Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.002 Å R factor = 0.063 wR factor = 0.162 Data-to-parameter ratio = 18.1

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

Zwitterionic 3-carboxylatomethyl-3-carboxymethyl-2-oxo-2,3-dihydroimidazo[1,2-*a*]pyridin-1-ium

The title compound, $C_{11}H_{10}N_2O_5$, exists in the solid state as a carboxylate-imidazolium zwitterion. Intermolecular $O-H\cdots O$ and $N-H\cdots O$ hydrogen bonds link the molecules into two-dimensional layers.

Received 19 August 2006 Accepted 14 September 2006

Comment

Imidazo[1,2-*a*]pyridine derivatives are important intermediates in organic synthesis (Baumann *et al.*, 1984), especially in the synthesis of biologically active and medicinally useful agents. For instance, they are widely used in the synthesis of cyclin-dependent kinase (CDK) inhibitors (Anderson *et al.*, 2003), sleep inducers (Hempel & Blaschke, 1996), anticonvulsant agents (Trapani *et al.*, 2003) and antiviral agents (Gueiffier *et al.*, 1998; Mavel *et al.*, 2002). We have exploited a new and convenient method to obtain the title compound, (I), which was prepared from the reaction of potassium chloroacetate with 2-aminopyridine in aqueous alkaline solution.



In the solid state, (I) exhibits a zwitterionic form (Fig. 1). Each zwitterionic molecule acts as T-shaped connector, forming hydrogen bonds to three other molecules. The carboxyl groups link to two other molecules through O- $H \cdots O$ hydrogen bonds (Table 1) to form chains extending along the *c* direction. The NH group of the imidazolium ring links these chains *via* N- $H \cdots O$ hydrogen bonds (Table 1) to form layers in the *bc* plane (Fig. 2). Between adjacent layers, the pyridine rings form π - π stacking contacts, with an interplanar separation of 3.50 (1) Å and a centroid-centroid distance of 4.03 (1) Å.

Experimental

A solution of KOH (0.6 mol, 33.6 g) in 75 ml of water was added dropwise to a solution of chloroacetic acid (0.3 mol, 28.4 g) in 75 ml of water cooled in an ice bath. To the resulting alkaline solution, 2-aminopyridine (0.1 mol, 9.4 g) was added slowly, and the mixture was refluxed at 359 K for 30 h. 6 *N* HCl (53 ml) was added to neutralize the solution, which was then concentrated by boiling and cooled in an ice bath to remove KCl. After the mother liquor (pH 2–3) had been allowed to stand in a refrigerator for three days, the desired acid was

© 2006 International Union of Crystallography All rights reserved separated out and collected by filtration. The solid was recrystallized in a hot ethanol–water (1:1) mixture, and the colourless product was collected by filtration and air-dried (yield 23%). Block-shaped crystals of (I) suitable for X-ray diffraction analysis were obtained by slow evaporation of a DMF/H₂O (1:1) solution.

Crystal data

 $\begin{array}{l} C_{11}H_{10}N_2O_5\\ M_r = 250.21\\ \text{Monoclinic, } C2/c\\ a = 13.735 \ (3) \ \text{\AA}\\ b = 9.862 \ (2) \ \text{\AA}\\ c = 18.218 \ (4) \ \text{\AA}\\ \beta = 110.35 \ (3)^\circ\\ V = 2313.7 \ (10) \ \text{\AA}^3 \end{array}$

Data collection

Siemens SMART CCD diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Bruker, 1998) $T_{\rm min} = 0.955, T_{\rm max} = 0.969$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.063$ $wR(F^2) = 0.162$ S = 1.133040 reflections 168 parameters H atoms treated by a mixture of independent and constrained refinement

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	$D - \mathbf{H} \cdot \cdot \cdot A$
O3-H3···O1 ⁱ	1.12 (3)	1.39 (3)	2.4724 (16)	162 (3)
$N1 - H1 \cdot \cdot \cdot O2^{ii}$	0.86	2.03	2.7734 (16)	145

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

H atoms bound to C atoms were placed in calculated positions and constrained to ride on their parent atoms, with C–H distances in the range 0.93–0.97 Å, and with $U_{\rm iso}({\rm H}) = 1.2 U_{\rm eq}({\rm C})$. Atom H1, attached to N1, was also placed geometrically, with N–H = 0.86 Å and allowed to ride with $U_{\rm iso}({\rm H}) = 1.2 U_{\rm eq}({\rm N})$. Atom H3, attached to O3, was located in a difference Fourier map and refined isotropically without restraint.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1998); 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, 1998); software used to prepare material for publication: *SHELXTL*.

We are grateful for the financial support from NSFC (No. 20472078).



11493 measured reflections 3040 independent reflections 2852 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.050$ $\theta_{\text{max}} = 29.7^{\circ}$

$$\begin{split} &w = 1/[\sigma^2(F_o^2) + (0.0914P)^2 \\ &+ 1.0573P] \\ &where \ P = (F_o^2 + 2F_c^2)/3 \\ (\Delta/\sigma)_{max} < 0.001 \\ \Delta\rho_{max} = 0.75 \ e \ \text{\AA}^{-3} \\ \Delta\rho_{min} = -0.69 \ e \ \text{\AA}^{-3} \\ &\text{Extinction correction: SHELXL97} \\ &\text{Extinction coefficient: 0.030 (3)} \end{split}$$



Figure 1

The molecular structure of (I), with displacement ellipsoids shown at the 50% probability level for non-H atoms.



Figure 2

Projection along *a* showing a single layer of molecules in the *bc* plane, linked by $O-H\cdots O$ and $N-H\cdots O$ hydrogen bonds (dashed lines). H atoms are omitted.

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