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## Key indicators

Single-crystal X-ray study
$T=293 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.003 \AA$
$R$ factor $=0.043$
$w R$ factor $=0.121$
Data-to-parameter ratio $=13.0$

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

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## 3-(2-Fluorophenyl)-2-phenylacrylic acid

The title compound, $\mathrm{C}_{15} \mathrm{H}_{11} \mathrm{FO}_{2}$, is a derivative of $\alpha$-phenylcinnamic acid. The two benzene rings in the molecule are approximately perpendicular to each other. $\mathrm{O}-\mathrm{H} \cdots \mathrm{O}$ hydrogen bonding occurs between molecules.

## Comment

Cinnamic acid derivatives are important building blocks in the production of lignins in higher plants (Mann, 1987). They are also key intermediates used in shikimic acid metabolic pathways of higher plants (Forgo et al., 2005). They are widely used as the starting materials for the synthesis of antimalarial drugs. The presence of the fluorine group shows an increase in the effectiveness of these compounds in prototype medicinals (Noddif et al., 1971). In these acids, the main structural feature is the strong hydrogen bonding between the carbonyl O atom of one molecule and H atom of another, which is responsible for the formation and stabilization of the dimer. In view of the importance of this class of compounds, the title compound, (I), has been synthesized, and its crystal structure is reported here.

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(I)

The molecular structure of (I) is shown in Fig. 1. Bond lengths and angles are within normal ranges (Allen et al., 1987). The bond lengths within the benzene rings range from 1.364 (4) A to 1.397 (3) A, typical of aromatic character. The $\mathrm{C} 2-\mathrm{C} 1-\mathrm{C} 6$ bond angle of $124.0(2)^{\circ}$ shows a small deviation from the ideal value of $120^{\circ}$. The bond length of 1.336 (2) $\AA$ shows $\mathrm{C} 7=\mathrm{C} 8$ to have double-bond character. Owing to the electronegative nature of fluorine, the $\mathrm{C} 1-\mathrm{F} 1$ bond length of 1.353 (2) $\AA$ is shorter than a normal single bond length (1.39 A).

Strong intermolecular hydrogen bonding is found between carboxyl groups in the crystal structure of (I) (Table 1).

## Experimental

The acid was synthesized according to a reported procedure (Noddif et al., 1971). A mixture of phenylacetic acid ( $4.08 \mathrm{~g}, 30 \mathrm{mmol}$ ), 2fluorobenzaldehyde $(3.13 \mathrm{ml}, 30 \mathrm{mmol})$, potassium carbonate $(2.346 \mathrm{~g}, 17 \mathrm{mmol})$ and acetic anhydride ( $7.07 \mathrm{ml}, 75 \mathrm{mmol}$ ) were
heated (see scheme). The temperature was slowly raised to 373 K and maintained for 24 h . Water ( 400 ml ) and hydrochloric acid ( 200 ml , $10 \%$ ) were added, and the reaction mixture was stirred at room temperature for 2 h and then filtered. The precipitate was washed with water $(100 \mathrm{ml} \times 3)$ to remove the impurities (yield $86 \%$ ). Colourless single crystals of (I) were obtained from an ethyl acetate solution.

## Crystal data

$\mathrm{C}_{15} \mathrm{H}_{11} \mathrm{FO}_{2}$
$M_{r}=242.24$
Monoclinic, $P 2_{\mathrm{k}} / c$
$a=9.1942(8) \AA$
$b=5.7823(5) \AA$
$c=23.126(2) \AA$
$\beta=96.398(2)^{\circ}$
$V=1221.78(18) \AA^{3}$

## Data collection

Bruker SMART CCD diffractometer
$\varphi$ and $\omega$ scans
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
$T_{\text {min }}=0.765, T_{\text {max }}=0.990$

$$
\begin{aligned}
& Z=4 \\
& D_{x}=1.317 \mathrm{Mg} \mathrm{~m}^{-3} \\
& \text { Mo } K \alpha \text { radiation } \\
& \mu=0.10 \mathrm{~mm}^{-1} \\
& T=293(2) \mathrm{K} \\
& \text { Rod, colourless } \\
& 0.30 \times 0.12 \times 0.10 \mathrm{~mm}
\end{aligned}
$$

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.043$
$w R\left(F^{2}\right)=0.121$
$S=1.04$
2181 reflections
168 parameters
H atoms treated by a mixture of independent and constrained refinement


## Figure 1

The molecular structure of (I) with $30 \%$ probability displacement ellipsoids (arbitrary spheres for H atoms).

Data collection: SMART (Bruker, 1998); cell refinement: SAINT (Bruker, 1999); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 1999); software used to prepare material for publication: SHELXTL.

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