

## A dimeric sesquiterpene, gochnatiolide A

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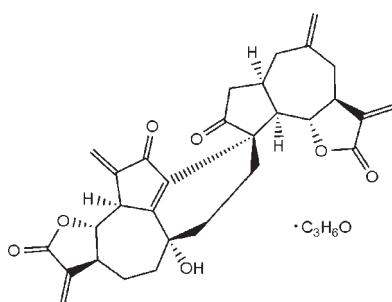
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Key indicators: single-crystal X-ray study;  $T = 294\text{ K}$ ; mean  $\sigma(\text{C}-\text{C}) = 0.006\text{ \AA}$ ; disorder in main residue;  $R$  factor = 0.047;  $wR$  factor = 0.119; data-to-parameter ratio = 7.1.

The title compound [systematic name: 5'a-hydroxy-1',3,6,8'-tetrakis(methylene)-3a,4,5,5',5'a,6,6',6a,7,7',7'a,8',9a,9b,10'a,-10'b-hexadecahydrospiro[azuleno[4,5-*b*]furan-9(2*H*),3'-[3*H*]-benz[1,8]azuleno[4,5-*b*]furan]-2,2',8,9'(1'*H*,3*H*,4'*H*)-tetron acetone 0.92-solvate],  $\text{C}_{30}\text{H}_{30}\text{O}_7 \cdot 0.92\text{C}_3\text{H}_6\text{O}$ , is a dimeric sesquiterpene formed by a cyclohexane system connecting two monomeric sesquiterpene lactone units of dehydro-zaluzanin C. It was isolated from *Ainsliaea henryi*.

## Related literature

For similar compounds and background information, see: *Chinese Materia Medica* (2007); Bohlmann & Zdero (1979); Bohlmann *et al.* (1981, 1982, 1983, 1984, 1986). For the pharmacological activity of a related compound, see: Wu *et al.* (2008).



## Experimental

## Crystal data

|   |  |
|---|--|
| $\text{C}_{30}\text{H}_{30}\text{O}_7 \cdot 0.92\text{C}_3\text{H}_6\text{O}$ | $V = 2853 (2)\text{ \AA}^3$              |
| $M_r = 555.94$  | $Z = 4$                                  |
| Orthorhombic, $P2_12_12_1$  | Mo $K\alpha$ radiation                   |
| $a = 8.709 (4)\text{ \AA}$  | $\mu = 0.09\text{ mm}^{-1}$              |
| $b = 12.652 (6)\text{ \AA}$   | $T = 294\text{ K}$                       |
| $c = 25.890 (12)\text{ \AA}$  | $0.15 \times 0.10 \times 0.08\text{ mm}$ |

## Data collection

|  |  |
|--|--|
| Bruker SMART APEX CCD area-detector diffractometer                   | 11789 measured reflections             |
| Absorption correction: multi-scan ( <i>SADABS</i> ; Sheldrick, 1996) | 2864 independent reflections           |
| $(SADABS$ ; Sheldrick, 1996)   | 1873 reflections with $I > 2\sigma(I)$ |
| $T_{\min} = 0.986$ , $T_{\max} = 0.993$                              | $R_{\text{int}} = 0.078$               |

## Refinement

|                                 |   |
|---------------------------------|---|
| $R[F^2 > 2\sigma(F^2)] = 0.047$ | 128 restraints                                |
| $wR(F^2) = 0.119$               | H-atom parameters constrained                 |
| $S = 0.93$                      | $\Delta\rho_{\max} = 0.32\text{ e \AA}^{-3}$  |
| 2864 reflections                | $\Delta\rho_{\min} = -0.12\text{ e \AA}^{-3}$ |
| 401 parameters                  |   |

Data collection: *SMART* (Bruker, 2005); cell refinement: *SAINT* (Bruker, 2005); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXL97*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: PK2191).

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# supporting information

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## A dimeric sesquiterpene, gochnatiolide A

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### S1. Comment

*Ainsliaea henryi* Diels is mainly distributed in south-west of China. The whole plant of *Ainsliaea henryi* has been used in Chinese folk medicine to treat cough, asthma and lumbago (Editorial committee of Chinese Mateia Medica, 2007). The chemical constituents of this plant have not been reported previously. Our chemical investigation of this plant for bioactive components resulted in the isolation of the title compound (I), which was previously obtained from the South American species, *Gochnatia paniculata* (Bohlmann *et al.*, 1983) and *Gochnatia polymorpha* (Bohlmann *et al.*, 1986). This kind of dimeric sesquiterpene has exhibited a remarkable inhibitory activity against the production of nitric oxide in RAW264.7 (Mouse leukaemic monocyte macrophage cell line) stimulated by LPS (Lipopolysaccharide) (Wu *et al.*, 2008).

The molecular struture of (I) is shown in Fig.1; bond lengths and angles are within normal ranges. Gochnatiolide A is a dimeric sequiterpene which was derived from two molecules of the compound dehydrozaluzanin C. The dehydrozaluzanin C molecule is composed of a seven-membered ring and two five-membered rings (A ring atoms C1—C5; B ring atoms C1—C10; C ring atoms C6/C7/C11/C12/O4). The connections between the two dehydrozaluzanin C molecules are the bonds C2—C24, and C10—C14—C35—C24. Ring A adopts an envelope conformation, ring B adopts a badly distorted chair conformation, while ring C exhibits an envelope conformation.

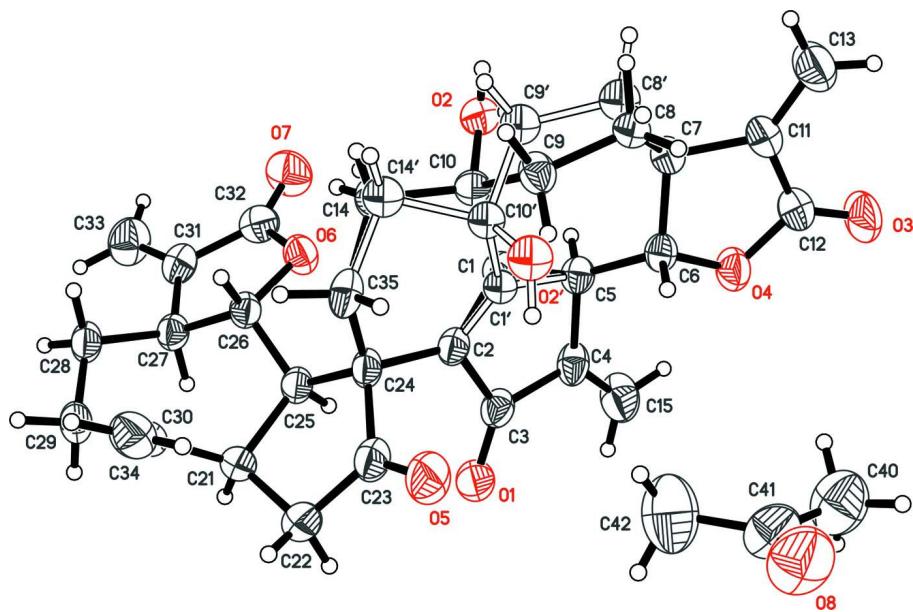
### S2. Experimental

Dry powders (5 kg) of the whole plant of *Ainsliaea henryi* were refluxed for 1 h with 95% ethanol (50L) three times. After removal of the ethanol under reduced pressure, the extract was suspended in water and then partitioned with petroleum ether, chloroform, ethyl acetate and n-butanol. The chloroform soluble fraction (30 g) was subjected to silica gel column chromatography using gradient elution (petroleum ether/acetone, 15:1 to 2:1, *v/v*). Gochnatiolide A was obtained from the fraction eluted by petroleum ether/acetone (5:1). Single crystals suitable for X-ray diffraction analysis were obtained by slow evaporation from petroleum ether/acetone (1:1) after two weeks at room temperature.

### S3. Refinement

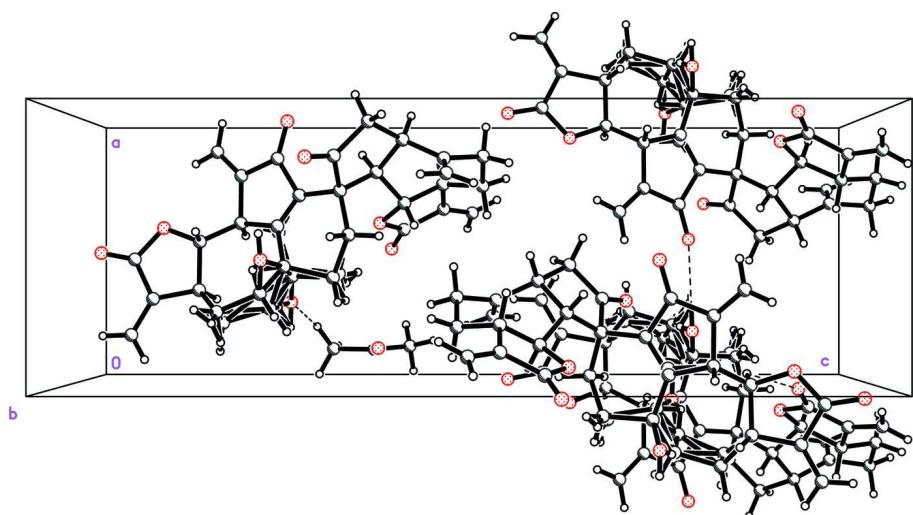
The hydroxyl H atoms attached at O2 was located in a difference Fourier map and isotropically refined using a riding model with O—H distance 0.82 Å. The remaining H atoms were placed in calculated positions with C—H distances in the range 0.93–0.98 Å. The  $U_{\text{iso}}$  values were set equal to  $1.5U_{\text{eq}}(\text{C})$  for methyl H atoms and  $1.5U_{\text{eq}}(\text{C})$  for the remaining H atoms. Friedel pairs were merged before the final refinement as there is no significant anomalous dispersion. As a consequence the absolute configuration of the compound is unknown. The stereochemistry of the title compound is known from the literature (Bohlmann *et al.* (1983)). Its structure was elucidated by highfield  $^1\text{H}$ -NMR spectroscopy. We have also confirmed its structure by  $^1\text{H}$ ,  $^{13}\text{C}$ , 2D NMR spectroscopy.

A number of restraints (128 in total) were required to ensure that the geometry (SADI and SAME in SHELXL-97) and displacement parameters (SIMU in SHELXL-97) retained chemically and physically reasonable values during the refinement.



**Figure 1**

The molecular structure of (I), showing the atom-labelling scheme with displacement ellipsoids drawn at the 30% probability level. H atoms are shown as small spheres of arbitrary radius.



**Figure 2**

The molecular packing of (I), viewed along the  $a$  axis.

5'a-hydroxy-1',3,6,8'-tetrakis(methylene)- 3a,4,5,5',5'a,6,6',6a,7,7',7'a,8',9a,9b,10'a,10'b-hexadecahydro  
spiro[azuleno[4,5-*b*]furan-9(2*H*),3'- [3*H*]benz[1,8]azuleno[4,5-*b*]furan]- 2,2',8,9'(1'*H*,3*H*,4'*H*)-tetrone acetone  
0.92-solvate

#### Crystal data

$C_{30}H_{30}O_7 \cdot 0.92C_3H_6O$

$M_r = 555.94$

Orthorhombic,  $P2_12_12_1$

Hall symbol: P 2ac 2ab

$a = 8.709 (4) \text{ \AA}$

$b = 12.652 (6) \text{ \AA}$

$c = 25.890 (12) \text{ \AA}$

$V = 2853 (2) \text{ \AA}^3$

$Z = 4$

$F(000) = 1182$

$D_x = 1.294 \text{ Mg m}^{-3}$

Mo  $K\alpha$  radiation,  $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 1001 reflections

$\theta = 2.5\text{--}24.3^\circ$

$\mu = 0.09 \text{ mm}^{-1}$

$T = 294 \text{ K}$

Block, colourless

$0.15 \times 0.10 \times 0.08 \text{ mm}$

#### Data collection

Bruker SMART APEX CCD area-detector  
diffractometer

Radiation source: sealed tube

Graphite monochromator

$\varphi$  and  $\omega$  scans

Absorption correction: multi-scan  
(*SADABS*; Sheldrick, 1996)

$T_{\min} = 0.986$ ,  $T_{\max} = 0.993$

11789 measured reflections

2864 independent reflections

1873 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.078$

$\theta_{\max} = 25.0^\circ$ ,  $\theta_{\min} = 1.6^\circ$

$h = -7 \rightarrow 10$

$k = -15 \rightarrow 14$

$l = -30 \rightarrow 30$

#### Refinement

Refinement on  $F^2$

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.047$

$wR(F^2) = 0.119$

$S = 0.93$

2864 reflections

401 parameters

128 restraints

0 constraints

Primary atom site location: structure-invariant  
direct methods

Secondary atom site location: difference Fourier  
map

Hydrogen site location: inferred from  
neighbouring sites

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0673P)^2]$

where  $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} < 0.001$

$\Delta\rho_{\max} = 0.32 \text{ e \AA}^{-3}$

$\Delta\rho_{\min} = -0.12 \text{ e \AA}^{-3}$

Extinction correction: *SHELXL*,  
 $F_c^* = kFc[1 + 0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$

Extinction coefficient: 0.0034 (11)

#### Special details

**Geometry.** All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

**Refinement.** Refinement of  $F^2$  against ALL reflections. The weighted  $R$ -factor  $wR$  and goodness of fit  $S$  are based on  $F^2$ , conventional  $R$ -factors  $R$  are based on  $F$ , with  $F$  set to zero for negative  $F^2$ . The threshold expression of  $F^2 > 2\sigma(F^2)$  is used only for calculating  $R$ -factors(gt) etc. and is not relevant to the choice of reflections for refinement.  $R$ -factors based on  $F^2$  are statistically about twice as large as those based on  $F$ , and  $R$ -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters ( $\text{\AA}^2$ )

|      | <i>x</i>     | <i>y</i>    | <i>z</i>     | $U_{\text{iso}}^*/U_{\text{eq}}$ | Occ. (<1) |
|------|--------------|-------------|--------------|----------------------------------|-----------|
| O1   | 0.4589 (3)   | 0.8430 (3)  | 0.72146 (11) | 0.0859 (10)                      |           |
| O3   | -0.0138 (4)  | 0.9219 (3)  | 0.95361 (11) | 0.1010 (12)                      |           |
| O4   | 0.0768 (3)   | 0.8957 (2)  | 0.87468 (10) | 0.0704 (8)                       |           |
| O5   | 0.3123 (4)   | 0.6000 (3)  | 0.68984 (12) | 0.0936 (10)                      |           |
| O6   | 0.0973 (3)   | 0.9794 (2)  | 0.61271 (10) | 0.0648 (8)                       |           |
| O7   | 0.0044 (4)   | 1.1419 (3)  | 0.60213 (13) | 0.0963 (11)                      |           |
| C2   | 0.1914 (4)   | 0.8169 (3)  | 0.70590 (13) | 0.0511 (9)                       |           |
| C3   | 0.3257 (5)   | 0.8549 (3)  | 0.73401 (15) | 0.0609 (10)                      |           |
| C4   | 0.2677 (5)   | 0.9100 (3)  | 0.78106 (14) | 0.0609 (11)                      |           |
| C5   | 0.0944 (4)   | 0.9005 (3)  | 0.78111 (13) | 0.0523 (9)                       |           |
| H5   | 0.0503       | 0.9716      | 0.7791       | 0.063*                           |           |
| C6   | 0.0263 (4)   | 0.8446 (3)  | 0.82731 (13) | 0.0510 (9)                       |           |
| H6   | 0.0602       | 0.7707      | 0.8274       | 0.061*                           |           |
| C7   | -0.1479 (5)  | 0.8484 (4)  | 0.82852 (14) | 0.0612 (10)                      |           |
| H7   | -0.1816      | 0.9131      | 0.8111       | 0.073*                           |           |
| C11  | -0.1797 (5)  | 0.8596 (4)  | 0.88515 (15) | 0.0695 (12)                      |           |
| C12  | -0.0371 (5)  | 0.8938 (4)  | 0.91013 (17) | 0.0737 (12)                      |           |
| C13  | -0.3092 (7)  | 0.8448 (5)  | 0.91028 (19) | 0.0967 (16)                      |           |
| H13A | -0.3133      | 0.8562      | 0.9457       | 0.116*                           |           |
| H13B | -0.3965      | 0.8230      | 0.8926       | 0.116*                           |           |
| C8   | -0.2239 (6)  | 0.7527 (4)  | 0.80139 (18) | 0.0570 (14)                      | 0.808 (5) |
| H8A  | -0.2150      | 0.6919      | 0.8240       | 0.068*                           | 0.808 (5) |
| H8B  | -0.3325      | 0.7677      | 0.7974       | 0.068*                           | 0.808 (5) |
| C9   | -0.1589 (6)  | 0.7225 (4)  | 0.74833 (17) | 0.0584 (13)                      | 0.808 (5) |
| H9A  | -0.2389      | 0.6861      | 0.7292       | 0.070*                           | 0.808 (5) |
| H9B  | -0.0761      | 0.6724      | 0.7537       | 0.070*                           | 0.808 (5) |
| C10  | -0.0979 (5)  | 0.8128 (4)  | 0.71442 (17) | 0.0512 (11)                      | 0.808 (5) |
| O2   | -0.1932 (4)  | 0.9031 (3)  | 0.71949 (13) | 0.0660 (10)                      | 0.808 (5) |
| H2   | -0.2835      | 0.8847      | 0.7197       | 0.099*                           | 0.808 (5) |
| C14  | -0.0889 (5)  | 0.7804 (5)  | 0.65696 (19) | 0.0609 (17)                      | 0.808 (5) |
| H14A | -0.1830      | 0.7440      | 0.6478       | 0.073*                           | 0.808 (5) |
| H14B | -0.0833      | 0.8440      | 0.6361       | 0.073*                           | 0.808 (5) |
| C1   | 0.0602 (6)   | 0.8438 (4)  | 0.7319 (3)   | 0.0468 (15)                      | 0.808 (5) |
| C8'  | -0.263 (2)   | 0.7876 (19) | 0.7957 (6)   | 0.062 (5)*                       | 0.192 (5) |
| H8'1 | -0.2629      | 0.7133      | 0.8049       | 0.075*                           | 0.192 (5) |
| H8'2 | -0.3661      | 0.8153      | 0.8008       | 0.075*                           | 0.192 (5) |
| C9'  | -0.214 (2)   | 0.8023 (17) | 0.7405 (7)   | 0.060 (3)*                       | 0.192 (5) |
| H9'1 | -0.2103      | 0.8778      | 0.7340       | 0.073*                           | 0.192 (5) |
| H9'2 | -0.2944      | 0.7736      | 0.7188       | 0.073*                           | 0.192 (5) |
| C10' | -0.0634 (19) | 0.7563 (13) | 0.7216 (7)   | 0.051 (3)*                       | 0.192 (5) |
| O2'  | -0.0418 (17) | 0.6540 (11) | 0.7438 (6)   | 0.070 (4)*                       | 0.192 (5) |
| H2'  | 0.0503       | 0.6411      | 0.7456       | 0.106*                           | 0.192 (5) |
| C14' | -0.102 (2)   | 0.754 (3)   | 0.6631 (7)   | 0.062 (4)*                       | 0.192 (5) |
| H14C | -0.1885      | 0.7082      | 0.6555       | 0.074*                           | 0.192 (5) |
| H14D | -0.1220      | 0.8243      | 0.6495       | 0.074*                           | 0.192 (5) |

|      |             |            |              |             |           |
|------|-------------|------------|--------------|-------------|-----------|
| C1'  | 0.081 (3)   | 0.817 (2)  | 0.7323 (14)  | 0.045 (4)*  | 0.192 (5) |
| C15  | 0.3579 (6)  | 0.9603 (4) | 0.81350 (18) | 0.0902 (17) |           |
| H15A | 0.4632      | 0.9626     | 0.8076       | 0.108*      |           |
| H15B | 0.3165      | 0.9935     | 0.8423       | 0.108*      |           |
| C21  | 0.3861 (4)  | 0.7640 (3) | 0.58050 (14) | 0.0600 (10) |           |
| H21  | 0.4744      | 0.8079     | 0.5709       | 0.072*      |           |
| C22  | 0.4426 (5)  | 0.6809 (4) | 0.61919 (16) | 0.0775 (13) |           |
| H22A | 0.4597      | 0.6139     | 0.6019       | 0.093*      |           |
| H22B | 0.5383      | 0.7034     | 0.6349       | 0.093*      |           |
| C23  | 0.3214 (5)  | 0.6699 (4) | 0.65901 (15) | 0.0693 (11) |           |
| C24  | 0.2052 (4)  | 0.7607 (3) | 0.65507 (13) | 0.0519 (9)  |           |
| C25  | 0.2759 (4)  | 0.8337 (3) | 0.61335 (13) | 0.0511 (9)  |           |
| H25  | 0.3408      | 0.8844     | 0.6318       | 0.061*      |           |
| C26  | 0.1675 (4)  | 0.8979 (3) | 0.58093 (12) | 0.0509 (9)  |           |
| H26  | 0.0862      | 0.8510     | 0.5682       | 0.061*      |           |
| C27  | 0.2387 (5)  | 0.9549 (3) | 0.53460 (13) | 0.0558 (10) |           |
| H27  | 0.3469      | 0.9696     | 0.5422       | 0.067*      |           |
| C28  | 0.2292 (5)  | 0.8915 (3) | 0.48538 (14) | 0.0612 (10) |           |
| H28A | 0.2586      | 0.9365     | 0.4567       | 0.073*      |           |
| H28B | 0.1232      | 0.8706     | 0.4800       | 0.073*      |           |
| C29  | 0.3292 (5)  | 0.7929 (4) | 0.48453 (14) | 0.0665 (11) |           |
| H29A | 0.3035      | 0.7524     | 0.4539       | 0.080*      |           |
| H29B | 0.4355      | 0.8149     | 0.4812       | 0.080*      |           |
| C30  | 0.3170 (4)  | 0.7206 (4) | 0.53086 (14) | 0.0585 (10) |           |
| C31  | 0.1522 (5)  | 1.0569 (3) | 0.53462 (16) | 0.0658 (11) |           |
| C32  | 0.0762 (5)  | 1.0678 (4) | 0.58487 (18) | 0.0683 (12) |           |
| C33  | 0.1407 (7)  | 1.1302 (4) | 0.49896 (19) | 0.0936 (16) |           |
| H33A | 0.0833      | 1.1908     | 0.5053       | 0.112*      |           |
| H33B | 0.1899      | 1.1215     | 0.4674       | 0.112*      |           |
| C34  | 0.2586 (5)  | 0.6260 (4) | 0.5270 (2)   | 0.0837 (14) |           |
| H34A | 0.2233      | 0.6017     | 0.4953       | 0.100*      |           |
| H34B | 0.2522      | 0.5829     | 0.5561       | 0.100*      |           |
| C35  | 0.0480 (5)  | 0.7089 (3) | 0.64304 (13) | 0.0637 (11) |           |
| H35A | 0.0434      | 0.6920     | 0.6065       | 0.076*      |           |
| H35B | 0.0397      | 0.6433     | 0.6622       | 0.076*      |           |
| O8   | 0.3631 (7)  | 0.4623 (4) | 0.8854 (2)   | 0.146 (2)   | 0.919 (7) |
| C40  | 0.3646 (10) | 0.6279 (6) | 0.9215 (3)   | 0.133 (3)   | 0.919 (7) |
| H40A | 0.3417      | 0.5887     | 0.9523       | 0.199*      | 0.919 (7) |
| H40B | 0.2852      | 0.6791     | 0.9155       | 0.199*      | 0.919 (7) |
| H40C | 0.4612      | 0.6634     | 0.9255       | 0.199*      | 0.919 (7) |
| C41  | 0.3730 (8)  | 0.5566 (5) | 0.8781 (3)   | 0.104 (2)   | 0.919 (7) |
| C42  | 0.3737 (11) | 0.6047 (7) | 0.8249 (3)   | 0.153 (3)   | 0.919 (7) |
| H42A | 0.4679      | 0.5870     | 0.8077       | 0.230*      | 0.919 (7) |
| H42B | 0.3649      | 0.6801     | 0.8276       | 0.230*      | 0.919 (7) |
| H42C | 0.2886      | 0.5775     | 0.8054       | 0.230*      | 0.919 (7) |

Atomic displacement parameters ( $\text{\AA}^2$ )

|     | $U^{11}$    | $U^{22}$    | $U^{33}$    | $U^{12}$     | $U^{13}$     | $U^{23}$     |
|-----|-------------|-------------|-------------|--------------|--------------|--------------|
| O1  | 0.0530 (19) | 0.142 (3)   | 0.0630 (18) | -0.0116 (19) | -0.0036 (14) | -0.0044 (19) |
| O3  | 0.105 (3)   | 0.149 (3)   | 0.0488 (18) | -0.008 (2)   | 0.0050 (17)  | -0.024 (2)   |
| O4  | 0.0734 (18) | 0.090 (2)   | 0.0480 (15) | -0.0099 (17) | -0.0003 (14) | -0.0065 (15) |
| O5  | 0.120 (3)   | 0.088 (2)   | 0.072 (2)   | 0.024 (2)    | 0.009 (2)    | 0.0258 (19)  |
| O6  | 0.0758 (18) | 0.0645 (17) | 0.0542 (16) | 0.0076 (15)  | 0.0078 (14)  | -0.0041 (15) |
| O7  | 0.114 (3)   | 0.071 (2)   | 0.104 (2)   | 0.021 (2)    | 0.007 (2)    | -0.0157 (19) |
| C2  | 0.055 (2)   | 0.057 (2)   | 0.041 (2)   | -0.0060 (19) | -0.0039 (18) | 0.0063 (17)  |
| C3  | 0.052 (2)   | 0.083 (3)   | 0.048 (2)   | -0.016 (2)   | -0.006 (2)   | 0.008 (2)    |
| C4  | 0.069 (3)   | 0.070 (3)   | 0.044 (2)   | -0.024 (2)   | -0.0062 (19) | 0.000 (2)    |
| C5  | 0.058 (2)   | 0.051 (2)   | 0.048 (2)   | -0.0062 (18) | -0.0026 (18) | -0.0016 (18) |
| C6  | 0.061 (2)   | 0.049 (2)   | 0.043 (2)   | -0.0018 (18) | -0.0023 (18) | -0.0045 (17) |
| C7  | 0.059 (2)   | 0.076 (3)   | 0.049 (2)   | 0.006 (2)    | 0.0055 (19)  | 0.002 (2)    |
| C11 | 0.068 (3)   | 0.088 (3)   | 0.053 (2)   | 0.003 (3)    | 0.012 (2)    | -0.003 (2)   |
| C12 | 0.078 (3)   | 0.088 (3)   | 0.055 (3)   | 0.006 (3)    | 0.010 (2)    | -0.006 (2)   |
| C13 | 0.091 (4)   | 0.128 (4)   | 0.071 (3)   | -0.009 (3)   | 0.017 (3)    | -0.014 (3)   |
| C8  | 0.044 (3)   | 0.074 (3)   | 0.053 (3)   | -0.002 (2)   | 0.002 (2)    | 0.002 (2)    |
| C9  | 0.059 (3)   | 0.065 (3)   | 0.051 (2)   | -0.013 (2)   | 0.004 (2)    | -0.004 (2)   |
| C10 | 0.047 (2)   | 0.056 (2)   | 0.051 (2)   | -0.003 (2)   | -0.0003 (19) | 0.000 (2)    |
| O2  | 0.060 (2)   | 0.076 (2)   | 0.062 (2)   | 0.0098 (19)  | -0.0026 (18) | -0.0013 (19) |
| C14 | 0.051 (3)   | 0.084 (4)   | 0.048 (3)   | -0.024 (3)   | 0.003 (2)    | -0.002 (3)   |
| C1  | 0.049 (3)   | 0.047 (3)   | 0.044 (2)   | -0.003 (3)   | -0.004 (2)   | 0.005 (3)    |
| C15 | 0.085 (3)   | 0.119 (4)   | 0.067 (3)   | -0.038 (3)   | 0.000 (3)    | -0.014 (3)   |
| C21 | 0.049 (2)   | 0.078 (3)   | 0.053 (2)   | 0.008 (2)    | 0.0035 (18)  | 0.002 (2)    |
| C22 | 0.065 (3)   | 0.106 (4)   | 0.061 (3)   | 0.016 (3)    | 0.000 (2)    | 0.014 (2)    |
| C23 | 0.082 (3)   | 0.077 (3)   | 0.048 (2)   | 0.012 (3)    | -0.005 (2)   | 0.008 (2)    |
| C24 | 0.059 (2)   | 0.062 (2)   | 0.0348 (18) | -0.0041 (19) | -0.0005 (17) | 0.0052 (17)  |
| C25 | 0.052 (2)   | 0.060 (2)   | 0.041 (2)   | -0.0039 (18) | -0.0013 (17) | 0.0007 (18)  |
| C26 | 0.052 (2)   | 0.058 (2)   | 0.0423 (19) | -0.006 (2)   | -0.0009 (17) | -0.0030 (18) |
| C27 | 0.059 (2)   | 0.064 (2)   | 0.044 (2)   | -0.0039 (19) | -0.0001 (18) | 0.0052 (19)  |
| C28 | 0.066 (3)   | 0.078 (3)   | 0.039 (2)   | -0.008 (2)   | -0.0010 (18) | 0.000 (2)    |
| C29 | 0.068 (3)   | 0.089 (3)   | 0.042 (2)   | -0.002 (3)   | 0.002 (2)    | -0.007 (2)   |
| C30 | 0.052 (2)   | 0.070 (3)   | 0.053 (2)   | 0.009 (2)    | 0.0046 (19)  | -0.006 (2)   |
| C31 | 0.074 (3)   | 0.067 (3)   | 0.056 (3)   | -0.007 (2)   | -0.010 (2)   | 0.008 (2)    |
| C32 | 0.067 (3)   | 0.062 (3)   | 0.076 (3)   | 0.001 (2)    | -0.007 (2)   | -0.007 (2)   |
| C33 | 0.122 (4)   | 0.074 (3)   | 0.084 (3)   | 0.009 (3)    | -0.005 (3)   | 0.017 (3)    |
| C34 | 0.085 (3)   | 0.083 (3)   | 0.083 (3)   | 0.010 (3)    | 0.007 (3)    | -0.014 (3)   |
| C35 | 0.081 (3)   | 0.067 (3)   | 0.043 (2)   | -0.018 (2)   | -0.010 (2)   | 0.0013 (19)  |
| O8  | 0.191 (6)   | 0.105 (4)   | 0.143 (4)   | 0.033 (4)    | -0.016 (4)   | -0.010 (3)   |
| C40 | 0.156 (7)   | 0.126 (6)   | 0.117 (5)   | -0.007 (5)   | -0.026 (5)   | -0.016 (5)   |
| C41 | 0.131 (6)   | 0.074 (4)   | 0.106 (5)   | 0.024 (4)    | -0.027 (4)   | -0.022 (4)   |
| C42 | 0.208 (9)   | 0.138 (7)   | 0.114 (5)   | -0.043 (7)   | 0.002 (6)    | -0.016 (5)   |

Geometric parameters ( $\text{\AA}$ ,  $\text{^{\circ}}$ )

|          |            |           |            |
|----------|------------|-----------|------------|
| O1—C3    | 1.214 (5)  | C10'—C1'  | 1.499 (10) |
| O3—C12   | 1.198 (5)  | C10'—C14' | 1.554 (17) |
| O4—C12   | 1.351 (5)  | O2'—H2'   | 0.8200     |
| O4—C6    | 1.454 (4)  | C14'—C35  | 1.518 (11) |
| O5—C23   | 1.193 (5)  | C14'—H14C | 0.9700     |
| O6—C32   | 1.344 (5)  | C14'—H14D | 0.9700     |
| O6—C26   | 1.454 (4)  | C15—H15A  | 0.9300     |
| O7—C32   | 1.212 (5)  | C15—H15B  | 0.9300     |
| C2—C1'   | 1.18 (3)   | C21—C30   | 1.522 (5)  |
| C2—C1    | 1.370 (8)  | C21—C22   | 1.533 (6)  |
| C2—C3    | 1.459 (5)  | C21—C25   | 1.556 (5)  |
| C2—C24   | 1.500 (5)  | C21—H21   | 0.9800     |
| C3—C4    | 1.492 (6)  | C22—C23   | 1.482 (6)  |
| C4—C15   | 1.315 (5)  | C22—H22A  | 0.9700     |
| C4—C5    | 1.514 (5)  | C22—H22B  | 0.9700     |
| C5—C1    | 1.491 (8)  | C23—C24   | 1.535 (6)  |
| C5—C6    | 1.511 (5)  | C24—C25   | 1.548 (5)  |
| C5—C1'   | 1.65 (3)   | C24—C35   | 1.549 (5)  |
| C5—H5    | 0.9800     | C25—C26   | 1.502 (5)  |
| C6—C7    | 1.518 (5)  | C25—H25   | 0.9800     |
| C6—H6    | 0.9800     | C26—C27   | 1.531 (5)  |
| C7—C11   | 1.499 (5)  | C26—H26   | 0.9800     |
| C7—C8'   | 1.525 (11) | C27—C31   | 1.494 (6)  |
| C7—C8    | 1.548 (6)  | C27—C28   | 1.509 (5)  |
| C7—H7    | 0.9800     | C27—H27   | 0.9800     |
| C11—C13  | 1.315 (6)  | C28—C29   | 1.522 (6)  |
| C11—C12  | 1.465 (7)  | C28—H28A  | 0.9700     |
| C13—H13A | 0.9300     | C28—H28B  | 0.9700     |
| C13—H13B | 0.9300     | C29—C30   | 1.512 (5)  |
| C8—C9    | 1.534 (6)  | C29—H29A  | 0.9700     |
| C8—H8A   | 0.9700     | C29—H29B  | 0.9700     |
| C8—H8B   | 0.9700     | C30—C34   | 1.304 (6)  |
| C9—C10   | 1.536 (6)  | C31—C33   | 1.312 (6)  |
| C9—H9A   | 0.9700     | C31—C32   | 1.466 (6)  |
| C9—H9B   | 0.9700     | C33—H33A  | 0.9300     |
| C10—O2   | 1.418 (6)  | C33—H33B  | 0.9300     |
| C10—C1   | 1.502 (6)  | C34—H34A  | 0.9300     |
| C10—C14  | 1.545 (6)  | C34—H34B  | 0.9300     |
| O2—H2    | 0.8200     | C35—H35A  | 0.9700     |
| C14—C35  | 1.539 (6)  | C35—H35B  | 0.9700     |
| C14—H14A | 0.9700     | O8—C41    | 1.211 (7)  |
| C14—H14B | 0.9700     | C40—C41   | 1.441 (8)  |
| C8'—C9'  | 1.505 (17) | C40—H40A  | 0.9600     |
| C8'—H8'1 | 0.9700     | C40—H40B  | 0.9600     |
| C8'—H8'2 | 0.9700     | C40—H40C  | 0.9600     |
| C9'—C10' | 1.513 (16) | C41—C42   | 1.507 (9)  |

|              |            |                |            |
|--------------|------------|----------------|------------|
| C9'—H9'1     | 0.9700     | C42—H42A       | 0.9600     |
| C9'—H9'2     | 0.9700     | C42—H42B       | 0.9600     |
| C10'—O2'     | 1.427 (15) | C42—H42C       | 0.9600     |
|              |            |                |            |
| C12—O4—C6    | 110.1 (3)  | H14C—C14'—H14D | 109.7      |
| C32—O6—C26   | 110.2 (3)  | C2—C1'—C10'    | 125 (2)    |
| C1'—C2—C3    | 111.3 (13) | C2—C1'—C5      | 112.8 (12) |
| C1—C2—C3     | 109.9 (4)  | C10'—C1'—C5    | 122 (2)    |
| C1'—C2—C24   | 125.1 (13) | C4—C15—H15A    | 120.0      |
| C1—C2—C24    | 128.1 (4)  | C4—C15—H15B    | 120.0      |
| C3—C2—C24    | 121.9 (3)  | H15A—C15—H15B  | 120.0      |
| O1—C3—C2     | 126.3 (4)  | C30—C21—C22    | 115.5 (4)  |
| O1—C3—C4     | 126.9 (4)  | C30—C21—C25    | 114.9 (3)  |
| C2—C3—C4     | 106.8 (3)  | C22—C21—C25    | 103.3 (3)  |
| C15—C4—C3    | 123.0 (4)  | C30—C21—H21    | 107.6      |
| C15—C4—C5    | 129.3 (4)  | C22—C21—H21    | 107.6      |
| C3—C4—C5     | 107.6 (3)  | C25—C21—H21    | 107.6      |
| C1—C5—C6     | 111.9 (3)  | C23—C22—C21    | 106.9 (3)  |
| C1—C5—C4     | 103.7 (4)  | C23—C22—H22A   | 110.3      |
| C6—C5—C4     | 115.4 (3)  | C21—C22—H22A   | 110.3      |
| C6—C5—C1'    | 106.2 (11) | C23—C22—H22B   | 110.3      |
| C4—C5—C1'    | 97.0 (8)   | C21—C22—H22B   | 110.3      |
| C1—C5—H5     | 108.5      | H22A—C22—H22B  | 108.6      |
| C6—C5—H5     | 108.5      | O5—C23—C22     | 125.6 (4)  |
| C4—C5—H5     | 108.5      | O5—C23—C24     | 123.7 (4)  |
| C1'—C5—H5    | 121.2      | C22—C23—C24    | 110.7 (4)  |
| O4—C6—C5     | 109.9 (3)  | C2—C24—C23     | 110.4 (3)  |
| O4—C6—C7     | 105.7 (3)  | C2—C24—C25     | 111.2 (3)  |
| C5—C6—C7     | 113.2 (3)  | C23—C24—C25    | 103.3 (3)  |
| O4—C6—H6     | 109.3      | C2—C24—C35     | 107.8 (3)  |
| C5—C6—H6     | 109.3      | C23—C24—C35    | 106.2 (3)  |
| C7—C6—H6     | 109.3      | C25—C24—C35    | 117.6 (3)  |
| C11—C7—C6    | 102.0 (3)  | C26—C25—C24    | 117.6 (3)  |
| C11—C7—C8'   | 118.1 (8)  | C26—C25—C21    | 112.9 (3)  |
| C6—C7—C8'    | 129.2 (10) | C24—C25—C21    | 106.8 (3)  |
| C11—C7—C8    | 116.0 (4)  | C26—C25—H25    | 106.3      |
| C6—C7—C8     | 113.1 (4)  | C24—C25—H25    | 106.3      |
| C11—C7—H7    | 108.4      | C21—C25—H25    | 106.3      |
| C6—C7—H7     | 108.4      | O6—C26—C25     | 109.4 (3)  |
| C8'—C7—H7    | 88.1       | O6—C26—C27     | 106.2 (3)  |
| C8—C7—H7     | 108.4      | C25—C26—C27    | 116.0 (3)  |
| C13—C11—C12  | 123.4 (4)  | O6—C26—H26     | 108.4      |
| C13—C11—C7   | 129.0 (4)  | C25—C26—H26    | 108.4      |
| C12—C11—C7   | 107.6 (4)  | C27—C26—H26    | 108.4      |
| O3—C12—O4    | 120.6 (4)  | C31—C27—C28    | 115.6 (3)  |
| O3—C12—C11   | 130.2 (4)  | C31—C27—C26    | 101.7 (3)  |
| O4—C12—C11   | 109.1 (4)  | C28—C27—C26    | 112.9 (3)  |
| C11—C13—H13A | 120.0      | C31—C27—H27    | 108.8      |

|               |            |               |            |
|---------------|------------|---------------|------------|
| C11—C13—H13B  | 120.0      | C28—C27—H27   | 108.8      |
| H13A—C13—H13B | 120.0      | C26—C27—H27   | 108.8      |
| C9—C8—C7      | 116.3 (4)  | C27—C28—C29   | 114.6 (3)  |
| C9—C8—H8A     | 108.2      | C27—C28—H28A  | 108.6      |
| C7—C8—H8A     | 108.2      | C29—C28—H28A  | 108.6      |
| C9—C8—H8B     | 108.2      | C27—C28—H28B  | 108.6      |
| C7—C8—H8B     | 108.2      | C29—C28—H28B  | 108.6      |
| H8A—C8—H8B    | 107.4      | H28A—C28—H28B | 107.6      |
| C8—C9—C10     | 117.0 (4)  | C30—C29—C28   | 116.4 (3)  |
| C8—C9—H9A     | 108.1      | C30—C29—H29A  | 108.2      |
| C10—C9—H9A    | 108.1      | C28—C29—H29A  | 108.2      |
| C8—C9—H9B     | 108.1      | C30—C29—H29B  | 108.2      |
| C10—C9—H9B    | 108.1      | C28—C29—H29B  | 108.2      |
| H9A—C9—H9B    | 107.3      | H29A—C29—H29B | 107.3      |
| O2—C10—C1     | 107.3 (4)  | C34—C30—C29   | 121.5 (4)  |
| O2—C10—C9     | 110.1 (4)  | C34—C30—C21   | 123.4 (4)  |
| C1—C10—C9     | 109.8 (4)  | C29—C30—C21   | 115.0 (4)  |
| O2—C10—C14    | 109.4 (4)  | C33—C31—C32   | 121.5 (4)  |
| C1—C10—C14    | 108.3 (4)  | C33—C31—C27   | 130.4 (4)  |
| C9—C10—C14    | 111.8 (4)  | C32—C31—C27   | 108.0 (3)  |
| C35—C14—C10   | 114.8 (4)  | O7—C32—O6     | 121.1 (4)  |
| C35—C14—H14A  | 108.6      | O7—C32—C31    | 129.3 (4)  |
| C10—C14—H14A  | 108.6      | O6—C32—C31    | 109.6 (4)  |
| C35—C14—H14B  | 108.6      | C31—C33—H33A  | 120.0      |
| C10—C14—H14B  | 108.6      | C31—C33—H33B  | 120.0      |
| H14A—C14—H14B | 107.5      | H33A—C33—H33B | 120.0      |
| C2—C1—C5      | 111.9 (4)  | C30—C34—H34A  | 120.0      |
| C2—C1—C10     | 123.4 (5)  | C30—C34—H34B  | 120.0      |
| C5—C1—C10     | 124.5 (5)  | H34A—C34—H34B | 120.0      |
| C9'—C8'—C7    | 106.1 (13) | C14'—C35—C24  | 122.2 (12) |
| C9'—C8'—H8'1  | 110.5      | C14—C35—C24   | 112.9 (4)  |
| C7—C8'—H8'1   | 110.5      | C14'—C35—H35A | 112.3      |
| C9'—C8'—H8'2  | 110.5      | C14—C35—H35A  | 109.0      |
| C7—C8'—H8'2   | 110.5      | C24—C35—H35A  | 109.0      |
| H8'1—C8'—H8'2 | 108.7      | C14'—C35—H35B | 94.8       |
| C8'—C9'—C10'  | 120.5 (16) | C14—C35—H35B  | 109.0      |
| C8'—C9'—H9'1  | 107.2      | C24—C35—H35B  | 109.0      |
| C10'—C9'—H9'1 | 107.2      | H35A—C35—H35B | 107.8      |
| C8'—C9'—H9'2  | 107.2      | C41—C40—H40A  | 109.5      |
| C10'—C9'—H9'2 | 107.2      | C41—C40—H40B  | 109.5      |
| H9'1—C9'—H9'2 | 106.8      | H40A—C40—H40B | 109.5      |
| O2'—C10'—C1'  | 106.4 (15) | C41—C40—H40C  | 109.5      |
| O2'—C10'—C9'  | 109.5 (14) | H40A—C40—H40C | 109.5      |
| C1'—C10'—C9'  | 117.9 (17) | H40B—C40—H40C | 109.5      |
| O2'—C10'—C14' | 113.8 (16) | O8—C41—C40    | 119.4 (7)  |
| C1'—C10'—C14' | 111.8 (18) | O8—C41—C42    | 122.7 (6)  |
| C9'—C10'—C14' | 97.7 (11)  | C40—C41—C42   | 117.4 (6)  |
| C10'—O2'—H2'  | 109.5      | C41—C42—H42A  | 109.5      |

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|                |           |               |       |
|----------------|-----------|---------------|-------|
| C35—C14'—C10'  | 98.8 (11) | C41—C42—H42B  | 109.5 |
| C35—C14'—H14C  | 112.0     | H42A—C42—H42B | 109.5 |
| C10'—C14'—H14C | 112.0     | C41—C42—H42C  | 109.5 |
| C35—C14'—H14D  | 112.0     | H42A—C42—H42C | 109.5 |
| C10'—C14'—H14D | 112.0     | H42B—C42—H42C | 109.5 |

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