Structural and magnetic properties of an asymmetric dicopper(ii) anticancer drug analogue

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The single-crystal X-ray structure and variable temperature magnetic properties of the novel dicopper(II) anticancer drug analogue $[\text{Cu}_2(\text{HL})(\text{H}_2\text{PO}_4)_2]_2][\text{NO}_3]_2.2\text{H}_2\text{O}$ **[H2L = bis(pyridine-2-aldehyde) thiocarbohydrazone] is determined; the complex shows a dimer of dimeric {Cu2(HL)}3+ units with dihydrogenphosphato and sulfur bridges between the inequivalent CuII centres, resulting in three antiferromagnetic exchange interactions: a dominant intradimer** *J* **of** -109 **to** -116 **cm⁻¹, and two weaker interdimer interactions.**

Thiosemicarbazones have been extensively studied because of their biological properties. Cu^{II} is necessary for activity for kethoxal bis(thiosemicarbazone), 1 with the mode of action proposed to involve inhibition of DNA synthesis and oxidative $phosphorylation²$ and *N*-methylisatin- β -thiosemicarbazone can inactivate tumour viruses.3 Attention has however focused on pyridine-2-aldehyde thiosemicarbazone (Hpt) and derivatives as the Cu^{II} complexes are more bioactive than the metal free ligands.2,4 The metal-free 5-OH analogue underwent clinical trials as an anticancer drug but side effects such as disruption of iron metabolism prevented clinical use.5 However, administration of a preformed metal complex may alleviate this. Adducts of $[Cu(pt)]^+$ with sulfur and nitrogen donors have been detected in biological fluids⁶ with stable model thiolato complexes isolated from aqueous solution.7 Crystallographic studies have revealed ternary nitrogen adduct formation,7,8 complexation of dihydrogen phosphate⁹ and pyrophosphate¹⁰ and the system's remarkable ability to form complexes of both the anionic and neutral ligand.¹¹ The H₂L ligand may be considered as an extension of Hpt, now with a possible extra metal binding domain. Antiviral activity has been reported for 2-acetylpyridine thiocarbohydrazones¹² and $H_2\hat{L}$ is antifungal¹³ and cytotoxic14 but little chemical and no structural work has been carried out on the Cu–H2L system. On the human leukemia cell line MOLT4 (at 10 μ m), H₂L is more cytotoxic than Hpt, $Cu₂(HL)³⁺$ and $Cu(pt)⁺$ show equal activity whereas, surprisingly $Cu(HL)$ ⁺ is the most cytotoxic.¹⁴ Here we report the first crystallographic study of H₂L with Cu^{II}, $[\{Cu_2(HL)(H_2-H_1)$ PO_4 ₂ $\left[\frac{1}{2}[NO_3]_2.2H_2O$, giving a dimer of dicopper(II) moieties with H_2PO_4 ⁻ and weak sulfur bridges. Variable temperature magnetic studies reveal three antiferromagnetic exchange interactions involving superexchange pathways across the planar ligand, a three-atom H_2PO_4 - bridge and direct $\text{Cu}^{\text{II}}\text{...}\text{Cu}^{\text{II}}$ and/or out-of-plane (*via* S) interactions.

The starting nitrate complex, $\left[\text{Cu}_2\text{L}(\text{NO}_3)_2\right]$ 3H₂O **1**, used in subsequent metathetical reactions with phosphate was prepared by addition of H_2L (0.616 g, 2.16 mmol) in hot ethanol (150 ml)

to $Cu(NO₃)₂·3H₂O$ (1.05 g, 4.34 mmol) in the same solvent (10 ml). After stirring for 1 h the product was filtered, washed with ethanol then dried *in vacuo* (913 mg, 72%). $[\text{Cu}_2(\text{HL})$ - $(H_2PO_4)_2$][NO_3]₂·2H₂O 2 was prepared by dissolving 1 (157) mg, 0.267 mmol) and $NaH_2PO_4·H_2O$ (37 mg, 0.268 mmol) in H3PO4 (2 ml, 2 m). Dark green crystals suitable for X-ray analysis were collected after 20 days (82 mg, 45%).

 X -Ray analysis of 2^{\dagger} shows the structure (Fig. 1) to be best described as a centrosymmetric dimer of dicopper(ii) moieties with dihydrogenphosphato and weak sulfur bridges. Cu(1) has a square-pyramidal geometry with the monoanionic HL bound through pyridyl and imine nitrogens and sulfur. The coordination sphere is completed by phosphato oxygens O(11) and O(21). Cu(2) adopts a tetragonal '4 $+$ 2' geometry with pyridyl, imine and deprotonated amide nitrogens from HL and phosphato oxygen in the plane. Much weaker axial interactions to $\rm O(11)$ and $\rm S(1A)$ 3.24 Å are from symmetry related molecules. The dimeric $Cu_2(HL)^{3+}$ units are linked *via* three-atom phosphato $[Cu(1)-O(21)-P(2)-O(1A)-Cu(2)]$ and one-atom sulfur bridges with adjacent dimers connected by the weak Cu(2)–O(11) interaction and a hydrogen-bonding network involving phosphato, water and nitrate oxygens and protonated amide [N(3)] nitrogen. Other metric parameters may be considered normal.7–12 The two phosphate ions adopt different coordination modes, with $P(2)$ chelating and $\overline{P}(1)$ being monodentate and P(2) binds through the keto oxygen $[P(2)-O(1)$ 1.468(4) Å].

Fig. 1 Molecular structure of the dimeric dicopper(II) ion $[\{Cu_2(HL)(H_2PO_4)_2\}_2]^{2+}$ of 2. Hydrogen atoms are omitted for clarity. Selected bond distances (\AA) and angles (°): Cu(1)–N(1) 2.053(4), Cu(1)– N(2) 1.954(4), Cu(1)–N(4) 2.076(4), Cu(1)–O(11) 1.933(3), Cu(1)–O(21) 2.159(3), Cu(2)–N(5) 1.966(3), Cu(2)–N(6) 2.004(4), Cu(2)–O(1) 1.890(3), Cu(2)–S(1) 2.263(2), Cu(2)–O(11C) 2.860(4), Cu(2)–S(1A) 3.242(2), Cu(1)···Cu(2) 4.996, Cu(1)···Cu(2A) 5.076, Cu(2)···Cu(2A) 3.905, Cu(2)···Cu(2C) 3.797, Cu(1)···Cu(1A) 9.326; O(11)–Cu(1)–N(1) 95.4(2), N(4)–Cu(1)–O(11) 140.5(1), O(21)–Cu(1)–N(1) 97.4(2), O(21)–Cu(1)– O(11) 95.0(1), N(1)–Cu(1)–N(4) 155.1(1), N(2)–Cu(1)–O(11) 160.5(2), O(1)–Cu(2)–N(6) 92.8(2), S(1)–Cu(2)–O(1) 100.7(1), S(1)–Cu(2)–N(6) 165.4(1), S(1)–Cu(2)–N(5) 84.9(1), O(1)–Cu(2)–N(5) 171.9(2), $O(11)$ –Cu(2)–S(1) 86.6(1), $O(11)$ –Cu(2)– $O(1)$ 107.4(1), $O(11)$ –Cu(2)– N(5) 78.6(1), O(11)–Cu(2)–S(1A) 159.2(2), S(1A)–Cu(2)–S(1) 94.7(1), $S(1A)$ –Cu(2)–O(1) 92.8(1).

Fig. 2 Temperature dependence of χ_{Cu} and μ_{Cu} *vs. T* for [{Cu₂(HL)(H₂-PO4)2}2](NO3)2·2H2O **2**. Solid lines represent the best fit calculated values.

The temperature dependence of magnetic susceptibility and moment, per Cu^{II}, for $\overline{2}$ is shown in Fig. 2. A maximum in χ at *ca*. 180 K is clearly indicative of medium strength antiferromagnetic exchange occurring. The rapid increase in χ at low temperature is due to monomer impurity. Use of a simple dinuclear Bleaney–Bowers model¹⁵ $(-2 JS_1S_2$ Hamiltonian) did not precisely reproduce the shape of the susceptibility in the region of χ_{max} but did yield an approximate J_{12} value of *ca*. -105 cm⁻¹, albeit with a low *g* value of 1.98 and a large interdimer θ parameter (in *T*– θ) of –25 K. Excellent fits could be obtained when the tetranuclear model of Hatfield¹⁶ was employed. The rhomboidal framework of the dimer of dimers is shown in Fig. 3 together with the *J* labels. The value of the long $Cu(1)\cdots Cu(1)$ parameter $J_{1,1A}$ was set at zero and the other parameters were varied widely for best-fit. It was found that two parameter sets gave equally excellent fits *i.e.* set A: $g = 2.00 \pm$ 0.02 , $N_\alpha = (60 \pm 3) \times 10^{-6}$ cm³ mol⁻¹, $J_{2,2A} = -75.4 \pm 0.2$ cm⁻¹, $J_{1,2} = -116.6 \pm 0.2$ cm⁻¹, $J_{1,2A} = -16.8 \pm 0.2$ cm⁻¹, % monomer 1.7 ± 0.1 . This fit is shown in Fig. 2. Set B: $g = 2.00 \pm 0.02$, $J_{2,2A} = -46.0 \pm 0.5$ cm⁻¹, $J_{1,2} = -109.5 \pm 0.02$ 0.2 cm⁻¹, $J_{1,2A} = -47.9 \pm 0.2$ cm⁻¹, % monomer 1.7 ± 0.1 . Thus the dominant coupling, $J_{1,2}$, remains virtually constant in both fits and occurs across the planar thiocarbohydrazonato moiety. It involves superexchange pathways in the *x*–*y* plane such as a two-atom *trans*-Cu–N–N–Cu pathway and three- or four-atom pathways Cu–S–C–N–Cu or Cu–S–C–N–N–Cu. The value of $J_{1,2}$ is similar to that of a structurally related carbohydrazonato dinuclear analogue, reported recently17 having $J = -106.6$ cm⁻¹ (there is a factor of two error in *J* in ref. 17). The size of the 'edge' and 'short-diagonal' parameters $J_{1,2A}$ and $J_{2,2A}$ could not be identified unambiguously. When using set B, but with $J_{2,2A}$ set at zero or slightly positive, {as anticipated from a previous study18 on a related S-bridged complex $[CuCl(S_2CNEt)]_4$ } much poorer fits were obtained, with the calculated and observed curves crossing each other in the region of χ_{max} . Thus it appears that the Cu(2)S(1)-Cu(2A)S(1A) pathways yield stronger antiferromagnetic coupling (minimum of -46 cm⁻¹) than do related Cu($\bar{S}R$)CuCl or $Cu(SR)Cu(SR)$ pathways¹⁸ in $[CuCl(S₂CNEt)]₄$ and $[Cu(S₂Cr)]₂$ $NEt)_{2}]_{2}$. The antiferromagnetic coupling across the three-atom H_2PO_4 ⁻ bridge, with *J* of at least -16.8 cm⁻¹, is weak as

Fig. 3 Rhomboidal framework for the CuII centres of **2** showing distances (\AA) together with the *J* labels.

anticipated and involves axial–equatorial d-orbital overlap, respectively on $Cu(1)$ and $Cu(2\overline{A})$. The uncertainties and correlation in the $J_{1,2A}$ parameters may also be influenced by the weak axial interaction $Cu(2)-O(11)$ and $Cu(2)-S$ occurring between teramers, although these are expected to be very weak. In this regard, the visible spectra of solid and H_3PO_4 solutions of **2** are nearly identical.

The X-band ESR spectrum of a neat powder of **2** at 298 K shows a broad, symmetrical signal at $g = 2.1$. At 77 K this signal is resolved into a typical axial lineshipe $(g_{||} = 2.19,$ A_{\parallel} = 184.6 gauss, g_{\perp} = 2.00) probably due to the monomer impurity (vide supra). Triplet state lines due to Cu^{II}...Cu^{II} interactions19 are evident as weak broad lines at 'half-field' (1500 G) and 3700 G. The same spectrum is observed in frozen 2 m H_3 PO₄ solution, with evidence for a weaker monomer lineshape ($g_{\parallel} = 2.33$, $A_{\parallel} = 175$ G, $g_{\perp} = 2.06$) superimposed.

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Notes and References

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 $\frac{1}{7}$ *Crystal data*: C₁₃H₁₇Cu₂N₇O₁₂P₂S, *M* = 684.4, dark green crystal, 0.15 \times 0.1 \times 0.1 mm, triclinic, space group *P* $\overline{1}$, *a* = 8.604(2), *b* = 10.719(2), $c = 14.268(3)$ Å, $\alpha = 109.57(3)$, $\beta = 90.11(3)$, $\gamma = 110.62(3)$ °, $U = 1149.6(4)$ \AA^3 , $D_c = 1.977$ g cm⁻³, $Z = 2$, $F(000) = 688$, μ (Mo- $K\alpha$) = 2.157 mm⁻¹; \overline{R} = 0.041, $\overline{R_w}$ = 0.122 using 3150 unique reflections with $I > 4\sigma(I)$. CCDC 182/711.

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