

## YC-1, an activation inductor of soluble guanylyl cyclase

Jana Sopková-de Oliveira Santos,\* Valérie Collot, Isabelle Bureau and Sylvain Rault

Centre d'Etudes et de Recherche sur le Médicament de Normandie (CERMN),  
Université de Caen, 1 Rue Vaubénard, 14032 Caen, France  
Correspondence e-mail: [sopkova@pharmacie.unicaen.fr](mailto:sopkova@pharmacie.unicaen.fr)

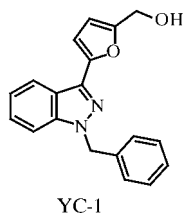
Received 17 March 2000

Accepted 17 May 2000

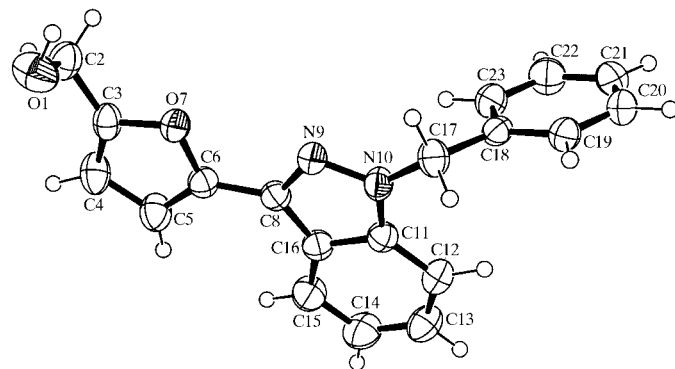
The crystal structure of 1-benzyl-3-(5-hydroxymethyl-2-furyl)indazole,  $C_{19}H_{16}N_2O_2$ , showed that the furan O and indazole N atoms lie on the same face of the molecule. The crystal packing consists of intermolecular hydrogen bonding, and indazole–indazole and indazole–phenyl interactions.

### Comment

The benzylindazole derivative YC-1 [1-benzyl-3-(5-hydroxymethyl-2-furyl)indazole] has been described as an inhibitor of platelet aggregation (Ko *et al.*, 1994; Wu *et al.*, 1995) and smooth muscle cell proliferation (Yu *et al.*, 1995). Recently, it has been demonstrated that YC-1 induces activation of soluble guanylyl cyclase (sGC) probably by binding to an allosteric site which sensitizes the sGC enzyme towards its gaseous activators (NO and CO) by reducing the ligand dissociation rate from the haem group (Friebe & Koesling, 1998; Denninger & Marletta, 1999; Kharitonov *et al.*, 1999; Friebe *et al.*, 1999; Schelvis *et al.*, 1999). Considering YC-1 as a



lead to design new sGC activators with potential therapeutic interest with respect to cardiovascular diseases, we are engaged in a program aiming to synthesize new analogues possessing the same structural features as YC-1 and in particular having the same spatial arrangement of the three putative interaction centres with the enzyme: two N atoms of the indazole ring, the O atom of the furan ring and the O atom of the alcohol group. In order to model these interactions, we needed to determine the three-dimensional structure of a stable conformation of YC-1. The first molecular modelling

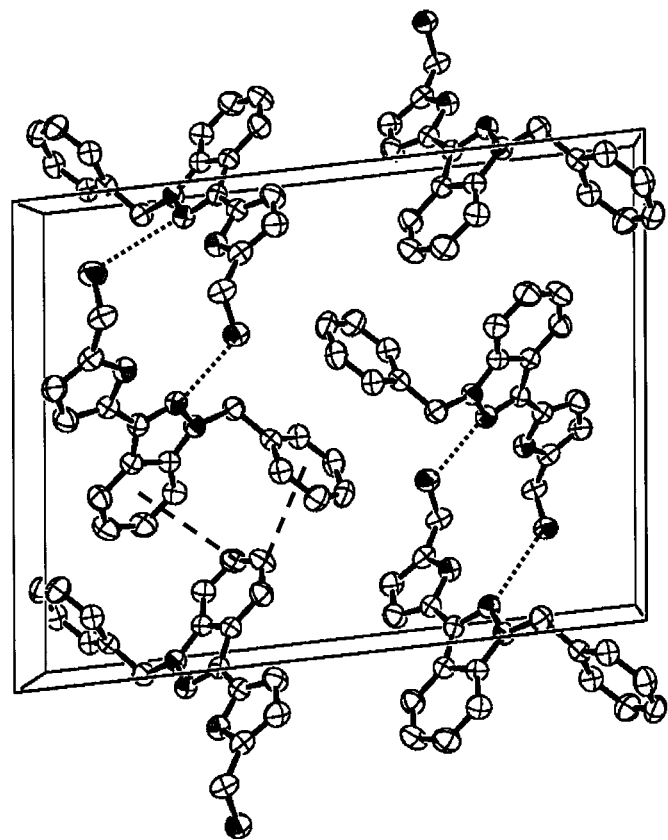


**Figure 1**

View of YC-1 showing the labelling of the non-H atoms. Displacement ellipsoids are shown at 50% probability levels and H atoms are drawn as small circles of arbitrary radii.

trials were unable to define the relative orientation of the furan and indazole rings, *i.e.* if the O atom of the furan lies *cis* or *trans* with respect to the N9 atom of the indazole nucleus with certainty.

In the resulting crystal structure, the furan and indazole rings are quasi-coplanar (Fig. 1). The ring heteroatoms, *i.e.* the furan O (O7) and indazole N atoms (N9, N10), lie on the same face of the molecule, *i.e.* in the *cis* conformation. The angle



**Figure 2**

The crystal packing of YC-1 molecules projected on the *ac* plane. H atoms have been omitted for clarity. Dashed lines indicate the hydrogen bonds and interactions between the aromatic rings.

between the planes of the two rings is about  $5.4 (1)^\circ$ . The third ring, the phenyl of the benzyl group, adopts an approximately perpendicular position with respect to the two rings mentioned above, the angle being  $77.87 (4)^\circ$ .

The crystal packing is stabilized by intermolecular hydrogen bonding and interactions between the aromatic rings of the molecule. An intermolecular hydrogen bond forms between the N9 atom of the indazole ring and the hydroxymethyl O1—H1 group of the furan ring of a neighbouring molecule  $[N9 \cdots H1(\frac{1}{2} - x, -\frac{1}{2} + y, \frac{1}{2} - z)] = 1.93 (3) \text{ \AA}$ ; Table 1 and Fig. 2]. The intermolecular distances observed between the aromatic rings of neighbouring molecules suggest two interactions as the most important: indazole–indazole and indazole–phenyl (Fig. 2), for which a T-shaped arrangement was observed. Such an arrangement has been shown to result in favourable intermolecular attractive forces (Koch & Egert, 1995). For indazole–indazole T-shaped arrangements, the shortest distance corresponds to a C13—H13 $\cdots$ Cg1 interaction, with an H13 $\cdots$ Cg1 distance of  $3.18 \text{ \AA}$  (Cg1 is the centroid of the indazole phenyl ring with symmetry code  $\frac{3}{2} - x, y - \frac{1}{2}, \frac{1}{2} - z$ ). For indazole–phenyl T-shaped arrangements, the shortest distance corresponds to a C14—H14 $\cdots$ Cg2 interaction, with an H14 $\cdots$ Cg2 distance of  $3.07 \text{ \AA}$  (Cg2 is the centroid of the phenyl ring with symmetry code  $\frac{3}{2} - x, \frac{1}{2} + y, \frac{1}{2} - z$ ).

## Experimental

Suitable red transparent crystals of the compound, synthesized according to Collot *et al.* (1999), were grown by slow evaporation from a hexane solution at room temperature.

### Crystal data

$C_{19}H_{16}N_2O_2$	$D_x = 1.331 \text{ Mg m}^{-3}$
$M_r = 304.34$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/n$	Cell parameters from 25 reflections
$a = 14.3450 (10) \text{ \AA}$	$\theta = 8\text{--}14^\circ$
$b = 5.6905 (7) \text{ \AA}$	$\mu = 0.088 \text{ mm}^{-1}$
$c = 18.7140 (10) \text{ \AA}$	$T = 293 (2) \text{ K}$
$\beta = 96.064 (9)^\circ$	Prism, translucent pale red
$V = 1519.1 (2) \text{ \AA}^3$	$0.6 \times 0.4 \times 0.3 \text{ mm}$
$Z = 4$	

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

$D\text{---}H \cdots A$	$D\text{---}H$	$H \cdots A$	$D \cdots A$	$D\text{---}H \cdots A$
O1—H1 $\cdots$ N9 <sup>i</sup>	0.96 (3)	1.93 (3)	2.831 (2)	156 (2)

Symmetry code: (i)  $\frac{1}{2} - x, \frac{1}{2} + y, \frac{1}{2} - z$ .

### Data collection

Enraf–Nonius CAD-4 diffractometer	$\theta_{\text{max}} = 27^\circ$
$\theta/2\theta$ scans	$h = -18 \rightarrow 18$
3412 measured reflections	$k = 0 \rightarrow 7$
3314 independent reflections	$l = 0 \rightarrow 23$
2522 reflections with $I > 2\sigma(I)$	3 standard reflections
$R_{\text{int}} = 0.012$	frequency: 60 min
	intensity decay: 3%

### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0797P)^2 + 0.1614P]$
$R[F^2 > 2\sigma(F^2)] = 0.045$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.140$	$(\Delta/\sigma)_{\text{max}} = 0.002$
$S = 1.060$	$\Delta\rho_{\text{max}} = 0.17 \text{ e \AA}^{-3}$
3314 reflections	$\Delta\rho_{\text{min}} = -0.26 \text{ e \AA}^{-3}$
272 parameters	
All H atoms refined	

All H atoms were found in the difference electron-density map with bond lengths and angles close to well established geometrical criteria [C—H  $0.92 (2)\text{--}1.02 (2) \text{ \AA}$ ]. They were refined with isotropic displacement parameters.

Data collection: *CAD-4-PC* (Enraf–Nonius, 1994); cell refinement: *CAD-4-PC*; data reduction: *Xtal3.2* (Hall *et al.*, 1992); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3* (Farrugia, 1997); software used to prepare material for publication: *SHELXL97*.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: GS1090). Services for accessing these data are described at the back of the journal.

## References

- Collot, V., Dallemagne, P., Bovy, P. R. & Rault, S. (1999). *Tetrahedron*, **55**, 6917–6922.
- Denninger, J. W. & Marletta, M. A. (1999). *B. B. A. Biochim. Biophys. Acta Libr.* **1411**, 334–350.
- Enraf–Nonius (1994). *CAD-4-PC*. Version 1.5. Enraf–Nonius, Delft, The Netherlands.
- Farrugia, L. J. (1997). *J. Appl. Cryst.* **30**, 565.
- Friebe, A. & Koesling, D. (1998). *Mol. Pharmacol.* **53**, 123–127.
- Friebe, A., Russwurm, M., Mergia, E. & Koesling, D. (1999). *Biochemistry*, **38**, 15253–15257.
- Hall, S. R., Flack, H. D. & Stewart, J. M. (1992). Editors. *Xtal3.2 Reference Manual*. Universities of Western Australia, Australia, Geneva, Switzerland, and Maryland, USA.
- Kharitonov, V. G., Sharma, V. S., Magde, D. & Koesling, D. (1999). *Biochemistry*, **38**, 10699–10706.
- Ko, F. N., Wu, C. C., Lee, F. Y. & Teng, C. M. (1994). *Blood*, **84**, 4226–4233.
- Koch, U. & Egert, E. (1995). *J. Comput. Chem.* **16**, 937–944.
- Schelvis, J. P. M., Kim, S., Zhao, Y., Marletta, M. A. & Babcock, G. T. (1999). *J. Am. Chem. Soc.* **121**, 7397–7400.
- Sheldrick, G. M. (1997). *SHELXS97* and *SHELXL97*. University of Göttingen, Germany.
- Wu, C. C., Ko, F. N., Kuo, S. C., Lee, F. Y. & Teng, C. M. (1995). *Br. J. Pharmacol.* **116**, 1973–1978.
- Yu, S. M., Cheng, Z. J., Guh, J. H., Lee, F. Y. & Kuo, S. C. (1995). *Biochem. J.* **306**, 787–792.