

3-[2-(3-Chlorophenyl)ethyl]pyridine-2-carbonitrile

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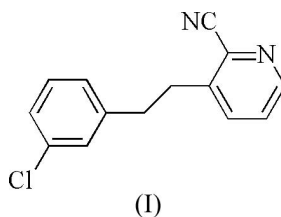
Key indicators

Single-crystal X-ray study
 $T = 293$ K
Mean $\sigma(\text{C}-\text{C}) = 0.003$ Å
 R factor = 0.048
 wR factor = 0.127
Data-to-parameter ratio = 13.5For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

The title compound, $\text{C}_{14}\text{H}_{11}\text{ClN}_2$, is the key intermediate in the synthesis of the antihistaminic drug loratadine. The benzene and pyridine rings are connected through the ethylene bridge and are coplanar. There are intermolecular $\pi-\pi$ and $\text{C}-\text{Cl}\cdots\text{N}$ interactions and weak hydrogen bonds stabilizing the packing.

Comment

The title compound, (I), which has been synthesized according to the literature method of Schumacher *et al.* (1989), is an intermediate in the synthesis of the antihistaminic drug loratadine. In the structure, shown in Fig. 1, the pyridine ring is connected to the benzene ring through the ethylene bridge. The benzene ring is coplanar with the pyridine ring. The benzene ring is planar, with an r.m.s. deviation of 0.003 Å, and the Cl atom deviates by only 0.004 (3) Å from this plane. The molecule adopts a stepped *trans* conformation, as shown by the torsion angles $\text{C}7-\text{C}8-\text{C}9-\text{C}10$ and $\text{C}5-\text{C}4-\text{C}7-\text{C}8$. The conformation of the molecule is similar to that of an earlier reported related compound, namely 2,2'-dinitrobenzyl (Yathirajan *et al.*, 2004). Several weak intermolecular bonds stabilize the packing, *viz.* the face-to-face contacts $\text{C}_g(\text{pyridine ring})\cdots\text{C}_g(\text{pyridine ring})^i$ at a distance of 4.240 (1) Å and $\text{C}_g(\text{pyridine ring})\cdots\text{C}_g(\text{benzene ring})^{ii}$ at a distance of 2.754 (1) Å [symmetry codes: (i) $-x, -y + 2, -z + 1$; (ii) $-1 + x, y, z$]. A further close contact is $\text{Cl}\cdots\text{N}1^{iii}$ [symmetry code: (iii) $x + \frac{3}{2}, -y + \frac{3}{2}, z + \frac{1}{2}$] of 3.280 (2) Å (Fig. 2).



Experimental

A solution containing 3-[2-(3-chlorophenyl)ethyl]-*N*-(1,1-dimethylethyl)pyridine-2-carboxamide (35 g, 0.11 mol) and POCl_3 (150 ml) was heated under reflux for 3.5 h. Excess POCl_3 was removed by distillation and the remaining solution was poured into ice-water. The pH of the solution was adjusted to 8 with 60% aqueous NaOH at room temperature. The mixture was stirred for 2.5 h, during which time the pH was maintained at 8. The product was collected by filtration, washed with water and dried to yield 25 g (93.9%) of crystalline product.

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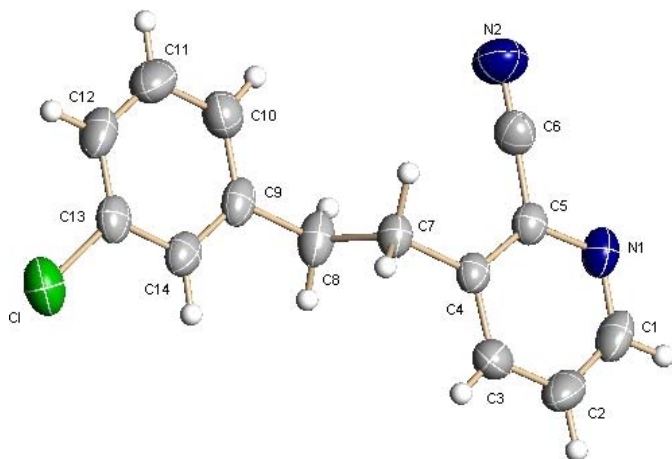


Figure 1
The molecular configuration and atom-numbering scheme for (I). Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as circles of arbitrary radius

Crystal data

$C_{14}H_{11}ClN_2$
 $M_r = 242.70$
 Monoclinic, $P2_1/n$
 $a = 7.5645$ (8) Å
 $b = 12.2332$ (12) Å
 $c = 13.5558$ (14) Å
 $\beta = 102.867$ (2)°
 $V = 1222.9$ (2) Å³
 $Z = 4$

$D_x = 1.318$ Mg m⁻³
 Mo $K\alpha$ radiation
 Cell parameters from 2106 reflections
 $\theta = 5.7$ – 52.7 °
 $\mu = 0.29$ mm⁻¹
 $T = 293$ (2) K
 Prism, colorless
 $0.51 \times 0.39 \times 0.22$ mm

Data collection

Bruker SMART 1K CCD area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick, 2002)
 $T_{\min} = 0.598$, $T_{\max} = 0.939$
 7051 measured reflections

2668 independent reflections
 1924 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.049$
 $\theta_{\text{max}} = 27.0$ °
 $h = -9 \rightarrow 9$
 $k = -10 \rightarrow 15$
 $l = -17 \rightarrow 17$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.048$
 $wR(F^2) = 0.127$
 $S = 1.01$
 2668 reflections
 198 parameters

All H-atom parameters refined
 $w = 1/[\sigma^2(F_o^2) + (0.0674P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 0.35$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.34$ e Å⁻³

Table 1

Selected torsion angles (°).

C5–C4–C7–C8	91.0 (2)	C7–C8–C9–C10	84.9 (2)
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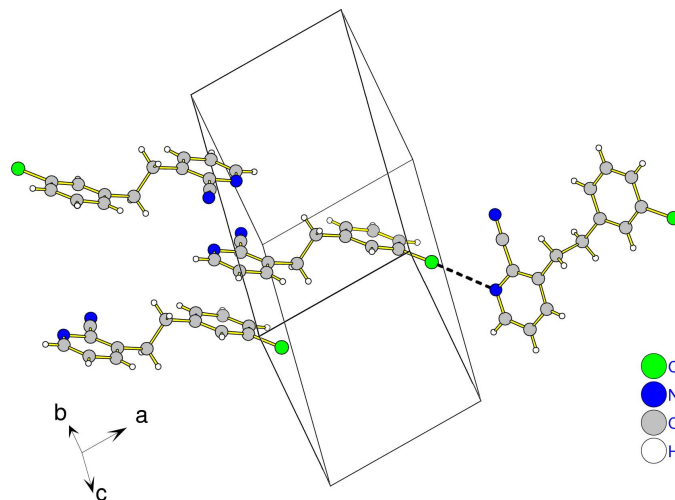


Figure 2
Part of the molecular packing of (I), showing the face-to-face contacts between the π ring systems. The short intermolecular Cl \cdots N1 bond is indicated with a broken line.

All H-atom coordinates were refined, as were the individual isotropic displacement parameters.

Data collection: *SMART* (Bruker, 1997); cell refinement: *SAINTE* (Bruker, 1997); data reduction: *SAINTE*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Sheldrick, 2000) and *DIAMOND* (Brandenburg, 1999); software used to prepare material for publication: *SHELXTL*.

X-ray data were collected at Shanghai Institute of Organic Chemistry, Chinese Academy of Sciences.

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