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## Structure Reports

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## 2-tert-Butylamino-1-(4-chlorophenyl)-4,4-dimethyl-1H-imidazolin-5(4H)-one

In the title compound, $\mathrm{C}_{15} \mathrm{H}_{20} \mathrm{ClN}_{3} \mathrm{O}$, the five-membered ring and the amino N atom are essentially coplanar.

## Key indicators

Single-crystal X-ray study
$T=292 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.003 \AA$
$R$ factor $=0.049$
$w R$ factor $=0.123$
Data-to-parameter ratio $=18.4$

For details of how these key indicators were
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## Crystal data

| $\mathrm{C}_{15} \mathrm{H}_{20} \mathrm{ClN}_{3} \mathrm{O}$ |
| :---: |
|  |
| Monoclinic, $P 2_{1} /$ $a=6.0758$ (8) A |
| $b=18.8340$ (2) A |
| $=13.8603$ (18) |
| $=98.530$ (2) ${ }^{\circ}$ |
| $V=1568.5$ (3) $\AA^{3}$ |
|  |

$D_{x}=1.244 \mathrm{Mg} \mathrm{m}^{-3}$
$M_{r}=293.79$
Monoclinic, $P 2_{1} / n$
$a=6.0758$ (8) $\AA$ 。
$b=18.8340$ (2) A
$\beta=98.530$ (2) ${ }^{\circ}$
$V=1568.5$ (3) $\AA^{3}$
$Z=4$
Mo $K \alpha$ radiation
Cell parameters from 2568 reflections
$\theta=2.6-22.7^{\circ}$
$\mu=0.24 \mathrm{~mm}^{-1}$
$T=292$ (2) K
Block, colourless
$0.40 \times 0.20 \times 0.10 \mathrm{~mm}$

## Data collection

Bruker SMART CCD area-detector
diffractometer
$\varphi$ and $\omega$ scans
Absorption correction: none
10046 measured reflections
3425 independent reflections
2208 reflections with $I>2 \sigma(I)$
$R_{\text {int }}=0.060$
$\theta_{\text {max }}=27.0^{\circ}$
$h=-7 \rightarrow 7$
$k=-22 \rightarrow 24$
$l=-17 \rightarrow 16$

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.049$
$w R\left(F^{2}\right)=0.123$
$S=0.98$
3425 reflections
186 parameters

H-atom parameters constrained
$w=1 /\left[\sigma^{2}\left(F_{\mathrm{o}}{ }^{2}\right)+(0.0571 P)^{2}\right.$
where $P=\left(F_{\mathrm{o}}{ }^{2}+2 F_{\mathrm{c}}{ }^{2}\right) / 3$
$(\Delta / \sigma)_{\max }<0.001$ 。
$\Delta \rho_{\max }=0.27 \mathrm{e}^{-3}$
$\Delta \rho_{\min }=-0.17 \mathrm{e}^{-3}$

Table 1
Selected geometric parameters ( $\left({ }^{\circ},{ }^{\circ}\right.$ ).

| C1-C2 | $1.366(3)$ | $\mathrm{C} 7-\mathrm{N} 1$ | $1.420(2)$ |
| :--- | :---: | :--- | ---: |
| C3-C4 | $1.378(3)$ | $\mathrm{C} 8-\mathrm{N} 2$ | $1.472(2)$ |
| C4-C6 | $1.378(3)$ | $\mathrm{C} 8-\mathrm{C} 9$ | $1.514(3)$ |
| C5-C6 | $1.373(2)$ | $\mathrm{C} 9-\mathrm{O} 1$ | $1.211(2)$ |
| C6-N1 | $1.426(2)$ | $\mathrm{C} 9-\mathrm{N} 1$ | $1.382(2)$ |
| C7-N2 | $1.272(2)$ | $\mathrm{C} 12-\mathrm{N} 3$ | $1.487(2)$ |
| C7-N3 | $1.354(2)$ |  |  |
| C5-C6-C4 | $119.37(17)$ | $\mathrm{O} 1-\mathrm{C} 9-\mathrm{N} 1$ | $125.24(18)$ |
| C5-C6-N1 | $120.14(16)$ | $\mathrm{N} 1-\mathrm{C} 9-\mathrm{C} 8$ | $105.81(15)$ |
| $\mathrm{C} 4-\mathrm{C} 6-\mathrm{N} 1$ | $120.46(16)$ | $\mathrm{C} 9-\mathrm{N} 1-\mathrm{C} 7$ | $107.06(15)$ |
| N2-C7-N3 | $127.10(16)$ | $\mathrm{C} 9-\mathrm{N} 1-\mathrm{C} 6$ | $125.00(14)$ |
| N2-C7-N1 | $114.80(15)$ | $\mathrm{C} 7-\mathrm{N} 1-\mathrm{C} 6$ | $127.84(14)$ |
| N3-C7-N1 | $118.09(16)$ | $\mathrm{C} 7-\mathrm{N} 2-\mathrm{C} 8$ | $107.38(14)$ |
| N2-C8-C9 | $104.76(14)$ |  |  |
| N2-C7-N1-C9 | $0.1(2)$ | $\mathrm{C} 5-\mathrm{C} 6-\mathrm{N} 1-\mathrm{C} 7$ | $117.96(19)$ |
| N3-C7-N1-C9 | $178.79(16)$ | $\mathrm{C} 4-\mathrm{C} 6-\mathrm{N} 1-\mathrm{C} 7$ | $-63.8(2)$ |
| N2-C7-N1-C6 | $176.62(16)$ | $\mathrm{N} 3-\mathrm{C} 7-\mathrm{N} 2-\mathrm{C} 8$ | $178.64(18)$ |
| N3-C7-N1-C6 | $-4.7(3)$ | $\mathrm{N} 1-\mathrm{C} 7-\mathrm{N} 2-\mathrm{C} 8$ | $-2.9(2)$ |
| C5-C6-N1-C9 | $-66.2(2)$ | $\mathrm{N} 2-\mathrm{C} 7-\mathrm{N} 3-\mathrm{C} 12$ | $-14.1(3)$ |
| C4-C6-N1-C9 | $112.1(2)$ | $\mathrm{N} 1-\mathrm{C} 7-\mathrm{N} 3-\mathrm{C} 12$ | $167.40(15)$ |

H atoms were placed at calculated positions and refined as riding $(\mathrm{C}-\mathrm{H}=0.93-0.96 \AA$ and $\mathrm{N}-\mathrm{H}=0.86 \AA)$, with $U_{\text {iso }}(\mathrm{H})=1.2(\mathrm{CH}$, $\mathrm{NH})$ or $1.5\left(\mathrm{CH}_{3}\right)$ times $U_{\text {eq }}$ (parent atom).

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

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Figure 1
View of the molecular structure of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the $50 \%$ probability level.


Figure 2
Packing diagram for (I), viewed down the $a$ axis.

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