

Triclinic, $P\bar{1}$	$V = 565.5 (5) \text{ \AA}^3$
$a = 5.308 (3) \text{ \AA}$	$Z = 2$
$b = 7.709 (4) \text{ \AA}$	Mo $K\alpha$ radiation
$c = 14.109 (7) \text{ \AA}$	$\mu = 0.09 \text{ mm}^{-1}$
$\alpha = 96.911 (8)^\circ$	$T = 296 \text{ K}$
$\beta = 99.210 (8)^\circ$	$0.30 \times 0.20 \times 0.20 \text{ mm}$
$\gamma = 90.511 (9)^\circ$	

Crystal structure of 4-methoxy-N-phenylbenzamide

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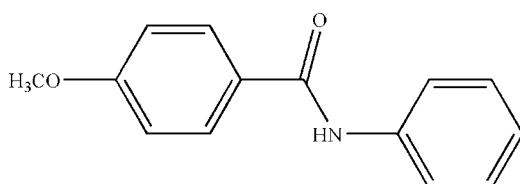
In the title molecule, $C_{14}H_{13}NO_2$, the dihedral angle between the planes of the benzene rings is $65.18 (4)^\circ$. The central amide group has about the same degree of twist with respect to both ring planes, as indicated by the dihedral angles of $34.70 (8)$ and $30.62 (8)^\circ$ between its plane and that of the phenyl and 4-methoxybenzene rings, respectively. The C atom of the methoxy group is close to being coplanar with its attached ring [deviation = $-0.112 (2) \text{ \AA}$]. In the crystal, molecules are linked by inter-amide N—H···O hydrogen bonds, which generate C(4) chains propagating in the [100] direction. Adjacent molecules in the chain are related by translational symmetry.

Keywords: crystal structure; amide; C(4) chain; hydrogen bonding.

CCDC reference: 1014108

1. Related literature

The background to this work has been described in earlier papers; see: Ren *et al.* (2010); Zhu *et al.* (2011). For related structures, see: Raza *et al.* (2010); Gowda *et al.* (2003).



2. Experimental

2.1. Crystal data

$C_{14}H_{13}NO_2$

$M_r = 227.25$

2.2. Data collection

Bruker SMART CCD diffractometer
Absorption correction: multi-scan (*SADABS*; Bruker, 2007)
 $T_{\min} = 0.974$, $T_{\max} = 0.982$

3188 measured reflections
2005 independent reflections
1605 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.013$

2.3. Refinement

$R[F^2 > 2\sigma(F^2)] = 0.036$
 $wR(F^2) = 0.097$
 $S = 1.03$
2005 reflections

156 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.16 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.13 \text{ e \AA}^{-3}$

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···O1 ⁱ	0.86	2.31	3.110 (2)	154

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

Data collection: *SMART* (Bruker, 2007); cell refinement: *SAINT* (Bruker, 2007); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL* and *PLATON* (Spek, 2009).

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Supporting information for this paper is available from the IUCr electronic archives (Reference: HB7253).

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supporting information

Acta Cryst. (2014). E70, o921 [doi:10.1107/S1600536814016420]

Crystal structure of 4-methoxy-N-phenylbenzamide

Zhijun Wang, Haiying Lei, Linhua Jin and Ruitao Zhu

S1. Experimental

To a 100 ml round flask fitted with a condenser was added aniline (0.93 g, 10 mmol), dichloromethane (15 ml) and triethylamine(0.5 ml) with magnetic stirring. 4-methoxybenzoyl chloride (1.70 g, 10 mmol) was added gradually. The reaction mixture was stirred at room temperature for 1 h and then refluxed for 2 h. The product precipitated as a white powder, which was washed three times with water and dichloromethane. Recrystallization from ethyl alcohol solution produced colourless prisms of the title compound.

S2. Refinement

H atoms were placed in idealized positions and allowed to ride on their respective parent atoms, with C—H = 0.93 Å, N—H = 0.86 Å and $U_{\text{iso}}(\text{H})=1.2U_{\text{eq}}(\text{C},\text{N})$.

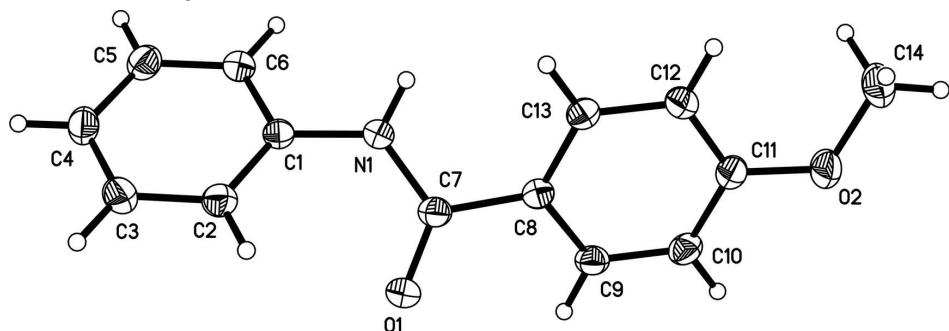
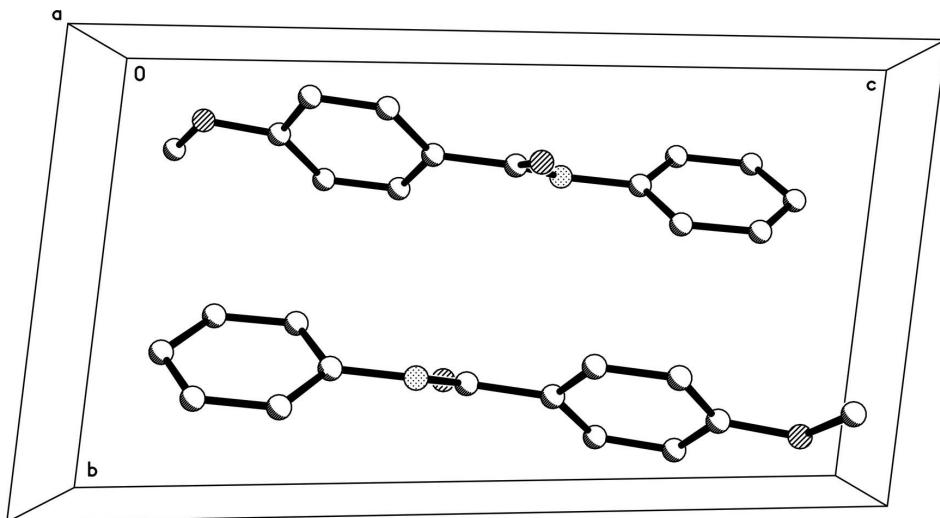


Figure 1

The molecular structure of the title compound with displacement ellipsoids drawn at the 30% probability level.

**Figure 2**

Part of the crystal structure of (I) with the donor-acceptor distances of hydrogen bonds drawn as dashed lines. H atoms are not shown.

4-Methoxy-N-phenylbenzamide

Crystal data

$C_{14}H_{13}NO_2$
 $M_r = 227.25$
Triclinic, $P\bar{1}$
Hall symbol: -P 1
 $a = 5.308 (3)$ Å
 $b = 7.709 (4)$ Å
 $c = 14.109 (7)$ Å
 $\alpha = 96.911 (8)^\circ$
 $\beta = 99.210 (8)^\circ$
 $\gamma = 90.511 (9)^\circ$

$V = 565.5 (5)$ Å³
 $Z = 2$
 $F(000) = 240$
 $D_x = 1.335$ Mg m⁻³
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
 $\theta = 1.5\text{--}25.1^\circ$
 $\mu = 0.09$ mm⁻¹
 $T = 296$ K
Prism, colorless
 $0.30 \times 0.20 \times 0.20$ mm

Data collection

Bruker SMART CCD
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 ω scans
Absorption correction: multi-scan
(SADABS; Bruker, 2007)
 $T_{\min} = 0.974$, $T_{\max} = 0.982$

3188 measured reflections
2005 independent reflections
1605 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.013$
 $\theta_{\max} = 25.1^\circ$, $\theta_{\min} = 1.5^\circ$
 $h = -5 \rightarrow 6$
 $k = -8 \rightarrow 9$
 $l = -16 \rightarrow 16$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.036$
 $wR(F^2) = 0.097$
 $S = 1.03$
2005 reflections
156 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

$$w = 1/[\sigma^2(F_o^2) + (0.045P)^2 + 0.1154P]$$

$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

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

$$\Delta\rho_{\max} = 0.16 \text{ e \AA}^{-3}$$

$$\Delta\rho_{\min} = -0.13 \text{ e \AA}^{-3}$$

Extinction correction: *SHELXL97* (Sheldrick, 2008), $F_c^* = kFc[1 + 0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$

Extinction coefficient: 0.037 (5)

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	$U_{\text{iso}}^*/U_{\text{eq}}$
N1	0.2321 (2)	0.27538 (17)	0.57341 (8)	0.0387 (3)
H1	0.0748	0.2753	0.5465	0.046*
O1	0.63875 (19)	0.25639 (17)	0.54382 (8)	0.0542 (4)
O2	0.0921 (2)	0.14685 (16)	0.11034 (7)	0.0520 (3)
C1	0.2818 (3)	0.29568 (19)	0.67586 (10)	0.0340 (3)
C2	0.4919 (3)	0.3893 (2)	0.72863 (11)	0.0427 (4)
H2	0.6085	0.4409	0.6971	0.051*
C3	0.5279 (3)	0.4060 (2)	0.82849 (12)	0.0504 (4)
H3	0.6693	0.4693	0.8638	0.060*
C4	0.3582 (3)	0.3307 (2)	0.87637 (11)	0.0483 (4)
H4	0.3849	0.3414	0.9436	0.058*
C5	0.1478 (3)	0.2391 (2)	0.82331 (11)	0.0468 (4)
H5	0.0306	0.1889	0.8551	0.056*
C6	0.1091 (3)	0.2210 (2)	0.72366 (11)	0.0401 (4)
H6	-0.0333	0.1585	0.6886	0.048*
C7	0.4093 (3)	0.2561 (2)	0.51365 (11)	0.0373 (4)
C8	0.3083 (3)	0.22995 (19)	0.40820 (10)	0.0345 (3)
C9	0.4522 (3)	0.1323 (2)	0.34765 (11)	0.0390 (4)
H9	0.6042	0.0850	0.3742	0.047*
C10	0.3727 (3)	0.1051 (2)	0.24959 (11)	0.0415 (4)
H10	0.4676	0.0365	0.2103	0.050*
C11	0.1516 (3)	0.1795 (2)	0.20900 (10)	0.0377 (4)
C12	0.0070 (3)	0.2788 (2)	0.26739 (11)	0.0395 (4)
H12	-0.1412	0.3297	0.2403	0.047*
C13	0.0852 (3)	0.3014 (2)	0.36659 (10)	0.0378 (4)
H13	-0.0139	0.3658	0.4060	0.045*
C14	-0.1445 (4)	0.2052 (3)	0.06486 (12)	0.0631 (5)
H14A	-0.2804	0.1623	0.0940	0.095*
H14B	-0.1707	0.1620	-0.0028	0.095*
H14C	-0.1425	0.3307	0.0726	0.095*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
N1	0.0288 (6)	0.0535 (8)	0.0333 (7)	0.0023 (5)	0.0034 (5)	0.0053 (6)
O1	0.0299 (6)	0.0892 (10)	0.0424 (7)	0.0060 (6)	0.0046 (5)	0.0050 (6)
O2	0.0563 (7)	0.0653 (8)	0.0327 (6)	0.0104 (6)	0.0067 (5)	0.0000 (5)
C1	0.0313 (7)	0.0364 (8)	0.0342 (8)	0.0065 (6)	0.0052 (6)	0.0035 (6)
C2	0.0360 (8)	0.0479 (10)	0.0434 (9)	-0.0027 (7)	0.0083 (7)	-0.0002 (7)
C3	0.0418 (9)	0.0583 (11)	0.0453 (10)	0.0005 (8)	0.0001 (7)	-0.0073 (8)
C4	0.0520 (10)	0.0587 (11)	0.0325 (8)	0.0115 (8)	0.0045 (7)	0.0016 (7)
C5	0.0471 (9)	0.0546 (11)	0.0418 (9)	0.0046 (8)	0.0140 (7)	0.0088 (8)
C6	0.0333 (8)	0.0477 (9)	0.0387 (8)	-0.0004 (7)	0.0055 (6)	0.0036 (7)
C7	0.0309 (8)	0.0416 (9)	0.0399 (8)	0.0030 (6)	0.0059 (6)	0.0066 (7)
C8	0.0314 (7)	0.0370 (8)	0.0358 (8)	-0.0006 (6)	0.0072 (6)	0.0051 (6)
C9	0.0313 (8)	0.0427 (9)	0.0437 (9)	0.0050 (6)	0.0072 (6)	0.0059 (7)
C10	0.0400 (8)	0.0446 (9)	0.0411 (9)	0.0060 (7)	0.0142 (7)	-0.0004 (7)
C11	0.0403 (8)	0.0390 (9)	0.0343 (8)	-0.0009 (7)	0.0087 (6)	0.0033 (6)
C12	0.0348 (8)	0.0451 (9)	0.0379 (8)	0.0062 (7)	0.0035 (6)	0.0052 (7)
C13	0.0357 (8)	0.0408 (9)	0.0372 (8)	0.0062 (6)	0.0094 (6)	0.0010 (7)
C14	0.0676 (12)	0.0819 (14)	0.0369 (9)	0.0150 (10)	0.0004 (8)	0.0059 (9)

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

N1—C7	1.3580 (19)	C6—H6	0.9300
N1—C1	1.4164 (19)	C7—C8	1.487 (2)
N1—H1	0.8600	C8—C13	1.385 (2)
O1—C7	1.2242 (18)	C8—C9	1.394 (2)
O2—C11	1.3689 (18)	C9—C10	1.370 (2)
O2—C14	1.419 (2)	C9—H9	0.9300
C1—C2	1.382 (2)	C10—C11	1.382 (2)
C1—C6	1.381 (2)	C10—H10	0.9300
C2—C3	1.381 (2)	C11—C12	1.384 (2)
C2—H2	0.9300	C12—C13	1.384 (2)
C3—C4	1.373 (2)	C12—H12	0.9300
C3—H3	0.9300	C13—H13	0.9300
C4—C5	1.378 (2)	C14—H14A	0.9600
C4—H4	0.9300	C14—H14B	0.9600
C5—C6	1.378 (2)	C14—H14C	0.9600
C5—H5	0.9300		
C7—N1—C1	126.16 (12)	C13—C8—C9	118.33 (14)
C7—N1—H1	116.9	C13—C8—C7	123.93 (13)
C1—N1—H1	116.9	C9—C8—C7	117.71 (13)
C11—O2—C14	118.22 (13)	C10—C9—C8	120.86 (14)
C2—C1—C6	119.58 (14)	C10—C9—H9	119.6
C2—C1—N1	122.48 (13)	C8—C9—H9	119.6
C6—C1—N1	117.92 (13)	C9—C10—C11	120.12 (14)
C1—C2—C3	119.65 (15)	C9—C10—H10	119.9

C1—C2—H2	120.2	C11—C10—H10	119.9
C3—C2—H2	120.2	O2—C11—C10	115.38 (13)
C4—C3—C2	121.03 (16)	O2—C11—C12	124.49 (14)
C4—C3—H3	119.5	C10—C11—C12	120.13 (14)
C2—C3—H3	119.5	C13—C12—C11	119.30 (14)
C3—C4—C5	119.01 (15)	C13—C12—H12	120.4
C3—C4—H4	120.5	C11—C12—H12	120.4
C5—C4—H4	120.5	C12—C13—C8	121.22 (14)
C4—C5—C6	120.70 (15)	C12—C13—H13	119.4
C4—C5—H5	119.6	C8—C13—H13	119.4
C6—C5—H5	119.6	O2—C14—H14A	109.5
C5—C6—C1	120.02 (15)	O2—C14—H14B	109.5
C5—C6—H6	120.0	H14A—C14—H14B	109.5
C1—C6—H6	120.0	O2—C14—H14C	109.5
O1—C7—N1	122.62 (14)	H14A—C14—H14C	109.5
O1—C7—C8	121.34 (13)	H14B—C14—H14C	109.5
N1—C7—C8	116.02 (13)		

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
N1—H1···O1 ⁱ	0.86	2.31	3.110 (2)	154

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