

For both compounds, data collection: *SMART* (Bruker, 2003); cell refinement: *SAINT* (Bruker, 2003); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL97*.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: GD3236). Services for accessing these data are described at the back of the journal.

organic compounds

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