1889 independent reflections 1245 reflections with $I > 2\sigma(I)$

3 standard reflections every 200 reflections

intensity decay: 2%

 $R_{\rm int} = 0.057$

mm

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Dimethyl 2-nitroterephthalate

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Key indicators: single-crystal X-ray study; T = 293 K; mean σ (C–C) = 0.006 Å; R factor = 0.078; wR factor = 0.201; data-to-parameter ratio = 12.1.

In the molecule of the title compound, C₁₀H₉NO₆, the two ester groups and the nitro group are inclined at 9.2 (2), 123.3 (6) and 135.2 $(5)^{\circ}$, respectively to the mean plane of the benzene ring. In the crystal structure, molecules are stacked along the *a* axis, without any $\pi - \pi$ interactions. The stacked columns are linked together by non-classical intermolecular interactions of the type $C-H \cdots O$

Related literature

For the use of the title compound in the preparation of 2amino-dimethyl-terephthalic acid, an intermediate for dyes, see: Niu et al. (2002). For related structures, see: Brisse & Pérez (1976); Huang & Liang (2007).



Experimental

Crystal data

$C_{10}H_9NO_6$	V = 1058.9 (4) Å ³
$M_r = 239.18$	Z = 4
Monoclinic, $P2_1/n$	Mo $K\alpha$ radiation
a = 6.9080 (14) Å	$\mu = 0.13 \text{ mm}^{-1}$
b = 12.662 (3) Å	T = 293 (2) K
c = 12.231 (2) Å	$0.30 \times 0.30 \times 0.10$
$\beta = 98.18 \ (3)^{\circ}$	

Data collection

Enrai–Nonius CAD-4
diffractometer
Absorption correction: ψ scan
(CAD-4 Software;
Enraf–Nonius, 1989)
$T_{\min} = 0.963, \ T_{\max} = 0.987$
2052 measured reflections

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.078$ 156 parameters $wR(F^2) = 0.201$ H-atom parameters constrained S = 1.00 $\Delta \rho_{\rm max} = 0.29 \text{ e} \text{ Å}^ \Delta \rho_{\rm min} = -0.31$ e Å⁻³ 1889 reflections

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

$D - H \cdot \cdot \cdot A$	D-H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$C1 - H1B \cdots O2^{i}$	0.96	2.59	3.523 (7)	164
$C4-H4A\cdots O2^{ii}$	0.93	2.54	3.185 (5)	127

Data collection: CAD-4 Software (Enraf-Nonius, 1989); cell refinement: CAD-4 Software; data reduction: XCAD4 (Harms & Wocadlo, 1995); 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: SHELXL97.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: PV2115).

References

Brisse, F. & Pérez, S. (1976). Acta Cryst. B32, 2110-2115.

Enraf-Nonius (1989). CAD-4 Software. Enraf-Nonius, Delft, The Netherlands

Harms, K. & Wocadlo, S. (1995). XCAD4. University of Marburg, Germany. Huang, J.-Y. & Liang, H.-Z. (2007). Acta Cryst. E63, 03019-03020.

Niu, T. S., Niu, X. Y., Yang, G. S. & Hou, J. Q. (2002). Appl. Chem. Ind. 34, 176-177

Sheldrick, G. M. (2008). Acta Cryst. A64, 112-122.

supplementary materials

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Dimethyl 2-nitroterephthalate

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Comment

The title compound, (I), is useful as a raw material for the preparation of 2-amino-dimethyl-terephthalic acid, which is used as an important intermediate for dyes (Niu *et al.*, 2002). The structures of dimethyl terephthalate (Brisse & Pérez, 1976) and dimethyl 2,3-dihydroxyterephthalate (Huang & Liang, 2007) which are closely related to the title compound have already been reported. In this article, we report the crystal structure of (I). A view of the molecule of (I) is presented in Fig. 1. The bond lengths and angles are within expected ranges. The C1/O1/C2/O2, C10/O6/C9/O5 and O3/N/O4 planes form dihedral angles of 9.2 (2), 123.3 (6) and 135.2 (5)°, respectively, with the C3—C8 plane. In the crystal structure, the molecules are stacked along the *a* axis, without any π - π interactions. The stacked columns are linked together by non-classical intermolecular interactions of the type C—H···O (Table 1).

Experimental

A sample of commercial 2-nitro-dimethyl-terephthalic acid (Aldrich) was crystalized by slow evaporation of a solution in methanol.

Refinement

Positional parameters of all the H atoms bonded to C atoms were calculated geometrically and were allowed to ride on the C atoms to which they are bonded, with H—C(aryl) = 0.93 Å and $U_{iso}(H) = 1.2U_{eq}(C)$ or with H—C(methyl) = 0.96 Å and $U_{iso}(H) = 1.5U_{eq}(C)$.

Figures



Fig. 1. A view of the molecule of (I) with the atomic numbering scheme. Displacement ellipsoids were drawn at the 30% probability level.

Dimethyl 2-nitroterephthalate

Crystal data

C₁₀H₉NO₆ $M_r = 239.18$ Monoclinic, $P2_1/n$ Hall symbol: -P 2yn a = 6.9080 (14) Å b = 12.662 (3) Å c = 12.231 (2) Å $\beta = 98.18$ (3)° V = 1058.9 (4) Å³ Z = 4

Data collection

Enraf–Nonius CAD-4 diffractometer	$R_{\rm int} = 0.057$
Radiation source: fine-focus sealed tube	$\theta_{\text{max}} = 25.3^{\circ}$
Monochromator: graphite	$\theta_{\min} = 2.3^{\circ}$
T = 293(2) K	$h = -8 \rightarrow 8$
$\omega/2\theta$ scans	$k = 0 \rightarrow 15$
Absorption correction: ψ scan (CAD-4 Software; Enraf–Nonius,1989)	$l = 0 \rightarrow 14$
$T_{\min} = 0.963, \ T_{\max} = 0.987$	3 standard reflections
2052 measured reflections	every 200 reflections
1889 independent reflections	intensity decay: 2%
1245 reflections with $I > 2\sigma(I)$	

Refinement

Refinement on F^2	Hydrogen site location: inferred from neighbouring sites
Least-squares matrix: full	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.078$	$w = 1/[\sigma^2(F_o^2) + (0.05P)^2 + 3.5P]$ where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.201$	$(\Delta/\sigma)_{\rm max} < 0.001$
<i>S</i> = 1.01	$\Delta \rho_{max} = 0.29 \text{ e} \text{ Å}^{-3}$
1889 reflections	$\Delta \rho_{min} = -0.31 \text{ e } \text{\AA}^{-3}$
156 parameters	Extinction correction: SHELXL97 (Sheldrick, 2008)
Primary atom site location: structure-invariant direct methods	Extinction coefficient: 0.053 (8)

 $F_{000} = 496$

 $D_{\rm x} = 1.500 \text{ Mg m}^{-3}$ Mo *K* α radiation

Cell parameters from 25 reflections

 $\lambda = 0.71073 \text{ Å}$

 $\theta = 10 - 13^{\circ}$ $\mu = 0.13 \text{ mm}^{-1}$

T = 293 (2) K

Block, colourless

 $0.30 \times 0.30 \times 0.10 \text{ mm}$

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 (A^2)

	x	У	Z		$U_{\rm iso}$ */ $U_{\rm eq}$	
06	0.0321 (5)	0.4552 (3)	0.88	00 (3)	0.0576 (9)	
02	0.7697 (5)	0.0539 (2)	1.01	89 (3)	0.0634 (10)	
01	0.8957 (5)	0.2101 (3)	1.07	87 (3)	0.0584 (9)	
05	0.0234 (5)	0.3733 (3)	0.71	73 (3)	0.0633 (10)	
04	0.1497 (6)	0.0806 (3)	0.73	85 (3)	0.0755 (12)	
Ν	0.1105 (6)	0.1519 (3)	0.80	06 (3)	0.0545 (11)	
C3	0.5871 (6)	0.2093 (3)	0.96	98 (3)	0.0397 (10)	
03	-0.0550 (5)	0.1733 (3)	0.81	56 (4)	0.0806 (13)	
C2	0.7594 (7)	0.1483 (3)	1.024	48 (4)	0.0445 (11)	
C8	0.5802 (7)	0.3193 (3)	0.98	06 (4)	0.0489 (11)	
H8A	0.6838	0.3561	1.02	01	0.059*	
C6	0.2593 (6)	0.3208 (3)	0.87	12 (3)	0.0412 (10)	
C7	0.4154 (7)	0.3717 (3)	0.93	09 (4)	0.0516 (12)	
H7A	0.4098	0.4447	0.93	32	0.062*	
C9	0.0911 (7)	0.3822 (3)	0.812	20 (4)	0.0470 (11)	
C10	-0.1157 (8)	0.5271 (4)	0.83	35 (5)	0.0629 (14)	
H10A	-0.1305	0.5819	0.88	59	0.094*	
H10B	-0.2373	0.4901	0.81	55	0.094*	
H10C	-0.0790	0.5579	0.76	77	0.094*	
C5	0.2728 (6)	0.2107 (3)	0.86	13 (3)	0.0386 (10)	
C4	0.4337 (7)	0.1556 (3)	0.90	90 (4)	0.0465 (11)	
H4A	0.4398	0.0827	0.90	06	0.056*	
C1	1.0681 (8)	0.1571 (5)	1.132	24 (5)	0.0769 (17)	
H1A	1.1662	0.2086	1.15	83	0.115*	
H1B	1.1174	0.1106	1.08	08	0.115*	
H1C	1.0357	0.1170	1.19	39	0.115*	
Atomic disp	lacement parameters	(\mathring{A}^2)				
	<i>U</i> ¹¹	1/22	1,33	<i>u</i> ¹²	<i>u</i> ¹³	<i>L</i> ²³
06	0 073 (2)	0.0433 (18)	0.060.(2)	0.0163(17)	0.0214(17)	-0.0000 (16)
Ω^{2}	0.075(2)	0.0435(10) 0.0235(18)	0.000(2)	0.0105(17)	0.0214(17) 0.0280(10)	-0.0090(10)
04	0.073 (2)	0.0555 (10)	0.007 (5)	0.0070(1/)	0.0207 (17)	0.00/0(1/)

supplementary materials

01	0.062 (2)	0.0383 (18)	0.076 (2)	0.0080 (16)	0.0165 (18)	0.0019 (17)
05	0.065 (2)	0.075 (3)	0.049 (2)	0.0151 (19)	0.0045 (17)	-0.0057 (18)
04	0.099 (3)	0.048 (2)	0.081 (3)	-0.005 (2)	0.017 (2)	-0.028 (2)
Ν	0.063 (3)	0.042 (2)	0.061 (3)	-0.017 (2)	0.019 (2)	-0.001 (2)
C3	0.051 (3)	0.029 (2)	0.044 (2)	0.0019 (19)	0.0248 (19)	0.0009 (18)
O3	0.052 (2)	0.083 (3)	0.113 (3)	-0.017 (2)	0.036 (2)	-0.010 (3)
C2	0.054 (3)	0.032 (2)	0.053 (3)	0.001 (2)	0.025 (2)	0.004 (2)
C8	0.064 (3)	0.027 (2)	0.058 (3)	-0.002 (2)	0.015 (2)	-0.005 (2)
C6	0.053 (3)	0.029 (2)	0.045 (2)	0.0025 (19)	0.020 (2)	-0.0016 (19)
C7	0.071 (3)	0.023 (2)	0.061 (3)	0.006 (2)	0.010 (2)	-0.004 (2)
C9	0.059 (3)	0.036 (2)	0.052 (3)	-0.002 (2)	0.025 (2)	-0.002 (2)
C10	0.068 (3)	0.052 (3)	0.071 (3)	0.018 (3)	0.021 (3)	0.002 (3)
C5	0.048 (2)	0.030 (2)	0.043 (2)	0.0010 (19)	0.0230 (19)	-0.0007 (18)
C4	0.059 (3)	0.023 (2)	0.065 (3)	-0.007 (2)	0.035 (2)	-0.005 (2)
C1	0.076 (4)	0.059 (4)	0.099 (5)	0.016 (3)	0.026 (3)	0.006 (3)

Geometric parameters (Å, °)

O6—C9	1.345 (5)	C8—H8A	0.9300
O6—C10	1.425 (6)	C6—C7	1.373 (6)
O2—C2	1.200 (5)	C6—C5	1.404 (6)
01—C2	1.325 (5)	C6—C9	1.497 (6)
O1—C1	1.442 (6)	C7—H7A	0.9300
О5—С9	1.191 (5)	C10—H10A	0.9600
O4—N	1.234 (5)	C10—H10B	0.9600
N—O3	1.214 (5)	C10—H10C	0.9600
N—C5	1.458 (6)	C5—C4	1.371 (6)
C3—C4	1.383 (6)	C4—H4A	0.9300
C3—C8	1.401 (6)	C1—H1A	0.9600
C3—C2	1.496 (6)	C1—H1B	0.9600
C8—C7	1.381 (6)	C1—H1C	0.9600
C9—O6—C10	117.2 (4)	O5—C9—C6	126.3 (4)
C2-01-C1	115.7 (4)	O6—C9—C6	109.9 (4)
O3—N—O4	123.5 (4)	O6-C10-H10A	109.5
O3—N—C5	118.7 (4)	O6-C10-H10B	109.5
O4—N—C5	117.8 (4)	H10A—C10—H10B	109.5
C4—C3—C8	120.4 (4)	O6-C10-H10C	109.5
C4—C3—C2	119.2 (4)	H10A—C10—H10C	109.5
C8—C3—C2	120.5 (4)	H10B-C10-H10C	109.5
O2—C2—O1	125.0 (4)	C4—C5—C6	121.9 (4)
O2—C2—C3	122.5 (4)	C4—C5—N	118.4 (4)
O1—C2—C3	112.5 (4)	C6—C5—N	119.7 (4)
C7—C8—C3	118.3 (4)	C5—C4—C3	119.4 (4)
С7—С8—Н8А	120.9	C5—C4—H4A	120.3
С3—С8—Н8А	120.9	C3—C4—H4A	120.3
C7—C6—C5	117.1 (4)	O1—C1—H1A	109.5
С7—С6—С9	120.7 (4)	O1—C1—H1B	109.5
С5—С6—С9	122.0 (4)	H1A—C1—H1B	109.5
С6—С7—С8	123.0 (4)	O1—C1—H1C	109.5

С6—С7—Н7А	118.5	H1A—C1—H1C	109.5
С8—С7—Н7А	118.5	H1B—C1—H1C	109.5
O5—C9—O6	123.7 (5)		
C1—O1—C2—O2	0.6 (7)	C7—C6—C9—O6	-48.0 (5)
C1—O1—C2—C3	-179.0 (4)	C5—C6—C9—O6	137.2 (4)
C4—C3—C2—O2	-0.5 (6)	C7—C6—C5—C4	0.1 (6)
C8—C3—C2—O2	178.9 (4)	C9—C6—C5—C4	175.1 (4)
C4—C3—C2—O1	179.1 (4)	C7—C6—C5—N	179.0 (4)
C8—C3—C2—O1	-1.4 (6)	C9—C6—C5—N	-6.1 (6)
C4—C3—C8—C7	1.3 (7)	O3—N—C5—C4	135.0 (5)
C2—C3—C8—C7	-178.2 (4)	O4—N—C5—C4	-43.2 (6)
C5—C6—C7—C8	-0.3 (7)	O3—N—C5—C6	-43.9 (6)
C9—C6—C7—C8	-175.3 (4)	O4—N—C5—C6	138.0 (4)
C3—C8—C7—C6	-0.4 (7)	C6—C5—C4—C3	0.7 (6)
C10—O6—C9—O5	-1.9 (7)	N—C5—C4—C3	-178.1 (4)
C10—O6—C9—C6	174.5 (4)	C8—C3—C4—C5	-1.5 (6)
C7—C6—C9—O5	128.2 (5)	C2—C3—C4—C5	178.0 (4)
C5—C6—C9—O5	-46.5 (7)		

Hydrogen-bond geometry (Å, °)

D—H···A	<i>D</i> —Н	H···A	$D \cdots A$	D—H···A
C1—H1B···O2 ⁱ	0.96	2.59	3.523 (7)	164
C4—H4A···O2 ⁱⁱ	0.93	2.54	3.185 (5)	127
Symmetry codes: (i) -x+2, -y, -z+2; (ii) -x+1, -y, -	- <i>z</i> +2.			

Fig. 1

