

4'-Amino-2,2''-dioxo-2,2'',3,3''-tetrahydro-1H-indole-3-spiro-1'-cyclopent-3'-ene-2'-spiro-3''-1H-indole-3',5',5'-tricarbonitrile dihydrate

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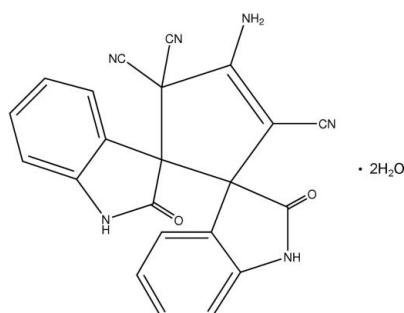
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Key indicators: single-crystal X-ray study; $T = 293\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$; R factor = 0.042; wR factor = 0.109; data-to-parameter ratio = 8.2.

In the title compound, $\text{C}_{22}\text{H}_{12}\text{N}_6\text{O}_2\cdot 2\text{H}_2\text{O}$, the cyclopentene ring adopts an envelope conformation, with the spiro C atom bonded to the dicyano-substituted C atom deviating by $0.437(2)\text{ \AA}$ from the plane of the remaining four atoms in the ring. The puckering and smallest displacement asymmetry parameters for the ring are $q_2 = 0.275(2)\text{ \AA}$, $\varphi = 212.4(4)^\circ$ and $\Delta_s(\text{C}_2) = 2.7(2)$. The dihedral angle between the two indole groups is $60.1(1)^\circ$. The structure contains intermolecular N—H···O hydrogen bonds involving the indole groups and O—H···O and O—H···N hydrogen bonds involving the water molecules.

Related literature

For related literature, see: Akai *et al.* (2004); Cremer & Pople (1975); Gallagher *et al.* (1985); Nagata *et al.* (2001); Nardelli (1983); Williams & Cox (2003); Zaveri *et al.* (2004).



Experimental

Crystal data

$\text{C}_{22}\text{H}_{12}\text{N}_6\text{O}_2\cdot 2\text{H}_2\text{O}$	$V = 2057.8(3)\text{ \AA}^3$
$M_r = 428.41$	$Z = 4$
Orthorhombic, $Pna2_1$	Mo $K\alpha$ radiation
$a = 17.1850(16)\text{ \AA}$	$\mu = 0.10\text{ mm}^{-1}$
$b = 8.9849(9)\text{ \AA}$	$T = 293(2)\text{ K}$
$c = 13.3275(13)\text{ \AA}$	$0.25 \times 0.24 \times 0.20\text{ mm}$

Data collection

Bruker SMART APEX CCD diffractometer	2551 independent reflections
Absorption correction: none	2367 reflections with $I > 2\sigma(I)$
16968 measured reflections	$R_{\text{int}} = 0.022$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.042$	H atoms treated by a mixture of independent and constrained refinement
$wR(F^2) = 0.108$	$\Delta\rho_{\text{max}} = 0.27\text{ e \AA}^{-3}$
$S = 1.06$	$\Delta\rho_{\text{min}} = -0.14\text{ e \AA}^{-3}$
2551 reflections	
313 parameters	
9 restraints	

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
N1—H1···O2 ⁱ	0.86	2.09	2.905 (2)	158
N2—H2···O3 ⁱ	0.86	2.22	3.032 (3)	157
O3—H3A···O1 ⁱⁱ	0.85 (1)	2.00 (2)	2.795 (3)	157 (4)
O3—H3B···N6 ⁱⁱⁱ	0.85 (1)	2.22 (5)	2.834 (3)	129 (5)
O3—H3B···O4 ^{iv}	0.85 (1)	2.62 (3)	3.381 (4)	149 (5)
O4—H4A···N4 ^v	0.85 (1)	2.38 (3)	3.171 (5)	155 (7)
O4—H4B···N3	0.85 (1)	2.68 (4)	3.392 (4)	142 (6)
N5—H5A···O3 ^{vi}	0.90 (1)	1.97 (1)	2.860 (3)	175 (3)
N5—H5B···O4 ^{vi}	0.90 (1)	2.17 (1)	3.061 (4)	171 (3)

Symmetry codes: (i) $-x + \frac{1}{2}, y - \frac{1}{2}, z - \frac{1}{2}$; (ii) $x, y + 1, z + 1$; (iii) $x, y, z + 1$; (iv) $-x, -y + 1, z + \frac{1}{2}$; (v) $-x, -y, z + \frac{1}{2}$; (vi) $-x, -y + 1, z - \frac{1}{2}$.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97* and *PARST* (Nardelli, 1995).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BI2273).

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supporting information

Acta Cryst. (2008). E64, o501–o502 [doi:10.1107/S1600536808002146]

4'-Amino-2,2''-dioxo-2,2'',3,3''-tetrahydro-1H-indole-3-spiro-1'-cyclopent-3'-ene-2'-spiro-3''-1H-indole-3',5',5'-tricarbonitrile dihydrate

D. Gayathri, D. Velmurugan, G. Shanthi, P. T. Perumal and K. Ravikumar

S1. Comment

The indole template is generally recognized as an important structure in medicinal chemistry, and in particular, oxindoles are important constituents of natural indole alkaloids as well as drugs in development and also in the clinic (Akai *et al.*, 2004). Among them, the spiro-oxindole framework is an important structural motif in biologically relevant compounds such as natural products and pharmaceuticals (Williams & Cox, 2003). The oxindole motif is present in the anti-Parkinson's drug ropinirole (Gallagher *et al.*, 1985), in non-opioid nociceptin receptor ligands (Zaveri *et al.*, 2004), and in the growth hormone secretagogues (Nagata *et al.*, 2001). As the title compound is biologically important, we have determined its crystal structure using X-ray diffraction.

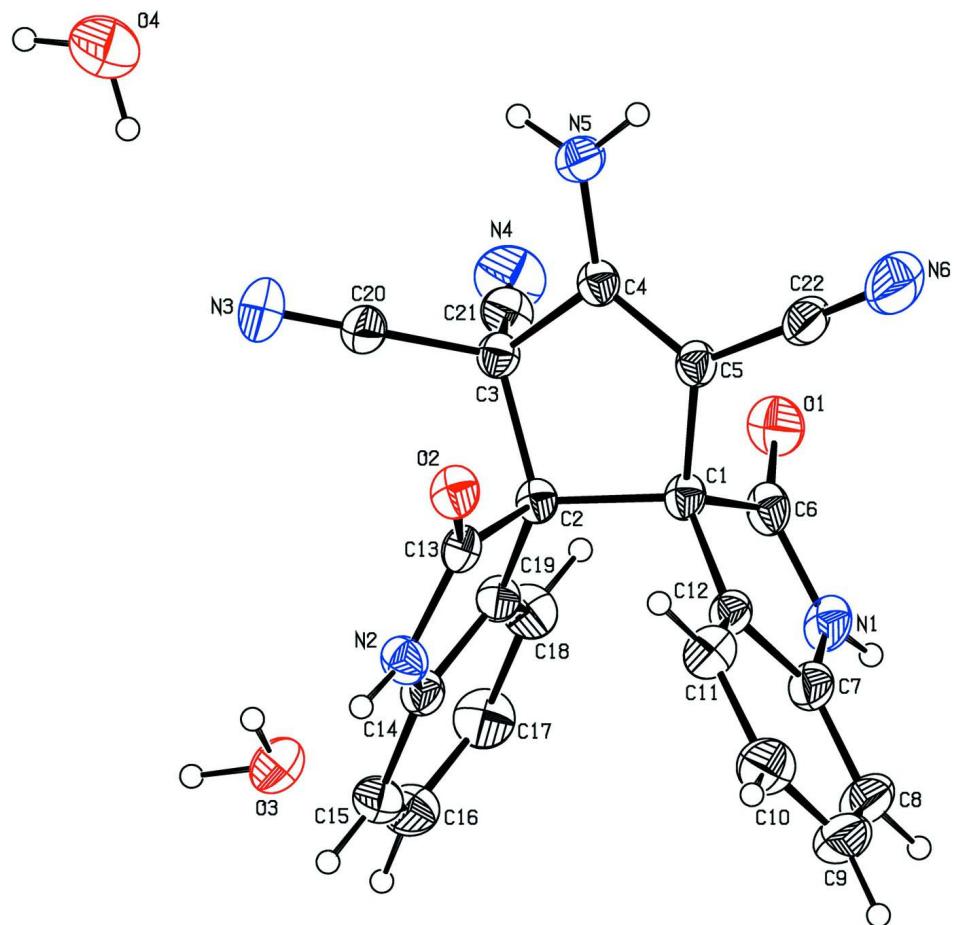
One of the oxindole moieties (C2/C13/N2/C14/C19) is planar with atom O2 deviating by 0.127 (1) Å from the plane of five-membered ring. The dihedral angle between the five- (C2/C13/N2/C14/C19) and six-membered (C14—C19) rings is 1.6 (1)°. The cyclopentene ring adopts an envelope conformation with C2 deviating by 0.437 (2) Å from the plane of the rest of the atoms in the ring. The five-membered ring (N1/C6/C1/C12/C7) in the oxindole moieties adopts a slightly twisted conformation. The puckering parameters (Cremer & Pople, 1975) and the smallest displacement asymmetry parameters (Nardelli, 1983) for the cyclopentene ring and the five-membered ring (N1/C6/C1/C12/C7) are $q_2 = 0.275$ (2) Å and 0.091 (2) Å, $\phi = 212.4$ (4)° and 56.3 (12)° and $\Delta_s(C_2) = 2.7$ (2) and $\Delta_2(C_7) = 0.5$ (2), respectively.

S2. Experimental

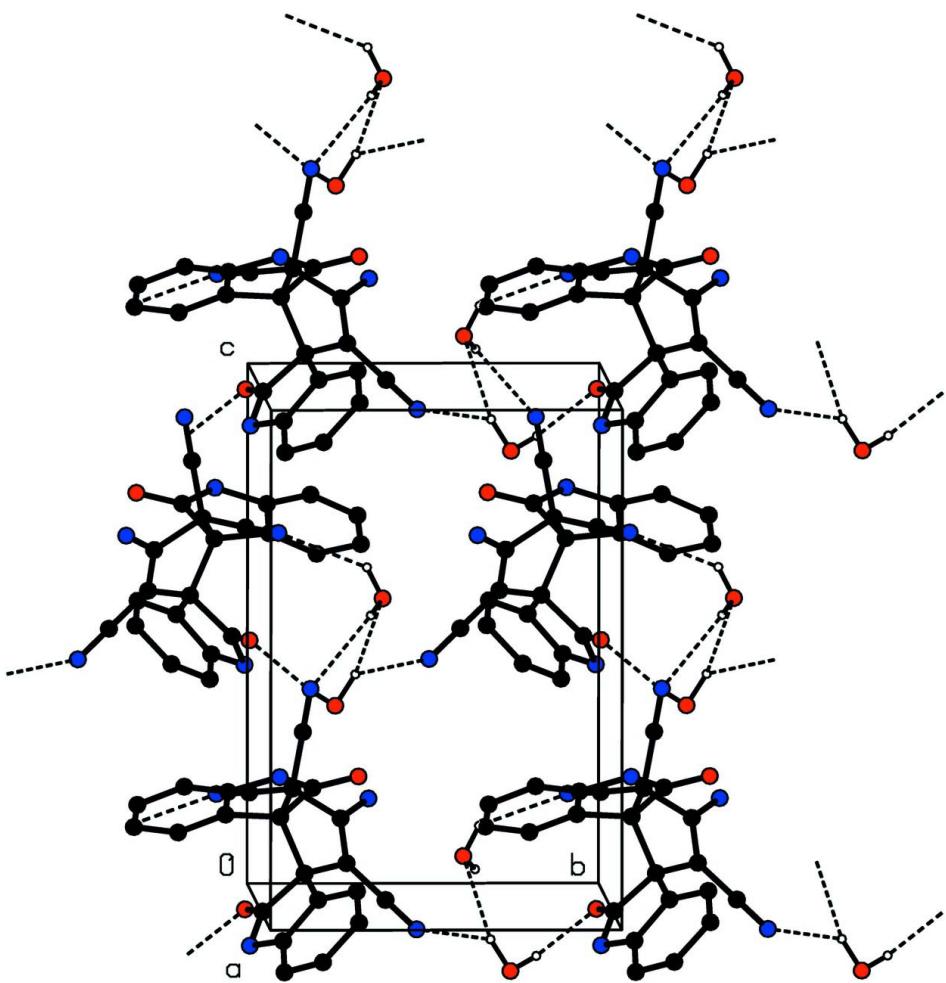
To a stirred mixture of isatylidene malononitrile (2 mmol) and Hantzsch dihydropyridine ester (1 mmol) in ethanol (10 ml), a catalytic amount of indium(III) chloride (20 mol%) was added and the mixture was stirred at room temperature for about 1–2 h. After complete conversion (as indicated by TLC), the reaction mixture was poured into water and extracted with ethyl acetate (2×15 ml). The combined extracts were dried over anhydrous Na_2SO_4 and concentrated *in vacuo*. The resulting product was purified by column chromatography on silica gel (Merck, 60–120 mesh, ethyl acetate-hexane, 4:6) to afford the pure product in 88% yield as a white solid. Crystals were grown by slow evaporation from ethanol.

S3. Refinement

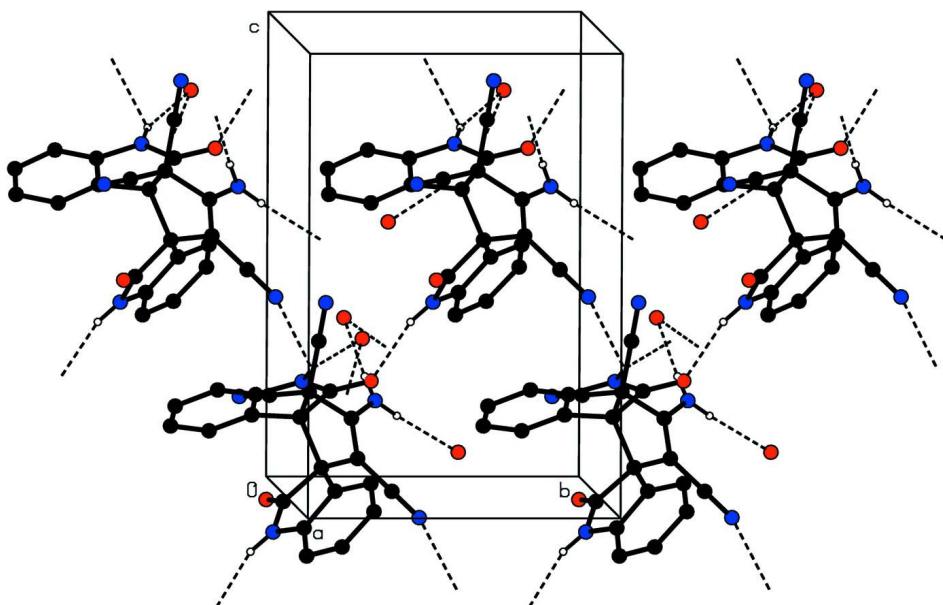
H atoms of the NH_2 group and water molecules were located in a difference Fourier map and refined with the N—H and O—H distances restrained to 0.90 (1) and 0.85 (1) Å, respectively. H atoms bound to C atoms and the N atoms of the indole rings were placed geometrically and refined using a riding model with $d(\text{C—H}) = 0.93$ Å, $d(\text{N—H}) = 0.86$ Å, and $U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{C}/\text{N})$. In the absence of significant anomalous scattering effects, 2271 Friedel pairs were merged as equivalent data.

**Figure 1**

The molecular structure of title compound, showing 30% probability displacement ellipsoids for non-H atoms.

**Figure 2**

The molecular packing viewed down the a axis, showing $\text{O}—\text{H}\cdots\text{N}$ and $\text{O}—\text{H}\cdots\text{O}$ intermolecular interactions.

**Figure 3**

The molecular packing viewed down the *a* axis, showing N—H···O intermolecular interactions.

4'-Amino-2,2''-dioxo-2,2'',3,3''-tetrahydro-1*H*-indole-3-spiro-1'-cyclopent- 3'-ene-2'-spiro-3''-1*H*-indole-3',5',5'-tricarbonitrile dihydrate

Crystal data



$M_r = 428.41$

Orthorhombic, $Pna2_1$

Hall symbol: P 2c -2n

$a = 17.1850(16)$ Å

$b = 8.9849(9)$ Å

$c = 13.3275(13)$ Å

$V = 2057.8(3)$ Å³

$Z = 4$

$F(000) = 888$

$D_x = 1.383$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 2386 reflections

$\theta = 2.4\text{--}28.0^\circ$

$\mu = 0.10$ mm⁻¹

$T = 293$ K

Block, colorless

0.25 × 0.24 × 0.20 mm

Data collection

Bruker SMART APEX CCD
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

ω scans

16968 measured reflections

2551 independent reflections

2367 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.022$

$\theta_{\max} = 28.0^\circ$, $\theta_{\min} = 2.4^\circ$

$h = -22 \rightarrow 22$

$k = -11 \rightarrow 11$

$l = -17 \rightarrow 17$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.042$

$wR(F^2) = 0.108$

$S = 1.06$

2551 reflections

313 parameters

9 restraints

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map

Hydrogen site location: inferred from
neighbouring sites

H atoms treated by a mixture of independent
and constrained refinement

$$w = 1/[\sigma^2(F_o^2) + (0.0751P)^2 + 0.1068P]$$

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

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

$$\Delta\rho_{\max} = 0.27 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.14 \text{ e } \text{\AA}^{-3}$$

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 (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.19406 (12)	0.1506 (2)	0.03645 (16)	0.0341 (4)
C2	0.19661 (11)	0.0810 (2)	0.14549 (15)	0.0320 (4)
C3	0.11505 (12)	0.1206 (2)	0.19253 (16)	0.0352 (4)
C4	0.09089 (12)	0.2598 (2)	0.13339 (17)	0.0349 (4)
C5	0.13624 (13)	0.2738 (2)	0.05081 (16)	0.0354 (4)
C6	0.16280 (14)	0.0331 (2)	-0.03919 (16)	0.0414 (5)
C7	0.28704 (15)	0.0835 (3)	-0.08495 (19)	0.0452 (5)
C8	0.35505 (18)	0.0831 (4)	-0.1392 (2)	0.0614 (7)
H8A	0.3636	0.0143	-0.1902	0.074*
C9	0.41025 (19)	0.1889 (4)	-0.1150 (3)	0.0679 (8)
H9A	0.4570	0.1906	-0.1501	0.082*
C10	0.39785 (17)	0.2922 (4)	-0.0399 (3)	0.0612 (7)
H10A	0.4358	0.3629	-0.0256	0.073*
C11	0.32867 (16)	0.2907 (3)	0.0144 (2)	0.0489 (6)
H11A	0.3198	0.3607	0.0645	0.059*
C12	0.27339 (13)	0.1839 (2)	-0.00708 (16)	0.0375 (4)
C13	0.25886 (12)	0.1685 (2)	0.20696 (15)	0.0340 (4)
C14	0.29683 (12)	-0.0747 (2)	0.20308 (17)	0.0378 (4)
C15	0.34057 (15)	-0.2019 (3)	0.2170 (2)	0.0493 (6)
H15A	0.3882	-0.1992	0.2501	0.059*
C16	0.31063 (17)	-0.3336 (3)	0.1795 (2)	0.0564 (7)
H16A	0.3388	-0.4211	0.1878	0.068*
C17	0.24070 (18)	-0.3388 (3)	0.1305 (2)	0.0581 (7)
H17A	0.2221	-0.4293	0.1065	0.070*
C18	0.19709 (16)	-0.2097 (2)	0.1161 (2)	0.0489 (6)
H18A	0.1497	-0.2129	0.0824	0.059*
C19	0.22569 (13)	-0.0769 (2)	0.15289 (16)	0.0364 (4)
C20	0.12114 (15)	0.1500 (3)	0.30204 (19)	0.0449 (5)
C21	0.05602 (14)	0.0018 (3)	0.1798 (2)	0.0470 (5)
C22	0.12327 (16)	0.3857 (3)	-0.0210 (2)	0.0506 (6)
N1	0.22129 (14)	-0.0067 (2)	-0.09965 (16)	0.0486 (5)

H1	0.2189	-0.0785	-0.1422	0.058*
N2	0.31446 (11)	0.0717 (2)	0.23268 (15)	0.0413 (4)
H2	0.3564	0.0962	0.2639	0.050*
N3	0.12535 (18)	0.1713 (4)	0.38515 (19)	0.0706 (7)
N4	0.01054 (15)	-0.0884 (3)	0.1731 (3)	0.0727 (8)
N5	0.03222 (13)	0.3429 (3)	0.16605 (18)	0.0498 (5)
H5A	0.0025 (15)	0.311 (3)	0.2169 (17)	0.048 (7)*
H5B	0.011 (2)	0.417 (3)	0.130 (3)	0.070 (10)*
N6	0.1106 (2)	0.4754 (4)	-0.0785 (3)	0.0916 (11)
O1	0.09597 (11)	-0.0125 (2)	-0.04096 (16)	0.0567 (5)
O2	0.25526 (10)	0.29951 (16)	0.22853 (13)	0.0429 (4)
O3	0.05768 (12)	0.7461 (2)	0.83632 (15)	0.0540 (4)
H3A	0.073 (3)	0.830 (3)	0.858 (3)	0.107 (15)*
H3B	0.054 (4)	0.692 (4)	0.888 (2)	0.15 (2)*
O4	0.0238 (2)	0.3832 (4)	0.5501 (3)	0.0938 (9)
H4A	0.020 (4)	0.322 (6)	0.599 (4)	0.16 (3)*
H4B	0.066 (2)	0.360 (6)	0.521 (4)	0.13 (2)*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0439 (10)	0.0289 (9)	0.0294 (9)	0.0008 (8)	-0.0003 (8)	-0.0036 (7)
C2	0.0387 (9)	0.0268 (9)	0.0304 (9)	-0.0010 (7)	-0.0022 (8)	-0.0033 (7)
C3	0.0384 (9)	0.0334 (9)	0.0338 (10)	-0.0009 (8)	0.0006 (8)	0.0015 (8)
C4	0.0404 (10)	0.0319 (9)	0.0325 (9)	-0.0002 (7)	-0.0024 (8)	-0.0019 (8)
C5	0.0428 (10)	0.0329 (10)	0.0305 (9)	0.0024 (8)	-0.0017 (8)	-0.0015 (8)
C6	0.0565 (13)	0.0354 (10)	0.0324 (10)	0.0013 (9)	-0.0081 (10)	-0.0082 (8)
C7	0.0582 (13)	0.0408 (11)	0.0367 (11)	0.0131 (10)	0.0031 (10)	-0.0020 (9)
C8	0.0700 (18)	0.0639 (17)	0.0503 (15)	0.0182 (13)	0.0193 (14)	-0.0015 (13)
C9	0.0570 (16)	0.083 (2)	0.0642 (18)	0.0135 (15)	0.0240 (14)	0.0145 (17)
C10	0.0539 (15)	0.0616 (16)	0.0682 (19)	-0.0052 (12)	0.0088 (14)	0.0140 (14)
C11	0.0552 (14)	0.0411 (12)	0.0504 (14)	-0.0037 (10)	0.0070 (11)	-0.0017 (10)
C12	0.0449 (11)	0.0353 (10)	0.0321 (10)	0.0060 (8)	0.0030 (9)	0.0003 (8)
C13	0.0407 (9)	0.0341 (9)	0.0271 (9)	-0.0040 (8)	0.0010 (8)	-0.0036 (7)
C14	0.0443 (10)	0.0351 (10)	0.0340 (10)	0.0029 (8)	0.0020 (9)	-0.0012 (8)
C15	0.0517 (13)	0.0486 (13)	0.0476 (13)	0.0143 (10)	0.0002 (11)	0.0010 (11)
C16	0.0687 (16)	0.0374 (11)	0.0632 (16)	0.0167 (11)	0.0081 (13)	0.0036 (11)
C17	0.0764 (18)	0.0286 (11)	0.0692 (18)	0.0006 (11)	0.0009 (14)	-0.0073 (11)
C18	0.0586 (14)	0.0320 (10)	0.0563 (14)	-0.0015 (9)	-0.0054 (12)	-0.0061 (10)
C19	0.0443 (10)	0.0300 (9)	0.0351 (10)	0.0021 (8)	0.0006 (9)	-0.0004 (8)
C20	0.0473 (12)	0.0483 (12)	0.0392 (13)	0.0044 (10)	0.0049 (10)	0.0056 (10)
C21	0.0447 (11)	0.0433 (12)	0.0530 (14)	-0.0024 (10)	-0.0003 (11)	0.0055 (10)
C22	0.0640 (15)	0.0478 (12)	0.0400 (12)	0.0167 (11)	0.0151 (11)	0.0054 (10)
N1	0.0696 (13)	0.0394 (10)	0.0368 (10)	0.0049 (9)	-0.0004 (10)	-0.0135 (8)
N2	0.0418 (9)	0.0414 (9)	0.0406 (10)	0.0006 (7)	-0.0076 (8)	-0.0047 (8)
N3	0.0812 (18)	0.0944 (19)	0.0363 (13)	0.0063 (14)	0.0041 (11)	0.0020 (12)
N4	0.0595 (13)	0.0613 (14)	0.097 (2)	-0.0217 (12)	-0.0013 (14)	0.0057 (15)
N5	0.0509 (11)	0.0526 (12)	0.0458 (11)	0.0136 (9)	0.0108 (10)	0.0073 (9)

N6	0.121 (3)	0.084 (2)	0.0696 (18)	0.0470 (19)	0.0373 (18)	0.0380 (16)
O1	0.0578 (11)	0.0560 (10)	0.0563 (11)	-0.0089 (8)	-0.0123 (9)	-0.0163 (9)
O2	0.0552 (9)	0.0330 (7)	0.0404 (8)	-0.0026 (6)	-0.0050 (7)	-0.0110 (6)
O3	0.0606 (11)	0.0517 (9)	0.0498 (10)	0.0029 (9)	-0.0094 (9)	-0.0077 (8)
O4	0.093 (2)	0.091 (2)	0.097 (2)	0.0319 (15)	-0.0022 (16)	0.0075 (18)

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

C1—C5	1.500 (3)	C11—H11A	0.930
C1—C12	1.511 (3)	C13—O2	1.213 (3)
C1—C6	1.556 (3)	C13—N2	1.337 (3)
C1—C2	1.583 (3)	C14—C15	1.380 (3)
C2—C19	1.508 (3)	C14—C19	1.394 (3)
C2—C13	1.560 (3)	C14—N2	1.406 (3)
C2—C3	1.576 (3)	C15—C16	1.384 (4)
C3—C21	1.482 (3)	C15—H15A	0.930
C3—C20	1.487 (3)	C16—C17	1.369 (4)
C3—C4	1.535 (3)	C16—H16A	0.930
C4—N5	1.328 (3)	C17—C18	1.394 (4)
C4—C5	1.354 (3)	C17—H17A	0.930
C5—C22	1.406 (3)	C18—C19	1.380 (3)
C6—O1	1.220 (3)	C18—H18A	0.930
C6—N1	1.337 (3)	C20—N3	1.126 (4)
C7—C8	1.374 (4)	C21—N4	1.130 (3)
C7—C12	1.395 (3)	C22—N6	1.133 (4)
C7—N1	1.404 (3)	N1—H1	0.860
C8—C9	1.381 (5)	N2—H2	0.860
C8—H8A	0.930	N5—H5A	0.90 (1)
C9—C10	1.382 (5)	N5—H5B	0.90 (1)
C9—H9A	0.930	O3—H3A	0.85 (1)
C10—C11	1.392 (4)	O3—H3B	0.85 (1)
C10—H10A	0.930	O4—H4A	0.85 (1)
C11—C12	1.380 (3)	O4—H4B	0.85 (1)
C5—C1—C12	120.04 (17)	C12—C11—H11A	120.4
C5—C1—C6	110.78 (17)	C10—C11—H11A	120.4
C12—C1—C6	101.37 (17)	C11—C12—C7	119.2 (2)
C5—C1—C2	101.12 (16)	C11—C12—C1	132.7 (2)
C12—C1—C2	113.93 (17)	C7—C12—C1	108.0 (2)
C6—C1—C2	109.64 (16)	O2—C13—N2	127.39 (19)
C19—C2—C13	102.28 (17)	O2—C13—C2	125.34 (19)
C19—C2—C3	118.76 (17)	N2—C13—C2	107.25 (16)
C13—C2—C3	106.69 (16)	C15—C14—C19	122.0 (2)
C19—C2—C1	116.18 (17)	C15—C14—N2	128.3 (2)
C13—C2—C1	107.57 (15)	C19—C14—N2	109.72 (18)
C3—C2—C1	104.56 (15)	C14—C15—C16	117.2 (2)
C21—C3—C20	106.75 (19)	C14—C15—H15A	121.4
C21—C3—C4	110.04 (18)	C16—C15—H15A	121.4

C20—C3—C4	112.24 (19)	C17—C16—C15	121.9 (2)
C21—C3—C2	113.62 (17)	C17—C16—H16A	119.1
C20—C3—C2	111.58 (18)	C15—C16—H16A	119.1
C4—C3—C2	102.72 (16)	C16—C17—C18	120.6 (2)
N5—C4—C5	130.6 (2)	C16—C17—H17A	119.7
N5—C4—C3	119.7 (2)	C18—C17—H17A	119.7
C5—C4—C3	109.72 (18)	C19—C18—C17	118.6 (3)
C4—C5—C22	121.9 (2)	C19—C18—H18A	120.7
C4—C5—C1	114.59 (18)	C17—C18—H18A	120.7
C22—C5—C1	123.10 (19)	C18—C19—C14	119.7 (2)
O1—C6—N1	127.3 (2)	C18—C19—C2	132.2 (2)
O1—C6—C1	124.5 (2)	C14—C19—C2	107.96 (18)
N1—C6—C1	108.2 (2)	N3—C20—C3	179.4 (3)
C8—C7—C12	122.4 (3)	N4—C21—C3	178.0 (3)
C8—C7—N1	127.5 (2)	N6—C22—C5	178.1 (3)
C12—C7—N1	110.0 (2)	C6—N1—C7	111.51 (19)
C7—C8—C9	117.3 (3)	C6—N1—H1	124.2
C7—C8—H8A	121.3	C7—N1—H1	124.2
C9—C8—H8A	121.3	C13—N2—C14	112.51 (18)
C8—C9—C10	121.7 (3)	C13—N2—H2	123.7
C8—C9—H9A	119.1	C14—N2—H2	123.7
C10—C9—H9A	119.1	C4—N5—H5A	119.9 (19)
C9—C10—C11	120.1 (3)	C4—N5—H5B	123 (3)
C9—C10—H10A	119.9	H5A—N5—H5B	115 (3)
C11—C10—H10A	119.9	H3A—O3—H3B	105.1 (17)
C12—C11—C10	119.1 (2)	H4A—O4—H4B	104.8 (17)
C5—C1—C2—C19	158.77 (17)	C9—C10—C11—C12	0.7 (4)
C12—C1—C2—C19	-71.0 (2)	C10—C11—C12—C7	-2.2 (4)
C6—C1—C2—C19	41.8 (2)	C10—C11—C12—C1	179.7 (2)
C5—C1—C2—C13	-87.37 (18)	C8—C7—C12—C11	2.5 (4)
C12—C1—C2—C13	42.8 (2)	N1—C7—C12—C11	-175.0 (2)
C6—C1—C2—C13	155.61 (17)	C8—C7—C12—C1	-179.0 (2)
C5—C1—C2—C3	25.79 (18)	N1—C7—C12—C1	3.5 (3)
C12—C1—C2—C3	155.99 (17)	C5—C1—C12—C11	48.3 (4)
C6—C1—C2—C3	-91.22 (18)	C6—C1—C12—C11	170.6 (2)
C19—C2—C3—C21	-37.6 (3)	C2—C1—C12—C11	-71.8 (3)
C13—C2—C3—C21	-152.32 (18)	C5—C1—C12—C7	-130.0 (2)
C1—C2—C3—C21	93.88 (19)	C6—C1—C12—C7	-7.7 (2)
C19—C2—C3—C20	83.1 (2)	C2—C1—C12—C7	110.0 (2)
C13—C2—C3—C20	-31.6 (2)	C19—C2—C13—O2	-173.2 (2)
C1—C2—C3—C20	-145.38 (18)	C3—C2—C13—O2	-47.8 (3)
C19—C2—C3—C4	-156.44 (18)	C1—C2—C13—O2	63.9 (3)
C13—C2—C3—C4	88.85 (18)	C19—C2—C13—N2	5.2 (2)
C1—C2—C3—C4	-24.95 (19)	C3—C2—C13—N2	130.60 (18)
C21—C3—C4—N5	72.4 (3)	C1—C2—C13—N2	-117.67 (18)
C20—C3—C4—N5	-46.3 (3)	C19—C14—C15—C16	-0.5 (4)
C2—C3—C4—N5	-166.3 (2)	N2—C14—C15—C16	-179.2 (2)

C21—C3—C4—C5	−106.6 (2)	C14—C15—C16—C17	0.1 (4)
C20—C3—C4—C5	134.7 (2)	C15—C16—C17—C18	0.4 (5)
C2—C3—C4—C5	14.7 (2)	C16—C17—C18—C19	−0.4 (5)
N5—C4—C5—C22	−3.5 (4)	C17—C18—C19—C14	0.0 (4)
C3—C4—C5—C22	175.3 (2)	C17—C18—C19—C2	175.9 (3)
N5—C4—C5—C1	−176.4 (2)	C15—C14—C19—C18	0.4 (4)
C3—C4—C5—C1	2.4 (3)	N2—C14—C19—C18	179.4 (2)
C12—C1—C5—C4	−144.5 (2)	C15—C14—C19—C2	−176.4 (2)
C6—C1—C5—C4	97.9 (2)	N2—C14—C19—C2	2.6 (3)
C2—C1—C5—C4	−18.3 (2)	C13—C2—C19—C18	179.1 (3)
C12—C1—C5—C22	42.7 (3)	C3—C2—C19—C18	62.1 (4)
C6—C1—C5—C22	−74.9 (3)	C1—C2—C19—C18	−64.0 (3)
C2—C1—C5—C22	168.9 (2)	C13—C2—C19—C14	−4.6 (2)
C5—C1—C6—O1	−41.8 (3)	C3—C2—C19—C14	−121.6 (2)
C12—C1—C6—O1	−170.3 (2)	C1—C2—C19—C14	112.2 (2)
C2—C1—C6—O1	68.9 (3)	O1—C6—N1—C7	171.7 (2)
C5—C1—C6—N1	138.14 (19)	C1—C6—N1—C7	−8.3 (3)
C12—C1—C6—N1	9.6 (2)	C8—C7—N1—C6	−174.1 (3)
C2—C1—C6—N1	−111.1 (2)	C12—C7—N1—C6	3.2 (3)
C12—C7—C8—C9	−1.1 (4)	O2—C13—N2—C14	174.4 (2)
N1—C7—C8—C9	176.0 (3)	C2—C13—N2—C14	−4.0 (2)
C7—C8—C9—C10	−0.5 (5)	C15—C14—N2—C13	179.9 (2)
C8—C9—C10—C11	0.7 (5)	C19—C14—N2—C13	1.0 (3)

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

$D—H\cdots A$	$D—H$	$H\cdots A$	$D\cdots A$	$D—H\cdots A$
N1—H1 \cdots O2 ⁱ	0.86	2.09	2.905 (2)	158
N2—H2 \cdots O3 ⁱ	0.86	2.22	3.032 (3)	157
O3—H3A \cdots O1 ⁱⁱ	0.85 (1)	2.00 (2)	2.795 (3)	157 (4)
O3—H3B \cdots N6 ⁱⁱⁱ	0.85 (1)	2.22 (5)	2.834 (3)	129 (5)
O3—H3B \cdots O4 ^{iv}	0.85 (1)	2.62 (3)	3.381 (4)	149 (5)
O4—H4A \cdots N4 ^v	0.85 (1)	2.38 (3)	3.171 (5)	155 (7)
O4—H4B \cdots N3	0.85 (1)	2.68 (4)	3.392 (4)	142 (6)
N5—H5A \cdots O3 ^{vi}	0.90 (1)	1.97 (1)	2.860 (3)	175 (3)
N5—H5B \cdots O4 ^{vi}	0.90 (1)	2.17 (1)	3.061 (4)	171 (3)

Symmetry codes: (i) $-x+1/2, y-1/2, z-1/2$; (ii) $x, y+1, z+1$; (iii) $x, y, z+1$; (iv) $-x, -y+1, z+1/2$; (v) $-x, -y, z+1/2$; (vi) $-x, -y+1, z-1/2$.