A Crystalline Cobalt(II)corrinate derived from Vitamin B₁₂: Preparation and X-Ray Crystal Structure

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The preparation and X-ray crystal structure of a monomeric cobalt(n)corrinate derived from vitamin B₁₂ is reported.

Detailed information on the structure of cobalt(II) corrinates is of interest, particularly in the context of studies on the coenzyme B₁₂ catalysed enzymatic reactions, where cob(II)alamin (1) has been detected as a reaction intermediate.¹ Hitherto, however, only one specimen of that class of Co(II) complexes has been studied by X-ray analysis,² the unusual, diamagnetic, iodonium bridged, Co(II) corrin dimer (2), prepared by Werthemann in Eschenmoser's laboratory.³ We report here the preparation and X-ray structural analysis of a related monomeric Co(II) corrinate derived from vitamin B₁₂.

Formic acid reduction⁴ of a crystalline sample of the vitamin B_{12} methanolysis product 'cobester' (**3**, dicyano-heptamethyl-Co(II)cobyrinate)³ cleanly furnished 'Co(II)cobester' (**4**, heptamethyl-Co(II) cobyrinate perchlorate).^{†5} Crystals of (**4**)

[†] The reaction mixture⁴ was dried at room temperature, taken up in toluene and shaken with 0.1 M aq. NaClO₄ under N₂. The crude (4) was recrystallised from toluene (*ca.* 70% yield). U.v.-visible and c.d. spectra of crystallised (4) were similar to ones published previously;⁵ the e.s.r. spectrum (CHCl₃: benzene 2:1; 77 K) indicated two Co(1) species (*ca.* 9:1, possibly co-ordination isomers). In CDCl₃ solution (4) has $\mu = 2.6$ B.M. (*ca.* 300K; Evans–Scheffold method⁶). The i.r. spectrum (CHCl₃) indicated ClO₄ co-ordination (bands at 1050 and 627 cm⁻¹).⁷



Figure 1. Structure of the cobalt(π)cobyrinate (4), as determined from the X-ray analysis (thermal spheres at 50% probability).



suitable for X-ray analysis \ddagger were obtained by recrystallising the crude Co(II) cobyrinate (4) twice from deaerated solutions in toluene.

In the Co(II) cobyrinate (4), the corrin ligand is identical in constitution and configuration with that in the parent cobalt(III) complex (3), and also shows a very similar geometry.⁸ The cobalt-nitrogen bond lengths in (4) [Co-N(21) 1.89, Co-N(22) 1.93, Co-N(23) 1.91, Co-N(24) 1.87 Å (av. 1.90 Å)] are not significantly different from the ones in (3) (av. 1.91 Å)⁸ [or in (2) (1.92), vitamin B₁₂ (1.90), or coenzyme B₁₂ (1.91 Å)²]. The structure of the corrin ligand in (4) differs [as does the one in (3)] from that of the Co(III) corrin vitamin B₁₂, particularly with respect to the 'upward folding' about the Co-C(10) axis (6° in (4), 18° in vitamin B₁₂)^{2b} and in the peripheral torsion angles, *e.g.* of rings B and C (-30.4 and -30.0°, respectively in (4), -21 and -26°, respectively in vitamin B₁₂^{2b}), affecting the orientation of the peripheral substituents.

A perchlorate ligand is directly bound via oxygen to a five-co-ordinate Co(II) centre§ of the Co(II) cobyrinate (4) (see Figure 1). The axial Co-O(2) bond in (4) (2.31 Å) is considerably longer than the one in aquo-Co(III) corrins (2.06 Å in Co_{β} -aquo- Co_{α} -cyanocobyrate^{2b}). Co-ordination of the ClO₄ group is accompanied by only a 0.048 Å movement of the Co(II) centre out of the best plane through the corrin nitrogen atoms towards the apical ligand [nitrogen root mean square deviation from plane: 0.075 Å (tetrahedral distortion)]. This is less than the out of plane displacement of the five-co-ordinate cobalt centre in (2) (0.12 Å)² or in several Co(II)porphinoids (average displacement 0.14 Å).9 The Co-O-Cl fragment lies close to the plane through Co and C(10)and vertical to the corrin ligand, but the Co-O bond is tilted from the normal by 9.8° towards C(10). The Co-O-Cl bond angle (133.5°) resembles that in other Co(II) perchlorate complexes.⁷ In crystalline (4) the perchlorate ligand is bound to the β side of the Co(II) corrin, thus to the face opposite the one to which the nucleotide is presumably bound in cob(II)alamin (1).

The structure of the Co(II) corrin (4) differs from that of Co(III) corrins^{2b} mainly by a reduction of the co-ordination number at the cobalt centre (from 6 to 5)§ and by an

[‡] Crystal data for (4): C₅₂H₇₃N₄O₁₈ClCo, M = 1136.5 space group $P2_12_12_1$, a = 13.215(13), b = 15.707(14), c = 25.484(23) Å, V = 5289.7(17) Å³, Z = 4, $D_c = 1.427$ g cm⁻¹, F(000) = 2404, 3914 unique reflections [2988 with $F > 4\sigma(F)$], R = 0.085 ($R_w = 0.081$) for 608 parameters and 3061 observables; data were collected on a Stoe 4-circle diffractometer (λ (Mo K_{α}) = 0.71069 Å) at 98 K for all independent reflections with 3° ≤ 20 ≤ 45°. Atomic co-ordinates, bond lengths and angles, and thermal parameters have been deposited at the Cambridge Crystallographic Data Centre. See Notice to Authors, Issue No. 1.

[§] The ester-O(49) of the symmetry equivalent molecule $(\frac{1}{2} + x, \frac{1}{2} - y, -z)$ occupies the sixth co-ordination site with a Co–O distance of 3.56 Å (in crystalline (2)² an ester methyl group is found at Co–C = 4.12 Å).

elongation of the remaining axial bond. The geometries of the corrin ligand in (4) and of its binding of the cobalt centre differ but little, particularly when compared to that in (3). The transition from Co(III) to Co(II) corrin (and *vice versa*) thus appears to concern mainly the geometry of the axial ligation, but to have little effect on the residual cobalt–corrin portion. Such a conservation of the geometry of the cobalt–corrin fragment would presumably be a factor in support of particularly low activation barriers in radical bond forming and cleavage reactions at cobalt, relevant to the coenzyme B₁₂ biochemistry.¹⁰ The long axial bond in (4) points to a structural feature of Co(II) cobyrinates that also might be taken into account when considering the modes of activation¹⁰ in the B₁₂-catalysed enzymatic radical reactions.

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