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2,2-Dichloro-*N*-(3-nitrophenyl)-acetamideB. Thimme Gowda,^{a*} Sabine Foro^b and Hartmut Fuess^b^aDepartment of Chemistry, Mangalore University, Mangalagangothri 574 199, Mangalore, India, and ^bInstitute of Materials Science, Darmstadt University of Technology, Petersenstrasse 23, D-64287 Darmstadt, Germany

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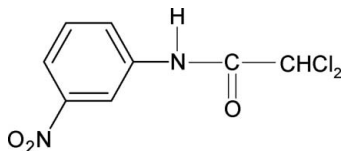
Received 27 December 2007; accepted 31 December 2007

Key indicators: single-crystal X-ray study; $T = 299$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.033; wR factor = 0.099; data-to-parameter ratio = 13.4.

The conformation of the N—H bond in the structure of the title compound (3NPDCa), $\text{C}_8\text{H}_6\text{Cl}_2\text{N}_2\text{O}_3$, is *anti* to the *meta*-nitro group, similar to that in the structures of 2-chloro-*N*-(3-nitrophenyl)acetamide (3NPCA) and 2,2,2-trichloro-*N*-(3-nitrophenyl)acetamide (3NPTCA), and the *meta*-chloro group in 2,2-dichloro-*N*-(3-chlorophenyl)acetamide (3CPDCA). The geometric parameters of 3NPDCa are similar to those of 2,2-dichloro-*N*-phenylacetamide, 3CPDCA, 3NPCA, 3NPTCA and other acetanilides. Intermolecular N—H...O hydrogen bonds link the molecules into chains running along the *b* axis.

Related literature

For related literature, see: Gowda & Weiss (1994); Gowda *et al.* (2000, 2006, 2007).



Experimental

Crystal data

 $\text{C}_8\text{H}_6\text{Cl}_2\text{N}_2\text{O}_3$ $M_r = 249.05$ Orthorhombic, *Pbca* $a = 9.6092$ (6) Å $b = 10.6487$ (7) Å $c = 19.868$ (1) Å $V = 2033.0$ (2) Å³ $Z = 8$ Mo $K\alpha$ radiation $\mu = 0.63$ mm⁻¹ $T = 299$ (2) K $0.60 \times 0.52 \times 0.24$ mm

Data collection

Oxford Diffraction Xcalibur diffractometer with Sapphire CCD Detector

Absorption correction: multi-scan (*SCALE3 ABSPACK*);

Oxford Diffraction, 2007)

 $T_{\min} = 0.706$, $T_{\max} = 0.865$

11267 measured reflections

2072 independent reflections

1614 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.022$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.032$ $wR(F^2) = 0.098$ $S = 1.10$

2072 reflections

155 parameters

1 restraint

H atoms treated by a mixture of independent and constrained refinement

 $\Delta\rho_{\text{max}} = 0.35$ e Å⁻³ $\Delta\rho_{\text{min}} = -0.36$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N1}-\text{H1N}\cdots\text{O1}^i$	0.833 (16)	2.081 (17)	2.907 (2)	171 (2)

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

Data collection: *CrysAlis CCD* (Oxford Diffraction, 2004); cell refinement: *CrysAlis RED* (Oxford Diffraction, 2007); data reduction: *CrysAlis RED*; 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: *SHELXS97*.

BTG thanks the Alexander von Humboldt Foundation, Bonn, Germany, for extensions of his research fellowship.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: DN2307).

References

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supplementary materials

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2,2-Dichloro-*N*-(3-nitrophenyl)acetamide

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Comment

As part of a study of the effect of ring and side chain substitutions on the solid state structures of acetanilides (Gowda *et al.*, 2000, 2006, 2007), in the present work, the crystal structure of 2,2-dichloro-*N*-(3-nitrophenyl)-acetamide (3NPDCa) has been determined to explore the effects of polar substituent groups on the structures of *N*-aromatic amides. The conformation of the N—H bond in the structure of 3NPDCa (Fig.1) is *anti* to the *meta* nitro group, similar to that in the structure of 2-chloro-*N*-(3-nitrophenyl)-acetamide (3NPCa) (Gowda *et al.*, 2007) and 2,2,2-trichloro-*N*-(3-nitrophenyl)-acetamide (3NPtCa)(Gowda *et al.*, 2000) and *meta* chloro group in 2,2-dichloro-*N*-(3-chlorophenyl)-acetamide (3CPDCa)(Gowda *et al.*, 2006). The geometric parameters in 3NPDCa are similar to those of 2,2-dichloro-*N*-(phenyl)-acetamide, 3CPDCa, 3NPCa, 3NPtCa and other acetanilides. The intermolecular N—H···O hydrogen bonds (Table 1) link the molecules into chains running along the *b* axis (Fig. 2).

Experimental

The title compound was prepared similar to the literature method (Gowda and Weiss, 1994). The purity of the compound was checked by determining its melting point. It was characterized by recording its infrared and NQR spectra (Gowda and Weiss, 1994). Single crystals of the title compound were obtained from an ethanolic solution and used for X-ray diffraction studies at room temperature.

Refinement

The H atoms were located in difference map with C—H = 0.89 (3)–0.96 (3) Å and N—H distance was restrained to 0.86 (2) %Å. All H atoms were refined with isotropic displacement parameters (set to 1.2 times of the U_{eq} of the parent atom).

Figures

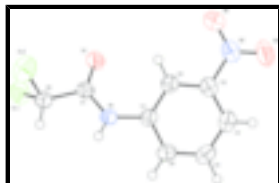


Fig. 1. Molecular structure of the title compound showing the atom labelling scheme. Displacement ellipsoids are drawn at the 50% probability level. H atoms are represented as small spheres of arbitrary radii.

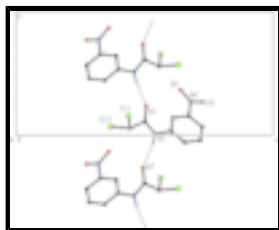


Fig. 2. Partial packing view showing the hydrogen bonding as dashed lines. H atoms not involved in hydrogen bonding have been omitted for clarity. [Symmetry code: (i) $x - 1/2, 1/2 - y, 1 - z$]

2,2-Dichloro-*N*-(3-nitrophenyl)acetamide

Crystal data

C₈H₆Cl₂N₂O₃

M_r = 249.05

Orthorhombic, *Pbca*

Hall symbol: -P 2ac 2ab

a = 9.6092 (6) Å

b = 10.6487 (7) Å

c = 19.868 (1) Å

V = 2033.0 (2) Å³

Z = 8

*F*₀₀₀ = 1008

D_x = 1.627 Mg m⁻³

Mo *K*α radiation

λ = 0.71073 Å

Cell parameters from 4893 reflections

θ = 2.8–27.8°

μ = 0.63 mm⁻¹

T = 299 (2) K

Prism, yellow

0.60 × 0.52 × 0.24 mm

Data collection

Oxford Diffraction Xcalibur
diffractometer with Sapphire CCD Detector

Radiation source: fine-focus sealed tube

Monochromator: graphite

T = 299(2) K

Rotation method data acquisition using ω and φ
scans.

Absorption correction: multi-scan
(SCALE3 ABSPACK; Oxford Diffraction, 2007)

T_{min} = 0.706, *T_{max}* = 0.865

11267 measured reflections

2072 independent reflections

1614 reflections with *I* > 2σ(*I*)

R_{int} = 0.022

θ_{max} = 26.4°

θ_{min} = 3.0°

h = -12→11

k = -11→13

l = -24→20

Refinement

Refinement on *F*²

Least-squares matrix: full

R [*F*² > 2σ(*F*²)] = 0.032

wR(*F*²) = 0.098

S = 1.10

2072 reflections

155 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 atoms treated by a mixture of
independent and constrained refinement

$$w = 1/[\sigma^2(F_o^2) + (0.0385P)^2 + 1.4567P]$$

where *P* = (*F_o*² + 2*F_c*²)/3

(Δ/σ)_{max} = 0.044

Δρ_{max} = 0.35 e Å⁻³

Δρ_{min} = -0.36 e Å⁻³

Extinction correction: SHELXL,

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

Extinction coefficient: 0.0230 (13)

Special details

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

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

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.03253 (19)	0.14098 (18)	0.60262 (10)	0.0344 (4)
C2	0.1408 (2)	0.05696 (19)	0.61012 (11)	0.0365 (4)
H2	0.204 (2)	0.043 (2)	0.5757 (12)	0.044*
C3	0.1543 (2)	-0.00244 (19)	0.67144 (11)	0.0380 (5)
C4	0.0653 (2)	0.0166 (2)	0.72466 (11)	0.0453 (5)
H4	0.075 (3)	-0.025 (2)	0.7632 (14)	0.054*
C5	-0.0418 (2)	0.1009 (2)	0.71576 (12)	0.0490 (6)
H5	-0.103 (3)	0.122 (2)	0.7521 (14)	0.059*
C6	-0.0584 (2)	0.1632 (2)	0.65542 (11)	0.0426 (5)
H6	-0.131 (3)	0.220 (2)	0.6503 (13)	0.051*
C7	0.1102 (2)	0.24984 (19)	0.50064 (11)	0.0357 (4)
C8	0.0552 (2)	0.3297 (2)	0.44287 (11)	0.0398 (5)
H8	-0.034 (3)	0.352 (2)	0.4488 (11)	0.048*
N1	0.01034 (17)	0.20745 (17)	0.54149 (9)	0.0381 (4)
H1N	-0.0711 (18)	0.227 (2)	0.5320 (12)	0.046*
N2	0.2700 (2)	-0.09083 (18)	0.67926 (10)	0.0491 (5)
O1	0.23372 (15)	0.22854 (16)	0.50684 (9)	0.0532 (5)
O2	0.3669 (2)	-0.08426 (19)	0.63930 (10)	0.0664 (5)
O3	0.2672 (2)	-0.16436 (19)	0.72637 (10)	0.0759 (6)
Cl1	0.15452 (6)	0.46874 (5)	0.43763 (3)	0.0529 (2)
Cl2	0.06927 (8)	0.24208 (7)	0.36778 (3)	0.0661 (2)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0270 (9)	0.0403 (10)	0.0360 (10)	-0.0056 (8)	-0.0003 (8)	0.0025 (8)
C2	0.0334 (10)	0.0388 (10)	0.0374 (10)	-0.0034 (8)	-0.0001 (8)	0.0008 (8)
C3	0.0363 (10)	0.0346 (10)	0.0430 (11)	-0.0033 (8)	-0.0046 (9)	0.0017 (8)
C4	0.0494 (13)	0.0483 (12)	0.0382 (11)	-0.0071 (10)	-0.0014 (10)	0.0065 (10)
C5	0.0460 (13)	0.0583 (14)	0.0428 (12)	-0.0043 (11)	0.0103 (10)	-0.0006 (10)
C6	0.0311 (10)	0.0487 (12)	0.0479 (12)	-0.0012 (9)	0.0036 (9)	0.0014 (10)
C7	0.0272 (9)	0.0389 (10)	0.0409 (10)	0.0013 (8)	-0.0009 (8)	0.0020 (8)

supplementary materials

C8	0.0307 (10)	0.0479 (12)	0.0409 (11)	0.0006 (9)	-0.0001 (9)	0.0045 (9)
N1	0.0239 (8)	0.0503 (10)	0.0401 (9)	0.0018 (7)	-0.0002 (7)	0.0076 (8)
N2	0.0507 (11)	0.0446 (10)	0.0521 (11)	0.0028 (9)	-0.0091 (10)	0.0056 (9)
O1	0.0258 (7)	0.0698 (11)	0.0641 (10)	0.0050 (7)	0.0044 (7)	0.0272 (8)
O2	0.0606 (11)	0.0718 (12)	0.0668 (12)	0.0250 (10)	0.0086 (10)	0.0109 (10)
O3	0.0786 (13)	0.0713 (12)	0.0778 (13)	0.0122 (11)	-0.0056 (11)	0.0361 (11)
Cl1	0.0558 (4)	0.0395 (3)	0.0635 (4)	-0.0031 (2)	-0.0001 (3)	0.0064 (2)
Cl2	0.0740 (5)	0.0778 (5)	0.0467 (4)	-0.0200 (4)	-0.0084 (3)	-0.0116 (3)

Geometric parameters (Å, °)

C1—C2	1.380 (3)	C6—H6	0.93 (3)
C1—C6	1.386 (3)	C7—O1	1.215 (2)
C1—N1	1.422 (2)	C7—N1	1.335 (3)
C2—C3	1.379 (3)	C7—C8	1.523 (3)
C2—H2	0.93 (2)	C8—Cl1	1.764 (2)
C3—C4	1.376 (3)	C8—Cl2	1.765 (2)
C3—N2	1.464 (3)	C8—H8	0.90 (3)
C4—C5	1.377 (3)	N1—H1N	0.833 (16)
C4—H4	0.89 (3)	N2—O3	1.220 (3)
C5—C6	1.380 (3)	N2—O2	1.226 (3)
C5—H5	0.96 (3)		
C2—C1—C6	120.30 (19)	C1—C6—H6	120.0 (16)
C2—C1—N1	121.86 (17)	O1—C7—N1	125.26 (19)
C6—C1—N1	117.83 (18)	O1—C7—C8	121.32 (19)
C3—C2—C1	117.65 (19)	N1—C7—C8	113.42 (17)
C3—C2—H2	120.9 (15)	C7—C8—Cl1	108.99 (14)
C1—C2—H2	121.5 (15)	C7—C8—Cl2	108.35 (15)
C4—C3—C2	123.5 (2)	Cl1—C8—Cl2	110.62 (12)
C4—C3—N2	119.03 (19)	C7—C8—H8	112.2 (15)
C2—C3—N2	117.42 (19)	Cl1—C8—H8	107.9 (16)
C3—C4—C5	117.6 (2)	Cl2—C8—H8	108.8 (15)
C3—C4—H4	121.6 (17)	C7—N1—C1	125.44 (17)
C5—C4—H4	120.8 (17)	C7—N1—H1N	116.8 (17)
C4—C5—C6	120.8 (2)	C1—N1—H1N	117.5 (17)
C4—C5—H5	120.9 (16)	O3—N2—O2	123.4 (2)
C6—C5—H5	118.2 (16)	O3—N2—C3	118.5 (2)
C5—C6—C1	120.2 (2)	O2—N2—C3	118.10 (18)
C5—C6—H6	119.8 (16)		
C6—C1—C2—C3	-0.2 (3)	N1—C7—C8—Cl1	131.85 (17)
N1—C1—C2—C3	179.62 (18)	O1—C7—C8—Cl2	71.9 (2)
C1—C2—C3—C4	0.8 (3)	N1—C7—C8—Cl2	-107.71 (18)
C1—C2—C3—N2	-179.58 (18)	O1—C7—N1—C1	6.6 (4)
C2—C3—C4—C5	-0.8 (3)	C8—C7—N1—C1	-173.76 (18)
N2—C3—C4—C5	179.6 (2)	C2—C1—N1—C7	-35.6 (3)
C3—C4—C5—C6	0.1 (3)	C6—C1—N1—C7	144.2 (2)
C4—C5—C6—C1	0.5 (3)	C4—C3—N2—O3	15.6 (3)
C2—C1—C6—C5	-0.4 (3)	C2—C3—N2—O3	-164.0 (2)
N1—C1—C6—C5	179.8 (2)	C4—C3—N2—O2	-162.0 (2)

O1—C7—C8—C11

-48.5 (3)

C2—C3—N2—O2

18.3 (3)

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

D—H \cdots *A*

D—H

H \cdots *A*

D \cdots *A*

D—H \cdots *A*

N1—H1N \cdots O1ⁱ

0.833 (16)

2.081 (17)

2.907 (2)

171 (2)

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

Fig. 1

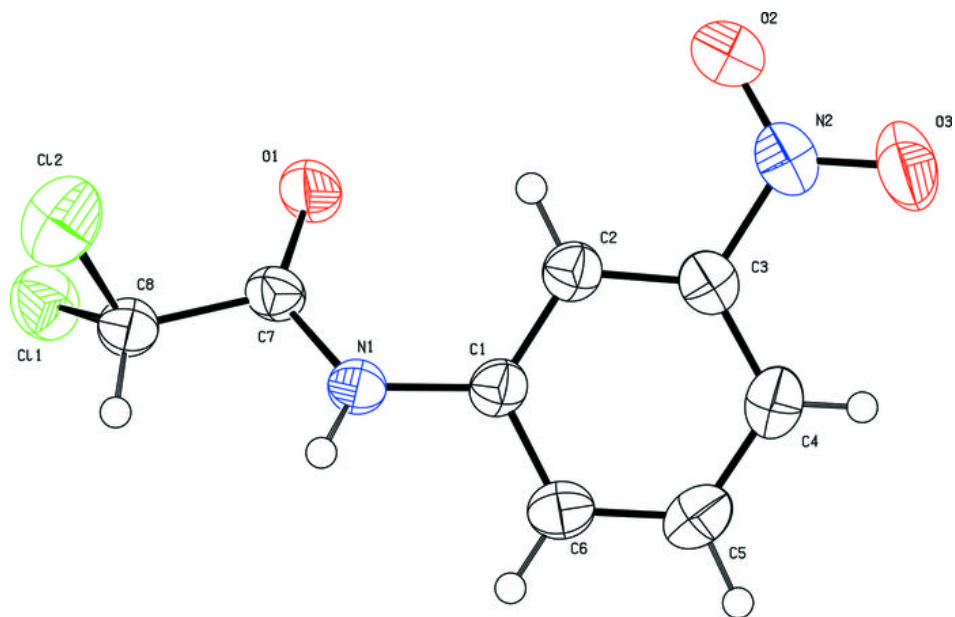


Fig. 2

