

## INDUCTION OF HEPATIC PHASE II DRUG-METABOLIZING ENZYMES BY 1,7-PHENANTHROLINE IN RATS IS ACCOMPANIED BY INDUCTION OF MRP3

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### ABSTRACT:

The purpose of the present study was to evaluate the effect of 1,7-phenanthroline (PH), which has been proposed to be a selective phase II enzyme inducer, on the gene expression of xenobiotic transporters, as well as hepatic and renal drug-metabolizing enzymes. After oral administration of PH for 3 days to male Sprague-Dawley rats, mRNA levels in liver (75 and 150 mg/kg doses) and kidney (75 mg/kg dose only) were determined using real-time quantitative polymerase chain reaction. At 150 mg/kg/day, PH treatment resulted in significant increases in hepatic mRNA levels of *Mrp3* (36-fold), *UGT1A6* (20-fold), *UGT2B1* (4-fold), and quinone reductase (*QR*, 5-fold), compared with the vehicle-treated group.

Similar increases in *Mrp3* (99-fold), *UGT1A6* (17-fold), *UGT2B1* (3-fold), and *QR* (11-fold) mRNA levels were observed in the liver after PH treatment of rats at 75 mg/kg/day. In contrast, the expression levels of *CYP2C11* and *Oatp2* were decreased by ~80 and 50%, respectively. In addition, PH (75 mg/kg/day) elicited statistically significant changes in renal gene expression of *CYP3A1*, *UGT1A6*, *QR*, and *Mrp3*, but the magnitude of renal *Mrp3* induction was less than 2-fold over control. Although PH is known to modulate hepatic glucuronidation in vivo, these data indicated that PH induced mRNA levels of the efflux transporter, *Mrp3*, which may also affect the disposition of xenobiotics.

Multidrug resistance proteins 1, 2, and 3 (*Mrp1*,<sup>1</sup> 2, and 3) have gained significance during the last few years because of their function as transporters of organic anions and conjugates, and their involvement in hepatic detoxification and tissue-specific distribution of drugs (Konig et al., 1999; Hirohashi et al., 1999; Borst et al., 2000). Notably, induction of hepatic rat *Mrp3* and another transporter, *Oatp2*, by drugs like phenobarbital was implicated in the altered disposition of acetaminophen and the enhanced uptake of digoxin in rats (Rausch-Derra et al., 2001; Xiong et al., 2002a). *Mrp3* induction has been shown also with oltipraz (Cherrington et al., 2002), a dithiolthione phase II enzyme inducer being investigated as a chemoprotectant against aflatoxin carcinogenicity and hepatotoxicity (Kwak et al., 2001a). Oltipraz and other selective phase II inducers, such as 1,7-phenanthroline (PH), offer a new mechanism of protection against carcinogenic and hepatotoxic compounds by inducing the enzymes involved in their metabolism [glutathione *S*-transferase (GST) and UDP-glucuronosyltransferase (UGT)], without increasing cytochrome P450-mediated bioactivation (Franklin and Moody, 1992; Franklin et al., 1993; Buetler et al., 1995; Vargas et al., 1998; Dong et al., 1999; Lamb and Franklin, 2000). Additionally, selective phase II inducers can be used to modulate in vivo exposure to acyl glucuronides. In this regard, a significant increase in the glucuronidation of benoxaprofen,

a nonsteroidal anti-inflammatory drug, was observed when rats were treated with PH at 75 mg/kg/day for 3 days, resulting in a 2-fold increase in the biliary excretion of the acyl glucuronide, and an 8-fold increase in the peak plasma concentration ( $C_{max}$ ) and area under the curve values of the acyl glucuronide (Dong et al., 1999).

In the present study, we report that PH significantly induced not only mRNA levels of hepatic phase II enzymes, but also mRNA levels of *Mrp3*, which is likely to contribute to the altered metabolism and disposition of xenobiotics and their metabolites in PH-treated rats.

### Materials and Methods

**Materials.** PH was purchased from Sigma-Aldrich (St. Louis, MO). The purity, verified by high-pressure liquid chromatography, was 99.5%. Solvents used for analysis were of analytical or high-pressure liquid chromatography grade (Fisher Scientific, Pittsburgh, PA).

**In Vivo Animal Studies.** All studies were reviewed and approved by the Merck Research Laboratories Institutional Animal Care and Use Committee. Male Sprague-Dawley rats were obtained from Charles River Laboratories Inc. (Wilmington, MA). After an overnight fast, five male Sprague-Dawley rats were dosed orally with PH at 150 mg/kg once daily for 3 days, and four male Sprague-Dawley rats were dosed with vehicle (0.1 M citric acid in 0.5% methylcellulose) in the same manner. Livers were removed from treated and vehicle control rats at 24 h after the last dose and stored at  $-70^{\circ}\text{C}$  for quantitation of mRNA. In a separate experiment, three rats were dosed with PH at 75 mg/kg/day for 3 days, and an additional three rats were dosed with vehicle in the same manner. At 24 h after the last dose, the livers and kidneys were removed and stored at  $-70^{\circ}\text{C}$  for mRNA quantification.

**Development of Specific Primers and Probes for Quantitative Real-Time Polymerase Chain Reaction.** Coding sequences for the genes listed in Table 1 were accessed from GenBank. Specific target regions within the coding sequences were determined through nucleotide sequence alignment comparisons of targets within multiple member gene families (e.g., *Oatp2* with *Oatps*, *Mrp2* with *Mrps*, *CYP2C11* with *CYP2C12*, etc.). Primers and probes

<sup>1</sup> Abbreviations used are: *Mrp*, multidrug resistance protein; *Oatp*, organic anion transporting polypeptide; PH, 1,7-phenanthroline; GST, glutathione *S*-transferase; UGT, UDP-glucuronosyltransferase; GAPDH, glyceraldehyde-3-phosphate dehydrogenase; Ct, cycle threshold; QR, quinone reductase; Nrf2, nuclear factor-erythroid 2-related factor.

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TABLE 1  
Rat primer-probe sets and gene abbreviations

Gene	GenBank	Forward Primer	Reverse Primer	Probe
<i>CYP3A1</i>	D13912	TGGTCATATTCATCTTATGCTC	GGCGAAATTCCTCAGGCTCT	TCACCGTGATCCACA
<i>CYP2C11</i>	NM_010184	CACCAGCTATCAGTGGATTTGG	GTCTGCCCTTTGACACAGGAA	CATTGCCCCCTTTTACGAGGCTTG
<i>GSTA2</i>	NM_01713	CGCTAGACAGTGAAGCAGTTTG	CCGGATGCACTCCATTCTGT	CCAGTGCTTCACTACTTCAATGCCCGG
<i>Mrp2</i>	X96393	TGCAGCCTCCATAACCATGAG	GATGCCTGCCATTGGACCTA	AGAGAGAACAGCTTTCGTCGAACACTTAGCC
<i>Mrp3</i>	AF072816	CCAACCCATGAACCCCAA	GCACATCTAGGTGACGTAGCAGG	CGCTACCAACAGGCTCTGGAGACGTGT
<i>Oatp2</i>	U95011	GTACTCAAGATTTCCAGGGAAA	AACATGCTGTTGAAAATGAAGG	CTTCTATAGTGGCCTTACCCTTCACCTTAAAGC
<i>QR</i>	NM_017000	CTCGCCTCATGCGTTTTTG	CCCCTAATCTGACCTCGTTTAT	TGCCACGGCTGTGACAACAA
<i>UGT1A1</i>	D38065	CCGTTGTCTACGTGCCC	TCACCCGCTGCAGGAAGT	AGAGTTTGTCTCGAACACAGATCGCATG
<i>UGT1A6</i>	NM_057105	CCGCTATCGCTCCTTTGG	CTGTACTCTTTAGAGGAGCCATCAG	AACAACCACTTTGCTGCTGCCAGTTCCCC
<i>UGT2B1</i>	M13506	GCTGCTTCCAGGAACCTG	TGAGGTCCCAACGCTGTCTT	TGTCATGCCAGTGCCCTTCCCTCTAAAA

were designed to the selected target using Applied Biosystems Inc. (Foster City, CA) Primer Express software (v.2.0). All primers and probes were submitted to the National Center for Biotechnological Information for nucleotide comparison using the basic logarithmic alignment search tool (BLASTn) search for short, nearly exact sequences to ensure specificity. Primers and probes were synthesized by QIAGEN Operon (Alameda, CA), where primers were 5'- and 3'-labeled with the 6-carboxyfluorescein and 6-carboxytetramethylrhodamine reporter dyes, respectively. The rodent glyceraldehyde-3-phosphate dehydrogenase (*GAPDH*) primer/probe set was purchased from Applied Biosystems Inc. and used per manufacturer's instructions. Each RNA sample was reverse-transcribed before analysis of different gene expression by PCR. Real-time quantitative PCR was performed using an ABI PRISM 7700 Sequence Detector instrument and Sequence Detector v.1.7 software (PerkinElmer Instruments, Skelton, CT).

**mRNA Isolation and Quantitative Real-Time Polymerase Chain Reaction.** Total RNA from rat tissues was isolated using the SV Total RNA Isolation System (Promega, Madison, WI) according to the manufacturer's instructions. Samples were quantitated by spectrophotometry and diluted to a concentration of 15 ng/ $\mu$ l. Aliquots (500 ng) of RNA were analyzed by agarose/formaldehyde gel electrophoresis to check RNA integrity. Samples were then assayed in triplicate 25- $\mu$ l reactions using 25 ng of RNA per reaction. Gene-specific primers were used at 7.5 pmol per reaction, and the gene-specific probes were used at 5 pmol per reaction. *GAPDH* was used to normalize gene expression in all samples since it is a highly expressed gene in rat liver and did not change in response to PH treatment [cycle threshold (Ct), control,  $21.8 \pm 0.2$ ; PH,  $22 \pm 0.3$ ]. Fold induction values were calculated by subtracting the mean difference of gene and *GAPDH* Ct number for each treatment group from the mean difference of gene and *GAPDH* Ct number for the vehicle group and raising this difference to the power of 2.

Statistical analyses were performed using a two-tailed Student's *t* test at  $\alpha = 0.01$  level of significance.

## Results and Discussion

As expected from the literature (Vargas et al., 1998; Lamb and Franklin, 2000), significant increases in the amount of mRNA for *UGT1A6* (20-fold) and, to a lesser extent, *UGT2B1* (4-fold) and *QR* (5-fold) were detected in rat liver in response to PH treatment of 150 mg/kg/day (Fig. 1). The mRNA levels of *UGT1A1* were not affected substantially (<2-fold) by this treatment. Similar increases in *UGT1A6* (17-fold), *UGT2B1* (3-fold), *GST* (6-fold), and *QR* (11-fold) mRNA were observed in liver after PH treatment of rats at 75 mg/kg/day for 3 days (Table 2). These results were similar to those reported by Vargas et al. (1998). The observed lack of dose proportionality in the increase of the mRNA of these enzymes between 75 and 150 mg/kg of PH needs further investigation to establish the time-dependence of the induction.

The transcription factor *Nrf2* has been shown to be important for the induction of phase II enzymes. Induction of *QR* and *UGT1A6* by oltipraz was observed in the wild-type, but not *nrf2*-deficient mice, which suggested that *Nrf2* plays a major role in the regulation of these genes (Kwak et al., 2001b). Given that PH causes induction of a

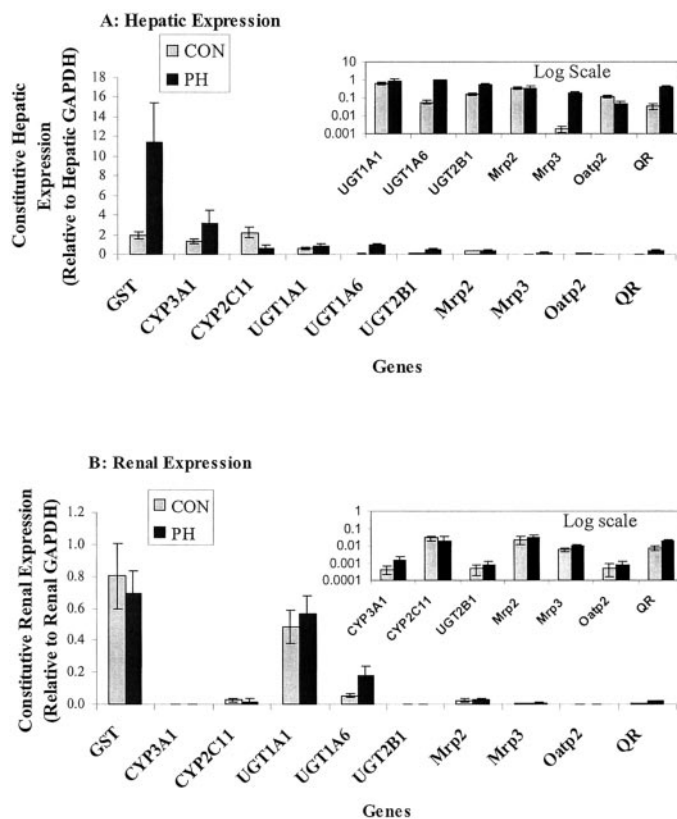


Fig. 1. Comparison of constitutive (gray) and 1,7-phenanthroline-inducible (75 mg/kg/day for 3 days; black) expression of hepatic (A) and renal (B) drug-metabolizing enzymes and xenobiotic transporters.

Three animals were in each group; three determinations were performed for each animal. Some of the gene expression levels in log scale are shown in the box on the right in the figure.

battery of genes similar to that of oltipraz, further studies are needed to elucidate the role of *Nrf2*, if any, in the inductive effects of PH.

PH caused a 3- to 4-fold induction of *CYP3A1* gene expression in both liver and kidney; however, this is considered minor in comparison to the induction of *CYP3A1* gene expression by pregnane X receptor agonists (~30-fold) (Hartley and Klaassen, 2000) and may not be associated with increased protein levels (Dong et al., 1999). Significant decreases in the mRNA levels of *CYP2C11* (17% of control) and *Oatp2* (50% of control) were observed by PH treatment at 150 mg/kg/day. Down-regulation of *CYP2C11* and *Oatp2* by some aryl hydrocarbon receptor agonists, such as 3-methylcholanthrene, has been reported previously (Lee and Riddick, 2000; Rausch-Derra et al., 2001; Guo et al., 2002). However, PH does not induce *CYP1A2* activity (Dong et al., 1999), suggesting that aryl hydrocarbon receptor

TABLE 2

Effects of 1,7-phenanthroline (75 mg/kg/day and 150 mg/kg/day, 3 days) on hepatic and kidney gene expression in male Sprague-Dawley rats

Data represent mean  $\pm$  S.D.;  $n = 5$  for the 150 mg/kg treatment group and  $n = 3$  for the 75 mg/kg group. Three determinations were performed for each animal. Statistically significant differences between treated and control groups were evaluated by the Student's *t* test.

Gene	Liver		Kidney (75 mg/kg treatment)
	150 mg/kg treatment	75 mg/kg treatment	
	fold induction over control		
<i>CYP3A1</i>	3.4 $\pm$ 0.5**	2.3 $\pm$ 0.7**	3.7 $\pm$ 1.5**
<i>CYP2C11</i>	0.2 $\pm$ 0.03**	0.3 $\pm$ 0.1**	0.7 $\pm$ 0.4
<i>UGT1A1</i>	0.7 $\pm$ 0.07**	1.4 $\pm$ 0.2**	1.2 $\pm$ 0.1
<i>UGT1A6</i>	20 $\pm$ 1.7**	17 $\pm$ 0.8**	3.2 $\pm$ 0.6**
<i>UGT2B1</i>	3.7 $\pm$ 0.6**	3.3 $\pm$ 0.3**	1.6 $\pm$ 0.3
<i>Mrp2</i>	1.2 $\pm$ 0.1	1.0 $\pm$ 0.2	1.4 $\pm$ 0.1
<i>Mrp3</i>	36 $\pm$ 4.9**	99 $\pm$ 13**	1.6 $\pm$ 0.2**
<i>Oatp2</i>	0.5 $\pm$ 0.1**	0.4 $\pm$ 0.1**	1.1 $\pm$ 0.4
<i>GSTA2</i>	N.D.	5.9 $\pm$ 1.3**	0.9 $\pm$ 0.1
<i>QR</i>	4.8 $\pm$ 1.1**	11 $\pm$ 1.9**	2.5 $\pm$ 0.3**

N.D., not determined.

\*\* Significantly different ( $P < 0.01$ ) from controls.

may not be involved in the PH down-regulation of *CYP2C11* and *Oatp2*.

In addition to the increases in *QR* and *UGT* mRNA levels, which are consistent with the previous report (Vargas et al., 1998), we also demonstrated that PH markedly increased hepatic *Mrp3* mRNA levels (35-fold at 150 mg/kg and 99-fold at 75 mg/kg), but *Mrp2* levels remained unaffected. *Mrp3* is located on the basolateral membrane of polarized cells, and it plays a role in the hepatic elimination by transporting organic anions from liver to blood, which could lead to increased levels of xenobiotics and their metabolites in plasma and possibly an increase in urinary excretion (Konig et al., 1999; Kool et al., 1999). In normal rat liver, constitutive expression of *Mrp3* mRNA is very low; however, the inducible nature of *Mrp3* is well documented (Ogawa et al., 2000; Cherrington et al., 2002). *Mrp3* has been shown to be induced by activators of the constitutive androstane receptor and an antioxidant/electrophile responsive element (Cherrington et al., 2002). However, a recent report by Xiong et al. (2002b) suggests that *Mrp3* regulation occurs independent of constitutive androstane receptor.

In contrast to the robust increase in gene expression levels for *UGT1A6* and *Mrp3* in liver of PH-treated rats, there were much smaller changes in these genes in kidney by PH treatment at 75 mg/kg for 3 days (Table 2). The lack of evidence for a robust renal induction of xenobiotic transporters in rats treated with known hepatic enzyme inducers has been reported previously (Brady et al., 2002; Cherrington et al., 2002). In addition, previous studies have shown minimal inductive effects on drug-metabolizing genes by PH in the small intestine (Vargas et al., 1998).

Thus, PH is a pleiotropic inducer of genes responsible for drug metabolism and transport. These results indicate that in vivo data from PH-induced rats should be interpreted with caution, since up-regula-

tion of *Mrp3* gene expression may result in increased efflux of glucuronides and other xenobiotics from liver into the plasma compartment (Gregus et al., 1990; Xiong et al., 2002a). Further studies are needed to investigate the mechanism by which PH induces *Mrp3* and phase II enzymes, and to determine whether these changes in mRNA levels will correlate with protein levels and/or activity.

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