

(23*R*,23¹*S*,25*S*)-23¹,26-Epoxy-23-ethylfurost-20(22)-en-3β-ol

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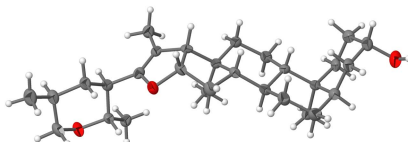
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Keywords: crystal structure; tetrahydropyran; sarsasapogenin; rearrangement reaction; hydrogen bond.

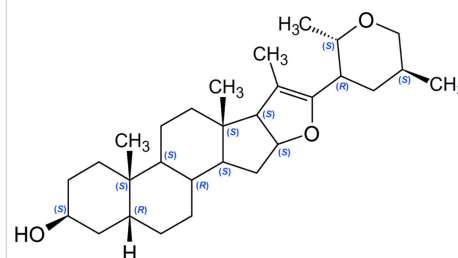
Structural data: full structural data are available from iucrdata.iucr.org

The title compound, C₂₉H₄₆O₃, is a steroid synthesized through a rearrangement of a sarsasapogenin derivative in acidic medium. The newly formed ring *F* is a tetrahydro-2*H*-pyran heterocycle substituted by two methyl groups placed in equatorial positions. This ring displays a chair conformation, while dihydrofuran ring *E*, to which it is bonded, has an envelope conformation. The molecules are associated by weak O—H···O hydrogen bonds to form chains running in the [101] direction in the crystal.

3D view



Chemical scheme



Structure description

The title steroid (**2**) was synthesized, starting from a derivative of sarsasapogenin (**1**), through a cleavage of ring *F* in acidic medium, followed by a Michael-type nucleophilic attack that affords a tetrahydro-2*H*-pyran ring bonded to the steroidal *E* ring (Fig. 1). The crystal structure of **1**·0.5H₂O has been reported previously (Viñas-Bravo *et al.*, 2003). On the other hand, the mechanism for the rearrangement **1**→**2** was described previously using a diosgenin derivative as substrate, instead of sarsasapogenin (del Río *et al.*, 2006). As expected from this mechanism, both methyl groups substituting the pyran ring in **2** are placed in equatorial positions, defining the stereochemistry for atoms C25 and C27 as *S,S* (Fig. 2). The pyran ring adopts a chair conformation, characterized by a puckering amplitude $q = 0.578$ (4) Å. Surprisingly, the Cambridge Structural Database (Version 5.43, with all updates; Groom *et al.*, 2016) does not contain any structure including the same heterocycle. However, many polysubstituted monocyclic tetrahydro-2*H*-pyran structures have been characterized by X-ray diffraction, showing that the chair confor-

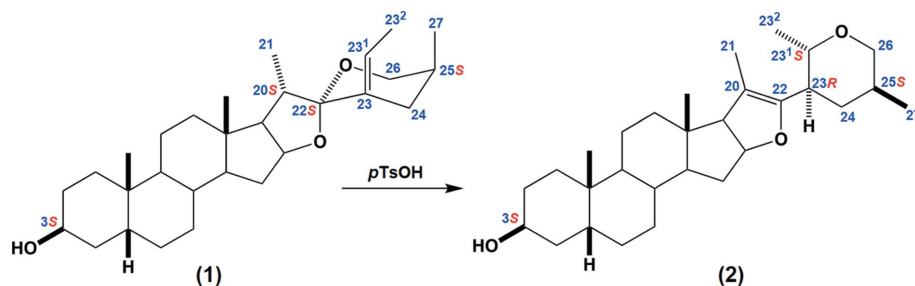


Figure 1
The synthesis of the title compound, **2**, starting from **1**. The atom-numbering scheme follows the recommendations of IUPAC (Moss, 1989). Key *R/S* configurations are displayed in red. *pTsOH* is *p*-toluenesulfonic acid.

Table 1
Hydrogen-bond geometry (Å, °).

| <i>D</i> —H... <i>A</i> | <i>D</i> —H | H... <i>A</i> | <i>D</i> ... <i>A</i> | <i>D</i> —H... <i>A</i> |
|---------------------------|-------------|---------------|-----------------------|-------------------------|
| O3—H3A...O26 ⁱ | 0.87 (4) | 2.11 (4) | 2.943 (4) | 159 (5) |

Symmetry code: (i) *x* + 1, *y*, *z* + 1.

mation is almost always stabilized (e.g. Burton *et al.*, 2007). Only a few exceptions to this rule are known, for some large molecules with steric hindrance issues (e.g. Aydilto *et al.*, 2013).

The 2,3-dihydrofuran ring *E* in **2** is close to being planar due to the presence of the C20=C22 double bond [1.324 (5) Å], also evidenced by a vibration at 1627 cm⁻¹ in the IR spectrum. The conformation can be described as an envelope with atom C16 as the flap, which belongs to the C—C bond fusing the *D* and *E* rings. The *E* ring has a small puckering amplitude, *q* = 0.184 (4) Å. A very similar conformation was observed in other steroids having the same *E* ring (Shen *et al.*, 2013; Jeong & Fuchs, 1994). The remainder of the molecular structure, i.e. the *A/B/C/D* steroidal nucleus, is identical to that of sarsasapogenin, with *cis*-fused *A/B* rings.

The crystal structure is very simple, since it is based on a single weak O—H...O hydrogen bond, involving the hydroxy group at C3 and the O-atom acceptor of the tetrahydro-2H-pyran ring, O26 (Table 1 and Fig. 3). The molecules form infinite chains, running in the [101] direction. Neighbouring chains in the crystal are related by the twofold screw axis parallel to [010] in the space group *P2*₁.

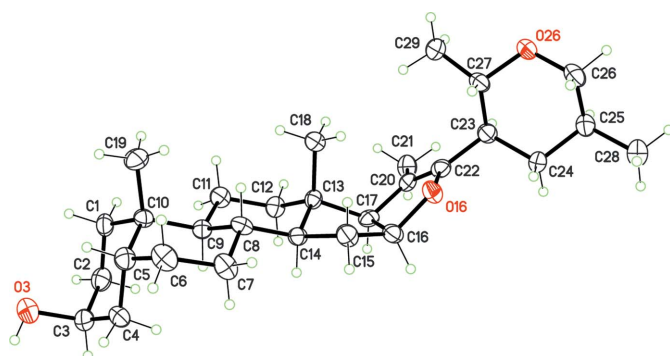


Figure 2
The molecular structure of the title compound, **2**, with displacement ellipsoids for non-H atoms drawn at the 50% probability level.

The molecular structure of **2** is embedded in a broader project aimed at targeting steroidal compounds which could interact with signalling pathways that control skeletal muscle atrophy and hypertrophy (Cohen *et al.*, 2015). Indeed, the web tool *SwissTargetPrediction* (Daina *et al.*, 2019) predicts that compound **2** presents a binding affinity for androgen and estrogen receptors, as well as for PI3K enzyme.

Synthesis and crystallization

In a round-bottomed flask was dissolved 275 mg (0.62 mmol) of **1** and 150 mg of *pTsOH* (ca 0.9 mmol) in 5 ml of benzene, and this mixture was refluxed for 30 min. The crude was then evaporated and the resulting solid dissolved in CH₂Cl₂, washed with distilled H₂O, dried over Na₂SO₄ and evaporated *in vacuo* to dryness. The residue was purified by chromatography over silica gel with hexane/EtOAc (4:1 *v:v*) as eluent, to give 268 mg of **2** (97% yield). IR (*ν*, cm⁻¹): 2998 (C—H), 1627 (C=C). ¹H NMR (500 MHz, CDCl₃): δ (ppm) 4.70 (1*H*, *m*, H-16), 4.08 (1*H*, *m*, H-3), 3.79 (1*H*, *ddd*, *J*_{26eq,26ax} = 11.0, *J*_{26eq,25} = 3.4 Hz, *J*_{26eq,24eq} = 1.4, H-26eq), 3.36 (1*H*, *dq*, *J*_{23(1),23} = 9.54, *J*_{23(1),23(2)} = 6.2 Hz, H-23¹), 2.98 (1*H*, *dd*, *J*_{26ax,26eq} = *J*_{26ax,25} = 11.0 Hz, H-26ax), 2.46 (1*H*, *d*, *J*_{17,16} = 10.0 Hz, H-17), 2.40 (1*H*, *ddd*, *J*_{23,23(1)} = 9.54, *J*_{23,24eq} = 3.4, *J*_{23,24ax} = 11.0 Hz, H-23), 1.57 (3*H*, *s*, CH₃-21), 1.11 (3*H*, *d*, *J*_{23(2),23(1)} = 6.5 Hz, C-23²), 0.95 (3*H*, *s*, CH₃-19), 0.76 (3*H*, *d*, *J*_{27,25} = 7.0 Hz, CH₃-27), 0.67 (3*H*, *s*, CH₃-18). ¹³C NMR (125 MHz, CDCl₃): δ (ppm) 152.5 (C-22), 105.0 (C-20), 85.0

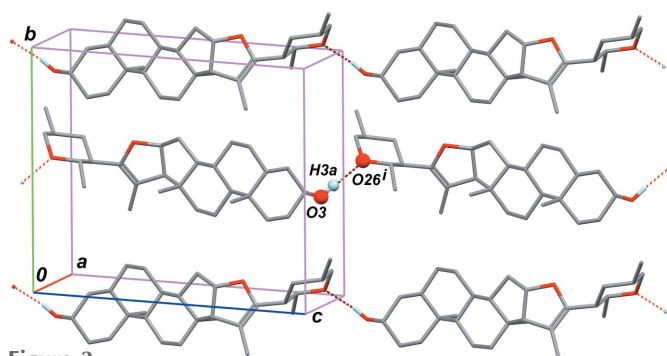


Figure 3
Part of the crystal structure of the title compound, **2**, showing chains formed *via* O—H...O hydrogen bonds (dashed bonds). For symmetry code (i), see Table 1.

Table 2
Experimental details.

| | |
|---|--|
| Crystal data | |
| Chemical formula | C ₂₉ H ₄₆ O ₃ |
| <i>M_r</i> | 442.66 |
| Crystal system, space group | Monoclinic, <i>P</i> 2 ₁ |
| Temperature (K) | 153 |
| <i>a</i> , <i>b</i> , <i>c</i> (Å) | 6.3829 (7), 12.6897 (8), 15.9539 (17) |
| β (°) | 101.308 (8) |
| <i>V</i> (Å ³) | 1267.1 (2) |
| <i>Z</i> | 2 |
| Radiation type | Ag <i>K</i> α , λ = 0.56083 Å |
| μ (mm ⁻¹) | 0.05 |
| Crystal size (mm) | 0.24 × 0.16 × 0.07 |
| Data collection | |
| Diffraction | Stoe Stadivari |
| Absorption correction | Multi-scan (<i>X-AREA</i> ; Stoe & Cie, 2018) |
| <i>T_{min}</i> , <i>T_{max}</i> | 0.519, 1.000 |
| No. of measured, independent and observed [<i>I</i> > 2 σ (<i>I</i>)] reflections | 15499, 4669, 2283 |
| <i>R_{int}</i> | 0.102 |
| ($\sin \theta/\lambda$) _{max} (Å ⁻¹) | 0.624 |
| Refinement | |
| <i>R</i> [<i>F</i> ² > 2 σ (<i>F</i> ²)], <i>wR</i> (<i>F</i> ²), <i>S</i> | 0.043, 0.077, 0.70 |
| No. of reflections | 4669 |
| No. of parameters | 298 |
| No. of restraints | 1 |
| H-atom treatment | H atoms treated by a mixture of independent and constrained refinement |
| $\Delta\rho_{\max}$, $\Delta\rho_{\min}$ (e Å ⁻³) | 0.16, -0.18 |

Computer programs: *X-AREA* (Stoe & Cie, 2018), *SHELXT2018* (Sheldrick, 2015a), *SHELXL2018* (Sheldrick, 2015b), *XP* in *SHELXTL-Plus* (Sheldrick, 2008), *Mercury* (Macrae *et al.*, 2020) and *pubCIF* (Westrip, 2010).

(C-16), 73.4 (C-23¹), 73.3 (C-26), 67.1 (C-3), 64.3 (C-17), 55.1 (C-14), 44.0 (C-9), 41.5 (C-10), 41.3 (C-23), 40.1 (C-5), 39.8 (C-4), 36.5 (C-6), 36.3 (C-15), 35.3 (C-1), 35.0 (C-12), 30.9 (C-2), 30.0 (C-13), 27.9 (C-7), 26.7 (C-8), 26.5 (C-25), 23.9 (C-24), 21.7 (C-11), 20.1 (CH₃-23²), 17.0 (CH₃-19), 14.8 (CH₃-27), 14.3 (CH₃-18), 11.9 (CH₃-21).

Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2. All H atoms were placed in calculated positions, with C–H = 1.00 (methine CH), 0.99

(methylene CH₂) or 0.98 Å (methyl CH₃). Atom H3A (of the hydroxy group) was refined with free coordinates. Isotropic displacement parameters for the H atoms were calculated as $U_{\text{iso}}(\text{H}) = xU_{\text{eq}}(\text{carrier atom})$, with $x = 1.5$ for methyl groups and the hydroxy H atom, and $x = 1.2$ for the other H atoms. The methyl groups were allowed to rotate but not to tip. Due to the absence of anomalous scatterers, the absolute configuration could not be determined and was set according to the starting material.

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full crystallographic data

IUCrData (2023). **8**, x230344 [https://doi.org/10.1107/S2414314623003449]

(23*R*,23¹*S*,25*S*)-23¹,26-Epoxy-23-ethylfurost-20(22)-en-3β-ol

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(23*R*,23¹*S*,25*S*)-23¹,26-Epoxy-23-ethylfurost-20(22)-en-3β-ol*Crystal data*

C₂₉H₄₆O₃

M_r = 442.66

Monoclinic, *P*2₁

a = 6.3829 (7) Å

b = 12.6897 (8) Å

c = 15.9539 (17) Å

β = 101.308 (8)°

V = 1267.1 (2) Å³

Z = 2

F(000) = 488

D_x = 1.160 Mg m⁻³

Ag *Kα* radiation, λ = 0.56083 Å

Cell parameters from 7481 reflections

θ = 2.4–27.7°

μ = 0.05 mm⁻¹

T = 153 K

Irregular, colourless

0.24 × 0.16 × 0.07 mm

Data collection

Stoe Stadivari

diffractometer

Radiation source: Sealed X-ray tube, Axo Astix-
f Microfocus source

Graded multilayer mirror monochromator

Detector resolution: 5.81 pixels mm⁻¹

ω scans

Absorption correction: multi-scan
(X-AREA; Stoe & Cie, 2018)

T_{min} = 0.519, *T_{max}* = 1.000

15499 measured reflections

4669 independent reflections

2283 reflections with *I* > 2σ(*I*)

R_{int} = 0.102

θ_{max} = 20.5°, θ_{min} = 2.4°

h = -7→7

k = -15→15

l = -19→19

Refinement

Refinement on *F*²

Least-squares matrix: full

R[*F*² > 2σ(*F*²)] = 0.043

wR(*F*²) = 0.077

S = 0.70

4669 reflections

298 parameters

1 restraint

0 constraints

Primary atom site location: dual

Secondary atom site location: difference Fourier
map

Hydrogen site location: mixed

H atoms treated by a mixture of independent
and constrained refinement

w = 1/[σ²(*F_o*²) + (0.0188*P*)²]

where *P* = (*F_o*² + 2*F_c*²)/3

(Δ/σ)_{max} < 0.001

Δρ_{max} = 0.16 e Å⁻³

Δρ_{min} = -0.17 e Å⁻³

Extinction correction: (SHELXL2018;

Sheldrick 2015*b*),

*F_c** = *kF_c*[1 + 0.001*xF_c*²λ³/sin(2θ)]^{-1/4}

Extinction coefficient: 0.0100 (10)

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

| | <i>x</i> | <i>y</i> | <i>z</i> | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|------|------------|--------------|--------------|----------------------------------|
| C1 | 0.7075 (7) | 0.3112 (3) | 0.7615 (3) | 0.0303 (10) |
| H1A | 0.621334 | 0.309403 | 0.806762 | 0.036* |
| H1B | 0.674723 | 0.246376 | 0.726844 | 0.036* |
| C2 | 0.9432 (7) | 0.3097 (3) | 0.8037 (3) | 0.0335 (11) |
| H2A | 0.972593 | 0.247124 | 0.841295 | 0.040* |
| H2B | 1.030808 | 0.303861 | 0.759143 | 0.040* |
| C3 | 1.0065 (6) | 0.4089 (3) | 0.8563 (2) | 0.0353 (10) |
| H3 | 1.165276 | 0.410355 | 0.875704 | 0.042* |
| O3 | 0.9070 (5) | 0.4036 (2) | 0.93021 (18) | 0.0447 (8) |
| H3A | 0.972 (8) | 0.449 (3) | 0.967 (3) | 0.067* |
| C4 | 0.9355 (6) | 0.5075 (3) | 0.8039 (2) | 0.0313 (10) |
| H4A | 1.024977 | 0.515500 | 0.760204 | 0.038* |
| H4B | 0.961535 | 0.569582 | 0.842168 | 0.038* |
| C5 | 0.7003 (7) | 0.5073 (3) | 0.7592 (2) | 0.0304 (10) |
| H5 | 0.612950 | 0.506869 | 0.804783 | 0.036* |
| C6 | 0.6443 (7) | 0.6089 (3) | 0.7069 (3) | 0.0369 (12) |
| H6A | 0.487038 | 0.614593 | 0.689524 | 0.044* |
| H6B | 0.695409 | 0.670465 | 0.743349 | 0.044* |
| C7 | 0.7433 (7) | 0.6119 (3) | 0.6272 (2) | 0.0332 (11) |
| H7A | 0.900717 | 0.615808 | 0.644573 | 0.040* |
| H7B | 0.693851 | 0.675862 | 0.593581 | 0.040* |
| C8 | 0.6827 (6) | 0.5142 (3) | 0.5712 (2) | 0.0257 (9) |
| H8 | 0.524350 | 0.514261 | 0.550355 | 0.031* |
| C9 | 0.7450 (6) | 0.4127 (3) | 0.6240 (2) | 0.0234 (9) |
| H9 | 0.902181 | 0.417847 | 0.646833 | 0.028* |
| C10 | 0.6379 (6) | 0.4079 (3) | 0.7034 (2) | 0.0255 (9) |
| C11 | 0.7125 (7) | 0.3130 (3) | 0.5679 (3) | 0.0287 (10) |
| H11A | 0.775839 | 0.252065 | 0.602689 | 0.034* |
| H11B | 0.557409 | 0.299583 | 0.549940 | 0.034* |
| C12 | 0.8121 (7) | 0.3202 (3) | 0.4879 (2) | 0.0283 (10) |
| H12A | 0.969689 | 0.322403 | 0.505309 | 0.034* |
| H12B | 0.773442 | 0.256685 | 0.452232 | 0.034* |
| C13 | 0.7352 (6) | 0.4182 (3) | 0.4354 (2) | 0.0229 (9) |
| C14 | 0.7897 (6) | 0.5144 (3) | 0.4943 (2) | 0.0235 (9) |
| H14 | 0.947049 | 0.511511 | 0.517207 | 0.028* |
| C15 | 0.7515 (7) | 0.6089 (3) | 0.4321 (2) | 0.0314 (11) |
| H15A | 0.597790 | 0.626844 | 0.416707 | 0.038* |
| H15B | 0.832343 | 0.671696 | 0.457074 | 0.038* |
| C16 | 0.8341 (7) | 0.5686 (3) | 0.3548 (2) | 0.0288 (10) |
| H16 | 0.975553 | 0.601243 | 0.352721 | 0.035* |
| O16 | 0.6841 (4) | 0.58617 (17) | 0.27418 (16) | 0.0291 (7) |
| C17 | 0.8563 (6) | 0.4468 (3) | 0.3631 (3) | 0.0251 (10) |
| H17 | 1.010420 | 0.427022 | 0.380546 | 0.030* |
| C18 | 0.4938 (5) | 0.4103 (3) | 0.3978 (2) | 0.0277 (9) |
| H18A | 0.466708 | 0.349055 | 0.359841 | 0.042* |

| | | | | |
|------|------------|--------------|--------------|-------------|
| H18B | 0.446279 | 0.474441 | 0.365412 | 0.042* |
| H18C | 0.415055 | 0.402417 | 0.444388 | 0.042* |
| C19 | 0.3914 (5) | 0.4007 (4) | 0.6764 (3) | 0.0373 (10) |
| H19A | 0.339983 | 0.455995 | 0.634456 | 0.056* |
| H19B | 0.326656 | 0.410169 | 0.726717 | 0.056* |
| H19C | 0.351771 | 0.331534 | 0.650887 | 0.056* |
| C20 | 0.7720 (5) | 0.4121 (3) | 0.2728 (2) | 0.0246 (9) |
| C21 | 0.7993 (7) | 0.3020 (3) | 0.2430 (3) | 0.0340 (11) |
| H21A | 0.720725 | 0.253044 | 0.273020 | 0.051* |
| H21B | 0.951313 | 0.283437 | 0.255138 | 0.051* |
| H21C | 0.743733 | 0.297329 | 0.181338 | 0.051* |
| C22 | 0.6775 (6) | 0.4928 (3) | 0.2281 (2) | 0.0261 (10) |
| C23 | 0.5667 (6) | 0.5036 (3) | 0.1365 (2) | 0.0268 (9) |
| H23 | 0.585841 | 0.436189 | 0.106392 | 0.032* |
| C24 | 0.6676 (6) | 0.5930 (3) | 0.0927 (2) | 0.0300 (10) |
| H24A | 0.662424 | 0.659391 | 0.124732 | 0.036* |
| H24B | 0.819359 | 0.576203 | 0.093234 | 0.036* |
| C25 | 0.5498 (7) | 0.6076 (3) | 0.0008 (3) | 0.0338 (10) |
| H25 | 0.565143 | 0.541937 | -0.032062 | 0.041* |
| C26 | 0.3142 (7) | 0.6251 (3) | 0.0011 (3) | 0.0381 (12) |
| H26A | 0.297184 | 0.690286 | 0.033231 | 0.046* |
| H26B | 0.235255 | 0.634484 | -0.058357 | 0.046* |
| O26 | 0.2251 (4) | 0.53738 (19) | 0.03962 (17) | 0.0344 (8) |
| C27 | 0.3263 (6) | 0.5230 (3) | 0.1284 (2) | 0.0276 (10) |
| H27 | 0.304525 | 0.587849 | 0.161257 | 0.033* |
| C28 | 0.6418 (7) | 0.6999 (3) | -0.0415 (3) | 0.0475 (13) |
| H28A | 0.612316 | 0.765851 | -0.014044 | 0.071* |
| H28B | 0.575716 | 0.702249 | -0.102335 | 0.071* |
| H28C | 0.796571 | 0.690815 | -0.035290 | 0.071* |
| C29 | 0.2096 (6) | 0.4323 (3) | 0.1596 (3) | 0.0335 (11) |
| H29A | 0.059098 | 0.451242 | 0.155687 | 0.050* |
| H29B | 0.274301 | 0.416708 | 0.219255 | 0.050* |
| H29C | 0.219447 | 0.370062 | 0.124274 | 0.050* |

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|-----|-----------|-------------|-------------|--------------|--------------|--------------|
| C1 | 0.036 (3) | 0.033 (2) | 0.022 (2) | -0.002 (2) | 0.008 (2) | 0.004 (2) |
| C2 | 0.037 (3) | 0.032 (2) | 0.033 (3) | 0.002 (2) | 0.010 (2) | 0.007 (2) |
| C3 | 0.031 (2) | 0.048 (2) | 0.027 (2) | -0.004 (2) | 0.005 (2) | 0.007 (2) |
| O3 | 0.050 (2) | 0.0554 (19) | 0.0292 (19) | -0.0094 (18) | 0.0075 (16) | -0.0026 (17) |
| C4 | 0.033 (3) | 0.035 (2) | 0.024 (2) | -0.004 (2) | 0.001 (2) | -0.002 (2) |
| C5 | 0.031 (3) | 0.032 (2) | 0.028 (2) | 0.002 (2) | 0.007 (2) | 0.000 (2) |
| C6 | 0.042 (3) | 0.032 (2) | 0.035 (3) | 0.008 (2) | 0.004 (2) | -0.006 (2) |
| C7 | 0.045 (3) | 0.0236 (19) | 0.029 (3) | 0.005 (2) | 0.002 (2) | 0.0007 (19) |
| C8 | 0.025 (2) | 0.0226 (19) | 0.026 (2) | 0.000 (2) | -0.0029 (19) | -0.003 (2) |
| C9 | 0.022 (2) | 0.0236 (17) | 0.025 (2) | 0.004 (2) | 0.0046 (18) | 0.001 (2) |
| C10 | 0.020 (2) | 0.031 (2) | 0.024 (2) | 0.002 (2) | 0.0025 (19) | 0.001 (2) |

| | | | | | | |
|-----|-------------|-------------|-------------|--------------|--------------|--------------|
| C11 | 0.033 (3) | 0.022 (2) | 0.030 (2) | 0.0007 (19) | 0.004 (2) | 0.002 (2) |
| C12 | 0.030 (3) | 0.026 (2) | 0.029 (3) | 0.0032 (18) | 0.007 (2) | 0.0023 (19) |
| C13 | 0.023 (2) | 0.0224 (18) | 0.024 (2) | 0.002 (2) | 0.0064 (18) | 0.0032 (19) |
| C14 | 0.022 (2) | 0.0216 (18) | 0.025 (2) | -0.0012 (19) | 0.0010 (19) | 0.004 (2) |
| C15 | 0.040 (3) | 0.0238 (19) | 0.029 (3) | -0.004 (2) | 0.001 (2) | 0.0005 (19) |
| C16 | 0.030 (3) | 0.026 (2) | 0.027 (2) | -0.0041 (18) | -0.003 (2) | 0.0039 (19) |
| O16 | 0.0371 (18) | 0.0245 (14) | 0.0235 (16) | 0.0008 (13) | 0.0006 (14) | 0.0011 (12) |
| C17 | 0.015 (2) | 0.028 (2) | 0.032 (3) | 0.0028 (16) | 0.004 (2) | 0.0017 (18) |
| C18 | 0.028 (2) | 0.0282 (18) | 0.026 (2) | 0.000 (2) | 0.0025 (18) | -0.001 (2) |
| C19 | 0.025 (2) | 0.049 (2) | 0.039 (3) | 0.001 (2) | 0.009 (2) | -0.004 (2) |
| C20 | 0.022 (2) | 0.0290 (19) | 0.025 (2) | -0.001 (2) | 0.0094 (18) | 0.001 (2) |
| C21 | 0.042 (3) | 0.031 (2) | 0.031 (3) | 0.004 (2) | 0.010 (2) | 0.003 (2) |
| C22 | 0.024 (2) | 0.029 (2) | 0.027 (2) | -0.0041 (19) | 0.008 (2) | -0.0045 (19) |
| C23 | 0.028 (2) | 0.0256 (19) | 0.026 (2) | 0.000 (2) | 0.0030 (19) | 0.000 (2) |
| C24 | 0.030 (3) | 0.033 (2) | 0.027 (2) | -0.0035 (19) | 0.006 (2) | 0.0073 (19) |
| C25 | 0.037 (3) | 0.035 (2) | 0.028 (2) | -0.005 (2) | 0.003 (2) | 0.006 (2) |
| C26 | 0.037 (3) | 0.043 (2) | 0.030 (3) | 0.001 (2) | -0.005 (2) | 0.009 (2) |
| O26 | 0.0336 (18) | 0.0410 (17) | 0.0252 (17) | -0.0081 (14) | -0.0027 (14) | 0.0052 (13) |
| C27 | 0.025 (2) | 0.034 (2) | 0.022 (2) | 0.0020 (19) | 0.0002 (19) | -0.0017 (19) |
| C28 | 0.050 (4) | 0.056 (3) | 0.036 (3) | -0.006 (3) | 0.008 (3) | 0.014 (2) |
| C29 | 0.030 (3) | 0.043 (3) | 0.029 (2) | -0.002 (2) | 0.010 (2) | 0.003 (2) |

Geometric parameters (Å, °)

| | | | |
|--------|-----------|----------|-----------|
| C1—C2 | 1.523 (5) | C15—H15A | 0.9900 |
| C1—C10 | 1.551 (5) | C15—H15B | 0.9900 |
| C1—H1A | 0.9900 | C16—O16 | 1.462 (4) |
| C1—H1B | 0.9900 | C16—C17 | 1.556 (4) |
| C2—C3 | 1.523 (5) | C16—H16 | 1.0000 |
| C2—H2A | 0.9900 | O16—C22 | 1.391 (4) |
| C2—H2B | 0.9900 | C17—C20 | 1.501 (5) |
| C3—O3 | 1.446 (4) | C17—H17 | 1.0000 |
| C3—C4 | 1.524 (5) | C18—H18A | 0.9800 |
| C3—H3 | 1.0000 | C18—H18B | 0.9800 |
| O3—H3A | 0.87 (4) | C18—H18C | 0.9800 |
| C4—C5 | 1.531 (5) | C19—H19A | 0.9800 |
| C4—H4A | 0.9900 | C19—H19B | 0.9800 |
| C4—H4B | 0.9900 | C19—H19C | 0.9800 |
| C5—C6 | 1.539 (5) | C20—C22 | 1.324 (5) |
| C5—C10 | 1.550 (5) | C20—C21 | 1.498 (5) |
| C5—H5 | 1.0000 | C21—H21A | 0.9800 |
| C6—C7 | 1.527 (5) | C21—H21B | 0.9800 |
| C6—H6A | 0.9900 | C21—H21C | 0.9800 |
| C6—H6B | 0.9900 | C22—C23 | 1.500 (5) |
| C7—C8 | 1.532 (5) | C23—C27 | 1.535 (5) |
| C7—H7A | 0.9900 | C23—C24 | 1.538 (5) |
| C7—H7B | 0.9900 | C23—H23 | 1.0000 |
| C8—C14 | 1.517 (5) | C24—C25 | 1.522 (5) |

| | | | |
|------------|-----------|---------------|-----------|
| C8—C9 | 1.548 (5) | C24—H24A | 0.9900 |
| C8—H8 | 1.0000 | C24—H24B | 0.9900 |
| C9—C11 | 1.540 (5) | C25—C26 | 1.521 (5) |
| C9—C10 | 1.554 (5) | C25—C28 | 1.525 (5) |
| C9—H9 | 1.0000 | C25—H25 | 1.0000 |
| C10—C19 | 1.550 (5) | C26—O26 | 1.441 (4) |
| C11—C12 | 1.537 (5) | C26—H26A | 0.9900 |
| C11—H11A | 0.9900 | C26—H26B | 0.9900 |
| C11—H11B | 0.9900 | O26—C27 | 1.449 (4) |
| C12—C13 | 1.525 (5) | C27—C29 | 1.507 (5) |
| C12—H12A | 0.9900 | C27—H27 | 1.0000 |
| C12—H12B | 0.9900 | C28—H28A | 0.9800 |
| C13—C14 | 1.538 (5) | C28—H28B | 0.9800 |
| C13—C18 | 1.543 (5) | C28—H28C | 0.9800 |
| C13—C17 | 1.552 (5) | C29—H29A | 0.9800 |
| C14—C15 | 1.545 (5) | C29—H29B | 0.9800 |
| C14—H14 | 1.0000 | C29—H29C | 0.9800 |
| C15—C16 | 1.521 (5) | | |
| C2—C1—C10 | 114.6 (3) | C16—C15—H15A | 111.2 |
| C2—C1—H1A | 108.6 | C14—C15—H15A | 111.2 |
| C10—C1—H1A | 108.6 | C16—C15—H15B | 111.2 |
| C2—C1—H1B | 108.6 | C14—C15—H15B | 111.2 |
| C10—C1—H1B | 108.6 | H15A—C15—H15B | 109.1 |
| H1A—C1—H1B | 107.6 | O16—C16—C15 | 113.0 (3) |
| C3—C2—C1 | 111.4 (3) | O16—C16—C17 | 105.0 (3) |
| C3—C2—H2A | 109.3 | C15—C16—C17 | 107.8 (3) |
| C1—C2—H2A | 109.3 | O16—C16—H16 | 110.3 |
| C3—C2—H2B | 109.3 | C15—C16—H16 | 110.3 |
| C1—C2—H2B | 109.3 | C17—C16—H16 | 110.3 |
| H2A—C2—H2B | 108.0 | C22—O16—C16 | 106.4 (3) |
| O3—C3—C2 | 107.4 (3) | C20—C17—C13 | 120.5 (3) |
| O3—C3—C4 | 110.7 (3) | C20—C17—C16 | 101.5 (3) |
| C2—C3—C4 | 111.0 (3) | C13—C17—C16 | 104.2 (3) |
| O3—C3—H3 | 109.2 | C20—C17—H17 | 109.9 |
| C2—C3—H3 | 109.2 | C13—C17—H17 | 109.9 |
| C4—C3—H3 | 109.2 | C16—C17—H17 | 109.9 |
| C3—O3—H3A | 107 (3) | C13—C18—H18A | 109.5 |
| C3—C4—C5 | 114.5 (3) | C13—C18—H18B | 109.5 |
| C3—C4—H4A | 108.6 | H18A—C18—H18B | 109.5 |
| C5—C4—H4A | 108.6 | C13—C18—H18C | 109.5 |
| C3—C4—H4B | 108.6 | H18A—C18—H18C | 109.5 |
| C5—C4—H4B | 108.6 | H18B—C18—H18C | 109.5 |
| H4A—C4—H4B | 107.6 | C10—C19—H19A | 109.5 |
| C4—C5—C6 | 110.7 (3) | C10—C19—H19B | 109.5 |
| C4—C5—C10 | 112.8 (3) | H19A—C19—H19B | 109.5 |
| C6—C5—C10 | 111.4 (3) | C10—C19—H19C | 109.5 |
| C4—C5—H5 | 107.2 | H19A—C19—H19C | 109.5 |

| | | | |
|---------------|-----------|---------------|-----------|
| C6—C5—H5 | 107.2 | H19B—C19—H19C | 109.5 |
| C10—C5—H5 | 107.2 | C22—C20—C21 | 128.3 (4) |
| C7—C6—C5 | 112.4 (3) | C22—C20—C17 | 109.3 (3) |
| C7—C6—H6A | 109.1 | C21—C20—C17 | 122.4 (3) |
| C5—C6—H6A | 109.1 | C20—C21—H21A | 109.5 |
| C7—C6—H6B | 109.1 | C20—C21—H21B | 109.5 |
| C5—C6—H6B | 109.1 | H21A—C21—H21B | 109.5 |
| H6A—C6—H6B | 107.9 | C20—C21—H21C | 109.5 |
| C6—C7—C8 | 111.5 (3) | H21A—C21—H21C | 109.5 |
| C6—C7—H7A | 109.3 | H21B—C21—H21C | 109.5 |
| C8—C7—H7A | 109.3 | C20—C22—O16 | 114.2 (3) |
| C6—C7—H7B | 109.3 | C20—C22—C23 | 132.4 (4) |
| C8—C7—H7B | 109.3 | O16—C22—C23 | 113.4 (3) |
| H7A—C7—H7B | 108.0 | C22—C23—C27 | 111.7 (3) |
| C14—C8—C7 | 111.6 (3) | C22—C23—C24 | 110.6 (3) |
| C14—C8—C9 | 109.5 (3) | C27—C23—C24 | 110.2 (3) |
| C7—C8—C9 | 110.3 (3) | C22—C23—H23 | 108.1 |
| C14—C8—H8 | 108.5 | C27—C23—H23 | 108.1 |
| C7—C8—H8 | 108.5 | C24—C23—H23 | 108.1 |
| C9—C8—H8 | 108.5 | C25—C24—C23 | 110.9 (3) |
| C11—C9—C8 | 112.1 (3) | C25—C24—H24A | 109.5 |
| C11—C9—C10 | 114.2 (3) | C23—C24—H24A | 109.5 |
| C8—C9—C10 | 111.6 (3) | C25—C24—H24B | 109.5 |
| C11—C9—H9 | 106.0 | C23—C24—H24B | 109.5 |
| C8—C9—H9 | 106.0 | H24A—C24—H24B | 108.1 |
| C10—C9—H9 | 106.0 | C26—C25—C24 | 108.5 (3) |
| C19—C10—C5 | 109.9 (3) | C26—C25—C28 | 110.9 (3) |
| C19—C10—C1 | 106.0 (3) | C24—C25—C28 | 111.1 (3) |
| C5—C10—C1 | 106.8 (3) | C26—C25—H25 | 108.8 |
| C19—C10—C9 | 111.2 (3) | C24—C25—H25 | 108.8 |
| C5—C10—C9 | 109.4 (3) | C28—C25—H25 | 108.8 |
| C1—C10—C9 | 113.5 (3) | O26—C26—C25 | 111.2 (3) |
| C12—C11—C9 | 113.7 (3) | O26—C26—H26A | 109.4 |
| C12—C11—H11A | 108.8 | C25—C26—H26A | 109.4 |
| C9—C11—H11A | 108.8 | O26—C26—H26B | 109.4 |
| C12—C11—H11B | 108.8 | C25—C26—H26B | 109.4 |
| C9—C11—H11B | 108.8 | H26A—C26—H26B | 108.0 |
| H11A—C11—H11B | 107.7 | C26—O26—C27 | 112.1 (3) |
| C13—C12—C11 | 111.3 (3) | O26—C27—C29 | 105.4 (3) |
| C13—C12—H12A | 109.4 | O26—C27—C23 | 110.4 (3) |
| C11—C12—H12A | 109.4 | C29—C27—C23 | 113.6 (3) |
| C13—C12—H12B | 109.4 | O26—C27—H27 | 109.1 |
| C11—C12—H12B | 109.4 | C29—C27—H27 | 109.1 |
| H12A—C12—H12B | 108.0 | C23—C27—H27 | 109.1 |
| C12—C13—C14 | 107.6 (3) | C25—C28—H28A | 109.5 |
| C12—C13—C18 | 110.5 (3) | C25—C28—H28B | 109.5 |
| C14—C13—C18 | 112.2 (3) | H28A—C28—H28B | 109.5 |
| C12—C13—C17 | 116.4 (3) | C25—C28—H28C | 109.5 |

| | | | |
|-----------------|------------|-----------------|------------|
| C14—C13—C17 | 100.5 (3) | H28A—C28—H28C | 109.5 |
| C18—C13—C17 | 109.3 (3) | H28B—C28—H28C | 109.5 |
| C8—C14—C13 | 114.1 (3) | C27—C29—H29A | 109.5 |
| C8—C14—C15 | 118.8 (3) | C27—C29—H29B | 109.5 |
| C13—C14—C15 | 103.6 (3) | H29A—C29—H29B | 109.5 |
| C8—C14—H14 | 106.5 | C27—C29—H29C | 109.5 |
| C13—C14—H14 | 106.5 | H29A—C29—H29C | 109.5 |
| C15—C14—H14 | 106.5 | H29B—C29—H29C | 109.5 |
| C16—C15—C14 | 102.9 (3) | | |
| C10—C1—C2—C3 | 56.8 (4) | C18—C13—C14—C15 | -70.0 (4) |
| C1—C2—C3—O3 | 70.1 (4) | C17—C13—C14—C15 | 46.1 (4) |
| C1—C2—C3—C4 | -51.1 (4) | C8—C14—C15—C16 | -166.3 (3) |
| O3—C3—C4—C5 | -68.4 (4) | C13—C14—C15—C16 | -38.5 (4) |
| C2—C3—C4—C5 | 50.8 (4) | C14—C15—C16—O16 | 131.1 (3) |
| C3—C4—C5—C6 | -179.1 (3) | C14—C15—C16—C17 | 15.5 (4) |
| C3—C4—C5—C10 | -53.6 (5) | C15—C16—O16—C22 | -135.1 (3) |
| C4—C5—C6—C7 | 71.5 (4) | C17—C16—O16—C22 | -17.8 (4) |
| C10—C5—C6—C7 | -54.9 (5) | C12—C13—C17—C20 | 95.8 (4) |
| C5—C6—C7—C8 | 54.7 (4) | C14—C13—C17—C20 | -148.4 (3) |
| C6—C7—C8—C14 | -177.2 (3) | C18—C13—C17—C20 | -30.2 (5) |
| C6—C7—C8—C9 | -55.3 (4) | C12—C13—C17—C16 | -151.3 (3) |
| C14—C8—C9—C11 | -49.8 (4) | C14—C13—C17—C16 | -35.5 (4) |
| C7—C8—C9—C11 | -173.0 (4) | C18—C13—C17—C16 | 82.6 (4) |
| C14—C8—C9—C10 | -179.5 (3) | O16—C16—C17—C20 | 17.8 (4) |
| C7—C8—C9—C10 | 57.4 (4) | C15—C16—C17—C20 | 138.5 (3) |
| C4—C5—C10—C19 | 167.6 (3) | O16—C16—C17—C13 | -108.1 (4) |
| C6—C5—C10—C19 | -67.1 (4) | C15—C16—C17—C13 | 12.7 (4) |
| C4—C5—C10—C1 | 53.1 (4) | C13—C17—C20—C22 | 102.3 (4) |
| C6—C5—C10—C1 | 178.3 (4) | C16—C17—C20—C22 | -12.0 (4) |
| C4—C5—C10—C9 | -70.1 (4) | C13—C17—C20—C21 | -78.2 (5) |
| C6—C5—C10—C9 | 55.1 (4) | C16—C17—C20—C21 | 167.5 (3) |
| C2—C1—C10—C19 | -173.3 (3) | C21—C20—C22—O16 | -178.3 (3) |
| C2—C1—C10—C5 | -56.1 (4) | C17—C20—C22—O16 | 1.2 (5) |
| C2—C1—C10—C9 | 64.5 (4) | C21—C20—C22—C23 | 0.7 (7) |
| C11—C9—C10—C19 | -64.1 (4) | C17—C20—C22—C23 | -179.9 (4) |
| C8—C9—C10—C19 | 64.5 (4) | C16—O16—C22—C20 | 11.1 (4) |
| C11—C9—C10—C5 | 174.4 (3) | C16—O16—C22—C23 | -168.1 (3) |
| C8—C9—C10—C5 | -57.0 (4) | C20—C22—C23—C27 | 112.5 (5) |
| C11—C9—C10—C1 | 55.3 (4) | O16—C22—C23—C27 | -68.5 (4) |
| C8—C9—C10—C1 | -176.2 (3) | C20—C22—C23—C24 | -124.5 (5) |
| C8—C9—C11—C12 | 49.6 (4) | O16—C22—C23—C24 | 54.5 (4) |
| C10—C9—C11—C12 | 177.9 (3) | C22—C23—C24—C25 | -177.5 (3) |
| C9—C11—C12—C13 | -53.6 (4) | C27—C23—C24—C25 | -53.6 (4) |
| C11—C12—C13—C14 | 56.6 (4) | C23—C24—C25—C26 | 54.9 (4) |
| C11—C12—C13—C18 | -66.2 (4) | C23—C24—C25—C28 | 177.0 (3) |
| C11—C12—C13—C17 | 168.4 (3) | C24—C25—C26—O26 | -58.8 (4) |
| C7—C8—C14—C13 | 180.0 (3) | C28—C25—C26—O26 | 179.0 (3) |

| | | | |
|-----------------|------------|-----------------|-----------|
| C9—C8—C14—C13 | 57.6 (4) | C25—C26—O26—C27 | 62.2 (4) |
| C7—C8—C14—C15 | -57.3 (4) | C26—O26—C27—C29 | 177.6 (3) |
| C9—C8—C14—C15 | -179.7 (3) | C26—O26—C27—C23 | -59.3 (4) |
| C12—C13—C14—C8 | -61.0 (4) | C22—C23—C27—O26 | 177.7 (3) |
| C18—C13—C14—C8 | 60.7 (4) | C24—C23—C27—O26 | 54.4 (4) |
| C17—C13—C14—C8 | 176.8 (3) | C22—C23—C27—C29 | -64.2 (4) |
| C12—C13—C14—C15 | 168.3 (3) | C24—C23—C27—C29 | 172.6 (3) |

Hydrogen-bond geometry (Å, °)

| <i>D</i> —H \cdots <i>A</i> | <i>D</i> —H | H \cdots <i>A</i> | <i>D</i> \cdots <i>A</i> | <i>D</i> —H \cdots <i>A</i> |
|----------------------------------|-------------|---------------------|----------------------------|-------------------------------|
| O3—H3A \cdots O26 ⁱ | 0.87 (4) | 2.11 (4) | 2.943 (4) | 159 (5) |

Symmetry code: (i) $x+1, y, z+1$.