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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 $=0.054$
$w R$ factor $=0.177$
Data-to-parameter ratio $=12.3$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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# $N$-(p-Tolyl)maleamic acid 

The title molecule, $\mathrm{C}_{11} \mathrm{H}_{11} \mathrm{NO}_{3}$, is nearly planar, with the mean planes through the $p$-tolyl and maleamic acid groups inclined at an angle of $5.45(3)^{\circ}$. The glide-related molecules are linked by $\mathrm{N}-\mathrm{H} \cdots \mathrm{O}$ and $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ intermolecular hydrogen bonds, forming infinite one-dimensional chains, which are assembled to form layers parallel to the $b c$ plane.

## Comment

The structure of $N$-(p-tolyl)maleamic acid, (I), was reported previously by Prasad \& Sinha (1978), using photographic X-ray diffraction data, with an $R$ value of 0.16 . The structure has now been refined using single-crystal X-ray diffraction data and the results are presented here.

(I)

The molecule of (I) is nearly planar, with C3 deviating by a maximum of 0.103 (3) $\AA$. The mean planes through the $p$-tolyl and maleamic acid groups are inclined at an angle of 5.45 (3) ${ }^{\circ}$ (Table 1). The bond lengths and angles in the maleamic acid group agree with those reported for the structure of maleic acid (James \& Williams, 1974). The carboxyl H atom is involved in an intramolecular $\mathrm{O}-\mathrm{H} \cdots \mathrm{O}$ hydrogen bond with carbonyl atom O3. In the solid state, the glide-related molecules are linked by $\mathrm{N}-\mathrm{H} \cdots \mathrm{O}$ and $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ hydrogen bonds (Table 2), forming infinite one-dimensional chains along the $c$-cell direction. These chains are arranged so as to form layers parallel to the $b c$ plane, approximately at $x=1 / 4$ and $3 / 4$. The mean interlayer separation of $3.406 \AA$ indicates significant interactions between the aryl and maleamic acid groups; the shortest interlayer contacts are $\mathrm{C} 6 \cdots \mathrm{C} 10(-x$, $1-y, 1-z$ ) of $3.319(5) \AA$ and $\mathrm{C} 4 \cdots \mathrm{C} 7(1-x, 1-y, 1-z)$ of 3.351 (4) A.

## Experimental

The title compound was prepared by a solid-state reaction between p-phenylaniline and maleic anhydride by Professor R. P. Rastogi (Gorakhpur University) and his co-workers (private communication) and was recrystallized from methanol at room temperature.


Figure 1
The molecular structure of the title compound, showing $50 \%$ probability displacement ellipsoids and the atom-numbering scheme.

Crystal data
$\mathrm{C}_{11} \mathrm{H}_{11} \mathrm{NO}_{3}$
$M_{r}=205.21$
Monoclinic, $P 2_{1} / c$
$a=6.769(2) \AA$
$b=12.109(1) \AA$
$c=12.606(1) \AA$
$\beta=95.73(1)^{\circ} \AA$
$V=1028.1(3) \AA^{3}$
$Z=4$
$D_{x}=1.326 \mathrm{Mg} \mathrm{m}^{-3}$
$\mathrm{Cu} K \alpha$ radiation
Cell parameters from 25 reflections
$\theta=28.2-36.3^{\circ}$
$\mu=0.81 \mathrm{~mm}^{-1}$
$T=293$ (2) K
Block, colourless
$0.38 \times 0.31 \times 0.25 \mathrm{~mm}$

## Data collection

| Enraf-Nonius CAD-4 | $R_{\text {int }}=0.029$ |
| :--- | :--- |
| diffractometer | $\theta_{\max }=75.5^{\circ}$ |
| $\omega-2 \theta$ scans | $h=0 \rightarrow 8$ |
| Absorption correction: $\psi$ scan | $k=0 \rightarrow 15$ |
| (North et al., 1968$)$ | $l=-15 \rightarrow 15$ |
| $T_{\text {min }}=0.733, T_{\text {max }}=0.817$ | 3 standard reflections |
| 2123 measured reflections | frequency: 60 min |
| 2047 independent reflections | intensity decay: none |

1200 reflections with $I$

## Refinement

Refinement on $F^{2}$

$$
\begin{aligned}
& w= 1 /\left[\sigma^{2}\left(F_{o}{ }^{2}\right)+(0.0899 P)^{2}\right. \\
&+0.2017 P] \\
& \text { where } P=\left(F_{o}{ }^{2}+2 F_{c}^{2}\right) / 3
\end{aligned}
$$

$w R\left(F^{2}\right)=0.178$
$S=1.04$
2047 reflections
167 parameters
H atoms treated by a mixture of independent and constrained refinement


Figure 2
A view of the molecular layer, close to $x=1 / 4$.

Table 2
Hydrogen-bonding geometry $\left(\AA^{\circ},{ }^{\circ}\right)$.

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| N-H1N $\cdots \mathrm{O} 2^{\mathrm{i}}$ | $0.93(3)$ | $1.96(3)$ | $2.885(3)$ | $177(2)$ |
| $\mathrm{O}^{\mathrm{H}}-\mathrm{H} 1 \mathrm{O} \cdots \mathrm{O} 3$ | 0.82 | 1.68 | $2.495(2)$ | 173 |
| C3-H3 $\cdots \mathrm{O} 3$ | $0.98(3)$ | $2.26(2)$ | $2.866(3)$ | $119(2)$ |
| C8-H8 $\mathrm{O}^{\mathrm{i}}$ |  | $0.97(3)$ | $2.55(2)$ | $3.498(3)$ |

Symmetry code: (i) $x, \frac{3}{2}-y, z-\frac{1}{2}$.
All H atoms apart from the methyl and hydroxyl H atoms were refined isotropically.

Data collection: CAD-4 EXPRESS (Enraf-Nonius, 1994); cell refinement: CAD-4 EXPRESS; data reduction: MolEN (Fair, 1990); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 (Farrugia, 1997); software used to prepare material for publication: SHELXL97.

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