

Biocatalysis

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Internationale Ausgabe: DOI: 10.1002/anie.20151033125-Hydroxyvitamin D₃ Synthesis by Enzymatic Steroid Side-Chain Hydroxylation with Water

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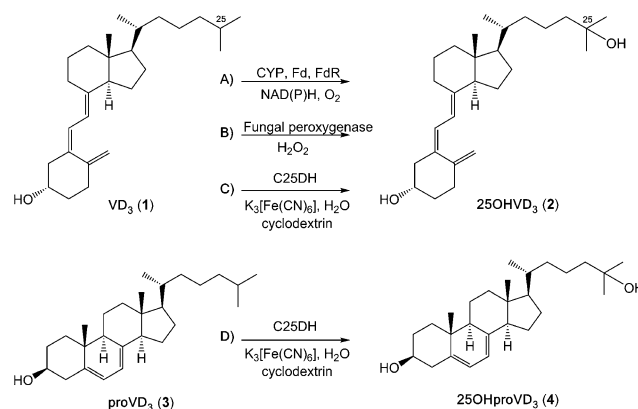
Dedicated to Georg Fuchs on the occasion of his 70th birthday

Abstract: The hydroxylation of vitamin D₃ (VD₃, cholecalciferol) side chains to give 25-hydroxyvitamin D₃ (25OHVD₃) is a crucial reaction in the formation of the circulating and biologically active forms of VD₃. It is usually catalyzed by cytochrome P450 monooxygenases that depend on complex electron donor systems. Cell-free extracts and a purified Mo enzyme from a bacterium anaerobically grown with cholesterol were employed for the regioselective, ferricyanide-dependent hydroxylation of VD₃ and proVD₃ (7-dehydrocholesterol) into the corresponding tertiary alcohols with greater than 99% yield. Hydroxylation of VD₃ strictly depends on a cyclodextrin-assisted isomerization of VD₃ into preVD₃, the actual enzymatic substrate. This facile and robust method developed for 25OHVD₃ synthesis is a novel example for the concept of substrate-engineered catalysis and offers an attractive alternative to chemical or O₂/electron-donor-dependent enzymatic procedures.

Vitamin D₃ (VD₃, cholecalciferol, **1**) is generally known for its regulatory function in calcium and phosphorous homeostasis, but it is also recognized as an antiproliferative factor in dividing cells and tissues.^[1] VD₃ is formed from proVD₃ (7-dehydrocholesterol, **3**) by UV-B irradiation to give the *seco* form, followed by a thermal [1,7] sigmatropic hydrogen shift. It is converted into its biologically active form by consecutively acting cytochrome P450 monooxygenases (CYPs): hydroxylation in the liver yields 25-hydroxy-VD₃ (25OHVD₃, calcidiol, **2**), which is converted into calcitriol (1 α ,25(OH)₂VD₃) in the kidney. Insufficient levels of biologically active VD₃ are linked to numerous forms of cancer but

also to cardiovascular diseases, immunodeficiency, and diabetes.^[2] 25OHVD₃ (**2**) represents the circulating form of VD₃, and its plasma concentration has been associated with VD₃-linked disorders.^[3]

Classical chemical 25OHVD₃ (**2**) synthesis procedures involve multiple steps with low yields and specificities,^[4] although recently more convenient procedures, for example, methods involving anchoring on a solid phase, have been developed.^[5] For enzymatic synthesis, CYPs either from liver, fungi, or actinomycetes have been employed.^[6] However, the dependence on complex electron-donor systems consisting of NAD(P)H, ferredoxin (Fd), and ferredoxin reductase (Fdr) has limited biotechnological applications; promising approaches employ whole cells expressing the genes encoding CYP, Fd, and Fdr (Scheme 1 A). As an alternative, an H₂O₂-dependent peroxygenase has been used for hydroxylation of low concentrations of VD₃ and ergocalciferol (VD₂; Scheme 1 B).^[7]



Scheme 1. Enzymatic conversions of VD₃ (**1**) and proVD₃ (**3**) into 25OHVD₃ (**2**) and 25OHproVD₃ (**4**), respectively. A) Hydroxylation of **1** by O₂-dependent CYPs in the presence of an electron-donor system including NAD(P)H, Fd, and Fdr. B) Hydroxylation of **1** by H₂O₂-dependent peroxygenase. C, D) Water- and K₃[Fe(CN)₆]-dependent hydroxylation of **1** and **3** by C25DH. (A) and (B) were previously established,^[6,7] (C) and (D) were studied in this work.

A fundamentally different mode of enzymatic steroid side-chain hydroxylation was identified during the anaerobic degradation of cholesterol to CO₂ in the denitrifying bacterium *Sterolibacterium denitrificans*: the regioselective hydroxylation of cholesterol into 25-hydroxycholesterol with

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water catalyzed by steroid C25 dehydrogenase (C25DH).^[8] This enzyme belongs to the DMSO reductase (DMSOR) family of Mo enzymes that catalyze O_2 -independent hydroxy- and oxo-transfer reactions.^[9] A C–H bond functionalization with water to a tertiary alcohol via a carbocation transition state has been proposed.^[8] C25DH transfers electrons to electrochemically regenerative electron acceptors such as $K_3[Fe(CN)_6]$ (Scheme 1C). The genome of *S. denitrificans* contains at least eight homologous genes encoding putative Mo-dependent steroid hydroxylases,^[8] thus making this organism a promising biological platform for water-dependent hydroxylations of various steroids. In this work, we explored the use of *S. denitrificans* enzyme(s) for regioselective synthesis of the biologically relevant hydroxylated forms of VD_3 (**1**) and $proVD_3$ (**3**). The hydroxylated $25OHproVD_3$ (**4**, Scheme 1D) can easily be converted into $25OHVD_3$ (**2**) by established UV irradiation methods.^[10]

In the presence of 2-hydroxypropyl- β -cyclodextrin (HPCD), crude extracts from *S. denitrificans* anaerobically grown with cholesterol catalyzed the $K_3[Fe(CN)_6]$ -dependent conversion of VD_3 (**1**) and $proVD_3$ (**3**) into the single products **2** and **4** (Figure 2A,B), respectively, with substrate conversions of >99% (**2**) and 97% (**4**; Figure 1A,B). After the

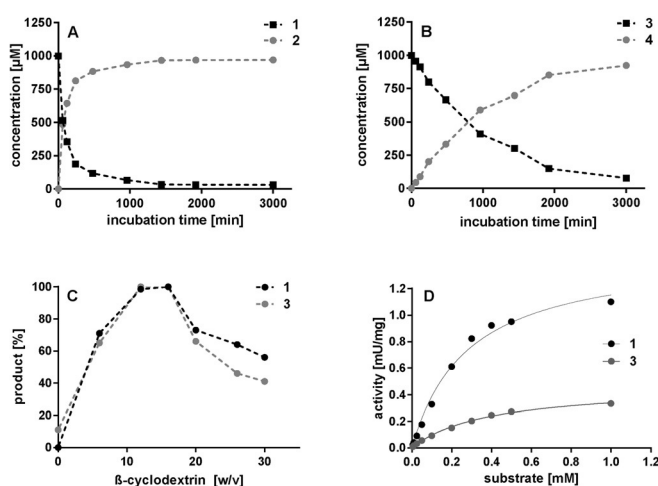


Figure 2. Properties of the enzymatic hydroxylation of **1** and **3**.

A) Time-dependent conversion of **1** into **2** in the presence of 0.5 mM $AgNO_3$. B) Time-dependent conversion of **3** into **4**. Both reactions contained 15% HPCD. C) HPCD dependence of the conversion of **1** and **3**. D) Substrate concentration dependence of the conversion of **1** and **3**.

dependent on electron acceptors with standard redox potentials higher than 200 mV, such as $[Fe(CN)_6]^{3-}$ (ferricyanide), with the highest initial rates between 5–10 mM $K_3[Fe(CN)_6]$. This rather high value can be explained by competing unspecific $K_3[Fe(CN)_6]$ oxidizing reactions in cell extracts.

The products formed from **1** and **3** were analyzed by chemical ionization mass spectrometry. For the product of **1** ($M_r = 384.65$), compound **2**, an ion with $m/z [M+H]^+$ of 401.3, was determined, thus indicating the addition of a hydroxy group (Figure S1 in the Supporting Information). Likewise, the conversion of **3** ($M_r = 384.65$) yielded a product with an identical $m/z [M+H]^+$ (Figure S2). Both results strongly suggest the hydroxylation of VD_3 (**1**) and $proVD_3$ (**3**).

To unambiguously determine the regioselectivity of the reaction, the products were characterized by NMR spectroscopy. The 1H and ^{13}C NMR spectra of authentic **1** and **3** standards were highly similar to their products **2** and **4**, respectively, with the exception of the chemical shifts for C26, C27 methyl groups (1H spectra), and C25 (^{13}C spectra). For **1** and **3**, the C26 and C27 methyl groups are observed as doublets at $\delta = 0.86$ ppm in the 1H spectra. For **2** and **4**, the 1H NMR signal for CH_3 -26,27 is observed as a singlet at $\delta = 1.23$ ppm. A downfield shift is observed in the ^{13}C NMR spectrum for the C25 signal ($\delta = 71.1$ ppm versus 22.8 ppm for **1** and **3**), thus indicating the hydroxylation of C25, as described for 25-hydroxycholest-4-en-3-one.^[11]

The 1H and ^{13}C NMR spectroscopic data are in perfect agreement with those published for $25OHVD_3$.^[14] The complete NMR spectroscopic data are presented in the Supporting Information (1H and ^{13}C NMR spectra/shifts, 2D NMR spectra for **2**, and 1H NMR spectrum of **4**; Figures S3–S5 and Table S1).

To identify the enzyme(s) involved in VD_3 (**1**) hydroxylation, we enriched the activity from extracts of cells grown with cholesterol. After four chromatographic steps, four

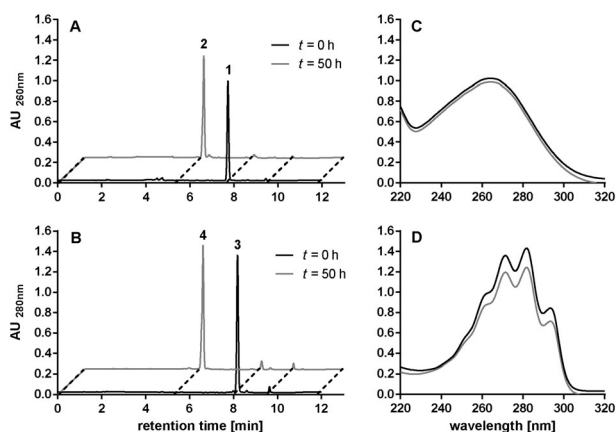


Figure 1. Enzyme assays for the hydroxylation of **1** and **3**. A, B) Ultra-performance liquid chromatography (UPLC) analysis of the enzymatic conversion of **1** and **3** into their hydroxylated products **2** and **4**, respectively. C, D) UV spectra of **1–4**.

conversion of VD_3 (**1**) reached 92% (16 h), the product **2** was slowly degraded by unspecific enzymatic reactions, which could be fully prevented by the addition of 0.5 mM $AgNO_3$, a known inhibitor of the downstream reactions of steroid catabolism.^[8] The UV/Vis spectra of the products were identical to those of the substrates, which is in agreement with side-chain hydroxylations distant from the conjugated triene (**1**) and diene (**3**) chromophors (Figure 1C,D).

Conversion of **1** and **3** was HPCD dependent, with an optimum at 15% (w/v; Figure 2C). The initial rate of the reaction followed Michaelis–Menten kinetics, with $v_{max} = 1.48 \pm 0.05$ nmol min^{−1} per mg crude extract protein for VD_3 (**1**), which is threefold higher than the rate of $proVD_3$ (**3**) conversion (0.47 ± 0.02 nmol min^{−1} per mg crude extract protein; Figure 2D). The conversions of **1** and **3** were strictly

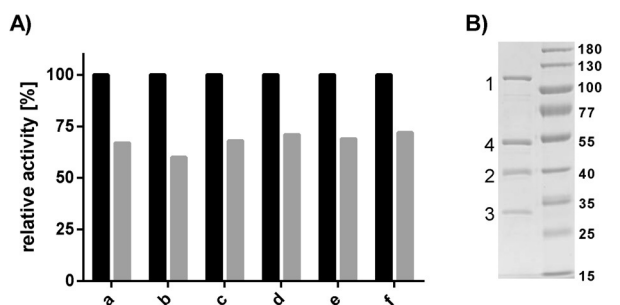
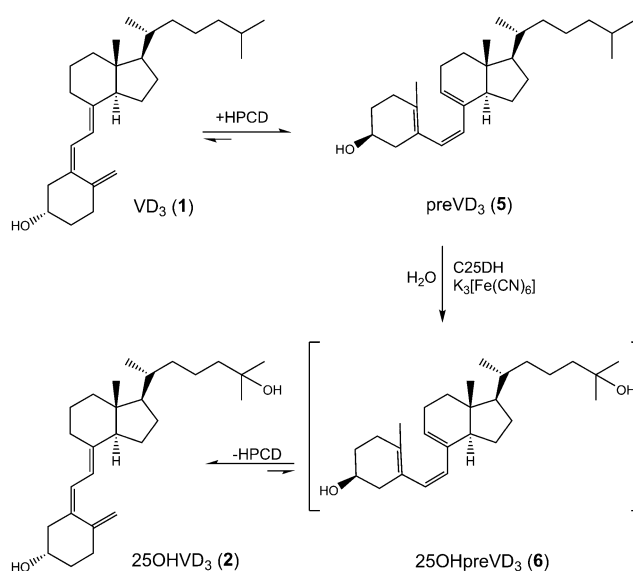


Figure 3. Enrichment of the VD₃ (1) hydroxylating enzyme. A) Relative specific conversion rates of VD₃ (1, grey columns) and cholest-4-en-3-one (black columns) obtained during the solubilization and purification steps. a) cell extract; b) solubilized membrane fraction; c–f) fractions obtained after chromatography on DEAE-Sephadex, ceramic hydroxyapatite, Reactive Red agarose, and Mono Q anion exchange chromatography, respectively. B) SDS gel of enriched enzyme after four chromatographic steps. Left lane: bands 1–3 were identified as αβγ-subunits of C25DH; band 4 is a degradation product of band 1 (Table S2). Right lane: Molecular mass standard with masses in kDa as indicated.

protein bands were highly enriched with a specific VD₃ (1) hydroxylation activity of 80 nmol min^{−1} mg^{−1} (Figure 3A). Mass spectrometric analysis of tryptic digests of the enriched protein bands identified the αβγ-subunits of C25DH (JQ292991-3) plus a fourth band corresponding to a degradation product of the α-subunit (JQ292991; Table S2). After each purification step, the relative specific hydroxylation rates of VD₃ (1) and cholest-4-en-3-one^[8] remained almost constant in all activity-containing fractions (around 70 % with 1), thus confirming that both reactions are catalyzed by the same enzyme (Figure 3B). VD₃ (1) hydroxylation activity was highest in extracts of cells grown with cholesterol (100 %), and significantly lower in cells grown with sitosterol (18 %), ergosterol (15 %), or stigmasterol (5 %), which is in line with the induction of C25DH during growth with cholesterol.

The conversion of cholest-4-en-3-one and preVD₃ (3) by C25DH was observed in the presence of various detergents and HPCD. In contrast, conversion of VD₃ (1) was negligible when HPCD was replaced by a number of commonly used detergents (Figure S6). This finding suggests that HPCD plays a specific role in VD₃ conversion rather than serving as a general solubilizing agent. The equilibrium preVD₃ (5) ⇌ VD₃ (1) through thermal [1,7] sigmatropic rearrangement lies far to the right in aqueous detergent solutions (Scheme 2, upper reaction).^[12] However, in the presence of HPCD, an inclusion complex is formed at a 2:1 ratio (HPCD/VD₃), in which the thermodynamically unfavorable *cZc* conformer of preVD₃ (5) is stabilized.^[12] This finding led us to the assumption that preVD₃ (5) rather than VD₃ (1) is the actual substrate of C25DH.

By replacing the commonly used ethyl acetate for steroid extraction by *n*-hexane, more than 80 % of VD₃ (1, 1 mM) added to the HPCD-containing assay mixture was identified as preVD₃ (5) by UPLC analysis. In the course of the enzymatic reaction, preVD₃ (5) was time-dependently converted into 25OHVD₃ (2), with virtually no formation of the expected 25OHpreVD₃ (6) intermediate (Figure 4). In con-



Scheme 2. Proposal for HPCD-dependent hydroxylation of VD₃ by C25DH. In the presence of HPCD, the equilibrium between VD₃ (1) and preVD₃ (5), which are interconverted through a thermal [1,7] hydrogen shift, is shifted to the right. The *cZc* conformer of 5 is likely the substrate for C25DH. The hydroxylated 6 was not detected owing to rapid isomerization to 2.

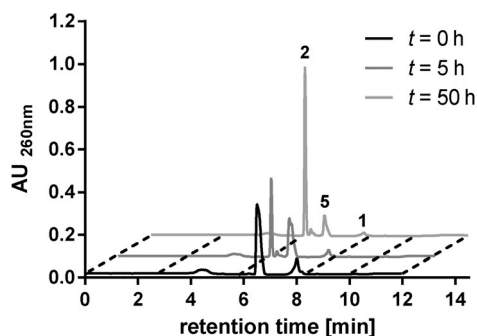


Figure 4. UPLC analysis of enzymatic VD₃ (1) hydroxylation in the presence of HPCD after steroid extraction with *n*-hexane. At *t* = 0 h, before enzyme addition, around 80 % of the added VD₃ was HPCD-dependently isomerized into preVD₃ (5).

trast, during extraction with ethyl acetate, preVD₃ (5) was never identified during VD₃ (1) hydroxylation (Figure 1A). To substantiate the HPCD-dependent conversion of VD₃ (1) into preVD₃ (5) as an essential process for enzymatic side-chain hydroxylation, we replaced VD₃ (1) by authentic preVD₃ (5) and observed an identical product pattern as during VD₃ (1) conversion (Figure S7). The rates of VD₃ (1) and preVD₃ (5) hydroxylation were nearly identical, thus suggesting that the HPCD-dependent conversion of VD₃ (1) into preVD₃ (5) is not rate limiting.

In summary, the results indicate that preVD₃ (5) but not VD₃ (1) is the substrate for C25DH. The failure to detect any trace of a 25-OH-preVD₃ intermediate indicates a rapid isomerization into 25OHVD₃ (Scheme 2).

The procedure presented in this work represents a new enzymatic route for the regioselective hydroxylation of VD₃

(1), proVD₃ (3), and preVD₃ (5) at C25, and expands our knowledge of tertiary alcohol synthesis.^[13] Since cell extracts of *S. denitrificans* can be used for these reactions, it circumvents tedious enzyme purification or genetic manipulation procedures and offers a facile method for a mechanistically demanding chemical reaction. The method strictly depends on HPCD for VD₃ (1) solubilization and isomerization and represents a novel example of substrate-range expansion. In comparison to CYP-dependent hydroxylations it does not depend on electron-donor systems. The only cofactor used, K₃[Fe(CN)₆], can be regenerated by electrodes as has been demonstrated for the related ethyl benzene dehydrogenase.^[14] Moreover, the method does not depend on oxygen and nor is it oxygen sensitive. The synthetic method for 25OHVD₃ synthesis may be useful for the treatment of 25OHVD₃ deficiencies and to satisfy the increasing demand for monitoring of its levels in blood plasma.

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- [1] a) M. F. Holick, *N. Engl. J. Med.* **2007**, 357, 266–281; b) S. V. Ramagopalan, A. Heger, A. J. Berlanga, N. J. Mauger, M. R. Lincoln, A. Burrell, L. Handunnetthi, A. E. Handel, G. Disanto, S. M. Orton, C. T. Watson, J. M. Morahan, G. Giovannoni, C. P. Ponting, G. C. Ebers, J. C. Knight, *Genome Res.* **2010**, 20, 1352–1360.
- [2] a) M. Wacker, M. F. Holick, *Nutrients* **2013**, 5, 111–148; b) A. Hosseini-nezhad, M. F. Holick, *Mayo Clin. Proc.* **2013**, 88, 720–755.
- [3] C. J. Farrell, S. Martin, B. McWhinney, I. Straub, P. Williams, M. Herrmann, *Clin. Chem.* **2012**, 58, 531–542.
- [4] G. D. Zhu, W. H. Okamura, *Chem. Rev.* **1995**, 95, 1877–1952.
- [5] D. Nicoletti, A. Mourino, M. Torneiro, *J. Org. Chem.* **2009**, 74, 4782–4786.
- [6] a) K. Hayashi, H. Sugimoto, R. Shinkyo, M. Yamada, S. Ikeda, S. Ikushiro, M. Kamakura, Y. Shiro, T. Sakaki, *Biochemistry* **2008**, 47, 11964–11972; b) K. Hayashi, K. Yasuda, H. Sugimoto, S. Ikushiro, M. Kamakura, A. Kittaka, R. L. Horst, T. C. Chen, M. Ohta, Y. Shiro, T. Sakaki, *FEBS J.* **2010**, 277, 3999–4009; c) D. J. Kang, J. H. Im, J. H. Kang, K. H. Kim, *Bioprocess Biosyst. Eng.* **2015**, 38, 1281–1290; d) D. J. Kang, J. H. Im, J. H. Kang, K. H. Kim, *Biotechnol. Lett.* **2015**, 37, 1895–1904; e) T. Sakaki, H. Sugimoto, K. Hayashi, K. Yasuda, E. Munetsuna, M. Kamakura, S. Ikushiro, Y. Shiro, *Biochim. Biophys. Acta Proteins Proteomics* **2011**, 1814, 249–256; f) K. Yasuda, M. Endo, S. Ikushiro, M. Kamakura, M. Ohta, T. Sakaki, *Biochem. Biophys. Res. Commun.* **2013**, 434, 311–315; g) Y. Yasutake, T. Nishioka, N. Imoto, T. Tamura, *ChemBioChem* **2013**, 14, 2284–2291.
- [7] E. D. Babot, J. C. Del Rio, M. Canellas, F. Sancho, F. Lucas, V. Guallar, L. Kalum, H. Lund, G. Grobe, K. Scheibner, R. Ullrich, M. Hofrichter, A. T. Martinez, A. Gutierrez, *Appl. Environ. Microbiol.* **2015**, 81, 4130–4142.
- [8] J. Dermer, G. Fuchs, *J. Biol. Chem.* **2012**, 287, 36905–36916.
- [9] R. Hille, J. Hall, P. Basu, *Chem. Rev.* **2014**, 114, 3963–4038.
- [10] S. Fuse, Y. Mifune, N. Tanabe, T. Takahashi, *Org. Biomol. Chem.* **2012**, 10, 5205–5211.
- [11] Y. R. Chiang, W. Ismail, M. Müller, G. Fuchs, *J. Biol. Chem.* **2007**, 282, 13240–13249.
- [12] X. Q. Tian, M. F. Holick, *J. Biol. Chem.* **1995**, 270, 8706–8711.
- [13] M. Müller, *ChemBioEng. Rev.* **2014**, 1, 14–26.
- [14] P. Kalimuthu, J. Heider, D. Knack, P. V. Bernhardt, *J. Phys. Chem. B* **2015**, 119, 3456–3463.

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