

Table I. Reaction of **1** + O₂ → **2**^a

expt	[1] × 10 ⁻² M	pH	t, h	% 2 (based on 1) ^b
1	1.02	1.0	1	5
	1.02	1.0	20	5
2	1.02	4.0	20	5
	3.07	7.0	20	6
4	1.03	7.0	1	4
	1.03	7.0	2	5
5	1.03	7.0	20	6
	0.77	9.0	20	48
6	1.03	9.0	0.25	20
	1.03	9.0	2.5	39
7	1.03	9.0	20	50
	1.53	9.0	20	61
8	3.07	9.0	20	66
	9	1.53	10.0	1.5
20			70	

^a Reaction was run at 25 °C in H₂O using 1 atm of O₂. ^b Determined by monitoring λ = 6220 Å. Since **2** is slowly oxidized by O₂, visible spectra samples were prepared by acidification with deoxygenated 0.01 M HTFA and treatment with zinc amalgam for 30 min.

deprotonation of a hydroxy ligand upon oxidation of [(bpy)₂pyRuOH]²⁺ to [(bpy)₂pyRu=O]²⁺ serves dramatically to lower the Ru(IV)/Ru(III) potential in this complex (+0.99 V vs. SCE) as compared with a similar complex which does not possess ligands which can deprotonate, e.g., [(bpy)₂-RuCl₂]²⁺, whose Ru(IV)/Ru(III) potential is +1.98 V vs. SCE.¹⁶

At the time **1** is oxidized by O₂ to **3** or **4**, 44% H₂O₂ based on **1** can be detected.¹⁷ Subsequently, **2** is formed slowly. The amount of **2** increases with higher pH (expt 4–9), concentration of **1** (expt 5–8), and time (expt 4, 6, 9). The dramatic difference between the results obtained at low and high pH can be understood in terms of Scheme I.

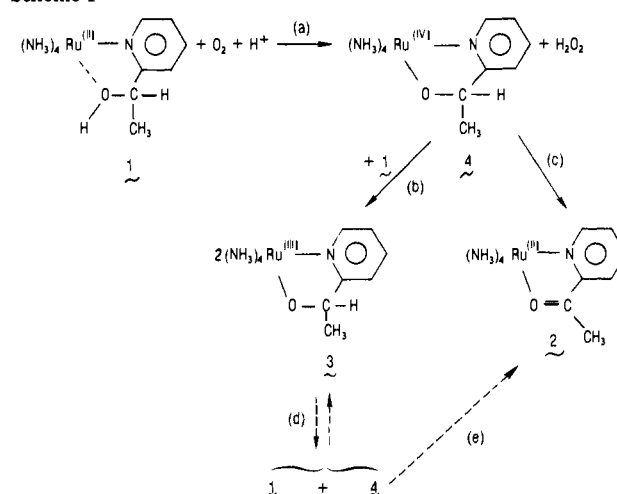
In acidic solution, reaction of **1** with O₂ forms **4** and H₂O₂ (step a).¹⁸ Complex **4** can then react in one of two ways: (1) with a molecule of **1** to form two molecules of **3** (step b), or (2) internal alcohol oxidation to form **2** (step c).¹⁹

In acidic solution, **3** is the predominant product (Table I) and is stable under these conditions. However, in basic solution, **3** disproportionates to give **1** and **4** (step d). Complex **1** can react with O₂ (step a) and reenter the cycle. Complex **4** can undergo internal redox reaction to give **2** (step e). Thus, in basic solution, **3** can be recycled because of disproportionation which facilitates formation of **2** in high yield.

Scheme I is supported by several facts. The dramatic effect of pH on the yield of **2** (Table I) reflects the fact that disproportionation (step d) of **3** to **4** (precursor to **2**) is pH dependent.⁸ Furthermore, the observed increase in the yield of **2** with the initial concentration of **1** and the observed second-order kinetics for disproportionation of similar Ru(III) complexes⁸ is again consistent with this scheme. This work supports the suggestion of Diamond et al.¹ concerning the intermediacy of Ru(IV) in the oxidation of coordinated primary and secondary amines. A similar reaction scheme can be invoked to explain the observations of Guengerich et al.⁵ on the conversion of Ru(III)-imine complexes into Ru(II)-nitrile complexes in the presence and absence of O₂.

These results clearly demonstrate that, when a two-electron oxidative pathway is provided (i.e., Ru(IV) → Ru(II)), the alcohol oxidation proceeds smoothly. On the other hand, the inability of Ru(III) complexes to form **2** suggests that a one-electron path is not feasible. The chemistry of the copper center in galactose oxidase as discussed by Hamilton²⁰ exhibits many features similar to those of this ruthenium system. In galactose oxidase a high-valent species, Cu(III) (compare with Ru(IV)

Scheme I



^a Solid lines refer to reactions taking place at low pH; broken lines represent additional reactions taking place at high pH.

complex, **4**), performs a two-electron oxidation of a primary alcohol, giving the corresponding aldehyde (compare with 2-acetylpyridine in Scheme I) and Cu(I) (compare with Ru(II) complex, **1**). A two-electron reduction of O₂ regenerates Cu(III) with the concomitant formation of H₂O₂ (compare step a in Scheme I). In both systems intermediate oxidation state complexes, i.e., Cu(II) in galactose oxidase or Ru(III) in **3**, are inactive for reaction with O₂ or alcohol. Although E° values for complexes **1**, **3**, or **4** are not known, the reactivities discussed in this work suggest that E°_{4/1} ≥ E°_{3/1} and E°_{3/1} ≤ 0.30 V. These estimates roughly coincide with the copper counterparts in the galactose oxidase system.²⁰

It is our intention to apply the principals discussed in this work to the generation of H₂O₂ and potent two-electron, specific oxidants from other metal complexes and O₂.

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References and Notes

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- Formed in reaction of (NH₃)₅Ru²⁺-OH₂ and 2-(1'-hydroxyethyl)pyridine at pH 5. In acidic solution the alcohol group rapidly displaces an NH₃ forming complex **1**.
- DeF. P. Rudd and H. Taube, *Inorg. Chem.*, **10**, 1543 (1971).
- Upon admission of O₂ to solutions of **1**, all resonances in the NMR become very broad. This is likely due to the formation of paramagnetic Ru(III) and Ru(IV) (one and two unpaired electrons, respectively) which serve to broaden the resonances of diamagnetic species such as **1** and **2**. For this reason, quantitative integration of the appearance of **2** is not possible.
- Prepared as described in ref 8.
- Although **3** does not have any distinctive visible characteristics, its presence is verified as follows. After a reaction for 2 h at pH 1 with O₂, the solution is degassed and the pH raised to 11. The rate and amount of **2** formed (correcting for 5% **2** formed in reaction with O₂) is identical with that for disproportionation of **3**.
- P. Ford, DeF. P. Rudd, G. Gaunder, and H. Taube, *J. Am. Chem. Soc.*, **90**, 1187 (1968).
- In this reaction, roughly 65% of the theoretical amount of H₂O₂ is found.
- H. S. Lim, D. J. Barclay, and F. C. Anson, *Inorg. Chem.*, **11**, 1460 (1972).
- For example, (NH₃)₅Ru²⁺ (E°_{Ru(III)/Ru(II)} = 0.05 V) reacts in seconds with O₂ to produce Ru(III) and 1/2 H₂O₂; J. R. Pladewicz, T. J. Meyer, J. A. Broomhead, and H. Taube, *Inorg. Chem.*, **12**, 639 (1973). These workers were the first to suggest formation of Ru(IV) from reaction of Ru(II) with O₂.
- B. A. Moyer and T. J. Meyer, *J. Am. Chem. Soc.*, **100**, 3601 (1978).

- (17) The H_2O_2 was determined by use of the iron(II) phenanthroline method as described previously (see ref 15).
- (18) Thermodynamically the two-electron reaction of 1 with O_2 (reaction a, Scheme 1) is favored ($E^\circ_{O_2/H_2O_2} = 0.2$ V, $E^\circ_{O_2/H_2O} = -0.4$ V at pH 9) (see ref 20). However, we do not have experimental evidence confirming that the reaction of 1 with O_2 gives 4 directly. An alternate possibility involves formation of 3 and $HO_2\cdot$ (or $O_2^{\cdot-}$) and subsequent reaction of $HO_2\cdot$ ($O_2^{\cdot-}$) with another molecule of 1 to give 3 and H_2O_2 .
- (19) In independent experiments, oxidation of 1 by an equivalent quantity of H_2O_2 yielded only small quantities of 2 (~5% after 2 h). Similarly, after 1 h only negligible quantities (<1%) of 2 were obtained in a reaction of 3 with $HO_2\cdot$ ($3/KO_2 = 1$; pH 2). These results strongly suggest that step c in Scheme 1 is the dominant pathway for production of 2 in acidic solution.
- (20) G. A. Hamilton, P. K. Adolf, J. deJessey, G. C. DuBois, G. R. Dyrkacz, and R. D. Libby, *J. Am. Chem. Soc.*, **100**, 1899 (1978).

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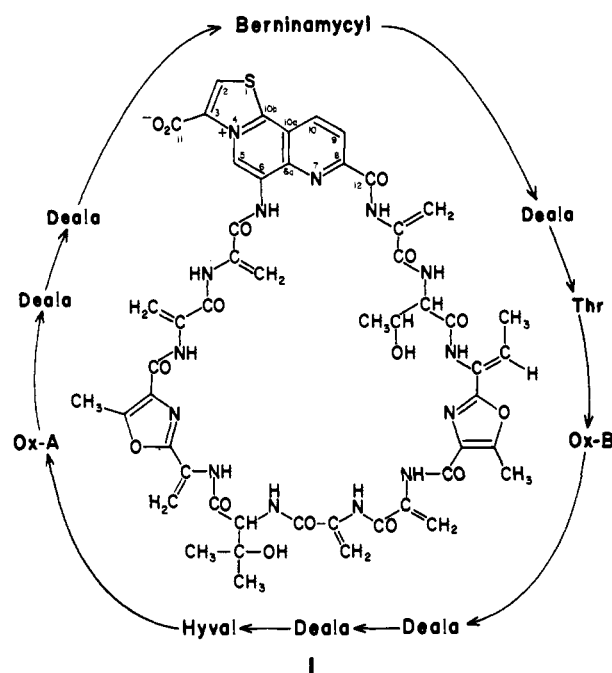
Berninamycin Biosynthesis. 1. Origin of the Dehydroalanine Residues

Sir:

Dehydro amino acid residues are a common feature of many microbial peptides. In particular, dehydroalanine occurs in a number of peptide antibiotics, inter alia, in nisin,¹ alternarolide,² subtilin,³ siomycin,⁴ thioestrepton,⁵ nosiheptide,⁶ and thiopeptin.⁷ Dehydro amino acids have also frequently been implicated as biochemical reaction intermediates. It has been suggested, for example, that dehydrocysteine- and dehydrovaline-containing peptides, produced from dehydrogenation of the relevant precursors, could be intermediates of penicillin and cephalosporin biosynthesis.⁸ In addition, enzyme-bound dehydroalanine has been proposed as an intermediate in the desulfuration of cysteine catalyzed by *S*-alkyl-L-cysteine lyase,⁹ in the dehydration of serine by serine dehydratase,¹⁰ and in the metabolism of *O*-acetylserine.¹¹

The biochemical origins of the peptide antibiotic dehydro amino acid residues have received little attention. There is some experimental evidence suggesting that the lanthionine and β -methylanthionine residues found in nisin are derived, in part, from serine and threonine, respectively, and it was proposed that dehydro amino acids play a role as reaction intermediates.¹² In addition, it was suggested that the dehydroalanine and dehydrobutyrine residues present in nisin may also be produced by dehydration of the relevant precursors. Furthermore, Bycroft¹³ has postulated that the biosynthesis of dehydroalanine residues found in peptide antibiotics results from either the dehydration of serine or the dehydrogenation of alanine, but until now no experimental evidence has provided a

sensible choice between the two possibilities. The present communication describes results which demonstrate that dehydroalanine residues arise by dehydration of serine, at least in berninamycin A (1), a polypeptide antibiotic produced by



*Streptomyces bernensis*¹⁴ which has been shown to inhibit protein synthesis at the ribosome level.¹⁵ The structure assigned in this laboratory¹⁶⁻¹⁸ contains five dehydroalanine residues; in addition, other dehydro amino acids are involved in the oxazoline A and B units and the berninamycinic acid residue contains a dehydrocysteine unit and yet another potential dehydroalanine unit.

Incubation of *S. bernensis* in the presence of DL-[1-¹⁴C]- or L-[U-¹⁴C]serine produced heavily labeled berninamycin (Table I), with incorporation from L-serine ~1.5 times that of DL-serine, demonstrating that the L isomer is preferentially utilized for berninamycin biosynthesis. On the other hand, similar experiments using DL-[1-¹⁴C]alanine indicated only a very small incorporation of label from that precursor ($1/100$ th the incorporation of L-serine) into the antibiotic, showing that alanine is not an effective precursor.

Samples of berninamycin produced by incubating *S. bernensis* with ¹⁴C-labeled serine were degraded, as described previously, to berninamycinic acid¹⁷ and pyruvic acid dinitrophenylhydrazide,¹⁷ with dehydroalanine residues being isolated as the latter derivatives. The specific activities of the fragments isolated are reported in Table I.

Table I. Incorporations of ¹⁴C-Labeled Amino Acids into Berninamycin and Subunits^a

	DL-[1- ¹⁴ C]serine added		L-[U- ¹⁴ C]serine added		DL-[1- ¹⁴ C]alanine added		L-[U- ¹⁴ C]cysteine added	
	48 h	72 h	48 h	72 h	48 h	72 h	48 h	72 h
incorporation into berninamycin, % ^b	2.8	1.9	4.2	2.6	0.042	0.018	1.0	0.27
specific activity of berninamycin ^c	0.497	0.384	0.755	0.369	ND	ND	0.138	0.031
specific activity of pyruvic acid dinitrophenylhydrazide ^c	0.056	0.056	0.074	0.061	ND	ND	0.0012	0.0013
specific activity of berninamycinic acid ^c	0.151	0.129	0.159	0.110	ND	ND	0.128	0.039

^a The labeled amino acids were added to 50 mL of *S. bernensis* cultures in 500-mL Erlenmeyer flasks. The culture medium contained glucose, 1%; L-glutamic acid, 0.2%; K_2HPO_4 , 0.1%; $MgSO_4 \cdot H_2O$, $FeSO_4 \cdot 7H_2O$, $ZnSO_4 \cdot 7H_2O$, $CaCl_2 \cdot 2H_2O$, 0.0025%; and distilled water to 50 mL, with a final pH of 7.2-7.3. This was inoculated using 0.5 mL of a 24-h culture of *S. bernensis* grown in Pharmamedia, 2.5%; glucose, 2.5%; distilled water to 50 mL, with a final pH of 7.2. The cultures were incubated on a rotary shaker at 30 °C and 250 rpm for 96 h. Berninamycin was isolated by the method previously described.¹⁴ Final purification was achieved by TLC over silica gel (CH_2Cl_2 - CH_3OH , 12:1). ^b The incorporation values have been adjusted for berninamycin recovery. There was some variation in the exact percent incorporation observed in individual experiments, related to the yield of berninamycin. In a series of strictly parallel experiments, the ratio of incorporation/mole berninamycin for L/DL-serine was 1.80 for 48-h addition and 1.73 for 72-h addition. ^c Carrier berninamycin was added prior to degradation. Specific activities are expressed in terms of $\mu Ci/mmol$; ND = not determined.