

(E)-3-(4-Fluorophenyl)-1-(2-nitrophenyl)prop-2-en-1-one

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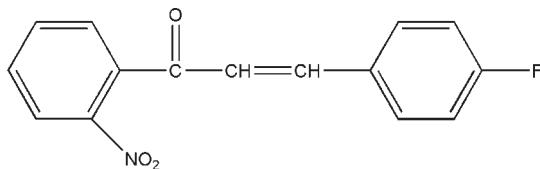
Received 2 May 2010; accepted 16 May 2010

Key indicators: single-crystal X-ray study; $T = 293\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.002\text{ \AA}$; R factor = 0.044; wR factor = 0.141; data-to-parameter ratio = 16.1.

The title compound, $\text{C}_{15}\text{H}_{10}\text{FNO}_3$, was prepared from 2-nitroacetophenone and 4-fluorobenzaldehyde by an Aldol condensation reaction. The dihedral angle formed by the two benzene rings is $67.37(2)^\circ$. The crystal structure is stabilized by weak intermolecular $\text{C}-\text{H}\cdots\text{O}$ and $\text{C}-\text{H}\cdots\text{F}$ hydrogen bonds.

Related literature

For the biological activities of chalcones, see: Hsieh *et al.* (1998); Anto *et al.* (1994); De Vincenzo *et al.* (2000); Dimmock *et al.* (1998). For related structures, see: Fun *et al.* (2008); Guo *et al.* (2009).



Experimental

Crystal data

$\text{C}_{15}\text{H}_{10}\text{FNO}_3$	$c = 9.759(2)\text{ \AA}$
$M_r = 271.24$	$\beta = 96.72(3)^\circ$
Monoclinic, $P2_1/n$	$V = 1285.6(5)\text{ \AA}^3$
$a = 7.7698(16)\text{ \AA}$	$Z = 4$
$b = 17.072(3)\text{ \AA}$	Mo $K\alpha$ radiation

 $\mu = 0.11\text{ mm}^{-1}$
 $T = 293\text{ K}$ $0.3 \times 0.3 \times 0.2\text{ mm}$

Data collection

Bruker SMART CCD
diffractometer
12293 measured reflections2921 independent reflections
2218 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.021$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.044$
 $wR(F^2) = 0.141$
 $S = 1.13$
2921 reflections181 parameters
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.24\text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.19\text{ e \AA}^{-3}$

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$H\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
C4—H4A \cdots O3 ⁱ	0.93	2.57	3.500 (2)	177
C5—H5A \cdots F1 ⁱⁱ	0.93	2.54	3.396 (2)	153
C9—H9A \cdots O3 ⁱⁱⁱ	0.93	2.57	3.411 (2)	150

Symmetry codes: (i) $x + 1, y, z$; (ii) $-x + 1, -y, -z + 2$; (iii) $-x, -y, -z + 1$.

Data collection: *SMART* (Bruker, 1997); cell refinement: *SAINT* (Bruker, 1997); 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*.

The authors would like to thank the National Natural Science Foundation of Shandong Province (2009ZRA07002).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: LH5041).

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