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1,2-Bis(8-quinolylthiomethyl)benzene

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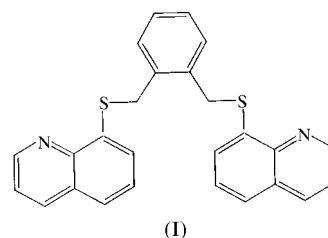
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The title potentially tetradentate *N,S,S,N*-donor ligand, C₂₆H₂₀N₂S₂, has been structurally characterized. The two S atoms adopt a *trans* conformation, lying above and below the benzene ring. The two quinoline rings are planar, with one parallel to the benzene ring and the other nearly perpendicular to it.

Comment

Among the tasks of bioinorganic chemistry, structural and functional modelling of the properties of biocomplexes has a special position. It is well known that in 'blue' or type I copper proteins, the Cu^{II} atom is coordinated by two histidine imidazole N atoms, a cysteine thiolate S atom and a methionine thioether S atom in a distorted tetrahedral geometry (Colman *et al.*, 1978). Biomodelling with the help of copper(II) coordination compounds with sulfur-containing ligands is used at present to explain the structural details of the coordination unit of copper of type I for their low *A* value in ESR spectra, abnormally high intensity absorption bands at *ca* 600 nm and the positive values of the redox potentials (Colman *et al.*, 1978). So far, no models have been obtained having these three properties simultaneously. In particular, the binding of thiolate groups to copper(II) is hard to achieve in low-molecular-weight compounds, the end product being an often insoluble, polymeric copper(I) compound and disulfide (Hughey *et al.*, 1979). So, in some literature, such models are tried with ligands containing thioether groups (Amundsen *et al.*, 1977; Addison *et al.*, 1984; Sakaguchi & Addison, 1979; Dockal *et al.*, 1976; Ritmeijer *et al.*, 1982; Pavlishchuk *et al.*, 1988). We are interested in models of blue copper proteins with low-molecular-weight complexes and report here the crystal structure of a model ligand, namely 1,2-bis(8-quinolylthiomethyl)benzene [or α,α' -bis(8-sulfanylquinoline)-*o*-xylene], (I).



The C—N and C—S distances show no remarkable features, with C—N distances in the range 1.315 (4)–1.374 (3) Å, and C—S distances in the ranges 1.774 (3)–1.757 (3) Å and 1.817 (3)–1.819 (3) Å. The C atoms in the shorter C—S bonds belong to the quinoline moieties and the C atoms in the longer C—S bonds belong to the methylene groups. The C11—C10—S1 angle [107.03 (17)°] is almost the same as C16—C17—S2 [107.78 (18)°], but the two dihedral angles between the two quinoline moieties and the *o*-xylene moiety are obviously not the same. This suggests that the two quinoline moieties can rotate freely about the two C atoms of the methylene groups even though the bridge between the two quinoline moieties is rigid. So, it is possible to form a distorted tetrahedral geometry which is necessary to model blue copper proteins when the ligand coordinates with Cu^{II} ions.

Experimental

To a hot ethanol (60 ml) solution of 8-mercaptoquinoline sodium salt (3.684 g, 10 mmol) was added under nitrogen a solution of α,α' -di-bromo-*o*-xylene (2.64 g, 10 mmol) in acetone (30 ml) with stirring. The mixture was refluxed under nitrogen for 6 h. After cooling to room temperature, the mixture was cooled in a refrigerator to give a white precipitate. The precipitate was extracted with chloroform (15 ml). The white crystals were obtained when acetonitrile (2 ml) was added carefully to a chloroform (2 ml) solution of α,α' -bis(8-sulfanylquinoline)-*o*-xylene (0.085 g, 0.2 mmol), which was then left to stand at room temperature for 2 d; yield 67%. Analysis found: C 73.59, H 4.69, N 6.60%; calculated for C₂₆H₂₀N₂S₂: C 73.55, H 4.75, N 6.60%.

Crystal data

C ₂₆ H ₂₀ N ₂ S ₂	<i>Z</i> = 2
<i>M_r</i> = 424.56	<i>D_x</i> = 1.344 Mg m ⁻³
Triclinic, <i>P</i> $\bar{1}$	Mo <i>K</i> α radiation
<i>a</i> = 9.230 (8) Å	Cell parameters from 26 reflections
<i>b</i> = 10.907 (6) Å	θ = 6–14°
<i>c</i> = 12.056 (12) Å	μ = 0.270 mm ⁻¹
α = 112.68 (6)°	<i>T</i> = 293 (2) K
β = 95.20 (7)°	Block, colourless
γ = 106.05 (6)°	0.54 × 0.35 × 0.24 mm
<i>V</i> = 1049.2 (15) Å ³	

Data collection

Siemens <i>R3m</i> diffractometer	<i>R</i> _{int} = 0.040
ω scans	θ _{max} = 26°
Absorption correction: empirical (North <i>et al.</i> , 1968)	<i>h</i> = 0 → 10
<i>T</i> _{min} = 0.869, <i>T</i> _{max} = 0.918	<i>k</i> = -13 → 12
3961 measured reflections	<i>l</i> = -14 → 14
3949 independent reflections	3 standard reflections
2680 reflections with <i>I</i> > 2 σ (<i>I</i>)	every 97 reflections
	intensity decay: 3%

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.046$
 $wR(F^2) = 0.111$
 $S = 1.030$
 3949 reflections
 271 parameters
 H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0432P)^2 + 0.2725P]$$

where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.18 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.17 \text{ e } \text{\AA}^{-3}$

Data collection: *XSCANS* (Siemens, 1996); cell refinement: *XSCANS*; data reduction: *XSCANS*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); software used to prepare material for publication: *SHELXTL*.

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