



Crystal structure and Hirshfeld surface analysis of a new polymorph of (*E*)-2-(4-bromophenyl)-1-[2,2-dibromo-1-(3-nitrophenyl)ethenyl]diazene

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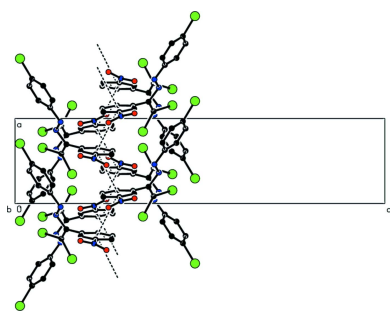
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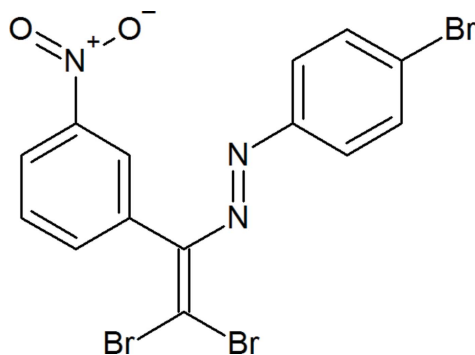
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A new polymorph of the title compound, C₁₄H₈Br₃N₃O₂, (form-2) was obtained in the same manner as the previously reported form-1 [Akkurt *et al.* (2022). *Acta Cryst. E* **78**, 732–736]. The structure of the new polymorph is stabilized by a C—H···O hydrogen bond that links molecules into chains. These chains are linked by face-to-face π – π stacking interactions, resulting in a layered structure. Short inter-molecular Br···O contacts and van der Waals interactions between the layers aid in the cohesion of the crystal packing. In the previously reported form-1, C—H···Br interactions connect molecules into zigzag chains, which are linked by C—Br··· π interactions into layers, whereas the van der Waals interactions between the layers stabilize the crystal packing of form-2. Hirshfeld molecular surface analysis was used to compare the intermolecular interactions of the polymorphs.

1. Chemical context

Aromatic azo compounds provide ubiquitous motifs in organic chemistry and are widely used as indicators, organic dyes, pigments, radical reaction initiators, food additives, therapeutic agents, etc. (Zollinger 1994, 1995; Gurbanov *et al.*, 2020*a,b*). Moreover, in azo dyes the ligands play a crucial role in coordination chemistry and in the construction of functional materials, such as ionophores, self-assembled layers, catalysts, antimicrobial agents, liquid crystals and semiconductors (Ma *et al.*, 2020, 2021; Mahmudov *et al.*, 2010, 2013). Depending on the attached functional groups, the chemical and physical properties of azo dyes and their transition-metal complexes can be improved. The *azo-to-hydrazo* tautomerization as well as *E/Z* isomerization of azo dyes are key phenomena in the synthesis and design of new functional materials (Shikhaliyev *et al.*, 2013, 2014). Moreover, an attachment of donor or acceptor centres of non-covalent bonds to the azo compounds can be applied as a synthetic strategy in the improvement of functional properties of their metal complexes (Mahmudov *et al.*, 2020, 2021, 2022). Thus, we have attached bromine and nitro substituents to the aryl rings leading to a new azo dye, (*E*)-1-(2,2-dibromo-1-(3-nitrophenyl)vinyl)-2-(4-bromophenyl)diazene, which can participate in intermolecular halogen and hydrogen bonds as well as in π -interactions.





2. Structural commentary

A view of the molecule of the new polymorph (henceforth referred to as form-2) is shown in Fig. 1. The central fragment of the molecule, C1/C2/N2/N3/C3/C9/Br1/Br2, is almost planar with the largest deviation from mean plane being 0.101 (1) Å for Br1. This plane forms dihedral angles of 13.51 (7) and 61.26 (7)° with the planes of the bromine- and nitro-substituted aromatic rings, respectively. In the previously reported polymorph (form-1), the corresponding angles were 26.35 (15) and 72.57 (14)° (Akkurt *et al.*, 2022). All bond lengths and angles in the title compound are in agreement with those reported for the related azo compounds discussed in the *Database survey* section.

3. Supramolecular features and Hirshfeld surface analysis

The crystal packing of the new polymorph is stabilized by a C—H···O hydrogen bond that links molecules into chains along the *b*-axis direction (Table 1, Figs. 2–4). These chains are joined by zigzag face-to-face π – π stacking interactions along the [100] direction [$Cg1 \cdots Cg1(-\frac{1}{2} + x, y, \frac{1}{2} - z) = 3.7305(11)$ Å, slippage: 2.057 Å; $Cg1 \cdots Cg1(\frac{1}{2} + x, y, \frac{1}{2} - z) = 3.7305(11)$ Å, slippage: 0.9775 Å; where *Cg1* is the centroid of

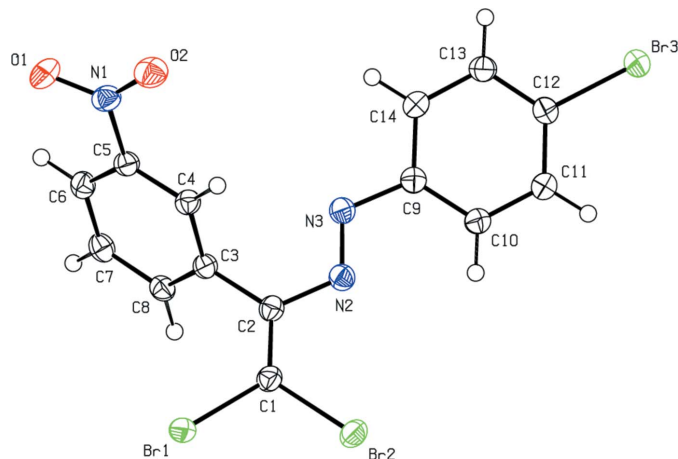


Figure 1
The molecular structure of the title compound. Displacement ellipsoids are drawn at the 50% probability level.

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

<i>D</i> —H··· <i>A</i>	<i>D</i> —H	H··· <i>A</i>	<i>D</i> ··· <i>A</i>	<i>D</i> —H··· <i>A</i>
C8—H8···O1 ⁱ	0.95	2.56	3.394 (2)	146

Symmetry code: (i) $-x, y - \frac{1}{2}, -z + \frac{1}{2}$.

the nitrophenyl ring], resulting in the layers parallel to (001) (Fig. 4). Short inter-molecular Br1···O2 contacts (Table 2) and van der Waals interactions between the layers help to keep the crystal packing together. In the previously reported

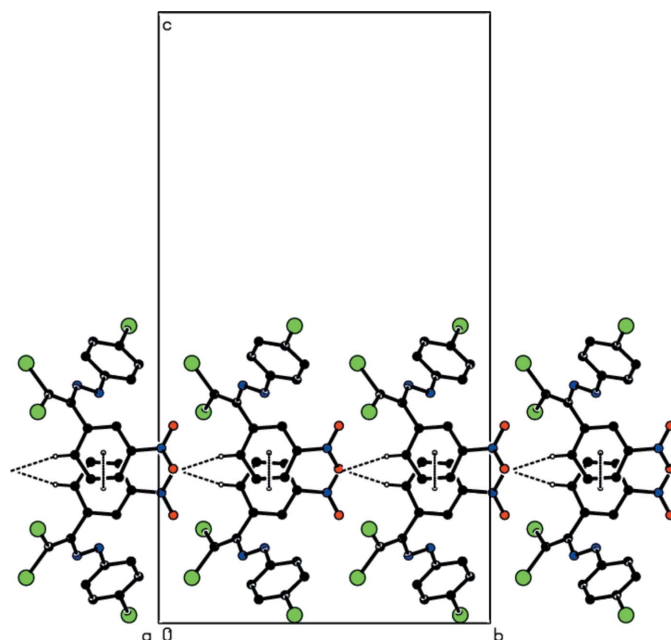


Figure 2
View down the *a*-axis of the C—H···O and π – π interactions (dashed lines) in the title compound.

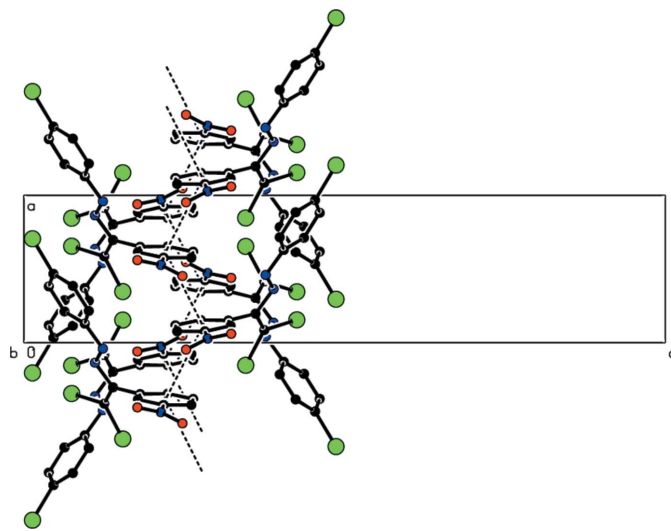


Figure 3
View down the *b*-axis of the C—H···O and π – π interactions (dashed lines) in the title compound.

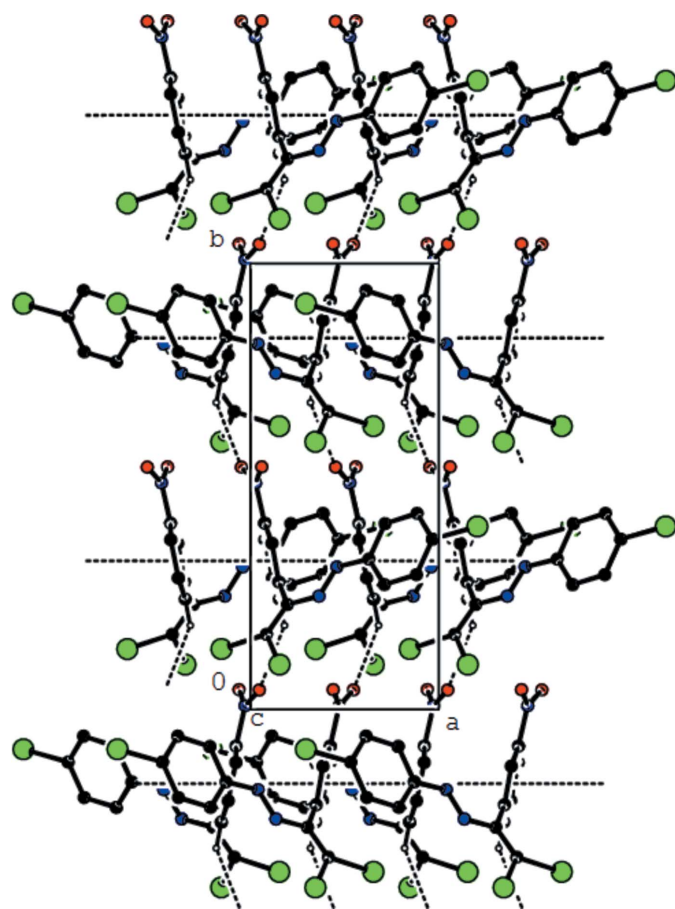


Figure 4
View down the *c*-axis of the C–H...O and π – π interactions (dashed lines) in the title compound.

form-1 of the title compound (Akkurt *et al.*, 2022), C–H...Br interactions connect molecules, generating zigzag *C*(8) chains along the [100] direction, which are linked by C–Br... π interactions into layers parallel to (001), and van der Waals interactions between layers contribute to the crystal cohesion.

Crystal Explorer 17.5 (Turner *et al.*, 2017) was used to perform a Hirshfeld surface analysis of form-2 and to generate the related two-dimensional fingerprint plots, with a standard resolution of the three-dimensional d_{norm} surfaces plotted over a fixed colour scale of -0.1471 (red) to $+1.1715$ (blue) a.u.

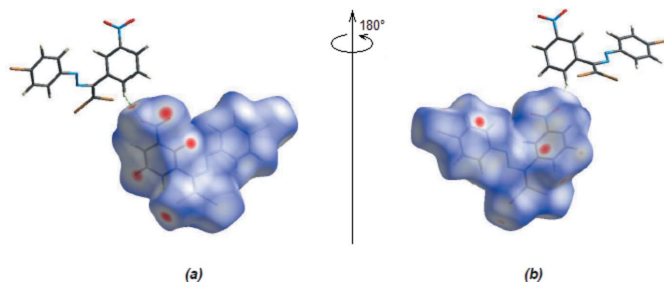
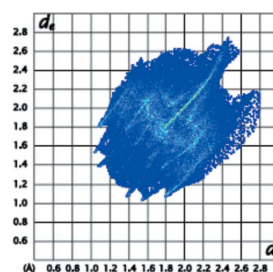


Figure 5
(*a*) Front and (*b*) back views of the three-dimensional Hirshfeld surface of the title compound plotted over d_{norm} in the range -0.1471 to 1.1715 a.u.

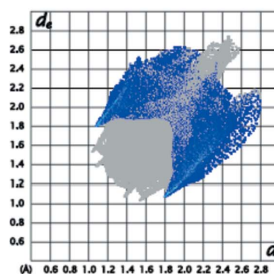
Table 2
Summary of short interatomic contacts (\AA) in the title compound.

H4...C13	2.67	$-1 + x, y, z$
H8...O1	2.56	$-x, -\frac{1}{2} + y, \frac{1}{2} - z$
H7...N3	2.78	$-\frac{1}{2} + x, y, \frac{1}{2} - z$
Br1...O2	3.137 (2)	$-\frac{1}{2} - x, -\frac{1}{2} + y, z$
Br1...H14	2.98	$-\frac{1}{2} - x, -\frac{1}{2} + y, z$
Br2...H13	3.15	$-x, -\frac{1}{2} + y, z$
Br3...H10	3.02	$+x, \frac{1}{2} - y, 1 - z$
C13...Br3	3.569 (2)	$2 - x, 1 - y, 1 - z$

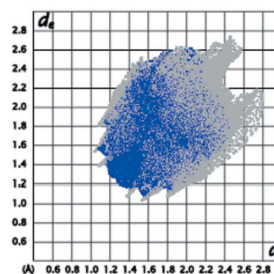
(Fig. 5). The red areas on the surface present short contacts and negative d_{norm} values, which correspond to the C–H...O hydrogen bonds mentioned above (Table 1). The red patch that appears around O1 is due to the C8–H8...O1 interaction, which is critical for the molecular packing of the title compound. In form-1, the C–H...Br interactions are also prominent (Akkurt *et al.*, 2022).



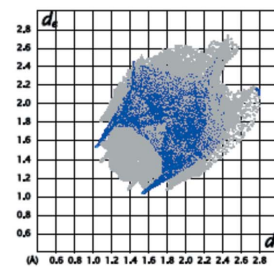
(*a*) All...All



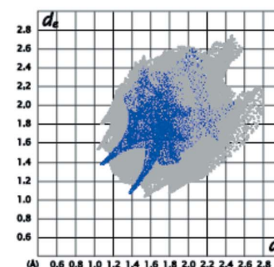
(*b*) Br...H / H...Br



(*c*) H...H



(*d*) C...H / H...C



(*e*) O...H / H...O

Figure 6
The full two-dimensional fingerprint plots for the title compound, showing (*a*) all interactions, and delineated into (*b*) Br...H/H...Br, (*c*) H...H, (*d*) C...H/H...C, and (*e*) O...H/H...O interactions. The d_i and d_e values are the closest internal and external distances (in \AA) from given points on the Hirshfeld surface.

The overall two-dimensional fingerprint plot for form-2 is given in Fig. 6a, and those delineated into Br···H/H···Br (26.5%), H···H (12.8%), C···H/H···C (11.5%) and O···H/H···O (10.6%) contacts are shown in Fig. 6b–e, while the numerical details for the shortest contacts are given in Table 2. Other contacts, such as Br···C/C···Br (7.7%), C···C (6.0%), Br···Br (5.8%), Br···O/O···Br (5.3%), N···H/H···N (5.3%), O···C/C···O (2.5%), Br···N/N···Br (2.3%), O···N/N···O (1.7%), O···O (1.3%) and N···C/C···N (0.8%), have little influence on the molecular packing. For form-1, the set includes only four types of interactions, viz. Br···H/H···Br, H···H, C···H/H···C and O···H/H···O contacts (Akkurt *et al.*, 2022). The predominant interactions in both cases are Br···H/H···Br and H···H, constituting 26.5% and 12.8%, respectively, in form-2 vs 20.9% and 15.2% in form-1.

4. Database survey

A search of the Cambridge Structural Database (CSD, Version 5.42, update of September 2021; Groom *et al.*, 2016) for similar structures with the (*E*)-1-(2,2-dibromo)-2-(4-bromophenyl)diazene unit showed that the ten closest are those of CSD refcodes HEHKEO (**I**) (Akkurt *et al.*, 2022), TAZDIL (**II**) (Atioğlu *et al.*, 2022), PAXDOL (**III**) (Çelikesir *et al.*, 2022), GUPHIL (**IV**) (Özkaraca *et al.*, 2020b), HONBUK (**V**) (Akkurt *et al.*, 2019), HONBOE (**VI**) (Akkurt *et al.*, 2019), HODQAV (**VII**) (Shikhaliyev *et al.*, 2019), XIZREG (**VIII**) (Atioğlu *et al.*, 2019), LEQXOX (**IX**) (Shikhaliyev *et al.*, 2018) and LEQXIR (**X**) (Shikhaliyev *et al.*, 2018).

C—H···Br interactions connect the molecules in the crystal of the form-1 polymorph of the title compound, (**I**), resulting in zigzag C(8) chains along the [100] direction. These chains are connected by C—Br··· π interactions into layers parallel to (001). van der Waals interactions between the layers contribute to the crystal cohesion.

The molecules in (**II**) are joined into layers parallel to (011) by C—H···O and C—H···F hydrogen bonds. C—Br··· π and C—F··· π contacts, as well as π – π stacking interactions, strengthen the crystal packing.

The molecules in the crystal of (**III**) are connected into chains running parallel to [001] by C—H···O hydrogen bonds. C—F··· π contacts and π – π stacking interactions help to consolidate the crystal packing, and short Br···O [2.9828 (13) Å] distances are also observed.

In the crystal of (**IV**), the molecules are linked into inversion dimers *via* short halogen–halogen contacts [Cl1···Cl1 = 3.3763 (9) Å, Cl16—Cl1···Cl1 = 141.47 (7)° compared to the van der Waals radii sum of 3.50 Å for two chlorine atoms]. No other directional contacts could be identified, and the shortest aromatic ring-centroid separation is greater than 5.25 Å.

In the crystals of (**V**) and (**VI**), the molecules are linked through weak X···Cl contacts [*X* = Cl for (**V**) and Br for (**VI**)], C—H···Cl and C—Cl··· π interactions into sheets lying parallel to (001).

In the crystal of (**VII**), the molecules are stacked in columns along [100] *via* weak C—H···Cl hydrogen bonds and face-to-

Table 3
Experimental details.

Crystal data	
Chemical formula	C ₁₄ H ₈ Br ₃ N ₃ O ₂
<i>M_r</i>	489.96
Crystal system, space group	Orthorhombic, <i>Pbca</i>
Temperature (K)	100
<i>a</i> , <i>b</i> , <i>c</i> (Å)	6.6579 (1), 15.7683 (3), 29.0301 (6)
<i>V</i> (Å ³)	3047.69 (10)
<i>Z</i>	8
Radiation type	Mo <i>K</i> α
μ (mm ^{−1})	7.95
Crystal size (mm)	0.34 × 0.06 × 0.05
Data collection	
Diffractometer	Bruker AXS D8 QUEST, Photon III detector
Absorption correction	Multi-scan (<i>SADABS</i> ; Krause <i>et al.</i> , 2015).
<i>T_{min}</i> , <i>T_{max}</i>	0.020, 0.058
No. of measured, independent and observed [<i>I</i> > 2σ(<i>I</i>)] reflections	67179, 5538, 4620
<i>R_{int}</i>	0.033
(sin θ/λ) _{max} (Å ^{−1})	0.758
Refinement	
<i>R</i> [<i>F</i> ² > 2σ(<i>F</i> ²)], <i>wR</i> (<i>F</i> ²), <i>S</i>	0.024, 0.064, 1.05
No. of reflections	5538
No. of parameters	199
H-atom treatment	H-atom parameters constrained
$\Delta\rho_{\max}$, $\Delta\rho_{\min}$ (e Å ^{−3})	0.68, −0.52

Computer programs: *APEX3* and *SAINT* (Bruker, 2018), *SHELXT* (Sheldrick, 2015a), *SHELXL2018* (Sheldrick, 2015b), *ORTEP-3 for Windows* (Farrugia, 2012) and *PLATON* (Spek, 2020).

face π – π stacking interactions. The crystal packing is further consolidated by short Cl···Cl contacts.

In (**VIII**), molecules are linked by C—H···O hydrogen bonds into zigzag chains running parallel to [001]. The crystal packing also features C—Cl··· π , C—F··· π and N—O··· π interactions.

In (**IX**), C—H···N and short Cl···Cl contacts are observed, and in (**X**), C—H···N and C—H···O hydrogen bonds and short Cl···O contacts occur.

5. Synthesis and crystallization

This dye was synthesized according to the reported method (Akkurt *et al.*, 2019; Maharramov *et al.*, 2018; Özkaraca *et al.*, 2020a,b). A 20 mL screw-neck vial was charged with DMSO (10 mL), (*E*)-1-(4-bromophenyl)-2-(3-nitrobenzylidene)-hydrazine (1 mmol), tetramethylethylenediamine (TMEDA; 295 mg, 2.5 mmol), CuCl (2 mg, 0.02 mmol) and CBr₄ (4.5 mmol). After 1–3 h (until TLC analysis showed complete consumption of corresponding Schiff base), the reaction mixture was poured into a 0.01 *M* solution of HCl (100 mL, pH = 2–3), and extracted with dichloromethane (3 × 20 mL). The combined organic phase was washed with water (3 × 50 mL), brine (30 mL), dried over anhydrous Na₂SO₄ and concentrated *in vacuo* using a rotary evaporator. The residue was purified by column chromatography on silica gel using appropriate mixtures of hexane and dichloromethane (3/1–1/1). Crystals suitable for X-ray analysis were obtained by slow

evaporation of an ethanol solution. Red solid (62%); m.p. 391 K. Analysis calculated for $C_{14}H_8Br_3N_3O_2$ ($M = 489.95$): C 34.32, H 1.65, N 8.58; found: C 34.29, H 1.66, N 8.55%. 1H NMR (300 MHz, $CDCl_3$) δ 7.90–7.44 (8H, Ar–H). ^{13}C NMR (75 MHz, $CDCl_3$) δ 150.88, 148.57, 148.12, 132.81, 132.47, 132.25, 130.04, 126.40, 125.30, 124.53, 123.57, 94.10. ESI-MS: m/z : 490.91 [$M + H$] $^+$.

6. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 3. All H atoms were positioned geometrically and allowed to ride on their parent atoms (C–H = 0.95 Å) with $U_{iso}(H) = 1.2U_{eq}(C)$.

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The author's contributions are as follows. Conceptualization, NQS, MA and AB; synthesis, NAM and GVB; X-ray analysis, ZA, VNK and MA; writing (review and editing of the manuscript) ZA, MA and AB; funding acquisition, NQS, NAM and GVB; supervision, NQS, MA and AB.

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supporting information

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Computing details

Data collection: *APEX3* (Bruker, 2018); cell refinement: *SAINTE* (Bruker, 2018); data reduction: *SAINTE* (Bruker, 2018); program(s) used to solve structure: *SHELXT* (Sheldrick, 2015a); program(s) used to refine structure: *SHELXL2018* (Sheldrick, 2015b); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 2012); software used to prepare material for publication: *PLATON* (Spek, 2020).

(*E*)-2-(4-Bromophenyl)-1-[2,2-dibromo-1-(3-nitrophenyl)ethenyl]diazene

Crystal data

$C_{14}H_8Br_3N_3O_2$

$M_r = 489.96$

Orthorhombic, *Pbca*

$a = 6.6579$ (1) Å

$b = 15.7683$ (3) Å

$c = 29.0301$ (6) Å

$V = 3047.69$ (10) Å³

$Z = 8$

$F(000) = 1872$

$D_x = 2.136$ Mg m⁻³

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

Cell parameters from 9783 reflections

$\theta = 2.7\text{--}34.8^\circ$

$\mu = 7.95$ mm⁻¹

$T = 100$ K

Needle, red

$0.34 \times 0.06 \times 0.05$ mm

Data collection

Bruker AXS D8 QUEST, Photon III detector
diffractometer

Radiation source: fine-focus sealed X-Ray tube

Graphite monochromator

Detector resolution: 7.31 pixels mm⁻¹

φ and ω shutterless scans

Absorption correction: multi-scan
(SADABS; Krause *et al.*, 2015).

$T_{\min} = 0.020$, $T_{\max} = 0.058$

67179 measured reflections

5538 independent reflections

4620 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.033$

$\theta_{\max} = 32.6^\circ$, $\theta_{\min} = 2.6^\circ$

$h = -10 \rightarrow 10$

$k = -23 \rightarrow 23$

$l = -43 \rightarrow 43$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.024$

$wR(F^2) = 0.064$

$S = 1.05$

5538 reflections

199 parameters

0 restraints

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

$$w = 1/[\sigma^2(F_o^2) + (0.0318P)^2 + 2.0035P]$$

where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.002$

$$\Delta\rho_{\max} = 0.68 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.52 \text{ e } \text{\AA}^{-3}$$

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds 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 (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
Br1	-0.15339 (3)	0.13530 (2)	0.34586 (2)	0.02722 (5)
Br2	0.15481 (3)	0.10023 (2)	0.42549 (2)	0.02936 (5)
Br3	1.20530 (3)	0.41027 (2)	0.48664 (2)	0.02853 (5)
O1	-0.0475 (3)	0.54340 (9)	0.25282 (6)	0.0379 (3)
O2	0.0602 (3)	0.54406 (9)	0.32311 (5)	0.0353 (3)
N1	0.0276 (2)	0.50824 (10)	0.28639 (6)	0.0272 (3)
N2	0.3641 (2)	0.25393 (10)	0.38915 (6)	0.0238 (3)
N3	0.4595 (2)	0.31953 (10)	0.37700 (5)	0.0237 (3)
C1	0.0855 (3)	0.16829 (11)	0.37477 (6)	0.0238 (3)
C2	0.1967 (3)	0.23522 (11)	0.36103 (6)	0.0226 (3)
C3	0.1509 (2)	0.28572 (11)	0.31884 (6)	0.0215 (3)
C4	0.1146 (3)	0.37242 (11)	0.32230 (6)	0.0221 (3)
H4	0.115991	0.400016	0.351386	0.027*
C5	0.0763 (3)	0.41751 (11)	0.28225 (6)	0.0224 (3)
C6	0.0770 (3)	0.38058 (12)	0.23897 (6)	0.0236 (3)
H6	0.050338	0.413167	0.212131	0.028*
C7	0.1178 (3)	0.29485 (13)	0.23606 (6)	0.0250 (3)
H7	0.121139	0.268071	0.206755	0.030*
C8	0.1540 (3)	0.24729 (11)	0.27555 (6)	0.0228 (3)
H8	0.180911	0.188307	0.273038	0.027*
C9	0.6294 (3)	0.33715 (12)	0.40511 (6)	0.0233 (3)
C10	0.7117 (3)	0.27990 (12)	0.43672 (7)	0.0254 (3)
H10	0.650813	0.226090	0.441376	0.030*
C11	0.8820 (3)	0.30204 (12)	0.46115 (7)	0.0267 (3)
H11	0.940184	0.263540	0.482488	0.032*
C12	0.9671 (3)	0.38174 (12)	0.45398 (6)	0.0239 (3)
C13	0.8856 (3)	0.43973 (12)	0.42367 (6)	0.0257 (3)
H13	0.944333	0.494183	0.419849	0.031*
C14	0.7154 (3)	0.41659 (12)	0.39883 (6)	0.0248 (3)
H14	0.657723	0.455267	0.377500	0.030*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.02466 (8)	0.02296 (8)	0.03404 (10)	-0.00303 (6)	-0.00384 (7)	0.00397 (7)
Br2	0.02669 (9)	0.02915 (9)	0.03225 (10)	0.00022 (7)	-0.00123 (7)	0.01012 (7)
Br3	0.02656 (9)	0.02931 (9)	0.02972 (9)	-0.00397 (7)	-0.00470 (7)	0.00276 (7)
O1	0.0417 (9)	0.0284 (7)	0.0437 (9)	0.0056 (6)	-0.0074 (7)	0.0110 (7)
O2	0.0422 (8)	0.0260 (7)	0.0378 (8)	0.0048 (6)	0.0005 (7)	-0.0047 (6)
N1	0.0234 (7)	0.0231 (7)	0.0352 (8)	0.0012 (6)	0.0018 (6)	0.0035 (6)
N2	0.0218 (6)	0.0251 (7)	0.0246 (7)	-0.0002 (5)	0.0002 (6)	0.0006 (6)
N3	0.0220 (6)	0.0243 (7)	0.0248 (7)	0.0003 (5)	-0.0002 (5)	0.0003 (6)
C1	0.0222 (7)	0.0227 (7)	0.0264 (8)	0.0017 (6)	-0.0009 (6)	0.0028 (6)
C2	0.0223 (7)	0.0222 (7)	0.0232 (8)	0.0019 (6)	-0.0003 (6)	0.0003 (6)
C3	0.0188 (7)	0.0219 (7)	0.0237 (8)	0.0000 (6)	0.0000 (6)	0.0010 (6)
C4	0.0212 (7)	0.0215 (7)	0.0237 (8)	0.0007 (6)	0.0006 (6)	0.0006 (6)
C5	0.0185 (7)	0.0214 (7)	0.0274 (8)	0.0001 (6)	0.0003 (6)	0.0016 (6)
C6	0.0177 (7)	0.0290 (8)	0.0242 (8)	-0.0007 (6)	0.0004 (6)	0.0036 (6)
C7	0.0210 (7)	0.0307 (9)	0.0232 (8)	-0.0013 (6)	0.0006 (6)	-0.0031 (7)
C8	0.0200 (7)	0.0223 (7)	0.0262 (8)	-0.0001 (6)	-0.0013 (6)	-0.0026 (6)
C9	0.0235 (7)	0.0235 (8)	0.0230 (8)	0.0007 (6)	0.0002 (6)	-0.0008 (6)
C10	0.0245 (8)	0.0241 (8)	0.0275 (8)	-0.0015 (6)	-0.0015 (7)	0.0032 (7)
C11	0.0274 (8)	0.0252 (8)	0.0275 (9)	0.0002 (7)	-0.0035 (7)	0.0030 (7)
C12	0.0225 (7)	0.0263 (8)	0.0227 (8)	-0.0011 (6)	-0.0006 (6)	-0.0012 (6)
C13	0.0281 (8)	0.0232 (8)	0.0259 (8)	-0.0024 (7)	-0.0004 (7)	0.0004 (6)
C14	0.0266 (8)	0.0235 (8)	0.0244 (8)	0.0014 (6)	-0.0004 (6)	0.0028 (6)

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

Br1—C1	1.8723 (18)	C6—C7	1.381 (3)
Br2—C1	1.8796 (18)	C6—H6	0.9500
Br3—C12	1.9016 (18)	C7—C8	1.391 (3)
O1—N1	1.227 (2)	C7—H7	0.9500
O2—N1	1.226 (2)	C8—H8	0.9500
N1—C5	1.472 (2)	C9—C14	1.389 (3)
N2—N3	1.264 (2)	C9—C10	1.399 (3)
N2—C2	1.413 (2)	C10—C11	1.382 (3)
N3—C9	1.422 (2)	C10—H10	0.9500
C1—C2	1.349 (2)	C11—C12	1.394 (3)
C2—C3	1.493 (2)	C11—H11	0.9500
C3—C4	1.392 (2)	C12—C13	1.380 (3)
C3—C8	1.395 (3)	C13—C14	1.392 (3)
C4—C5	1.386 (2)	C13—H13	0.9500
C4—H4	0.9500	C14—H14	0.9500
C5—C6	1.385 (3)		
O2—N1—O1	123.67 (17)	C6—C7—H7	119.6
O2—N1—C5	118.68 (16)	C8—C7—H7	119.6
O1—N1—C5	117.64 (17)	C7—C8—C3	120.37 (17)

N3—N2—C2	113.96 (15)	C7—C8—H8	119.8
N2—N3—C9	113.53 (15)	C3—C8—H8	119.8
C2—C1—Br1	123.43 (14)	C14—C9—C10	120.46 (17)
C2—C1—Br2	122.92 (14)	C14—C9—N3	115.35 (16)
Br1—C1—Br2	113.64 (9)	C10—C9—N3	124.18 (17)
C1—C2—N2	115.18 (16)	C11—C10—C9	119.65 (17)
C1—C2—C3	123.21 (16)	C11—C10—H10	120.2
N2—C2—C3	121.60 (15)	C9—C10—H10	120.2
C4—C3—C8	119.61 (16)	C10—C11—C12	118.98 (17)
C4—C3—C2	120.02 (16)	C10—C11—H11	120.5
C8—C3—C2	120.30 (16)	C12—C11—H11	120.5
C5—C4—C3	118.36 (17)	C13—C12—C11	122.18 (17)
C5—C4—H4	120.8	C13—C12—Br3	119.28 (14)
C3—C4—H4	120.8	C11—C12—Br3	118.53 (14)
C6—C5—C4	122.99 (17)	C12—C13—C14	118.48 (17)
C6—C5—N1	118.92 (16)	C12—C13—H13	120.8
C4—C5—N1	118.07 (16)	C14—C13—H13	120.8
C7—C6—C5	117.88 (17)	C9—C14—C13	120.23 (17)
C7—C6—H6	121.1	C9—C14—H14	119.9
C5—C6—H6	121.1	C13—C14—H14	119.9
C6—C7—C8	120.75 (17)		
C2—N2—N3—C9	-179.14 (15)	C4—C5—C6—C7	0.1 (3)
Br1—C1—C2—N2	175.57 (13)	N1—C5—C6—C7	-178.28 (16)
Br2—C1—C2—N2	-3.5 (2)	C5—C6—C7—C8	0.9 (3)
Br1—C1—C2—C3	-5.5 (3)	C6—C7—C8—C3	-0.4 (3)
Br2—C1—C2—C3	175.44 (13)	C4—C3—C8—C7	-1.0 (3)
N3—N2—C2—C1	-176.40 (16)	C2—C3—C8—C7	-177.89 (16)
N3—N2—C2—C3	4.6 (2)	N2—N3—C9—C14	-168.04 (16)
C1—C2—C3—C4	121.4 (2)	N2—N3—C9—C10	13.1 (3)
N2—C2—C3—C4	-59.7 (2)	C14—C9—C10—C11	-1.3 (3)
C1—C2—C3—C8	-61.7 (2)	N3—C9—C10—C11	177.51 (18)
N2—C2—C3—C8	117.13 (19)	C9—C10—C11—C12	0.6 (3)
C8—C3—C4—C5	1.9 (2)	C10—C11—C12—C13	0.8 (3)
C2—C3—C4—C5	178.81 (16)	C10—C11—C12—Br3	-178.42 (15)
C3—C4—C5—C6	-1.5 (3)	C11—C12—C13—C14	-1.5 (3)
C3—C4—C5—N1	176.87 (15)	Br3—C12—C13—C14	177.71 (14)
O2—N1—C5—C6	-167.63 (17)	C10—C9—C14—C13	0.6 (3)
O1—N1—C5—C6	13.8 (2)	N3—C9—C14—C13	-178.32 (17)
O2—N1—C5—C4	13.9 (2)	C12—C13—C14—C9	0.8 (3)
O1—N1—C5—C4	-164.68 (17)		

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C8—H8 \cdots O1 ⁱ	0.95	2.56	3.394 (2)	146

Symmetry code: (i) $-x, y-1/2, -z+1/2$.