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## Key indicators

Single-crystal X-ray study  
 $T = 223$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.003$  Å  
Disorder in main residue  
 $R$  factor = 0.058  
 $wR$  factor = 0.163  
Data-to-parameter ratio = 15.4For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.A rare *syn-anti* catemer in 4-nitrophenyl-  
propionic acidThe crystal structure of 4-nitrophenylpropionic acid,  $\text{C}_9\text{H}_5\text{NO}_4$ , is described. The carboxylic acid functional group adopts the rare *syn-anti* conformation and the molecules are arranged in a catemeric chain.

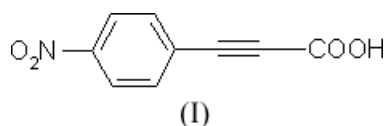
Received 11 February 2005

Accepted 27 April 2005

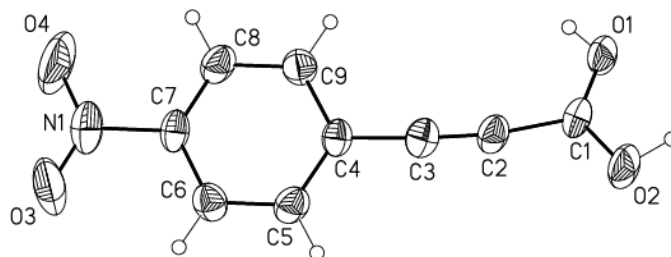
Online 7 May 2005

## Comment

Most carboxylic acids without other hydrogen-bonding substituents form centrosymmetric dimers in the solid state (Kolotuchin *et al.*, 1995). This functional group can, however, also be arranged as catemers (Leiserowitz, 1976). The catemer motif is uncommon. Even rarer are structures where the carboxylic acid group adopts both *syn* and *anti* conformations in the crystal structure, these conformations alternating along an infinite  $\text{O}-\text{H}\cdots\text{O}$  mediated array. This is referred to as the *syn-anti* conformation. In such a situation, the carboxyl groups may be either ordered or disordered (Kuduva *et al.*, 1999; Das *et al.*, 2003), but the general packing of the molecules is similar in the two cases. In a continuation of our previous work on 4-halosubstituted phenylpropionic acids (Das *et al.*, 2003), we studied the title compound, (I).



Molecule (I) crystallizes in space group  $P2_1/n$  with one molecule in the crystallographic asymmetric unit. The electron-withdrawing substituent in the *para* position of the benzene ring directs the structure towards the catemer because of the activation of the aromatic CH groups towards  $\text{C}-\text{H}\cdots\text{O}$  hydrogen-bond formation, which supports the formation of such a catemer. The molecular geometry of (I) is shown in Fig. 1. The carboxylic acid H atom is disordered between O1 and O2, with 50% occupancy at each site. Such



**Figure 1**  
An ORTEP view (Johnson, 1976) and the atom-numbering scheme for (I). Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as small spheres of arbitrary radii. Both components of the disordered carboxylic acid H atom are shown.

disorder was also observed in 4-chlorophenylpropionic acid (Desiraju *et al.*, 1990). The nitro group is coplanar with the benzene ring, but the carboxyl group deviates from this plane. Notable interactions are C9—H9 $\cdots$ O2, C6—H6 $\cdots$ O3 and C5—H5 $\cdots$ O4, as detailed in Table 1.

## Experimental

The title compound, (I), was prepared by standard procedures from the corresponding cinnamic acid. Single crystals for the X-ray diffraction study were obtained by slow evaporation of a solution in a 1:1 mixture of CHCl<sub>3</sub> and MeCN.

### Crystal data

C <sub>9</sub> H <sub>5</sub> NO <sub>4</sub>	$D_x = 1.514 \text{ Mg m}^{-3}$
$M_r = 191.14$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/n$	Cell parameters from 2334 reflections
$a = 3.726 (2) \text{ \AA}$	$\theta = 3.0\text{--}26.9^\circ$
$b = 7.511 (5) \text{ \AA}$	$\mu = 0.12 \text{ mm}^{-1}$
$c = 29.983 (18) \text{ \AA}$	$T = 223 (2) \text{ K}$
$\beta = 91.937 (12)^\circ$	Block, yellow
$V = 838.6 (9) \text{ \AA}^3$	$0.32 \times 0.21 \times 0.12 \text{ mm}$
$Z = 4$	

### Data collection

Siemens SMART CCD area-detector diffractometer	2065 independent reflections
$\varphi$ and $\omega$ scans	1200 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan (SADABS; Blessing, 1995)	$R_{\text{int}} = 0.055$
$T_{\text{min}} = 0.962$ , $T_{\text{max}} = 0.986$	$\theta_{\text{max}} = 28.4^\circ$
5168 measured reflections	$h = -4 \rightarrow 4$
	$k = -10 \rightarrow 9$
	$l = -21 \rightarrow 40$

### Refinement

Refinement on $F^2$	H atoms treated by a mixture of independent and constrained refinement
$R[F^2 > 2\sigma(F^2)] = 0.058$	$w = 1/[\sigma^2(F_o^2) + (0.0814P)^2]$
$wR(F^2) = 0.163$	where $P = (F_o^2 + 2F_c^2)/3$
$S = 1.03$	$(\Delta/\sigma)_{\text{max}} < 0.001$
2065 reflections	$\Delta\rho_{\text{max}} = 0.29 \text{ e \AA}^{-3}$
134 parameters	$\Delta\rho_{\text{min}} = -0.31 \text{ e \AA}^{-3}$

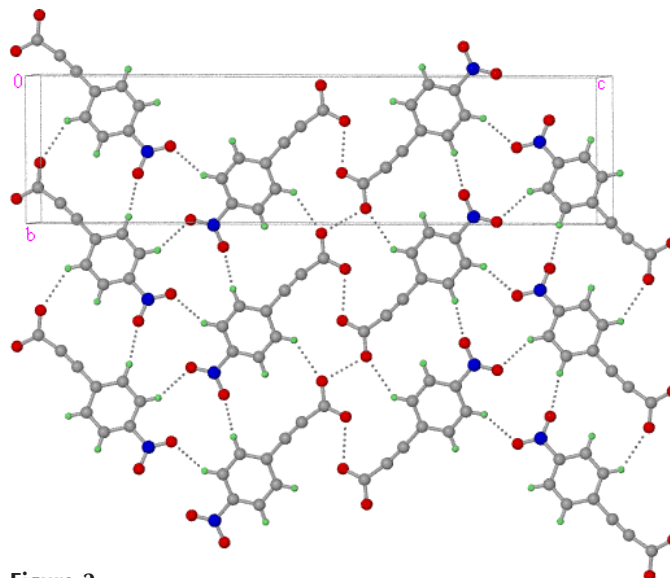
**Table 1**

Hydrogen-bonding geometry ( $\text{\AA}$ ,  $^\circ$ ).

$D\text{---}H\cdots A$	$D\text{---}H$	$H\cdots A$	$D\cdots A$	$D\text{---}H\cdots A$
O1—H1 $\cdots$ O1 <sup>i</sup>	0.75 (5)	1.90 (5)	2.635 (3)	168 (6)
O2—H2 $\cdots$ O2 <sup>ii</sup>	0.92 (6)	1.78 (6)	2.660 (3)	158 (6)
C5—H5 $\cdots$ O4 <sup>iii</sup>	0.94	2.56	3.345 (4)	142
C6—H6 $\cdots$ O3 <sup>iv</sup>	0.94	2.57	3.367 (4)	143
C9—H9 $\cdots$ O2 <sup>v</sup>	0.94	2.70	3.425 (3)	134

Symmetry codes: (i)  $-1 - x, -y, -z$ ; (ii)  $-1 - x, -1 - y, -z$ ; (iii)  $x, y - 1, z$ ; (iv)  $\frac{3}{2} - x, y - \frac{1}{2}, \frac{1}{2} - z$ ; (v)  $x, 1 + y, z$ .

All H atoms except carboxylic acid H atoms were positioned geometrically ( $C\text{---}H = 0.94 \text{ \AA}$ ) and refined with riding constraints and with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ . Carboxylic acid H-atom positions were



**Figure 2**

Packing diagram of (I). For clarity, carboxylic acid H atoms have been omitted. Hydrogen bonds are shown as dashed lines.

found in a difference Fourier map and refined with  $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$ .

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINTE* (Bruker, 1998); data reduction: *SAINTE*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 2000) and *X-SEED* (Barbour, 2001); software used to prepare material for publication: *SHELXTL*.

DD thanks the CSIR for fellowship support. RKRJ and RB are grateful to the Deutsche Forschungsgemeinschaft for financial assistance.

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