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## Interplay between Vitamin D and the Drug Metabolizing Enzyme CYP3A4

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### Abstract

Cytochrome P450 3A4 (CYP3A4) is a multifunctional enzyme involved in both xenobiotic and endobiotic metabolism. This review focuses on two aspects: regulation of *CYP3A4* expression by vitamin D and metabolism of vitamin D by CYP3A4. Enterohepatic circulation of vitamin D metabolites and their conjugates will be also discussed. The interplay between vitamin D and CYP3A4 provides new insights into our understanding of how enzyme induction can contribute to vitamin D deficiency.

### Keywords

Cytochrome P450 3A4; Vitamin D; Drug metabolism; Enterohepatic circulation; Osteomalacia

### 1. Vitamin D as a Hormone

Vitamin D consists of two isoforms, vitamin D<sub>2</sub> and vitamin D<sub>3</sub>. Vitamin D<sub>3</sub> is the major source of the hormone in humans and is essential for the proper maintenance of calcium and phosphate homeostasis [1]. Once synthesized in skin after sunlight exposure, or absorbed in the gut from an oral diet, vitamin D<sub>3</sub> is delivered to the liver where it undergoes 25-hydroxylation by CYP2R1 and CYP27A1 (and other enzymes) to form 25-hydroxyvitamin D<sub>3</sub> (25OHD<sub>3</sub>). Although 25OHD<sub>3</sub> is the major circulating form of vitamin D<sub>3</sub>, it must be further oxidized at the 1 $\alpha$  position by the enzyme CYP27B1 to 1 $\alpha$ ,25-dihydroxyvitamin D<sub>3</sub> [1 $\alpha$ ,25(OH)<sub>2</sub>D<sub>3</sub>] to become fully active at regulating gene transcription and cell function. 1 $\alpha$ ,25(OH)<sub>2</sub>D<sub>3</sub> initiates or suppresses gene transcription by binding to the vitamin D receptor (VDR) [2]. Binding to VDR triggers hetero-dimerization of VDR with retinoid X receptor (RXR). The heterodimer then translocates to the nucleus where the complex binds to vitamin D response elements and alters gene transcription. In addition to this classical pathway of controlling cell function, several “non-genomic” pathways of cell regulation have also been proposed, involving extranuclear 1 $\alpha$ ,25(OH)<sub>2</sub>D<sub>3</sub> and multiple growth factors [3].

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CYP3A4 is considered to be the most important of the family of drug-metabolizing cytochrome P450 enzyme, contributing importantly to the clearance of perhaps half of therapeutic agents that undergo metabolic biotransformation [4]. It is highly expressed in the liver and small intestine [5] and several transcription factors interacting with multiple transcriptional elements regulate its expression. The induction of *CYP3A4* gene expression in the liver and small intestine is mainly regulated through activation of the pregnane X receptor (PXR) and, in the case of the liver, also by the constitutive androstane receptor [6]. However, it has been demonstrated that  $1\alpha,25(\text{OH})_2\text{D}_3$  can also enhance the transcription of *CYP3A4* by a VDR-mediated pathway. Treatment of Caco-2 cells [7–9], LS180 cells [10, 11], HepG2 cells [12], and human hepatocytes [13] with  $1\alpha,25(\text{OH})_2\text{D}_3$  results in increased *CYP3A4* mRNA levels and accumulation of the encoded, functional enzyme. In addition, treatment of rats with  $1\alpha,25(\text{OH})_2\text{D}_3$  increases intestinal *CYP3A23* (a homolog of *CYP3A4*) expression and CYP3A metabolic activity [14, 15].

*CYP3A4* transcription is induced by  $1\alpha,25(\text{OH})_2\text{D}_3$  through the binding of ligand-VDR-RXR heterodimer to the same proximal ER6 (–169/–152), distal DR3 (–7733/–7719) and DR4 (–7618/–7603) response elements to which a ligand-PXR-RXR complex binds [16–18]. VDR also functions as a receptor for the secondary bile acid lithocholic acid (LCA) and binds (as an activated heterodimer) to the same *CYP3A4* ER6 and DR3 response elements as does the  $1\alpha,25(\text{OH})_2\text{D}_3$ -VDR-RXR complex [19]. Moreover, activation of VDR by LCA can induce CYP3A expression *in vivo* indicating that LCA may also contribute to CYP3A4 induction in the enteric system [20]. Finally, C-jun-N-terminal kinase has been suggested to be another mediator of  $1\alpha,25(\text{OH})_2\text{D}_3$ -induced *CYP3A4* expression, presumably without involvement of VDR [8, 21].

The primary sites of CYP3A4 expression are the liver and mucosa of the small intestine. Thus, it has been suggested that VDR helps regulate CYP3A4 enzyme content at these tissue sites [16, 22]. While this is clearly plausible for intestinal expression, it has been noted that the expression of VDR in human liver is much lower than that found in the small intestine and kidney [23]. Indeed, an immunohistochemical analysis of cross-sections of the human liver indicated that VDR is essentially absent from hepatocytes, but instead expressed in selective hepatic cell populations such as Kupffer, stellate and endothelial cells [24]. Therefore, considering the cellular composition of the liver, induction of hepatic *CYP3A4* expression via VDR in non-parenchymal cells may not be that relevant to overall hepatic drug clearance [25], although additional work is needed to clarify this important regulatory issue.

Induction of CYP3A4 by  $1\alpha,25(\text{OH})_2\text{D}_3$  appears to affect the systemic exposure of orally administered drugs that are substrates of CYP3A4 (Figure 1). Humans taking vitamin D supplementation show an increased clearance of atorvastatin, a substrate of CYP3A4 [26]. In addition, blood levels of CYP3A substrates, tacrolimus and sirolimus, showed cyclic variation throughout the year that correlated with ultraviolet light exposure and serum levels of vitamin D, whereas no significant difference was observed for mycophenolic acid, a non-substrate of CYP3A4 [27]. Similarly, Thirumaran et al [25] reported that intestinal *CYP3A4* expression *in vivo* varied seasonally, correlating with seasonal vitamin D levels. It is known that intestinal CYP3A4 contributes to the first-pass metabolism of many orally administered drugs [28]. Thus, intra- and inter-individual differences in circulating vitamin D levels and associated intestinal CYP3A4 activity may contribute to variability in oral drug bioavailability.

## 2. Enterohepatic Circulation of Vitamin D

Delivery of  $1\alpha,25(\text{OH})_2\text{D}_3$  to the intestinal mucosa can in theory occur by both vascular and biliary routes [29]. In the absence of active uptake across the basolateral membrane of intestinal enterocytes, one would expect unbound intracellular  $1\alpha,25(\text{OH})_2\text{D}_3$  concentrations to be the same as that found in plasma, driving transcriptional activation of VDR gene targets such as *CYP3A4*, transient receptor potential cation channel, subfamily V, member 6 (*TRPV6*) and calbindin D9K. However, studies utilizing radiolabeled  $1\alpha,25(\text{OH})_2\text{D}_3$  in humans [30, 31] and animals [32] have revealed biliary excretion and intestinal reabsorption of  $1\alpha,25(\text{OH})_2\text{D}_3$  (or its polar conjugate), suggesting the possibility of functionally significant enterohepatic cycling of  $1\alpha,25(\text{OH})_2\text{D}_3$  (Figure 1). Although biliary vitamin D metabolites have not been fully characterized, chromatography indicates small amounts of  $1\alpha,25(\text{OH})_2\text{D}_3$  and larger amounts of its more polar metabolites [31, 33]. Similar findings have been observed in man [34] and rats [35] after administration with radiolabeled  $25\text{OHD}_3$ . A considerable portion of the radiolabeled  $1\alpha,25(\text{OH})_2\text{D}_3$  and  $25\text{OHD}_3$  dose found in the bile is subsequently absorbed from the intestine and presumably delivered back to the liver [32, 34]. Thus, although biliary excretion may not be an efficient pathway of vitamin D clearance, it may still expose enterocytes to higher unbound concentrations of active hormone (through vectorial transport through the enterocytes) than would otherwise occur by vascular delivery and represent an important element of gene regulation [34, 36].

A key initial step in enterohepatic cycling of  $1\alpha,25(\text{OH})_2\text{D}_3$  and  $25\text{OHD}_3$  is conversion to a polar metabolite suitable for canalicular excretion. Glucuronidation of different vitamin D species has been reported [37–41]. In humans, the conjugation of  $1\alpha,25(\text{OH})_2\text{D}_3$  is catalyzed primarily by UGT1A4 and to a much lesser extent by UGT2B4 and UGT2B7 [41]. Three isomers are produced, with the 25-O-glucuronide isomer being the dominant product. Delivery of 25-O-glucuronide of  $1\alpha,25(\text{OH})_2\text{D}_3$  stimulates gene expression in mouse duodenum and colon tissues [42]. Glucuronides deposited in the gastrointestinal lumen are presumably subject to hydrolysis by  $\beta$ -glucuronidases (human epithelial or bacterial) after being secreted from the liver into the bile. Intraluminal release of  $1\alpha,25(\text{OH})_2\text{D}_3$  and  $25\text{OHD}_3$  and subsequent reabsorption may serve as a critical tissue-specific hormone delivery pathway, giving explanation for a similar pattern of preferential expression of *CYP3A4* [28], *TRPV6* and calbindin D9K (Figure 2) in proximal sections of the small intestine, compared to more distal regions.

## 3. Biotransformation of Vitamin D by CYP3A4

The bioactivation of  $25\text{OHD}_3$  involves  $1\alpha$ -hydroxylation, a reaction catalyzed primarily by CYP27B1 in the renal tubular epithelium [43]. The renal production of  $1\alpha,25(\text{OH})_2\text{D}_3$  is tightly regulated by other hormones such as parathyroid hormone (PTH) and fibroblast growth factor-23 (FGF23), which up- and down-regulate expression of the *CYP27B1* gene, respectively [44]. Furthermore, a variety of cell types in different organs of the human body contain the enzyme CYP27B1 and have the capacity to synthesize  $1\alpha,25(\text{OH})_2\text{D}_3$ . It has been shown that local production of  $1\alpha,25(\text{OH})_2\text{D}_3$  within nonrenal tissues is regulated in a manner different from that seen in the kidney, and thus,  $1\alpha,25(\text{OH})_2\text{D}_3$  production and action can be cell specific [45, 46]. Hydroxylation at the alkyl side chain of  $1\alpha,25(\text{OH})_2\text{D}_3$  or  $25\text{OHD}_3$  is considered to be a critical first step in the hormone inactivation pathway [47] (Figure 3). Mitochondrial CYP24A1 is recognized as a key enzyme for hydroxylation at the C-24 and C-23 positions of both  $25\text{OHD}_3$  and  $1\alpha,25(\text{OH})_2\text{D}_3$  [48, 49] and is found in kidney, intestine, lymphocytes, fibroblasts, bone, skin, macrophages and possibly other tissues where  $1\alpha,25(\text{OH})_2\text{D}_3$  exerts its biological effects [50]. However, Gupta et al [51, 52] reported that CYP3A4 exhibits significant 24- and 25-hydroxylation activities for  $1\alpha$ -

OHD<sub>3</sub>, 1 $\alpha$ -OHD<sub>2</sub> and vitamin D<sub>2</sub>, after screening 16 major human hepatic P450s expressed in baculovirus infected insect cells. Moreover, Xu et al [53] reported that CYP3A4 can catalyze the 23- and 24-hydroxylation of 1 $\alpha$ ,25(OH)<sub>2</sub>D<sub>3</sub> in the human liver and small intestine. In addition, it was recently shown that CYP3A4 can catalyze 25OHD<sub>3</sub> monohydroxylation, generating nine different isomeric metabolites [54]. Interestingly, the major product generated by CYP3A4, 4 $\beta$ ,25(OH)<sub>2</sub>D<sub>3</sub>, was detected in human plasma at concentrations comparable to that of 1 $\alpha$ ,25(OH)<sub>2</sub>D<sub>3</sub> and its formation was induced by rifampin [54, 55]. Thus, intestinal and hepatic CYP3A4 may also contribute to the metabolic clearance of 1 $\alpha$ ,25(OH)<sub>2</sub>D<sub>3</sub> and 25OHD<sub>3</sub>, particularly under conditions of induced *CYP3A4* expression.

#### 4. Potential Clinical Implications

Osteomalacia can be a serious, debilitating side effect from certain drug therapies. It has been associated most strongly with chronic administration of anti-epileptic drugs, such as phenobarbital, carbamazepine, phenytoin and valproic acid [56–58], as well as the antimicrobial drug rifampin [59, 60]. Although the molecular mechanism underlying this event is not fully understood, a number of theories have been proposed to explain why anti-epileptic drugs affect bone; this includes reduced levels of vitamin D metabolites, reduced calcium absorption, inhibition of the cellular response to PTH, hyperparathyroidism, vitamin K deficiency and calcitonin deficiency [61]. Among them, vitamin D deficiency is thought to be the principal pathway to drug-induced osteomalacia [62, 63]. For example, daily administration of rifampin to healthy volunteers lowered the plasma concentration of 25OHD<sub>3</sub> by 70% [60, 64]. Similar effects were also described for patients receiving phenobarbital and phenytoin for the treatment of epilepsy [65]. It has been suggested that alterations in vitamin D metabolism result mainly from induction of hepatic P450 enzymes [66, 67]. Pascussi et al [68] proposed that induction of CYP24A1 accelerates vitamin D catabolism, and that PXR is responsible for the regulation of CYP24A1 enzyme. In contrast, Zhou et al [69] reported that CYP3A4, and not CYP24A1, is the dominant enzyme catalyzing hydroxylation of 1 $\alpha$ ,25(OH)<sub>2</sub>D<sub>3</sub> in human liver and small intestine. Thus, the relative contribution of these two enzymes to vitamin D catabolism may be tissue-specific. In the healthy kidney, CYP24A1 activity likely dominates 1 $\alpha$ ,25(OH)<sub>2</sub>D<sub>3</sub> and 25OHD<sub>3</sub> catabolism; while in the liver and small intestine, CYP3A4 activity most likely dominates vitamin D catabolism because of a much greater level of basal and induced enzyme expression [69]. This interpretation is supported by the observation that treatment of healthy volunteers with rifampin causes preferential increases in human duodenal *CYP3A4*, but not *CYP24A1*, mRNA content [69]. Importantly, induction of CYP3A4 in the small intestine could cause local tissue vitamin D deficiency and possibly directly affect intestinal calcium absorption [11, 70]. On the other hand, induction of hepatic CYP3A4 may alter systemic circulating levels of 25OHD<sub>3</sub> through activation of the 4 $\beta$ -hydroxylation pathway [54]. Therefore, inhibition of *CYP3A4* activity in the enterohepatic circuit may constitute a viable therapeutic approach for prevention or reversal of drug-induced osteomalacia in at-risk patients.

Liver function is crucial for physiological vitamin D metabolism. Patients with chronic liver disease and cirrhosis have very high prevalence of vitamin D deficiency and insufficiency [71]. Low vitamin D levels in these patients are possibly due to limited sunlight exposure, impaired intestinal absorption and decreased hepatic 25-hydroxylation activity [71]. Given the correlation between plasma levels of vitamin D metabolites and intestinal CYP3A4 content/activity, low levels of vitamin D metabolites in patients with cirrhosis might result in a reduction in CYP3A protein expression. Indeed, it has been noted that intestinal CYP3A4 expression and function are reduced in patients with cirrhosis [72]. Although reduced metabolism of drugs is typically attributed to decreased liver function, decreased

intestinal CYP3A activity may also contribute to greater drug exposure in these patients. Whether vitamin D supplementation could rescue intestinal CYP3A expression in these patients is something to be considered.

## 5. Concluding remarks

Hormonal control of *CYP3A4* expression by vitamin D represents the foundation of a potentially important interplay between xenobiotic and vitamin D metabolism. Enterohepatic cycling of vitamin D could be a functionally important pathway for delivery of active hormone to the upper intestine, resulting in preferentially higher levels of expression of VDR target genes, such as *TRPV6*, calbindin D9K and *CYP3A4*, in the duodenum and jejunum, in comparison to the ileum and colon. Intra- and inter-individual differences in vitamin D levels may contribute to the considerable variability in intestinal CYP3A4 content that affects drug disposition and pharmacological response. Interestingly, CYP3A4 catalyzes vitamin D biotransformation down pathways that appear catabolic in nature. Certain drugs, such as anti-epileptic drugs, that can induce *CYP3A4* expression in the liver and small intestine, accelerate vitamin D catabolism and may contribute to vitamin D deficiency, although a causal mechanistic link between CYP3A4 induction and vitamin D deficiency requires further evaluation.

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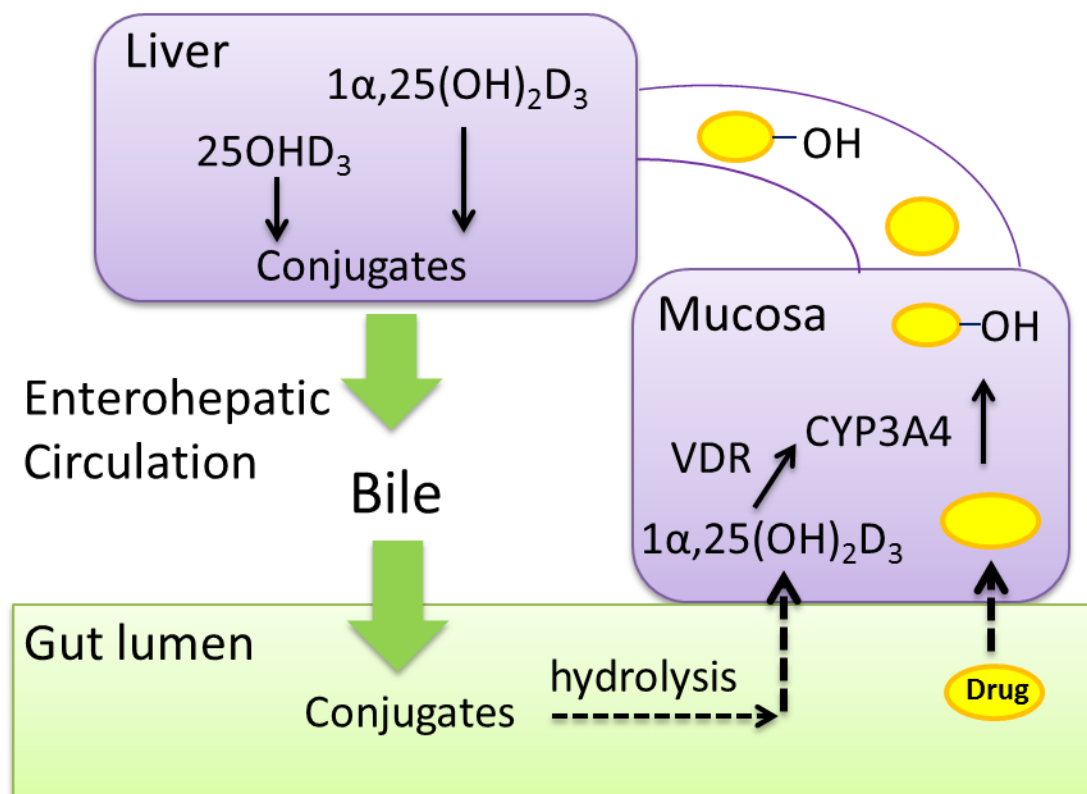
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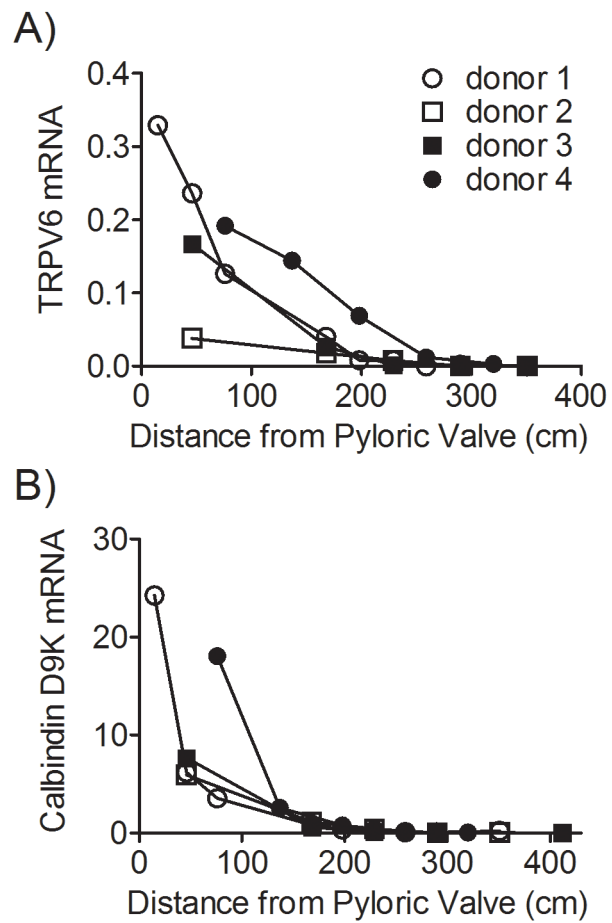
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### Highlights

1. Regulation of *CYP3A4* gene expression by vitamin D
2. Enterohepatic circulation of vitamin D metabolites
3. Metabolism of vitamin D by CYP3A4
4. Vitamin D deficiency and clinical implications

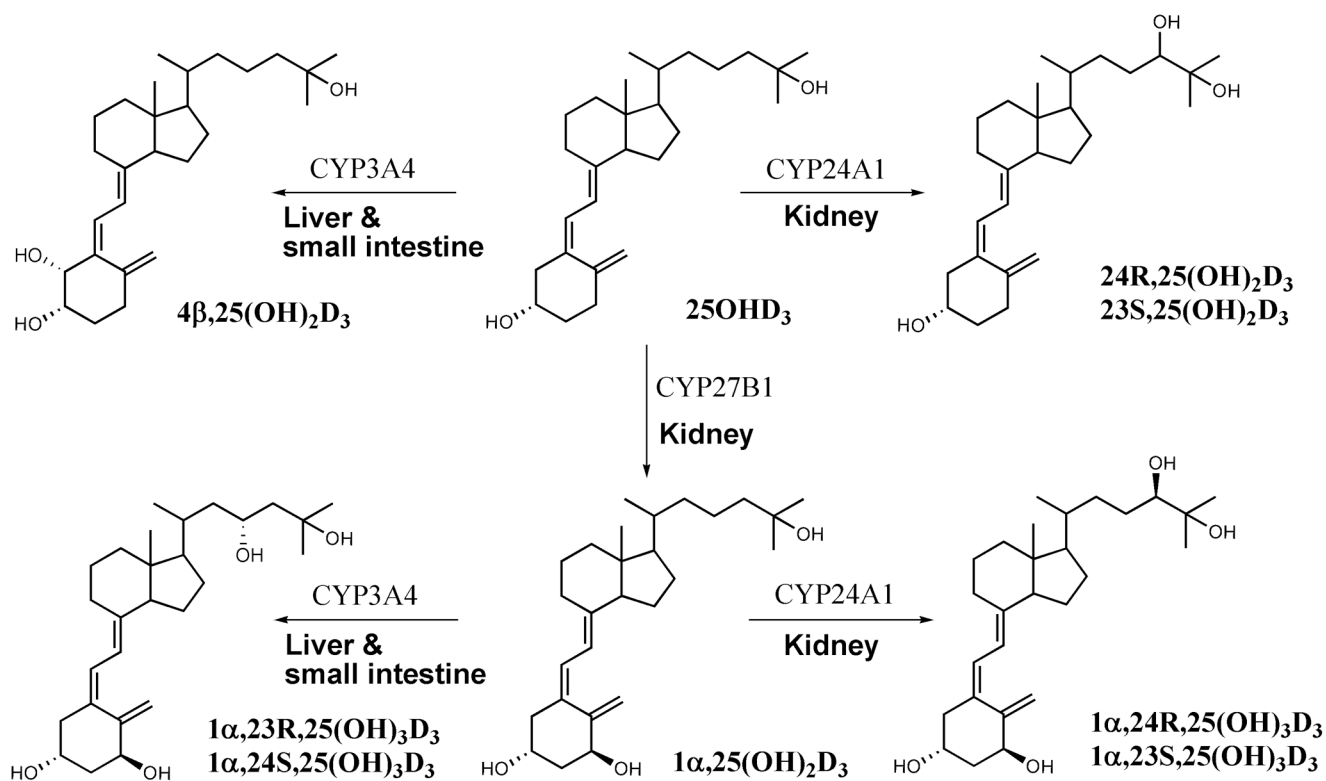


**Figure 1.** Proposed mechanism that enterohepatic circulation delivers vitamin D derivatives to the intestine and  $1\alpha,25(\text{OH})_2\text{D}_3$  induces intestinal *CYP3A4* expression via VDR-mediated pathway.



**Figure 2. Longitudinal distribution of TRPV6 and calbindin D9k mRNA in the human small intestine**

Tissue samples from four different donors were presented. Isolation of total RNA and real-time quantitative PCR were performed as previously described [53]. A) the levels of TRPV6 mRNA were normalized to the corresponding GAPDH mRNA; B) the levels of Calbindin D9k mRNA were normalized to villin mRNA.



**Figure 3. Tissue-specific metabolism of vitamin D metabolites**

Under the constitutive condition in the kidney, CYP27B1 catalyzes  $1\alpha$  hydroxylation of  $25\text{OHD}_3$  for vitamin D activation. While CYP24A1 catalyzes 24R- and 23S-hydroxylation of  $25\text{OHD}_3$  and  $1\alpha,25(\text{OH})_2\text{D}_3$  for vitamin D deactivation. In the liver, CYP3A4 is the most abundant enzyme that catalyzes  $4\beta$ -hydroxylation of  $25\text{OHD}_3$ , and 23R- and 24S-hydroxylation of  $1\alpha,25(\text{OH})_2\text{D}_3$  for vitamin D deactivation.