

ACCELERATED COMMUNICATION

Di(2-ethylhexyl) phthalate Is a Highly Potent Agonist for the Human Constitutive Androstane Receptor Splice Variant CAR2

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ABSTRACT

The human constitutive androstane receptor (CAR, CAR1) regulates the expression of genes involved in xenobiotic metabolism in the liver. The *CAR* gene uses multiple alternative splicing events during pre-mRNA processing, thereby enhancing the CAR transcriptome. Previous reports have identified two prominent human CAR variants, CAR2 and CAR3, that possess four- and five-amino acid insertions in their ligand binding domains, respectively. Unlike the constitutively active reference form of the receptor, we now demonstrate that CAR2 is a ligand-activated receptor and comprises approximately 30% of the reference transcript level in human liver tissues in human hepatocytes. Furthermore, we identify the common plasticizer, di(2-ethylhexyl) phthalate (DEHP), as a highly potent and uniquely

selective agonist of CAR2. Results from reporter transactivation and mammalian two-hybrid assays reveal that DEHP activates CAR2 at low nanomolar concentrations, results further supported by analysis of CAR target gene expression in primary human hepatocytes. In addition, comparative genomic analyses show that the typical mouse, rat, and marmoset models of DEHP toxicity cannot accurately profile potential human toxicity because of these species' inability to generate a CAR2-like transcript. The discovery that CAR2 is an ultimate human DEHP receptor identifies a novel pathway modulating human DEHP toxicity with potential clinical implications for a subset of patients undergoing critical care medical interventions.

The constitutive androstane receptor (CAR, NR1I3) is a member of the nuclear receptor superfamily that is expressed primarily in the liver (Baes et al., 1994). In conjunction with the pregnane X receptor (NR1I2), CAR regulates hepatic genes involved in all three phases of xenobiotic metabolism

(Wei et al., 2000). The products of these genes play a role in modulating the metabolism of a variety of pharmaceuticals, such as acetaminophen (Zhang et al., 2002) and cyclophosphamide (Roy et al., 1999), as well as endogenous compounds, including thyroid hormone (Maglich et al., 2004), bile acids (Guo et al., 2003), and steroid hormones (Xie et al., 2003).

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In human livers, the *CAR* gene expresses a number of differentially spliced mRNA transcripts, generated through the use of alternative splice acceptor sites during pre-mRNA processing (Auerbach et al., 2003; Savkur et al., 2003; Arnold et al., 2004; Jinno et al., 2004; Lamba et al., 2004). Previous reports from our laboratory have focused on functional analysis of two prominent human CAR variants, termed CAR2 (Auerbach et al., 2007) and CAR3 (Auerbach et al., 2005).

ABBREVIATIONS: CAR, constitutive androstane receptor; CITCO, 6-(4-chlorophenyl)imidazo[2,1-b][1,3]thiazole-5-carbaldehyde *O*-3,4-dichlorobenzyl oxime; DEHP, di(2-ethylhexyl) phthalate; DMSO, dimethyl sulfoxide; RXR, retinoid X receptor; CMV, cytomegalovirus; FBS, fetal bovine serum; SRC1, steroid receptor coactivator 1; MEHP, mono(ethylhexyl) phthalate.

Results from these and other studies demonstrated that both CAR2 and CAR3 activate reporters containing response elements derived from the endogenous promoters of the *CYP2B6* and *CYP3A4* genes (Auerbach et al., 2005, 2007). Unlike CAR1, CAR3 functions as a ligand-dependent receptor (Auerbach et al., 2005), activating transcription in the presence of the human CAR ligand 6-(4-chlorophenyl)imidazo[2,1-*b*] [1,3]thiazole-5-carbaldehyde *O*-3,4-dichlorobenzyl oxime (CITCO) (Maglich et al., 2003). Our recent study of CAR2 suggested that it may have a ligand binding profile that is distinct from both CAR1 and CAR3 (Auerbach et al., 2007).

Based on these data and results from molecular modeling analyses, we hypothesized that CAR2 possesses distinct biological properties. In this report, we demonstrate that CAR2 is a ligand-activated receptor that comprises approximately one third of the total CAR transcriptome in human hepatocytes. Furthermore, we identify the common plasticizer, di(2-ethylhexyl) phthalate (DEHP), as a highly potent and uniquely selective agonist of CAR2. Results from reporter transactivation and mammalian two-hybrid assays reveal that DEHP activates CAR2 at low nanomolar concentrations. These results are further supported by analysis of CAR target gene expression in primary human hepatocytes, where DEHP activation of CAR2 markedly induced the *CYP2B6* and *CYP3A4* transcripts. Furthermore, using genomic analyses, we demonstrate that the alternative splice site used to generate the CAR2 transcript is not conserved in mice, rats, or marmosets—widely used animal models of DEHP toxicity.

DEHP is a plasticizer with an annual production volume approaching 260 million pounds (Kavlock et al., 2006). Human exposure to DEHP is ubiquitous: it is used in the manufacturing of flexible vinyl found in medical tubing, other medical devices, commercial floorings, food product wraps, and many additional consumer products (Hauser and Calafat, 2005). Concerns regarding the human toxicology of DEHP are prevalent; for example, during medical interventions, patients can be exposed to relatively high doses of DEHP that leaches from plastic medical devices into infusate (United States Food and Drug Administration, 2001). Our finding that DEHP is a potent and selective activator of CAR2 is likely to have important implications for predictive human toxicity.

In summary, in this report, we describe how the naturally occurring NR1I3 splice variant, CAR2, is unique in that it 1) is ligand activated, unlike the previously recognized constitutively active reference form of the receptor, 2) comprises ~30% of the reference transcript level in human liver tissues, and, 3) is selectively activated by low nanomolar concentrations of DEHP, levels achieved in human serum and urine of certain patients. Furthermore, these studies yield new mechanistic insight for the toxicological effects of DEHP in human liver, findings that are particularly relevant because the conventional rodent and more recent marmoset models for DEHP toxicity do not reflect that of the human DEHP receptor, CAR2.

Materials and Methods

Chemicals. DEHP, 5 α -androstane-3 α -ol (androstanol), and dimethyl sulfoxide (DMSO) were purchased from Sigma (St. Louis, MO). CITCO

was obtained from BIOMOL Research Laboratories (Plymouth Meeting, PA).

Plasmids. The vectors CMV2-CAR1, CMV2-CAR2, 3.1-RXR α , 3.1-RXR α LBD, VP16-CAR1LBD, VP16-CAR2LBD, 2B6XREM, and 3A4XREM were described previously (Auerbach et al., 2007). The vectors CMV2-CAR3 and VP16-CAR3LBD were also reported previously (Auerbach et al., 2005) as well as the pM-steroid receptor coactivator 1 (SRC1) clone (Chang et al., 1999). All clones were derived from human genes.

Bioinformatics. Protein sequences were retrieved from the NCBI database. Amino acids 212 to 348 of the human CAR1 protein (NCBI RefSeq no. NP_005113.1) were used as the reference sequence and aligned using the blast matrix of the ClustalW2 program from the EBI website (<http://www.ebi.ac.uk/Tools/clustalw2/>). Genomic alignments used the University of California Santa Cruz (UCSC) genome browser (<http://genome.ucsc.edu/>) with the mouse *CAR* gene as the reference sequence.

Cell Culture. Culture conditions for the COS-1 and HepG2 cell lines were described previously (Auerbach et al., 2007), except that the cells were maintained in media containing 8% fetal bovine serum (FBS) and transfected in media containing 8% dextran/charcoal-treated FBS (HyClone, Logan, UT). Culturing conditions for primary human hepatocytes have been described previously (Olsavsky et al., 2007). Available donor information is presented in Table 1. Preparations enriched for hepatocytes were received plated in collagen-coated six-well culture dishes. Upon arrival the media was removed and replaced with 2 ml of fresh media. Matrigel was added along with a 2-ml media change within 48 h of plating. Two-milliliter changes of media occurred once 24 h after the addition of matrigel and then again approximately 48 h later. Cells were harvested 24 h subsequently if no treatment was added. For cells that underwent chemical exposures, media was replaced in the morning (72 h after Matrigel) and then again ~4 h later with fresh media that included the chemical agents. Cells were harvested 24 h after treatments. If not specified otherwise, all culturing materials were obtained from Invitrogen (Carlsbad, CA).

RNA Isolation and Quantitative PCR Analysis. Total RNA was extracted from the hepatocytes using TRIzol Reagent (Invitrogen) according to the manufacturer's protocol. RNA concentrations were assessed by UV absorbance at 260 nm using a DU 800 spectrophotometer (Beckman Coulter, Fullerton, CA) and RNA integrity was ascertained using an Agilent Bioanalyzer (Agilent, Santa Clara, CA). A high-capacity cDNA archive kit (Applied Biosystems, Foster City, CA) was used to reverse transcribe RNA into cDNA, according to the manufacturer's instructions. TaqMan Assays-on-Demand Gene Expression assays (Applied Biosystems) were performed according to manufacturer's directions for a 50- μ l reaction volume subsequently divided into duplicate 25- μ l reactions in a 96-well plate, and run in a real-time PCR system (7300; Applied Biosystems, Foster City, CA). Taqman probes were custom designed to detect the presence of the CAR2 insertion at the exon 6–7 junction. A multiplex reaction was performed containing the REF assay (Applied Biosystems) and CAR2 probe. The REF probe sequence is

TABLE 1
Hepatocyte donor information

Donor	Age	Sex	Race
	<i>years</i>		
A	0.75	Male	White
B	57	Male	White
C	49	Male	White
D	61	Male	White
E	63	Male	White
F	3	Female	White
G	61	Male	White
H	60	Female	Unknown
I	69	Female	White
J	57	Female	White

5'-AGATGGAGCCCGTGTGGGGTTCCAG-3' (5-carboxyfluorescein reporter). The CAR2 probe sequence is 5'-TGTATCTCCCACAGTGGG-3' (VIC Reporter). Absolute quantification was determined through the use of a standard curve generated using plasmid DNA containing either the CAR1 (REF) or CAR2 variant. Six point standard curves were used, encompassing the range from 2 to 6000 fg of target sequence. Applied Biosystems inventoried assays were used for detection of 18S, CYP2B6, and CYP3A4.

Transient Transfection Assays. The details of the luciferase reporter assays were described previously (Auerbach et al., 2007). Likewise, the mammalian two-hybrid assays were conducted using 40 ng of pVP16 expression plasmid, 10 ng of pM (GAL4) expression plasmid, 100 ng of pFR-luc reporter, and 10 ng pRL-CMV (Auerbach et al., 2007). When the pcDNA3.1 expression plasmid, containing RXR-LBD, was incorporated in to the two-hybrid assay, 10 ng was used. The pM-SRC1 (Chang et al., 1999) vector was substituted in place of the pM-SRC1 RID vector in the mammalian two-hybrid experiments.

Statistical Analysis. Unless stated otherwise, statistics, and EC₅₀ values were obtained using Prism (version 4.00; GraphPad Software, San Diego, CA); specific details are provided in the figure legends.

Results

Mice, Rats, and Marmosets Cannot Generate the CAR2 Transcript. Protein sequences were retrieved from

the NCBI database and aligned as described under *Materials and Methods*. Helix-6 and helix-7 of the CAR ligand-binding domain (Xu et al., 2004) are shown (Fig. 1A). The result shows that the rhesus monkey sequence includes a four-amino-acid insertion very similar to CAR2 (shaded boxes) suggesting that the splice variant is conserved across multiple species. To determine whether other species could generate similar transcripts a genomic alignment was performed for three species of rodents and six primates using the UCSC genome browser (Kent et al., 2002), with mouse CAR as the reference sequence (Fig. 1B). Although the splice acceptor site (AG) for CAR1 is conserved across each species, the CAR2 site is not conserved in marmoset, mouse, or rat, indicating that these latter species are incapable of generating a CAR2-like, four-amino acid insertion protein.

The CAR2 Insertion Is Found in ~30.5% of Total CAR Transcripts in Human Hepatocytes. To determine the frequency of the CAR2 insertion we used a TaqMan qPCR approach as described under *Materials and Methods*. The frequency of the CAR2 insertion was found to be very consistent across all donors with a minimum of 25.9%, a maximum of 34.3%, and an average of 30.5% (Fig. 2). These data show that a significant portion of CAR transcripts in hepatocytes contain the CAR2 insertion.

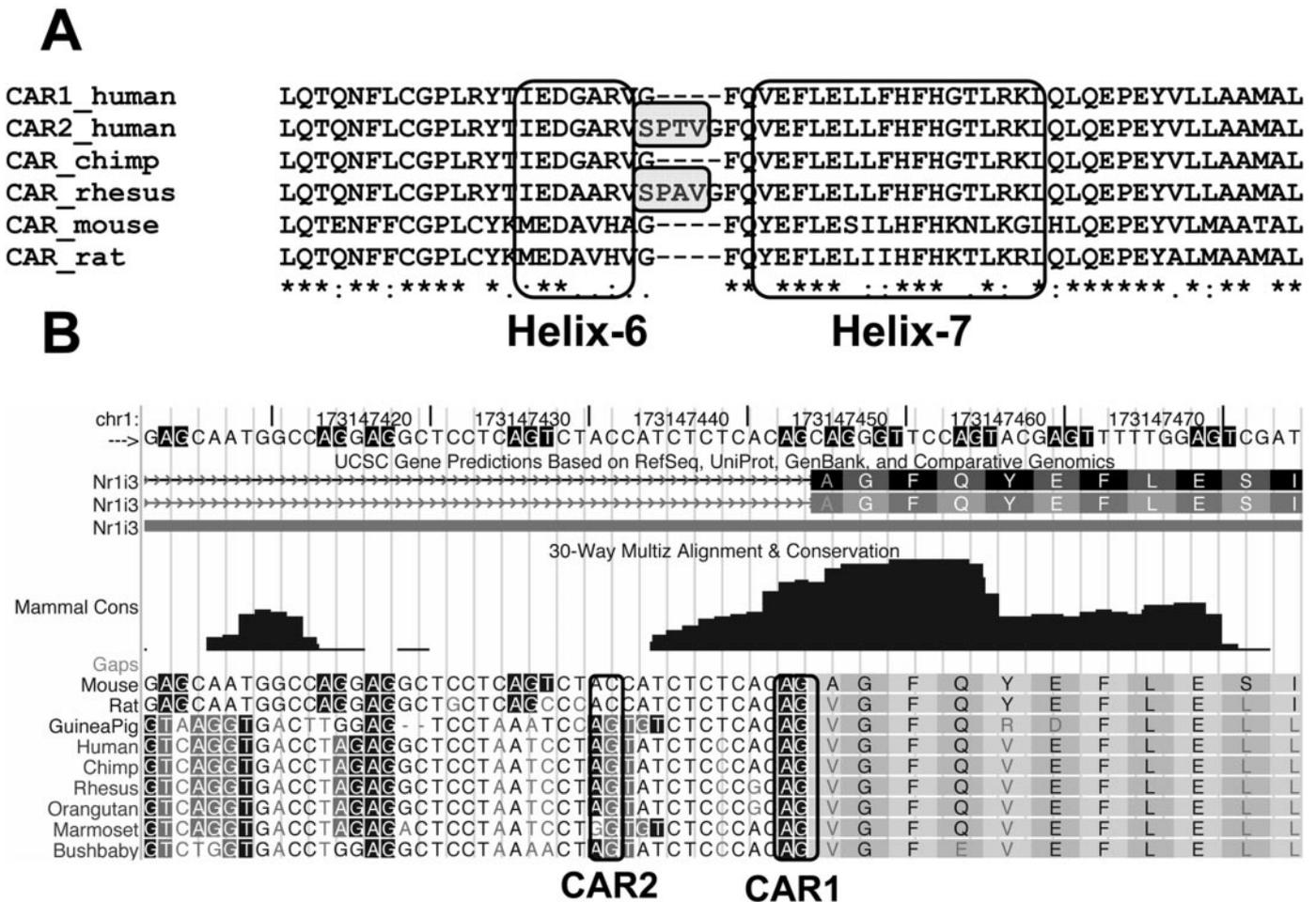


Fig. 1. Rodent and primate alignments of the amino acid and genomic nucleotide sequences flanking the CAR2 insertion. A, the protein alignment of helix-6 and helix-7 of the ligand-binding domain. Protein sequences were retrieved from the NCBI database. Amino acids 212 to 348 of the human CAR1 protein (NP_005113.1) were used as the reference sequence and aligned using the blosum matrix of ClustalW2. B, CAR genomic alignment based on the mouse CAR gene using the UCSC genome browser. The intron 6/exon 7 junction is shown (relative to human CAR). The splice acceptor sites that are used to generate CAR1 or CAR2 are boxed.

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DEHP Is a Contaminant of Fetal Bovine Serum and a Potent Agonist of CAR2. Previous studies conducted in our laboratory used media containing untreated FBS. Results from these studies implied that CAR2 was constitutively active, like CAR1, whereas CAR3 was uniquely ligand-activated (Auerbach et al., 2003, 2005, 2007). Although no previous reports indicated an effect caused by using dextran/charcoal-treated FBS for the study of CAR activity, our initial data suggested that CAR2 may possess a unique ligand-binding profile (Auerbach et al., 2007). To examine this property further, we tested charcoal/dextran-treated FBS to reduce the likelihood of potential background effects that may be contributed from endogenous compounds.

HepG2 cells were transfected in media containing either normal FBS or charcoal/dextran-treated FBS and reporter assays were conducted for activation of the 2B6XREM reporter by the CAR variants (Fig. 3A). In cells that were cultured using media containing the untreated FBS, CAR1, and CAR2 activated the reporter both in the presence and absence of the specific CAR ligand CITCO, whereas CAR3 was ligand-activated as detected previously (Auerbach et al., 2005). We were surprised to find that the use of dextran/charcoal-treated FBS resulted in CAR2 activation only in the presence of CITCO, whereas CAR1 and CAR3 were relatively unaffected by the differences in media formulation. This result suggested that FBS contains a specific CAR2 agonist that is removed by dextran/charcoal treatment.

It should be noted that the data in Fig. 3A were obtained using HepG2 cells, a human hepatoma cell line. We have previously shown that the HepG2 and COS-1 cell lines obtain similar results in our system (Auerbach et al., 2007). In our hands, COS-1 cells generally exhibit a much higher transfection efficiency and consistency than HepG2 cells, therefore further experiments were conducted using the COS-1 cell line.

Using methods to be described elsewhere, we identified DEHP as a contaminant of FBS that is efficiently removed by charcoal/dextran treatment. We assessed the ability of both DEHP and CITCO to activate CAR2 using the 2B6XREM reporter (Fig. 3B). DEHP (Fig. 3C) activated the reporter through CAR2 with an EC_{50} of 211 nM, more than 10-fold greater than the EC_{50} of CITCO, a highly potent CAR ligand.

DEHP Shows Strong Specificity for CAR2 Compared with CAR1 and CAR3. Activity of the CAR variants was assessed by 2B6XREM reporter activity after treatment (Fig. 4A). CAR1 is active under all conditions tested because of its constitutive nature and ligand-independent nuclear accumulation after overexpression in cell lines. Under normal physiologic conditions, CAR1 is sequestered in the cytoplasm and translocates to the nucleus upon activation (Kawamoto et al., 1999). This complex mechanism of CAR1 makes in vitro assessment of ligand activation difficult. The CAR3 insertion is predicted not to significantly alter the ligand binding pocket; therefore, CAR3 is probably an accurate proxy of CAR1 ligand interactions (Faucette et al., 2007). CAR3 shows only a minimal response to DEHP at a dose of 25.2 μ M but is markedly activated by 5 μ M CITCO, as previously reported (Auerbach et al., 2005). CAR2 again displayed a strong dose-response to DEHP; no effect is seen in the empty expression vector (CMV2) control group.

Transactivation of target genes by CAR occurs through the recruitment of coactivators when the receptor is in its active conformation. SRC1 has been shown previously to be recruited by CAR1, CAR2, and CAR3 (Moore et al., 2000; Auerbach et al., 2005, 2007). To assess the interaction of SRC1 with CAR2 and CAR3 in the presence of DEHP, we employed a mammalian two-hybrid assay as described under *Materials and Methods*. Both CAR2 and CAR3 interact with SRC1 in the presence of 5 μ M CITCO (Fig. 4B). CAR2 also shows a dose-dependent interaction with SRC1 in response to DEHP, an effect that is not seen with CAR3. No effect resulted when CAR2 or CAR3 were cotransfected with a pM vector that did not include SRC1 (data not shown).

To better evaluate the interaction between CAR1 and DEHP, we applied the mammalian two-hybrid system used in Fig. 4B to CAR1. The dynamic range of this system is greater than that obtained using the 2B6XREM reporter construct, allowing a more sensitive approach for evaluating physical interactions of chemical entities with the ligand binding pocket of CAR1, as determined by their abilities to compete away the inverse agonist androstanol (Forman et al., 1998). The interaction between the CAR1LBD and SRC1 was measured with and without 10 μ M androstanol. In the presence of 10 μ M androstanol, the reporter activity is ap-

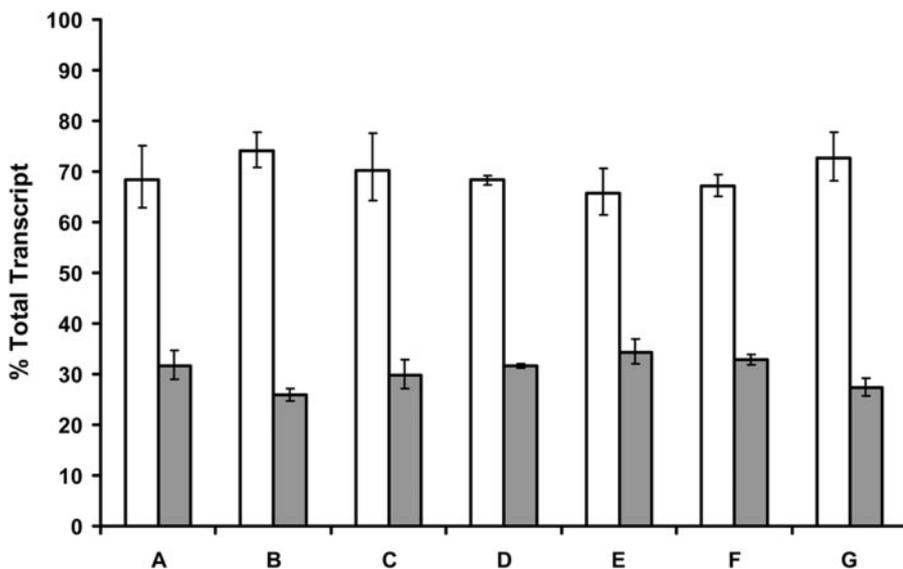


Fig. 2. Frequency of the CAR2 insertion in primary human hepatocytes. TaqMan qPCR was performed on total RNA obtained from seven primary human hepatocyte cultures. Hepatocytes were plated into collagen-coated six-well culture dishes and cultured for approximately six days before harvesting. Quantification of each variant was determined with standard curves generated from plasmid DNA containing either the CAR1 or CAR2 coding sequence. The data are expressed as a percentage of total transcript, defined as the sum of the REF and CAR2 probes divided by the individual probe of interest. The error bars indicate the range based on the standard deviation of the technical replicates using the square root of sum-of-squares technique.

proximately 30% of the constitutive activity seen in the absence of androstanol (Fig. 4C). When CITCO is titrated into the experiment reporter activity is readily restored, with a clearly pronounced effect at 100 nM. On the other hand, DEHP acts as only a weak competitor of androstanol at 10 μ M. Similarly weak competition of androstanol was also observed in separate analysis of mCAR (data not shown). To better illustrate this effect, dose-response curves were generated using the data in Fig. 4C, inset. The result shows an EC_{50} of 571 nM for CITCO and 299 μ M for DEHP. In a previous report, using a CAR-Förster resonance energy transfer assay, the EC_{50} for CITCO was published as 49 nM (Maglich et al., 2003), 8.6% of that determined in our competition study. If the competition among CITCO, androstanol, DEHP, and androstanol is linear, we estimate the EC_{50} of DEHP for CAR1 to be \sim 25 μ M, an estimate that is consistent with the data obtained for the CAR3 variant (Fig. 4A).

DEHP Increases CYP2B6 and CYP3A4 Transcript Levels in Primary Human Hepatocytes. Hepatocytes were isolated from three separate donors and cultured as described under *Materials and Methods*. Cells were treated with 100 nM CITCO or increasing concentrations of DEHP for 24 h and total RNA was isolated for use in qPCR analysis for activation of two CAR target genes, *CYP2B6* and *CYP3A4* (Fig. 5). All three donors showed an increase in CYP2B6 and

CYP3A4 transcripts in response to 50 μ M DEHP. Donor H displayed a strong dose-response to DEHP with an increase in transcript seen at a 10 nM dose for both CYP2B6 and CYP3A4. The other two donors exhibited more moderate responses to DEHP, although the responses from all donors indicated a dose-response enhancement of transcript levels with DEHP. The more modest responses noted for donors I and J probably reflect either interindividual differences in effect, or the apparent higher basal level of expression for both CYP2B6 and CYP3A4 existing before treatments.

Discussion

The report of the crystal structure of CAR1 demonstrated that helix-6 and helix-7 line a large portion of the ligand binding pocket and that a F234A mutation of CAR1 results in loss of the receptor's constitutive activity (Xu et al., 2004). CAR2 contains a four-amino acid insertion that resides in the highly conserved loop between helix-6 and helix-7 of the ligand binding domain, and lies just 1 amino acid upstream of Phe234 (Fig. 1A). These data provide support for a hypothesis that CAR2 lacks constitutive activity and may possess a unique ligand binding profile. We now demonstrate that CAR2, unlike CAR1, is a ligand-dependent receptor and that DEHP is a highly sensitive and selective activator of this

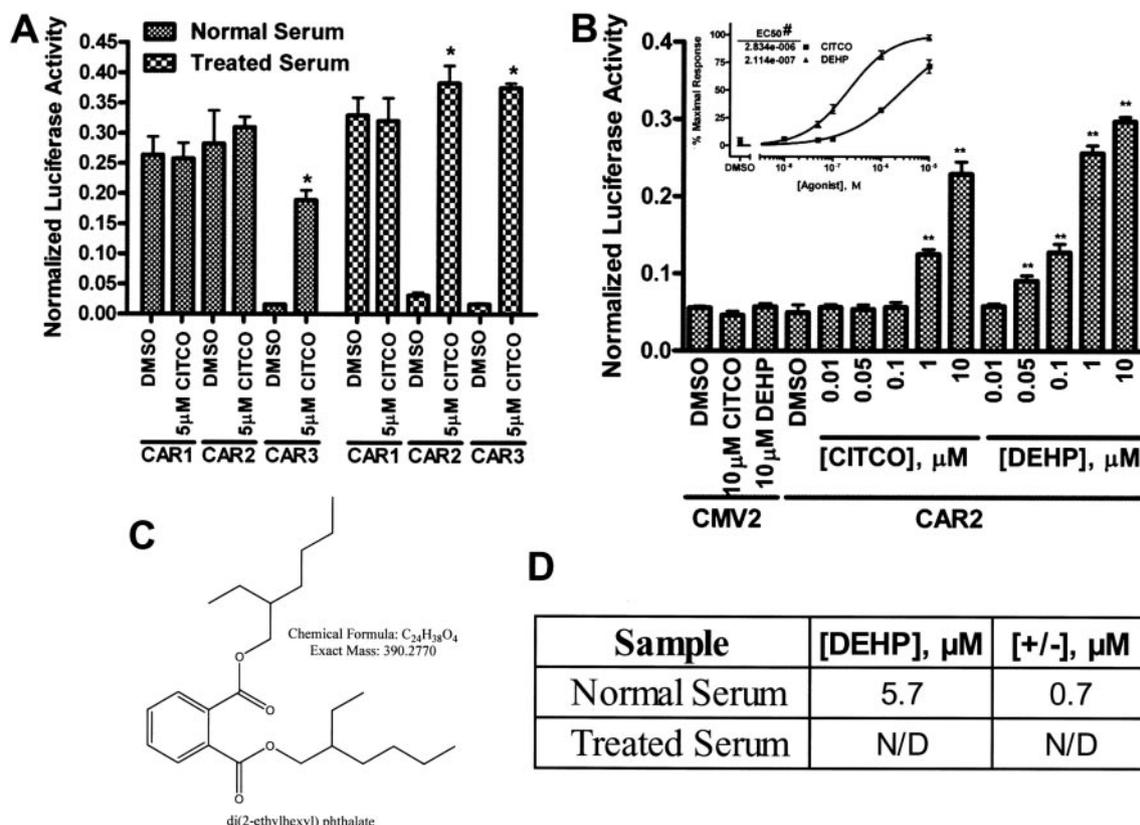


Fig. 3. Differential activation of the CAR variants in response to dextran/charcoal-treated FBS and activation of CAR2 by DEHP. All transfections included the 3.1-RXR α expression vector, the 2B6XREM reporter and the pRL-CMV vector for normalization of transfection efficiency. Data are represented as normalized luciferase values and each data point represents the mean (\pm S.D.) of four separate transfections. A, HepG2 cells were transfected with CMV2-CAR1, CMV2-CAR2, or CMV2-CAR3 in media containing 8% FBS or 8% charcoal/dextran-treated FBS and then treated for 24 h. B, COS-1 cells were transfected with CMV2 or CMV2-CAR2 and treated for 24 h in media containing 8% charcoal/dextran-treated FBS. The inset shows the CAR2 data plotted as a dose-response curve. The DMSO group defined the 0% response and the 10 μ M DEHP group was set to 100% response. C, chemical structure and properties of DEHP. *, data points that were significantly different from their respective DMSO control as determined by unpaired two-tailed *t* test ($p < 0.0001$). **, data points that were significantly different from their respective DMSO control as determined by ANOVA in combination with a Dunnett's multiple comparison post test ($p < 0.01$). #, the difference in the Log EC_{50} values was statistically significant as determined by *F* test ($p < 0.0001$). D, DEHP concentrations ascertained using quantitative mass spectroscopy.

variant CAR. In separate experiments (DeKeyser et al., manuscript in preparation), we demonstrated that no other member of the human NR1I subfamily possess a similar affinity for DEHP, establishing a potency of DEHP for CAR2

that is 40-fold greater than that of any other human NR1I receptor.

As a consequence of their widespread use and environmental prevalence, there has been a great deal of concern noted

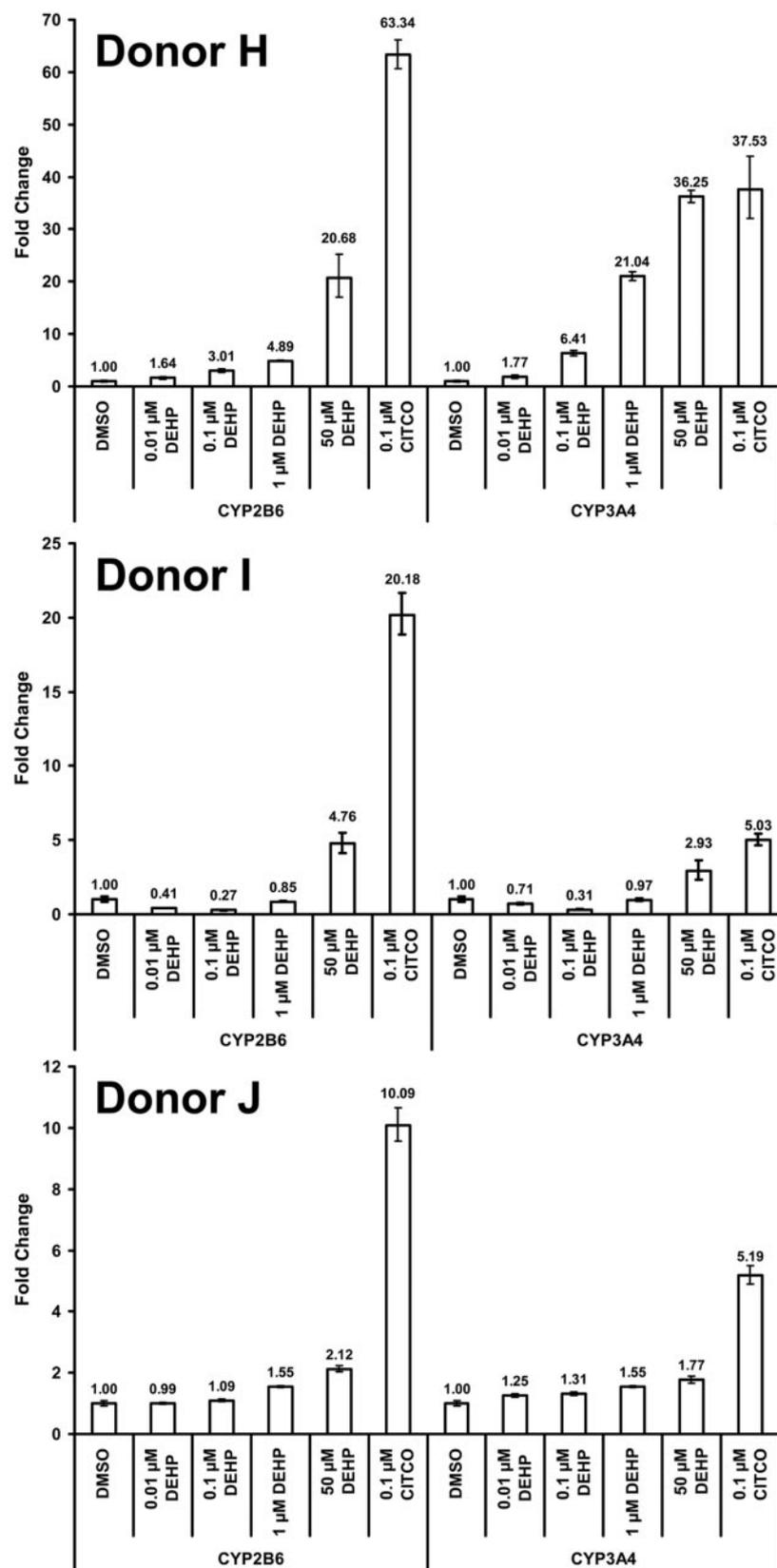


Fig. 5. Relative levels of CYP2B6 and CYP3A4 transcript in primary human hepatocytes after treatment with DEHP or CITCO. Hepatocytes were plated into collagen-coated six-well culture dishes and cultured for approximately 5 days before being treated for 24 h, after which they were harvested for TaqMan qPCR analysis. All data have been normalized to the expression of 18s RNA and are presented as the -fold change in transcript relative to the DMSO-treated group. The error bars indicate the range based on the standard deviation of the technical replicates using the square root of sum-of-squares technique.

for potential human toxicity resulting from exposure to phthalates, concerns that have been focused primarily on reproductive toxicity, stemming from the antiandrogenic properties of DEHP. A recent report from the National Toxicology Program Center for the Evaluation of Risks to Human Reproduction (NTP-CERHR) underscores the issues over potential adverse effects of DEHP on the developing male reproductive tract, with infants, particularly those who have faced critical illness with associated medical device interventions, at the greatest risk (Kavlock et al., 2006). It has been suggested that blood phthalate levels in patients undergoing critical care procedures may approach micromolar concentrations (McKee, 2004), with DEHP levels of 1.19 $\mu\text{g/ml}$ reported in cord blood (Latini et al., 2003). DEHP exposure in male subjects may compromise sperm quality and quantity (Hoppin, 2003; Duty et al., 2005; Kavlock et al., 2006) and disrupt circulating levels of thyroid hormone (Meeker et al., 2007). It is noteworthy that the metabolism of thyroid hormone is enhanced through CAR regulation (Maglich et al., 2004). Furthermore, the human CYP2B6 and CYP3A4 enzymes metabolize testosterone (Imaoka et al., 1996). It is interesting to speculate that phthalate levels in certain tissues and perhaps hepatocytes may actually exceed circulating blood levels, implying that CAR2 may be activated by DEHP at even lower blood concentrations than achieved in critical care interventions, and that CAR1 or CAR3 may additionally take on a target role under such circumstances.

Mono(ethylhexyl) phthalate (MEHP), a major metabolite of DEHP, has been implicated as a mediator of the reproductive and developmental toxicity associated with DEHP (Heindel and Powell, 1992; Kavlock et al., 2006). We tested MEHP as a CAR2 activator, but even at 10 μM , this agent exhibited only weak activity in our in vitro reporter assays (data not shown), results that further support our conclusion that the DEHP parent compound itself is the most active modulator of CAR2. It is noteworthy that most of the previous conclusions regarding phthalate toxicology (Kavlock et al., 2006) have been drawn from high-dose studies in rodent or marmoset models, species that are not capable of generating a CAR2-like transcript (Fig. 1B). Furthermore, it has been suggested that DEHP is metabolized to MEHP to a lesser degree in primates compared with rodents (Ito et al., 2005; Rusyn et al., 2006). In light of these considerations, the potential role of CAR2 as a mediator of the developmental and reproductive toxicities associated with DEHP in humans warrants further investigation.

In this report, we demonstrate that the CAR2 transcript comprises ~30% of the CAR reference level in human hepatocytes (Fig. 2), and that exposure to DEHP induces transcription of the CAR target genes, *CYP2B6* and *CYP3A4*. In particular, one donor exhibited a DEHP response at a dose that is likely to activate CAR2 exclusively (Fig. 5). The discovery that low levels of DEHP can induce xenobiotic metabolism in human hepatocytes may have immediate clinical relevance for some patients, especially those receiving large amounts of intravenous fluids or blood transfusions, as well as medications that have contraindications associated with CAR target genes.

Furthermore, the results of our study strongly imply that CAR2 has the capacity to interact with a set of ligands that are highly distinct from that of other nuclear receptors. This finding will probably serve to redirect future evaluation of

drug and chemical safety data and should set new directions underscoring the importance of CAR in human biology, including for the emerging role of CAR as a regulator of hepatic energy and lipid metabolism (Konno et al., 2008; Masson et al., 2008).

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