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2-Amino-6-chloro-4-(1-phenylethylamino)pyrimidine

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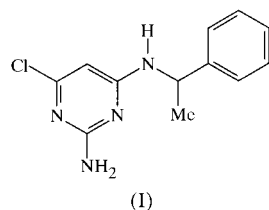
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The structure of the title compound, C₁₂H₁₃ClN₄, (I), comprises a racemic mixture of chiral molecules associated by N—H···N hydrogen-bonding interactions. The dihedral angle between the two rings is 77.90 (6)°.



Experimental

Crystals of (I) were obtained from Spa Contract Synthesis.

Crystal data

C₁₂H₁₃ClN₄
M_r = 248.71
Tetragonal, I4₁/a
a = 17.4851 (7) Å
c = 16.2878 (9) Å
V = 4979.6 (4) Å³
Z = 16
D_x = 1.327 Mg m⁻³

Mo Kα radiation
Cell parameters from 10256 reflections
θ = 2.91–27.48°
μ = 0.290 mm⁻¹
T = 150 (2) K
Block, yellow
0.15 × 0.10 × 0.10 mm

Data collection

Enraf–Nonius KappaCCD area-detector diffractometer
φ and ω scans
Absorption correction: multi-scan (SORTAV; Blessing, 1995)
T_{min} = 0.958, T_{max} = 0.972
15 828 measured reflections
2828 independent reflections

1893 reflections with I > 2σ(I)
R_{int} = 0.137
θ_{max} = 27.43°
h = -22 → 22
k = -19 → 19
l = -17 → 21
Intensity decay: none

Refinement

Refinement on F²
R[F² > 2σ(F²)] = 0.051
wR(F²) = 0.134
S = 1.012
2828 reflections
156 parameters
H-atom parameters constrained

w = 1/[σ²(F_o²) + (0.0658P)² + 0.1078P]
where P = (F_o² + 2F_c²)/3
(Δ/σ)_{max} < 0.001
Δρ_{max} = 0.24 e Å⁻³
Δρ_{min} = -0.35 e Å⁻³
Extinction correction: SHELXL97
Extinction coefficient: 0.0014 (3)

Table 1

Hydrogen-bonding geometry (Å, °).

D—H···A	D—H	H···A	D···A	D—H···A
N21—H21···N3 ⁱ	0.88	2.23	3.098 (2)	169
N21—H22···N1 ⁱⁱ	0.88	2.17	3.043 (2)	172

Symmetry codes: (i) $\frac{3}{4} - y, x - \frac{1}{4}, \frac{3}{4} - z$; (ii) $\frac{1}{2} - x, \frac{1}{2} - y, \frac{1}{2} - z$.

All H atoms were included in the refinement at calculated positions as riding, with the C—H distance set to either 0.98 (for CH₃) or 0.95 Å (for aryl H) and the N—H distance set to 0.88 Å.

Data collection: DENZO (Otwinowski & Minor, 1997) and COLLECT (Hooft, 1998); cell refinement: DENZO and COLLECT; data reduction: DENZO and COLLECT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); software used to prepare material for publication: SHELXL97.

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References

- Blessing, R. H. (1995). *Acta Cryst.* **A51**, 33–37.
Hooft, R. (1998). *COLLECT*. Nonius BV, Delft, The Netherlands.
Otwinowski, Z. & Minor, W. (1997). *Methods Enzymol.* **276**, 307–326.
Sheldrick, G. M. (1997). *SHELXS97* and *SHELXL97*. University of Göttingen, Germany.

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