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## **Sulfotransferase genes: Regulation by nuclear receptors in response to xeno/endo-biotics**

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### **Abstract**

Pregnane X receptor (PXR) and constitutive active/androstane receptor (CAR), members of the nuclear receptor superfamily, are two major xeno-sensing transcription factors. They can be activated by a broad range of lipophilic xenobiotics including therapeutics drugs. In addition to xenobiotics, endogenous compounds such as steroid hormones and bile acids can also activate PXR and/or CAR. These nuclear receptors regulate genes that encode enzymes and transporters that metabolize and excrete both xenobiotics and endobiotics. Sulfotransferases (SULTs) are a group of these enzymes and sulfate xenobiotics for detoxification. In general, inactivation by sulfation constitutes the mechanism to maintain homeostasis of endobiotics. Thus, deciphering the molecular mechanism by which PXR and CAR regulate *SULT* genes is critical for understanding the roles of SULTs in the alterations of physiological and pathophysiological processes caused by drug treatment or environmental exposures.

### **Keywords**

Constitutive active/androstane receptor; gene regulation; pregnane X receptor; sulfotransferase; xeno-sensing nuclear; receptor

### **Introduction**

Constant exposure to numerous xenobiotics (e.g. therapeutics, agricultural and industrial chemicals) often causes significant impacts on human health. To counter toxicity, xenobiotic metabolizing enzymes such as cytochrome P450s (CYPs) and sulfotransferases (SULTs) are

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#### **Declaration of interest**

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coordinately induced to eliminate them from the body (Omiecinski et al., 2011). Pregnane X receptor (PXR, NR1I2) and constitutive active/androstane receptor (CAR, NR1I3) are two major transcription factors that regulate these genes in response to xenobiotics (Willson & Kliewer, 2002). In addition, farnesoid X receptor (FXR, NR1H4), liver X receptor (LXR, NR1H2/3), peroxisome proliferator-activated receptors (PPARs, NR1C1/2/3) and vitamin D receptor (VDR, NR1I1) are also involved in the regulation of genes that encode for xenobiotic metabolizing enzymes (Bouillon et al., 2008; Jakobsson et al., 2012; Modica et al., 2010; Pyper et al., 2010). These nuclear receptors can be co-regulated by xenobiotics and endobiotics and are involved in various physiological and pathophysiological processes such as cell differentiation and development, drug and energy metabolism, immune response and tumorigenesis as well as endocrine homeostasis. Sulfate conjugation is an important reaction in both xenobiotic and endobiotic metabolism and is catalyzed by cytosolic SULTs (Gamage et al., 2006). In most cases, sulfation inactivates both xenobiotics and endobiotics, increasing their water solubility to accelerate excretion (Alnouti, 2009; Gamage et al., 2006). Expression of *SULT* genes is primarily regulated by the above-mentioned nuclear receptors, in particular by PXR and CAR.

### Xeno-sensing nuclear receptors

Liver, kidney and intestines are three major organs that metabolize and excrete xenobiotics and endobiotics. Upon xenobiotic exposures, these organs activate nuclear receptors such as PXR and CAR that regulate SULT expressions.

#### PXR

PXR (NR1I2) was first cloned from a mouse liver cDNA library (Kliewer et al., 1998). Subsequently PXR orthologs were cloned from various species including human, rat, rabbit and monkey (Blumberg et al., 1998; Moore et al., 2002; Savas et al., 2000; Zhang et al., 1999). The name PXR was given based on its ability to be activated by various natural and synthetic pregnanes. The *CYP3A* genes were identified as the first PXR target since PXR activators and *CYP3A* inducers overlapped. It was then determined that ligand specificity was markedly different among various species (Kliewer et al., 2002). PXR adapts a large and flexible structure for ligand binding to accommodate a broad range of hydrophobic man-made and naturally occurring xenobiotics at effective concentrations, which are usually the low  $\mu\text{M}$  range. Endobiotics such as steroids and bile acids can also activate PXR. However, it remains elusive as to whether or not these endobiotics actually play any role in regulating PXR *in vivo* in organs. Un-liganded PXR is predominantly localized in the cytoplasm (Squires et al., 2004). A co-chaperon dubbed cytoplasmic CAR retention protein (CCRP) forms a complex with CAR to enable sequestration of PXR in the cytoplasm (Squires et al., 2004). Liganded PXR translocates into the nucleus, heterodimerizes with retinoid acid X receptor (RXR, NR2B), and binds to response sequences (i.e. DR3, DR4, ER6 and ER8 motifs), thereby activating transcription of targeted genes. Co-activators such as steroid receptor coactivator 1 $\alpha$  and PPAR- $\gamma$  coactivator 1 $\alpha$  and co-repressors such as nuclear receptor corepressor 1 and 2 co-regulate PXR-mediated transcription (Bhalla et al., 2004; Sugatani et al., 2005; Synold et al., 2001). A recent report has demonstrated that cell signaling can also activate PXR in the absence of xenobiotic activators (Sivertsson et al.,

2013). Cyclin-dependent kinase 2 (CDK2) phosphorylates PXR and sequesters it in the cytoplasm of human hepatocellular carcinoma Huh7 cells. Continuous culture of confluent Huh7 cells decreases CDK2 levels, resulting in the dephosphorylation of PXR. Non-phosphorylated PXR translocates into the nucleus and activates the *CYP3A4* gene. Although there are potential phosphorylation sites by protein kinase C, protein kinase A, or p70 S6 kinase in PXR, phosphorylation of these sites in endogenous PXR has not been confirmed (Ding & Staudinger, 2005; Lichti-Kaiser et al., 2009a, b; Pondugula et al., 2009). In addition to phosphorylation, PXR may undergo ubiquitylation, acetylation or SUMOylation (Biswas et al., 2011; Hu et al., 2010; Staudinger et al., 2011). Current understanding of PXR activation is summarized in Figure 1.

PXR is also known to be expressed in lung, ovary, normal and cancerous breast tissues and peripheral blood mononuclear cells (PMBCs) (Masuyama et al., 2001; Schote et al., 2007; Siest et al., 2008). PXR appears to induce CYPs in the ovary and PMBCs. PXR may regulate growth and death of breast cancer cells, in addition to increasing drug resistance by inducing metabolism. Activation of PXR may affect various metabolic diseases such as acetaminophen and bilirubin toxicities, steatosis, cholestasis, diabetes and osteomalacia (Kakizaki et al., 2008; Konno et al., 2008; Staudinger et al., 2001a; Zhou et al., 2009). In the other cases, activation can be beneficial acting as an anti-inflammatory factor to bowel disease and biliary primary cirrhosis (Shah et al., 2007; Wallace et al., 2010).

## CAR

CAR (NR1H3), nominally called constitutive activate receptor, constitutive androstane receptor or CAR, was cloned from human liver cDNA library and initially named as MB67 (Baes et al., 1994). Because of its ability to transactivate retinoic acid responsive elements constitutively, the term constitutive activator of retinoid response was first given to CAR, later shortened to constitutive active receptor (Choi et al., 1997). Subsequently, after two androstane metabolites, 5 $\alpha$ -androst-16-en-3 $\alpha$ -ol and 5 $\alpha$ -androstan-3 $\alpha$ -ol were identified as inverse agonists that repress this constitutive activity in cell-based reporter assays, constitutive androstane receptor was conferred to CAR (Forman et al., 1998). However, in actuality it has not been confirmed that androstanes repress CAR in *in vivo* environments, such as liver and primary hepatocytes, thereby regulating CAR activation. The only known mechanism to repress CAR *in vivo* is phosphorylation of threonine 38 (Mutoh et al., 2009). Phosphorylated CAR at threonine 38 is an inactive form of CAR that is sequestered in the cytoplasm in a complex with heat shock protein 90 and CCRP (Kawamoto et al., 1999; Kobayashi et al., 2003; Mutoh et al., 2009; Yoshinari et al., 2003). In response to activators, threonine 38 is dephosphorylated to activate CAR and translocate it into the nucleus. This dephosphorylation is catalyzed by protein phosphatase 2A which utilizes receptor for activated C kinase 1 (RACK1) as the specific regulatory subunit (Mutoh et al., 2013). The first CAR response sequence characterized was the phenobarbital (PB)-responsive enhancer module (PBREM) characterized within the *Cyp2b10* gene (Honkakoski et al., 1998a,b). There are two different types of CAR activators, direct and indirect activators. PB and phenytoin indirectly activate CAR without direct binding, represent one group. Another group is represented by TCPOBOP for mouse CAR and CITCO for human CAR; these are ligands of CAR (Maglich et al., 2003; Sueyoshi et al., 1999; Tzamelis et al., 2000). As to the

indirect activation mechanism, PB binds to and represses epidermal growth factor receptor (EGFR) signaling in mouse hepatocytes. Repression of EGFR transduces the PB induction signal to promote the RACK1-regulated dephosphorylation of threonine 38, thereby activating CAR and translocating it into the nucleus (Mutoh et al., 2013). How ligands activate phosphorylated CAR remains unknown. However, the fact is that, as long as threonine 38 is phosphorylated, ligands are not able to activate CAR (Mutoh et al., 2009). Current understanding of CAR activation is summarized in Figure 1.

## SULTs in biological homeostasis

SULTs catalyze transfer of a sulfonate ( $\text{SO}_3^-$ ) group from the universal sulfate donor 3'-phosphoadenosine 5'-phosphosulfate (PAPS) to hydroxyl or amino-groups of acceptor substrates (Gamage et al., 2006; Strott, 2002). SULTs sulfate both xenobiotics (e.g. *p*-nitrophenol, diethylstilbestrol,  $\alpha$ -hydroxytamoxifen, 6-hydroxymethylbenzo[*a*]pyrene) and endobiotics such as steroid hormones (Gamage et al., 2006; Glatt, 1997). Their expression is robustly influenced by xenobiotics and regulated by PXR and CAR, thereby altering biological homeostasis (Maglich et al., 2002; Rosenfeld et al., 2003).

### Bile acid homeostasis

Bile acids, the end product of hepatic cholesterol catabolism, emulsify dietary fats and vitamins to promote their absorption in the small intestine. Bile acids also play important roles in the energy homeostasis of endobiotics such as glucose and lipids (Li & Chiang, 2012). Bile acids, the secondary bile acid lithocholic acid (LCA) in particular, are toxic and can cause cholestasis in rodent models (Fisher et al., 1971; Javitt, 1966). LCA levels are elevated in patients with chronic cholestatic liver disease and are suggested to be pathogenic in colorectal cancer (Fischer et al., 1996; Kishida et al., 1997). PXR activating drugs have been used to treat cholestatic liver disease in humans (Kliwer et al., 2002). Bile acid homeostasis is tightly maintained through synthesis and degradation. Nuclear receptors PXR, CAR and FXR play various roles in regulating bile acid homeostasis. In synthesis, CYP7A1 is the key enzyme that synthesizes bile acids from cholesterol. Activation of PXR represses hepatic CYP7A1 and protects mouse livers from developing LCA-induced hepatotoxicity and mortality (Kliwer & Willson, 2002; Staudinger et al., 2001b; Xie et al., 2001).

For degradation processes, SULTs are the enzymes that play a major role in the detoxification and elimination of LCA and other bile acids (Alnouti, 2009). Sulfated bile acids are less toxic and more easily excreted than un-sulfated ones. In fact, 40–75% of LCA is sulfated in human bile acids (Alnouti, 2009). SULT2A, also known as dehydroepiandrosterone SULT, is the major enzyme to sulfate LCA (Radomska et al., 1990). PAPS synthase 2 (PAPSS2) is the enzyme that synthesizes sulfate donor PAPS and can also play a role in this sulfation-mediated degradation of LCA. PXR and CAR regulate expression of these enzymes to regulate the hepatic ability of sulfating LCA. Treatment with PXR activator PCN induces hepatic expression of SULT2A in rodents (Liu & Klaassen, 1996; Sonoda et al., 2002). Treatment also co-induces PAPSS2, thereby increasing hepatic LCA sulfation capability. Transgenic mice expressing VP-CAR increase resistance to developing LCA-induced hepatotoxicity. This CAR-mediated protection appears to be

largely associated with LCA sulfation, although it remains unknown which SULT sulfates LCA in mouse liver.

Over decades, PB has been used for treatment of pruritus associated with intrahepatic cholestasis (Jenkins & Boothby, 2002). Therefore, what the role of CAR observed in mice may be conserved in humans. However, whether or not PXR regulates SULT2A1 in humans as observed in mice remains controversial (Duanmu et al., 2002; Echchgadda et al., 2007; Fang et al., 2007). Rifampicin treatment induced SULT2A1 at least 1.5-fold in only 12 out of 23 primary hepatocytes (Fang et al., 2007). Eleven other primary hepatocytes either repressed SULT2A1 or did not alter levels. In fact, PXR repressed the *SULT2A1* promoter in HepG2-based assays *in vitro*. PXR, on the other hand, was shown to activate the *SULT2A1* expression in human colon adenocarcinoma cells (Echchgadda et al., 2007).

FXR, can be activated by cholic acid, chenodeoxycholic acid, or their conjugated derivatives at their physiological concentrations, is a physiological bile acid sensor to maintain bile acid homeostasis by controlling bile acid synthesis, metabolism and excretion (Modica et al., 2010). Activated FXR induces small heterodimer partner (SHP, NR0B2) and SHP then inactivates transcription factors that activate the *CYP7A1* gene such as liver receptor homolog 1 (LRH1, NR5A2) and hepatocyte nuclear factor 4 $\alpha$  (HNF4 $\alpha$ , NR2A1) (Chiang, 2002; Goodwin et al., 2000). FXR also prompts bile acid elimination by up-regulating SULT2A1 and transporters (Song et al., 2001). FXR, PXR and CAR appear to exert coordinated efforts to regulate bile acid homeostasis. In this, FXR acts as physiological bile acid sensor, while PXR works as pathophysiological bile acid sensor. CAR, over-expressed in FXR-KO mice and FXR/PXR-double KO mice, compensates the functions of FXR and PXR (Guo et al., 2003). In addition to these three nuclear receptors, LXRs, VDR and PPARs were also reported to up-regulate SULTs (Echchgadda et al., 2004b; Fang et al., 2005b; Uppal et al., 2007).

### Thyroid hormone homeostasis

3,5,3'-Triiodothyronine (T3) is the active form of thyroid hormone (TH) (Brent, 2012; Gereben et al., 2008). By binding TH receptors TR $\alpha$  (NR1A1) and TR $\beta$  (NR1A2), T3 plays important roles in liver regeneration and overall energy expenditure (Brent, 2012; Lopez-Fontal et al., 2010). The pro-hormone 3,5,3',5'-tetraiodothyronine (T4) is synthesized in and secreted from the thyroid gland. Thyroid-stimulating hormone (TSH) stimulates T4 synthesis and secretion. In peripheral target organs such as liver, T4 is converted to T3 through outer ring deiodination by type 1 deiodinase (D1). The levels of free T3 are *in vivo* a diagnostic indicator for TH activity in the body. T3 can be inactivated by conversion to diiodothyroxines (T2s), 3,3',5-triiodothyronine (rT3; reverse T3) or sulfated or glucuronide-conjugates (Gereben et al., 2008; Kaptein et al., 1997; Visser, 1994; Visser et al., 1993, 1998). In particular, T4 sulfation by SULTs is the quite important step that could determine the fate of T4 metabolism. T4 sulfation inhibits the outer ring deiodination of T4 while it stimulates the inner ring deiodination of T4, resulting in a rapid and irreversible inactivation of THs (Visser, 1994).

Serum rT3 levels were significantly increased after partial hepatectomy (PH) in both wild-type and CAR-KO mice (Tien et al., 2007). PB treatment decreased these levels in wild-type

but not CAR-KO mice. No significant changes were observed in the levels of total T3, free T3, total T4 or TSH in either wild-type or CAR-KO mice. While D1 was repressed in PH liver of both wild-type and CAR-KO mice, PB treatment induced D1 in wild type but not CAR-KO mice. Therefore, CAR-regulated induction of D1 appears to be a major pathway to increase rT3 levels, thereby modulating expression of TR-targeted genes in PH livers (Tien et al., 2007).

Treatment with CAR activator TCPOBOP is reported to stimulate thyroid follicular cell proliferation in rodents (Diwan et al., 1996; Qatanani et al., 2005). CAR-mediated increase of SULTs and serum TSH levels are suggested to cause a TH-related like disorder (Qatanani et al., 2005). Caloric restriction is shown to decrease serum total T3 and T4 levels only in wild-type but not in CAR-KO mice (Maglich et al., 2004). This induction of SULTs is speculated to decrease T4 and T3. However, neither were the levels of sulfated T3 and T4 determined in that report nor was the particular SULT enzyme responsible for T3 and T4 sulfation described (Maglich et al., 2004; Qatanani et al., 2005).

### **Steroid hormone homeostasis**

Steroid hormones are essential in various normal physiological processes. Steroid hormones are inactivated by sulfation and reactivated by desulfation. Hence, the expression of SULTs could have significant impact on the levels of active steroid hormones.

### **Androgen homeostasis**

Androgens, testosterone, dihydrotestosterone and androstenedione, activate androgen receptor (AR, NR3C4) to regulate the development and maintenance of male characteristics (Wang et al., 2009). Dysregulation of the androgen/AR signal is associated with the development and progression of prostate cancer (Balk & Knudsen, 2008). SULT2A1 is responsible for sulfation of androgens. SULT2A1 is predominantly expressed in the liver, intestine, adrenal glands prostate and ovary in humans (Chen et al., 2003; Falany, 1997a,b; Falany et al., 1995; Javitt et al., 2001). SULT2A1 is expressed in the mouse liver (Alnouti & Klaassen, 2006). Overexpression of rat SULT2A1 attenuated AR-mediated trans-activation in human prostate cancer-derived PC-3 cells (Chan et al., 1998).

Activation of PXR has been implicated in androgen deprivation in mice *in vivo* (Zhang et al., 2010). The study utilized genetically engineered mice either expressing constitutively activated PXR (VP-PXR) in the liver and small intestine or lacking the expression of PXR. Both VP-PXR and activation of PXR by PCN partially inhibited prostate regeneration by the exogenous testosterone administration in castrated mice, in which the serum testosterone levels were lowered in a PXR-dependent manner. Activation of PXR significantly inhibited the proliferation of human androgen-responsive prostate cancer cells. Activation of PXR was reported to inactivate androgens by up-regulating the expression of SULT2A1 for sulfation. Thus, PXR may antagonize the dysregulation caused by the androgen/AR signal, particularly in the development and progression of prostate cancer.

LXR $\alpha$  and LXR $\beta$  are sterol sensors and are activated by endogenous oxysterols, oxidized derivatives of cholesterol (Zhao & Dahlman-Wright, 2010). LXR $\alpha$  is highly expressed in

organs such as liver, adipose tissue, kidney and macrophages, whereas LXR $\beta$  is ubiquitously expressed. In experiments with VP-LXR $\alpha$  transgenic and LXR $\alpha$ /LXR $\beta$  double-KO mice demonstrated that LXRs directly activate the hepatic expression of SULT2A1 and attenuate hepatotoxicity and cholestasis caused by LCA (Uppal et al., 2007). The LXRSULT2A1 pathway is linked to androgen deprivation and also to decreasing serum testosterone levels and decrease prostate epithelial proliferation in castrated mice and in human androgen-responsive prostate cancer cells (Lee et al., 2008). Moreover, LXRs also promote androgen inactivation by down-regulating the expression of steroid sulfatase in mouse prostate and cancer cells.

### Estrogen homeostasis

By activating estrogen receptors ER $\alpha$  (NR3A1) and ER $\beta$  (NR3A2), estrogens play key roles in a wide range of physiological aspects throughout life; the development and maintenance of female characteristics, male characteristics such as spermatogenesis and libido and energy homeostasis (Barros & Gustafsson, 2011; Carreau & Hess, 2010; Nilsson & Gustafsson, 2011; Robertson et al., 1999). Aromatase synthesizes active estrogens, while SULTs sulfates estrogens to inactivate them. Sulfated estrogens can be re-activated by sulfatases (Suzuki et al., 2003a). Estrogen SULT (SULT1E1) is the major enzyme that sulfates estrogens in reproductive tissues (Alnouti & Klaassen, 2006). SULT1E1 is considered to be a potent prognostic factor in human breast cancer (Suzuki et al., 2003b). SULT1E1-KO mice develop placental thrombosis and spontaneous fetal loss in female mice (Tong et al., 2005) and an age-dependent structural and functional lesions in the male reproductive systems (Qian et al., 2001). SULT1E1 is also expressed in human livers (Miki, 2002). Both PXR and CAR repress SULT1E1 expression in human primary hepatocytes (Kodama et al., 2011; Lambert et al., 2009). SULT1E1, expressed at low levels in normal mouse livers, increases its expression levels in diabetogenic db/db mice (Song et al., 1995). CAR activates the *Sult1e1* gene in mouse livers (Aleksunes & Klaassen, 2012; Sueyoshi et al., 2011). Although CAR-mediated increase of SULT1E1 enzyme facilitates clearance of exogenously administered estrogen, levels of circulating estrogens remain constant (Sueyoshi et al., 2011). Whether or not PXR also regulates the *Sult1e1* gene in mouse livers has not been confirmed. Utilizing VP-LXR $\alpha$  transgenic and LXR $\alpha$ /LXR $\beta$  double-KO mice, LXR $\alpha$  is shown to activate directly the *Sult1e1* gene and induce estrogen SULT activity in livers (Gong et al., 2007). VP-LXR $\alpha$  transgenic females exhibit resistance against uterine epithelial cell proliferation induced by estrogen at pharmacological doses as well as growth of xenografted estrogen-responsive breast cancer cells. These functions of LXR and SULT1E1 have not been confirmed in humans.

### Regulation of the *SULT* genes

#### Activation mechanism

The regulatory mechanism by which nuclear receptors regulate the *SULT2A* genes has been investigated in both rodents and humans (Runge-Morris & Kocarek, 2005; Runge-Morris et al., 2013). The proximal promoters of *Sult2a* genes are well conserved in rodents and contain binding sites for liver-enriched transcription factors HNF1 and CCAAT/enhancer-binding proteins (C/EBPs) (Song et al., 1998). Androgen receptor indirectly represses the rat

*Sult2a1* gene by preventing the OCT-1 transcription factor and C/EBPs from activating its promoter (Song et al., 1998). Glucocorticoid induces expression of C/EBP $\alpha$  and C/EBP $\beta$ , which in turn activate the rat *Sult2a1* promoter in primary hepatocytes (Fang et al., 2005a). The *Sult2a* promoter conserves an IR0 (inverted repeat without a spacing nucleotide) motif, through which PXR, CAR, VDR, FXR or LXR activate the *Sult2a* genes (Echchgadda et al., 2004a,b; Saini et al., 2004; Sonoda et al., 2002; Uppal et al., 2007). Figure 2(A) summarizes current views of how *Sult2a* genes are regulated.

The proximal promoter of human *SULT2A1* gene contains an IR2 (inverted repeat with two spacing nucleotides), DR4 (direct repeat with four spacing nucleotides), C/EBP and DR1 (direct repeat with a spacing nucleotide) motifs (Echchgadda et al., 2007; Song et al., 2006). PXR, CAR or VDR activates the *SULT2A1* gene through these motifs in human colon adenocarcinoma-derived cells (Echchgadda et al., 2007; Song et al., 2006). For instance, vitamin D<sub>3</sub> treatment recruits VDR and C/EBP $\alpha$  to the human *SULT2A1* promoter, thereby activating the *SULT2A1* gene (Song et al., 2006). PXR binds to both IR2 and DR4 motifs to activate this gene (Echchgadda et al., 2007). On the other hand, CAR can activate the promoter by binding to either IR2 or DR4 (Echchgadda et al., 2007). HNF4 $\alpha$  binds to DR1 motif and determines the basal promoter activity of the *SULT2A1* gene as well as synergizes the PXR- and CAR-mediated activation (Echchgadda et al., 2007). PPAR $\alpha$  constitutively binds to a distal DR1 motif (at -5949 bp upstream) and activates *SULT2A* genes in human but not rat primary hepatocytes (Fang et al., 2005b).

### Repression mechanism

Rifampicin treatment represses expression of the *SULT1E1* gene in human primary hepatocytes (Kodama et al., 2011). Activity of PXR-mediated repression was delineated to the distal enhancer (-1000/-901) that contains overlapping DR1 and DR2 (direct repeat with two spacing nucleotides) motifs and the proximal promoter within the *SULT1E1* promoter (Kodama et al., 2011). When HNF4 $\alpha$  binds to distal DR1 and DR2 motifs, the promoter loops to locate distal enhancer to juxtapose the proximal promoter, thereby forming a transcriptionally active chromatin structure. This looping associates with an increase histone H3 acetylation in both distal and proximal promoter regions. Upon rifampicin activation, PXR disrupts this active chromatin structure and represses the *SULT1E1* gene (Figure 2B). Chromatin immunoprecipitation assays did not detect PXR binding to either regions of the promoter. PXR may transiently interact with HNF4 $\alpha$  and dissociate from the promoter together with HNF4 $\alpha$ . In human primary hepatocytes, rifampicin treatment results in either induction or repression of *SULT2A1* (Fang et al., 2007). Although reasons for these different responses are not known, studies with *in vitro* cell-based analyses suggest that PXR prevents HNF4 $\alpha$  from activating the *SULT2A1* promoter via both distal (at -6160 bp) and proximal DR1 motifs, thereby repressing the *SULT2A1* gene. Therefore, HNF4 $\alpha$  can be a common target when PXR represses *SULT* genes. CITCO treatment also decreases *SULT1E1* mRNA levels in human primary hepatocytes. Whether or not CAR utilizes the same mechanism as described for the PXR-mediated repression remains as a possibility at the present time.



## Conclusion

By cross talking with various nuclear receptors (e.g. HNF4 $\alpha$ , FXR, LXRs, PPARs and VDR), PXR and CAR regulate *SULT* genes in response to both xeno- and endobiotics. Through the regulation of *SULT* expression, these nuclear receptors are involved not only in metabolic detoxification and excretion of xenobiotics but also in maintaining biological homeostasis (Figure 3). The molecular mechanism by which *SULT* genes can be regulated by these nuclear receptors is beginning to be understood but further investigations for complete understanding is required.

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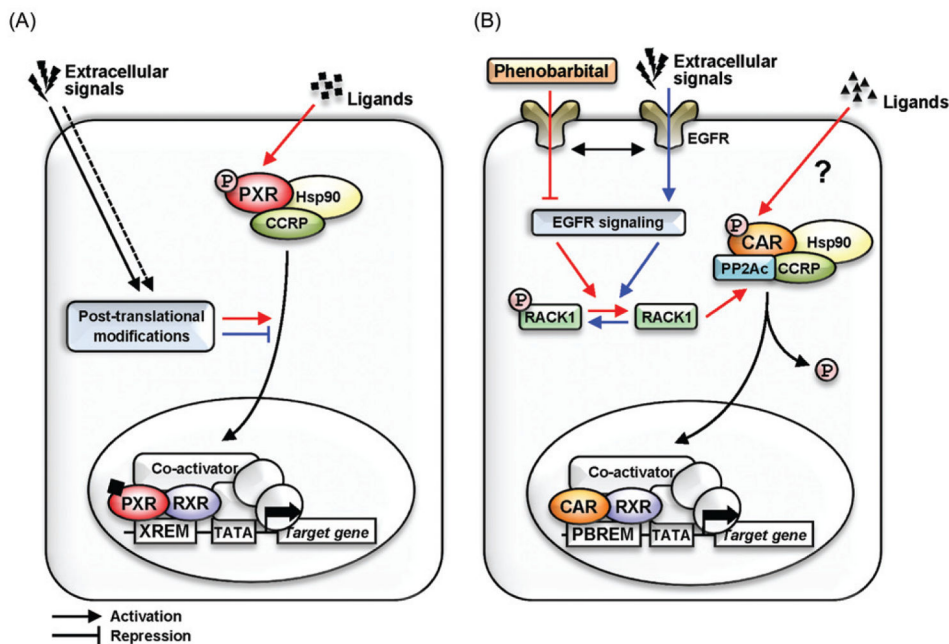
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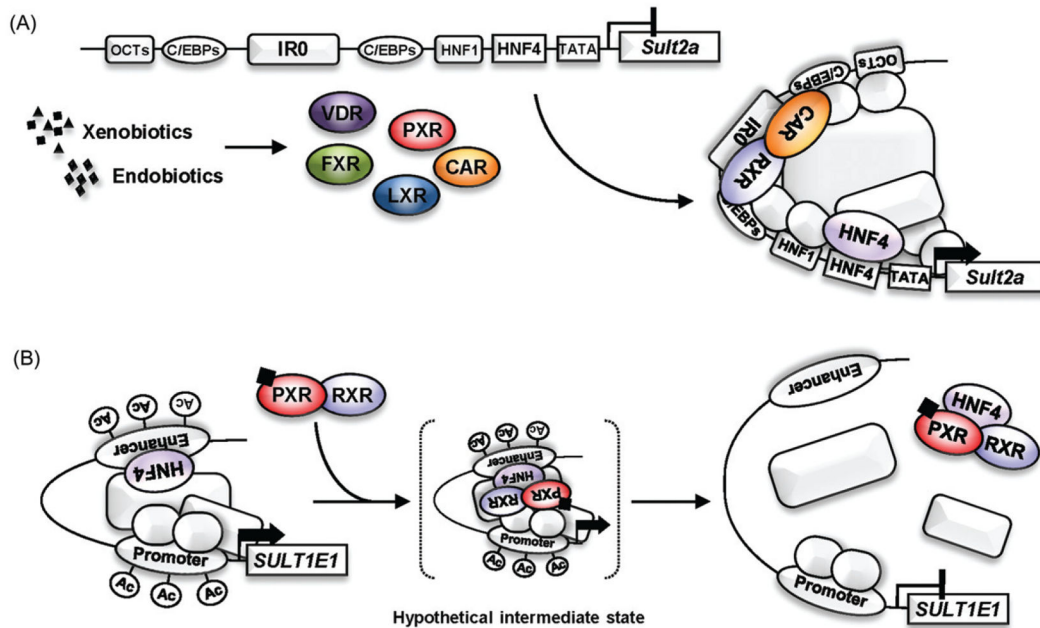
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**Figure 1.**

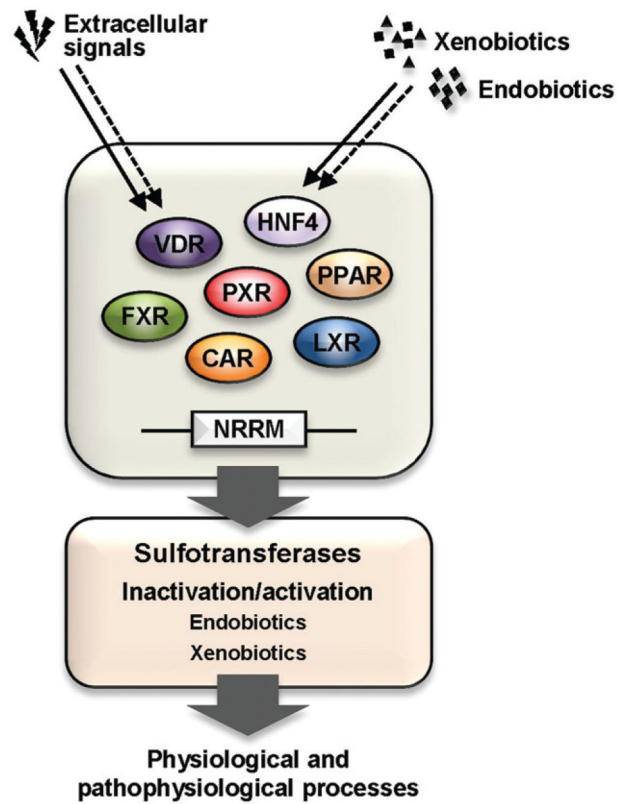
Activation mechanism of PXR and CAR. (A) Current understanding of PXR activation. Being part of the Hsp90–CCR<sub>P</sub> complex, PXR is predominantly localized in the cytoplasm. Upon ligand binding, PXR translocates into the nucleus, heterodimerizes with RXR, and transactivates response sequences such as XREM in the *CYP3A4* gene. Cell signaling pathways have been implicated to modulate PXR activity, either positively or negatively, where PXR undergoes post-translational modifications including phosphorylation, acetylation and SUMOylation. (B) Current understanding of CAR activation. Under unstimulated condition, threonine 38 of CAR is phosphorylated. Phosphorylation status of threonine 38 determines the intercellular localization of CAR, and this phosphorylated form is inactive and mostly localized in the cytoplasm as a part of Hsp90–CCR<sub>P</sub> complex. Dephosphorylation of threonine 38 allows CAR to translocate into the nucleus, which in turn heterodimerizes with RXR and transactivates response sequences such as PBREM in the *Cyp2b10* gene. Upon stimulation, PP2A catalyzes dephosphorylation of threonine 38, whereas RACK1 serves as the specific regulatory subunit. Phenobarbital, a typical indirect CAR activator, binds to EGF receptor antagonizes EGFR signaling to promote the RACK1-regulated dephosphorylation. On the other hand, how ligands activate CAR remains unknown. XREM: xenobiotic responsive enhancer module; PBREM: phenobarbital responsive enhancer module; PP2A: protein phosphatase 2A; RACK1: receptor for activated protein C kinase 1; CCR<sub>P</sub>: cytoplasmic CAR retention protein; EGFR: epidermal growth factor receptor.



**Figure 2.**

Regulation mechanism of the *SULT* genes. (A) Activation of the rodent *Sult2a* promoters by nuclear receptors. In rodents, the proximal promoters of *Sult2a* genes are well conserved. Nuclear receptors (e.g. PXR, CAR, VDR, FXR and LXR) bind to an IR0 and activate the *Sult2a* genes in concert with liver-enriched transcription factors. (B) Repression of the human *SULT1E1* promoter by PXR. HNF4 $\alpha$  is a transcription factor that determines the basal expression level of *SULT1E1* gene in human primary hepatocytes. The binding of HNF4 $\alpha$  to the enhancer prompts the *SULT1E1* promoter to form a transcriptionally active chromatin structure by looping it position close to the proximal promoter. Upon activation by ligand, PXR interacts with HNF4 $\alpha$  and dissociate it from the enhancer, resulting in disruption of the active chromatin structure, concomitant with deacetylation of histone H3 on both the enhancer and proximal promoter regions.





**Figure 3.** Biological role of nuclear receptors in regulation of the *SULT* genes. Through the regulation of SULT expression, nuclear receptors are involved in xenobiotic metabolism and endobiotic homeostasis, and thereby play important roles in physiological and pathophysiological processes. NRRM: nuclear receptor responsive module.