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## Structure Reports

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## Daniel E. Lynch ${ }^{\text {a }}$ and Ian McClenaghan ${ }^{\text {b }} \dagger$

${ }^{\mathrm{a}}$ School of Natural and Environmental Sciences, Coventry University, Coventry CV1 5FB,
England, and ${ }^{\mathbf{b}}$ Spa Contract Synthesis, School of Natural and Environmental Sciences, Coventry University, Coventry CV1 5FB, England

+ E-mail: 106355.1670@CompuServe.com.
Correspondence e-mail:
apx106@coventry.ac.uk


## Key indicators

Single-crystal X-ray study
$T=150 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.002 \AA$
$R$ factor $=0.043$
$w R$ factor $=0.115$
Data-to-parameter ratio $=17.2$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## 2-Chloro-4-methylquinoline

The structure of the title compound, $\mathrm{C}_{10} \mathrm{H}_{8} \mathrm{ClN}$, comprises planar molecules that form stacked columns parallel to the $b$ cell direction. Two symmetry-related molecules associate via $\mathrm{C}-\mathrm{H} \cdots \mathrm{N}$ interactions to form a non-planar $R_{2}^{2}(8)$ dimer.

## Experimental

The title compound, (I), was prepared by Spa Contract Synthesis. Crystals of (I) were grown from a methanol solution.

(I)

Crystal data
$\mathrm{C}_{10} \mathrm{H}_{8} \mathrm{ClN}$
$M_{r}=177.62$
Orthorhombic, Pbca
$a=12.721$ (3) $\AA$ 。
$b=7.8961(16) \AA$
$c=16.617$ (3) $\AA$
$V=1669.1(6) \AA^{3}$
$Z=8$
$D_{x}=1.414 \mathrm{Mg} \mathrm{m}^{-3}$
Data collection

| Enraf-Nonius KappaCCD area- | 1892 independent reflections |
| :--- | :--- |
| $\quad$ detector diffractometer | 1466 reflections with $I>2 \sigma(I)$ |
| $\varphi$ and $\omega$ scans | $R_{\text {int }}=0.074$ |
| Absorption correction: multi-scan | $\theta_{\max }=27.5^{\circ}$ |
| $\quad(S O R T A V ;$ Blessing, 1995 $)$ | $h=-16 \rightarrow 16$ |
| $T_{\min }=0.926, T_{\max }=0.973$ | $k=-10 \rightarrow 10$ |
| 7671 measured reflections | $l=-21 \rightarrow 17$ |

## Refinement

| Refinement on $F^{2}$ | H-atom parameters constrained |
| :--- | :--- |
| $R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.043$ | $w=1 /\left[\sigma^{2}\left(F_{o}{ }^{2}\right)+(0.0699 P)^{2}\right]$ |
| $w R\left(F^{2}\right)=0.115$ | where $P=\left(F_{o}^{2}+2 F_{c}^{2}\right) / 3$ |
| $S=1.02$ | $(\Delta / \sigma)_{\max }=0.001$ |
| 1892 reflections | $\Delta \rho_{\max }=0.33 \mathrm{e} \AA^{-3}$ |
| 110 parameters | $\Delta \rho_{\min }=-0.36 \mathrm{e}^{-3}$ |

Table 1
Hydrogen-bonding geometry $\left(\AA^{\circ},^{\circ}\right)$.

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{C} 8-\mathrm{H} 8 \cdots \mathrm{~N} 1^{\mathrm{i}}$ | 0.95 | 2.61 | $3.517(2)$ | 160 |

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

All H atoms were included in the refinement at calculated positions as riding models, with $\mathrm{C}-\mathrm{H}$ set to $0.95(\mathrm{Ar}-\mathrm{H})$ and $0.98 \AA$ $\left(\mathrm{CH}_{3}\right)$.

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Data collection: DENZO (Otwinowski \& Minor, 1997) and COLLECT (Hooft, 1998); cell refinement: DENZO and COLLECT; data reduction: $D E N Z O$ 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

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The molecular configuration and atom-numbering scheme for (I), showing $30 \%$ probability displacement ellipsoids.

