

Acta Crystallographica Section E

## Structure Reports

Online

ISSN 1600-5368

# (Z)-3 $\alpha$ -(1,3-Dioxoisindolin-2-yl)-17(20)-pregnene

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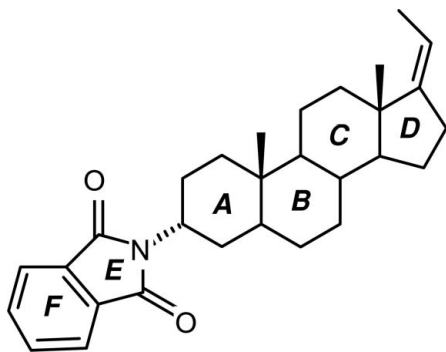
Received 21 June 2011; accepted 11 July 2011

 Key indicators: single-crystal X-ray study;  $T = 113$  K; mean  $\sigma(\text{C}-\text{C}) = 0.003$  Å;  $R$  factor = 0.039;  $wR$  factor = 0.096; data-to-parameter ratio = 9.7.

The title compound,  $\text{C}_{29}\text{H}_{37}\text{NO}_2$ , crystallized with two independent molecules in an asymmetric unit in which the conformation of the cyclohexyl ring of the pregnene moiety bonded to the 3 $\alpha$ -(1,3-dioxoisindolin-2-yl)- ring system differs: in one molecule it is in a chair conformation, while in the other it exhibits a half-chair conformation. The other six-membered rings in the pregnene moiety are in chair conformations and the five-membered rings are in envelope forms in both molecules. In both molecules, the 3 $\alpha$ -(1,3-dioxoisindolin-2-yl)- ring systems are individually approximately planar, with r.m.s. deviations 0.0148 and 0.0264 Å. The structure is consolidated by intermolecular C—H $\cdots$ O hydrogen-bonding interactions involving the carbonyl O atoms and methyl, methylene and methyldyne groups, resulting in a two-dimensional structure.

## Related literature

The title compound was synthesized from epiandrosterone, a pregnane alkaloid isolated from *Pachysandra axillaris*, a traditional Chinese medicine. For the biological activity of *Pachysandra axillaris*, see: Sun *et al.* (2010). For the synthesis of the title compound, see: Batcho *et al.* (1981). For a related structure, see: Ishida *et al.* (1981). For the absolute structure, see: Pollard & Ahmed (1971).



## Experimental

## Crystal data

|   |                                   |
|---|-----------------------------------|
| $\text{C}_{29}\text{H}_{37}\text{NO}_2$ | $V = 2350.4$ (5) Å <sup>3</sup>   |
| $M_r = 431.60$                          | $Z = 4$                           |
| Monoclinic, $P2_1$                      | Mo $K\alpha$ radiation            |
| $a = 7.5895$ (10) Å                     | $\mu = 0.08$ mm <sup>-1</sup>     |
| $b = 31.355$ (3) Å                      | $T = 113$ K                       |
| $c = 9.8912$ (12) Å                     | $0.22 \times 0.20 \times 0.12$ mm |
| $\beta = 93.056$ (6)°                   |                                   |

## Data collection

|  |  |
|--|--|
| Rigaku Saturn724 CCD diffractometer  | 17383 measured reflections             |
| Absorption correction: multi-scan ( <i>CrystalClear</i> ; Rigaku/MS, 2005) | 5642 independent reflections           |
| $T_{\min} = 0.984$ , $T_{\max} = 0.991$                                    | 5296 reflections with $I > 2\sigma(I)$ |
|  | $R_{\text{int}} = 0.052$               |

## Refinement

|                                 |   |
|---------------------------------|---|
| $R[F^2 > 2\sigma(F^2)] = 0.039$ | 1 restraint   |
| $wR(F^2) = 0.096$               | H-atom parameters constrained                       |
| $S = 1.04$                      | $\Delta\rho_{\text{max}} = 0.26$ e Å <sup>-3</sup>  |
| 5642 reflections                | $\Delta\rho_{\text{min}} = -0.27$ e Å <sup>-3</sup> |
| 583 parameters                  |   |

Table 1

Hydrogen-bond geometry (Å, °).

| $D-H\cdots A$                       | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|-------------------------------------|-------|-------------|-------------|---------------|
| C1—H1B $\cdots$ O2                  | 0.99  | 2.25        | 3.027 (3)   | 134           |
| C5—H5 $\cdots$ O4 <sup>i</sup>      | 1.00  | 2.59        | 3.237 (3)   | 122           |
| C21—H21C $\cdots$ O3 <sup>ii</sup>  | 0.98  | 2.60        | 3.501 (3)   | 153           |
| C31—H31A $\cdots$ O2 <sup>iii</sup> | 0.99  | 2.47        | 3.361 (3)   | 150           |

 Symmetry codes: (i)  $x, y, z + 1$ ; (ii)  $x - 1, y, z$ ; (iii)  $x + 1, y, z - 1$ .

Data collection: *CrystalClear* (Rigaku/MS, 2005); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; 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: *CrystalStructure* (Rigaku/MS, 2005).

This work was supported by the National Natural Science Foundation of China (No. 81072540). The authors are grateful to the Central Laboratory of Nankai University for the data collection. Special thanks go to Dr Xie Chengzhi (School of Pharmaceutical Sciences, Tianjin Medical University) for his invaluable support.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: PV2421).

## References

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## supporting information

*Acta Cryst.* (2011). E67, o2065 [doi:10.1107/S1600536811027632]

**(Z)-3 $\alpha$ -(1,3-Dioxoisindolin-2-yl)-17(20)-pregnene**

Yue Qi, Nan Qin and Hong-Quan Duan

**S1. Comment**

The title compound was synthesized from epiandrosterone which is a pregnane alkaloids isolated from *Pachysandra axillaris*, a Traditional Chinese Medicine (TCM). The pregnane alkaloids from *Pachysandra axillaris* had been reported to be effective as anticancer (Sun *et al.*, 2010).

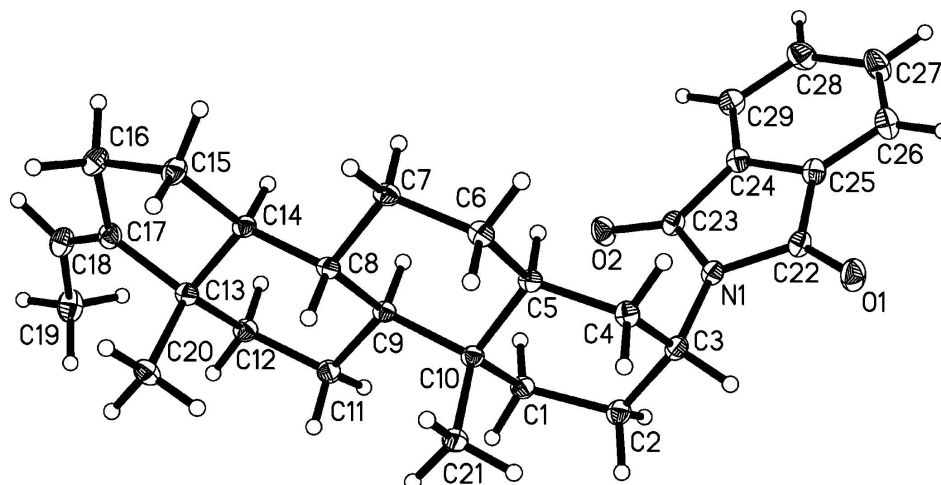
The title compound, crystallizes with two independent molecules in an asymmetric unit. The cyclohexyl rings (*A*) of the pregnene moiety bonded to the 3 $\alpha$ -(1,3-dioxoisindolin-2-yl)- ring system in both molecules differ in conformations; in one molecule (Fig. 1) it is in a chair conformation while in the other molecule (Fig. 2) it exhibits a half chair conformation. The conformations of the rings B–D in the two molecules are identical with six membered rings (B & C) in chair conformations and the five membered rings in envelope forms in both molecules. In both molecules, the 3 $\alpha$ -(1,3-dioxoisindolin-2-yl)- ring systems are individually planar with rms deviations 0.0148 and 0.0264 Å. The structure is consolidated by intermolecular hydrogen bonding interactions of the type C—H $\cdots$ O involving carbonyl O atoms and methyl, methylene and methylidyne groups, resulting in a two-dimensional structure (Fig. 3).

**S2. Experimental**

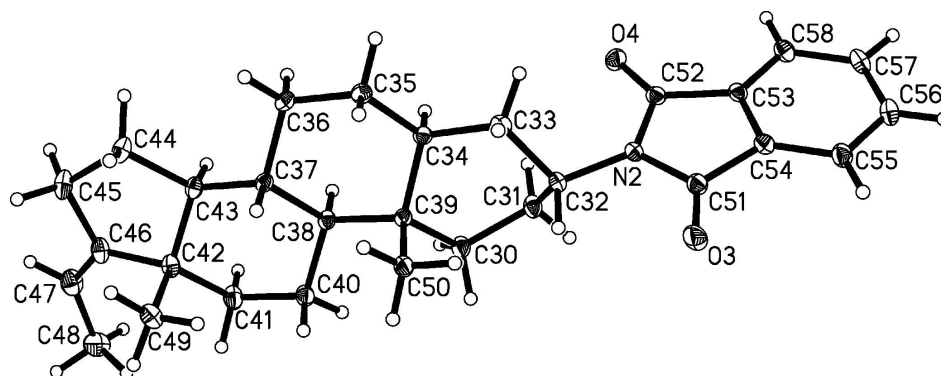
The title compound was prepared by following a literature procedure (Batcho *et al.*, 1981). A solution of potassium *t*-butoxide (*t*-BuOK) (2.28 g, 20.0 mmol) in 15 ml THF was added to a suspension of Ph<sub>3</sub>PEtBr (7.58 g, 20.0 mmol), in 25 ml dry THF, and was stirred for 1 h at room temperature. The mixture was refluxed for another 4 h after adding epiandrosterone (1.45 g, 5.0 mmol). The residue thus obtained was purified by recrystallization from MeOH to afford **1** (675.8 mg, 45%; Fig. 4). To a solution of **1** (86.9 mg, 0.287 mmol) and triphenylphosphine (85.1 mg, 0.324 mmol) in THF (12 ml), phthalimide (46.5 mg, 0.316 mmol) and diisopropyl azodicarboxylate (63.9 mg, 0.316 mmol) were added, and the mixture was stirred for 18 h at room temperature. The residue was purified by crystallization from MeOH to afford the title compound, **2** (78.0 mg, 63%) as a white solid. The crystals of **2** were obtained by slow evaporation of its solution using a mixed solvent MeOH/CH<sub>2</sub>Cl<sub>2</sub> (1:1).

**S3. Refinement**

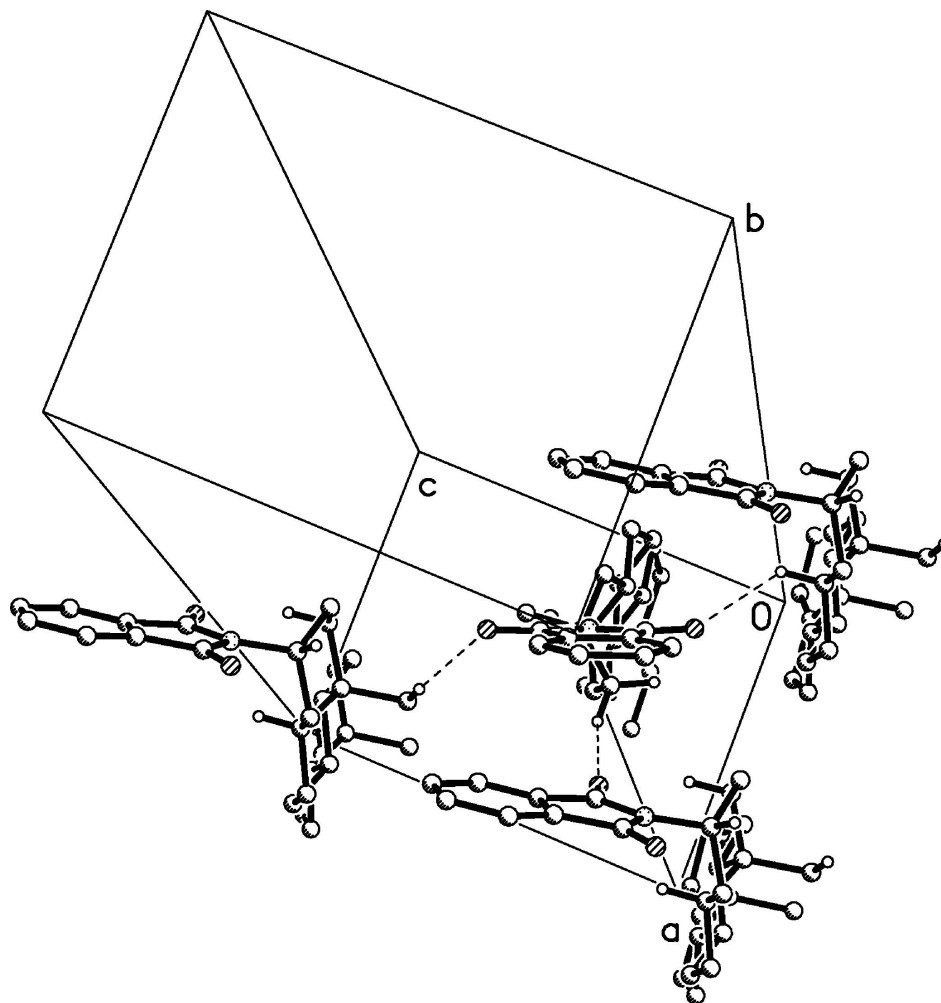
H atoms were placed at calculated positions with C—H = 0.95 Å (aryl), 0.98 (methylene), 0.99 (methyl) and 1.00 Å (methyne) and were refined in the riding-model approximation with  $U_{\text{iso}} = 1.2\text{--}1.5$  times  $U_{\text{eq}}$  of the parent atoms. As the structure has no anomalous scatterer, an absolute structure could not be established in this analysis; the Friedel-pairs (3600) of reflections were merged.

**Figure 1**

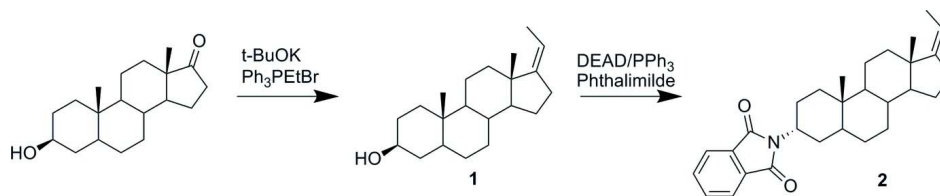
One independent molecule in the asymmetric unit of the title compound, with 30% probability displacement ellipsoids.

**Figure 2**

The other independent molecule in the asymmetric unit of the title compound, with 30% probability displacement ellipsoids.

**Figure 3**

The packing of the title compound, showing the two-dimensional structure, with intermolecular hydrogen bonds (dashed lines); for clarity H atoms not involved in H-bonds have been omitted.

**Figure 4**

The scheme of synthesis of the title compound.

**(Z)-3 $\alpha$ -(1,3-Dioxisoindolin-2-yl)-17 (20)-pregnene**

*Crystal data*

$C_{29}H_{37}NO_2$

$M_r = 431.60$

Monoclinic,  $P2_1$

Hall symbol: P 2yb

$a = 7.5895 (10) \text{ \AA}$

$b = 31.355 (3) \text{ \AA}$

$c = 9.8912 (12) \text{ \AA}$

$\beta = 93.056 (6)^\circ$

$V = 2350.4 (5) \text{ \AA}^3$   
 $Z = 4$   
 $F(000) = 936$   
 $D_x = 1.220 \text{ Mg m}^{-3}$   
 Mo  $K\alpha$  radiation,  $\lambda = 0.71073 \text{ \AA}$   
 Cell parameters from 7024 reflections

$\theta = 1.9\text{--}28.4^\circ$   
 $\mu = 0.08 \text{ mm}^{-1}$   
 $T = 113 \text{ K}$   
 Prism, colorless  
 $0.22 \times 0.20 \times 0.12 \text{ mm}$

*Data collection*

Rigaku Saturn724 CCD  
 diffractometer  
 Radiation source: rotating anode  
 Multilayer monochromator  
 Detector resolution:  $14.22 \text{ pixels mm}^{-1}$   
 $\omega$  and  $\phi$  scans  
 Absorption correction: multi-scan  
 (*CrystalClear*; Rigaku/MSC, 2005)  
 $T_{\min} = 0.984$ ,  $T_{\max} = 0.991$

17383 measured reflections  
 5642 independent reflections  
 5296 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.052$   
 $\theta_{\max} = 27.9^\circ$ ,  $\theta_{\min} = 2.1^\circ$   
 $h = -9 \rightarrow 9$   
 $k = -41 \rightarrow 35$   
 $l = -12 \rightarrow 13$

*Refinement*

Refinement on  $F^2$   
 Least-squares matrix: full  
 $R[F^2 > 2\sigma(F^2)] = 0.039$   
 $wR(F^2) = 0.096$   
 $S = 1.04$   
 5642 reflections  
 583 parameters  
 1 restraint  
 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  
 $w = 1/[\sigma^2(F_o^2) + (0.0522P)^2 + 0.1053P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} < 0.001$   
 $\Delta\rho_{\max} = 0.26 \text{ e \AA}^{-3}$   
 $\Delta\rho_{\min} = -0.27 \text{ e \AA}^{-3}$

*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  $F^2 > \sigma(F^2)$  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 ( $\text{\AA}^2$ )*

|     | $x$         | $y$         | $z$          | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|-----|-------------|-------------|--------------|----------------------------------|
| O1  | 0.0919 (2)  | 0.42410 (5) | 0.94269 (15) | 0.0332 (4)                       |
| O2  | -0.0642 (2) | 0.30027 (5) | 1.15244 (15) | 0.0323 (4)                       |
| O3  | 0.6594 (2)  | 0.33129 (5) | 0.49697 (15) | 0.0363 (4)                       |
| O4  | 0.4062 (2)  | 0.30622 (5) | 0.07334 (14) | 0.0282 (3)                       |
| N1  | -0.0051 (2) | 0.35800 (5) | 1.01138 (16) | 0.0212 (3)                       |
| N2  | 0.5259 (2)  | 0.30729 (5) | 0.29473 (16) | 0.0209 (3)                       |
| C1  | -0.2120 (3) | 0.27009 (6) | 0.8787 (2)   | 0.0226 (4)                       |
| H1A | -0.3231     | 0.2561      | 0.8459       | 0.027*                           |
| H1B | -0.1964     | 0.2647      | 0.9772       | 0.027*                           |
| C2  | -0.2298 (3) | 0.31824 (7) | 0.8555 (2)   | 0.0245 (4)                       |

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|      |             |             |              |            |
|------|-------------|-------------|--------------|------------|
| H2A  | -0.3138     | 0.3296      | 0.9197       | 0.029*     |
| H2B  | -0.2824     | 0.3229      | 0.7629       | 0.029*     |
| C3   | -0.0590 (3) | 0.34439 (6) | 0.87136 (19) | 0.0224 (4) |
| H3   | -0.0820     | 0.3712      | 0.8188       | 0.027*     |
| C4   | 0.0929 (3)  | 0.32200 (6) | 0.80395 (19) | 0.0222 (4) |
| H4A  | 0.0725      | 0.3237      | 0.7044       | 0.027*     |
| H4B  | 0.2045      | 0.3371      | 0.8286       | 0.027*     |
| C5   | 0.1118 (3)  | 0.27542 (6) | 0.84529 (18) | 0.0194 (4) |
| H5   | 0.1287      | 0.2749      | 0.9462       | 0.023*     |
| C6   | 0.2767 (3)  | 0.25537 (7) | 0.78951 (19) | 0.0218 (4) |
| H6A  | 0.2656      | 0.2555      | 0.6893       | 0.026*     |
| H6B  | 0.3814      | 0.2726      | 0.8183       | 0.026*     |
| C7   | 0.3024 (2)  | 0.20970 (6) | 0.83963 (19) | 0.0209 (4) |
| H7A  | 0.3314      | 0.2101      | 0.9384       | 0.025*     |
| H7B  | 0.4033      | 0.1968      | 0.7951       | 0.025*     |
| C8   | 0.1381 (2)  | 0.18215 (6) | 0.81060 (19) | 0.0188 (4) |
| H8   | 0.1181      | 0.1790      | 0.7103       | 0.023*     |
| C9   | -0.0256 (2) | 0.20368 (6) | 0.86728 (18) | 0.0181 (4) |
| H9   | 0.0025      | 0.2076      | 0.9664       | 0.022*     |
| C10  | -0.0571 (3) | 0.24924 (6) | 0.80799 (18) | 0.0193 (4) |
| C11  | -0.1900 (3) | 0.17485 (7) | 0.8555 (2)   | 0.0233 (4) |
| H11A | -0.2851     | 0.1884      | 0.9052       | 0.028*     |
| H11B | -0.2312     | 0.1730      | 0.7591       | 0.028*     |
| C12  | -0.1595 (3) | 0.12943 (7) | 0.9107 (2)   | 0.0224 (4) |
| H12A | -0.1375     | 0.1306      | 1.0102       | 0.027*     |
| H12B | -0.2667     | 0.1120      | 0.8909       | 0.027*     |
| C13  | -0.0016 (3) | 0.10837 (6) | 0.84645 (19) | 0.0200 (4) |
| C14  | 0.1594 (2)  | 0.13797 (6) | 0.87296 (18) | 0.0192 (4) |
| H14  | 0.1701      | 0.1425      | 0.9732       | 0.023*     |
| C15  | 0.3180 (3)  | 0.11070 (7) | 0.8388 (2)   | 0.0252 (4) |
| H15A | 0.4285      | 0.1218      | 0.8837       | 0.030*     |
| H15B | 0.3313      | 0.1095      | 0.7399       | 0.030*     |
| C16  | 0.2700 (3)  | 0.06657 (7) | 0.8955 (2)   | 0.0263 (4) |
| H16A | 0.3051      | 0.0436      | 0.8336       | 0.032*     |
| H16B | 0.3308      | 0.0619      | 0.9853       | 0.032*     |
| C17  | 0.0697 (3)  | 0.06678 (7) | 0.90705 (19) | 0.0237 (4) |
| C18  | -0.0176 (3) | 0.03466 (7) | 0.9622 (2)   | 0.0291 (5) |
| H18  | 0.0521      | 0.0113      | 0.9946       | 0.035*     |
| C19  | -0.2132 (3) | 0.03103 (8) | 0.9792 (3)   | 0.0352 (5) |
| H19A | -0.2483     | 0.0516      | 1.0474       | 0.053*     |
| H19B | -0.2412     | 0.0021      | 1.0085       | 0.053*     |
| H19C | -0.2772     | 0.0371      | 0.8927       | 0.053*     |
| C20  | -0.0439 (3) | 0.10017 (7) | 0.6943 (2)   | 0.0266 (4) |
| H20A | 0.0583      | 0.0869      | 0.6546       | 0.040*     |
| H20B | -0.0710     | 0.1273      | 0.6486       | 0.040*     |
| H20C | -0.1459     | 0.0811      | 0.6832       | 0.040*     |
| C21  | -0.0997 (3) | 0.24812 (7) | 0.65418 (19) | 0.0241 (4) |
| H21A | -0.1986     | 0.2286      | 0.6341       | 0.036*     |

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|      |             |             |              |            |
|------|-------------|-------------|--------------|------------|
| H21B | 0.0040      | 0.2382      | 0.6083       | 0.036*     |
| H21C | -0.1316     | 0.2768      | 0.6222       | 0.036*     |
| C22  | 0.0686 (3)  | 0.39855 (7) | 1.0316 (2)   | 0.0234 (4) |
| C23  | -0.0088 (3) | 0.33577 (7) | 1.13435 (19) | 0.0225 (4) |
| C24  | 0.0676 (3)  | 0.36517 (7) | 1.24055 (19) | 0.0228 (4) |
| C25  | 0.1115 (3)  | 0.40315 (7) | 1.1793 (2)   | 0.0246 (4) |
| C26  | 0.1802 (3)  | 0.43709 (7) | 1.2540 (2)   | 0.0327 (5) |
| H26  | 0.2100      | 0.4632      | 1.2120       | 0.039*     |
| C27  | 0.2040 (4)  | 0.43139 (8) | 1.3936 (2)   | 0.0387 (6) |
| H27  | 0.2503      | 0.4542      | 1.4480       | 0.046*     |
| C28  | 0.1616 (3)  | 0.39320 (9) | 1.4547 (2)   | 0.0361 (5) |
| H28  | 0.1806      | 0.3902      | 1.5499       | 0.043*     |
| C29  | 0.0919 (3)  | 0.35916 (8) | 1.3792 (2)   | 0.0295 (5) |
| H29  | 0.0623      | 0.3330      | 1.4207       | 0.035*     |
| C30  | 0.6180 (3)  | 0.18753 (7) | 0.2866 (2)   | 0.0267 (4) |
| H30A | 0.7183      | 0.1847      | 0.3541       | 0.032*     |
| H30B | 0.6414      | 0.1685      | 0.2097       | 0.032*     |
| C31  | 0.6114 (3)  | 0.23375 (7) | 0.2351 (2)   | 0.0266 (4) |
| H31A | 0.7330      | 0.2448      | 0.2291       | 0.032*     |
| H31B | 0.5527      | 0.2346      | 0.1433       | 0.032*     |
| C32  | 0.5104 (3)  | 0.26194 (6) | 0.3308 (2)   | 0.0220 (4) |
| H32  | 0.5656      | 0.2582      | 0.4240       | 0.026*     |
| C33  | 0.3173 (3)  | 0.24763 (7) | 0.3335 (2)   | 0.0226 (4) |
| H33A | 0.2446      | 0.2651      | 0.2685       | 0.027*     |
| H33B | 0.2748      | 0.2528      | 0.4250       | 0.027*     |
| C34  | 0.2920 (3)  | 0.20021 (6) | 0.29740 (19) | 0.0194 (4) |
| H34  | 0.2892      | 0.1982      | 0.1964       | 0.023*     |
| C35  | 0.1129 (3)  | 0.18443 (7) | 0.3409 (2)   | 0.0239 (4) |
| H35A | 0.1073      | 0.1880      | 0.4401       | 0.029*     |
| H35B | 0.0184      | 0.2021      | 0.2967       | 0.029*     |
| C36  | 0.0799 (3)  | 0.13791 (7) | 0.3046 (2)   | 0.0233 (4) |
| H36A | 0.0666      | 0.1351      | 0.2049       | 0.028*     |
| H36B | -0.0318     | 0.1286      | 0.3425       | 0.028*     |
| C37  | 0.2294 (2)  | 0.10888 (6) | 0.35822 (18) | 0.0186 (4) |
| H37  | 0.2339      | 0.1096      | 0.4594       | 0.022*     |
| C38  | 0.4080 (3)  | 0.12490 (6) | 0.30987 (19) | 0.0192 (4) |
| H38  | 0.3969      | 0.1247      | 0.2087       | 0.023*     |
| C39  | 0.4472 (2)  | 0.17192 (6) | 0.35221 (18) | 0.0184 (4) |
| C40  | 0.5592 (3)  | 0.09406 (7) | 0.3504 (2)   | 0.0278 (4) |
| H40A | 0.6672      | 0.1036      | 0.3071       | 0.033*     |
| H40B | 0.5829      | 0.0956      | 0.4497       | 0.033*     |
| C41  | 0.5217 (3)  | 0.04757 (7) | 0.3106 (2)   | 0.0287 (5) |
| H41A | 0.6196      | 0.0293      | 0.3470       | 0.034*     |
| H41B | 0.5159      | 0.0450      | 0.2107       | 0.034*     |
| C42  | 0.3478 (3)  | 0.03193 (7) | 0.3651 (2)   | 0.0233 (4) |
| C43  | 0.2021 (3)  | 0.06299 (6) | 0.31204 (18) | 0.0209 (4) |
| H43  | 0.2081      | 0.0634      | 0.2113       | 0.025*     |
| C44  | 0.0288 (3)  | 0.04034 (7) | 0.3408 (2)   | 0.0282 (5) |

|      |            |              |              |            |
|------|------------|--------------|--------------|------------|
| H44A | -0.0691    | 0.0504       | 0.2787       | 0.034*     |
| H44B | -0.0030    | 0.0451       | 0.4354       | 0.034*     |
| C45  | 0.0704 (3) | -0.00700 (7) | 0.3158 (2)   | 0.0329 (5) |
| H45A | 0.0266     | -0.0249      | 0.3893       | 0.039*     |
| H45B | 0.0138     | -0.0166      | 0.2286       | 0.039*     |
| C46  | 0.2709 (3) | -0.01020 (7) | 0.3130 (2)   | 0.0275 (5) |
| C47  | 0.3507 (4) | -0.04530 (8) | 0.2710 (2)   | 0.0346 (5) |
| H47  | 0.2742     | -0.0674      | 0.2378       | 0.042*     |
| C48  | 0.5431 (4) | -0.05482 (8) | 0.2688 (3)   | 0.0404 (6) |
| H48A | 0.5838     | -0.0492      | 0.1782       | 0.061*     |
| H48B | 0.5637     | -0.0849      | 0.2919       | 0.061*     |
| H48C | 0.6082     | -0.0367      | 0.3349       | 0.061*     |
| C49  | 0.3627 (3) | 0.02932 (7)  | 0.5214 (2)   | 0.0294 (5) |
| H49A | 0.2481     | 0.0214       | 0.5550       | 0.044*     |
| H49B | 0.3989     | 0.0571       | 0.5587       | 0.044*     |
| H49C | 0.4508     | 0.0078       | 0.5496       | 0.044*     |
| C50  | 0.4707 (3) | 0.17570 (7)  | 0.5074 (2)   | 0.0251 (4) |
| H50A | 0.4909     | 0.2056       | 0.5324       | 0.038*     |
| H50B | 0.5720     | 0.1585       | 0.5401       | 0.038*     |
| H50C | 0.3640     | 0.1654       | 0.5484       | 0.038*     |
| C51  | 0.6074 (3) | 0.33777 (7)  | 0.3812 (2)   | 0.0241 (4) |
| C52  | 0.4818 (3) | 0.32479 (6)  | 0.16710 (19) | 0.0215 (4) |
| C53  | 0.5465 (3) | 0.36949 (6)  | 0.17163 (19) | 0.0199 (4) |
| C54  | 0.6196 (3) | 0.37753 (6)  | 0.30028 (19) | 0.0225 (4) |
| C55  | 0.6941 (3) | 0.41651 (7)  | 0.3343 (2)   | 0.0278 (4) |
| H55  | 0.7417     | 0.4223       | 0.4233       | 0.033*     |
| C56  | 0.6962 (3) | 0.44707 (7)  | 0.2317 (2)   | 0.0303 (5) |
| H56  | 0.7488     | 0.4740       | 0.2508       | 0.036*     |
| C57  | 0.6231 (3) | 0.43903 (7)  | 0.1023 (2)   | 0.0282 (5) |
| H57  | 0.6266     | 0.4605       | 0.0349       | 0.034*     |
| C58  | 0.5448 (3) | 0.39987 (7)  | 0.0701 (2)   | 0.0256 (4) |
| H58  | 0.4927     | 0.3943       | -0.0175      | 0.031*     |

*Atomic displacement parameters (Å<sup>2</sup>)*

|    | $U^{11}$    | $U^{22}$    | $U^{33}$   | $U^{12}$    | $U^{13}$    | $U^{23}$    |
|----|-------------|-------------|------------|-------------|-------------|-------------|
| O1 | 0.0471 (10) | 0.0241 (8)  | 0.0284 (8) | -0.0045 (7) | 0.0028 (7)  | 0.0046 (6)  |
| O2 | 0.0435 (9)  | 0.0258 (8)  | 0.0274 (7) | -0.0104 (7) | 0.0000 (6)  | 0.0038 (6)  |
| O3 | 0.0557 (11) | 0.0276 (9)  | 0.0240 (7) | -0.0018 (8) | -0.0127 (7) | 0.0044 (6)  |
| O4 | 0.0325 (8)  | 0.0286 (8)  | 0.0228 (7) | -0.0053 (6) | -0.0053 (6) | 0.0008 (6)  |
| N1 | 0.0242 (8)  | 0.0197 (8)  | 0.0198 (7) | 0.0024 (6)  | 0.0026 (6)  | 0.0013 (6)  |
| N2 | 0.0229 (8)  | 0.0184 (8)  | 0.0211 (8) | -0.0003 (7) | -0.0004 (6) | 0.0033 (6)  |
| C1 | 0.0161 (9)  | 0.0247 (11) | 0.0270 (9) | 0.0017 (7)  | 0.0010 (7)  | -0.0031 (7) |
| C2 | 0.0233 (10) | 0.0246 (11) | 0.0257 (9) | 0.0049 (8)  | 0.0021 (8)  | -0.0024 (7) |
| C3 | 0.0258 (10) | 0.0201 (10) | 0.0211 (9) | 0.0035 (8)  | 0.0006 (8)  | 0.0012 (7)  |
| C4 | 0.0238 (10) | 0.0222 (10) | 0.0208 (9) | -0.0004 (8) | 0.0035 (7)  | -0.0012 (7) |
| C5 | 0.0180 (9)  | 0.0224 (10) | 0.0177 (8) | 0.0015 (7)  | 0.0017 (7)  | -0.0021 (7) |
| C6 | 0.0194 (9)  | 0.0253 (10) | 0.0212 (9) | -0.0001 (8) | 0.0038 (7)  | -0.0031 (7) |



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|     |             |             |             |              |              |              |
|-----|-------------|-------------|-------------|--------------|--------------|--------------|
| C7  | 0.0148 (9)  | 0.0272 (10) | 0.0207 (9)  | -0.0001 (7)  | 0.0015 (7)   | -0.0030 (7)  |
| C8  | 0.0163 (9)  | 0.0226 (10) | 0.0174 (8)  | 0.0008 (7)   | 0.0009 (7)   | -0.0031 (7)  |
| C9  | 0.0149 (9)  | 0.0220 (10) | 0.0176 (8)  | 0.0013 (7)   | 0.0016 (7)   | -0.0019 (7)  |
| C10 | 0.0190 (9)  | 0.0212 (10) | 0.0176 (8)  | 0.0021 (7)   | -0.0001 (7)  | -0.0013 (7)  |
| C11 | 0.0171 (9)  | 0.0238 (10) | 0.0288 (10) | 0.0010 (8)   | 0.0004 (7)   | -0.0032 (8)  |
| C12 | 0.0170 (9)  | 0.0250 (10) | 0.0253 (9)  | 0.0000 (8)   | 0.0020 (7)   | 0.0002 (7)   |
| C13 | 0.0191 (9)  | 0.0225 (10) | 0.0184 (8)  | 0.0003 (8)   | 0.0018 (7)   | -0.0014 (7)  |
| C14 | 0.0168 (9)  | 0.0227 (10) | 0.0181 (8)  | 0.0033 (7)   | 0.0014 (7)   | -0.0008 (7)  |
| C15 | 0.0206 (9)  | 0.0247 (10) | 0.0307 (10) | 0.0052 (8)   | 0.0052 (8)   | -0.0007 (8)  |
| C16 | 0.0254 (11) | 0.0265 (11) | 0.0277 (10) | 0.0062 (8)   | 0.0069 (8)   | 0.0031 (8)   |
| C17 | 0.0255 (10) | 0.0237 (10) | 0.0222 (9)  | 0.0023 (8)   | 0.0038 (8)   | -0.0028 (7)  |
| C18 | 0.0324 (12) | 0.0257 (11) | 0.0299 (10) | 0.0039 (9)   | 0.0081 (9)   | 0.0009 (8)   |
| C19 | 0.0364 (13) | 0.0272 (12) | 0.0432 (13) | -0.0030 (10) | 0.0126 (10)  | 0.0018 (9)   |
| C20 | 0.0301 (11) | 0.0282 (11) | 0.0211 (9)  | -0.0044 (8)  | -0.0017 (8)  | -0.0038 (7)  |
| C21 | 0.0280 (11) | 0.0249 (10) | 0.0190 (8)  | 0.0017 (8)   | -0.0030 (8)  | -0.0016 (7)  |
| C22 | 0.0269 (10) | 0.0203 (10) | 0.0233 (9)  | 0.0032 (8)   | 0.0040 (8)   | 0.0011 (7)   |
| C23 | 0.0251 (10) | 0.0218 (10) | 0.0208 (9)  | 0.0019 (8)   | 0.0038 (7)   | 0.0017 (7)   |
| C24 | 0.0222 (10) | 0.0235 (10) | 0.0229 (9)  | 0.0008 (8)   | 0.0038 (7)   | -0.0011 (7)  |
| C25 | 0.0253 (10) | 0.0231 (10) | 0.0260 (10) | 0.0011 (8)   | 0.0056 (8)   | -0.0007 (8)  |
| C26 | 0.0400 (13) | 0.0256 (12) | 0.0329 (11) | -0.0053 (10) | 0.0055 (10)  | -0.0038 (8)  |
| C27 | 0.0498 (15) | 0.0345 (13) | 0.0316 (12) | -0.0098 (11) | 0.0010 (10)  | -0.0110 (9)  |
| C28 | 0.0423 (14) | 0.0419 (14) | 0.0240 (10) | -0.0056 (11) | -0.0005 (9)  | -0.0034 (9)  |
| C29 | 0.0341 (12) | 0.0315 (12) | 0.0232 (10) | -0.0018 (9)  | 0.0034 (8)   | 0.0023 (8)   |
| C30 | 0.0186 (10) | 0.0220 (11) | 0.0401 (12) | -0.0002 (8)  | 0.0071 (9)   | 0.0036 (8)   |
| C31 | 0.0219 (10) | 0.0211 (10) | 0.0376 (11) | -0.0012 (8)  | 0.0098 (9)   | 0.0030 (8)   |
| C32 | 0.0244 (10) | 0.0167 (9)  | 0.0249 (9)  | -0.0010 (8)  | 0.0000 (8)   | 0.0047 (7)   |
| C33 | 0.0212 (10) | 0.0231 (10) | 0.0238 (9)  | 0.0030 (8)   | 0.0046 (7)   | 0.0033 (7)   |
| C34 | 0.0179 (9)  | 0.0211 (10) | 0.0191 (8)  | -0.0001 (7)  | 0.0012 (7)   | 0.0019 (7)   |
| C35 | 0.0178 (9)  | 0.0231 (11) | 0.0310 (10) | 0.0017 (8)   | 0.0041 (8)   | 0.0011 (8)   |
| C36 | 0.0160 (9)  | 0.0252 (11) | 0.0288 (10) | -0.0026 (8)  | 0.0016 (7)   | 0.0009 (8)   |
| C37 | 0.0173 (9)  | 0.0204 (9)  | 0.0183 (8)  | -0.0020 (7)  | 0.0027 (7)   | 0.0001 (7)   |
| C38 | 0.0174 (9)  | 0.0196 (9)  | 0.0211 (8)  | -0.0013 (7)  | 0.0050 (7)   | 0.0006 (7)   |
| C39 | 0.0163 (9)  | 0.0182 (9)  | 0.0209 (8)  | -0.0007 (7)  | 0.0022 (7)   | 0.0008 (7)   |
| C40 | 0.0198 (10) | 0.0204 (10) | 0.0434 (12) | 0.0008 (8)   | 0.0042 (9)   | 0.0048 (9)   |
| C41 | 0.0260 (11) | 0.0217 (11) | 0.0395 (12) | 0.0014 (8)   | 0.0105 (9)   | 0.0008 (8)   |
| C42 | 0.0275 (10) | 0.0193 (10) | 0.0238 (9)  | -0.0016 (8)  | 0.0069 (8)   | -0.0012 (7)  |
| C43 | 0.0228 (10) | 0.0214 (10) | 0.0191 (8)  | -0.0044 (8)  | 0.0052 (7)   | -0.0024 (7)  |
| C44 | 0.0254 (11) | 0.0276 (11) | 0.0324 (11) | -0.0084 (8)  | 0.0078 (8)   | -0.0069 (8)  |
| C45 | 0.0370 (13) | 0.0246 (11) | 0.0383 (12) | -0.0105 (9)  | 0.0132 (10)  | -0.0057 (9)  |
| C46 | 0.0373 (12) | 0.0219 (10) | 0.0241 (10) | -0.0054 (9)  | 0.0092 (9)   | -0.0010 (8)  |
| C47 | 0.0450 (14) | 0.0272 (12) | 0.0318 (11) | -0.0028 (10) | 0.0030 (10)  | -0.0057 (9)  |
| C48 | 0.0478 (15) | 0.0281 (13) | 0.0442 (13) | 0.0077 (11)  | -0.0080 (11) | -0.0132 (10) |
| C49 | 0.0389 (12) | 0.0245 (11) | 0.0248 (10) | 0.0028 (9)   | 0.0027 (9)   | 0.0015 (8)   |
| C50 | 0.0298 (11) | 0.0212 (10) | 0.0235 (9)  | -0.0036 (8)  | -0.0040 (8)  | 0.0029 (7)   |
| C51 | 0.0283 (11) | 0.0215 (10) | 0.0224 (9)  | 0.0000 (8)   | -0.0013 (8)  | 0.0004 (7)   |
| C52 | 0.0203 (10) | 0.0227 (10) | 0.0215 (9)  | 0.0007 (7)   | 0.0008 (7)   | 0.0037 (7)   |
| C53 | 0.0197 (9)  | 0.0181 (9)  | 0.0221 (9)  | 0.0026 (7)   | 0.0023 (7)   | -0.0001 (7)  |
| C54 | 0.0242 (10) | 0.0213 (10) | 0.0219 (9)  | 0.0025 (8)   | 0.0015 (7)   | 0.0002 (7)   |

|     |             |             |             |             |             |             |
|-----|-------------|-------------|-------------|-------------|-------------|-------------|
| C55 | 0.0345 (12) | 0.0207 (10) | 0.0280 (10) | 0.0008 (9)  | -0.0001 (8) | -0.0031 (8) |
| C56 | 0.0376 (13) | 0.0192 (10) | 0.0343 (11) | -0.0020 (9) | 0.0048 (9)  | -0.0035 (8) |
| C57 | 0.0392 (13) | 0.0189 (10) | 0.0274 (10) | 0.0041 (9)  | 0.0085 (9)  | 0.0039 (7)  |
| C58 | 0.0324 (11) | 0.0231 (10) | 0.0215 (9)  | 0.0028 (8)  | 0.0043 (8)  | 0.0018 (7)  |

*Geometric parameters (Å, °)*

|          |           |          |           |
|----------|-----------|----------|-----------|
| O1—C22   | 1.210 (2) | C27—C28  | 1.387 (4) |
| O2—C23   | 1.207 (3) | C27—H27  | 0.9500    |
| O3—C51   | 1.208 (2) | C28—C29  | 1.391 (3) |
| O4—C52   | 1.213 (2) | C28—H28  | 0.9500    |
| N1—C22   | 1.399 (3) | C29—H29  | 0.9500    |
| N1—C23   | 1.403 (2) | C30—C31  | 1.536 (3) |
| N1—C3    | 1.486 (2) | C30—C39  | 1.559 (3) |
| N2—C52   | 1.401 (2) | C30—H30A | 0.9900    |
| N2—C51   | 1.404 (3) | C30—H30B | 0.9900    |
| N2—C32   | 1.472 (2) | C31—C32  | 1.531 (3) |
| C1—C2    | 1.532 (3) | C31—H31A | 0.9900    |
| C1—C10   | 1.544 (3) | C31—H31B | 0.9900    |
| C1—H1A   | 0.9900    | C32—C33  | 1.534 (3) |
| C1—H1B   | 0.9900    | C32—H32  | 1.0000    |
| C2—C3    | 1.534 (3) | C33—C34  | 1.539 (3) |
| C2—H2A   | 0.9900    | C33—H33A | 0.9900    |
| C2—H2B   | 0.9900    | C33—H33B | 0.9900    |
| C3—C4    | 1.532 (3) | C34—C35  | 1.530 (3) |
| C3—H3    | 1.0000    | C34—C39  | 1.549 (3) |
| C4—C5    | 1.521 (3) | C34—H34  | 1.0000    |
| C4—H4A   | 0.9900    | C35—C36  | 1.520 (3) |
| C4—H4B   | 0.9900    | C35—H35A | 0.9900    |
| C5—C6    | 1.529 (3) | C35—H35B | 0.9900    |
| C5—C10   | 1.550 (3) | C36—C37  | 1.527 (3) |
| C5—H5    | 1.0000    | C36—H36A | 0.9900    |
| C6—C7    | 1.525 (3) | C36—H36B | 0.9900    |
| C6—H6A   | 0.9900    | C37—C43  | 1.521 (3) |
| C6—H6B   | 0.9900    | C37—C38  | 1.545 (2) |
| C7—C8    | 1.531 (3) | C37—H37  | 1.0000    |
| C7—H7A   | 0.9900    | C38—C40  | 1.537 (3) |
| C7—H7B   | 0.9900    | C38—C39  | 1.557 (3) |
| C8—C14   | 1.521 (3) | C38—H38  | 1.0000    |
| C8—C9    | 1.546 (2) | C39—C50  | 1.540 (3) |
| C8—H8    | 1.0000    | C40—C41  | 1.533 (3) |
| C9—C11   | 1.540 (3) | C40—H40A | 0.9900    |
| C9—C10   | 1.558 (3) | C40—H40B | 0.9900    |
| C9—H9    | 1.0000    | C41—C42  | 1.533 (3) |
| C10—C21  | 1.539 (2) | C41—H41A | 0.9900    |
| C11—C12  | 1.538 (3) | C41—H41B | 0.9900    |
| C11—H11A | 0.9900    | C42—C46  | 1.523 (3) |
| C11—H11B | 0.9900    | C42—C43  | 1.544 (3) |

|            |             |               |             |
|------------|-------------|---------------|-------------|
| C12—C13    | 1.535 (3)   | C42—C49       | 1.547 (3)   |
| C12—H12A   | 0.9900      | C43—C44       | 1.534 (3)   |
| C12—H12B   | 0.9900      | C43—H43       | 1.0000      |
| C13—C17    | 1.523 (3)   | C44—C45       | 1.540 (3)   |
| C13—C20    | 1.543 (3)   | C44—H44A      | 0.9900      |
| C13—C14    | 1.546 (3)   | C44—H44B      | 0.9900      |
| C14—C15    | 1.529 (3)   | C45—C46       | 1.527 (3)   |
| C14—H14    | 1.0000      | C45—H45A      | 0.9900      |
| C15—C16    | 1.544 (3)   | C45—H45B      | 0.9900      |
| C15—H15A   | 0.9900      | C46—C47       | 1.333 (3)   |
| C15—H15B   | 0.9900      | C47—C48       | 1.492 (4)   |
| C16—C17    | 1.530 (3)   | C47—H47       | 0.9500      |
| C16—H16A   | 0.9900      | C48—H48A      | 0.9800      |
| C16—H16B   | 0.9900      | C48—H48B      | 0.9800      |
| C17—C18    | 1.338 (3)   | C48—H48C      | 0.9800      |
| C18—C19    | 1.507 (3)   | C49—H49A      | 0.9800      |
| C18—H18    | 0.9500      | C49—H49B      | 0.9800      |
| C19—H19A   | 0.9800      | C49—H49C      | 0.9800      |
| C19—H19B   | 0.9800      | C50—H50A      | 0.9800      |
| C19—H19C   | 0.9800      | C50—H50B      | 0.9800      |
| C20—H20A   | 0.9800      | C50—H50C      | 0.9800      |
| C20—H20B   | 0.9800      | C51—C54       | 1.487 (3)   |
| C20—H20C   | 0.9800      | C52—C53       | 1.485 (3)   |
| C21—H21A   | 0.9800      | C53—C58       | 1.384 (3)   |
| C21—H21B   | 0.9800      | C53—C54       | 1.384 (3)   |
| C21—H21C   | 0.9800      | C54—C55       | 1.381 (3)   |
| C22—C25    | 1.487 (3)   | C55—C56       | 1.397 (3)   |
| C23—C24    | 1.491 (3)   | C55—H55       | 0.9500      |
| C24—C25    | 1.384 (3)   | C56—C57       | 1.391 (3)   |
| C24—C29    | 1.387 (3)   | C56—H56       | 0.9500      |
| C25—C26    | 1.381 (3)   | C57—C58       | 1.393 (3)   |
| C26—C27    | 1.394 (3)   | C57—H57       | 0.9500      |
| C26—H26    | 0.9500      | C58—H58       | 0.9500      |
| C22—N1—C23 | 110.72 (17) | C24—C29—C28   | 116.9 (2)   |
| C22—N1—C3  | 118.85 (16) | C24—C29—H29   | 121.5       |
| C23—N1—C3  | 130.38 (17) | C28—C29—H29   | 121.5       |
| C52—N2—C51 | 111.04 (16) | C31—C30—C39   | 114.93 (17) |
| C52—N2—C32 | 125.32 (17) | C31—C30—H30A  | 108.5       |
| C51—N2—C32 | 123.24 (16) | C39—C30—H30A  | 108.5       |
| C2—C1—C10  | 114.37 (17) | C31—C30—H30B  | 108.5       |
| C2—C1—H1A  | 108.7       | C39—C30—H30B  | 108.5       |
| C10—C1—H1A | 108.7       | H30A—C30—H30B | 107.5       |
| C2—C1—H1B  | 108.7       | C32—C31—C30   | 110.37 (16) |
| C10—C1—H1B | 108.7       | C32—C31—H31A  | 109.6       |
| H1A—C1—H1B | 107.6       | C30—C31—H31A  | 109.6       |
| C1—C2—C3   | 116.35 (17) | C32—C31—H31B  | 109.6       |
| C1—C2—H2A  | 108.2       | C30—C31—H31B  | 109.6       |

|            |             |               |             |
|------------|-------------|---------------|-------------|
| C3—C2—H2A  | 108.2       | H31A—C31—H31B | 108.1       |
| C1—C2—H2B  | 108.2       | N2—C32—C31    | 111.04 (16) |
| C3—C2—H2B  | 108.2       | N2—C32—C33    | 112.05 (17) |
| H2A—C2—H2B | 107.4       | C31—C32—C33   | 110.65 (17) |
| N1—C3—C4   | 111.16 (16) | N2—C32—H32    | 107.6       |
| N1—C3—C2   | 115.98 (16) | C31—C32—H32   | 107.6       |
| C4—C3—C2   | 111.20 (17) | C33—C32—H32   | 107.6       |
| N1—C3—H3   | 105.9       | C32—C33—C34   | 112.69 (16) |
| C4—C3—H3   | 105.9       | C32—C33—H33A  | 109.1       |
| C2—C3—H3   | 105.9       | C34—C33—H33A  | 109.1       |
| C5—C4—C3   | 112.65 (16) | C32—C33—H33B  | 109.1       |
| C5—C4—H4A  | 109.1       | C34—C33—H33B  | 109.1       |
| C3—C4—H4A  | 109.1       | H33A—C33—H33B | 107.8       |
| C5—C4—H4B  | 109.1       | C35—C34—C33   | 110.41 (16) |
| C3—C4—H4B  | 109.1       | C35—C34—C39   | 112.75 (15) |
| H4A—C4—H4B | 107.8       | C33—C34—C39   | 112.90 (16) |
| C4—C5—C6   | 111.40 (16) | C35—C34—H34   | 106.8       |
| C4—C5—C10  | 112.30 (16) | C33—C34—H34   | 106.8       |
| C6—C5—C10  | 112.24 (16) | C39—C34—H34   | 106.8       |
| C4—C5—H5   | 106.8       | C36—C35—C34   | 112.41 (16) |
| C6—C5—H5   | 106.8       | C36—C35—H35A  | 109.1       |
| C10—C5—H5  | 106.8       | C34—C35—H35A  | 109.1       |
| C7—C6—C5   | 111.21 (15) | C36—C35—H35B  | 109.1       |
| C7—C6—H6A  | 109.4       | C34—C35—H35B  | 109.1       |
| C5—C6—H6A  | 109.4       | H35A—C35—H35B | 107.9       |
| C7—C6—H6B  | 109.4       | C35—C36—C37   | 112.34 (17) |
| C5—C6—H6B  | 109.4       | C35—C36—H36A  | 109.1       |
| H6A—C6—H6B | 108.0       | C37—C36—H36A  | 109.1       |
| C6—C7—C8   | 112.34 (16) | C35—C36—H36B  | 109.1       |
| C6—C7—H7A  | 109.1       | C37—C36—H36B  | 109.1       |
| C8—C7—H7A  | 109.1       | H36A—C36—H36B | 107.9       |
| C6—C7—H7B  | 109.1       | C43—C37—C36   | 111.88 (16) |
| C8—C7—H7B  | 109.1       | C43—C37—C38   | 108.82 (15) |
| H7A—C7—H7B | 107.9       | C36—C37—C38   | 110.27 (15) |
| C14—C8—C7  | 111.70 (16) | C43—C37—H37   | 108.6       |
| C14—C8—C9  | 108.70 (15) | C36—C37—H37   | 108.6       |
| C7—C8—C9   | 110.37 (15) | C38—C37—H37   | 108.6       |
| C14—C8—H8  | 108.7       | C40—C38—C37   | 111.70 (16) |
| C7—C8—H8   | 108.7       | C40—C38—C39   | 113.34 (17) |
| C9—C8—H8   | 108.7       | C37—C38—C39   | 112.50 (15) |
| C11—C9—C8  | 112.36 (16) | C40—C38—H38   | 106.2       |
| C11—C9—C10 | 113.73 (16) | C37—C38—H38   | 106.2       |
| C8—C9—C10  | 111.96 (15) | C39—C38—H38   | 106.2       |
| C11—C9—H9  | 106.0       | C50—C39—C34   | 110.48 (15) |
| C8—C9—H9   | 106.0       | C50—C39—C38   | 110.56 (15) |
| C10—C9—H9  | 106.0       | C34—C39—C38   | 108.50 (15) |
| C21—C10—C1 | 109.42 (16) | C50—C39—C30   | 109.65 (16) |
| C21—C10—C5 | 112.02 (15) | C34—C39—C30   | 107.93 (15) |

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| C1—C10—C5     | 107.91 (15) | C38—C39—C30   | 109.66 (15) |
| C21—C10—C9    | 111.78 (16) | C41—C40—C38   | 113.94 (18) |
| C1—C10—C9     | 108.84 (15) | C41—C40—H40A  | 108.8       |
| C5—C10—C9     | 106.73 (15) | C38—C40—H40A  | 108.8       |
| C12—C11—C9    | 114.24 (16) | C41—C40—H40B  | 108.8       |
| C12—C11—H11A  | 108.7       | C38—C40—H40B  | 108.8       |
| C9—C11—H11A   | 108.7       | H40A—C40—H40B | 107.7       |
| C12—C11—H11B  | 108.7       | C40—C41—C42   | 111.44 (17) |
| C9—C11—H11B   | 108.7       | C40—C41—H41A  | 109.3       |
| H11A—C11—H11B | 107.6       | C42—C41—H41A  | 109.3       |
| C13—C12—C11   | 110.97 (16) | C40—C41—H41B  | 109.3       |
| C13—C12—H12A  | 109.4       | C42—C41—H41B  | 109.3       |
| C11—C12—H12A  | 109.4       | H41A—C41—H41B | 108.0       |
| C13—C12—H12B  | 109.4       | C46—C42—C41   | 118.69 (17) |
| C11—C12—H12B  | 109.4       | C46—C42—C43   | 100.25 (17) |
| H12A—C12—H12B | 108.0       | C41—C42—C43   | 107.07 (16) |
| C17—C13—C12   | 118.33 (16) | C46—C42—C49   | 107.39 (16) |
| C17—C13—C20   | 106.99 (16) | C41—C42—C49   | 110.45 (19) |
| C12—C13—C20   | 110.40 (17) | C43—C42—C49   | 112.71 (16) |
| C17—C13—C14   | 100.73 (15) | C37—C43—C44   | 119.13 (16) |
| C12—C13—C14   | 107.40 (16) | C37—C43—C42   | 114.27 (16) |
| C20—C13—C14   | 112.77 (15) | C44—C43—C42   | 104.49 (17) |
| C8—C14—C15    | 119.19 (16) | C37—C43—H43   | 106.0       |
| C8—C14—C13    | 114.32 (16) | C44—C43—H43   | 106.0       |
| C15—C14—C13   | 104.56 (16) | C42—C43—H43   | 106.0       |
| C8—C14—H14    | 105.9       | C43—C44—C45   | 103.37 (17) |
| C15—C14—H14   | 105.9       | C43—C44—H44A  | 111.1       |
| C13—C14—H14   | 105.9       | C45—C44—H44A  | 111.1       |
| C14—C15—C16   | 102.55 (16) | C43—C44—H44B  | 111.1       |
| C14—C15—H15A  | 111.3       | C45—C44—H44B  | 111.1       |
| C16—C15—H15A  | 111.3       | H44A—C44—H44B | 109.1       |
| C14—C15—H15B  | 111.3       | C46—C45—C44   | 106.23 (17) |
| C16—C15—H15B  | 111.3       | C46—C45—H45A  | 110.5       |
| H15A—C15—H15B | 109.2       | C44—C45—H45A  | 110.5       |
| C17—C16—C15   | 106.14 (17) | C46—C45—H45B  | 110.5       |
| C17—C16—H16A  | 110.5       | C44—C45—H45B  | 110.5       |
| C15—C16—H16A  | 110.5       | H45A—C45—H45B | 108.7       |
| C17—C16—H16B  | 110.5       | C47—C46—C42   | 130.5 (2)   |
| C15—C16—H16B  | 110.5       | C47—C46—C45   | 122.0 (2)   |
| H16A—C16—H16B | 108.7       | C42—C46—C45   | 107.50 (17) |
| C18—C17—C13   | 129.17 (19) | C46—C47—C48   | 129.1 (2)   |
| C18—C17—C16   | 122.9 (2)   | C46—C47—H47   | 115.4       |
| C13—C17—C16   | 107.93 (16) | C48—C47—H47   | 115.4       |
| C17—C18—C19   | 127.9 (2)   | C47—C48—H48A  | 109.5       |
| C17—C18—H18   | 116.0       | C47—C48—H48B  | 109.5       |
| C19—C18—H18   | 116.0       | H48A—C48—H48B | 109.5       |
| C18—C19—H19A  | 109.5       | C47—C48—H48C  | 109.5       |
| C18—C19—H19B  | 109.5       | H48A—C48—H48C | 109.5       |

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| H19A—C19—H19B | 109.5        | H48B—C48—H48C   | 109.5       |
| C18—C19—H19C  | 109.5        | C42—C49—H49A    | 109.5       |
| H19A—C19—H19C | 109.5        | C42—C49—H49B    | 109.5       |
| H19B—C19—H19C | 109.5        | H49A—C49—H49B   | 109.5       |
| C13—C20—H20A  | 109.5        | C42—C49—H49C    | 109.5       |
| C13—C20—H20B  | 109.5        | H49A—C49—H49C   | 109.5       |
| H20A—C20—H20B | 109.5        | H49B—C49—H49C   | 109.5       |
| C13—C20—H20C  | 109.5        | C39—C50—H50A    | 109.5       |
| H20A—C20—H20C | 109.5        | C39—C50—H50B    | 109.5       |
| H20B—C20—H20C | 109.5        | H50A—C50—H50B   | 109.5       |
| C10—C21—H21A  | 109.5        | C39—C50—H50C    | 109.5       |
| C10—C21—H21B  | 109.5        | H50A—C50—H50C   | 109.5       |
| H21A—C21—H21B | 109.5        | H50B—C50—H50C   | 109.5       |
| C10—C21—H21C  | 109.5        | O3—C51—N2       | 125.04 (19) |
| H21A—C21—H21C | 109.5        | O3—C51—C54      | 128.65 (19) |
| H21B—C21—H21C | 109.5        | N2—C51—C54      | 106.28 (16) |
| O1—C22—N1     | 124.81 (19)  | O4—C52—N2       | 125.56 (19) |
| O1—C22—C25    | 128.1 (2)    | O4—C52—C53      | 128.19 (18) |
| N1—C22—C25    | 107.07 (16)  | N2—C52—C53      | 106.25 (17) |
| O2—C23—N1     | 127.59 (19)  | C58—C53—C54     | 121.97 (19) |
| O2—C23—C24    | 126.24 (18)  | C58—C53—C52     | 129.63 (19) |
| N1—C23—C24    | 106.16 (17)  | C54—C53—C52     | 108.38 (17) |
| C25—C24—C29   | 121.8 (2)    | C55—C54—C53     | 121.63 (18) |
| C25—C24—C23   | 108.49 (17)  | C55—C54—C51     | 130.32 (18) |
| C29—C24—C23   | 129.69 (19)  | C53—C54—C51     | 107.99 (17) |
| C26—C25—C24   | 121.4 (2)    | C54—C55—C56     | 116.76 (19) |
| C26—C25—C22   | 131.0 (2)    | C54—C55—H55     | 121.6       |
| C24—C25—C22   | 107.53 (18)  | C56—C55—H55     | 121.6       |
| C25—C26—C27   | 117.1 (2)    | C57—C56—C55     | 121.7 (2)   |
| C25—C26—H26   | 121.4        | C57—C56—H56     | 119.2       |
| C27—C26—H26   | 121.4        | C55—C56—H56     | 119.2       |
| C28—C27—C26   | 121.4 (2)    | C56—C57—C58     | 120.95 (19) |
| C28—C27—H27   | 119.3        | C56—C57—H57     | 119.5       |
| C26—C27—H27   | 119.3        | C58—C57—H57     | 119.5       |
| C27—C28—C29   | 121.3 (2)    | C53—C58—C57     | 116.98 (19) |
| C27—C28—H28   | 119.4        | C53—C58—H58     | 121.5       |
| C29—C28—H28   | 119.4        | C57—C58—H58     | 121.5       |
|               |              |                 |             |
| C10—C1—C2—C3  | -45.7 (2)    | C39—C30—C31—C32 | 37.1 (3)    |
| C22—N1—C3—C4  | 90.0 (2)     | C52—N2—C32—C31  | 55.2 (3)    |
| C23—N1—C3—C4  | -87.3 (2)    | C51—N2—C32—C31  | -117.0 (2)  |
| C22—N1—C3—C2  | -141.66 (18) | C52—N2—C32—C33  | -69.1 (2)   |
| C23—N1—C3—C2  | 41.1 (3)     | C51—N2—C32—C33  | 118.7 (2)   |
| C1—C2—C3—N1   | -85.1 (2)    | C30—C31—C32—N2  | 171.34 (17) |
| C1—C2—C3—C4   | 43.2 (2)     | C30—C31—C32—C33 | -63.6 (2)   |
| N1—C3—C4—C5   | 81.2 (2)     | N2—C32—C33—C34  | 150.33 (16) |
| C2—C3—C4—C5   | -49.6 (2)    | C31—C32—C33—C34 | 25.8 (2)    |
| C3—C4—C5—C6   | -173.49 (16) | C32—C33—C34—C35 | 163.46 (16) |

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| C3—C4—C5—C10    | 59.6 (2)     | C32—C33—C34—C39 | 36.2 (2)     |
| C4—C5—C6—C7     | 175.86 (16)  | C33—C34—C35—C36 | 178.74 (16)  |
| C10—C5—C6—C7    | -57.2 (2)    | C39—C34—C35—C36 | -53.9 (2)    |
| C5—C6—C7—C8     | 53.6 (2)     | C34—C35—C36—C37 | 53.5 (2)     |
| C6—C7—C8—C14    | -174.54 (14) | C35—C36—C37—C43 | -175.55 (15) |
| C6—C7—C8—C9     | -53.5 (2)    | C35—C36—C37—C38 | -54.3 (2)    |
| C14—C8—C9—C11   | -50.5 (2)    | C43—C37—C38—C40 | -51.4 (2)    |
| C7—C8—C9—C11    | -173.29 (16) | C36—C37—C38—C40 | -174.50 (16) |
| C14—C8—C9—C10   | -179.90 (15) | C43—C37—C38—C39 | 179.77 (15)  |
| C7—C8—C9—C10    | 57.3 (2)     | C36—C37—C38—C39 | 56.7 (2)     |
| C2—C1—C10—C21   | -71.3 (2)    | C35—C34—C39—C50 | -67.3 (2)    |
| C2—C1—C10—C5    | 50.8 (2)     | C33—C34—C39—C50 | 58.7 (2)     |
| C2—C1—C10—C9    | 166.25 (16)  | C35—C34—C39—C38 | 54.08 (19)   |
| C4—C5—C10—C21   | 62.4 (2)     | C33—C34—C39—C38 | -179.93 (14) |
| C6—C5—C10—C21   | -64.1 (2)    | C35—C34—C39—C30 | 172.84 (16)  |
| C4—C5—C10—C1    | -58.15 (19)  | C33—C34—C39—C30 | -61.2 (2)    |
| C6—C5—C10—C1    | 175.42 (16)  | C40—C38—C39—C50 | -62.6 (2)    |
| C4—C5—C10—C9    | -174.99 (14) | C37—C38—C39—C50 | 65.3 (2)     |
| C6—C5—C10—C9    | 58.58 (19)   | C40—C38—C39—C34 | 176.08 (15)  |
| C11—C9—C10—C21  | -64.7 (2)    | C37—C38—C39—C34 | -55.99 (19)  |
| C8—C9—C10—C21   | 64.1 (2)     | C40—C38—C39—C30 | 58.4 (2)     |
| C11—C9—C10—C1   | 56.3 (2)     | C37—C38—C39—C30 | -173.65 (16) |
| C8—C9—C10—C1    | -174.97 (15) | C31—C30—C39—C50 | -97.8 (2)    |
| C11—C9—C10—C5   | 172.55 (14)  | C31—C30—C39—C34 | 22.6 (2)     |
| C8—C9—C10—C5    | -58.75 (18)  | C31—C30—C39—C38 | 140.61 (18)  |
| C8—C9—C11—C12   | 50.1 (2)     | C37—C38—C40—C41 | 51.1 (2)     |
| C10—C9—C11—C12  | 178.62 (15)  | C39—C38—C40—C41 | 179.42 (16)  |
| C9—C11—C12—C13  | -53.2 (2)    | C38—C40—C41—C42 | -54.0 (2)    |
| C11—C12—C13—C17 | 168.86 (17)  | C40—C41—C42—C46 | 168.13 (19)  |
| C11—C12—C13—C20 | -67.4 (2)    | C40—C41—C42—C43 | 55.8 (2)     |
| C11—C12—C13—C14 | 55.9 (2)     | C40—C41—C42—C49 | -67.3 (2)    |
| C7—C8—C14—C15   | -54.9 (2)    | C36—C37—C43—C44 | -54.6 (2)    |
| C9—C8—C14—C15   | -176.95 (17) | C38—C37—C43—C44 | -176.72 (17) |
| C7—C8—C14—C13   | -179.57 (14) | C36—C37—C43—C42 | -179.08 (15) |
| C9—C8—C14—C13   | 58.41 (19)   | C38—C37—C43—C42 | 58.84 (19)   |
| C17—C13—C14—C8  | 174.11 (14)  | C46—C42—C43—C37 | 174.66 (15)  |
| C12—C13—C14—C8  | -61.45 (19)  | C41—C42—C43—C37 | -60.9 (2)    |
| C20—C13—C14—C8  | 60.4 (2)     | C49—C42—C43—C37 | 60.8 (2)     |
| C17—C13—C14—C15 | 42.02 (17)   | C46—C42—C43—C44 | 42.74 (17)   |
| C12—C13—C14—C15 | 166.46 (15)  | C41—C42—C43—C44 | 167.20 (16)  |
| C20—C13—C14—C15 | -71.7 (2)    | C49—C42—C43—C44 | -71.1 (2)    |
| C8—C14—C15—C16  | -168.54 (17) | C37—C43—C44—C45 | -165.23 (18) |
| C13—C14—C15—C16 | -39.30 (19)  | C42—C43—C44—C45 | -36.2 (2)    |
| C14—C15—C16—C17 | 20.9 (2)     | C43—C44—C45—C46 | 15.0 (2)     |
| C12—C13—C17—C18 | 34.9 (3)     | C41—C42—C46—C47 | 31.1 (3)     |
| C20—C13—C17—C18 | -90.4 (2)    | C43—C42—C46—C47 | 147.1 (2)    |
| C14—C13—C17—C18 | 151.6 (2)    | C49—C42—C46—C47 | -95.0 (3)    |
| C12—C13—C17—C16 | -145.12 (17) | C41—C42—C46—C45 | -149.3 (2)   |

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| C20—C13—C17—C16 | 89.52 (19)   | C43—C42—C46—C45 | -33.23 (19)  |
| C14—C13—C17—C16 | -28.51 (18)  | C49—C42—C46—C45 | 84.7 (2)     |
| C15—C16—C17—C18 | -174.9 (2)   | C44—C45—C46—C47 | -168.5 (2)   |
| C15—C16—C17—C13 | 5.1 (2)      | C44—C45—C46—C42 | 11.8 (2)     |
| C13—C17—C18—C19 | 0.0 (4)      | C42—C46—C47—C48 | 3.4 (4)      |
| C16—C17—C18—C19 | -179.9 (2)   | C45—C46—C47—C48 | -176.2 (2)   |
| C23—N1—C22—O1   | 178.1 (2)    | C52—N2—C51—O3   | -179.8 (2)   |
| C3—N1—C22—O1    | 0.3 (3)      | C32—N2—C51—O3   | -6.6 (3)     |
| C23—N1—C22—C25  | -1.5 (2)     | C52—N2—C51—C54  | -1.4 (2)     |
| C3—N1—C22—C25   | -179.25 (16) | C32—N2—C51—C54  | 171.68 (17)  |
| C22—N1—C23—O2   | 179.7 (2)    | C51—N2—C52—O4   | -178.0 (2)   |
| C3—N1—C23—O2    | -2.8 (4)     | C32—N2—C52—O4   | 9.0 (3)      |
| C22—N1—C23—C24  | 0.6 (2)      | C51—N2—C52—C53  | 2.2 (2)      |
| C3—N1—C23—C24   | 178.00 (18)  | C32—N2—C52—C53  | -170.77 (17) |
| O2—C23—C24—C25  | -178.5 (2)   | O4—C52—C53—C58  | -3.6 (4)     |
| N1—C23—C24—C25  | 0.6 (2)      | N2—C52—C53—C58  | 176.2 (2)    |
| O2—C23—C24—C29  | -0.2 (4)     | O4—C52—C53—C54  | 178.1 (2)    |
| N1—C23—C24—C29  | 179.0 (2)    | N2—C52—C53—C54  | -2.1 (2)     |
| C29—C24—C25—C26 | -0.6 (3)     | C58—C53—C54—C55 | 0.3 (3)      |
| C23—C24—C25—C26 | 177.9 (2)    | C52—C53—C54—C55 | 178.75 (19)  |
| C29—C24—C25—C22 | -179.99 (19) | C58—C53—C54—C51 | -177.17 (19) |
| C23—C24—C25—C22 | -1.5 (2)     | C52—C53—C54—C51 | 1.3 (2)      |
| O1—C22—C25—C26  | 3.0 (4)      | O3—C51—C54—C55  | 1.1 (4)      |
| N1—C22—C25—C26  | -177.5 (2)   | N2—C51—C54—C55  | -177.2 (2)   |
| O1—C22—C25—C24  | -177.7 (2)   | O3—C51—C54—C53  | 178.3 (2)    |
| N1—C22—C25—C24  | 1.8 (2)      | N2—C51—C54—C53  | 0.1 (2)      |
| C24—C25—C26—C27 | 0.2 (3)      | C53—C54—C55—C56 | -1.6 (3)     |
| C22—C25—C26—C27 | 179.4 (2)    | C51—C54—C55—C56 | 175.3 (2)    |
| C25—C26—C27—C28 | 0.4 (4)      | C54—C55—C56—C57 | 1.4 (3)      |
| C26—C27—C28—C29 | -0.6 (4)     | C55—C56—C57—C58 | -0.1 (3)     |
| C25—C24—C29—C28 | 0.4 (3)      | C54—C53—C58—C57 | 1.0 (3)      |
| C23—C24—C29—C28 | -177.8 (2)   | C52—C53—C58—C57 | -177.01 (19) |
| C27—C28—C29—C24 | 0.2 (4)      | C56—C57—C58—C53 | -1.2 (3)     |

Hydrogen-bond geometry ( $\text{\AA}$ ,  $^\circ$ )

| <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> |
|---|-------------|---------------------|----------------------------|-------------------------------|
| C1—H1 <i>B</i> $\cdots$ O2                  | 0.99        | 2.25                | 3.027 (3)                  | 134                           |
| C3—H3 $\cdots$ O1                           | 1.00        | 2.41                | 2.823 (3)                  | 104                           |
| C5—H5 $\cdots$ O4 <sup>i</sup>              | 1.00        | 2.59                | 3.237 (3)                  | 122                           |
| C21—H21 <i>C</i> $\cdots$ O3 <sup>ii</sup>  | 0.98        | 2.60                | 3.501 (3)                  | 153                           |
| C31—H31 <i>A</i> $\cdots$ O2 <sup>iii</sup> | 0.99        | 2.47                | 3.361 (3)                  | 150                           |
| C31—H31 <i>B</i> $\cdots$ O4                | 0.99        | 2.58                | 3.144 (3)                  | 116                           |
| C32—H32 $\cdots$ O3                         | 1.00        | 2.49                | 2.917 (3)                  | 105                           |

Symmetry codes: (i)  $x, y, z+1$ ; (ii)  $x-1, y, z$ ; (iii)  $x+1, y, z-1$ .