Acta Crystallographica Section E
Structure Reports Online
ISSN 1600-5368

2,4,6-Trinitro-*N*-[4-(phenyldiazenyl)-phenyl]aniline

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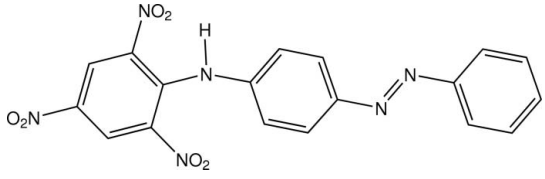
Received 10 March 2011; accepted 17 March 2011

Key indicators: single-crystal X-ray study; T = 200 K; mean σ(C–C) = 0.004 Å; R factor = 0.037; wR factor = 0.058; data-to-parameter ratio = 8.5.

The title compound, $C_{18}H_{12}N_6O_6$, was prepared from the reaction of 4-(phenyldiazenyl)aniline (aniline yellow) with picrylsulfonic acid. The dihedral angle formed by the two benzene rings of the diphenyldiazenyl ring system is 6.55 (13)° and that formed by the rings of the picrate–aniline ring system is 48.76 (12)°. The molecule contains an intramolecular aniline–nitro N—H···O hydrogen bond.

Related literature

For the reaction of picryl chloride with isomeric aminobenzoic acids, see: Crocker & Matthews (1911). For the application of the title compound in dyeing technology, see: Beretta (1926);. For structural data on *N*-picryl-substituted anilines, see: Forlani *et al.* (1992); Pan *et al.* (2007); Smith *et al.* (2007); Braun *et al.* (2008); Smith *et al.* (2009). For diazenyl-protonated salts of aniline yellow, see: Mahmoudkhani & Langer (2001); Smith *et al.* (2009, 2011).



Experimental

Crystal data

$C_{18}H_{12}N_6O_6$
M_r = 408.34
Monoclinic, $P2_1$
a = 7.4255 (4) Å
b = 7.6613 (4) Å
c = 16.1510 (9) Å
β = 98.160 (5)°
V = 909.51 (9) Å^3
Z = 2
Mo $K\alpha$ radiation
μ = 0.12 mm^{-1}
T = 200 K
0.30 × 0.30 × 0.15 mm

Data collection

Oxford Diffraction Gemini-S CCD-detector diffractometer
Absorption correction: multi-scan (*CrysAlis PRO*; Oxford Diffraction, 2010)
T_{min} = 0.920, T_{max} = 0.990
6768 measured reflections
2297 independent reflections
1407 reflections with $I > 2\sigma(I)$
R_{int} = 0.036

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.037
$wR(F^2)$ = 0.058
S = 0.86
2297 reflections
271 parameters
1 restraint
H-atom parameters not refined
$\Delta\rho_{max}$ = 0.15 e Å^{-3}
$\Delta\rho_{min}$ = −0.14 e Å^{-3}

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

D—H···A	D—H	H···A	D···A	D—H···A
N1—H1···O21A	0.86	1.98	2.607 (3)	129

Data collection: *CrysAlis PRO* (Oxford Diffraction, 2010); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis PRO*; program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008) within *WinGX* (Farrugia, 1999); molecular graphics: *PLATON* (Spek, 2009); software used to prepare material for publication: *PLATON*.

The authors acknowledge financial support from the Australian Research Council, the Faculty of Science and Technology and the University Library, Queensland University of Technology.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: LH5220).

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supplementary materials

Acta Cryst. (2011). E**67**, o948 [**doi:10.1107/S160053681101004X**]

2,4,6-Trinitro-*N*-[4-(phenyldiazenyl)phenyl]aniline

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Comment

The diazo-dye precursor aniline yellow [4-(phenyldiazenyl)aniline] reacts with strong acids to form purple-black to red-black diazenyl-protonated salts (Mahmoudkhani & Langer, 2001; Smith *et al.*, 2009, 2011). However, our 1:1 stoichiometric reaction of aniline yellow with 2,4,6-trinitrobenzenesulfonic acid (picrylsulfonic acid) in 50% ethanol-water atypically gave orange-red crystals. This indicated a substitution reaction typical of this acid with anilines, giving *N*-picryl products with elimination of the sulfonate group. Reaction of picryl chloride with the isomeric aniline carboxylates to give similar products was reported by Crocker & Matthews (1911) while the application of picryl substituted azoanilines including the title compound, (I), in dyeing, was discussed by Beretta (1926). A number of structures of picryl-substituted anilines and other aromatic amines have been reported (*e.g.* Forlani *et al.*, 1992; Braun *et al.*, 2008; Pan *et al.*, 2007; Smith *et al.*, 2007).

In the title compound, (I), picryl-substitution of the aniline group of the 4-(phenyldiazenyl)aniline molecule has occurred. The molecular structure of (I) is shown Fig. 1. The diphenyldiazenyl ring system is non-planar [torsion angles C3—C4—N4—N41 and C51—C41—N41—N4: 175.4 (2) and -169.5 (2)°, respectively] as is the picrate to aniline ring system [torsion angles C2A—C1A—N1—C1 and C6—C1—N1—C1A: 152.5 (2) and 156.3 (3)° respectively]. Within the picrate moiety, one of the two *ortho*-related nitro groups is rotated out of the benzene plane [torsion angle C5A—C6A—N6A—O62A, 147.1 (2)°], while the other, which is associated with an intramolecular hydrogen bond [N1—H···O21A (Table 1)] is close to coplanar [C1A—C2A—N2A—O22A, -173.3 (2)°]. The *para*-related nitro group is also essentially coplanar with the ring [C3A—C4A—N4A—O42A, 172.0 (2)°]. There is one short intermolecular non-bonding nitro group interaction [O41A···O42A^{i}, 2.860 (3) Å: symmetry code (i) -*x* + 1, *y* - 1/2, -*z*]. In addition, there are weak π···π ring interactions with a centroid to centroid distance 3.7744 (16) Å.

Experimental

The title compound was synthesized by heating together under reflux for 10 minutes, 1 mmol quantities of 4-(phenyldiazenyl)aniline (aniline yellow) and 2,4,6-trinitrobenzenesulfonic acid (picrylsulfonic acid) in 50 ml of 50% ethanol-water. After concentration to *ca* 30 ml, partial room temperature evaporation of the hot-filtered solution gave orange-red prisms of (I) from which a specimen was cleaved for the X-ray analysis.

Refinement

All H-atoms were included in the refinement in calculated positions and were allowed to ride with C—H = 0.93 Å or N—H = 0.86 Å and with U_{iso}(H) = 1.2U_{eq}(C, N). In the absence of significant anomalous dispersion effects Friedel pairs were merged for the final cycles of refinement.

Figures

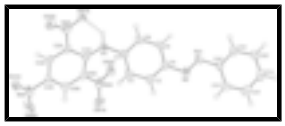


Fig. 1. The molecular structure of the title compound. The intramolecular hydrogen bond is shown as a dashed line and displacement ellipsoids are drawn at the 40% probability level.

2,4,6-Trinitro-*N*-[4-(phenyldiazenyl)phenyl]aniline

Crystal data

$C_{18}H_{12}N_6O_6$	$F(000) = 420$
$M_r = 408.34$	$D_x = 1.491$ Mg m^{-3}
Monoclinic, $P2_1$	Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Hall symbol: P 2yb	Cell parameters from 2105 reflections
$a = 7.4255$ (4) Å	$\theta = 3.5$–$28.5°$
$b = 7.6613$ (4) Å	$\mu = 0.12$ mm^{-1}
$c = 16.1510$ (9) Å	$T = 200$ K
$\beta = 98.160$ (5)°	Plate, orange-red
$V = 909.51$ (9) Å^3	0.30 × 0.30 × 0.15 mm
$Z = 2$	

Data collection

Oxford Diffraction Gemini-S CCD-detector diffractometer	2297 independent reflections
Radiation source: Enhance (Mo) X-ray source	1407 reflections with $I > 2\sigma(I)$
graphite	$R_{int} = 0.036$
Detector resolution: 16.077 pixels mm^{-1}	$\theta_{max} = 28.5°$, $\theta_{min} = 3.5°$
ω scans	$h = -9 \rightarrow 9$
Absorption correction: multi-scan (*CrysAlis PRO*; Oxford Diffraction, 2010)	$k = -10 \rightarrow 10$
$T_{min} = 0.920$, $T_{max} = 0.990$	$l = -21 \rightarrow 21$
6768 measured reflections	

Refinement

Refinement on F^2	Primary atom site location: structure-invariant direct methods
Least-squares matrix: full	Secondary atom site location: difference Fourier map
$R[F^2 > 2\sigma(F^2)] = 0.037$	Hydrogen site location: inferred from neighbouring sites
$wR(F^2) = 0.058$	H-atom parameters not refined
$S = 0.86$	$w = 1/[\sigma^2(F_o^2) + (0.021P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$
2297 reflections	$(\Delta/\sigma)_{max} < 0.001$
271 parameters	$\Delta\rho_{max} = 0.15$ e Å^{-3}

1 restraint $\Delta\rho_{min} = -0.14$ e Å^{-3}

Special details

Geometry. Bond distances, angles *etc*. have been calculated using the rounded fractional coordinates. All su's are estimated from the variances of the (full) variance-covariance matrix. The cell e.s.d.'s are taken into account in the estimation of distances, angles and torsion angles

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 (Å^2)

	x	*y*	*z*	U_{iso}*/U_{eq}
O21A	0.7894 (3)	0.0405 (2)	0.33042 (15)	0.0557 (8)
O22A	0.8395 (3)	−0.0131 (3)	0.20499 (15)	0.0673 (10)
O41A	0.7046 (3)	0.3737 (3)	−0.01934 (12)	0.0524 (8)
O42A	0.5573 (3)	0.6113 (3)	−0.00428 (13)	0.0733 (10)
O61A	0.4815 (3)	0.8150 (3)	0.26498 (14)	0.0582 (9)
O62A	0.4238 (2)	0.6124 (3)	0.35215 (12)	0.0467 (7)
N1	0.6941 (3)	0.3543 (3)	0.36913 (14)	0.0379 (8)
N2A	0.7860 (3)	0.0827 (3)	0.25675 (17)	0.0438 (10)
N4	0.8515 (3)	0.7986 (3)	0.64591 (15)	0.0376 (8)
N4A	0.6334 (3)	0.4775 (3)	0.02299 (15)	0.0403 (9)
N6A	0.4931 (3)	0.6655 (3)	0.29240 (16)	0.0397 (9)
N41	0.7964 (3)	0.7460 (3)	0.71112 (15)	0.0388 (8)
C1	0.7256 (3)	0.4761 (4)	0.43607 (17)	0.0343 (10)
C1A	0.6660 (3)	0.3836 (3)	0.28559 (17)	0.0304 (9)
C2	0.8086 (3)	0.6339 (3)	0.42983 (17)	0.0367 (10)
C2A	0.7143 (3)	0.2547 (3)	0.22900 (18)	0.0323 (9)
C3	0.8466 (3)	0.7381 (4)	0.50001 (17)	0.0363 (10)
C3A	0.7034 (3)	0.2855 (3)	0.14441 (17)	0.0325 (10)
C4	0.8030 (3)	0.6843 (4)	0.57641 (18)	0.0357 (10)
C4A	0.6397 (3)	0.4442 (4)	0.11242 (17)	0.0304 (9)
C5	0.7209 (4)	0.5221 (4)	0.58222 (18)	0.0420 (11)
C5A	0.5793 (3)	0.5688 (3)	0.16261 (17)	0.0321 (10)
C6	0.6810 (4)	0.4184 (4)	0.51302 (18)	0.0429 (11)
C6A	0.5887 (3)	0.5373 (3)	0.24652 (17)	0.0301 (9)
C11	0.9043 (4)	1.0521 (4)	0.92620 (19)	0.0475 (11)
C21	0.9363 (3)	1.1240 (4)	0.8506 (2)	0.0455 (11)
C31	0.9046 (3)	1.0274 (4)	0.77855 (17)	0.0361 (10)
C41	0.8401 (3)	0.8580 (3)	0.78142 (17)	0.0313 (10)
C51	0.8078 (3)	0.7876 (4)	0.85652 (18)	0.0391 (10)
C61	0.8423 (4)	0.8851 (4)	0.92962 (19)	0.0437 (11)
H1	0.69280	0.24650	0.38380	0.0450*
H2	0.83930	0.67080	0.37880	0.0440*

H3	0.90220	0.84590	0.49580	0.0430*
H3A	0.73870	0.19990	0.10920	0.0390*
H5	0.69290	0.48390	0.63360	0.0500*
H5A	0.53250	0.67380	0.14000	0.0390*
H6	0.62490	0.31080	0.51700	0.0510*
H11	0.92550	1.11840	0.97480	0.0570*
H21	0.97920	1.23780	0.84890	0.0540*
H31	0.92630	1.07520	0.72800	0.0430*
H51	0.76280	0.67450	0.85820	0.0470*
H61	0.82320	0.83690	0.98050	0.0520*

Atomic displacement parameters (Å^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O21A	0.0839 (16)	0.0368 (12)	0.0427 (14)	−0.0079 (11)	−0.0033 (12)	0.0064 (11)
O22A	0.0965 (18)	0.0423 (14)	0.0679 (18)	0.0221 (13)	0.0283 (14)	−0.0003 (12)
O41A	0.0661 (14)	0.0584 (14)	0.0361 (14)	−0.0002 (12)	0.0191 (11)	−0.0091 (11)
O42A	0.0923 (18)	0.0854 (17)	0.0449 (15)	0.0435 (16)	0.0196 (13)	0.0222 (14)
O61A	0.0700 (16)	0.0370 (14)	0.0701 (18)	0.0057 (11)	0.0187 (12)	−0.0105 (11)
O62A	0.0385 (11)	0.0683 (14)	0.0349 (13)	−0.0036 (11)	0.0107 (10)	−0.0101 (11)
N1	0.0475 (14)	0.0359 (13)	0.0305 (15)	−0.0133 (12)	0.0065 (12)	0.0030 (12)
N2A	0.0492 (16)	0.0339 (16)	0.0470 (19)	−0.0023 (12)	0.0022 (14)	0.0000 (14)
N4	0.0358 (13)	0.0478 (15)	0.0286 (16)	−0.0031 (12)	0.0022 (11)	−0.0014 (12)
N4A	0.0354 (14)	0.0543 (17)	0.0318 (16)	0.0028 (13)	0.0067 (12)	0.0020 (14)
N6A	0.0276 (13)	0.0496 (18)	0.0414 (17)	−0.0032 (12)	0.0031 (12)	−0.0160 (14)
N41	0.0429 (14)	0.0470 (15)	0.0268 (15)	−0.0013 (13)	0.0058 (11)	−0.0008 (13)
C1	0.0327 (16)	0.0389 (18)	0.0303 (18)	−0.0076 (14)	0.0014 (13)	−0.0048 (15)
C1A	0.0253 (15)	0.0353 (16)	0.0302 (18)	−0.0072 (13)	0.0028 (13)	−0.0062 (14)
C2	0.0347 (15)	0.0505 (19)	0.0246 (17)	−0.0113 (14)	0.0031 (12)	0.0061 (15)
C2A	0.0312 (15)	0.0283 (15)	0.0365 (18)	−0.0030 (13)	0.0019 (13)	−0.0032 (14)
C3	0.0359 (15)	0.0401 (17)	0.0318 (19)	−0.0123 (14)	0.0012 (13)	−0.0004 (15)
C3A	0.0278 (15)	0.0359 (17)	0.0343 (18)	−0.0073 (13)	0.0057 (13)	−0.0111 (14)
C4	0.0337 (16)	0.0433 (18)	0.0280 (18)	−0.0039 (14)	−0.0027 (13)	0.0004 (15)
C4A	0.0273 (14)	0.0397 (17)	0.0244 (16)	−0.0027 (13)	0.0039 (12)	−0.0024 (14)
C5	0.0560 (19)	0.0454 (19)	0.0251 (18)	−0.0117 (16)	0.0073 (14)	0.0015 (14)
C5A	0.0227 (14)	0.0357 (17)	0.0370 (19)	−0.0007 (13)	0.0010 (13)	0.0007 (14)
C6	0.0527 (18)	0.0401 (18)	0.0359 (19)	−0.0138 (14)	0.0065 (15)	0.0029 (14)
C6A	0.0229 (13)	0.0359 (16)	0.0323 (18)	−0.0024 (13)	0.0063 (12)	−0.0076 (14)
C11	0.0380 (17)	0.063 (2)	0.040 (2)	0.0134 (17)	0.0005 (15)	−0.0199 (17)
C21	0.0331 (17)	0.0445 (18)	0.057 (2)	−0.0013 (15)	0.0001 (15)	−0.0113 (17)
C31	0.0280 (15)	0.0444 (18)	0.035 (2)	0.0026 (14)	0.0014 (14)	0.0023 (15)
C41	0.0304 (15)	0.0346 (17)	0.0277 (18)	0.0030 (14)	−0.0004 (13)	−0.0009 (14)
C51	0.0387 (16)	0.0432 (19)	0.0345 (19)	0.0078 (14)	0.0018 (14)	−0.0046 (15)
C61	0.0434 (17)	0.053 (2)	0.0338 (19)	0.0094 (16)	0.0025 (14)	−0.0003 (17)

Geometric parameters (Å, °)

O21A—N2A	1.230 (4)	C3A—C4A	1.378 (4)
O22A—N2A	1.221 (3)	C4—C5	1.393 (4)

O41A—N4A	1.217 (3)	C4A—C5A	1.369 (4)
O42A—N4A	1.222 (3)	C5—C6	1.369 (4)
O61A—N6A	1.227 (3)	C5A—C6A	1.369 (4)
O62A—N6A	1.226 (3)	C11—C61	1.364 (4)
N1—C1	1.422 (4)	C11—C21	1.390 (4)
N1—C1A	1.354 (4)	C21—C31	1.371 (4)
N2A—C2A	1.467 (3)	C31—C41	1.386 (4)
N4—N41	1.250 (3)	C41—C51	1.379 (4)
N4—C4	1.429 (4)	C51—C61	1.390 (4)
N4A—C4A	1.461 (4)	C2—H2	0.9300
N6A—C6A	1.472 (3)	C3—H3	0.9300
N41—C41	1.423 (3)	C3A—H3A	0.9300
N1—H1	0.8600	C5—H5	0.9300
C1—C6	1.402 (4)	C5A—H5A	0.9300
C1—C2	1.367 (4)	C6—H6	0.9300
C1A—C2A	1.426 (4)	C11—H11	0.9300
C1A—C6A	1.418 (3)	C21—H21	0.9300
C2—C3	1.382 (4)	C31—H31	0.9300
C2A—C3A	1.378 (4)	C51—H51	0.9300
C3—C4	1.382 (4)	C61—H61	0.9300
C1—N1—C1A	129.4 (2)	C1—C6—C5	119.3 (3)
O21A—N2A—O22A	122.7 (2)	N6A—C6A—C1A	121.6 (2)
O21A—N2A—C2A	119.2 (2)	N6A—C6A—C5A	114.9 (2)
O22A—N2A—C2A	118.0 (3)	C1A—C6A—C5A	123.2 (2)
N41—N4—C4	112.9 (2)	C21—C11—C61	120.5 (3)
O41A—N4A—O42A	124.1 (2)	C11—C21—C31	120.1 (3)
O41A—N4A—C4A	119.1 (2)	C21—C31—C41	119.7 (3)
O42A—N4A—C4A	116.8 (2)	N41—C41—C51	114.7 (2)
O61A—N6A—O62A	125.3 (2)	N41—C41—C31	125.3 (2)
O61A—N6A—C6A	117.1 (2)	C31—C41—C51	120.0 (3)
O62A—N6A—C6A	117.5 (2)	C41—C51—C61	120.2 (3)
N4—N41—C41	114.4 (2)	C11—C61—C51	119.5 (3)
C1—N1—H1	115.00	C1—C2—H2	120.00
C1A—N1—H1	115.00	C3—C2—H2	120.00
N1—C1—C2	123.5 (2)	C2—C3—H3	120.00
C2—C1—C6	120.6 (3)	C4—C3—H3	120.00
N1—C1—C6	115.7 (3)	C2A—C3A—H3A	120.00
C2A—C1A—C6A	114.4 (2)	C4A—C3A—H3A	120.00
N1—C1A—C6A	125.2 (2)	C4—C5—H5	120.00
N1—C1A—C2A	120.4 (2)	C6—C5—H5	120.00
C1—C2—C3	119.5 (2)	C4A—C5A—H5A	120.00
C1A—C2A—C3A	122.2 (2)	C6A—C5A—H5A	120.00
N2A—C2A—C1A	122.7 (2)	C1—C6—H6	120.00
N2A—C2A—C3A	115.1 (2)	C5—C6—H6	120.00
C2—C3—C4	120.9 (3)	C21—C11—H11	120.00
C2A—C3A—C4A	119.5 (2)	C61—C11—H11	120.00
N4—C4—C5	123.9 (3)	C11—C21—H21	120.00
N4—C4—C3	117.0 (3)	C31—C21—H21	120.00
C3—C4—C5	119.1 (3)	C21—C31—H31	120.00

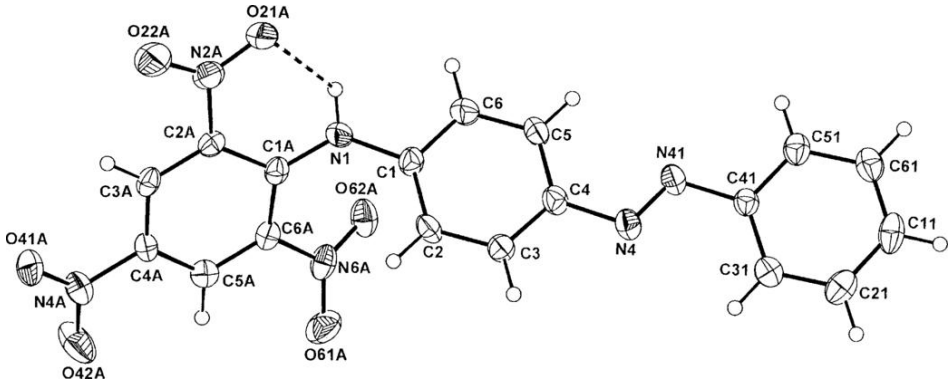
N4A—C4A—C5A	119.8 (2)	C41—C31—H31	120.00
N4A—C4A—C3A	119.1 (2)	C41—C51—H51	120.00
C3A—C4A—C5A	121.1 (3)	C61—C51—H51	120.00
C4—C5—C6	120.6 (3)	C11—C61—H61	120.00
C4A—C5A—C6A	119.3 (2)	C51—C61—H61	120.00
C1A—N1—C1—C2	−29.4 (4)	C6A—C1A—C2A—C3A	6.3 (3)
C1A—N1—C1—C6	156.3 (3)	N1—C1A—C6A—N6A	−14.0 (4)
C1—N1—C1A—C2A	152.5 (2)	N1—C1A—C6A—C5A	173.3 (2)
C1—N1—C1A—C6A	−27.7 (4)	C2A—C1A—C6A—N6A	165.7 (2)
O21A—N2A—C2A—C1A	7.0 (4)	C2A—C1A—C6A—C5A	−7.0 (3)
O21A—N2A—C2A—C3A	−175.4 (2)	C1—C2—C3—C4	0.5 (4)
O22A—N2A—C2A—C1A	−173.3 (2)	N2A—C2A—C3A—C4A	−179.3 (2)
O22A—N2A—C2A—C3A	4.3 (3)	C1A—C2A—C3A—C4A	−1.6 (4)
C4—N4—N41—C41	−179.2 (2)	C2—C3—C4—N4	178.5 (2)
N41—N4—C4—C3	175.4 (2)	C2—C3—C4—C5	0.5 (4)
N41—N4—C4—C5	−6.7 (4)	C2A—C3A—C4A—N4A	178.3 (2)
O41A—N4A—C4A—C3A	−8.9 (4)	C2A—C3A—C4A—C5A	−3.0 (4)
O41A—N4A—C4A—C5A	172.4 (2)	N4—C4—C5—C6	−178.9 (3)
O42A—N4A—C4A—C3A	172.0 (2)	C3—C4—C5—C6	−1.1 (4)
O42A—N4A—C4A—C5A	−6.7 (3)	N4A—C4A—C5A—C6A	−178.9 (2)
O61A—N6A—C6A—C1A	157.5 (2)	C3A—C4A—C5A—C6A	2.4 (4)
O61A—N6A—C6A—C5A	−29.3 (3)	C4—C5—C6—C1	0.8 (4)
O62A—N6A—C6A—C1A	−26.1 (3)	C4A—C5A—C6A—N6A	−170.2 (2)
O62A—N6A—C6A—C5A	147.1 (2)	C4A—C5A—C6A—C1A	2.9 (4)
N4—N41—C41—C31	12.6 (4)	C61—C11—C21—C31	0.4 (4)
N4—N41—C41—C51	−169.5 (2)	C21—C11—C61—C51	−1.2 (4)
N1—C1—C2—C3	−174.9 (2)	C11—C21—C31—C41	0.2 (4)
C6—C1—C2—C3	−0.8 (4)	C21—C31—C41—N41	178.0 (2)
N1—C1—C6—C5	174.7 (3)	C21—C31—C41—C51	0.1 (4)
C2—C1—C6—C5	0.2 (4)	N41—C41—C51—C61	−179.0 (2)
N1—C1A—C2A—N2A	3.5 (3)	C31—C41—C51—C61	−1.0 (4)
N1—C1A—C2A—C3A	−174.0 (2)	C41—C51—C61—C11	1.5 (4)
C6A—C1A—C2A—N2A	−176.3 (2)		

Hydrogen-bond geometry (Å, °)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*
N1—H1···O21A	0.86	1.98	2.607 (3)	129
C3A—H3A···O22A	0.93	2.30	2.633 (3)	100
C5—H5···O61A^{i}	0.93	2.58	3.453 (4)	157
C11—H11···O41A^{ii}	0.93	2.56	3.068 (4)	115
C21—H21···O22A^{iii}	0.93	2.56	3.425 (4)	155

Symmetry codes: (i) −*x*+1, *y*−1/2, −*z*+1; (ii) *x*, *y*+1, *z*+1; (iii) −*x*+2, *y*+3/2, −*z*+1.

Fig. 1


O21A
O22A
N2A
C2A
C1A
N1
C1
C6
C5
C4
N41
C41
C51
C61
C11
C21
C31
N4
C2
C3
C3A
O62A
C6A
N6A
O61A
O41A
C4A
C5A
N4A
O42A