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## Structure Reports

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## Key indicators

Single-crystal X-ray study
$T=293 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.002 \AA$
$R$ factor $=0.050$
$w R$ factor $=0.155$
Data-to-parameter ratio $=16.0$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## Spiro-[2-3']oxindole-spiro-[3-3"]oxindole-4[p-methylbenzyl]pyrrolidizine

The oxindole moieties of the title compound, $\mathrm{C}_{29} \mathrm{H}_{25} \mathrm{~N}_{3} \mathrm{O}_{3}$, are planar. The pyrrolidine ring adopts a half-chair conformation. The structure is stabilized by $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ intramolecular interactions. It is remarkable that only one of the two NH groups forms a hydrogen bond.

## Comment

Pyrrolidine compounds are capable of exhibiting antimicrobial and antifungal activity, as reported by Amal Raj et al. (2003). The pyrrolidine moiety occurs in biologically important compounds (Baldwin et al., 1994). Several unusual amino acids which contain pyrrolidine moieties were investigated by Gallazzi et al. (1999). The spiro ring system is a frequently encountered structural motif in many pharmacologically relevant alkaloids (Cordel, 1981). Several optically active pyrrolidines have been used as intermediates in controlled asymmetric synthesis (Suzuki et al., 1994). In view of these important aspects, the crystal structure of the title compound, (I), has been determined.


Fig. 1 shows a displacement ellipsoid plot of the molecule, with the atomic numbering scheme. Selected geometric parameters are given in Table 1.

In the benzene rings of the oxindole systems, the endocyclic angles at C9 and C6 are 120.7 (2) ${ }^{\circ}, 122.1$ (2) ${ }^{\circ}$ and at C 24 and C27 are $122.4(2)^{\circ}, 120.9(2)^{\circ}$. The endocyclic angles at C 10 and C7 are 119.2 (2) ${ }^{\circ}, 117.7$ (2) and at C25 and C28 117.4 (2) ${ }^{\circ}$ and $118.7(2)^{\circ}$. Similar values are observed in related structures (Seshadri et al., 2002).

The bond geometries correlate with a variety of N -phenylsubstituted pyrrolidine-2-one systems (Billing et al., 1991). The bond lengths of the pyrrolidine moiety (Table 1) differ slightly from normal values, but are comparable with those of reported structures (Jeyabharathi et al., 2001; Gzella \& Wrzeciono, 1990). This may be due to steric forces of bulky substituents at the pyrrolidine moiety.

The oxindole moieties are planar. The asymmetry parameters (Nardelli, 1995; $q_{2}=0.4621, \varphi=62.27^{\circ}, \Delta_{s}\left[\mathrm{C} 4=0.0648^{\circ}\right]$

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and $\left.\Delta \mathrm{C}_{2}[\mathrm{C} 34=0.0343]\right)$ reveal a half-chair conformation for the pyrolidine ring $(B)$. The unsubstituted ring $(C)$ of the pyrrolidizine ring system adopts a half-chair conformation. This is confirmed by the puckering parameters $q_{2}=0.3643, \varphi=$ $\left.-89.66^{\circ}, \Delta_{s}[\mathrm{C} 33=0.0878)^{\circ}\right]$ and $\Delta C_{2}[\mathrm{~N} 30=0.0019]$ (Cremer \& Pople, 1975).

In addition to van der Waals interactions, the crystal structure is stabilized by $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ intramolecular hydrogen bonds. It is remarkable that only one of the two NH groups forms a hydrogen bond.

## Experimental

A mixture of (E)-3-p-(methylphenacylidine)oxindole, isatin and pyrroline was refluxed in aqueous methanol for 2-3 h. On completion of the reaction the solvent was evaporated in vacuo and the resulting crude product was purified by column chromatography, using $n$-hexane-ethyl acetate (7:3) as eluent.

Crystal data

$$
\begin{aligned}
& \mathrm{C}_{29} \mathrm{H}_{22} \mathrm{~N}_{3} \mathrm{O}_{3} \\
& M_{r}=463.52 \\
& \text { Triclinic, } P \overline{1} \\
& a=9.7632(6) \AA \\
& b=10.3841(7) \AA \\
& c=11.6874(8) \AA \\
& \alpha=99.735(1)^{\circ} \\
& \beta=96.480(1)^{\circ} \\
& \gamma=98.892(1)^{\circ} \\
& V=1141.98(13) \AA^{\circ}
\end{aligned}
$$

$$
\begin{aligned}
& Z=2 \\
& D_{x}=1.348 \mathrm{Mg} \mathrm{~m}^{-3} \\
& \text { Mo } K \alpha \text { radiation } \\
& \text { Cell parameters from } 3128 \\
& \quad \text { reflections } \\
& \theta=2.6-27.5^{\circ} \\
& \mu=0.09 \mathrm{~mm}^{-1} \\
& T=293(2) \mathrm{K} \\
& \text { Block, colourless } \\
& 0.40 \times 0.30 \times 0.20 \mathrm{~mm}
\end{aligned}
$$

## Data collection

Bruker SMART APEX CCD areadetector diffractometer
$\omega$ scans
Absorption correction: multi-scan (SADABS; Sheldrick, 2001)
$T_{\text {min }}=0.966, T_{\text {max }}=0.983$
7283 measured reflections
5057 independent reflections 4178 reflections with $I>2 \sigma(I)$
$R_{\text {int }}=0.016$
$\theta_{\text {max }}=28.0^{\circ}$
$h=-12 \rightarrow 12$
$k=-13 \rightarrow 5$
$l=-14 \rightarrow 15$

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.050$
$w R\left(F^{2}\right)=0.155$
$S=0.95$
5057 reflections
316 parameters
H -atom parameters constrained


View of (I) (50\% probability displacement ellipsoids).

Table 2
Hydrogen-bonding and short-contact geometry $\left(\AA,{ }^{\circ}\right)$.

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| C11-H11 $\cdots$ O22 | 0.98 | 2.31 | $2.949(2)$ | 122 |
| C34-H34 $\cdots$ O13 | 0.98 | 2.38 | $2.787(2)$ | 104 |
| N23-H23 $\cdots$ O3 $^{\mathrm{i}}$ | 0.86 | 2.05 | $2.892(2)$ | 168 |
| Symmetry code: (i) $1-x, 1-y, 1-z$ |  |  |  |  |

All H atoms were positioned geometrically and allowed to ride on their parent atoms, with $\mathrm{C}-\mathrm{H}=0.93-0.98 \AA$ and $U_{\text {iso }}(\mathrm{H})=1.5$ times $U_{\text {eq }}(\mathrm{C})$ for methyl H atoms and 1.2 times $U_{\text {eq }}(\mathrm{C})$ for other H atoms.

Data collection: SMART (Bruker, 2001); cell refinement: SAINT (Bruker, 2001); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP3 (Farrugia, 1997) and PLATON (Spek, 1990); software used to prepare material for publication: SHELXL97 and PARST (Nardelli, 1995).

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