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Key indicators

Single-crystal X-ray study
 $T = 100$ K
Mean $\sigma(\text{C}-\text{C}) = 0.001$ Å
 R factor = 0.030
 wR factor = 0.068
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>.

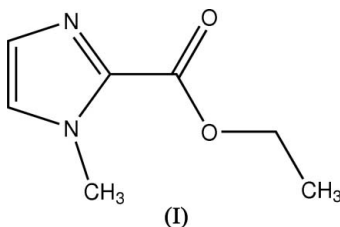
Ethyl 1-methylimidazole-2-carboxylate

The title compound, $\text{C}_7\text{H}_{10}\text{N}_2\text{O}_2$, crystallizes as a nearly planar molecule. The carboxyethyl group is twisted by 4.43 (14)° relative to the plane that includes the imidazole group. The carbonyl unit of the carboxyethyl group is oriented *anti* to the *N*-methyl group of the imidazole.

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Comment

The title compound, (I), has been used to synthesize polyamide anticancer agents that contain *N*-methylimidazole and *N*-methylpyrrole groups (Baird & Dervan, 1996; Baraldi *et al.*, 2003; Krowicki & Lown, 1987; Zaffaroni *et al.*, 2002), which bind to the minor groove of DNA and display G–C and C–G base pair recognition (Moser & Dervan, 1987; Marques *et al.*, 2002). Compound (I) is a potential synthon for synthesizing amide-functionalized imidazole chelates (Cheruzel *et al.*, 2002).



The structure of the related ethyl 1-methyl-4-nitroimidazole-2-carboxylate (Wu *et al.*, 2004), has been reported recently and contains disordered carboxyethyl and *N*-methyl groups, unlike the structure of (I). The carbonyl unit is oriented *anti* to the *N*-methyl group in (I), whereas in ethyl 1-methyl-4-nitroimidazole-2-carboxylate, the carbonyl group adopts a *syn* configuration.

Molecules of (I) are stacked along the crystallographic *a* axis (Fig. 2), the closest contacts between molecules being 3.4391 (13) Å, between N1 at (x, y, z) and C4' at $(-x, 1 - y, -z)$, and 3.4921 (12) Å, between C5 at (x, y, z) and O5'' at $(1 - x, 1 - y, 1 - z)$. The carboxyethyl groups in alternating layers of the stack are oriented in opposite directions, minimizing steric interactions between symmetry-related molecules in adjacent stacks. The imidazole ring in (I) is planar and the torsion angle (N2–C1–C5–O2) associated with the carboxyethyl group is 4.43 (14)°, compared with 15.0 (1)° reported for ethyl 1-methyl-4-nitroimidazole-2-carboxylate (Wu *et al.*, 2004).

Experimental

Compound (I) was synthesized following a previously reported procedure (Krowicki & Lown, 1987). Ethyl chloroformate (0.28 mol)

was added to an acetonitrile solution (60 ml) of *N*-methylimidazole (0.12 mol) and triethylamine (0.22 mol) at 253 K. After 24 h the solution was filtered and the solvent was removed. The resulting residue was dissolved in water and extracted with chloroform. A white solid obtained by column chromatography (silica gel, ethyl acetate) was recrystallized by slow evaporation of the solution in ethyl acetate.

Crystal data

$C_7H_{10}N_2O_2$	$D_x = 1.378 \text{ Mg m}^{-3}$
$M_r = 154.17$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 5362 reflections
$a = 7.1040$ (8) Å	$\theta = 2.7\text{--}28.0^\circ$
$b = 15.2244$ (18) Å	$\mu = 0.10 \text{ mm}^{-1}$
$c = 7.4511$ (9) Å	$T = 100$ (2) K
$\beta = 112.797$ (2)°	Plate, colorless
$V = 742.92$ (15) Å ³	$0.37 \times 0.29 \times 0.09 \text{ mm}$
$Z = 4$	

Data collection

Bruker SMART APEX diffractometer	1728 independent reflections
ω scans	1636 reflections with $I > 2\sigma(I)$
Absorption correction: multi-scan (SADABS; Sheldrick, 2001)	$R_{\text{int}} = 0.013$
$T_{\text{min}} = 0.960$, $T_{\text{max}} = 0.988$	$\theta_{\text{max}} = 28.0^\circ$
6360 measured reflections	$h = -9 \rightarrow 9$
	$k = -19 \rightarrow 19$
	$l = -9 \rightarrow 9$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0213P)^2 + 0.3603P]$
$R[F^2 > 2\sigma(F^2)] = 0.030$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.068$	$(\Delta/\sigma)_{\text{max}} = 0.001$
$S = 1.02$	$\Delta\rho_{\text{max}} = 0.36 \text{ e \AA}^{-3}$
1728 reflections	$\Delta\rho_{\text{min}} = -0.22 \text{ e \AA}^{-3}$
140 parameters	
All H-atom parameters refined	

All H atoms were located in difference electron-density maps and refined isotropically [C—H = 0.948 (13)–0.991 (15) Å].

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 2001); software used to prepare material for publication: *SHELXTL*.

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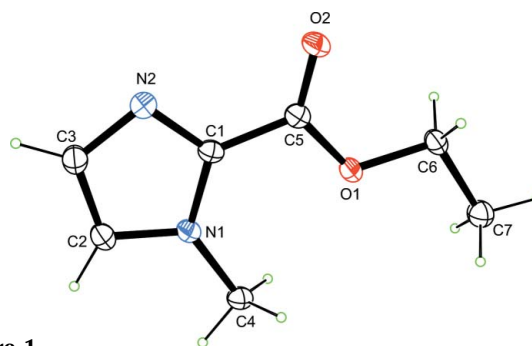


Figure 1
ORTEP-3 drawing (Farrugia, 1997) showing 50% probability displacement ellipsoids. H atoms are shown as spheres of arbitrary radii.

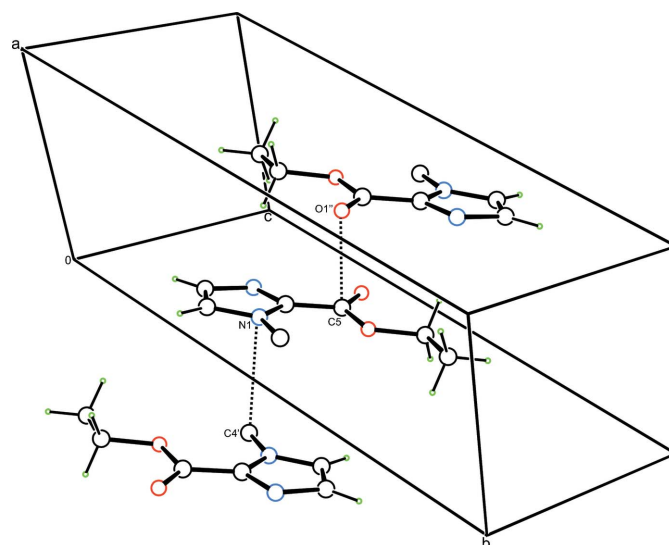


Figure 2
A partial packing diagram, showing molecules of (I) stacked along the crystallographic *a* axis. H atoms of the methyl C atoms have been omitted for clarity. Dashed lines indicate short contacts. [Symmetry codes: (') $-x, 1 - y, -z$; (") $1 - x, 1 - y, 1 - z$.]

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