

## Short Communication

### C-JUN N-TERMINAL KINASE MODULATES 1,25-DIHYDROXYVITAMIN D<sub>3</sub>-INDUCED CYTOCHROME P450 3A4 GENE EXPRESSION

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#### ABSTRACT:

1,25-Dihydroxyvitamin D<sub>3</sub> (1,25(OH)<sub>2</sub>D<sub>3</sub>) is known to induce the expression of cytochrome P450 3A4 (CYP3A4) in human colon carcinoma Caco-2 cells. Recently, it was demonstrated that the vitamin D receptor (VDR) regulates 1,25(OH)<sub>2</sub>D<sub>3</sub>-induced CYP3A4 gene expression through the xenobiotic-responsive element and the vitamin D-responsive element located on the 5'-flanking region of the CYP3A4 gene. On the other hand, we previously reported that protein kinases such as protein kinase C and tyrosine kinases contribute to the induction of CYP3A4 mRNA by 1,25(OH)<sub>2</sub>D<sub>3</sub>. In the present study, we examined the involvement of mitogen-activated protein kinases (MAPKs) in the 1,25(OH)<sub>2</sub>D<sub>3</sub>-induced CYP3A4 gene expression using MAPK inhibitors. Curcumin, a c-Jun N-terminal kinase (JNK) pathway

inhibitor, and anthra[1,9-cd]pyrazole-6(2H)-one (SP600125), a JNK inhibitor, suppressed the induction of CYP3A4 mRNA by 1,25(OH)<sub>2</sub>D<sub>3</sub>, but not 2'-amino-3'-methoxyflavone (PD098059), a mitogen-activated protein kinase kinase-extracellular signal-regulated kinase (ERK) pathway inhibitor, or 4-(4-fluorophenyl)-2-(4-methylsulfinylphenyl)-5-(4-pyridyl)1H-imidazole (SB203580), a p38 inhibitor. In addition, we demonstrated that SP600125 dose-dependently inhibited the CYP3A4 promoter activity induced by 1,25(OH)<sub>2</sub>D<sub>3</sub> using the reporter plasmid of the CYP3A4 promoter. However, SP600125 did not affect 1,25(OH)<sub>2</sub>D<sub>3</sub>-induced transactivation of the DR3 via VDR. These results indicate that JNK, but not ERK or p38, is required for the optimal activation of the CYP3A4 gene induced by 1,25(OH)<sub>2</sub>D<sub>3</sub>.

Cytochrome P450 (P450) plays an important role in the oxidative metabolism of numerous endogenous and exogenous compounds. In humans, cytochrome P450 3A4 (CYP3A4) is the predominant P450 isoform in the liver and small intestinal epithelial cells (Watkins et al., 1987; Paine et al., 1997) and is responsible for the metabolism of more than 60% of therapeutic drugs. Intestinal CYP3A4 is thought to contribute to the first-pass metabolism of orally administered drugs (Paine et al., 1997). The CYP3A4 gene is inducible by many xenobiotics, including rifampicin, dexamethasone, and phenobarbital (Pichard et al., 1990). The nuclear receptor, pregnane X receptor, is known to contribute to the CYP3A4 gene induction by these drugs (Kliewer et al., 1998; Lehmann et al., 1998; Goodwin et al., 1999). It has recently been demonstrated that 1,25-dihydroxyvitamin D<sub>3</sub> (1,25(OH)<sub>2</sub>D<sub>3</sub>) is a potent inducer of the CYP3A4 gene in the human colon carcinoma cell line Caco-2, which has been extensively used as an experimental model of small intestinal cells (Schmiedlin-Ren et al., 1997; Hara et al., 2000).

1,25(OH)<sub>2</sub>D<sub>3</sub>, the most active metabolite of vitamin D, functions to regulate cellular proliferation and differentiation and calcium homeostasis in the intestine, bone, and kidney (Christakos et al., 1996). Most of these physiological activations are mediated by the vitamin D

receptor (VDR), which belongs to the nuclear hormone receptor superfamily. The VDR acts as a ligand-inducible transcription factor through heterodimerization with the retinoid X receptor and binding to the vitamin D response element (VDRE) within the vitamin D-inducible genes (Christakos et al., 1996).

Recently, a direct repeat separated by three nucleotides (DR3) and an everted repeat separated by six nucleotides (ER6) within the 5'-flanking region of the CYP3A4 gene were identified as VDRE (Thummel et al., 2001; Drocourt et al., 2002). We and others demonstrated that the VDR is an essential factor for 1,25(OH)<sub>2</sub>D<sub>3</sub>-induced CYP3A4 expression (Thummel et al., 2001; Drocourt et al., 2002; Hara et al., 2002). On the other hand, several reports have demonstrated that the phosphorylation step is critical for the expression of xenobiotic-induced P450 genes such as CYP1A1 and CYP3A (Chen and Tukey, 1996; Galisteo et al., 1999). We also showed that alteration of the cellular phosphorylation state mediated by protein kinase C (PKC) and protein tyrosine kinases affects the 1,25(OH)<sub>2</sub>D<sub>3</sub>-mediated induction of CYP3A4 mRNA in Caco-2 cells (Hara et al., 2002). In addition, the inhibition of the phosphorylation reduced the VDR-mediated enhancement of osteocalcin gene transcription (Desai et al., 1995). These results suggest that the phosphorylation step is critical for the complete CYP3A4 induction by 1,25(OH)<sub>2</sub>D<sub>3</sub> via VDR. In the present study, we examined whether MAPKs are involved in the 1,25(OH)<sub>2</sub>D<sub>3</sub>-induced expression of the CYP3A4 gene via VDR.

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**ABBREVIATIONS:** P450, cytochrome P450; 1,25(OH)<sub>2</sub>D<sub>3</sub>, 1,25-dihydroxyvitamin D<sub>3</sub>; VDR, vitamin D receptor; VDRE, vitamin D response element; DR3, direct repeat separated by three nucleotides; ER6, everted repeat separated by six nucleotides; PKC, protein kinase C; MAPK, mitogen-activated protein kinase; PD098059, 2'-amino-3'-methoxyflavone; SB203580, 4-(4-fluorophenyl)-2-(4-methylsulfinylphenyl)-5-(4-pyridyl)1H-imidazole; SP600125, anthra[1,9-cd]pyrazole-6(2H)-one; ERK, extracellular signal-regulated kinase; JNK, c-Jun N-terminal kinase; RT-PCR, reverse transcription-polymerase chain reaction; TK, thymidine kinase.



CYP3A4 mRNA expression in Caco-2 cells increased 2.2-fold in the presence of 100 nM  $1,25(\text{OH})_2\text{D}_3$  as shown in Fig. 1A. The CYP3A4 mRNA induction was inhibited by treatment with curcumin, but not by PD098059 or SB203580. Curcumin, a JNK pathway inhibitor, is known to partly inhibit ERK at the concentration used (Chen and Tan, 1998). However, the possibility of ERK also being involved is eliminated from the results that PD098059, a specific mitogen-activated protein kinase kinase-ERK pathway inhibitor, had no effect on  $1,25(\text{OH})_2\text{D}_3$ -induced CYP3A4 mRNA expression. To confirm the involvement of JNK in the CYP3A4 mRNA induction, we used SP600125, a specific JNK inhibitor. As shown in Fig. 1B, SP600125 significantly inhibited the induction in a dose-dependent manner.

As shown in our previous study (Hara et al., 2002), VDR plays an important role in  $1,25(\text{OH})_2\text{D}_3$ -induced CYP3A4 mRNA expression. Therefore, we confirmed whether the MAP kinase inhibitors used affect the constitutive expression of VDR mRNA. Treatment with PD098059, curcumin, or SB203580 had no effect on the constitutive expression of VDR mRNA, whereas SP600125 reduced VDR mRNA expression at high concentration (Fig. 1). These results indicate that JNK inhibition by SP600125 not only suppresses the induction of CYP3A4 mRNA by  $1,25(\text{OH})_2\text{D}_3$ , but it also reduces the basal expression of VDR. Thus, it is possible that the reduction of VDR expression by SP600125 might be involved in the suppression of CYP3A4 mRNA induction in part.

To verify the mechanism involved in the transcriptional activation of the CYP3A4 gene via VDR, we constructed the CYP3A4 reporter plasmid, 3A4(-7836/-315)-pGL3. The 3A4(-7836/-315)-pGL3 was transfected into Caco-2 cells with the VDR expression vector or the empty expression vector. After the cells had been treated with 100 nM  $1,25(\text{OH})_2\text{D}_3$  for 24 h, the CYP3A4 promoter activity was measured. As shown in Fig. 2A, the treatment with  $1,25(\text{OH})_2\text{D}_3$  markedly increased the CYP3A4 promoter activity in cells transfected with the VDR expression vector ( $18.7 \pm 2.4$ -fold), whereas the same treatment hardly increased the promoter activity in cells transfected with the empty vector. These results indicate that  $1,25(\text{OH})_2\text{D}_3$  via VDR activates the chimera CYP3A4 promoter.

To determine whether MAPKs regulate  $1,25(\text{OH})_2\text{D}_3$ -induced CYP3A4 gene expression at the transcriptional level, we examined the effect of MAP kinase inhibitors on the activation of the CYP3A4 promoter using 3A4(-7836/-315)-pGL3. As shown in Fig. 2, B and C, treatment with PD098059 or SB203580 did not suppress  $1,25(\text{OH})_2\text{D}_3$ -induced activation of 3A4(-7836/-315)-pGL3, whereas with SP600125, it dose-dependently inhibited the activation. These results were consistent with the results shown in Fig. 1, and indicate that JNK contributes to the induction of the CYP3A4 gene by  $1,25(\text{OH})_2\text{D}_3$ . The treatment with SP600125 reduced the basal levels of CYP3A4 promoter activity by approximately 40% (Fig. 2, B and C), whereas it reduced the induced CYP3A4 promoter activity by over 80%. These results suggested that JNK is involved especially in CYP3A4 promoter activation induced by  $1,25(\text{OH})_2\text{D}_3$ .

Recent studies reveal that the DR3 and the ER6 in the 5'-flanking region of the CYP3A4 gene are functional VDREs (Thummel et al., 2001; Drocourt et al., 2002), and VDR plays a key role in  $1,25(\text{OH})_2\text{D}_3$ -induced CYP3A4 gene expression (Thummel et al., 2001; Drocourt et al., 2002; Hara et al., 2002). On the other hand, it was demonstrated that the phosphorylation of VDR itself by PKC and casein kinase II specifically modulates its transcriptional ability (Hsieh et al., 1991; Jurutka et al., 1996). Therefore, it is possible that the phosphorylation of VDR itself by JNK would modulate the transcriptional ability of VDR. To verify this possibility, we investigated whether SP600125 suppresses the activation of the promoter driven by VDR through the transactivation of the DR3 using (DR3)<sub>3</sub>-

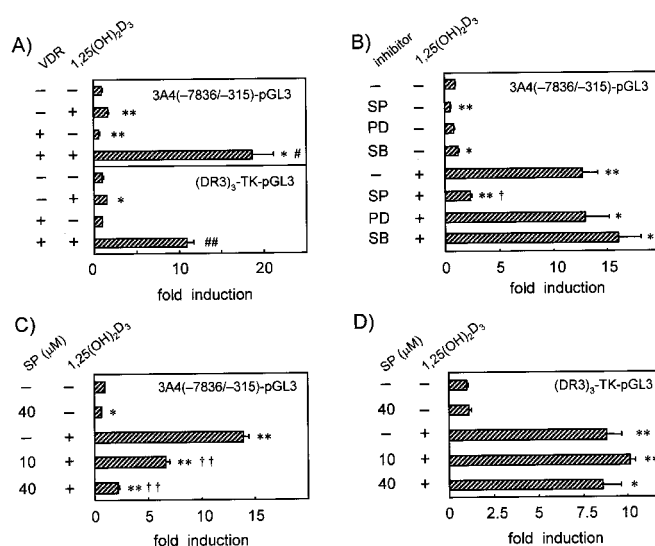


FIG. 2. Reporter analysis of the CYP3A4 promoter and the DR3 element. A,  $1,25(\text{OH})_2\text{D}_3$  activates the CYP3A4 promoter and the DR3 element. Caco-2 cells were cotransfected with the VDR expression vector (1.5  $\mu\text{g}$ ) or the empty expression vector (1.5  $\mu\text{g}$ ), the 3A4(-7836/-315)-pGL3 (1  $\mu\text{g}$ ) or the (DR3)<sub>3</sub>-TK-pGL3 (1  $\mu\text{g}$ ), and the pRL-TK vector (0.1  $\mu\text{g}$ ), and then treated with  $1,25(\text{OH})_2\text{D}_3$  (100 nM) for 24 h. B, effects of MAPK inhibitors on the transcriptional activation of the CYP3A4 promoter by  $1,25(\text{OH})_2\text{D}_3$ . Caco-2 cells were cotransfected with the VDR expression vector (1.5  $\mu\text{g}$ ), the 3A4(-7836/-315)-pGL3 (1  $\mu\text{g}$ ), and the pRL-CMV vector (0.02  $\mu\text{g}$ ). Cells were pretreated with SP600125 (SP, 40  $\mu\text{M}$ ), PD098059 (PD, 25  $\mu\text{M}$ ), and SB203580 (SB, 10  $\mu\text{M}$ ) for 30 min, followed by incubation with or without  $1,25(\text{OH})_2\text{D}_3$  (100 nM) for 24 h. C, SP600125 dose-dependently decreases transcriptional activation of the CYP3A4 promoter by  $1,25(\text{OH})_2\text{D}_3$ . Caco-2 cells were cotransfected with the VDR expression vector (1.5  $\mu\text{g}$ ), the 3A4(-7836/-315)-pGL3 (1  $\mu\text{g}$ ), and the pRL-CMV vector (0.02  $\mu\text{g}$ ). Cells were pretreated with SP600125 (SP, 10 or 40  $\mu\text{M}$ ) for 30 min, followed by incubation with  $1,25(\text{OH})_2\text{D}_3$  (100 nM) for 24 h. D, effect of SP600125 on the transcriptional activation of the DR3 element by  $1,25(\text{OH})_2\text{D}_3$ . Caco-2 cells were cotransfected with the VDR expression vector (0.5  $\mu\text{g}$ ), the (DR3)<sub>3</sub>-TK-pGL3 (1  $\mu\text{g}$ ), and the pRL-CMV vector (0.02  $\mu\text{g}$ ). The cells were then pretreated with SP600125 (SP, 10 or 40  $\mu\text{M}$ ) for 30 min, followed by incubation with  $1,25(\text{OH})_2\text{D}_3$  (100 nM) for 24 h. Values (mean  $\pm$  S.E.,  $n = 3$ ) are expressed as the fold induction relative to untreated cells. \* and \*\*, significant differences ( $P < 0.05$  and  $P < 0.01$ , respectively) compared with untreated cells. # and ##, significant differences ( $P < 0.05$  and  $P < 0.01$ , respectively) compared with vehicle-treated cells transfected the VDR expression vector. † and ††, significant differences ( $P < 0.05$  and  $P < 0.01$ , respectively) compared with  $1,25(\text{OH})_2\text{D}_3$ -treated cells.

TK-pGL3. As shown in Fig. 2A, (DR3)<sub>3</sub>-TK-pGL3 was markedly activated by  $1,25(\text{OH})_2\text{D}_3$  in cells cotransfected with the VDR expression vector, but not in empty expression vector-cotransfected cells, indicating that the transactivation of this construct is VDR-dependent. As shown in Fig. 2D, the activity of (DR3)<sub>3</sub>-TK-pGL3 increased 8.8-fold in the presence of 100 nM of  $1,25(\text{OH})_2\text{D}_3$ , and treatment with SP600125 did not inhibit the activity of (DR3)<sub>3</sub>-TK-pGL3 induced by  $1,25(\text{OH})_2\text{D}_3$ . Taken together, SP600125 decreased the induction of 3A4(-7836/-315)-pGL3 activity by  $1,25(\text{OH})_2\text{D}_3$ , whereas it did not affect the transactivation of the DR3 element by  $1,25(\text{OH})_2\text{D}_3$  via VDR. These results suggest that JNK does not modulate the transcriptional activity of VDR.

The detailed mechanism by which JNK regulates  $1,25(\text{OH})_2\text{D}_3$ -induced CYP3A4 expression remains unknown from this study. However, previous reports demonstrated that the xenobiotic-induced CYP2B1 expression mediated by the constitutive androstane receptor requires the transcription factor Sp1 for optimal expression (Muangmoonchai et al., 2001). In addition, it has been reported that the synergism between VDR and other transcription factors, including AP-1 and Sp1, activates the induction of VDR-mediated transactivation (Liu and Freedman, 1994). These observations indicate that the synergism between nuclear receptors and other transcription factors

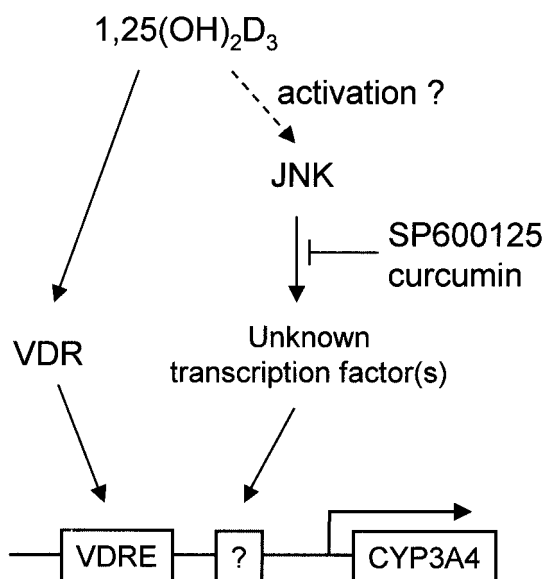


FIG. 3. Schema of regulatory mechanism of *CYP3A4* promoter activation caused by  $1,25(\text{OH})_2\text{D}_3$ .  $1,25(\text{OH})_2\text{D}_3$ -induced *CYP3A4* promoter activation might be regulated by the synergism between VDR and transcription factor(s) that is modulated by JNK.

coordinates nuclear receptor-mediated responses to xenobiotics. On the other hand,  $1,25(\text{OH})_2\text{D}_3$  is known to rapidly activate some protein kinases, including PKC, ERK, and JNK, through a non-genomic signaling pathway in various cell types, including Caco-2 cells (Chen et al., 1999). Therefore, it is likely that JNK activation caused by  $1,25(\text{OH})_2\text{D}_3$  might be involved in the induction of the *CYP3A4* gene. Taken together, the mechanism appears to be that  $1,25(\text{OH})_2\text{D}_3$ -induced *CYP3A4* gene activation is synergistically regulated by VDR and transcription factor(s) such as AP-1 and Sp1 that are modulated by JNK (Fig. 3).

In summary, our data indicate that JNK, but not ERK or p38, is an important regulator involved in  $1,25(\text{OH})_2\text{D}_3$ -induced *CYP3A4* gene activation. MAPKs are activated by various extracellular stimuli and regulate many gene expressions through phosphorylation of transcription factors. Therefore, elucidation of the involvement of MAPKs in the xenobiotic-induced expression of *P450* genes may be instrumental in understanding the induction of *P450*s under disease conditions such as inflammatory disorders and cancer.

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