

2,6-Di-*tert*-butyl-4-[(1-methyl-1*H*-benzimidazol-2-yl)-imino]cyclohexa-2,5-dien-1-one

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## Key indicators

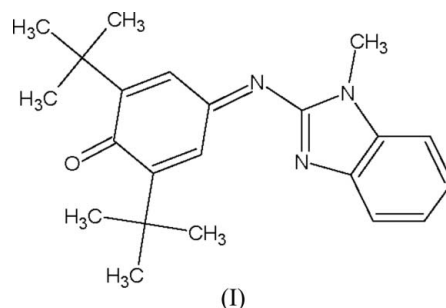
Single-crystal X-ray study  
 $T = 293$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.006$  Å  
 $R$  factor = 0.064  
 $wR$  factor = 0.214  
Data-to-parameter ratio = 12.9For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.

Except for two pairs of methyl groups, all non-H atoms of the title compound,  $\text{C}_{22}\text{H}_{27}\text{N}_3\text{O}$ , occupy special positions on a mirror plane. The molecules are packed in a herring-bone manner and form stacks along [010].

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## Comment

Compounds containing a benzimidazole group have potential pharmaceutical applications. Some of them exhibit a broad spectrum of pharmacological actions, ranging from anthelmintic activity (Geary *et al.*, 1999) to anticancer therapy (Lage *et al.*, 2006).



In the title compound, (I), all non-H atoms, except for C16 and C19, together with their symmetry-equivalents, occupy special positions on a mirror plane (Fig. 1). All bond lengths and angles are close to standard values (Allen *et al.*, 1987). Molecules of (I) are packed in stacks extending along [010], within which adjacent molecules are related by inversion operations (symmetry codes:  $-x, -y, 1 - z$  and  $-x, 1 - y, 1 - z$ ). The interplanar distance within the stacks is 3.39 (1) Å. The stacks are arranged in a herring-bone manner (Fig. 2), with no significant C—H...O or C—H...N interactions between them.

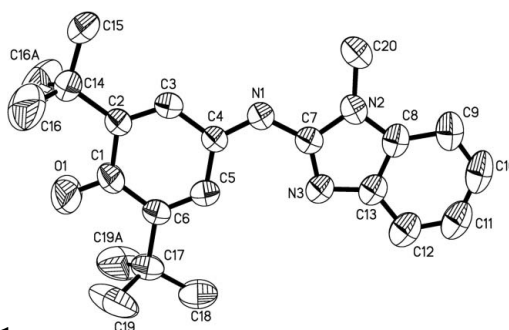


Figure 1

The molecular structure of (I) with displacement ellipsoids drawn at the 50% probability level. Atoms C16A and C19A are generated by the symmetry operation  $(x, \frac{1}{2} - y, z)$ . H atoms have been omitted.

## Experimental

Crystals of (I) were grown by slow evaporation of a solution in acetonitrile.

### Crystal data

$C_{22}H_{27}N_3O$	$Z = 4$
$M_r = 349.47$	$D_x = 1.107 \text{ Mg m}^{-3}$
Orthorhombic, $Pnma$	Mo $K\alpha$ radiation
$a = 18.372 (4) \text{ \AA}$	$\mu = 0.07 \text{ mm}^{-1}$
$b = 6.7816 (14) \text{ \AA}$	$T = 293 (2) \text{ K}$
$c = 16.833 (3) \text{ \AA}$	Prism, yellow
$V = 2097.3 (7) \text{ \AA}^3$	$0.50 \times 0.30 \times 0.30 \text{ mm}$

### Data collection

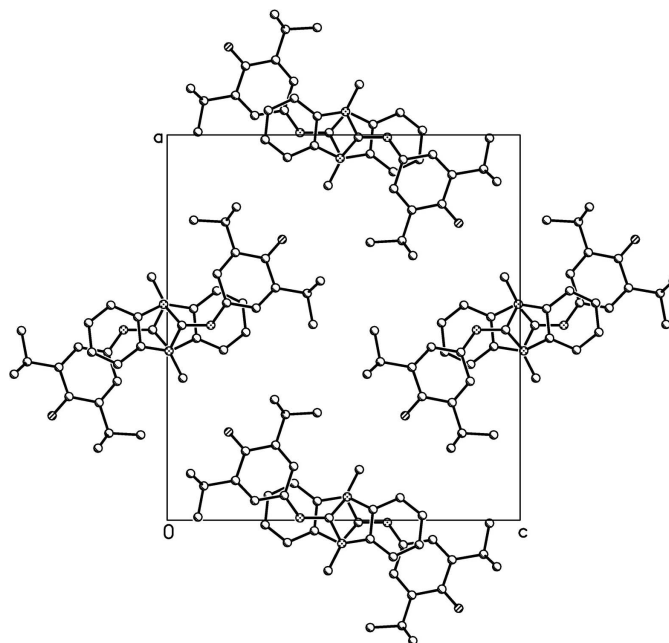
Enraf–Nonius CAD-4 diffractometer	996 reflections with $I > 2\sigma(I)$
$\omega/2\theta$ scans	$\theta_{\text{max}} = 25.0^\circ$
Absorption correction: none	2 standard reflections
1985 measured reflections	every 98 reflections
1985 independent reflections	intensity decay: 3.7%

### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.1333P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.064$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.214$	$(\Delta/\sigma)_{\text{max}} < 0.001$
$S = 1.00$	$\Delta\rho_{\text{max}} = 0.28 \text{ e \AA}^{-3}$
1985 reflections	$\Delta\rho_{\text{min}} = -0.23 \text{ e \AA}^{-3}$
154 parameters	Extinction correction: <i>SHELXL97</i>
H-atom parameters constrained	Extinction coefficient: 0.019 (4)

All H atoms were placed in calculated positions, with C–H = 0.93 (CH) or 0.96 Å (CH<sub>3</sub>), and refined using a riding model, with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$  (CH) or  $1.5U_{\text{eq}}(\text{C})$  (methyl). The methyl groups were allowed to rotate about their local threefold axes.

Data collection: *CAD-4 Software* (Enraf–Nonius, 1989); cell refinement: *CAD-4 Software*; data reduction: *CAD-4 Software*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1997); software used to prepare material for publication: *SHELXL97*.



**Figure 2**

Projection of the crystal structure of (I) along the  $b$  axis. H atoms have been omitted.

The CAD-4 diffractometer and software were accessed *via* the X-ray Structural Centre at Moscow, Russia. The author thanks Dr Olekhovich for providing crystals of (I).

## References

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