

Crystallization and preliminary X-ray analysis of the antimalarial and cytotoxic alkaloid cryptolepine complexed with the DNA fragment d(CCTAGG)<sub>2</sub>J. N. Lisgarten,<sup>a,b</sup> J. Pous,<sup>a</sup>  
M. Coll,<sup>a</sup> C. W. Wright<sup>c</sup> and  
J. Aymami<sup>a,d\*</sup><sup>a</sup>Institut de Biologia Molecular de Barcelona, CSIC, Jordi Girona 18, E-08034 Barcelona, Spain, <sup>b</sup>Department of Crystallography, Birkbeck College, University of London, Malet Street, London WC1E 7HX, England, <sup>c</sup>The School of Pharmacy, University of Bradford, West Yorkshire BD7 4ER, England, and <sup>d</sup>Department d'Enginyeria Química, Universitat Politècnica de Catalunya, Diagonal 647, E-08028 Barcelona, Spain

Correspondence e-mail: aymami@eq.upc.es

Crystals of the indoloquinoline alkaloid cryptolepine complexed with the DNA fragment d(CCTAGG)<sub>2</sub> have been grown by the hanging-drop technique at 293 K using ammonium sulfate as the precipitating agent. Over a period of three weeks, yellow tapering bullet-shaped crystals grew to maximum dimensions of 0.2 × 0.1 × 0.1 mm. The crystals belong to space group *P*6<sub>4</sub>, with unit-cell parameters *a* = *b* = 29.960, *c* = 39.64 Å,  $\alpha = \beta = 90^\circ$ ,  $\gamma = 120^\circ$ , and diffract to 1.4 Å.

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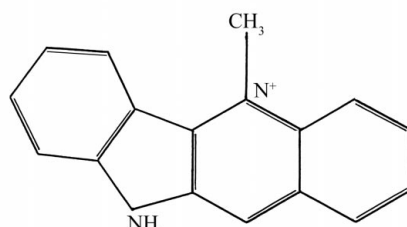
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## 1. Introduction

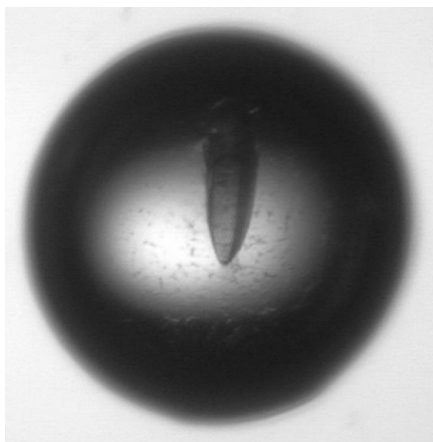
Cryptolepine (Fig. 1) [5-methylquinolo-(2',3',3,2)-indole] is an indoloquinoline alkaloid isolated from the roots of the West African shrub *Cryptolepis sanguinolenta*. This natural product was first isolated from the roots of *C. triangularis* collected in Kisantu (Democratic Republic of Congo). Extracts of the roots of *C. sanguinolenta* are used clinically in Ghana for the treatment of malaria (Boye & Ampofo, 1983). Extracts have also been used as a remedy against colic, as a stomach ulcer tonic and cryptolepine itself has been found to produce a variety of pharmacological effects; these include hypotensive and antipyretic properties (Raymond-Hamet, 1937, 1938; Noamesi & Bamgbose, 1980), presynaptic  $\alpha$ -adrenoreceptor blocking action (Noamesi & Bamgbose, 1982), anti-muscarinic properties (Rauwald *et al.*, 1992), anti-inflammatory properties (Bamgbose & Noamesi, 1981) and antibacterial effects (Boakye-Yiadom & Heman-Ackah, 1979; Paulo, Duarte *et al.*, 1994; Paulo, Pimentel *et al.*, 1994; Cimanga *et al.*, 1996).

Cryptolepine has potent *in vitro* activity against malaria parasites (*Plasmodium falciparum*) (Kirby *et al.*, 1995; Grellier *et al.*, 1996;

Wright *et al.*, 1996; Cimanga *et al.*, 1997) and possesses cytotoxic activity, inhibiting DNA synthesis in B16 melanoma cells (Bonjean *et al.*, 1998). The compound has also been shown to form a complex with haematin in cell-free systems, suggesting that it has a quinine-like mode of action (Wright *et al.*, 1996). An understanding of the way in which cryptolepine combines with DNA will assist in the rational design of antimalarial compounds which inhibit  $\beta$ -haematin formation but which, unlike cryptolepine, do not intercalate into DNA. Derivatives of cryptolepine are currently being evaluated as leads to selective antimalarial agents (Wright *et al.*, 1997). More recently, the results of various studies (Bonjean *et al.*, 1998) have revealed that the alkaloid binds tightly to DNA and behaves as a typical intercalating agent. It was found that the drug interacts preferentially with GC-rich sequences and discriminates against homo-oligomeric runs of A and T. The study also led to the discovery that cryptolepine is a potent topoisomerase II inhibitor and a promising antitumour agent. It stabilizes topoisomerase II–DNA covalent complexes and stimulates the cutting of DNA at a subset of pre-existing topoisomerase II cleavage sites. The intercalating properties of cryptolepine are worthy of investigation as this compound may be a potential lead to new anticancer drugs. The determination of its structure was based on chemical reactions of a degradative nature and on ultraviolet spectroscopy in neutral and basic media. <sup>1</sup>H NMR, <sup>13</sup>C NMR and MS data have confirmed the proposed structure (Dwumabadu *et al.*, 1978; Ablordeppey *et al.*, 1990; Tackie *et al.*, 1991). A recent crystallographic study of the cryptolepine–tetraphenyl borate complex has also confirmed the structure and revealed a unique packing propensity (Wright *et al.*, 1999).



**Figure 1**  
Structure of cryptolepine [5-methylquinolo-(2',3',3,2)-indole].



**Figure 2**  
Crystal of the cryptolepine-(CCTAGG)<sub>2</sub> complex used for data collection.

## 2. Methods and results

Cryptolepine was isolated from the roots of *C. sanguinolenta* as described in detail elsewhere (Wright *et al.*, 1996). Crystallization conditions were screened using the Matrix Screen (Hampton Research, USA) and the Nucleic Acid Mini Screen (Berger *et al.*, 1996). Crystals were grown at 293 K by the hanging-drop vapour-diffusion technique using Linbro multiwell tissue-culture plates. Crystals grew to a maximum size after about three weeks on mixing 0.5  $\mu$ l 5 mM cryptolepine hydrochloride and 0.5  $\mu$ l 3 mM d(CCTAGG)<sub>2</sub> with either 1.0  $\mu$ l of crystallization solution containing 5 mM magnesium acetate, 25 mM MES pH 6.5 and 1.25 M ammonium sulfate or 2  $\mu$ l of crystallization solution containing 5 mM magnesium sulfate, 25 mM sodium cacodylate pH 6.5 and 1 M ammonium sulfate. Yellow bullet-shaped crystals grew to maximum dimensions of 0.2  $\times$  0.1  $\times$  0.1 mm (Fig. 2) and diffracted weakly on an in-house rotating-anode generator. Synchrotron radiation was therefore essential for the success of this

**Table 1**

Summary of the cryptolepine–DNA data set obtained at cryogenic temperature.

Values in parentheses are for the last resolution shell (1.45–1.40 Å).

Unit-cell parameters (Å, °)	$a = b = 29.96$ , $c = 39.647$ , $\alpha = \beta = 90$ , $\gamma = 120$
Space group	$P6_4$
Resolution of data (Å)	1.4
No. of data collected	26108
No. of unique data	3356
Completeness (%)	87.3 (73.2)
$R_{\text{merge}}^{\dagger}$ (%)	3.8 (33)
Mean $\langle I/\sigma(I) \rangle$	29.1 (2.5)

$\dagger R_{\text{merge}}(I) = \sum_h \sum_i |I_i - I| / \sum_h \sum_i I$ , where  $I$  is the mean intensity of  $i$  reflections  $h$ .

project. Crystals were flash-frozen in a stream of evaporating liquid nitrogen at 120 K. Diffraction data were collected at EMBL beamline BW7A (Hamburg, Germany) to 1.4 Å resolution using a MAR Research image-plate detector and a wavelength of 1.1 Å (see Table 1).

Determination of unit-cell parameters and space group and the integration of reflection intensities were performed using *DENZO* (Otwinowski, 1993) and the data were scaled using *SCALEPACK* (Otwinowski, 1993). The structure determination of the cryptolepine–d(CCTAGG)<sub>2</sub> complex is in progress.

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