

Biosynthesis of Vitamin B₁₂: the Fate of 8-H as Precorrin-2 is Enzymically converted into Cobyric Acid

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The dimethylated B₁₂-precursor, precorrin-2 **2**, is prepared strongly labelled with tritium at 3-H and 8-H to allow proof that the 3-H label is retained whereas the 8-H label is lost during the biosynthetic conversion of precorrin-2 into cobyrinic acid **3**.

The corrin, cobyrinic acid **3**, is a late biosynthetic precursor of vitamin B₁₂ in *Propionibacterium shermanii*¹ and the related metal-free form, hydrogenobyric acid **4**, is on the biosynthetic pathway to B₁₂ in *Pseudomonas denitrificans*.² Both **3** and **4** are produced *in vivo* from uro'gen III **1** by a sequence of steps which includes eight C-methylations.¹ The first two C-methylations are carried out by one enzyme^{3,4} which is well characterised⁴ and the structure of the product **2** named⁵ precorrin-2 was fully established by direct isolation.⁶

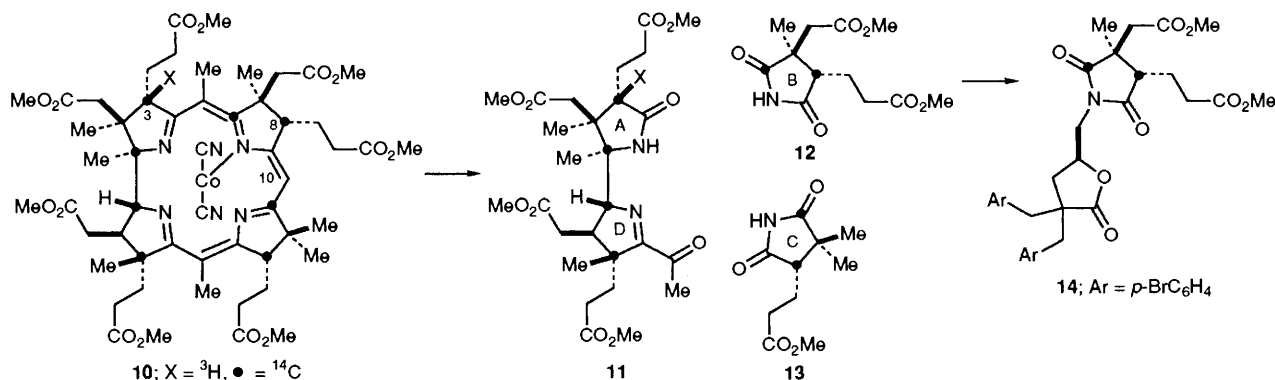
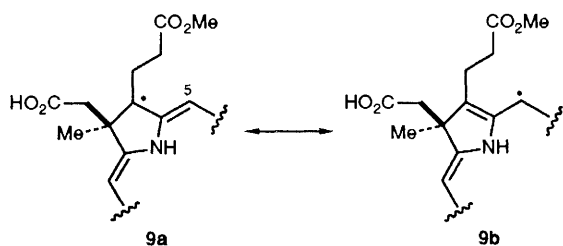
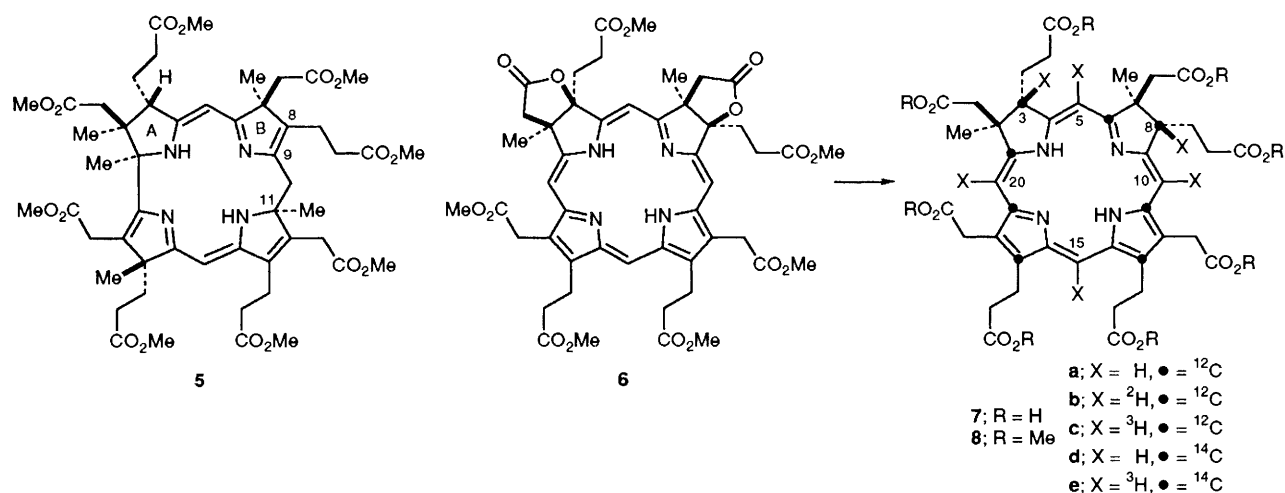
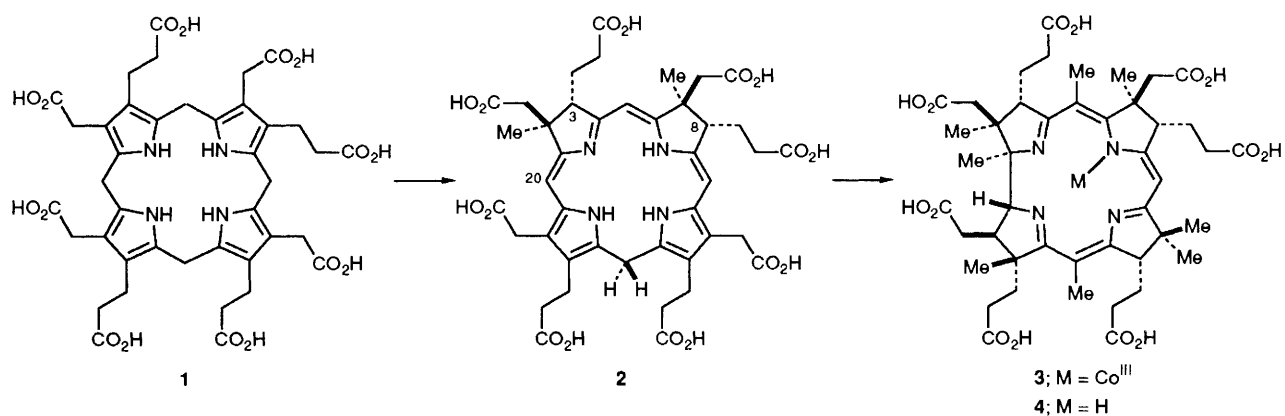
Precorrin-6x, the product of six C-methylations from **1**, was recently isolated⁷ from a *P. denitrificans* strain, an event that changed the direction of research on the biosynthesis of vitamin B₁₂ and related corrinoids. The reasons can be seen in the structure **5** established^{7,8} for the octamethyl ester of precorrin-6x which reveals several unexpected features, especially the C-11 methyl group. The quaternary centre at C-11 generates two separated chromophores (*cf.* the conjugated systems **2** and **3**). The chromophore of ring A—ring B in the ester of precorrin-6x has a double bond at C-8 to C-9 which may or may not be located there in the original enzymic product, precorrin-6x itself. Because the dimethylated precursor **2** is the same for both the anaerobic *P. shermanii* and the aerobic *P. denitrificans* and the corrins **3** and **4** produced differ only in the presence or absence of cobalt, and bearing in mind the C-8 to C-9 double bond in **5**, it was of considerable interest to study the fate of 8-H of precorrin-2 **2** as it is enzymically converted into cobyrinic acid **3** in *P. shermanii*.

Precorrin-2 **2** can be generated *in situ* from sirohydrochlorin **7a** in the enzymic incubation mixture¹ and the method for

labelling at C-8 of the corresponding ester **8a** was first developed as follows using deuterium. The lactone rings of the bis-lactone⁹ **6** were reductively cleaved¹⁰ with chromium(II) chloride in a mixture of deuterium oxide and tetrahydrofuran (THF) and the resultant dicarboxylic acid was esterified to afford ²H-labelled sirohydrochlorin octamethyl ester **8b**, 60% yield. Repeated integration of the ¹H-NMR spectrum of this product allowed the extent of ²H-labelling to be determined at the positions listed in Table 1. Two points need comment: (*a*) heavy labelling has been achieved at positions 3 and 8 but not complete deuteration, (*b*) positions 5, 10, 15 and 20 are significantly labelled. We believe both (*a*) and (*b*) are due to the radical nature of chromium(II) reductions.¹¹ Thus, an initial radical **9a** can abstract deuterium from D₂O or protium from the THF whilst reaction through the alternative limiting form **9b** labels C-5; the labelling at 10, 15 and 20 is explained in the same way.

Table 1 Distribution of ²H in **8b** and of ³H in **8c**

Site	8b % ² H at each site	8c % of total ³ H
3	66	29
5	46	20
8	66	29
10	7	3
15	28	12
20	15	7



The entire process was then repeated under the same conditions save that a small amount of high activity tritiated water was added to the D₂O–THF mixture at the outset. Note that D₂O was used as solvent rather than H₂O because transfer of tritium from DOT is more efficient than from HOT due to the smaller isotope effect in the former case. The ³H-labelled sirohydrochlorin octamethyl ester **8c** was isolated in 50% yield, total activity 6.7 μCi, specific activity 0.93 mCi

mmol⁻¹. Finally, the expected distribution of ³H-activity at the various labelled sites† of the ³H-sample **8c** was calculated from the deuterium values in Table 1.

† Because tritium is used at tracer level, the sample is an assembly of six singly ³H-labelled species which together are equivalent to the multiply labelled molecule illustrated. Each labelled site also carries deuterium which for simplicity is not shown.

Table 2 Enzymic formation and subsequent degradation of labelled cobester **10**

	³ H: ¹⁴ C ratio		³ H: ¹⁴ C ratio calc. from Table 1		¹⁴ C-Specific activity /dpm mmol ⁻¹		³ H-Specific activity /dpm mmol ⁻¹		
	1	2	1	2	1	2	1	2	
Experiment									
Precorrin-2 2		0.57	5.6	—	—	2.44 × 10 ⁹	4.22 × 10 ⁸	1.39 × 10 ⁹	2.38 × 10 ⁹
Cobester 10	(i) ^a	0.17	1.8	0.17	1.6	1.16 × 10 ⁷	2.06 × 10 ⁷	2.03 × 10 ⁶	3.63 × 10 ⁷
	(ii) ^a	0.17	1.7	0.17	1.6	2.46 × 10 ⁵	4.51 × 10 ⁵	4.30 × 10 ⁴	7.74 × 10 ⁵
Ring-A/ring-D 11		0.33	^b	0.34	—	1.19 × 10 ⁵	—	3.93 × 10 ⁴	—
Ring-B imide 12		<0.01	<0.01	0	0	5.57 × 10 ⁴	1.14 × 10 ⁵	^c	^c
Derivative 14		<0.01	<0.01	0	0	5.23 × 10 ⁴	—	^c	^c
Ring-C imide 13		<0.01	<0.01	0	0	5.34 × 10 ⁴	1.11 × 10 ⁵	^c	^c

^a Cobester (i) before and (ii) after substantial dilution for degradation. ^b The yield of ring-A/ring-D fragment from this degradation was too low to allow rigorous purification of a sufficiently active sample. ^c ³H-Activity below detection level.

¹⁴C-Labelled **7d** was biosynthesised¹² from 5-amino[4-¹⁴C]laevulinic acid and was isolated as its ester **8d**, specific activity 3.3 mCi mmol⁻¹. This was mixed with an appropriate amount of the ³H-sample **8c** and the radioactivity values are given in Table 2. Hydrolysis of this ³H: ¹⁴C-labelled sample **8e** and enzymic conversion of the resultant octa-acid into coobyric acid (as **3**) was carried out as usual using the cell-free enzyme system prepared¹³ from *P. shermanii* cells. Unlabelled coobyric acid was then added as carrier and the derived cobester **10** was recrystallised to constant specific activity and ³H: ¹⁴C ratio (incorporation 18.3%). This product was further diluted with unlabelled cobester for degradation by ozonolysis¹⁴ to yield the ring-B imide **12**, the ring-C imide **13** and the ring-A/ring-D fragment¹⁵ **11** which were carefully purified for assay of radioactivity. Since it was known¹⁶ that the ring-B imide **12** produced in this way contains some of the imide derived from ring-D of **10** and **11**, the ring-B imide was converted into the derivative **14** which is readily separated from the corresponding derivative of the ring-D imide¹⁷; the derivative **14** also serves to characterise further the ring-B fragment **12**. Table 2 collects the results for the set of labelled materials. All the foregoing studies were then repeated starting with a sample of **8e** having a higher ³H: ¹⁴C ratio especially to check the foregoing findings for the ring-B imide **12**: the incorporation was 27.5% and the activity values from this independent study are also in Table 2.

The key results from Table 2 are (a) the ring-C imide **13** carries the appropriate ¹⁴C-activity but, as expected, no ³H, (b) the ring-A/ring-D fragment **11** carries the required ¹⁴C- and ³H-activity corresponding to retention of 3-H but (necessarily) loss of 20-H of precorrin-2 **2**, (c) the ring-B imide **12** shows that all the ³H has been lost from C-8 of **2** but the ¹⁴C-activity of **12** matches the expected value, (d) the ³H: ¹⁴C ratio in the original cobester **10** interlocks with the results under (b) and (c) by agreeing with the loss of ³H from C-5, C-8, C-10, C-15 and C-20 of precorrin-2 **2** but with retention of ³H at C-3.

Loss of the small amount of ³H at C-10 of cobester **10** is included in (d) above because the acidic methanol used for esterification of coobyric acid to give **10** is known to cause complete exchange at this position.¹⁸ Possible similar exchange at C-8 was examined by using CD₃OD in a control esterification of **3** and NMR of the ester confirmed that there was no significant exchange at C-8.

Thus, the facts are that as precorrin-2 **2** is converted enzymically into coobyric acid **3**, 3-H is retained whereas 8-H is lost. These findings are consistent with there being a C-8 to C-9 double bond (as in **5**) at some stage of the biosynthesis but it remains to be established whether this double bond position is a requirement of one of the transforming enzymes. The present results will form part of the total picture when the structures are determined for the presently unknown intermediates on the pathway to coobyric acid **3**.

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