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(S)-2-(2-Chloroquinolin-3-yl)-2-[(S)- α -methylbenzylamino]acetonitrile

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Key indicators

Single-crystal X-ray study
 T = 293 K
 Mean σ (C–C) = 0.005 Å
 R factor = 0.055
 wR factor = 0.093
 Data-to-parameter ratio = 14.4

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

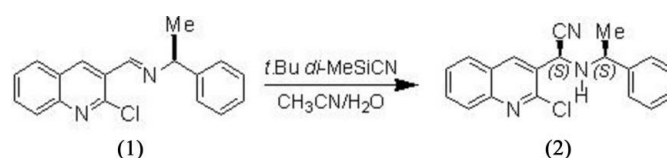
The title compound, C₁₉H₁₆ClN₃, crystallizes with two independent molecules in the asymmetric unit. The structure is stabilized by C–H...N, N–H...N and C–H...Cl hydrogen bonds.

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Comment

Quinolines are an important group of heterocyclic compounds. Among these, 2-chloro-3-formylquinolines occupy a prominent position as they are key intermediates for further (β)-annulation of a wide variety of rings and for various functional group interconversions (Meth-Cohn, 1993). Particular interest in quinoline derivatives arises owing to their biological activity, namely as antibiotics (Jackson & Meth-Cohn, 1995; Kansagra *et al.*, 2000), anti-inflammatories (Schroderet, 1989), anti-tumourals (Joseph *et al.*, 2002), anti-oxidants (Laalaoui *et al.*, 2003) and analgesics (Heide *et al.*, 1986; Solomon, 1970). In the same way, α -aminoacids are of great biological and economic importance (Williams, 1989). The asymmetric Strecker reaction is one of the most important methods for the synthesis of enantiomerically pure α -amino-nitrile derivatives, which are useful intermediates for the synthesis of α -aminoacids. The use of (*S*)-(–)- α -methylbenzylamine-derived aldimines has a significant role in the diastereoselective Strecker synthesis (Bhanu-Prasad *et al.*, 2004).

In recent years, we have developed a programme devoted to the synthesis and biological evaluation of quinolyl derivatives (Moussaoui *et al.*, 2002; Kedjadja *et al.*, 2004; Menasra *et al.*, 2004, 2005; Rezig *et al.*, 2000). In a continuation of our efforts in this area, we report here a short and efficient procedure for the preparation of the title α -aminonitrile, (2), containing a quinolyl ring system, and its crystal structure determination.



The crystallographic asymmetric unit of (2) contains two independent molecules, labelled *a* and *b* (Fig. 1). The analysis shows that atoms C11 and C12 each have an *S* configuration in both independent molecules. The geometric parameters of (2) (Table 1) are in agreement with those of other structures containing similar molecular connectivity (Benali-Cherif, Cherouana *et al.*, 2002; Benali-Cherif, Dokhane & Abdaoui, 2002). The 11 atoms defining the chloroquinolyl planes, *i.e.* N1 and C11–C10, have maximum deviations of 0.0037 (3) Å for

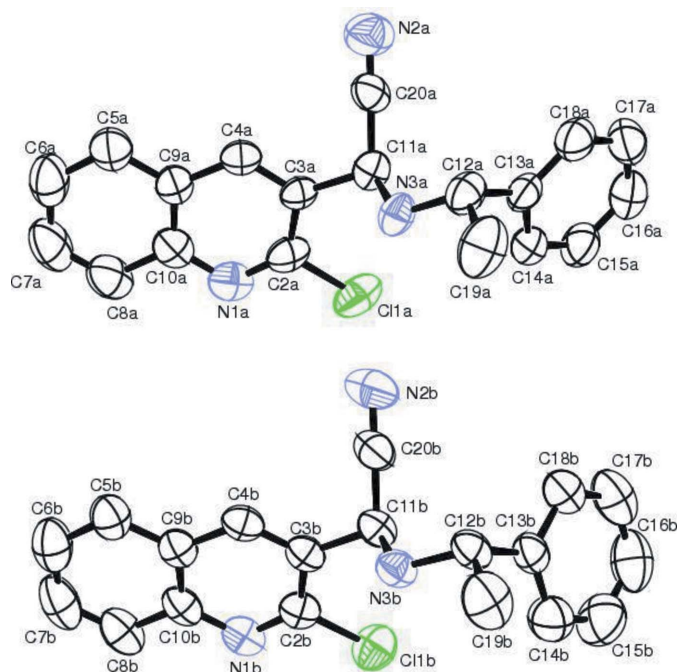


Figure 1
Views of the two independent molecules of (2), showing 50% probability displacement ellipsoids and the atom-numbering scheme. H atoms have been omitted for clarity.

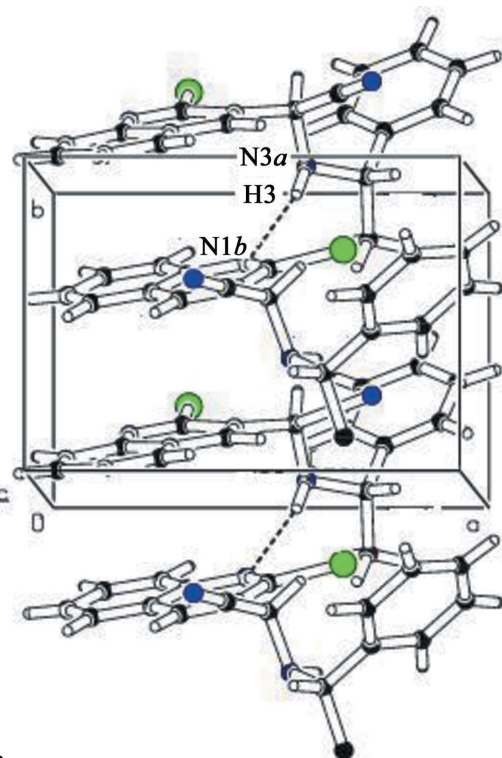


Figure 2
The unit-cell contents of (2), highlighting the N—H...N hydrogen bonding (dashed lines).

atom C7a and 0.0035 (3) Å for atom C7b for the two independent molecules. In the crystal structure, these planes are almost parallel, forming a dihedral angle of 7.40 (4)°. The clear difference between molecules *a* and *b* is noted in the dihedral

angles formed between the chloroquinolyl and phenyl groups of 27.83 (6) and 50.53 (8)°, respectively.

The three-dimensional crystal structure of (2) is stabilized via a variety of hydrogen-bonding interactions of the types C—H...N, N—H...N and C—H...Cl (Steiner, 1996), as analysed by PARST (Nardelli, 1995) and summarized in Fig. 2 and Table 2.

Experimental

Chiral imine (1) was prepared by condensation of optically active (*S*)-(-)- α -methylbenzylamine with 2-chloro-3-formylquinoline, according to the literature procedure of Meth-Cohn *et al.* (1981). Treatment of (1) with *tert*-butyldimethylsilyl cyanide at room temperature in CH₃CN solution with a few drops of water provided a mixture of two diastereoisomers as a yellow solid (yield 84%). Crystals of (2) were obtained by fractional crystallization from a hexane—CH₂Cl₂ (9:1) solution of this mixture. The isomeric ratio of (2) (63%) was determined from the ¹H NMR spectrum of the crude product (m.p. 383 K).

Crystal data

C₁₉H₁₆ClN₃
M_r = 321.8
 Monoclinic, *P*2₁
a = 9.8540 (1) Å
b = 7.1090 (1) Å
c = 23.9330 (3) Å
 β = 91.590 (2)°
V = 1675.91 (4) Å³
Z = 4

D_x = 1.275 Mg m⁻³
 Mo *K* α radiation
 Cell parameters from 2548 reflections
 θ = 1.7–28.0°
 μ = 0.23 mm⁻¹
T = 293 (2) K
 Prism, yellow
 0.2 × 0.15 × 0.1 mm

Data collection

Nonius KappaCCD area-detector diffractometer
 φ and ω scans
 Absorption correction: none
 35588 measured reflections
 7662 independent reflections

4482 reflections with *I* > 2 σ (*I*)
*R*_{int} = 0.065
 θ _{max} = 28.0°
h = -13 → 12
k = -8 → 9
l = -31 → 31

Refinement

Refinement on *F*²
R[*F*² > 2 σ (*F*²)] = 0.055
wR(*F*²) = 0.093
S = 1.03
 7662 reflections
 531 parameters
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0312P)^2 + 0.1354P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.002$
 $\Delta\rho_{\max} = 0.15 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.15 \text{ e \AA}^{-3}$
 Absolute structure: Flack (1983),
 3303 Friedel pairs
 Flack parameter: -0.01 (5)

Table 1

Selected geometric parameters (Å, °).

N2a—C20a	1.113 (3)	N2b—C20b	1.134 (3)
C17a—C18a	1.374 (4)	C17b—C18b	1.394 (5)
N3a—C11a—C20a	114.4 (2)	N3b—C11b—C20b	109.6 (2)
C11a—N3a—C12a	117.1 (2)	C11b—N3b—C12b	114.8 (2)
C3a—C11a—N3a—C12a	-169.6 (2)	C3b—C11b—N3b—C12b	178.9 (2)
C4a—C3a—C11a—N3a	100.8 (3)	C4b—C3b—C11b—N3b	89.2 (3)
C19a—C12a—C13a—C18a	-99.7 (3)	C19b—C12b—C13b—C18b	-125.7 (3)

Table 2

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$N3a-H3\cdots N1b^i$	0.90 (3)	2.31 (3)	3.189 (3)	165.6 (2)
$N3b-H3'\cdots N2a$	0.87 (2)	2.68 (2)	3.508 (3)	159.6 (19)
$C11b-H11'\cdots C11b$	0.98 (2)	2.641 (19)	3.067 (3)	106.3 (13)
$C11a-H11\cdots C11a$	1.00 (3)	2.76 (2)	3.044 (3)	97.4 (14)

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

All H atoms were freely refined except for the methyl H atoms bonded to atom C19, for which $C-H = 0.96$ Å and $U_{iso}(H) = 1.5U_{iso}(C)$.

Data collection: *KappaCCD Server Software* (Nonius, 1998); cell refinement: *DENZO* and *SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *DENZO* and *SCALEPACK*; program(s) used to solve structure: *SIR2004* (Burla *et al.*, 2005); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3* (Farrugia, 1997) and *PLATON* (Spek, 2003); software used to prepare material for publication: *WinGX* (Farrugia, 1999) and *enCIFer* (Allen *et al.*, 2004).

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