2-Chloro-*N*-phenylacetamide

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In the title compound, C_8H_8ClNO, the conformations of the N—H and C=O bonds are *anti* to each other, but the C—Cl and C=O bonds in the side chain are *syn*. The molecules are linked by N—H⋯O hydrogen bonds into infinite chains running in the [101] direction.

Related literature

For the synthesis, see: Gowda *et al.* (2003). For related structures, see: Gowda *et al.* (2007, 2008).
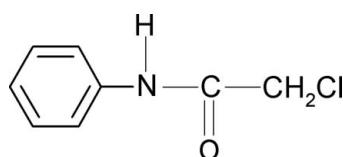
Experimental

Crystal data

C_8H_8ClNO
$M_r = 169.6$
Monoclinic, Cc
$a = 5.0623$ (15) Å
$b = 18.361$ (6) Å
$c = 9.115$ (2) Å
$\beta = 102.13$ (3)°

$V = 828.3$ (4) Å^3
$Z = 4$
Mo $K\alpha$ radiation
$\mu = 0.40$ mm^{-1}
$T = 297$ (2) K
$0.41 \times 0.24 \times 0.17$ mm

Data collection

Oxford Diffraction Xcalibur diffractometer
Absorption correction: analytical [*CrysAlis RED* (Oxford Diffraction, 2006), using a multi-faceted crystal model based on expressions derived by Clark &
Reid (1995)]
$T_{min} = 0.905$, $T_{max} = 0.938$
2388 measured reflections
1067 independent reflections
385 reflections with $I > 2\sigma(I)$
$R_{int} = 0.046$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.036$
$wR(F^2) = 0.086$
$S = 0.96$
1067 reflections
106 parameters
2 restraints

H-atom parameters constrained
$\Delta\rho_{max} = 0.1$ e Å^{-3}
$\Delta\rho_{min} = -0.11$ e Å^{-3}
Absolute structure: Flack (1983), 254 Friedel pairs
Flack parameter: 0.04 (11)

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

D—H⋯A	D—H	H⋯A	D⋯A	D—H⋯A
N1—H1N⋯O1^i	0.86	2.05	2.848 (5)	155

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

Data collection: *CrysAlis CCD* (Oxford Diffraction, 2006); cell refinement: *CrysAlis RED* (Oxford Diffraction, 2006); data reduction: *CrysAlis RED*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997) and *DIAMOND* (Brandenburg, 2002); software used to prepare material for publication: *SHELXL97*, *PLATON* (Spek, 2003) and *WinGX* (Farrugia, 1999).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB2728).

References

Brandenburg, K. (2002). *DIAMOND*. Bonn, Germany.
Clark, R. C. & Reid, J. S. (1995). *Acta Cryst.* A**51**, 887–897.
Farrugia, L. J. (1997). *J. Appl. Cryst.* **30**, 565.
Farrugia, L. J. (1999). *J. Appl. Cryst.* **32**, 837–838.
Flack, H. D. (1983). *Acta Cryst.* A**39**, 876–881.
Gowda, B. T., Foro, S. & Fuess, H. (2007). *Acta Cryst.* E**63**, o4611.
Gowda, B. T., Svoboda, I., Foro, S., Dou, S. & Fuess, H. (2008). *Acta Cryst.* E**64**, o208.
Gowda, B. T., Usha, K. M. & Jayalakshmi, K. L. (2003). *Z. Naturforsch. Teil A*, **58**, 801–806.
Oxford Diffraction (2006). *CrysAlis CCD* and *CrysAlis RED*. Oxford Diffraction Ltd, Abingdon, Oxfordshire, England.
Sheldrick, G. M. (2008). *Acta Cryst.* A**64**, 112–122.
Spek, A. L. (2003). *J. Appl. Cryst.* **36**, 7–13.

supporting information

2-Chloro-*N*-phenylacetamide

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S1. Comment

In the present work, the structure of the title compoud, (I), 2-chloro-*N*-(phenyl)-acetamide (NPCA) has been determined, as part of a study of the effect of ring and side chain substitutions on the solid state geometry of aromatic amides (Gowda *et al.*, 2007; 2008). The conformations of the N—H and C=O bonds are *anti* to each other, but the C—Cl and C=O bonds in the side chain are *syn* to each other (Fig. 1), similar to that observed in 2-chloro-*N*-(2-chlorophenyl)-acetamide (Gowda *et al.*, 2007)and 2-chloro-*N*-(3-methylphenyl)-acetamide (Gowda *et al.*, 2008) with similar bond parameters. Further, the amide group –NHCO– in (I) makes a dihedral angle of 16.0 (8)° with the phenyl ring.

   Part of the packing for (I) viewed down the *b* axis is shown in Fig. 2. Infinite chains running along the base vector [101] are formed by N-H···O hydrogen bonds (Table 1).

S2. Experimental

The title compound was prepared according to the literature method (Gowda *et al.*, 2003) and colourless prisms of (I) were recrystallised from an ethanol solution.

S3. Refinement

The H atoms were placed in calculated positions (C-H = 0.93Å, N-H = 0.86Å) and refined as riding with U_{iso}(H) = 1.2U_{eq}(C,*N*).
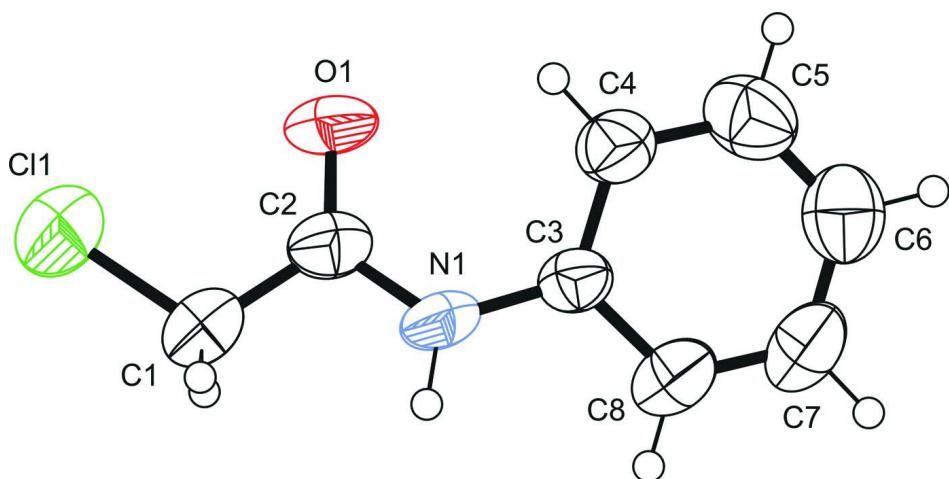
Figure 1

Molecular structure of (I) with displacement ellipsoids for the non-hydrogen atoms drawn at the 50% probability level.
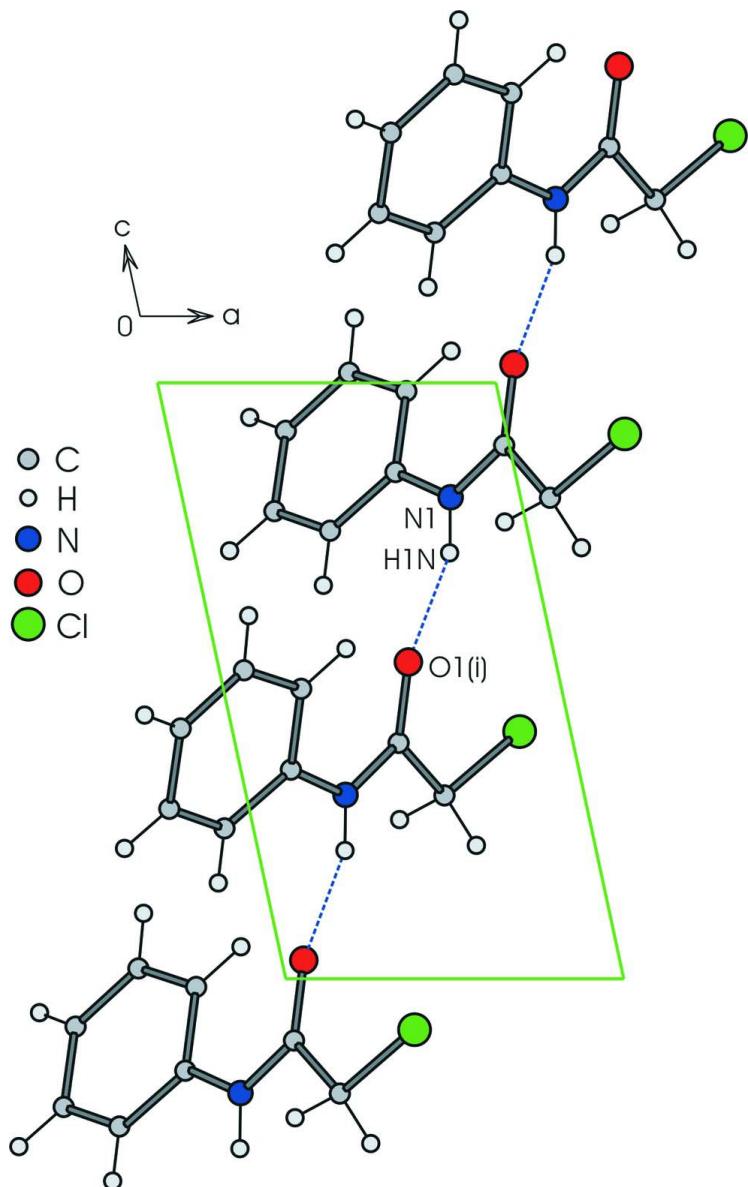
Figure 2

Part of the packing for (I) viewed down the *b* axis showing the chains arising from N-H···O hydrogen bonds Symmetry code (i): *x* - 1/2,-*y* + 1/2,*z* - 1/2.

2-Chloro-*N*-phenylacetamide

Crystal data

C_8H_8ClNO	$V = 828.3 (4)$ Å^3
$M_r = 169.6$	$Z = 4$
Monoclinic, *Cc*	$F(000) = 352$
Hall symbol: C -2yc	$D_x = 1.36$ Mg m^{-3}
$a = 5.0623 (15)$ Å	Mo *Kα* radiation, $λ = 0.71073$ Å
$b = 18.361 (6)$ Å	Cell parameters from 159 reflections
$c = 9.115 (2)$ Å	$θ = 4.9–25.1°$
$β = 102.13 (3)°$	$μ = 0.40$ mm^{-1}

T = 297 K

Prism, colorless

0.41 × 0.24 × 0.17 mm

Data collection

Oxford Diffraction Xcalibur System diffractometer	T_{min} = 0.905, T_{max} = 0.938
Radiation source: Enhance (Mo) X-ray Source	2388 measured reflections
Graphite monochromator	1067 independent reflections
Detector resolution: 10.4340 pixels mm^{-1}	385 reflections with $I > 2\sigma(I)$
ω scans	R_{int} = 0.046
Absorption correction: analytical	θ_{max} = 26°, θ_{min} = 4.3°
[*CrysAlis RED* (Oxford Diffraction, 2006),	$h = -6\rightarrow6$
using a multifaceted crystal model based on	$k = -22\rightarrow22$
expressions derived by Clark & Reid (1995)]	$l = -9\rightarrow11$

Refinement

Refinement on F^2	Hydrogen site location: inferred from neighbouring sites
Least-squares matrix: full	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)]$ = 0.036	$[\exp(3.70(\sin\theta/\lambda)^2)]/[\sigma^2(F_o^2) + (0.035P)^2]$
$wR(F^2)$ = 0.086	where $P = 0.33333F_o^2 + 0.66667F_c^2$
S = 0.96	$(\Delta/\sigma)_{max} < 0.001$
1067 reflections	$\Delta\rho_{max}$ = 0.1 e Å^{-3}
106 parameters	$\Delta\rho_{min}$ = −0.11 e Å^{-3}
2 restraints	Absolute structure: Flack (1983), 254 Friedel pairs
Primary atom site location: structure-invariant direct methods	Absolute structure parameter: 0.04 (11)
Secondary atom site location: difference Fourier map	

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 (Å^2)

	x	y	z	U_{iso}*/U_{eq}
Cl1	1.3492 (4)	0.15000 (9)	0.9151 (2)	0.1082 (7)
C1	1.0871 (11)	0.1998 (3)	0.8076 (6)	0.0824 (18)
H1A	1.1488	0.2215	0.7237	0.099*
H1B	0.939	0.1671	0.7672	0.099*
C2	0.9849 (11)	0.2595 (3)	0.8961 (6)	0.0629 (17)
N1	0.7939 (8)	0.3008 (2)	0.8086 (4)	0.0642 (13)
H1N	0.7565	0.2897	0.7149	0.077*
O1	1.0653 (7)	0.26846 (19)	1.0314 (3)	0.0833 (13)
C3	0.6481 (10)	0.3597 (3)	0.8507 (6)	0.0517 (13)
C4	0.7302 (11)	0.3975 (3)	0.9862 (6)	0.0670 (17)
H4	0.888	0.3845	1.0536	0.08*

C5	0.5727 (15)	0.4542 (3)	1.0177 (7)	0.082 (2)
H5	0.6222	0.478	1.1094	0.099*
C6	0.3490 (16)	0.4762 (3)	0.9197 (10)	0.0824 (18)
H6	0.2513	0.516	0.9428	0.099*
C7	0.2640 (13)	0.4399 (4)	0.7850 (7)	0.082 (2)
H7	0.1068	0.4538	0.7183	0.099*
C8	0.4153 (10)	0.3835 (4)	0.7525 (6)	0.0676 (16)
H8	0.3614	0.3598	0.6609	0.081*

Atomic displacement parameters (Å^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cl1	0.1120 (13)	0.1207 (13)	0.0829 (10)	0.0402 (13)	−0.0004 (9)	0.0083 (13)
C1	0.079 (4)	0.091 (4)	0.067 (4)	0.022 (4)	−0.006 (3)	0.006 (4)
C2	0.070 (4)	0.071 (4)	0.045 (3)	−0.003 (3)	0.005 (3)	0.004 (4)
N1	0.066 (3)	0.084 (3)	0.035 (3)	0.012 (3)	−0.006 (2)	0.003 (3)
O1	0.102 (3)	0.097 (3)	0.039 (2)	0.010 (2)	−0.013 (2)	−0.005 (2)
C3	0.051 (4)	0.062 (4)	0.041 (3)	−0.001 (3)	0.008 (3)	0.002 (3)
C4	0.055 (4)	0.084 (4)	0.059 (4)	0.005 (4)	0.007 (3)	−0.004 (3)
C5	0.083 (5)	0.097 (5)	0.073 (5)	−0.005 (5)	0.029 (4)	−0.014 (4)
C6	0.078 (5)	0.071 (4)	0.101 (5)	0.002 (5)	0.027 (4)	0.005 (5)
C7	0.066 (5)	0.094 (5)	0.081 (5)	0.019 (5)	0.005 (4)	0.020 (5)
C8	0.053 (4)	0.090 (5)	0.058 (4)	0.003 (3)	0.006 (3)	0.014 (3)

Geometric parameters (Å, º)

Cl1—C1	1.735 (5)	C4—C5	1.378 (7)
C1—C2	1.515 (6)	C4—H4	0.93
C1—H1A	0.97	C5—C6	1.349 (8)
C1—H1B	0.97	C5—H5	0.93
C2—O1	1.226 (6)	C6—C7	1.384 (9)
C2—N1	1.350 (6)	C6—H6	0.93
N1—C3	1.407 (6)	C7—C8	1.357 (7)
N1—H1N	0.86	C7—H7	0.93
C3—C8	1.392 (6)	C8—H8	0.93
C3—C4	1.401 (7)		
C2—C1—Cl1	112.8 (4)	C5—C4—C3	118.7 (6)
C2—C1—H1A	109	C5—C4—H4	120.7
Cl1—C1—H1A	109	C3—C4—H4	120.7
C2—C1—H1B	109	C6—C5—C4	122.0 (6)
Cl1—C1—H1B	109	C6—C5—H5	119
H1A—C1—H1B	107.8	C4—C5—H5	119
O1—C2—N1	124.3 (6)	C5—C6—C7	120.3 (6)
O1—C2—C1	123.7 (6)	C5—C6—H6	119.8
N1—C2—C1	112.0 (5)	C7—C6—H6	119.8
C2—N1—C3	128.5 (5)	C8—C7—C6	118.5 (6)
C2—N1—H1N	115.7	C8—C7—H7	120.8

C3—N1—H1N	115.7	C6—C7—H7	120.8
C8—C3—C4	117.8 (5)	C7—C8—C3	122.6 (6)
C8—C3—N1	119.2 (5)	C7—C8—H8	118.7
C4—C3—N1	123.0 (5)	C3—C8—H8	118.7
Cl1—C1—C2—O1	−4.8 (7)	N1—C3—C4—C5	−179.6 (5)
Cl1—C1—C2—N1	175.8 (4)	C3—C4—C5—C6	−2.8 (9)
O1—C2—N1—C3	−1.1 (9)	C4—C5—C6—C7	2.7 (9)
C1—C2—N1—C3	178.3 (5)	C5—C6—C7—C8	−2.0 (9)
C2—N1—C3—C8	−164.2 (5)	C6—C7—C8—C3	1.7 (8)
C2—N1—C3—C4	17.8 (8)	C4—C3—C8—C7	−1.8 (8)
C8—C3—C4—C5	2.3 (8)	N1—C3—C8—C7	−180.0 (5)

Hydrogen-bond geometry (Å, º)

D—H⋯*A*	*D*—H	H⋯*A*	*D*⋯*A*	*D*—H⋯*A*
N1—H1*N*⋯O1[i]	0.86	2.05	2.848 (5)	155

Symmetry code: (i) *x*−1/2, −*y*+1/2, *z*−1/2.