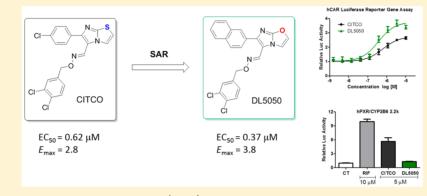
DL5050, a Selective Agonist for the Human Constitutive Androstane Receptor

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S Supporting Information



ABSTRACT: The constitutive androstane receptor (CAR) is a xenobiotic sensor governing the transcription of genes involved in drug disposition, energy homeostasis, and cell proliferation. However, currently available human CAR (hCAR) agonists are nonselective, which commonly activate hCAR along with other nuclear receptors, especially the closely related human pregnane X receptor (hPXR). Using a well-known hCAR agonist CITCO as a template, we report our efforts in the discovery of a potent and highly selective hCAR agonist. Two of the new compounds of the series, 18 and 19 (DL5050), demonstrated excellent potency and selectivity for hCAR over hPXR. DL5050 preferentially induced the expression of CYP2B6 (target of hCAR) over CYP3A4 (target of hPXR) on both the mRNA and protein levels. The selective hCAR agonist DL5050 represents a valuable tool molecule to further define the biological functions of hCAR, and may also be used as a new lead in the discovery of hCAR agonists for various therapeutic applications.

KEYWORDS: Nuclear receptor, constitutive androstane receptor, structure-activity relationship, pregnane X receptor

he constitutive androstane receptor (CAR, NR1I3) belongs to the nuclear receptor (NR) subfamily 1I that also includes the pregnane X receptor (PXR, NR1I2) and vitamin D receptor (VDR, NR1I1).^{1,2} CAR plays an important role orchestrating the metabolism and detoxification of potentially toxic chemicals,^{3,4} by upregulating genes that are involved in the clearance of both xenobiotics and endobiotics.^{5,6} Although CAR-mediated metabolism of drugs (e.g., acetaminophen) can lead to the production of toxic metabolites,' induction of CAR regulated genes has emerged as a therapeutic strategy for the treatment of various diseases. For instance, phenobarbital, a prototypical CAR activator, has been used to treat neonatal jaundice through its induction of UDP glucuronosyltransferase 1A1 (UGT1A1), the ratelimiting enzyme responsible for the metabolism of bilirubin.⁸ Maglich and co-workers have reported an important regulatory role of CAR in the metabolism of thyroid hormones during caloric restriction,⁹ while other studies revealed that activation of CAR ameliorates high-fat diet-induced obesity and diabetes

in mice, suggesting that CAR modulators might be potential therapeutics for metabolic disorders.^{10,11} It is well-known that cyclophosphamide (CPA), a widely used chemotherapeutic prodrug, is converted to a therapeutically active metabolite 4hydroxy-CPA primarily by CYP2B6 (target of hCAR) and to an inactive metabolite N-DCE-CPA predominantly by CYP3A4 (target of hPXR).^{12,13} Recent studies from our laboratory have shown that activation of CAR sensitizes cyclophosphamide (CPA)-based cancer treatment by enhancing biotransformation of CPA to its activate metabolite 4hydroxy CPA.^{14,15}

The activation mechanism of CAR begins with a ligandtriggered dissociation of CAR from its cochaperone partners, cytoplasmic CAR retention protein (CCRP), and heat shock protein 90 (Hsp90), in the cytosol (Figure 1).¹⁶ Dephosphor-

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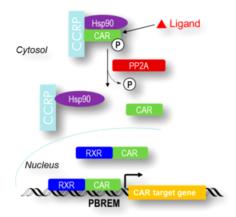


Figure 1. Activation mechanism of CAR.

ylation of CAR at Thr38 by protein phosphatase 2A (PP2A) generates free CAR,¹⁷ which then translocates into the nucleus and forms a heterodimer with the retinoid X receptor (RXR). The complex of RXR-CAR regulates its target genes upon binding to cis-acting responsive enhancer modules upstream of target genes.¹⁸ Different from most known NRs, CAR demonstrates constitutive and ligand-independent activity.¹⁹

Similar to most classical steroid receptors, CAR contains multiple functional domains including an *N*-terminal DNAbinding domain (DBD) that directly interacts with the promoters of its target genes, a hinge domain, and a multifunctional ligand-binding domain (LBD). The LBD contains a ligand binding pocket, a dimerization interface that binds to RXR, and a *C*-terminal ligand-dependent activation helix (AF2).²⁰ The LBD of human CAR (hCAR) can bind to a wide range of structurally diversified ligands.^{21–23} Earning its name as an orphan nuclear receptor, physiologically relevant endogenous ligands of hCAR have not been identified as of yet.

Within the NR1I subfamily, hCAR shares a modestly conserved LBD and highly conserved DBD with human PXR (hPXR).²⁴ As a result, these two nuclear receptors bind to multiple common ligands²⁵ and regulate an overlapping set of genes (e.g., CYP2B6 and CYP3A4).^{24,26,27} Compared to hPXR, hCAR has a much smaller LBD (675 Å³ vs ~1200 Å³ for hPXR).²³ Unlike hPXR to which selective agonists (e.g., SR12813²⁸) have been identified, no highly selective hCAR activator has been disclosed. The availability of potent and highly selective ligands for hCAR will not only benefit the confirmation of target genes for this important nuclear receptor but also verify whether hCAR is a viable therapeutic target.

In the past decade, efforts in both screens^{29–32} and medicinal chemistry³³ have been devoted to the identification and development of agonists for hCAR. Among all known hCAR agonists, the GSK high-throughput screening hit CITCO is the most well-known for its good potency and selectivity.^{5,20} CITCO potentiates constitutive activity of hCAR by stabilizing the constitutive AF2-coactivator interaction.²³ Although CITCO only indicates moderate selectivity for hCAR over hPXR, it is by far the most widely used chemical tool in elucidating the function of hCAR.^{34–37}

Herein, we report the development of hCAR agonists based on the structure of CITCO (Figure 2). Specifically, the 4-Clphenyl, imidazothiazole, and 3,4-diCl-phenyl groups of CITCO (1) were modified using compounds 2-29. Seven compounds were selected and further evaluated for their

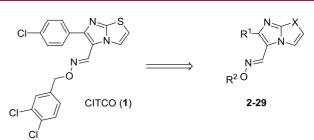
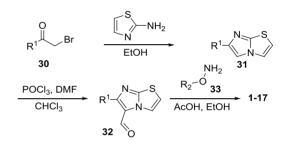


Figure 2. Design of compounds 2-29 from CITCO.

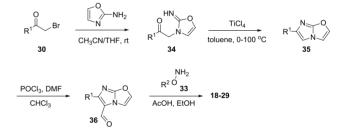
selectivity over hPXR. Two of the new compounds, **18** and **19** (DL5050), demonstrated excellent selectivity for hCAR over hPXR while maintaining similar hCAR agonistic potency to CITCO. In particular, compound DL5050 selectively induced the expression of cytochrome P450 CYP2B6 over CYP3A4 at both the mRNA and protein levels (Figure 4).

The synthetic strategies for compounds 1-29 were outlined in Schemes 1 and 2. Condensation of bromomethyl ketone

Scheme 1. Synthesis of Compounds 1-17



Scheme 2. Synthesis of Compounds 18-29



(30) with thiazol-2 amine provided imidazothiazole 31 (Scheme 1). Vilsmeier formylation of compound 31 gave aldehyde 32 in good yields. Treatment of aldehyde 32 with hydroxylamine 33 in EtOH led to the formation of final products 1-17.

As shown in Scheme 2, a modified four-step route was used to synthesize compounds 18-29. Condensation of bromomethyl ketone (30) with oxazol-2-amine in the mixture of THF and MeCN generated intermediate 34. TiCl₄-mediated cyclization of compound 34 yielded imidazooxazole 35. Vilsmeier reaction of compound 35 using POCl₃ in chloroform gave aldehyde 36 in excellent yields. Oximation of aldehyde 35 with substituted hydroxylamine 33 in the presence of AcOH led to the final compounds 18-29.

The new compounds were first evaluated for their potential modulation of hCAR activity using a cell-based quantitative high-throughput screening (qHTS) luciferase reporter assay on a double stable HepG2-CYP2B6-hCAR cell line.³¹ The potency was expressed as 50% effective concentration

(EC₅₀), and the maximum luciferase signal fold increase of a new compound over vehicle control as E_{max} . In addition, cytotoxicity of new compounds was evaluated in a luciferase-coupled ATP quantitation cell viability assay (Table S1).

In order to optimize \mathbb{R}^1 , a collection of compounds (1–17) with a common (*E*)-imidazo[2,1-*b*]thiazole-5-carbaldehyde *O*-(3,4-dichlorobenzyl)oxime scaffold were tested (Table 1). The

Table 1	. EC50	and E_{max}	of Compounds	$1 - 17^{a}$
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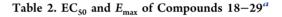
Cmpd	\mathbb{R}^1	EC50 (µM)	$E_{\rm max}$	$E_{\text{max}}/\text{EC}_{50}$
1	4-Cl-Ph	0.62 ± 0.08	2.8	4.5
2	Ph	0.58 ± 0.36	2.3	4.0
3	4-F-Ph	0.67 ± 0.10	2.6	3.9
4	4-Br-Ph	0.48 ± 0.22	2.4	5.0
5	4-Me-Ph	0.49 ± 0.23	2.7	5.5
6	4-CF ₃ -Ph	0.88 ± 0.09	2.3	2.6
7	4-MeO-Ph	0.49 ± 0.27	2.4	4.9
8	4- <i>i</i> Pr-Ph	0.59 ± 0.13	3.0	5.1
9	4- <i>t</i> Bu-Ph	2.4 ± 0.30	2.0	0.83
10	4-CN-Ph	3.4 ± 0.29	2.2	0.65
11	4-Cy-Ph	> 100	2.3	< 0.03
12	4-Ph-Ph	ND	1.1	-
13	$\bigwedge \rightarrow$	ND	1.2	-
14	3-Cl-Ph	1.1 ± 0.08	2.3	2.1
15	2-Naph	0.38 ± 0.31	2.3	6.1
16	1-Naph	0.94 ± 0.73	2.2	2.3
17	Ethyl	2.6 ± 0.16	3.6	1.4

^{*a*}EC₅₀ and E_{max} values were calculated by nonlinear regression. Data are presented as mean \pm SEM of at least three independent experiments in quadruplicate. ND (if the maximum tested concentration produced no effect).

 $E_{\rm max}$ values were shown as relative values compared to the basal value (negative control). Under our assay conditions, CITCO (1) indicated an EC₅₀ of 0.62 μ M and E_{max} of 2.8, which led to an E_{max} /EC₅₀ value of 4.5. Removing the Cl atom, the analogue 2 showed a slightly increased potency. However, the E_{max} of compound 2 was also decreased. While substitution of the Cl atom with an F led to a new compound (3) with similar EC_{50} and E_{max} values, a larger Br group (4) at this position brought increased agonistic activity ($E_{max}/EC_{50} = 5.0$). Similarly improved activities were also observed when substituting the Cl with an Me group (5). Addition of a more polar CF_3 group (6) yielded a weaker agonist. These results suggested that there was no demand for a hydrogen bond receptor at this position, instead, a small hydrophobic pocket might be available for new compound development. OMe (7) and *i*-Pr (8) groups at this position led to new compounds with slightly improved agonist activity. Large substituents, such as t-Bu (9),

CN (10), Cy (11), and Ph (12), were also explored. Compounds 9 and 10 demonstrated weak agonistic activity for hCAR ($E_{max}/EC_{50} < 1.0$), and compounds 11 and 12 showed essentially no activity toward hCAR. Our results demonstrated that the hCAR agonist activity decreases while the size of substituents increase. Compound 13, with a bulky substitution on the 3,4-position also indicated no activity for hCAR, which is in agreement with our previous discovery. Shifting the Cl atom to the 3-position from the 4-position yielded compound 14 with a 2.6-fold decreased EC_{50} and slightly lower E_{max} value, implying that 3-substitution can also be tolerated by the agonist. Replacement of the 4-Cl-phenyl group with naphthalen-2-yl group led to compound 15 with an improved potency although its E_{max} was also decreased. The $E_{\rm max}/{\rm EC}_{50}$ value of compound 15 was 1.4-fold better than that of CITCO. The regioisomer naphthalen-1-yl 16, however, indicated significantly lower EC50 and Emax values. Replacement of the aromatic group with an ethyl group led to compound 17 having a weak potency, highlighting the importance of an aromatic ring at this position. The cytotoxicity of new compounds was evaluated (Table S1). Similar to CITCO, our new compounds started to show toxicity at concentrations above 10 μ M.

Next, in an attempt to increase the E_{max} value of the compounds, we synthesized compounds 18-29 (Table 2).



Cmpd	\mathbb{R}^1	\mathbb{R}^2	$EC_{50}(\mu M)$	$E_{\rm max}$	$E_{\text{max}}/\text{EC}_{50}$
18	4-Cl-Ph	3,4-diCl-Bn	0.41 ± 0.09	4.2	10
19	2-Naph	3,4-diCl-Bn	0.37 ± 0.11	3.8	10
20	2-Naph	2,4-diCl-Bn	1.5 ± 0.12	3.5	2.3
21	2-Naph	2-Cl-Bn	1.1 ± 0.11	3.8	3.5
22	2-Naph	4-Cl-Bn	0.99 ± 0.15	4.2	4.2
23	2-Naph	Bn	0.84 ± 0.10	4.4	5.2
24	2-Naph	$\mathcal{O}\mathcal{H}$	0.98 ± 0.10	3.7	3.8
25	2-Naph	perF-Bn	1.7 ± 0.13	3.1	1.8
26	2-Naph	4-CF ₃ -Bn	0.47 ± 0.18	2.8	6.0
27	2-Naph	4-MeO-Bn	4.4 ± 0.44	3.3	0.75
28	2-Naph	4- <i>t</i> Bu-Bn	7.9 ± 0.48	2.3	0.29
29	2-Naph	4-PhO-Bn	ND	1.0	-

 ${}^{a}\text{EC}_{50}$ and E_{max} values were calculated by nonlinear regression. Data are presented as mean \pm SEM of at least three independent experiments in quadruplicate. ND (if the maximum tested concentration produced no effect).

Replacing the imidazothiazole core with imidazoxazole, in CITCO and **15**, yielded two new compounds, **18** and **19**, with similar EC₅₀ values but significantly larger E_{max} values (E_{max} / EC₅₀ = 10 for both **18** and **19**). These results indicated that the imidazoxazole core is preferred over the imidazothiazole for generating larger E_{max} values. Using compound **19** as a template, we studied the substitution effects on R². When the 3-Cl atom was moved to the 2-position, the resulting 2,4-diCl-Bn analogue **20** indicated a 4.1-fold decreased EC₅₀ value.

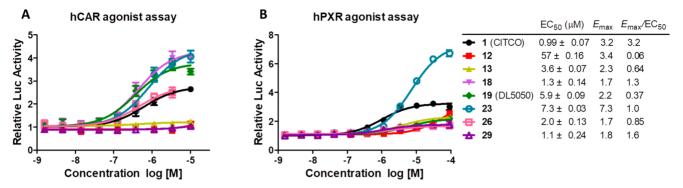


Figure 3. Selectivity of eight compounds 12, 13, 18, 19 (DL5050), 23, 26, 29, and CITCO (1) for hCAR over hPXR. (A) Dose-dependent activation of hCAR by CITCO, 12, 13, 18, 19, 23, 26, and 29 in high throughput hCAR agonist luciferase gene reporter assays. (B) Dose-dependent activation of hPXR by CITCO, 12, 13, 18, 19, 23, 26, and 29 in high throughput hPXR agonist luciferase gene reporter assays.

After removal of one of the two Cl atoms from 20, the resulting compounds, 21 and 22, indicated 3.0- and 2.7-fold decreased EC₅₀ values, respectively. When both Cl atoms of compound 19 were removed, the resulting compound 23 was 2.3-fold weaker; however, this compound also indicated a large $E_{\rm max}$ value of 4.4. Replacement of the ethylene linker with a structurally rigid five-membered ring generated a new compound (24) with diminished values for both EC_{50} and $E_{\rm max}$. This result indicated the necessity of a flexible linker. The analogue with a pentafluorophenyl group (25) was less potent and efficacious in comparison to the unsubstituted phenyl analogue 23. In addition, different substitutions at the paraposition of the Bn group, including CF_3 (26), OMe (27), t-Bu (28), and PhO (29) were introduced to probe the depth of the binding pocket for R². While compound 26 showed a comparable EC_{50} and E_{max} values to those of CITCO, compounds 27 and 28 indicated weak agonistic activities; in particular, the phenoxy analogue 29 demonstrated no agonistic activity toward hCAR. These results highlighted that large substitutions on the phenyl ring could not be tolerated in the pocket to which R²was bound. It is worth noting that the imidazoxazole-based compounds displayed less cytotoxicity compared to the ones listed in Table 1. In particular, compounds 18, 19, 23, and 26 showed no obvious cytotoxicity at the maximum tested concentration (92 μ M) (Table S1).

Based on the predicted hCAR activity, we selected seven new compounds including four potent hCAR agonists (18, 19 [DL5050], 23, and 26) and three weak hCAR agonists (12, 13, and 29) and tested their capacity in hPXR activation using a luciferase reporter assay (Figure 3B). The three weak hCAR agonists (12, 13, and 29) also indicated minimum agonistic activity for hPXR (Figure 3B). These results showed that CITCO analogs with large hydrophobic substitutions on either aromatic tail could abolish the potency in both hCAR and hPXR activation. These results also provided valuable information to define the size limitation of the LBD for both hCAR and hPXR based on the scaffold of CITCO. Among the four potent hCAR agonists, 23 exhibited potent hPXR activation at high concentrations with an E_{max} of 7.3 and EC_{50} of 7.3 μ M (Figure 3B). Notably, compounds 18, 19, and 26 indicated negligible agonistic activity of hPXR and thereby represent selective agonists for hCAR over hPXR.

Given that 26 demonstrated an hCAR activity that is markedly lower than that of 18 and 19 (DL5050), we next evaluated the effects of 18 and DL5050 on the induction of CYP2B6 and CYP3A4 genes in cultured human primary hepatocytes (HPH). As shown in Figure 4A, both 18 and

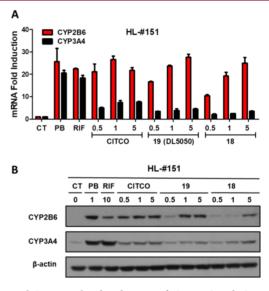


Figure 4. hCAR regulated induction of CYP2B6 and CYP3A4 in HPHs. (A) CYP2B6 and CYP3A4 mRNA levels were measured in HPHs prepared from human liver donors (HL-#151). The HPHs were treated with CITCO (0.5, 1, and 5 μ M), PB (1 mM), RIF (10 μ M), DL5050 (0.5, 1, and 5 μ M), **18** (0.5, 1, and 5 μ M), or vehicle control. (B) Representative immunoblots of CYP2B6 and CYP3A4 proteins in HPHs from liver donors #151. RT-PCR data obtained from three independent experiments were expressed as mean \pm SD normalized against vehicle control.

DL5050 induced CYP2B6 mRNA expression in a concentration-dependent manner while only moderately altered the expression of CYP3A4. This was significantly different to the hCAR and hPXR dual activator (PB), the selective hPXR agonist (RIF), and even the moderately selective hCAR agonist (CITCO). At the protein level, the naphthyl analog DL5050 also demonstrated a similar concentration-dependent selective induction of CYP2B6 over CYP3A4 (Figure 4B). Conversely, compound **18** only showed marginal induction of CYP2B6 protein.

The binding mode of compound DL5050 in hCAR was also studied (Figure 5). The dichloro-phenyl ring of DL5050, which is the same as in CITCO, rests in the hydrophobic pocket flanked by Phe132, Tyr224, and Tyr 217. The imidazo[2,1-*b*]oxazole core of DL5050 fit deeper at the junction of helixes H5 and H3 than the imidazothiazole core of CITCO. Finally, the naphthyl ring of DL5050 establishes a $\pi-\pi$ stacking interaction with Phe234, which is not present in

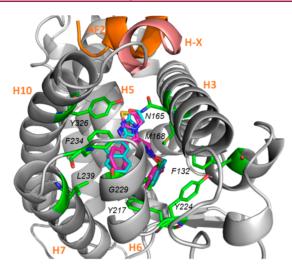


Figure 5. Docking result of compound DL5050 (orange) in the ligand-binding domain of hCAR in comparison with CITCO (magenta) using AutoDock Vina.³⁸ Key hCAR residues that are expected to mediate interactions with ligands are highlighted in green. Helix X is showed in light pink. The activation function 2 helix (AF2) is shown in orange.

CITCO, potentially providing a basis for the heightened activity of DL5050.

In summary, structure optimization of CITCO has yielded new hCAR agonists **18** and **19** (DL5050). While maintaining a similar hCAR activation potency as that of CITCO, these new compounds indicated a high specificity for hCAR over its closely related nuclear receptor, hPXR. One of the two new compounds, DL5050, exhibited potent induction of CYP2B6 gene at both the mRNA and protein levels while only resulted in minimal induction of CYP3A4. DL5050 displayed a strong induction preference in CYP2B6 over CYP3A4, which exceeds that of the known hCAR activator CITCO. The unique profile of compound DL5050 indicates this compound can be a tool for elucidating biological functions of hCAR with minimal hPXR interference. Additionally, the selectiveness of DL5050 for hCAR suggests it may be used as a therapeutic agent for various human diseases.

ASSOCIATED CONTENT

S Supporting Information

The Supporting Information is available free of charge on the ACS Publications website at DOI: 10.1021/acsmedchem-lett.9b00079.

General experimental procedures, supplementary data, and copies of ¹H and ¹³C NMR spectra and HPLC of compounds (PDF)

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Notes

The authors declare no competing financial interest.

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ABBREVIATIONS

CAR, constitutive androstane receptor; NR, nuclear receptors; PXR, pregnane X receptor; VDR, vitamin D receptor;; CCRP, cytoplasmic CAR retention protein; RXR, retinoid X receptor; ER, estrogen receptor; GR, glucocorticoid receptor; DBD, DNA-binding domain; LBD, ligand binding domain; qHTS, quantitative high-throughput screening

REFERENCES

(1) Wei, P.; Zhang, J.; Egan-Hafley, M.; Liang, S.; Moore, D. D. The nuclear receptor CAR mediates specific xenobiotic induction of drug metabolism. *Nature* **2000**, *407*, 920–923.

(2) Timsit, Y. E.; Negishi, M. CAR and PXR: the xenobiotic-sensing receptors. *Steroids* **2007**, *72*, 231–246.

(3) Xie, W.; Radominska-Pandya, A.; Shi, Y.; Simon, C. M.; Nelson, M. C.; Ong, E. S.; Waxman, D. J.; Evans, R. M. An essential role for nuclear receptors SXR/PXR in detoxification of cholestatic bile acids. *Proc. Natl. Acad. Sci. U. S. A.* **2001**, *98*, 3375–3380.

(4) Tzameli, I.; Moore, D. D. Role reversal: new insights from new ligands for the xenobiotic receptor CAR. *Trends Endocrinol. Metab.* **2001**, *12*, 7–10.

(5) Maglich, J. M.; Parks, D. J.; Moore, L. B.; Collins, J. L.; Goodwin, B.; Billin, A. N.; Stoltz, C. A.; Kliewer, S. A.; Lambert, M. H.; Willson, T. M.; Moore, J. T. Identification of a novel human constitutive androstane receptor (CAR) agonist and its use in the identification of CAR target genes. *J. Biol. Chem.* **2003**, *278*, 17277–17283.

(6) Stanley, L. A.; Horsburgh, B. C.; Ross, J.; Scheer, N.; Wolf, C. R. Drug Metab. Rev. 2006, 38, 515–597.

(7) Zhang, J.; Huang, W.; Chua, S. S.; Wei, P.; Moore, D. D. Modulation of acetaminophen-induced hepatotoxicity by the xenobiotic receptor CAR. *Science* **2002**, *298*, 422–424.

(8) Huang, W.; Zhang, J.; Chua, S. S.; Qatanani, M.; Han, Y.; Granata, R.; Moore, D. D. Induction of bilirubin clearance by the constitutive androstane receptor (CAR). *Proc. Natl. Acad. Sci. U. S. A.* **2003**, *100*, 4156–4161.

(9) Maglich, J. M.; Watson, J.; McMillen, P. J.; Goodwin, B.; Willson, T. M.; Moore, J. T. The nuclear receptor CAR is a regulator of thyroid hormone metabolism during caloric restriction. *J. Biol. Chem.* **2004**, *279*, 19832–19838.

(10) Gao, J.; He, J.; Zhai, Y.; Wada, T.; Xie, W. The constitutive androstane receptor is an anti-obesity nuclear receptor that improves insulin sensitivity. *J. Biol. Chem.* **2009**, *284*, 25984–25992.

(11) Dong, B.; Sara, P. K.; Huang, W.; Chen, W.; Abu-Elheiga, L. A.; Wakil, S. J.; Stevens, R. D.; Ilkayeva, O.; Newgard, C. B.; Chan, L.; Moore, D. D. Activation of nuclear receptor CAR ameliorates diabetes and fatty liver disease. *Proc. Natl. Acad. Sci. U. S. A.* **2009**, *106*, 18831–18836. (12) Huang, Z.; Roy, P.; Waxman, D. J. Role of human liver microsomal CYP3A4 and CYP2B6 in catalyzing *N*-dechloroethylation of cyclophosphamide and ifosfamide. *Biochem. Pharmacol.* **2000**, *59*, 961–972.

(13) Code, E. L.; Crespi, C. L.; Penman, B. W.; Gonzalez, F. J.; Chang, T. K.; Waxman, D. J. Human cytochrome P4502B6: interindividual hepatic expression, substrate specificity, and role in procarcinogen activation. *Drug Metab. Dispos.* **1997**, *25*, 985–993.

(14) Wang, D.; Li, L.; Yang, H.; Ferguson, S. S.; Baer, M. R.; Gartenhaus, R. B.; Wang, H. The constitutive androstane receptor is a novel therapeutic target facilitating cyclophosphamide-based treatment of hematopoietic malignancies. *Blood* **2013**, *121*, 329–338.

(15) Hedrich, W. D.; Xiao, J.; Heyward, S.; Zhang, Y.; Zhang, J.; Baer, M. R.; Hassan, H. E.; Wang, H. Activation of the constitutive androstane receptor increases the therapeutic index of CHOP in lymphoma treatment. *Mol. Cancer Ther.* **2016**, *15*, 392–401.

(16) Kobayashi, K.; Sueyoshi, T.; Inoue, K.; Moore, R.; Negishi, M. Cytoplasmic accumulation of the nuclear receptor CAR by a tetratricopeptide repeat protein in HepG2 cells. *Mol. Pharmacol.* **2003**, *64*, 1069–1075.

(17) Honkakoski, P.; Negishi, M. Protein serine/threonine phosphatase inhibitors suppress phenobarbital-induced Cyp2b10 gene transcription in mouse primary hepatocytes. *Biochem. J.* **1998**, 330, 889–895.

(18) Swales, K.; Kakizaki, S.; Yamamoto, Y.; Inoue, K.; Kobayashi, K.; Negishi, M. Novel CAR-mediated mechanism for synergistic activation of two distinct elements within the human cytochrome P450 2B6 gene in HepG2 cells. *J. Biol. Chem.* **2005**, *280*, 3458–3466. (19) Mackowiak, B.; Wang, H. 2016. Mechanisms of xenobiotic receptor activation: Direct vs. indirect. *Biochim. Biophys. Acta, Gene Regul. Mech.* **2016**, *1859*, 1130–1140.

(20) Xu, R. X.; Lambert, M. H.; Wisely, B. B.; Warren, E. N.; Weinert, E. E.; Waitt, G. M.; Williams, J. D.; Collins, J. L.; Moore, L. B.; Willson, T. M.; Moore, J. T. A structural basis for constitutive activity in the human CAR/RXRalpha heterodimer. *Mol. Cell* **2004**, *16*, 919–928.

(21) Xie, W.; Barwick, J. L.; Downes, M.; Blumberg, B.; Simon, C. M.; Nelson, M. C.; Neuschwander-Tetri, B. A.; Brunt, E. M.; Guzelian, P. S.; Evans, R. M. Humanized xenobiotic response in mice expressing nuclear receptor SXR. *Nature* **2000**, *406*, 435–439.

(22) Moore, L. B.; Parks, D. J.; Jones, S. A.; Bledsoe, R. K.; Consler, T. G.; Stimmel, J. B.; Goodwin, B.; Liddle, C.; Blanchard, S. G.; Willson, T. M.; Collins, J. L.; Kliewer, S. A. Orphan nuclear receptors constitutive androstane receptor and pregnane X receptor share xenobiotic and steroid ligands. *J. Biol. Chem.* **2000**, *275*, 15122–15127.

(23) Moore, L. B.; Maglich, J. M.; McKee, D. D.; Wisely, B.; Willson, T. M.; Kliewer, S. A.; Lambert, M. H.; Moore, J. T. Pregnane X receptor (PXR), constitutive androstane receptor (CAR), and benzoate X receptor (BXR) define three pharmacologically distinct classes of nuclear receptors. *Mol. Endocrinol.* **2002**, *16*, 977–986.

(24) Willson, T. M.; Kliewer, S. A. PXR, CAR and drug metabolism. *Nat. Rev. Drug Discovery* **2002**, *1*, 259–266.

(25) Cherian, M. T.; Chai, S. C.; Chen, T. Small-molecule modulators of the constitutive androstane receptor. *Expert Opin. Drug Metab. Toxicol.* **2015**, *11*, 1099–1114.

(26) Faucette, S. R.; Zhang, T. C.; Moore, R.; Sueyoshi, T.; Omiecinski, C. J.; LeCluyse, E. L.; Negishi, M.; Wang, H. Relative activation of human pregnane X receptor versus constitutive androstane receptor defines distinct classes of CYP2B6 and CYP3A4 inducers. J. Pharmacol. Exp. Ther. **2006**, 320, 72–80.

(27) Maglich, J. M.; Stoltz, C. M.; Goodwin, B.; Hawkins-Brown, D.; Moore, J. T.; Kliewer, S. A. Nuclear pregnane X receptor and constitutive androstane receptor regulate overlapping but distinct sets of genes involved in xenobiotic detoxification. *Mol. Pharmacol.* **2002**, *62*, 638–646.

(28) Kliewer, S. A.; Goodwin, B.; Willson, T. M. The nuclear pregnane X receptor: a key regulator of xenobiotic metabolism. *Endocr. Rev.* **2002**, *23*, 687–702.

(29) Molnar, F.; Kublbeck, J.; Jyrkkarinne, J.; Prantner, V.; Honkakoski, P. An update on the constitutive androstane receptor (CAR). *Drug Metabol. Drug Interact.* **2013**, *28*, 79–93.

(30) Lynch, C.; Pan, Y.; Li, L.; Ferguson, S. S.; Xia, M.; Swaan, P. W.; Wang, H. Identification of novel activators of constitutive androstane receptor from FDA-approved drugs by integrated computational and biological approaches. *Pharm. Res.* **2013**, *30*, 489–501.

(31) Lynch, C.; Zhao, J.; Huang, R.; Xiao, J.; Li, L.; Heyward, S.; Xia, M.; Wang, H. Quantitative high-throughput identification of drugs as modulators of human constitutive androstane receptor. *Sci. Rep.* **2015**, *5*, 10405.

(32) Lynch, C.; Mackowiak, B.; Huang, R.; Li, L.; Heyward, S.; Sakamuru, S.; Wang, H.; Xia, M. Identification of modulators that activate the constitutive androstane receptor from the Tox21 10K compound library. *Toxicol. Sci.* **2019**, *167*, 282–292.

(33) Smutny, T.; Nova, A.; Drechslerova, M.; Carazo, A.; Hyrsova, L.; Hruskova, Z. R.; Kunes, J.; Pour, M.; Spulak, M.; Pavek, P. 2-(3-Methoxyphenyl)quinazoline derivatives: a new class of direct constitutive androstane receptor (CAR) agonists. *J. Med. Chem.* **2016**, *59*, 4601–4610.

(34) Tschuor, C.; Kachaylo, E.; Limani, P.; Raptis, D. A.; Linecker, M.; Tian, Y.; Herrmann, U.; Grabliauskaite, K.; Weber, A.; Columbano, A.; Graf, R.; Humar, B.; Clavien, P. A. Constitutive androstane receptor (Car)-driven regeneration protects liver from failure following tissue loss. *J. Hepatol.* **2016**, *65*, 66–74.

(35) Shizu, R.; Min, J.; Sobhany, M.; Pedersen, L. C.; Mutoh, S.; Negishi, M. Interaction of the phosphorylated DNA-binding domain in nuclear receptor CAR with its ligand-binding domain regulates CAR activation. *J. Biol. Chem.* **2018**, *293*, 333–344.

(36) Shizu, R.; Osabe, M.; Perera, L.; Moore, R.; Sueyoshi, T.; Negishi, M. Phosphorylated nuclear receptor CAR forms a homodimer to repress its constitutive activity for ligand activation. *Mol. Cell. Biol.* **2017**, *37*, No. e00649-16.

(37) Kandel, B. A.; Thomas, M.; Winter, S.; Damm, G.; Seehofer, D.; Burk, O.; Schwab, M.; Zanger, U. M. Genomewide comparison of the inducible transcriptomes of nuclear receptors CAR, PXR and PPAR α in primary human hepatocytes. *Biochim. Biophys. Acta, Gene Regul. Mech.* **2016**, *1859*, 1218–1227.

(38) Trott, O.; Olson, A. J. AutoDock Vina: improving the speed and accuracy of docking with a new scoring function, efficient optimization and multithreading. *J. Comput. Chem.* **2009**, *31*, 455–461.