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Crystal structure of 2-methyl-3-nitrobenzoic anhydride

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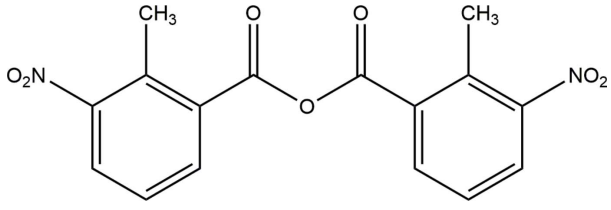
The title molecule, $C_{16}H_{12}N_2O_7$, lies on a twofold rotation axis which bisects the central O atom. The dihedral angle between two symmetry-related benzene rings is 48.54 (9)°. In the crystal, molecules are linked by weak C—H···O hydrogen bonds which generate $C(13)$ chains running parallel to [31$\bar{1}$].

Keywords: crystal structure; benzoic acid derivative; anhydrous compound; hydrogen bonding.

CCDC reference: 1404417

1. Related literature

For related structures, see: Schmitt *et al.* (2011); Liu *et al.* (2009); Huelgas *et al.* (2006); Glówka *et al.* (1990). For hydrogen-bond details, see: Nardelli (1995).



2. Experimental

2.1. Crystal data

$C_{16}H_{12}N_2O_7$
M_r = 344.28
Monoclinic, $C2/c$
a = 10.6332 (5) Å
b = 11.6961 (4) Å
c = 12.7934 (6) Å
β = 111.930 (6)°
V = 1475.95 (12) Å^3
Z = 4
Cu $K\alpha$ radiation
μ = 1.06 mm^{-1}
T = 123 K
0.45 × 0.40 × 0.16 mm

2.2. Data collection

Oxford Diffraction Gemini S diffractometer
Absorption correction: multi-scan (*CrysAlis PRO*; Oxford Diffraction, 2010)
T_{min} = 0.550, T_{max} = 1.000
5209 measured reflections
1466 independent reflections
1384 reflections with $I > 2\sigma(I)$
R_{int} = 0.101

2.3. Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.059
$wR(F^2)$ = 0.163
S = 1.08
1466 reflections
115 parameters
H-atom parameters constrained
$\Delta\rho_{max}$ = 0.40 e Å^{-3}
$\Delta\rho_{min}$ = −0.33 e Å^{-3}

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

D—H···A	D—H	H···A	D···A	D—H···A
C3—H3···O1^{i}	0.95	2.52	3.204 (2)	129

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

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: *SHELXL2014* (Sheldrick, 2015); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 2012) and *Mercury* (Macrae *et al.*, 2006); software used to prepare material for publication: *WinGX* (Farrugia, 2012).

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Supporting information for this paper is available from the IUCr electronic archives (Reference: LH5768).

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S1. Comment

In the synthesis of phenyl-benzamides performed in our research group for quite some time, the untimely production of 2-methyl-3-nitrobenzoic anhydride (I) as a product of the reaction system was given. A small excess in moles of 2-methyl-3-nitrobenzoic acid in the presence of thionyl chloride in the reaction and the subsequent addition of the o-nitro-aniline in dry acetonitrile, allowed the formation of two different types of crystals: the corresponding amide and the 2-methyl-3-nitrobenzoic anhydride. The excess addition of 2-methyl-3-nitrobenzoic acid possibly yield the benzyl halide formation which subsequently reacts with another molecule of acid, forming the anhydride system in dry condition. A number of anhydrous compounds, from benzoic acid derivatives are reported in the literature. Some with halogen substituents on the rings, encounter a widespread use as chelate ligands in coordination chemistry (Schmitt *et al.*, 2011). Similar compounds to (I) have been reported in the literature: N-phenylanthranilic anhydride (II) (Liu *et al.*, 2009), o-nitrobenzoic acid anhydride (III) (Huelgas *et al.*, 2006) and m-nitrobenzoic acid anhydride (IV) (Glówka *et al.*, 1990). The molecular structure of (I) is shown in Fig. 1. The central anhydride moiety C6-C8(═O3)-O4 shows a C8^{i}-O4-C8-O3 torsion angle (symmetry code: (i) -x, y, -z+3/2) of 25.06 (14)°. The twofold rotation axis passes through atom O4. Bond lengths and bond angles in the molecule are in a good agreement with those found in the related compounds (II), (III) and (IV), with the exception of the C6—C8 bond length. The title structure exhibits strong elongation in the C6—C8 bond length [1.494 (2) Å] if compared to the similar bond length presented in (III) [1.402 (2) Å]. In the crystal structure (Fig. 2), molecules are linked by weak C—H···O hydrogen bonds (see Table 1, Nardelli, 1995). The C3—H3 group in the molecule at (x,y,z) acts as hydrogen bond donor to O1 atom of the nitro group in the molecule at (-x-1/2,+y-1/2,-z+1/2). These interactions generate C(13) chains of molecules parallel to [3 1$\bar{1}$].

S2. Experimental

A mass of 0.380 g (1.104 mmol) of 2-methyl-3-nitrobenzoic acid was refluxed with 2 ml of thionyl chloride for one hour. Then 0.125 g (0.906 mmol) of 2-nitroaniline was added and dissolved in 10 ml of dry acetonitrile and it was placed under reflux and constant stirring for 3 hours. Subsequently, the final solvent was slowly evaporated to obtain colorless blocks of the title compound [m.p. 428 (1)K], and other yellow crystals of the amide show a melting point of 399 (1)K.

S3. Refinement

All H-atoms were located in difference Fourier maps and were positioned geometrically [C—H = 0.95 Å for aromatic, C—H= 0.98 Å for methyl] and were refined using a riding-model approximation with U_{iso}(H) constrained to 1.2 times U_{eq} of the respective parent atom or 1.2 times U_{eq}(C_{methyl}).

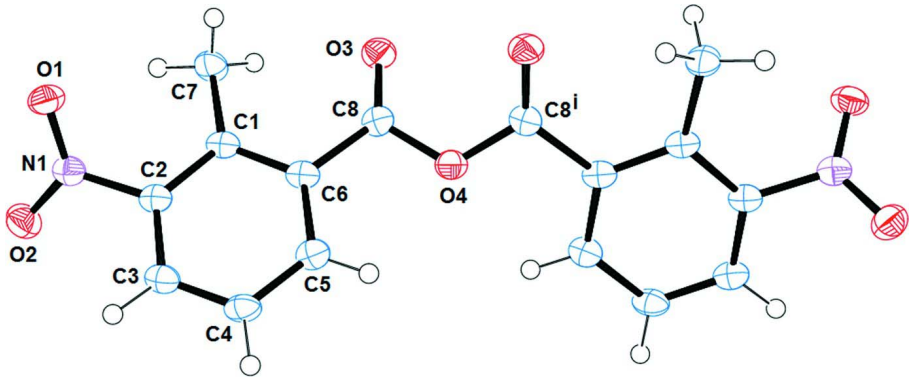

Figure 1

The molecular structure of (I) with displacement ellipsoids drawn at the 50% probability level. H atoms are shown as spheres of arbitrary radius (symmetry code: (i) -*x*, *y*, -*z* + 3/2).

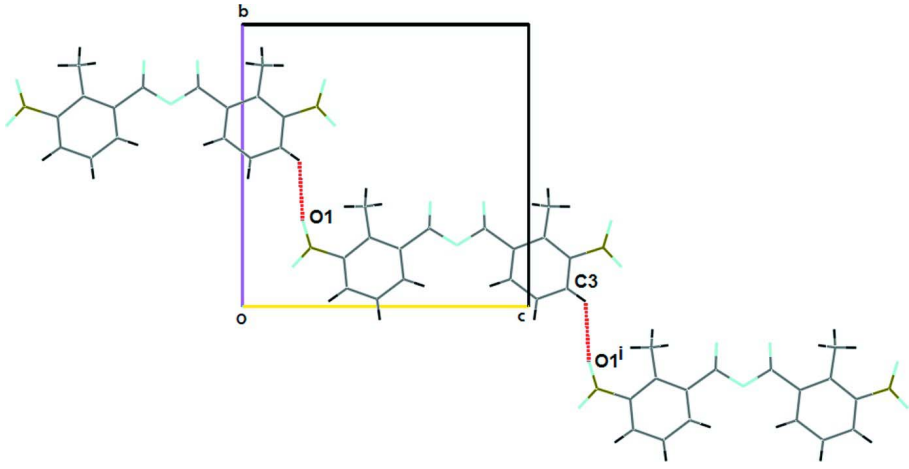

Figure 2

Part of the crystal structure of (I), showing the formation of a hydrogen-bonded *C*(13) chain parallel to [31$\bar{1}$] (symmetry code: (i) -*x* - 1/2 ,+*y* - 1/2, -*z* + 1/2).

2-Methyl-3-nitrobenzoic anhydride

Crystal data

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M_r = 344.28
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β = 111.930 (6)°
V = 1475.95 (12) Å^3
Z = 4
F(000) = 712

D_x = 1.549 Mg m^{-3}
Melting point: 428(1) K
Cu *Kα* radiation, *λ* = 1.54180 Å
Cell parameters from 5209 reflections
θ = 5.9–73.2°
μ = 1.06 mm^{-1}
T = 123 K
Block, colourless
0.45 × 0.40 × 0.16 mm

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Radiation source: fine-focus sealed tube
Graphite monochromator
ω scans
Absorption correction: multi-scan (*CrysAlis PRO*; Oxford Diffraction, 2010)
T_{min} = 0.550, T_{max} = 1.000
5209 measured reflections
1466 independent reflections
1384 reflections with $I > 2\sigma(I)$
R_{int} = 0.101
θ_{max} = 73.2°, θ_{min} = 5.9°
h = −12→13
k = −14→14
l = −14→15

Refinement

Refinement on F^2
Least-squares matrix: full
$R[F^2 > 2\sigma(F^2)]$ = 0.059
$wR(F^2)$ = 0.163
S = 1.08
1466 reflections
115 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
$w = 1/[\sigma^2(F_o^2) + (0.0933P)^2 + 1.5216P]$ where $P = (F_o^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{max} < 0.001$
$\Delta\rho_{max}$ = 0.40 e Å^{-3}
$\Delta\rho_{min}$ = −0.33 e Å^{-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 $F^2 > \sigma(F^2)$ 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 (Å²)

	x	y	z	U_{iso}*/U_{eq}
O1	−0.22053 (15)	0.30472 (11)	0.21092 (12)	0.0323 (4)
O2	−0.15352 (15)	0.13946 (12)	0.17543 (12)	0.0348 (4)
O3	0.03327 (14)	0.37124 (12)	0.65687 (11)	0.0303 (4)
O4	0.0000	0.21684 (16)	0.7500	0.0270 (5)
N1	−0.17462 (15)	0.20860 (13)	0.23928 (13)	0.0247 (4)
C1	−0.08126 (16)	0.24457 (15)	0.44518 (15)	0.0214 (4)
C2	−0.14895 (17)	0.17107 (15)	0.35513 (15)	0.0223 (4)
C3	−0.19730 (18)	0.06382 (15)	0.36540 (16)	0.0254 (4)
H3	−0.2393	0.0172	0.3009	0.031*
C4	−0.18356 (18)	0.02561 (16)	0.47094 (16)	0.0275 (4)
H4	−0.2150	−0.0482	0.4804	0.033*
C5	−0.12307 (18)	0.09647 (16)	0.56351 (15)	0.0252 (4)
H5	−0.1164	0.0716	0.6361	0.030*
C6	−0.07208 (17)	0.20339 (14)	0.55154 (15)	0.0219 (4)
C7	−0.0176 (2)	0.35437 (16)	0.42891 (16)	0.0284 (5)
H7A	−0.0092	0.3536	0.3552	0.043*

H7B	0.0724	0.3623	0.4883	0.043*
H7C	−0.0747	0.4189	0.4326	0.043*
C8	−0.00896 (18)	0.27684 (16)	0.65334 (15)	0.0231 (4)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0356 (8)	0.0225 (7)	0.0357 (8)	0.0027 (6)	0.0095 (6)	0.0041 (5)
O2	0.0382 (9)	0.0322 (8)	0.0336 (8)	0.0025 (6)	0.0130 (6)	−0.0055 (5)
O3	0.0288 (7)	0.0253 (7)	0.0315 (7)	−0.0084 (5)	0.0051 (6)	−0.0010 (5)
O4	0.0288 (10)	0.0222 (9)	0.0279 (9)	0.000	0.0082 (8)	0.000
N1	0.0188 (8)	0.0233 (8)	0.0299 (8)	−0.0014 (6)	0.0067 (6)	−0.0012 (6)
C1	0.0126 (7)	0.0194 (8)	0.0307 (9)	0.0006 (6)	0.0062 (6)	−0.0009 (7)
C2	0.0150 (8)	0.0207 (9)	0.0293 (9)	0.0012 (6)	0.0061 (7)	0.0002 (7)
C3	0.0170 (8)	0.0218 (9)	0.0327 (9)	−0.0007 (7)	0.0038 (7)	−0.0026 (7)
C4	0.0216 (8)	0.0199 (8)	0.0362 (10)	−0.0046 (7)	0.0052 (7)	0.0012 (7)
C5	0.0178 (8)	0.0247 (9)	0.0301 (9)	−0.0017 (7)	0.0054 (7)	0.0023 (7)
C6	0.0121 (7)	0.0202 (8)	0.0310 (10)	0.0002 (6)	0.0054 (7)	−0.0008 (6)
C7	0.0292 (10)	0.0253 (9)	0.0312 (9)	−0.0093 (7)	0.0118 (8)	−0.0030 (7)
C8	0.0146 (8)	0.0247 (9)	0.0272 (9)	0.0002 (7)	0.0047 (6)	0.0018 (7)

Geometric parameters (Å, º)

O1—N1	1.225 (2)	C3—C4	1.377 (3)
O2—N1	1.228 (2)	C3—H3	0.9500
O3—C8	1.186 (2)	C4—C5	1.391 (3)
O4—C8	1.3939 (19)	C4—H4	0.9500
O4—C8^{i}	1.3939 (19)	C5—C6	1.394 (2)
N1—C2	1.470 (2)	C5—H5	0.9500
C1—C2	1.402 (2)	C6—C8	1.494 (2)
C1—C6	1.412 (3)	C7—H7A	0.9800
C1—C7	1.502 (2)	C7—H7B	0.9800
C2—C3	1.380 (2)	C7—H7C	0.9800
C8—O4—C8^{i}	119.5 (2)	C4—C5—C6	121.08 (17)
O1—N1—O2	123.96 (16)	C4—C5—H5	119.5
O1—N1—C2	118.41 (15)	C6—C5—H5	119.5
O2—N1—C2	117.56 (15)	C5—C6—C1	121.47 (16)
C2—C1—C6	114.35 (16)	C5—C6—C8	119.12 (16)
C2—C1—C7	122.01 (16)	C1—C6—C8	119.39 (15)
C6—C1—C7	123.55 (15)	C1—C7—H7A	109.5
C3—C2—C1	125.05 (17)	C1—C7—H7B	109.5
C3—C2—N1	115.58 (15)	H7A—C7—H7B	109.5
C1—C2—N1	119.37 (15)	C1—C7—H7C	109.5
C4—C3—C2	118.77 (16)	H7A—C7—H7C	109.5
C4—C3—H3	120.6	H7B—C7—H7C	109.5
C2—C3—H3	120.6	O3—C8—O4	122.37 (16)
C3—C4—C5	119.17 (17)	O3—C8—C6	127.50 (16)

C3—C4—H4	120.4	O4—C8—C6	110.09 (15)
C5—C4—H4	120.4		
C6—C1—C2—C3	−3.8 (3)	C4—C5—C6—C1	0.8 (3)
C7—C1—C2—C3	172.93 (16)	C4—C5—C6—C8	179.52 (16)
C6—C1—C2—N1	175.09 (14)	C2—C1—C6—C5	2.1 (2)
C7—C1—C2—N1	−8.2 (2)	C7—C1—C6—C5	−174.58 (16)
O1—N1—C2—C3	132.42 (17)	C2—C1—C6—C8	−176.64 (14)
O2—N1—C2—C3	−44.8 (2)	C7—C1—C6—C8	6.7 (3)
O1—N1—C2—C1	−46.6 (2)	C8^{i}—O4—C8—O3	25.06 (14)
O2—N1—C2—C1	136.26 (17)	C8^{i}—O4—C8—C6	−157.20 (15)
C1—C2—C3—C4	2.4 (3)	C5—C6—C8—O3	−176.29 (18)
N1—C2—C3—C4	−176.46 (16)	C1—C6—C8—O3	2.5 (3)
C2—C3—C4—C5	0.8 (3)	C5—C6—C8—O4	6.1 (2)
C3—C4—C5—C6	−2.3 (3)	C1—C6—C8—O4	−175.15 (14)

Symmetry code: (i) −*x*, *y*, −*z*+3/2.

Hydrogen-bond geometry (Å, º)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*
C3—H3···O1^{ii}	0.95	2.52	3.204 (2)	129

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