

Crystal structure of 2-(4-methylbenzylidene)malononitrile

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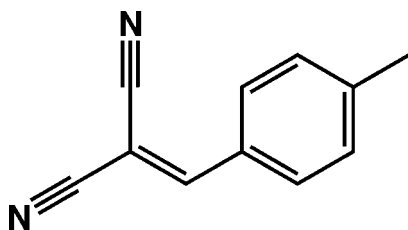
The molecule of the title compound, C₁₁H₈N₂, is approximately planar (r.m.s. deviation for all non-H atoms = 0.023 Å). The malononitrile C—C—C angle is 113.54 (13)°. In the crystal, molecules stack head-to-tail along [010]. There are no significant intermolecular interactions present.

Keywords: crystal structure; benzylidene; malononitrile; tyrphostins; benzylidenemalononitrile derivatives.

CCDC reference: 1033522

1. Related literature

For the pharmacological activity of benzylidenemalononitriles, see: Gazit *et al.* (1989); Levitzki & Mishani (2006). For the use of benzylidenemalononitrile derivatives in the preparation of heterocyclic compounds, see: Kolla & Lee (2011); Gao & Du (2012); Li *et al.* (2006).



2. Experimental

2.1. Crystal data

C ₁₁ H ₈ N ₂	$\gamma = 105.204 (4)^\circ$
$M_r = 168.19$	$V = 454.75 (5) \text{ \AA}^3$
Triclinic, $P\bar{1}$	$Z = 2$
$a = 7.0043 (5) \text{ \AA}$	Mo $K\alpha$ radiation
$b = 7.5270 (5) \text{ \AA}$	$\mu = 0.08 \text{ mm}^{-1}$
$c = 9.5396 (6) \text{ \AA}$	$T = 296 \text{ K}$
$\alpha = 106.757 (4)^\circ$	$0.40 \times 0.34 \times 0.30 \text{ mm}$
$\beta = 96.592 (4)^\circ$	

2.2. Data collection

Bruker X8 APEX diffractometer	7629 measured reflections
Absorption correction: multi-scan (SADABS; Bruker, 2009)	1923 independent reflections
$T_{\min} = 0.637$, $T_{\max} = 0.746$	1535 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.022$

2.3. Refinement

$R[F^2 > 2\sigma(F^2)] = 0.046$	118 parameters
$wR(F^2) = 0.132$	H-atom parameters constrained
$S = 1.06$	$\Delta\rho_{\text{max}} = 0.20 \text{ e \AA}^{-3}$
1923 reflections	$\Delta\rho_{\text{min}} = -0.18 \text{ e \AA}^{-3}$

Data collection: APEX2 (Bruker, 2009); cell refinement: SAINT (Bruker, 2009); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: ORTEP-3 for Windows (Farrugia, 2012); software used to prepare material for publication: PLATON (Spek, 2009) and publCIF (Westrip, 2010).

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Supporting information for this paper is available from the IUCr electronic archives (Reference: SU5017).

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S1. Comment

Pharmacological effects of benzylidenemalononitrile (BMN) compounds have been examined since the 1990's when several of their derivatives, referred to as tyrphostins, were recognized as specific inhibitors of epidermal growth factor tyrosine kinase (Gazit, *et al.*, 1989). Subsequent design and testing of a series of BMNs revealed new specific inhibitors of various protein tyrosine kinases (Levitzki & Mishani, 2006). It is well known that the benzylidenemalononitrile derivatives are very useful reagents for the preparation of heterocyclic compounds (Kolla & Lee, 2011; Gao & Du, 2012; Li *et al.*, 2006).

The title molecule is almost planar, Fig. 1, with an r.m.s. deviation = 0.023 Å; the maximum deviation of -0.037 (2) Å was observed for atom C5. The malononitrile angle C10–C9–C11 is 113.58 (12)°.

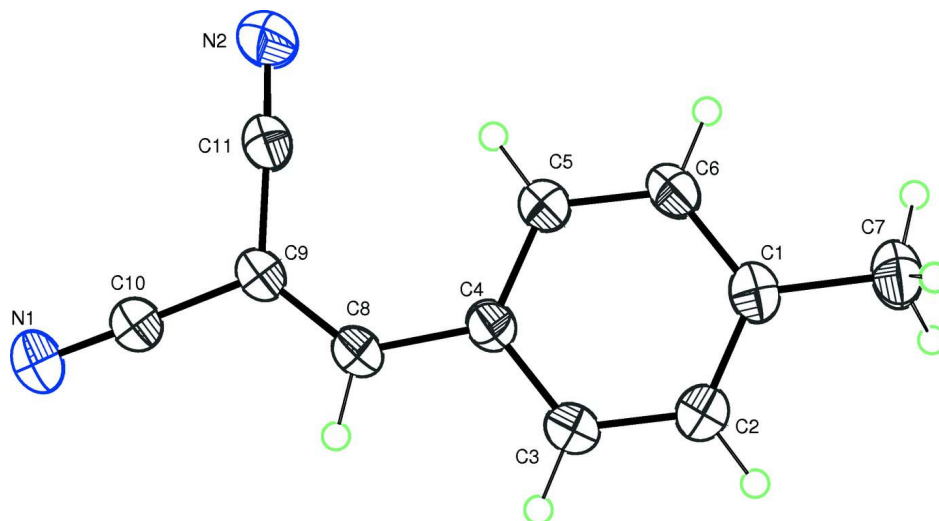
In the crystal, molecules stack head-to-tail along [010]. There are no significant intermolecular interactions present.

S2. Experimental

In a 250 ml round bottom flask, 4-methylbenzaldehyde (10 mmol), malononitrile (10 mmol) and phosphorus pentoxide (3.54 mmol) have stirred mechanically for ten minutes in 25 ml absolute ethanol. The resulting reaction mixture was heated at reflux using a water bath. The reaction mixture was poured onto crushed ice after the completion of the reaction monitored by TLC. On stirring separation of the desired product took place. The solid was filtered, washed with petroleum ether, dried and recrystallized from ethanol (yield: 68%, m.p.: 404 K), yielding block-like colourless crystals.

S3. Refinement

H atoms were located in a difference Fourier map and treated as riding: C–H = 0.93 - 0.96 Å with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{e}}(\text{C})$ for methyl H atoms and = $1.2U_{\text{eq}}(\text{C})$ for other H atoms.

**Figure 1**

The molecular structure of the title compound, with atom labelling. Displacement ellipsoids are drawn at the 50% probability level.

2-(4-Methylbenzylidene)malononitrile

Crystal data

$C_{11}H_8N_2$
 $M_r = 168.19$
 Triclinic, $P\bar{1}$
 $a = 7.0043$ (5) Å
 $b = 7.5270$ (5) Å
 $c = 9.5396$ (6) Å
 $\alpha = 106.757$ (4)°
 $\beta = 96.592$ (4)°
 $\gamma = 105.204$ (4)°
 $V = 454.75$ (5) Å³
 $Z = 2$

$F(000) = 176$
 $D_x = 1.229$ Mg m⁻³
 Melting point: 404 K
 Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
 Cell parameters from 1923 reflections
 $\theta = 3.0$ – 27.1 °
 $\mu = 0.08$ mm⁻¹
 $T = 296$ K
 Block, colourless
 $0.40 \times 0.34 \times 0.30$ mm

Data collection

Bruker X8 APEX
 diffractometer
 Radiation source: fine-focus sealed tube
 Graphite monochromator
 φ and ω scans
 Absorption correction: multi-scan
 (SADABS; Bruker, 2009)
 $T_{\min} = 0.637$, $T_{\max} = 0.746$

7629 measured reflections
 1923 independent reflections
 1535 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.022$
 $\theta_{\max} = 27.1$ °, $\theta_{\min} = 3.0$ °
 $h = -8 \rightarrow 8$
 $k = -9 \rightarrow 9$
 $l = -12 \rightarrow 12$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.046$
 $wR(F^2) = 0.132$
 $S = 1.06$
 1923 reflections
 118 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
 $w = 1/[\sigma^2(F_o^2) + (0.0567P)^2 + 0.1308P]$
 where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.20 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.18 \text{ e } \text{\AA}^{-3}$

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)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.3051 (2)	-0.0064 (2)	0.22452 (16)	0.0427 (4)
C2	0.1662 (3)	0.0403 (2)	0.30984 (17)	0.0488 (4)
H2	0.0285	-0.0170	0.2695	0.059*
C3	0.2306 (2)	0.1710 (2)	0.45410 (17)	0.0461 (4)
H3	0.1352	0.2004	0.5095	0.055*
C4	0.4357 (2)	0.25984 (19)	0.51829 (15)	0.0365 (3)
C5	0.5753 (2)	0.2128 (2)	0.43179 (17)	0.0448 (4)
H5	0.7131	0.2701	0.4714	0.054*
C6	0.5089 (2)	0.0816 (2)	0.28791 (17)	0.0476 (4)
H6	0.6036	0.0516	0.2321	0.057*
C7	0.2353 (3)	-0.1472 (3)	0.06642 (17)	0.0555 (4)
H7A	0.3430	-0.1963	0.0365	0.083*
H7B	0.1977	-0.0811	0.0006	0.083*
H7C	0.1209	-0.2539	0.0615	0.083*
C8	0.4888 (2)	0.3945 (2)	0.67091 (15)	0.0389 (3)
H8	0.3790	0.4133	0.7126	0.047*
C9	0.6697 (2)	0.4962 (2)	0.76167 (15)	0.0386 (3)
C10	0.6819 (2)	0.6205 (2)	0.91126 (16)	0.0435 (4)
C11	0.8637 (2)	0.4944 (2)	0.72682 (17)	0.0476 (4)
N1	0.6955 (2)	0.7184 (2)	1.03069 (15)	0.0601 (4)
N2	1.0205 (2)	0.4968 (3)	0.70532 (18)	0.0734 (5)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0507 (9)	0.0391 (7)	0.0347 (7)	0.0136 (7)	0.0069 (7)	0.0079 (6)
C2	0.0390 (9)	0.0520 (9)	0.0436 (8)	0.0077 (7)	0.0051 (7)	0.0059 (7)
C3	0.0398 (9)	0.0523 (9)	0.0417 (8)	0.0131 (7)	0.0140 (7)	0.0080 (7)
C4	0.0388 (8)	0.0379 (7)	0.0320 (7)	0.0119 (6)	0.0091 (6)	0.0098 (6)
C5	0.0381 (9)	0.0524 (9)	0.0377 (8)	0.0135 (7)	0.0090 (6)	0.0061 (6)
C6	0.0467 (10)	0.0570 (9)	0.0368 (8)	0.0210 (8)	0.0132 (7)	0.0060 (7)
C7	0.0619 (12)	0.0541 (9)	0.0381 (8)	0.0160 (8)	0.0038 (8)	0.0012 (7)

C8	0.0406 (8)	0.0428 (7)	0.0331 (7)	0.0139 (6)	0.0128 (6)	0.0096 (6)
C9	0.0419 (9)	0.0405 (7)	0.0326 (7)	0.0146 (6)	0.0110 (6)	0.0083 (6)
C10	0.0412 (9)	0.0464 (8)	0.0384 (8)	0.0118 (7)	0.0102 (7)	0.0084 (6)
C11	0.0442 (10)	0.0522 (9)	0.0367 (8)	0.0146 (7)	0.0066 (7)	0.0014 (6)
N1	0.0602 (10)	0.0649 (9)	0.0403 (7)	0.0147 (7)	0.0136 (7)	-0.0016 (6)
N2	0.0460 (9)	0.0935 (13)	0.0613 (10)	0.0225 (9)	0.0134 (7)	-0.0043 (8)

Geometric parameters (Å, °)

C1—C6	1.384 (2)	C6—H6	0.9300
C1—C2	1.389 (2)	C7—H7A	0.9600
C1—C7	1.508 (2)	C7—H7B	0.9600
C2—C3	1.382 (2)	C7—H7C	0.9600
C2—H2	0.9300	C8—C9	1.341 (2)
C3—C4	1.395 (2)	C8—H8	0.9300
C3—H3	0.9300	C9—C11	1.438 (2)
C4—C5	1.3998 (19)	C9—C10	1.4419 (18)
C4—C8	1.4535 (18)	C10—N1	1.1420 (19)
C5—C6	1.381 (2)	C11—N2	1.137 (2)
C5—H5	0.9300		
C6—C1—C2	118.20 (14)	C5—C6—H6	119.1
C6—C1—C7	121.04 (14)	C1—C6—H6	119.1
C2—C1—C7	120.75 (15)	C1—C7—H7A	109.5
C3—C2—C1	120.64 (15)	C1—C7—H7B	109.5
C3—C2—H2	119.7	H7A—C7—H7B	109.5
C1—C2—H2	119.7	C1—C7—H7C	109.5
C2—C3—C4	121.29 (14)	H7A—C7—H7C	109.5
C2—C3—H3	119.4	H7B—C7—H7C	109.5
C4—C3—H3	119.4	C9—C8—C4	130.79 (13)
C3—C4—C5	117.90 (13)	C9—C8—H8	114.6
C3—C4—C8	117.33 (12)	C4—C8—H8	114.6
C5—C4—C8	124.77 (14)	C8—C9—C11	126.43 (13)
C6—C5—C4	120.21 (15)	C8—C9—C10	120.02 (13)
C6—C5—H5	119.9	C11—C9—C10	113.54 (13)
C4—C5—H5	119.9	N1—C10—C9	178.50 (17)
C5—C6—C1	121.76 (14)	N2—C11—C9	177.20 (17)