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Analysis of 1,25-dihydroxyvitamin D₃ genomic action reveals calcium regulating and
 calcium independent effects in mouse intestine and human enteroids

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37 Abstract

Although vitamin D is critical for the function of the intestine, most studies have focused on the 38 39 duodenum. We show that transgenic expression of the vitamin D receptor (VDR) only in the 40 distal intestine of VDR null mice (KO/TG mice) results in the normalization of serum calcium 41 and rescue of rickets. Although it had been suggested that calcium transport in the distal intestine involves a paracellular process, we found that the $1,25(OH)_2D_3$ activated genes in the 42 proximal intestine associated with active calcium transport (Trpv6, S100g, Atp2b1) are also 43 44 induced by $1,25(OH)_2D_3$ in the distal intestine of KO/TG mice. In addition, *Slc30a10*, a manganese efflux transporter, was one of the genes most induced by $1,25(OH)_2D_3$ in both 45 46 proximal and distal intestine. Both villus and crypt were found to express Vdr and VDR target genes. RNA-seq analysis of human enteroids indicated that the effects of 1,25(OH)₂D₃ observed 47 in mice are conserved in humans. Using Slc30a10^{-/-} mice, a loss of cortical bone and a marked 48 decrease in S100g and Trpv6 in the intestine was observed. Our findings suggest an 49 interrelationship between vitamin D and intestinal Mn efflux and indicate the importance of 50 distal intestinal segments to vitamin D action. 51

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68 Introduction

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The essential role of vitamin D is to mediate calcium homeostasis through its regulatory 70 effects on intestine, bone and kidney (1). Vitamin D has also been reported to affect numerous 71 72 other physiological processes not involved in calcium regulation including immunomodulatory 73 effects and inhibition of cancer progression (1). The skeletal and extraskeletal effects of 74 $1,25(OH)_2D_3$ (the biologically active from of vitamin D; calcitriol) are mediated by the vitamin D receptor (VDR), a nuclear protein which heterodimerizes with the retinoid X receptor (RXR). 75 VDR/RXR interacts with DNA sequences, vitamin D response elements (VDREs) in target 76 genes and leads to activation or repression of transcription. 1,25(OH)₂D₃ also recruits 77 coregulatory complexes that participate in the regulation of transcription of VDR target genes. 78 79 VDR-mediated gene transcription is a multi-step process requiring the spatial and sequential combination of transcriptional coactivators and is influenced by the particular subcellular 80 environment in a gene specific manner (2,3). 81

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A principal target tissue of vitamin D action is the intestine (1.4.5). However, little is 83 known about the molecular targets mediating the effects of vitamin D on intestinal biology. The 84 classic role of $1,25(OH)_2D_3$ is regulation of intestinal calcium absorption which is crucial for 85 bone health. When the need for calcium increases, for example during periods of habitual low 86 87 calcium intake and during growth, vitamin D mediated intestinal calcium absorption occurs predominantly by an active transcellular process. The most pronounced effects of $1,25(OH)_2D_3$ 88 during the active calcium transport process are increased synthesis of the epithelial calcium 89 channel TRPV6 and the intracellular calcium binding protein calbindin- D_{9k} (4,5). In addition to 90 the active transcellular pathway, calcium can also be absorbed via a non-saturable paracellular 91 process (4,5). Whether $1,25(OH)_2D_3$ regulates calcium absorption by modulating genes 92 associated solely with the facilitated diffusion model or whether $1,25(OH)_2D_3$ affects the 93 paracellular process has been a matter of debate (6-9). Most research on calcium absorption has 94 95 utilized the proximal small intestine and early studies reported that 1,25(OH)₂D₃ regulated active calcium transport was localized only in the duodenum (9). It had been suggested that calcium 96

97 absorption in the distal intestine reflects vitamin D independent passive diffusion (10). However 98 previous studies also noted that $1,25(OH)_2D_3$ and low dietary calcium can regulate active 99 calcium absorption in the distal intestine of rats (11-13). In addition, we recently showed that 100 transgenic expression of VDR only in the distal intestine (distal ileum, cecum and colon) at 101 levels equivalent to wild type (WT) mice results in normal calcium homeostasis and reverses the 102 VDR dependent rickets of the VDR null mice (14). These findings indicate the importance of the 103 distal intestinal segments to vitamin D action.

104 Although the classical intestinal role of 1,25(OH)₂D₃ is regulation of calcium absorption, $1,25(OH)_2D_3$ has been reported to have other important effects on the intestine including 105 protection of the integrity of the mucosal barrier, regulation of pathogen invasion, apoptosis, 106 107 nutrient transport and cellular differentiation (7,15-19). However, the molecular responses to 108 $1,25(OH)_2D_3$ resulting in these effects in proximal vs. distal intestine are poorly understood. In 109 addition, although intestinal epithelial cells at the villi have been identified as targets for $1,25(OH)_2D_3$, little is known about the effect of $1,25(OH)_2D_3$ in crypts and whether $1,25(OH)_2D_3$ 110 111 responses occur in crypts has been a matter of debate (20-22). Thus, the diversity and complexity 112 of vitamin D signaling in the intestine have not been fully clarified.

In this study in order to further understand the role of vitamin D in the intestine, we 113 114 examined the expression of VDR targets using transgenic mice with expression of VDR restricted to the distal intestine (KO/TG mice) (14) and compared our findings in these 115 116 transgenic mice to vitamin D targets in both proximal and distal intestine of vitamin D deficient 117 mice treated with $1,25(OH)_2D_3$. The classic $1,25(OH)_2D_3$ activated genes found in the proximal intestine and associated with active calcium transport (Trpv6, S100g, Atp2b1) were also induced 118 119 in the distal intestine. In addition, one of the genes most induced by $1,25(OH)_2D_3$ in the proximal 120 and distal intestine was *Slc30a10*, a manganese (Mn) efflux transporter which has recently been found to be critical for protection against neurotoxicity and liver injury which occur in the 121 122 presence of elevated Mn levels (23). Both villus and crypts were found to express high levels of VDR and result in 1,25(OH)₂D₃ mediated target gene induction. RNA-seq analysis of human 123 124 enteroids confirmed the induction of the same genes in mice and humans (TRPV6, SLC30A10, 125 ATP2B1 and CYP24A1) in both villus and crypts indicating for the first time direct 126 transcriptomic responses to $1,25(OH)_2D_3$ in human enteroids in both crypt and villus-like

compartments. Studies in Slc30a10^{-/-} mice showed a loss of cortical bone and a marked decrease 127 in S100g and Trpv6 in the intestine of these mice. Our findings suggest that $1,25(OH)_2D_3$ may 128 have a role not only to maintain calcium homeostasis but also in the cellular homeostasis of other 129 130 divalent ions. Our findings emphasize the importance of the distal as well as the proximal intestine in order to understand intestinal effects of $1.25(OH)_2D_3$ and that the effects of 131 1,25(OH)₂D₃ involve intestinal epithelial cells in both villus and crypt. 132

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Results 134

RNA-seq Analysis of Transcriptomic Responses to 1,25(OH)₂D₃ in the Colon of Mice with 135 136 **Transgenic Expression of VDR Only in the Distal Intestine**

137 Although most studies have focused on calcium absorption in the proximal intestine, our 138 previous study demonstrated that transgenic expression of VDR only in the distal intestine rescued VDR-dependent rickets observed in the VDR KO mouse. To understand mechanisms by 139 which $1,25(OH)_2D_3$ acts to regulate intestinal biology not only in the proximal but also in the 140 141 distal intestine, transcriptomic profiling was done to examine the expression of genes induced by 142 1,25(OH)₂D₃ in the distal intestine of KO/TG mice (KO/TG line 1, noted as TG1) which express VDR exclusively in the distal intestine (distal ileum, cecum and colon) (14; Fig. 1A, B). We 143 focused on the colon as a possible unappreciated target for regulation of calcium homeostasis. As 144 145 noted in **Fig. 1C** the classic $1,25(OH)_2D_3$ activated genes in the proximal intestine were also the genes most markedly induced by $1,25(OH)_2D_3$ in the colon of the TG mice (Cyp24a1, Trpv6 and 146 147 Sl00g). Induction of Sl00g, Trpv6 as well as Atp2b, genes involved in active calcium transport (1), suggests that active calcium transport in the distal intestine is involved in the normalization 148 149 of mineral homeostasis in these mice. Genes associated with passive calcium transport, Cldn2, 150 Cldn12 or Cacnald (Cav1.3 gene) and Trpm7 which have been suggested as channels other than Trpv6 for regulation of intestinal calcium absorption (6,24,25), were not significantly regulated 151 152 by $1,25(OH)_2D_3$ in the colon. In addition to the classic $1,25(OH)_2D_3$ activated genes, *Slc30a10*, a manganese efflux transporter located in the apical/luminal domain, was also one of the genes 153 154 most induced by $1,25(OH)_2D_3$ and was the principal Slc transporter gene affected by 1,25(OH)₂D₃. Other Slc transporters (Slc30a1 (a Zn transporter) as well as Slc30a4 and Slc30a5 155 (also Zn transporters; not shown), Slc39a14 (a Mn transporter which mediates Mn reuptake from 156 157 the basolateral membrane) and *Slc37a2* (a phosphate linked glucose 6 phosphate antiporter) were

not significantly affected by $1,25(OH)_2D_3$. RT-PCR analysis of intestinal tissues from the same transgenic mouse line (KO/TG1) (**Fig.2A**) as well as from a second independent KO/TG line (KO/TG2, noted as TG2) (**Fig. 2B**) validated the RNA-seq analyses indicating that the genes most responsive to $1,25(OH)_2D_3$ in the distal intestine of KO/TG mice include *Cyp24a1*, *Trpv6*, *Sl00g* and *Slc30a10*. Other *Slc* transporters, *Slc30a1* and *Slc37a2* were unaffected by $1,25(OH)_2D_3$ (**Fig. 2**). Due to the expression of VDR only in the distal intestine, the duodenum of KO/TG mice is unresponsive to $1,25(OH)_2D_3$ (**Fig. 2**; duo).

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166 Regulation of Vitamin D Target Genes in the Intestine

In addition to the induction of Slc30a10 in the distal intestine of transgenic mice, Slc30a10 167 was found to be regulated by $1.25(OH)_2D_3$ in different intestinal segments (duodenum, ileum and 168 169 colon) of vitamin D deficient mice similar to other vitamin D target genes (Cyp24a1, S100g). 170 Slc37a2 although not regulated by $1,25(OH)_2D_3$ in the distal intestine, was induced by 171 $1,25(OH)_2D_3$ in the duodenum (Fig. 3). In addition, in the duodenum expression of Slc30a4, Slc30a5, Slc39a14, Trmp7, Cldn12 and Cacna1d, similar to findings in the colon (Fig. 1) was 172 unaffected by $1,25(OH)_2D_3$ treatment of vitamin D deficient mice (n = 4, vehicle vs. 173 $1,25(OH)_2D_3$ treated; p > 0.5). Cldn2, however was significantly induced by $1,25(OH)_2D_3$ in the 174 duodenum of these mice (3.8 fold; n=4 vehicle vs. $1,25(OH)_2D_3$ treated; p < 0.05). 175

176 To gain insight into a possible role of the vitamin D in the regulation of Mn transport, further studies were done comparing the regulation of *Slc30a10* in the intestine to the regulation 177 of known vitamin D target genes. As shown in Fig. 4A, Slc30a10 and Trpv6 showed a similar 178 179 time course of induction by 1,25(OH)₂D₃. Gene expression for these transporters was 180 significantly induced in the duodenum at 4h after injection and then decreased to control levels at 24 h after $1.25(OH)_2D_3$ administration. Similar findings were observed for Slc30a10 and Trpv6 181 182 in colon (data not shown). These findings suggest a rapid response of Trpv6 and Slc30a10 to 1,25(OH)₂D₃ for intestinal calcium or Mn transport. 183

In addition to regulation by $1,25(OH)_2D_3$, effects of dietary calcium on the expression of these genes was also examined. Four-week-old mice were fed either a high calcium (1%) or low calcium diet (0.02%) diet for 4 weeks. As shown in **Fig. 4B**, under conditions of low dietary calcium (which results in increased circulating $1,25(OH)_2D_3$ levels) (1) an induction in both *Slc30a10* and *Trpv6* is observed in the duodenum. A similar induction of these genes in response to low dietary calcium was also observed in the colon (not shown). *S100g* was also significantly
increased in duodenum and colon in response to low dietary calcium (data not shown). These
findings suggest coregulation of these manganese and calcium transporters under conditions of
dietary calcium alteration.

In additional studies developmental changes in the expression of Slc30a10 and Trpv6 193 were compared. Trpv6 was not expressed significantly in the intestine until after birth and was 194 195 induced at three weeks of age, the time of weaning in mice which coincides with the increase in 196 active intestinal calcium transport (Fig. 4C) (26). Unlike Trpv6, Slc30a10 was detected in the 197 fetal intestine. The most marked induction in the expression of Slc30a10 was observed between the expression in fetal intestine and at one week postpartum. Thus, developmental changes in 198 intestinal Slc30a10 and Trpv6 do not coincide, suggesting that different factors modulate the 199 200 developmental expression of these transporters.

To further understand a possible interrelationship between vitamin D and manganese 201 202 homeostasis, vitamin D target genes associated with $1,25(OH)_2D_3$ mediated calcium transport were examined in the intestine of the $Slc30a10^{-/-}$ mouse which has manganese levels in blood, 203 brain and liver 20-60 fold higher than controls (27). Expression of SlOOg and Trpv6 was 204 markedly decreased in the duodenum of the $Slc30a10^{-/-}$ mice (> 10-fold Fig. 5A) but not 205 significantly changed in the colon. Also, no change was observed in the duodenum and colon in 206 the expression of Tip1 (ZO-1 gene; previous reports suggested regulation by $1.25(OH)_2D_3$ of 207 intestinal tight junction protein ZO-1) (28). In order to determine the specificity for vitamin D 208 209 regulated calcium transporters in the intestine, Trpv5 and S100g were examined in the kidney of the $Slc30a10^{-/-}$ mice. S100g and Trpv5 were unaffected in the kidney (Fig. 5B). 210

Micro-computed tomography (µCT) analysis of the tibiae was performed to assess 211 trabecular and cortical parameters in control and Slc30a10^{-/-} mice. Trabecular bone mass was 212 slightly, although not significantly, reduced in male *Slc30a10^{-/-}* mice versus control mice, 213 whereas no differences in trabecular bone mass were observed in female mice (Table 1). 214 215 However, a strong cortical bone phenotype was observed in both male and female mice. Tibias of *Slc30a10^{-/-}* mice were smaller in diameter than tibias of control mice as evidenced by their 216 217 reduced cross-sectional tissue area. In addition, cortical thickness as well as cortical porosity were significantly reduced in $Slc30a10^{-/-}$ mice at the mid-diaphysis (Table 1). Serum calcium 218 levels were unchanged between $Slc30a10^{-/-}$ and control mice (12.5 ± 1.3 vs. 12.3 ± 0.4 ng/ml 219

220 respectively; n = 5, p > 0.5) Thus, in addition to the reduction in *Trpv6* and *Sl00g*, there is a significant bone phenotype in these mice. 221

222 VDR and Vitamin D Target Genes in Mouse Crypts and Villi and in Differentiated and

223 Undifferentiated Human Enteroids

224 To provide insight into the mechanisms involved in the regulation of vitamin D target genes 225 in the intestine, including the ion transporters, we examined both mouse villi and crypts. 226 Additionally, human enteroid cultures were used to examine both undifferentiated (crypt-like) 227 and differentiated (villus-like) responses to $1,25(OH)_2D_3$ and to determine whether the effects of $1,25(OH)_2D_3$ in the intestine observed in mice are conserved in humans. Although the 228 229 importance of vitamin D regulated proteins in intestinal villi has been reported, little is known about effects of $1,25(OH)_2D_3$ in crypts and few findings related to human intestinal VDR targets 230 231 have been reported. Crypts and villi were isolated from mouse duodenum using the EDTA 232 chelation method (Fig. 6A). Expression of Vdr as well as VDR protein in mouse villus and crypt 233 was equivalent (Fig. 6B). Both villus and crypt respond to $1,25(OH)_2D_3$ by inducing expression of Cyp24a1 (Fig. 6C). Slc30a10 and Trpv6, although more pronounced in villi were also 234 235 expressed in crypts of WT mice (Fig. 6D).

236 RNA-seq analysis of human duodenal enteroids (Fig. 7A-C) showed that intestinal vitamin D 237 target genes are conserved in humans. Similar to the findings in mice, expression of VDR under 238 basal conditions was equivalent in villus and crypt compartments. The expression of VDR was not significantly affected by 1.25(OH)₂D₃ However, TRPV6, SLC30a10, CYP24A1 and ATP2B1 239 were significantly upregulated upon $1,25(OH)_2D_3$ treatment in both villus and crypt (Fig. 7C). 240 241 S100G was significantly induced by $1,25(OH)_2D_3$ only in villus like cultures. A similar 242 regulation of vitamin D targets genes was observed in crypt-like and villus-like human enteroids 243 derived from the colon (data not shown). Gene expression of Slc transporters in addition to 244 SLC30A10 in differentiated and undifferentiated human duodenal enteroids treated with 1,25(OH)₂D₃ or vehicle is shown in **Table 2**. *SLC30A4* and *SLC30A5* were not significantly 245 246 affected by $1,25(OH)_2D_3$. SLC37A2 was induced in villus but not in crypts. The intestinal 247 phosphate transporter, SLC34A2, was upregulated by $1,25(OH)_2D_3$ in both villus and crypt. *TRPM7*, a non-*Slc* transporter, was unaffected by $1,25(OH)_2D_3$ treatment. Gene Ontology (GO) 248 249 analysis was performed to identify biological processes, cellular components and molecular

250 function terms that are enriched in human duodenal enteroids treated with $1,25(OH)_2D_3$. The enrichment score [-log10 (p value)] of the terms were plotted to indicate the significance of the 251 252 enrichment of each function. Regulation of epithelial cell differentiation represented a major category for enrichment of genes by $1,25(OH)_2D_3$ in crypts (Fig. 8). Sonic hedgehog (SHH), 253 which controls the proliferation of stem cells, was a $1,25(OH)_2D_3$ target gene in this ontology 254 group as well as response to nutrient, cell migration and cell motility, indicating a role for 255 $1,25(OH)_2D_3$ in the function of stem cells. Pathway analysis indicated enrichment of genes by 256 257 1,25(OH)₂D₃ involved ion transport, oxidation reduction, vitamin D 24-hydroxylase activity, 258 steroid metabolic processes, as well as calcium ion binding and microsomes and vesicular fraction were categories shared in both villus and crypt (Fig. 8). $1,25(OH)_2D_3$ target gene 259 members of the gene ontology group identifying microsomes and vesicular fraction include 260 261 CYP3A4 which is involved in vitamin D metabolism and CYP2C19 and CYP2C18 involved in 262 drug metabolism.

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264 Discussion

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Vitamin D plays a major role in maintaining the integrity and function of the intestine as 266 267 well as in the control of calcium homeostasis. However, the mechanisms by which $1,25(OH)_2D_3$ 268 regulates these processes are not fully understood. Our findings indicate the importance of the distal as well as the proximal segments of the intestine to understand the intestinal effects of 269 270 $1,25(OH)_2D_3$ and that $1,25(OH)_2D_3$ mediated responses involve epithelial cells in the crypts as well as the villus. This study also shows that the effects of $1,25(OH)_2D_3$ on intestine are complex, 271 272 are conserved in humans for active calcium uptake and include calcium regulating as well as 273 calcium independent effects in both proximal and distal intestine.

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Previous investigations reported that $1,25(OH)_2D_3$ mediated active calcium transport was localized only in the duodenum where the most pronounced effects of $1,25(OH)_2D_3$ during the active calcium transport process were induction of calbindin-D9k (*S100g*) and *Trpv6* (4,5,9). Our current findings indicate that the classic $1,25(OH)_2D_3$ activated genes (*S100g* and *Trpv6*) are also the two genes most markedly induced by $1,25(OH)_2D_3$ in the distal intestine (similar to previous findings in the proximal intestine). In addition, our recent study showing for the first

281 time a similar VDR-dependent calcium absorption efficiency between proximal colon and 282 duodenum in the same mouse further establishes a role for the distal intestine in VDR-mediated 283 intestinal calcium absorption (29). In that study calcitriol glycosides and glucuronides, which 284 target calcium absorption in the distal intestine, were also found to upregulate Trpv6 and S100g (the genes involved in active calcium transport) as well as calcium absorption in the colon. These 285 results which provide direct in vivo evidence for the importance of VDR in the distal intestine, 286 suggest, together with earlier studies (11-14), that active calcium transport is a vitamin D-287 288 dependent process in the distal intestine which plays a role in calcium homeostasis. Calcitriol 289 glucuronide treatment may be a potential means to target calcitriol to the distal intestine and thus compensate for the loss of calcium absorption for example by gastric bypass surgery or small 290 bowel resection. In the current investigation, specific genes previously associated with 291 292 paracellular calcium transport were not significantly changed by 1,25(OH)₂D₃ in mouse distal 293 intestine. However, Cldn2 was significantly induced by $1,25(OH)_2D_3$ in the duodenum. Although 294 it is possible that $1.25(OH)_2D_3$ may be involved in paracellular as well as active calcium 295 transport, the physiological relevance of the reported regulation of genes by $1,25(OH)_2D_3$ 296 involved in paracellular transport remains to be determined (6,7,18).

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As compared to other tissues, the highest levels of VDR are found in the small and large 298 intestine. Studies related to $1,25(OH)_2D_3$ mediated responses have focused on effects in 299 differentiated enterocytes and have supported a role for $1,25(OH)_2D_3$ in intestinal calcium 300 absorption as well as barrier function (4-7). However, the effect of $1,25(OH)_2D_3$ on intestinal 301 302 stem cells has not been well investigated and has been a matter of debate (20-22). Previous studies using oral treatment, reported that 1,25(OH)₂D₃ stimulates Cyp24a1 in intestinal 303 epithelial cells at the villus or tip region and not at the crypts (20). Early microscopic 304 305 autoradiographic studies however noted that tritium labeled 1,25(OH)₂D₃ concentrated in both 306 absorptive and crypt epithelial cells in small and large intestine (30,31). Using a highly specific 307 and sensitive VDR antibody (D6) (32), we also noted, similar to the autoradiographic studies, 308 that VDR is present in villus and crypts. Our studies using isolated mouse villus and crypt as 309 well as human enteroids with either a crypt-like phenotype (high proliferation, undifferentiated) 310 or a villus-like phenotype (low proliferation, differentiated) identified $1,25(OH)_2D_3$ induced 311 target genes, including Cyp24a1, in both villus and crypts. The importance of vitamin D in the

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315 cell functions of the Lgr5+ cells (22). Studies in human colon crypt organoids have also provided evidence that VDR is expressed in intestinal stem cells and has a regulatory role as $1,25(OH)_2D_3$ 316 was found to induce stemness-related genes and to inhibit genes involved in cell proliferation 317 318 (33). We are now only beginning to understand the regulatory role of vitamin D in intestinal 319 stem cells (22,33,34). It is of interest that both in our in vivo studies in mice as well as in our 320 studies using human enteroids, 1,25(OH)₂D₃ was found to regulate classic target genes in both villus and crypt. These findings suggest, similar to early studies showing the time dependency of 321 322 $1,25(OH)_2D_3$ action in enterocytes during their differentiation (35), that $1,25(OH)_2D_3$ action in 323 the intestine involves not only calcium absorption but also cellular differentiation and intestinal 324 homeostasis. Although there have been many studies related to the intestinal effects of vitamin 325 D in animals, our findings are one of the few studies examining human vitamin D target genes. Previous studies have been done using human intestinal cell lines derived from colon (19,36-38) 326 327 as well as human biopsies and duodenal explants (39,40). The limitation of the studies in 328 explants is the inability of the cells to renew as well as cell death in the explants. The enteroid culture system from human tissue allows for growth and maintenance over time and provides an 329 330 important in vitro model to examine both stem and non-stem cell responses to $1,25(OH)_2D_3$. These epithelial cultures of the intestinal tract will provide technology enabling the verification 331 of findings observed in animals and the identification of key components involved in 332 333 1,25(OH)₂D₃ action in human intestinal biology. 334 In addition to Cyp24a1, Trpv6 and S100g, Slc30a10, a manganese efflux transporter is 335 336

function of Lgr5+ stem cells which are sensitive to Wnt stimulation and contribute to intestinal homeostatic regeneration, was previously noted by L. Augenlicht's lab (22). They showed that

inactivation of the VDR in mouse Lgr5+ intestinal crypt base columnar cells compromised stem

also a 1,25(OH)₂D₃ target gene in both proximal and distal intestine. Slc30a10 is localized in the apical/luminal domain of the intestine, in liver and brain (41). Although initially thought to act as 337 338 a Zn efflux transporter, recent studies have identified SLC30A10 as a mediator of Mn efflux 339 which lowers cellular Mn levels and protects against Mn toxicity (23,27,42,43). In clinical 340 studies, patients carrying homozygous mutations in SLC30A10 show enhanced Mn levels, 341 neurotoxicity and liver injury (43,44). It is of interest that Mn levels in the brain are minimally 342 elevated in liver specific Slc30a10 KO mice (41,45). However endoderm specific KO mice

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Mn levels in the brain, indicating a critical role for the digestive system in the regulation of brain 344 345 manganese (41,45). Our results suggest that 1,25(OH)₂D₃, by inducing Slc30a10 in the intestine, may increase Mn excretion and thus may have therapeutic benefit without direct effects on the 346 brain. Preliminary data comparing mice injected with MnCl₂ (15 µg/g bw) alone or MnCl₂ + 347 $1,25(OH)_2D_3$ (1ng/g bw) 3 times per week for 3 week showed that $1,25(OH)_2D_3$ treatment results 348 349 in decreased levels of blood Mn (control 30 ± 2.3 ; Mn, 403 ± 91 ; Mn $\pm 1,25$ (OH)₂D₃, 288 ± 3.3 350 ng Mn/ml; S. Mukhopadhyay and S. Christakos, preliminary results). Further studies will be 351 needed to determine effects of $1,25(OH)_2D_3$ on neurotoxicity, on levels of Mn in different tissues as well as effects of vitamin D deficiency, dietary calcium and high dose dietary vitamin D on 352 353 Mn toxicity. Although elevated levels of Mn are neurotoxic, Mn is an essential element which 354 plays a role in many cellular processes (46). With regard to bone homeostasis, it has been 355 reported that the skeleton is positively influenced by Mn and that deficiency in Mn affects bone 356 mass. Studies in rats have shown that Mn supplementation inhibits ovariectomy induced bone loss (47). In addition, plasma levels of Mn were reported to be significantly decreased in groups 357 of osteoporotic patients (48). More recent studies have noted that local manganese chloride 358 359 treatment accelerates fracture healing in a rat model (49). However, excess of Mn can adversely affect bone quality. In our study using the Slc30a10 global KO mouse, which exhibits marked 360 361 elevations in blood Mn, significant cortical bone abnormalities were noted. These findings 362 further indicate the importance of maintaining Mn levels at an optimal range and not exceeding the levels required to accomplish essential functions. The mechanisms involved in the adverse 363 364 effects on bone observed in the Slc30a10 KO mouse are not known. Whether the reduced 365 thyroxine production observed in the Slc30a10 KO mouse plays a role in the bone phenotype remains to be determined (50). The precise mechanism whereby Mn affects thyroxine 366 367 biosynthesis is not known. It is of interest that a patient with a loss of function of SLC30A10 mutation was also identified with hypothyroidism (51). 368 369

(lacking Slc30a10 in liver and the gastrointestinal tract) were found to have markedly elevated

370 Due to the necessity of maintaining Mn levels in a narrow range in order to prevent 371 toxicity, understanding the regulation of SLC30A10 is critical. The regulation of Slc30a10 in the 372 intestine by 1,25(OH)₂D₃ suggests that 1,25(OH)₂D₃ not only affects absorption of calcium but 373 may also affect the transport of other divalent cations. The time course of response to

374 1,25(OH)₂D₃ showed that expression of *Trpv6* and *Slc30a10* is significantly elevated 4 h after 1,25(OH)₂D₃. 24 h after administration of 1,25(OH)₂D₃ the expression of the genes for both 375 376 transporters is decreased to levels observed prior to 0 h, consistent with a rapid response of both transporters to $1,25(OH)_2D_3$ as previously noted for *Trpv6* expression (52). The early induction 377 of these transporters would be needed for rapid calcium uptake or Mn efflux. Although putative 378 379 vitamin D response elements have been noted in the human and mouse gene which encode 380 SLC30A10 (18,19), studies are needed to determine VDR binding sites which are functionally 381 linked to the expression of these genes. The significance of a possible inhibitory effect of 382 unliganded VDR also requires further studies (53,54).

To further understand the role of vitamin D in manganese homeostasis, the expression of 383 384 the classic vitamin D target genes was examined in the intestine of the Slc30a10 KO mouse. In 385 the Slc30a10 KO mouse a marked decrease in the expression of Trpv6 and S100g in the 386 duodenum, a primary route of manganese excretion (55), was consistently observed. These findings suggest that TRPV6, S100G and SLC30A10 may work together in manganese efflux 387 388 transport. Previous studies using HEK cells transfected with human (h) TRPV6 showed that 389 TRPV6 is permeable to Mn as well as other heavy metal cations, indicating that TRPV6 is not only involved in calcium transport (56). Since intracellular calcium concentrations have been 390 reported to correlate with the level of expression of TRPV6 (56), our findings of a decrease in 391 392 Trpv6 (and S100g) in the intestine of mice with Slc30a10 deficiency and defective Mn excretion suggest the possibility of intracellular calcium mediated regulation of intestinal Slc30a10. 393 394 Kovacs et al observed an inhibition of calcium uptake mediated by TRPV6 in the presence of high concentrations of Mn (56). Therefore, it is also possible that low levels of expression of the 395 396 genes involved in calcium homeostasis may be due to the high Mn levels in the Slc30a10 KO 397 mice. Although not previously identified *in vivo* in the intestine, an interrelationship between 398 calcium and Mn transport has been noted in other reports including studies in mast cells, in 399 mitochondria and in brain (57-62). It has been suggested that Mn may interfere with brain 400 calcium homeostasis (62). In the mitochondria, Mn transport has been reported to be activated by 401 calcium (58,60). However calcium transport in the mitochondria is inhibited in the presence of Mn (58). In recent studies using HEK cells, it was noted that Slc30a10 extrusion of Mn was 402 403 calcium-dependent (63). The authors suggested that Slc30a10 uses a calcium gradient for active 404 counter ion exchange (63). Whether Mn efflux transport is regulated in part by calcium in the

405 intestine (as suggested by our findings) and whether intestinal calcium transport is inhibited in 406 the presence of high blood Mn levels (which are observed in the Slc30a10 KO mouse) remains to 407 be determined. It is possible that the role of the distal intestine, where Slc30a10 is also induced 408 by $1.25(OH)_2D_3$, may be compensatory when Mn efflux from the proximal intestine is disrupted. The relationship between calcium, the vitamin D endocrine system and Mn efflux in the intestine 409 is an unexplored topic. Since the digestive system has recently been identified as a critical 410 411 regulator of Mn toxicity (41,45), further studies are needed to provide new insight on the role of 412 calcium and the vitamin D on Mn homeostasis and protection against neurotoxicity.

Findings in our lab as well as prior reports have noted that $1,25(OH)_2D_3$ treatment affects 413 the gene expression of intestinal transporters in addition to Slc30a10. Expression of Slc37a2, 414 which encodes a phosphate linked glucose 6 phosphate antiporter, was previously shown to be 415 416 induced by $1,25(OH)_2D_3$ in mouse intestine (18). Although not regulated by $1,25(OH)_2D_3$ in the mouse distal intestine, we found that SLC37A2 was induced by $1,25(OH)_2D_3$ in human enteroids 417 from duodenum (Table 2) and colon (data not shown). Regulation of SLC37A2 by $1,25(OH)_2D_3$ 418 419 is not specific to the intestine. SLC37A2 has also been suggested as a gene target and marker of 420 vitamin D status in hematopoietic cells (64). Although SLC37A2 can transport glucose 6 phosphate (G6P), recent findings suggested that SLC37A2 does not have a role in the regulation 421 of blood glucose (65). Since the functional role of SLC37A2 is not understood at this time, the 422 423 physiological significance of regulation of gene expression by $1,25(OH)_2D_3$ in both mouse and human intestine remains to be determined. Although we did not find that 1,25(OH)₂D₃ affected 424 425 major Zn transporters, previous studies noted regulation of zinc transporters by $1,25(OH)_2D_3$ in mouse intestine (18). Both calcium and zinc are known to play important roles in normal 426 427 development and bone homeostasis (66-68). Early studies in chicks, however, noted that vitamin 428 D or $1,25(OH)_2D_3$ does not affect zinc absorption in zinc deficient or replete animals (69). Due to the importance of zinc as well as calcium for skeletal integrity, further studies are needed to 429 430 determine an interrelationship between vitamin D and zinc whether vitamin D has an effect on zinc homeostasis. In addition to regulation of SLC30A10 as well as SLC37A2, SLC34A2, the 431 intestinal cotransporter NaPi-IIb was noted to be a gene target induced by $1,25(OH)_2D_3$ in human 432 enteroids derived from duodenum (SLC34A2 is expressed in the proximal intestine in humans 433 and primarily in the ileum in mice). $1,25(OH)_2D_3$ is known to stimulate intestinal phosphate 434 435 absorption (70) and SLC34A2 has been reported to protect bone when dietary phosphate is

436 restricted (71). However, the role of $1,25(OH)_2D_3$ in the regulation of intestinal SLC34A2 (NaPi-IIb) has been a matter of debate (72-74). Further studies are needed to examine the regulation of 437 438 this phosphate transporter in human intestine, its physiological significance in bone and phosphate homeostasis in man and whether it has a role in the dysregulation of phosphate 439 440 observed in chronic kidney disease.

441 In a recent study intestine specific knockout of *Trpm7*, which encodes a channel kinase 442 suggested to control magnesium levels, resulted in decreased serum zinc and calcium levels as 443 well as deceased serum magnesium by post-natal day 5 (mice died by post-natal day 10) (25). The authors suggested that TRPM7 is the common gatekeeper for these ions in the intestine and 444 that TRPM7 and not TRPV6 is the key factor in intestinal calcium absorption. However, these 445 findings were observed in mice prior to the onset of vitamin D receptor mediated active intestinal 446 447 calcium absorption (which occurs at weaning). In addition, results of our transcriptomic analyses did not indicate regulation of Trpm7 by $1,25(OH)_2D_3$ in the intestine of mice and TRPM7448 expression was unaffected by 1,25(OH)₂D₃ treatment in human enteroids. Thus, although 449 450 TRPM7 is important in early post-natal life, TRPV6 and not TRPM7 is involved in the process 451 of 1,25(OH)₂D₃ regulated intestinal calcium absorption.

452 In summary, our findings suggest the importance of the distal as well as the proximal 453 intestinal segments in order to understand intestinal effects of vitamin D and that $1,25(OH)_2D_3$ effects involve intestinal epithelial cells in both the villus and crypt. In addition, our findings 454 show direct transcriptomic responses to 1,25(OH)₂D₃ in human enteroids in both villus and crypt 455 456 and that effects observed in mice are conserved in humans. Since SLC30A10 is the first reported transporter that uses a calcium gradient for active counter ion exchange, our findings suggest that 457 458 TRPV6, S100G and SLC30A10 work together in Mn transport and that $1,25(OH)_2D_3$ may have a 459 role not only to maintain calcium homeostasis but also in the cellular homeostasis of other divalent ions. 460

461

462 **Experimental procedures**

Animals - Studies including the use of wild type (C57BL/6J), KO/TG (mice expressing 463 VDR only in the distal intestine) and Vdr knockout (KO) mice were approved by the Rutgers, 464

465 New Jersey Medical School Animal Care and Use Committee. Mice were maintained in a virus
466 and parasite-free barrier facility and exposed to a 12h-light, 12h-dark cycle. Food and water were
467 given *ad libitum*.

Transgenic mice expressing VDR only in the distal intestine were generated as previously 468 469 described (14) using a 9.5 kb fragment from the CDX2 promoter which directs transgene 470 expression specially to the distal intestine (75). The pBSKS-9.5kb CDX2-human(h)VDR poly A 471 transgene was injected into pronuclei of fertilized mouse oocyte at the Rutgers New Jersey 472 Medical School Genome Editing Core Facility. Mice heterozygous for transgene integration and ablation of the endogenous VDR (TG^{+/-}, KO^{+/-}) were bred to obtain mice with distal intestine 473 specific hVDR expression (TG/KO mice) (Fig. 1A). All TG/KO mice were fed with standard 474 475 rodent chow diet (Rodent Laboratory Chow 5001; Ralston Purina Co.) and were used at 12-14 476 weeks of age. Genotyping was performed by PCR using DNA extracted from tail biopsies and mouse and human specific VDR primers (14). VDR KO mice were obtained from the Jackson 477 Laboratory (originally from M. Demay, Harvard Medical School). 478

Findings in the TG/KO mouse intestine were compared to results obtained using vitamin D deficient mice. To examine the response to 1,25(OH)₂D₃ in both TG/KO mice and vitamin Ddeficient mice, each animal was administered three intraperitoneal injections of vehicle or 1,25(OH)₂D₃ (Cayman Chemical Company, Ann Arbor, MI) at a concentration of 1 ng/g body weight (bw) in a 9:1 mix of propylene glycol and ethanol at 48, 24 and 6 hours before termination. The three-dose protocol was used to study short-term and long-term effects of 1,25(OH)₂D₃ administration.

The vitamin D-deficient mice were generated from C57BL/6J female mice fed a vitamin D deficient diet (-D, 0.47%Ca, 0.3%P; TD89123; Teklad diet from Envigo) for 3-4 weeks before mating, during pregnancy and lactation. Their offspring were provided the same diet starting directly after weaning until the tissues were harvested at 12-14 weeks of age.

490 Slc30a10 KO (*Slc30a10^{-/-}*) mice were generated in the lab of Somshuvra Mukhopadhyay
491 as previously described (27). Tissues from *Slc30a10^{-/-}* mice were obtained at the University of
492 Texas (UT) at Austin under protocols approved by the Institutional Animal Care and Use
493 Committee at UT Austin. Animals were fed regular rodent chow containing ~84 ug Mn/g chow

and tissues harvested at 5-6 weeks as knockouts develop severe Mn toxicity beyond 8 weeks ofage.

496 In order to compare the regulation of Slc30a10 to the regulation of classic $1,25(OH)_2D_3$ target genes a time course of response to $1.25(OH)_2D_3$ as well as the response to high or low 497 498 calcium diets and developmental changes in the expression of Slc30a10 were examined. The 499 time course study was conducted using vitamin D-deficient mice injected with a single dose of 1,25(OH)₂D₃ (ip 10 ng/g bw) and killed at 1,4 or 24h after injection. The dietary study was done 500 501 using 4-week-old mice fed either a high calcium (1% Ca, TD92309) or low calcium (0.02% Ca 502 TD86162) for 4 weeks. For the developmental study time pregnant, neonatal and adult mice were fed a standard chow diet. Intestine was harvested from 18-day-old fetus and 1-, 3-, and 6-week-503 504 old mice (n = 5-6 per group for 1-, 3- and 6-week-old mice; for fetus n = 3 pooled intestinal samples of 3 individual intestines). For the mice studies both male and female mice were studied. 505 506 No sexual dimorphism was observed in the vitamin D target genes at these ages.

507 To determine whether classical as well as novel responses to $1,25(OH)_2D_3$ occur in both 508 villus and crypt, intestinal epithelial cells were isolated from both crypts and villi. The duodenum 509 from 3-month-old mice was washed in cold phosphate buffered saline (PBS), cut into 1 cm pieces and rotated in 3 mM EDTA in PBS at 4 °C. After vigorous shaking to release the 510 511 epithelium, crypts were depleted of contaminating villi by passing through a 70 µm cell strainer (BD Falcon, Tewksbury, MA). Villus epithelium was collected from the top of the 70 µm cell 512 513 strainer (76). To test whether VDR target genes are conserved in humans, human enteroid 514 cultures were used with either a crypt-like phenotype (ie high proliferation, undifferentiated cells) or a villus like phenotype (ie. low proliferation, differentiated cells). Human intestinal 515 516 tissue specimens were obtained from endoscopy biopsy performed at Baylor College of 517 Medicine (BCM) from adult patients of both sexes (3 females and 3 males; before endoscopy biopsies informed consent was obtained from the patient and the Institutional Review Board at 518 519 BCM approved this study). Crypts were isolated using the EDTA chelation method described above. For human enteroids crypts were resuspended in Matrigel and plated in a 24 well plate. 520 521 Enteroids were maintained in complete media with growth factors, CMGF+ with media changes 522 every 2-3 days (77). Enteroids were split every week by passing them through an insulin syringe 523 and replated in Matrigel. For treatment of enteroids with $1,25(OH)_2D_3$ or vehicle, enteroids were

529 RNA isolation and expression analysis – Total RNA was isolated from mouse tissues, isolated mouse villus and crypt epithelium and human organoids using RiboZol RNA extraction 530 reagent (Amresco, Solon, OH) or TRIzol reagent (Invitrogen, Carlsbad, CA) according to 531 manufacturer's instructions and subsequently purified with RNeasy Plus Universal Kit (Qiagen, 532 Hilden, Germany) using on column DNAse digestion (Qiagen, Hilden, Germany). RNA 533 534 concentration was measured with a NanoDrop spectrophotometer ND-1000 (Isogen Life 535 Science, Utrecht, The Netherlands). RNA integrity was assessed using a denaturing agarose gel stained with ethidium bromide (EtBr) or by using an Agilent (Agilent Technologies, Santa Clara, 536 CA) Bioanalyzer nanochip. For quantitative real time PCR (qRT-PCR), 2 µg of total RNA were 537 538 used to synthesize complementary DNA using Superscript III First Strand synthesis system 539 (Invitrogen, Carlsbad, CA) following the manufacturer's instructions. Relative quantification of target gene expression was performed using Taqman analyses. Taqman Gene Expression Probes 540 (Applied Biosystems, Foster City, CA) used for qRT-PCR are the following: Cyp24a1 541 542 (Mm00487244-m1), S100g (Mm00486654-m1), Slc30a1 (Mm00437377_m1), Slc30a10 (Mm00437377-m1), *Slc37a2* (Mm00451435_m1), *Trpv5* (Mm01166037_m1), *Trpv6* 543 544 (Mm00499069-m1) and Tjp1 (ZO-1 gene; Mm01320638_m1). Reactions were performed in a 7500 Real-Time PCR System (Applied Biosystems, Foster City, CA). The cycle steps were as 545 follow: an initial 3-min incubation at 95 °C, followed by 40 cycles of 95 °C for 10 s, 60 °C for 546 547 30 s, and 72 °C for 30 s. Expressions of gene of interest were normalized to Gapdh (Mm999999g1). The $2^{-\Delta\Delta Ct}$ method was used to calculate relative gene expression. 548

maintained in high Wnt3A (undifferentiated) medium (CMGF+) or differentiation medium (CMGF+ without Wnt3A, nicotinamide, SB202 and R-Spondin) 3 days before treatment, with

one media change between (77). Subsequently enteroids were treated with $100 \text{ nM} 1,25(\text{OH})_2\text{D}_3$ (Cayman Chemical Company) or vehicle (ethanol) at equal volumes in media. 24h following

treatment enteroids were collected and processed

549 For global assessment of mouse RNA levels, total RNA was subjected to two rounds of poly(A) selection using oligo-d(T)25 magnetic beads (New England Biolabs (NEB), Ipswich, 550 551 MA) Illumina compatible RNA-seq libraries were constructed using the NEBNext® Ultra[™] II 552 RNA Library Prep with Sample Purification Beads (Catalog no. E7775) and NEBNext® Multiplex Oligos for Illumina® (Dual Index Primers Set 1; Catalog No. E7600) according to the 553

554 manufacturer's instructions. Poly A selection and library quality were assessed using Tapestation 2200 (Agilent Technologies, Santa Clara, CA) and libraries were quantified using Qubit 4.0 555 556 fluorometer (Thermofisher, Waltham, MA). The prepared libraries were sequenced on an Illumina NextSeq 500 instrument. (Illumina, San Diego, CA). CLC Genomics Workbench 11.0.1 557 version (http://www.clcbio.com/products/clc-genomics-workbench/:Qiagen) was used for RNA-558 seq analysis. De-multiplexed fastq files from RNA-Seq libraries were imported into the CLC 559 software. Bases with low quality were trimmed and reads were aligned to the Mus musculus 560 561 reference genome build 9 (mm9) using Kallisto. The aligned reads were obtained using the 562 RNA-Seq Analysis tool CLC Genomics Workbench. Kallisto (version 45) was utilized to quantify the transcript abundances of the RNA-Seq samples through pseudoalignment, using 563 single-end reads and an Ensembl mm9 transcriptome build index. The tximport (version 1.14.0) 564 565 package (78) was run in R (version 3.6.2) to create gene-level count matrices for use with DESeq2 (version 1.26) (79) by importing quantification data obtained from Kallisto. DESeq2 566 567 was then used to generate transcript levels in each tissue sample. RNA-seq data (GSE 144978) 568 were deposited in the Gene Expression Omnibus of the National Center for Biotechnology.

For global assessment of RNA levels from human enteroids samples, RNA-seq data were aligned to the reference human genome (hg38) using Kallsito. Subsequently, differential gene expressions for both mouse and human were assessed using DESeq2 in R and transcript levels were obtained using a false discovery rate of 0.05 and a cut off of 1.5-fold change. RNA-seq data (GSE 159811) were deposited in the Gene Expression Omnibus of the National Center for Biotechnology.

575 For GO term analyses transcript IDS were used with the biomaRt package (version 2.42) 576 (80) along with RDavidWebService package (version 1.24) (81) as well as DAVID 6.8 (82) to 577 identify biological processes, cellular components and molecular functions in which input genes 578 are enriched.

Western blot analysis – For Western bolt analysis of VDR, total cellular protein was
extracted with a lysis buffer containing 50 mM of Tris-HCl (pH 7.5), 150 mM of NaCl, 0.1%
sodium dodecyl sulfate, 1.0% NP-40, and protease inhibitors. Protein content was measured
using Bradford assay (83) or Reducing Agent and Detergent Compatible Protein Assay (Bio-Rad
Laboratories, Inc., Hercules, CA). 50 µg of denatured protein was subjected to sodium dodecyl

sulfate-polyacrylamide gel electrophoresis (4-20% gradient gel; Bio-Rad Laboratories, Inc.,
Hercules, CA) and transferred onto a polyvinylidene difluoride membrane (Bio-Rad
Laboratories, Inc., Hercules, CA) for Western blot analysis using an enhanced chemiluminescent
detection system (Denville Scientific, Inc., Holliston, MA). β-actin immunoblotting was used for
sample normalization. Anti-VDR (D-6) [research Resource Identified (RRID):AB_628040] and
anti-b-actin (RRID:AB_626632) as well as secondary antibodies were obtained from Santa Cruz
Biotechnology, Inc. (Dallas, TX).

Bone analysis – μ CT analysis of the left tibiae was performed *ex vivo* using a high resolution SkyScan 1172 system (50 kV, 200 μ A, 0.5 mm aluminum filter) to examine trabecular and cortical bone parameters (84). Serial tomographs, reconstructed from raw data using the conebeam reconstruction software (NRecon, v.1.4.4.0; Skyscan), were used to compute trabecular and cortical parameters, respectively from the proximal metaphyseal and mid-diaphyseal area. Analysis was performed according to the guidelines of the American Society for Bone and Mineral Research (85).

Serum analysis – Serum calcium, assayed using a colorimetric assay (Pointe Scientific, Inc.
 Canton Mi), was determined by Heartland Laboratories, Ames, Iowa.

600 *Statistical analysis* – Results are displayed as mean \pm standard error (SEM). Data were analyzed 601 using student t-test or ANOVA and additionally with Benjamini and Hochberg in a post-hoc test 602 to consider significant difference between groups (p < 0.05).

603

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607

608 Author contributions

SL, SC, SM, NFS, MPV and JCF designed the study. SL, JDLC, SH (UT), ZKC, RA, OPC, JH,
PD, LV performed experiments. SL, SC, SM, NFS, ZKC, MPV, JCF, GC, LV, PD, PS, SH
provided interpretation of results. SL and SC wrote the paper with comments from all authors.

612

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617 **Conflict of interest**

618 The authors disclose no conflicts

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857 Footnotes

The abbreviations used are: 1,25(OH)₂D₃, 1,25-dihydroxyvitamin D3; 25(OH)D₃, 25-858 859 hydroxyvitamin D_3 ; ANOVA, Analysis of variance; bw, body weight; CYP24A1, 25hydroxyvitamin 24-Hydroxylase; DNA, deoxyribonucleic acid; 860 D_3 EDTA, ethylenediaminetetraacetate; g, gram; HCL, hydrogen chloride; hVDR, human VDR; kb, 861 862 kilobases; kDa, kilodaltons; KO, knockout; KO/TG, knockout/transgenic; µCT, micro-computed 863 tomography; min, minute(s); Mn, manganese; mRNA, messenger RNA; NaPi-IIb, Sodiumdependent phosphate transport protein 2B; ng, nanogram; PBS, phosphate-buffered saline; RNA-864 Seq, RNA sequencing; S100g, RT-qPCR, quantitative reverse transcription PCR; S100 calcium 865 binding protein G; SEM, standard error of the mean; Slc30a1, zinc transporter 1; SLC30A10, 866 Solute Carrier Family 30 Member 10; SLC30A4, Solute Carrier Family 30 Member 4; 867 868 SLC30A5, solute carrier family 30 member 5; SLC34A2, Sodium-dependent phosphate transport 869 protein 2B; SLC37A2, Glucose-6-phosphate exchanger; Tris, tris(hydroxymethyl)aminomethane; 870 TRPV6, Transient receptor potential cation channel subfamily V member 6; VDR, vitamin D receptor; VDRE, vitamin D response element; Zn, zinc 871

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905Figure 1. Gene expression measured by RNA-seq in the colon of KO/TG1 mice treated with906 $1,25(OH)_2D_3$ (A) KO/TG mice express VDR only in the distal intestine. Expression of hVDR is907restricted to the distal intestine of KO/TG mice. Mouse (m) VDR is absent in KO/TG mice. (B)908Rescue of VDR dependent rickets in the KO/TG mice (Von Kossa staining; see ref 14). (C) Gene909expression in the colon of KO/TG mice treated with $1,25(OH)_2D_3$ (TG1 +D, black bar) compared910to VDR KO (open bar). Gene expression, measured by RNA-seq analysis, is shown as the mean911of RPKM ± SEM. n = 4 per group * p < 0.05, TG1 + D compared to VDR KO.

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913 Figure 2. Response to $1,25(OH)_2D_3$ treatment in KO/TG mouse intestine as measured by 914 RT-qPCR. Gene expression in the duodenum (Duo) and colon (Col) of KO/TG1 mice (TG1) 915 (A) or KO/TG2 mice [mice from a second transgenic line (TG2)] (B) treated with vehicle (black bar) or 1,25(OH)₂D₃ (1 ng/g bw at 48, 24 and 6 h before termination; TG1+ D or TG2 + D, 916 917 stripped bar). Results are compared to VDR KO mice (open bar). n = 6 per experimental group. 918 Gene expression was normalized to Gapdh and expressed as mean \pm SEM (+ p < 0.05 compared 919 to vehicle (TG1 or TG2), # $p < 0.05 \ 1,25(OH)_2D_3$ treated (TG1 or TG2 +D) compared to VDR 920 KO; * p < 0.05 vehicle (TG1 or TG2) compared to VDR KO. The duodenum was included as a 921 control. Note the lack of responsiveness of the duodenum to $1,25(OH)_2D_3$ treatment due to the 922 absence of VDR in the proximal intestine of these transgenic mice. Col, colon; Duo, duodenum; 923 ND, not detected.

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Figure 3. Gene expression in intestinal segments in response to repeated administration of 1,25(OH)₂D₃ to vitamin D deficient mice. 12- to 14-week-old vitamin D-deficient mice were injected with 1,25(OH)₂D₃ (+D, black bar) or vehicle (open bar) three times over 48 h (at 48, 24 and 6 h prior to termination (ip. 1ng/g bw per injection). Data was normalized to *Gapdh* and expressed as mean \pm SEM. n = 6 per experimental group. * Significantly different from vehicle treated group at *p* < 0.05. Duo, duodenum; Ile, ileum; Col, colon; ND, not detected.

932 Figure 4. Regulation of intestinal $1,25(OH)_2D_3$ target gene expression: time course of 933 induction by 1,25(OH)₂D₃, effects of dietary calcium and developmental changes. (A) RT-934 qPCR analysis of *Slc30a10* and *Trpv6* expression in the duodenum at 0, 4 and 24 h after a single 935 injection of $1,25(OH)_2D_3$ (10 ng/g bw) to vitamin D deficient mice. n = 5 per time point. * $p < 10^{-1}$ 936 0.05 compared to 0 time point. (B) RT-qPCR analysis of Slc30a10 and Trpv6 expression in duodenum of mice fed a low calcium (0.02%) or high calcium (1%) diet for 4 weeks. n = 7 per 937 938 group. * Significantly different from the mice fed the high calcium diet at p < 0.05. (C) 939 Developmental changes in gene expression in mice raised under standard conditions and killed at

Molecular and Cellular Biology 94018 days gestation and at 1, 3, and 6 weeks of age. Intestine was harvested (for fetus n = 5, pooled941intestinal samples of 3 - 4 individual intestines; n = 5 - 6 for 1-, 3-, and 6-week-old mice). Since942the small intestine and not individual segments were harvested and analyzed in the fetus, for943comparison small intestine was also analyzed at different post-natal stages. Data are expressed as944mean \pm SEM. Data were normalized using *Gapdh* or *18S rRNA*. # Significantly different from945fetus at p < 0.05. * Significantly different from 1 week old at p < 0.05. Duo, duodenum.

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Figure 5. 1,25(OH)₂D₃ target gene expression in the intestine and kidney of *Slc30a10^{-/-}* and control mice. 1,25(OH)₂D₃ target genes in the intestine (A) and kidney (B) in *Slc30a10^{-/-}* and control mice. Data are expressed as mean \pm SEM and were normalized using *Gapdh*. n = 4 - 6 in each group. * *p* < 0.05 significantly different between two experimental groups. Duo, duodenum; Col, colon.

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Figure 6. VDR and 1,25(OH)₂D₃ target genes in mouse epithelial cells in villi and crypts. 953 954 (A) Intestinal epithelial cells were isolated from crypts (left panel) and villi (right panel) of 3month-old mouse duodenum. (B) Vdr gene expression (left panel) and VDR protein (Western 955 956 blot analysis; right panel) in mouse duodenal crypts and villi from 3-month-old WT mice compared to VDR KO mice. For Western blot analysis the relative optical density (OD) obtained 957 using the VDR antibody was divided by the OD obtained after β -actin staining to produce VDR-958 relative OD. n=3 - 4 (C and D) RT-qPCR analysis of $1,25(OH)_2D_3$ target genes in mouse 959 960 duodenal villi and crypts. (C) Cyp24a1 expression was examined in 3-month-old mice injected 961 ip. with vehicle or 1,25(OH)₂D₃ (1 ng/g bw; injected at 48, 24, and 6 h prior to euthanasia). * 962 significantly different from WT + vehicle at p < 0.05. (D) Trpv6 and Slc30a10 gene expression 963 in intestinal epithelial cells from crypts and villi of 3-month-old WT mice. RT-qPCR reactions 964 were normalized to *Gapdh* or *18S rRNA*. n = 3 - 4. ND, not detected.

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966 Figure 7. Gene expression measured by RNA-seq in crypt-like and villus-like human 967 duodenal organoids. (A) Bright field images of crypt-like (undifferentiated) and villus-like 968 (differentiated) duodenal human enteroids. (B) Venn diagram indicating genes significantly 969 upregulated at least 1.5 folds and with FDR < 0.05 upon $1,25(OH)_2D_3$ treatment (100 nM for 24 970 h) in crypt-like (undifferentiated) and villus-like (differentiated) human enteroids and overlapping genes. (C) Expression counts of SLC30A10, VDR and classic 1,25(OH)₂D₃ target 971 972 genes in control and $1,25(OH)_2D_3$ treated villus and crypt-like human enteroids. Data is from enteroids from 6 patients (3 females and 3 males). 973

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Figure 8. Top GO terms for genes enriched in human duodenal enteroids treated with
1,25(OH)₂D₃ (A) Crypt only (B) Villus only (C) Crypt and Villus shared.

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Trpv6

VDR ' TG1 KO +D Colon

SIc39a14

4×10

3×10²

2×10²

1×10²

2×101

1×10

5×10

0

0





1.5×10²

S100g

VDR KO +D Colon

Slc30a1

VDR KO

VDR KO

TG1 +D Colon

TG1 +D

TG1 +D

Colon

Cacna1d

2.5×10⁴

2×10

1.5×104

1×104

5×103

2×101

2×101

0

Average RPKM

SIc30a10



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Α

10.0



Cyp24a1



Trpv6

12.5

10.0

S100g

VDR KO TG1 TG1 +D

Col

VDR TG1 TG1 KO +D

Col

TG2 +D

TG2

Col

10.0-

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Relative expression



С





D







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Α Intestine



Kidney

В







12.5-Relative expression 10.0 levels 7.5 5.0 2.5

0.0

Control SIc30a10-/-

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Α







Villi





Villi

MCB

С

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D





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В

С



	Male		Female		
	Control	Slc30a10 KO	Control	Slc30a10 KO	
Trabecular bone					
Ν	8	5	10	9	
BV/TV (%)	6.55 ± 1.60	4.44 ± 1.29	5.41 ± 2.33	4.72 ± 1.20	
Trabecular number (1/mm)	2.07 ± 0.39	$1.48 \pm 0.36^*$	1.62 ±0.57	1.54 ± 0.35	
Trabecular thickness (mm)	0.031 ± 0.002	0.030 ± 0.002	0.033 ± 0.002	0.031 ± 0.001*	
Cortical bone					
Ν	8	4	10	8	
Cross-sectional					
tissue area	2.11 ± 0.30	$1.35 \pm 0.39^*$	1.92 ± 0.27	$1.33 \pm 0.16*$	
(\mathbf{mm}^2)					
Medullary area (mm ²)	1.44 ± 0.23	$1.00 \pm 0.34^*$	1.29 ± 0.23	$1.03 \pm 0.13^*$	
Cortical thickness (mm)	0.124 ± 0.012	$0.092 \pm 0.023*$	0.140 ± 0.011	0.086 ± 0.010*	
Porosity (%)	12.07 ± 2.68	$6.12 \pm 1.12^*$	9.22 ± 2.07	$6.24 \pm 1.55^*$	

NOTE. Trabecular bone parameters and cortical bone parameters as measured by μ CT. Data are presented as mean ± SEM (unpaired, two-tailed t test KO vs WT for each sex). BV/TV, bone volume/tissue volume. *p < 0.05.

(pt-like	Vill	lus-like	
Gene	Fold Change	p Value	Fold Change	<i>p</i> Value	_
SLC30A10	5.38	2.05E-19*	3.20	2.99E-03*	Mn efflux transporter
SLC30A4	0.82	5.51E-01	0.93	8.44E-01	Zn transporter
SLC30A5	1.02	9.56E-01	1.05	7.89E-01	Zn transporter
SLC34A2	6.94	5.17E-22*	3.02	3.00E-03*	Phosphate transporter
SLC37A2	1.06	9.63E-01	13.13	8.34E-49*	G6P transporter
TRPM7	0.94	9.22E-01	0.78	1.23E-01	non Slc transporter of divalent cations

Table 2 Gene expression of transporters in crypt-like and villus-like human enteroids derived from duodenum treated with $1,25(OH)_2D_3$ or vehicle

**p* < 0.05.