

Novel Hydroperoxidation of the α -Position in Pyridoxylidene-Amino Acid Schiff Base–Cobalt Complex by Hydrogen Peroxide

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The reaction of pyridoxylidene-L-phenylalanine with Co^{II} ion in the presence of H_2O_2 generates a novel Co^{III} ternary complex, $[\text{Co}(\text{pyridoxylidene-}\alpha\text{-hydroxyphenylalaninato})(\text{pyridoxylidene-}\alpha\text{-hydroperoxyphenylalaninato})]^-$, which has been characterized on the basis of the analysis of its ^1H NMR, positive-ion FAB MS and X-ray structure.

Amino acid transformations catalysed by pyridoxal coenzyme have been non-enzymatically mimicked by various systems of metal ions and pyridoxylidene-amino acid Schiff bases.^{1–5} The metal ions act as the centre for the fixation of Schiff base intermediate and the electronic activation of the α -position of the amino acid moiety through the aldimine–ketimine equilibrium.¹ Recently we reported a novel transformation of an amino acid mediated by Co^{III} ion under aerobic conditions, in which the α -position of pyridoxylidene-amino acid was hydroxylated by oxygen, species which originated from molecular oxygen to generate the $[\text{Co}(\text{pyridoxylidene-}\alpha\text{-hydroxyamino acidato})_2]$.⁶ In further studies of such an amino acid transformation, an attractive Co^{III} ternary complex with pyridoxylidene- α -hydroperoxyamino and pyridoxylidene- α -hydroxyamino acids has been synthesized by the reaction of pyridoxylidene-L-amino acid Schiff base and Co^{II} ion in the presence of H_2O_2 . The resulting complex has been characterized on the basis of the analysis of its ^1H NMR, positive-ion FAB mass spectra and X-ray structure. This is a novel type of hydroperoxidation of pyridoxylidene-amino acid.

To 150 ml of a yellowish aqueous solution of pyridoxylidene-L-phenylalanine Schiff base (PFS) [prepared from L-phenylalanine (1.65 g, 10 mmol), pyridoxal hydrochloride (2.04 g, 10 mmol) and NaOH (0.40 g, 10 mmol)] was added $\text{Co}(\text{NO}_3)_2 \cdot 6\text{H}_2\text{O}$ (1.46 g, 5 mmol), and then 10 ml of 30% H_2O_2 solution was added dropwise at room temperature under nitrogen atmosphere. The resulting mixture was stirred overnight at pH 4 to give a dark-brown solution, and the solution was passed through a cation exchange column with SP-Sephadex C-25 resin (H^+ form, 3.6 cm ϕ \times 25 cm) pretreated to pH 3–4 with acetic acid. After the elution of neutral and anionic compounds with water, a single adsorbed band was fractionated with a 1 mol dm^{-3} NaCl solution, and the fraction was allowed to stand for several weeks to give single crystals of **1** suitable for X-ray structure analysis. The electronic absorption spectrum of **1**, showing λ_{max} at 515 nm with a shoulder at 655 nm, reveals that the Co^{II} ion has been oxidized to Co^{III} species.

The ^1H NMR spectrum of **1** showed the absence of any peak assignable to the α -proton of phenylalanine of the PFS ligand in the spectral range usually expected for the Schiff base complex (δ ca. 5),⁶ although all other proton peaks in **1** were assignable to the Schiff base complex. In order to elucidate the structural details of **1**, we tried to analyse its X-ray structure. The crystal structure established (Fig. 1) demonstrated that the two Schiff base ligands coordinated to the Co^{III} ion in *mer-trans*(*N*) configuration.[†] Judging from the double bond character of the azomethine bond [$\text{C}(4\text{M})=\text{N}(7)$ 1.33(1), 1.27(1) Å], the coordination structures of these two Schiff base ligands are both in aldimine form, this is also supported by the electronic absorption spectrum of **1**, being similar to that of the binary Co^{III} complex of PFS with a typical aldimine form.⁶

The most striking structural feature of **1** is found at the two α -positions of the phenylalanine moiety. The positions, as suggested from the ^1H NMR spectrum, have already been substituted by some other groups, whose difference Fourier map showed one and two larger peaks at the respective positions. These substituents were assigned to hydroxy and hydroperoxy groups from the magnitude of electron density of

the located peaks and their bond lengths; the C–O bond in the former was 1.41(1) Å and the C–O and O–O bonds in the latter were 1.40(1) and 1.44(1) Å, respectively. The length of the diatomic bond characterized as O–O is consistent with that reported for alkyhydroperoxide [1.477(3) Å].⁷ The structure of **1** was also confirmed by its positive-ion FAB mass spectrum; the parent ion peak found at $m/z = 733$ was assignable to $[\text{Co}(\text{pyridoxylidene-}\alpha\text{-hydroxyphenylalaninato})(\text{pyridoxylidene-}\alpha\text{-hydroperoxyphenylalaninato})]^-$ ion ($\{\text{H}[\text{Co}(\text{pfs})(\text{pohfs})] + \text{H}\}^+$).

Complex **1** was synthesized by the reaction of PFS with Co^{II} ion in the presence of H_2O_2 under a nitrogen atmosphere. In the absence of H_2O_2 , however, a Co^{II} binary complex with pyridoxylidene-phenylalanine was produced (positive-ion FAB mass spectrum $m/z = 683$). The reaction of the binary Co^{III} complex, $[\text{Co}(\text{pfs})_2]^-$, with H_2O_2 did not give such a peroxidation product. Therefore, the formation mechanism of complex **1** is clearly different from that of the $[\text{Co}(\text{pfs})_2]^-$ complex previously reported.⁶ The present complex **1** might have been formed by the attack of OH and/or OOH radical species on the α -position of PFS, because the reaction of H_2O_2 with Co^{II} ion generates the OH and OOH radicals with oxidation to Co^{III} ion.⁸ Interestingly, in the positive-ion FAB mass spectrum of the reaction product, the peak assignable to the bis(hydroperoxidized ligand) complex, $[\text{Co}(\text{pohfs})_2]^-$, was not detected, which may suggest that such a complex is unstable for isolation or that one of the two hydroperoxy

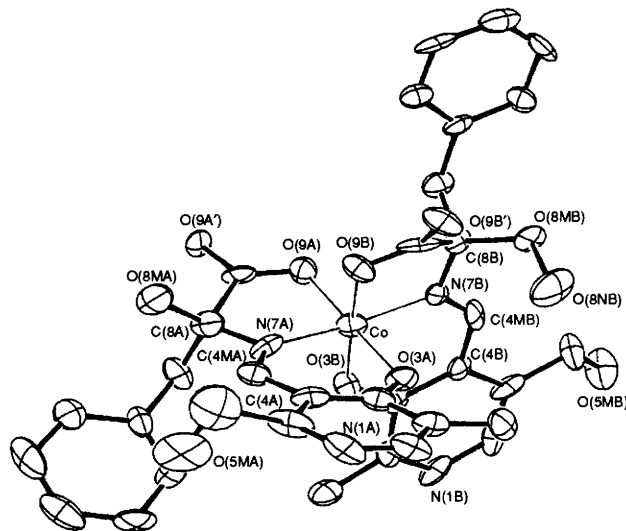
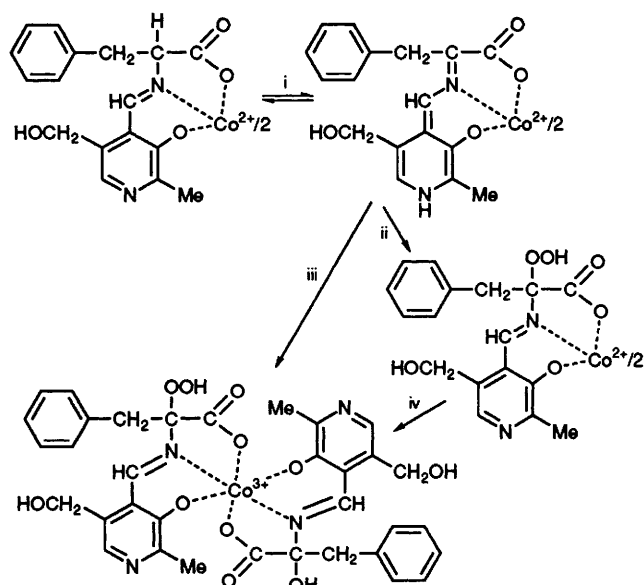


Fig. 1 Molecular structure of the complex anion, $[\text{Co}(\text{pfs})(\text{pohfs})]^-$. Only the important interatomic distances (Å) and angles ($^\circ$) are given: Co–O(3A) 1.880(5), Co–O(3B) 1.882(5), Co–N(7A) 1.858(7), Co–N(7B) 1.886(6), Co–O(9A) 1.886(6), Co–O(9B) 1.890(6), C(4MA)–N(7A) 1.33(1), C(4MB)–N(7B) 1.27(1), C(8A)–N(7A) 1.50(1), C(8B)–N(7B) 1.47(1), C(8A)–O(8MA) 1.41(1), C(8B)–O(8MB) 1.40(1), O(8BM)–O(8BN) 1.44(1), O(3A)–Co–N(7A) 94.9(3), O(3B)–Co–N(7B) 94.7(2), O(3A)–Co–O(9A) 178.7(2), O(3B)–Co–O(9B) 176.2(2), N(7A)–Co–O(9A) 86.2(3), N(7B)–Co–O(9B) 85.8(3), C(4A)–C(4MA)–N(7A) 120.5(8), C(4B)–C(4MB)–N(7B) 124.7(8), C(4MA)–N(7A)–C(8A) 119.4(7), C(4MB)–N(7B)–C(8B) 121.2(7).



Scheme 1 i, aldimine–ketimine equilibrium; ii, H_2O_2 attack at α -position; iii, OH, OOH radical attack at α -position; iv, intramolecular oxidation

groups initially substituted has been supplied by the oxidation of the central metal ion.

The present α -hydroperoxidation of pyridoxylidene-amino acid, to our knowledge, is a novel type of fixation of active oxygen species onto the Schiff base ligand. Such a fixation does not occur in the reaction of Co^{II} ion with the analogous Schiff bases containing salicylaldehyde in place of pyridoxal. Pyridoxal may play an important role for the hydroxylation and hydroperoxidation through activation of the C–H bond at the α -position of the amino acid, which is of interest in connection with a model for the enzymatic transformation of amino acid.^{9,10} From the above results, we presume that, at the initial stage of this reaction, peroxide species generated from hydrogen peroxide would attack at the α -position of the Schiff base, which is in equilibrium between aldimine and ketimine forms (Scheme 1). Further details of the mechanism will be reported elsewhere.

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Footnote

† Crystal data for 1: $\text{C}_{34}\text{H}_{32}\text{CoN}_4\text{NaO}_{11} \cdot 5\text{H}_2\text{O}$, $M = 844.65$, monoclinic, $P2_1/c$, $a = 11.505(3)$, $b = 22.266(4)$, $c = 14.732(3)$ Å, $\beta = 94.27(2)^\circ$, $Z = 4$, $V = 3763.4$ Å³, $D_c = 1.49$ g cm⁻³; Mo-K α ($\lambda = 0.71073$ Å); $\mu = 5.95$ cm⁻¹, $3 < 2\theta < 55^\circ$. A total of 6352 unique reflections were collected on an Enraf-Nonius CAD4-EXPRESS four-circle diffractometer. 2466 reflections having $I > 3\sigma(I_0)$ were used in the structure solution and refinement using the SDP-MoLEN program system. Absorption correction was applied by DIFABS. The structure was solved by the heavy-atom method and refined anisotropically. The hydrogen atoms, except those of hydroxy, hydroperoxy, and a part of water molecules, were located on Fourier difference maps, which were not refined, but included in the calculation. Final R and R_w factors were 0.063 and 0.069, respectively. Atomic coordinates, bond lengths and angles, and thermal parameters have been deposited at the Cambridge Crystallographic Data Centre. See Information for Authors, Issue No. 1.

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