

Acta Crystallographica Section E **Structure Reports** Online

ISSN 1600-5368

4-Acetylpyridine–fumaric acid (2/1)

Kan Xu,^a Bing-Yu Zhang,^b Jing-Jing Nie^b and Duan-Jun Xu^b*

^aDepartment of Orthopaedics, Second Affiliated Hospital, School of Medicine, Zhejiang University, People's Republic of China, and ^bDepartment of Chemistry, Zhejiang University, People's Republic of China Correspondence e-mail: xudj@mail.hz.zj.cn

Received 26 May 2009; accepted 29 May 2009

Key indicators: single-crystal X-ray study; T = 294 K; mean σ (C–C) = 0.004 Å; R factor = 0.040; wR factor = 0.143; data-to-parameter ratio = 12.8.

In the crystal structure of the title cocrystal, 2C₇H₇NO-- $C_4H_4O_4$, the complete fumaric acid molecule is generated by a crystallographic inversion centre. The two components of the cocrystal are linked by an O−H···N hydrogen bond.

Related literature

For biological and medicinal applications of 4-acetylpyridine and fumaric acid, see: Fidler et al. (2003); Thomas et al. (2007). For molecular complexes of neutral pyridine derivatives and neutral fumaric acid, see: Bowes et al. (2003); Aakeroy et al. (2002, 2006, 2007); Haynes et al. (2006); Bu et al. (2007). For literature on C-O bond distances in fumaric acid, see: Liu et al. (2003). For metal complexes of 4-acetylpyridine, see: Steffen & Palenik (1977); Pang et al. (1994). For a 4-acetylpyridinium salt, see: Kochel (2005).



Experimental

Crystal data

2C7H7NO·C4H4O4 $M_r = 358.34$ Triclinic $P\overline{1}$ a = 3.9062 (5) Åb = 8.6809 (13) Å c = 13.0909 (18) Å $\alpha = 87.925 \ (4)^{\circ}$ $\beta = 89.941 \ (3)^{\circ}$

$\gamma = 83.141 \ (4)^{\circ}$
$V = 440.44 (11) \text{ Å}^3$
Z = 1
Mo $K\alpha$ radiation
$\mu = 0.10 \text{ mm}^{-1}$
T = 294 K
$0.30 \times 0.11 \times 0.08 \text{ mm}$

organic compounds

1589 independent reflections

798 reflections with $I > 2\sigma(I)$

 $R_{\rm int} = 0.030$

Data collection

Rigaku R-AXIS RAPID IP diffractometer Absorption correction: none 3600 measured reflections

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.040$	H atoms treated by a mixture of
$wR(F^2) = 0.143$	independent and constrained
S = 1.18	refinement
1589 reflections	$\Delta \rho_{\rm max} = 0.19 \ {\rm e} \ {\rm \AA}^{-3}$
124 parameters	$\Delta \rho_{\rm min} = -0.20 \ {\rm e} \ {\rm \AA}^{-3}$

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	D-H	$H \cdots A$	$D \cdots A$	$D - H \cdots A$
$D3-H3A\cdots N1$	0.98 (4)	1.64 (4)	2.599 (3)	166 (4)

Data collection: PROCESS-AUTO (Rigaku, 1998); cell refinement: PROCESS-AUTO; data reduction: CrystalStructure (Rigaku/ MSC, 2002); program(s) used to solve structure: SIR92 (Altomare et al., 1993); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: ORTEP-3 for Windows (Farrugia, 1997); software used to prepare material for publication: WinGX (Farrugia, 1999).

The work was supported by the Scientific Foundation of the Department of Education of Zhejiang Province, China (grant No. Y200700867).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: NG2588).

References

- Aakeroy, C. B., Beatty, A. M. & Helfrich, B. A. (2002). J. Am. Chem. Soc. 124, 14425-14432.
- Aakeroy, C. B., Hussain, I. & Desper, J. (2006). Cryst. Growth Des. 6, 474-480. Aakeroy, C. B., Hussain, I., Forbes, S. & Desper, J. (2007). CrystEngComm, 9, 46-54
- Altomare, A., Cascarano, G., Giacovazzo, C. & Guagliardi, A. (1993). J. Appl. Cryst. 26, 343-350.
- Bowes, K. F., Ferguson, G., Lough, A. J. & Glidewell, C. (2003). Acta Cryst. B59, 100-117.
- Bu, T.-J., Li, B. & Wu, L.-X. (2007). Acta Cryst. E63, 03466.
- Farrugia, L. J. (1997). J. Appl. Cryst. 30, 565.
- Farrugia, L. J. (1999). J. Appl. Cryst. 32, 837-838.
- Fidler, M. C., Davidsson, L., Zeder, C., Walczyk, T. & Hurrell, R. F. (2003). Br. J. Nutr. 90, 1081-1085.
- Haynes, D. A., Jones, W. & Motherwell, W. D. S. (2006). CrystEngComm, 8, 830-840
- Kochel, A. (2005). Acta Cryst. E61, 0926-0927.
- Liu, Y., Xu, D.-J. & Hung, C.-H. (2003). Acta Cryst. E59, m297-m299.
- Pang, L., Whitehead, M. A., Bermardinelli, G. & Lucken, E. A. C. (1994). J. Chem. Crystallogr. 24, 203-211.
- Rigaku (1998). PROCESS-AUTO. Rigaku Corporation, Tokyo, Japan.
- Rigaku/MSC (2002). CrystalStructure. Rigaku/MSC, The Woodlands, Texas, USA.
- Sheldrick, G. M. (2008). Acta Cryst. A64, 112-122.
- Steffen, W. L. & Palenik, G. L. (1977). Inorg. Chem. 16, 1119-1128.
- Thomas, J. S., Muharrem, A. K. & Ulrich, M. (2007). Bioorg. Med. Chem. 15, 333-342.

supporting information

Acta Cryst. (2009). E65, o1467 [doi:10.1107/S1600536809020480]

4-Acetylpyridine-fumaric acid (2/1)

Kan Xu, Bing-Yu Zhang, Jing-Jing Nie and Duan-Jun Xu

S1. Comment

The fumaric acid and acetylpyridine have been widely used in the biological and medicine fileds (Thomas *et al.* 2007; Fidler *et al.* 2003). In the medicine composition the carboxyl group of the fumaric acid is usually deprotonated while the pyridine derivatives are protonated. But some crystal structure determinations showed the neutral pyridine derivatives and fumaric acid in the crystal structures, *i.e.* the pyridine derivatives are not protonated while the fumaric acid is also not deprotonated in these crystal structures (Bowes *et al.* 2003; Aakeroy *et al.*, 2002, 2006, 2007; Haynes *et al.* 2006; Bu *et al.* 2007). Herein we report the crystal structure of the new compound containing pyridine derivative and fumaric acid components.

The crystal structure of the title compond consists of fumaric acid and 4-acetylpyridine molecules (Fig. 1). The planar fumaric acid molecule is centrosymmetric with the mid-point of the C=C double bond located at an inversion center. The C8—O2 bond distance of 1.204 (3) Å is much shorter than the C8—O3 bond distance of 1.297 (3) Å, it suggests that the carboxyl group is not deprotonated in the crystal structure (Liu *et al.* 2003).

The acetylpyridine molecule is not protonated in the crystal structure, which contrasts with that found in the crystal structure of the 4-acetylpyridinium chloride (Kochel, 2005). The geometry data of the acetylpyridine is consistent with those found in metal complexes of acetylpyridine (Steffen & Palenik, 1977; Pang *et al.*, 1994). The planar acetylpyridine molecule is twisted to the fumaric acid with a dihedral angle of 25.97 (11)° in the crystal structure.

The intermolecular classic O—H···N hydrogen bonding and weak C—H···O hydrogen bonding help to stabilize the crystal structure (Table 1).

S2. Experimental

Reagents and solvent were used as purchased without further purification. 4-Acetylpyridine (2 mmol) and fumaric acid (1 mmol) were dissolved in water–ethanol (6 ml, 1:5) at room temperature. The single crystals were obtained from the solution after 3 d.

S3. Refinement

The carboxyl H atom was located in a difference Fourier map and refined isotropically. Methyl H atoms were placed in calculated positions with C—H = 0.96 Å and the torsion angle was refined to fit the electron density, $U_{iso}(H) = 1.5U_{eq}(C)$. Other H atoms were placed in calculated positions with C—H = 0.93 Å and refined in riding mode with $U_{iso}(H) = 1.2U_{eq}(C)$.



Figure 1

The molecular structure of the title compound with 40% probability displacement (arbitrary spheres for H atoms). Dashed lines indicate hydrogen bonding.

Z = 1

F(000) = 188 $D_{\rm x} = 1.351 \text{ Mg m}^{-3}$

 $\theta = 3.2-24.6^{\circ}$ $\mu = 0.10 \text{ mm}^{-1}$ T = 294 K

Needle, colourless $0.30 \times 0.11 \times 0.08 \text{ mm}$

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å Cell parameters from 2308 reflections

4-Acetylpyridine-fumaric acid (2/1)

Crystal data
$2C_7H_7NO\cdot C_4H_4O_4$
$M_r = 358.34$
Triclinic, $P\overline{1}$
Hall symbol: -P 1
a = 3.9062 (5) Å
b = 8.6809 (13) Å
c = 13.0909 (18) Å
$\alpha = 87.925 \ (4)^{\circ}$
$\beta = 89.941 \ (3)^{\circ}$
$\gamma = 83.141 \ (4)^{\circ}$
$V = 440.44 (11) \text{ Å}^3$
Data collection

Rigaku R-AXIS RAPID IP	798 reflections with $I > 2\sigma(I)$
diffractometer	$R_{\rm int} = 0.030$
Radiation source: fine-focus sealed tube	$\theta_{\rm max} = 25.2^{\circ}, \ \theta_{\rm min} = 3.1^{\circ}$
Graphite monochromator	$h = -4 \rightarrow 4$
ω scans	$k = -10 \rightarrow 10$
3600 measured reflections	$l = -15 \rightarrow 15$
1589 independent reflections	

Refinement

Refinement on F^2	Hydrogen site location: inferred from
Least-squares matrix: full	neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.040$	H atoms treated by a mixture of independent
$wR(F^2) = 0.143$	and constrained refinement
S = 1.18	$w = 1/[\sigma^2(F_o^2) + (0.0527P)^2 + 0.048P]$
1589 reflections	where $P = (F_o^2 + 2F_c^2)/3$
124 parameters	$(\Delta/\sigma)_{\rm max} = 0.001$
0 restraints	$\Delta \rho_{\rm max} = 0.19 \text{ e} \text{ Å}^{-3}$
Primary atom site location: structure-invariant	$\Delta ho_{ m min} = -0.20 \ m e \ { m \AA}^{-3}$
direct methods	Extinction correction: SHELXL97 (Sheldrick,
Secondary atom site location: difference Fourier	2008), $Fc^* = kFc[1+0.001xFc^2\lambda^3/sin(2\theta)]^{-1/4}$
map	Extinction coefficient: 0.032 (8)

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 (\hat{A}^2)

N1	0.4401.(6)			1	
111	0.4491 (6)	0.4588 (3)	0.69775 (18)	0.0630 (7)	
01	0.8832 (6)	0.7499 (3)	0.98158 (16)	0.0900 (8)	
O2	0.1660 (6)	0.1037 (2)	0.67034 (15)	0.0775 (7)	
03	0.2584 (6)	0.2870 (3)	0.55473 (16)	0.0764 (7)	
C1	0.5832 (8)	0.4178 (4)	0.7899 (2)	0.0733 (9)	
H1	0.6122	0.3130	0.8097	0.088*	
C2	0.6807 (7)	0.5230 (3)	0.8573 (2)	0.0651 (9)	
H2	0.7765	0.4894	0.9204	0.078*	
C3	0.6342 (6)	0.6786 (3)	0.82954 (19)	0.0494 (7)	
C4	0.4891 (6)	0.7228 (3)	0.73526 (19)	0.0546 (7)	
H4	0.4508	0.8270	0.7145	0.065*	
C5	0.4019 (7)	0.6088 (4)	0.6724 (2)	0.0612 (8)	
Н5	0.3048	0.6391	0.6089	0.073*	
C6	0.7453 (7)	0.7954 (3)	0.9018 (2)	0.0562 (8)	
C7	0.6902 (7)	0.9624 (3)	0.8712 (2)	0.0653 (9)	
H7A	0.7914	1.0212	0.9215	0.098*	
H7B	0.4474	0.9961	0.8664	0.098*	
H7C	0.7961	0.9784	0.8061	0.098*	
C8	0.1658 (7)	0.1535 (3)	0.5833 (2)	0.0539 (7)	
С9	0.0643 (7)	0.0651 (3)	0.4951 (2)	0.0561 (8)	
H9	0.0951	0.1057	0.4295	0.067*	
H3A	0.335 (9)	0.337 (5)	0.615 (3)	0.131 (14)*	

supporting information

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
N1	0.0771 (16)	0.0500 (17)	0.0641 (16)	-0.0142 (12)	-0.0009 (13)	-0.0083 (13)
01	0.1244 (18)	0.0731 (17)	0.0703 (14)	-0.0004 (13)	-0.0407 (14)	-0.0077 (12)
O2	0.1167 (18)	0.0640 (15)	0.0548 (13)	-0.0234 (12)	-0.0138 (12)	0.0004 (11)
03	0.1198 (18)	0.0541 (15)	0.0607 (13)	-0.0313 (13)	-0.0048 (12)	-0.0076 (11)
C1	0.096 (2)	0.046 (2)	0.077 (2)	-0.0096 (17)	-0.0064 (19)	0.0019 (17)
C2	0.082 (2)	0.052 (2)	0.0610 (19)	-0.0053 (15)	-0.0123 (16)	0.0030 (15)
C3	0.0523 (15)	0.0448 (18)	0.0513 (16)	-0.0058 (12)	-0.0004 (13)	-0.0030 (13)
C4	0.0682 (17)	0.0451 (17)	0.0512 (16)	-0.0100 (13)	-0.0046 (14)	-0.0014 (13)
C5	0.0704 (18)	0.057 (2)	0.0566 (18)	-0.0100 (15)	-0.0086 (15)	-0.0024 (15)
C6	0.0577 (16)	0.057 (2)	0.0541 (18)	-0.0048 (13)	-0.0047 (14)	-0.0051 (14)
C7	0.0738 (19)	0.054 (2)	0.070 (2)	-0.0163 (15)	-0.0083 (16)	-0.0077 (16)
C8	0.0604 (16)	0.0446 (18)	0.0567 (18)	-0.0048 (13)	-0.0084 (14)	-0.0063 (14)
C9	0.0680 (17)	0.0469 (18)	0.0534 (16)	-0.0063 (13)	-0.0072 (14)	-0.0016 (14)

Atomic displacement parameters $(Å^2)$

Geometric parameters (Å, °)

N1—C5	1.323 (4)	C3—C6	1.512 (4)
N1—C1	1.335 (4)	C4—C5	1.383 (4)
O1—C6	1.207 (3)	C4—H4	0.9300
O2—C8	1.204 (3)	С5—Н5	0.9300
O3—C8	1.297 (3)	C6—C7	1.481 (4)
O3—H3A	0.97 (4)	C7—H7A	0.9600
C1—C2	1.378 (4)	С7—Н7В	0.9600
C1—H1	0.9300	С7—Н7С	0.9600
C2—C3	1.377 (4)	C8—C9	1.488 (4)
C2—H2	0.9300	C9—C9 ⁱ	1.293 (5)
C3—C4	1.381 (3)	С9—Н9	0.9300
C5—N1—C1	117.3 (3)	C4—C5—H5	118.2
С8—О3—НЗА	108 (2)	O1—C6—C7	121.8 (3)
N1—C1—C2	123.2 (3)	O1—C6—C3	119.3 (3)
N1-C1-H1	118.4	C7—C6—C3	118.9 (2)
C2—C1—H1	118.4	С6—С7—Н7А	109.5
C3—C2—C1	118.9 (3)	С6—С7—Н7В	109.5
С3—С2—Н2	120.5	H7A—C7—H7B	109.5
C1—C2—H2	120.5	С6—С7—Н7С	109.5
C2—C3—C4	118.4 (2)	H7A—C7—H7C	109.5
C2—C3—C6	119.6 (2)	H7B—C7—H7C	109.5
C4—C3—C6	122.0 (3)	O2—C8—O3	124.8 (3)
C3—C4—C5	118.6 (3)	O2—C8—C9	123.1 (3)
C3—C4—H4	120.7	O3—C8—C9	112.2 (3)
C5—C4—H4	120.7	C9 ⁱ —C9—C8	123.6 (3)

N1—C5—C4	123.5 (3)	С9 ^і —С9—Н9	118.2
N1—C5—H5	118.2	С8—С9—Н9	118.2

Symmetry code: (i) -x, -y, -z+1.

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	Н…А	D····A	<i>D</i> —H… <i>A</i>
O3—H3A…N1	0.98 (4)	1.64 (4)	2.599 (3)	166 (4)
C4—H4···O2 ⁱⁱ	0.93	2.57	3.471 (3)	164
C7—H7C···O2 ⁱⁱⁱ	0.96	2.58	3.489 (3)	158

Symmetry codes: (ii) *x*, *y*+1, *z*; (iii) *x*+1, *y*+1, *z*.