



Regulation of Gene Expression of Various Phase I and Phase II Drug-Metabolizing Enzymes by Tamoxifen in Rat Liver

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ABSTRACT. The objective of the present investigation was to evaluate the effect of tamoxifen (TAM) on the gene expression of different phase I and phase II drug-metabolizing enzymes. Groups of male and female F344/NCr rats were administered either corn oil or TAM (2.8 to 45 mg/kg body wt \times 14 days) dissolved in corn oil by gavage. An additional group of rats received a diet supplemented with phenobarbital (PB, 500 ppm). Northern blot analyses of total liver RNA were conducted using [32 P]-labeled cDNA or oligonucleotide probes coding for different sulfotransferase (ST), UDP-glucuronosyltransferase (UGT), glutathione S-transferase (GST), epoxide hydrolase (EPH) or cytochrome P450 (CYP) mRNA transcripts. In male rats, TAM increased the levels of ST_{el}, ST_a and ST_{pl} mRNAs, whereas PB increased only the ST_{el} mRNA. In female rats, there was no expression of ST_{el} and ST_{HA} mRNA in either control or TAM-treated animals. TAM and PB increased UGT_{br/p} mRNAs in all rats, whereas UGT_{ml} mRNA was elevated only in PB-treated animals. EPH mRNA was elevated markedly in all rats treated with TAM and PB, whereas GST_{ya/yc} mRNA was highly increased by PB, but only marginally increased by TAM. Finally, TAM increased CYP3A1 mRNA, and slightly increased CYP2B1 mRNA, whereas PB highly elevated mRNAs for both of these CYP genes. In conclusion, treatments of rats with TAM increased the mRNA levels of many phase I and phase II drug-metabolizing enzymes, and this pleiotypic response to TAM seems to be different from other prototype inducers such as PB or dioxin (TCDD). *BIOCHEM PHARMACOL* 52;10:1561–1568, 1996. Copyright © 1996 Elsevier Science Inc.

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The anti-estrogen drug TAM^{||} is presently the most widely used therapeutic agent for the treatment of breast cancer in post-menopausal women [1, 2]. Recently, TAM has shown a great deal of promise for use as a chemopreventive agent [3–6], and large-scale clinical trials are currently underway to evaluate its efficacy as a prophylactic for breast cancer [7, 8].

TAM is presumed to exert its antitumor and chemoprotective effects by competitively antagonizing the binding of estradiol to estrogen receptors, which ultimately leads to the inhibition of gene transcription and protein synthesis [9]. However, recent evidence suggests that the effects of TAM may additionally involve mechanisms that are unrelated to its abilities to interact with the estrogen receptor [10, 11]. Presently, there are a number of questions remaining with respect to the ability of TAM to influence various phase I and phase II drug-metabolizing enzymes. First, it is possible that some portion of the pharmacological effects of TAM may be mediated, in part, via the induction of phase I and phase II drug-metabolizing enzymes. Previous reports have shown that induction of phase II detoxifying enzymes by certain xenobiotics, i.e. phenolic antioxidants, confers protection against chemically induced carcinogenesis [12–14]. In addition, it has been shown that induction of both phase I and phase II drug-metabolizing enzymes by com-

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^{||} Abbreviations: TAM, tamoxifen; PB, phenobarbital; ST, sulfotransferase; UGT, UDP-glucuronosyltransferase; GST, glutathione S-transferase; EPH, epoxide hydrolase; CYP, cytochrome P450; and SSC, 0.15 M sodium chloride + 0.015 M sodium citrate.

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pounds such as indole-3-carbinol sufficiently alters endogenous hormone metabolism to inhibit "spontaneous" mammary carcinogenesis in mice [15] or tumor "promotion and progression" in an *N*-methyl-*N*-nitrosourea (NMU) induced rat mammary model that does not require carcinogen activation [16]. Second, TAM has proven to be a potent liver carcinogen in both rats and hamsters [17]. This effect is presumably attributable, at least in part, to the fact that TAM can readily form DNA adducts in the livers of treated rodents [18, 19]. However, prior reports of both the promoting effects of TAM [20] and its ability to induce CYP2B implied that some portion of its carcinogenic activity may be related to its abilities to induce phase I drug-metabolizing enzymes. Finally, the common use of TAM in humans opens the possibility of certain drug-drug interactions that might either enhance or diminish its biologic activity. In view of these facts, the primary objectives of the present investigation were to examine the effects of TAM on the regulation of gene expression of various phase I and phase II drug-metabolizing enzymes.

In the present report, we focused specifically on the mRNA expression of different STs, UGTs, GST and microsomal EPH isoenzymes, as well as certain major inducible forms of CYPs. ST, UGT, GST, and EPH belong to large multigene families of metabolizing enzymes that are distributed in many organs throughout the body including the liver, lung, and kidney. These enzymes catalyze the metabolism of a number of xenobiotics and endogenous compounds. The STs, for example, are responsible for sulfoconjugating compounds such as biogenic amines (i.e. norepinephrine), steroid hormones (i.e. β -estradiol, dehydroepiandrosterone), bile acids (i.e. hyodeoxycholic acid), drugs (i.e. acetaminophen, propranolol) and carcinogens (i.e. *N*-hydroxy-2-acetylaminofluorene) [21]. UGTs conjugate compounds such as bilirubin, bile acids (i.e. hyodeoxycholic acid), drugs (i.e. acetaminophen, morphine) and carcinogens (i.e. *N*-hydroxy-2-acetylaminofluorene) [22]. The GSTs play an important role in the conjugation of numerous electrophilic drugs (i.e. melphalan, ethacrynic acid) and carcinogens (i.e. 4-nitropyridine-*N*-oxide, benzo[*a*]pyrene, aflatoxin B) [23], whereas EPH catalyzes the conversion of arene and alkene oxides to vicinal dihydrodiols [24]. Conjugation by these phase II enzymes increases the polarity and water solubility of the parent compounds, which facilitates biliary and/or urinary excretion and detoxification. Finally, we examined the effect of TAM on certain of the major inducible families of cytochrome P450 genes (CYP2B and CYP3A) that are involved with the metabolism of various endogenous ligands as well as activation and detoxification of various carcinogens.

MATERIALS AND METHODS

Chemicals

TAM and the free acid of PB were purchased from the Sigma Chemical Co. (St. Louis, MO). All other chemicals were of reagent grade and were purchased from the Fisher Scientific Co. (Pittsburgh, PA).

Animals and Treatments

Male and female F344/NCr rats were obtained at 6 weeks of age from the Animal Production Area, Frederick Cancer Research and Development Center (FCRDC, Frederick, MD). The animals were maintained on hardwood chips at 68–72°F and ~50% humidity in an American Association for Laboratory Animal Care-certified laboratory. The experimental protocols were reviewed and approved by the FCRDC Animal Care and Use Committee, in accordance with the Declaration of Helsinki and the NIH Guide for the Care and Use of Laboratory Animals. After a 2-week acclimation period, rats of the same sex were randomized by body weight into control and TAM-treated groups (N = 3 rats/group). Rats in the control group were maintained on a standard diet of Purina Lab Chow (No. 5010) and administered corn oil (0.4 mL/day) via oral gavage as a vehicle control. TAM-treated groups received TAM dissolved in corn oil at doses of 2.8 (low dose), 11 (medium dose), or 45 mg/kg body weight (high dose). An additional group of rats receiving the control diet supplemented with PB (500 ppm) served as the positive control for the experiment. This dose of PB would yield a daily dose of about 40 mg/kg body weight. Animals in all treatment groups were allowed free access to food and water throughout the experiment. After 14 days of treatment, individual body weights were recorded, and the rats were killed between 9:00 and 10:30 a.m. by CO₂ asphyxiation. Immediately thereafter, the livers were removed, trimmed free of extraneous tissue, and stored at -70° pending analysis of mRNA.

Isolation of Total RNA and Northern Blot Analysis

Total cellular RNA was isolated from a pooled sample of about 0.7 g (equal amount of liver tissue from each animal per treatment group) by the guanidine-thiocyanate method as described by Chomczynski and Sacchi [25] using the RNeasy Total RNA Isolation System (Promega, Madison, WI). For studies on the phase II genes (STs, UGTs, GST and EPH) 15 μ g of total RNA was electrophoretically separated on 1% agarose gels containing 5% formaldehyde and transferred onto HybondTM.N nylon membranes (Amersham, Arlington Heights, IL). The blot was prehybridized in 48% formamide, 5x SSC, 20 mM Tris-HCl (pH 7.6), 0.1% SDS and 1x Denhardt's solution at 42° for 2 hr. After prehybridization, the blot was incubated overnight at 42° in fresh prehybridization solution containing 100 μ g/mL sheared salmon sperm DNA and eight different ³²P-labeled cDNA probes coding for various ST, UGT and EPH mRNA transcripts. The specific cDNA probes utilized

¶ ST(s) is used to denote sulfotransferase protein(s) and italic ST with subscript, e.g. ST_{pl}, indicates the gene that encodes the protein. The names of individual enzymes/genes are abbreviated ST_a (alcohol/hydroxysteroid ST), ST_{pl} (phenol ST), ST_{HA} (*N*-hydroxyarylamine ST), ST_{et} (estrogen ST), UGT_{B₁/p} (family 1-bilirubin/phenol UGT), and UGT_{mt} (family 2-bile acid/hydroxysteroid UGT).

were: rST_{el} [26], rST_{HA} [27], rST_a [28], mST_{pl} [29], $mUGT_{Br/p}$ [30], $mUGT_{ml}$ [31] and $pEPH_{302}$ [32]. A separate blot was obtained for each cDNA probe. After hybridization, the blots were initially washed for 15 min at room temperature in 2x SSC and 0.1% SDS, and then for an additional 15 min at 65° in 0.2x SSC and 0.1% SDS. Finally, the washed blots were autoradiographed at -70°. For normalization of the amount of RNA in each lane, blots were stripped of radioactivity and reprobed with a human β -actin cDNA probe (ATCC, Rockville, MD). The methodologies employed to probe for CYP2B1, CYP3A1, EPH and $GST_{ya/yc}$ in the slot-blot analysis have been described previously [33].

Preparation of cDNA Probes

The rST_{el} [26], rST_{HA} [27] and rST_a [28] cDNAs were prepared from total RNA that was reverse-transcribed to cDNA and subsequently amplified using standard polymerase chain reaction techniques as described previously [34]. The $rGST_{ya/yc}$ [35] and $pEPH_{302}$ [32] cDNAs were obtained from Dr. Cecil B. Pickett (Schering-Plough Research Institute, Kenilworth, NJ). The $mUGT_{ml}$ [31] cDNA was provided by Dr. Ida S. Owens (NIH, Bethesda, MD). The mST_{pl} [29] and $mUGT_{Br/p}$ [30] were described previously. All cDNA probes were labeled with [α - 32 P]dCTP (New England Nuclear, Boston, MA) using the Multiprime DNA labeling system (Amersham). The specific oligonucleotide probes for CYP2B1 and CYP3A1 have been described previously [33].

Densitometry

The density of the bands of the northern blots was visualized by autoradiography and quantitated with a phosphorimager (AMBIS, Inc., San Diego, CA).

RESULTS

Effect of Tamoxifen on Levels of Sulfotransferase mRNA in Rat Liver

The effect of TAM on hepatic sulfotransferase mRNA is shown in Fig. 1. Hybridization of the northern blot to an rST_{el} cDNA probe (Fig. 1A) revealed a prominent RNA transcript of about 1.4 kb in male control rats (lane 1). As shown in Table 1, this transcript was 0.7, 1.6, and 1.9 times that of control in male rats treated with low, medium, and high doses of TAM, respectively. An increase in ST_{el} mRNA (2.4-fold) was also observed in male rats treated with PB. In contrast to male rats, female rats lacked both basal and drug-induced expression of ST_{el} mRNA.

Figure 1B shows an autoradiogram from a northern blot hybridized to an rST_{HA} cDNA probe. A prominent transcript of approximately 1.4 kb was observed in male control rats (lane 1), which was relatively unchanged in male rats treated with low, medium, or high doses of TAM or with

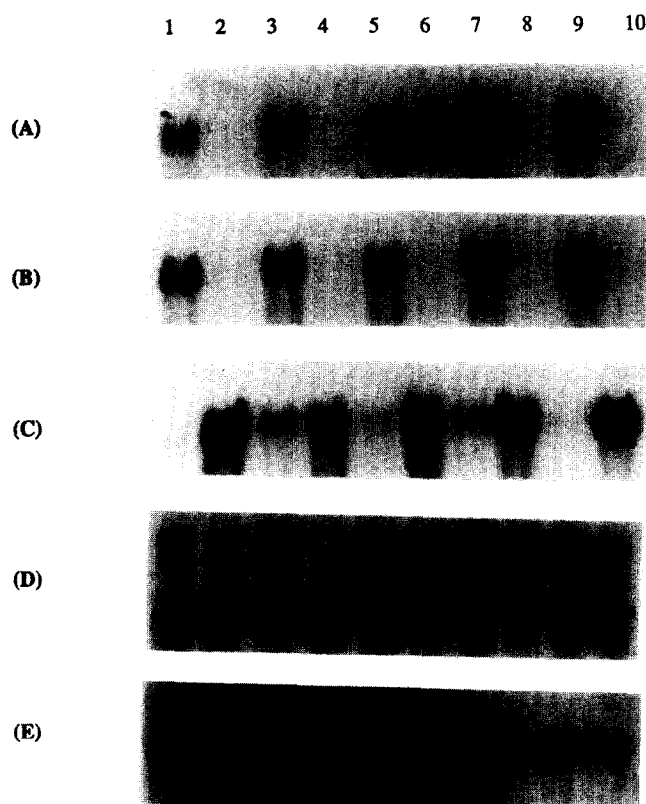


FIG. 1. Northern blot analysis of the effect of TAM on sulfotransferase mRNA levels in rat liver. Total hepatic RNA was isolated from a tissue sample pooled from groups of three rats administered either corn oil, TAM (2.8 to 45 mg/kg body wt) dissolved in corn oil, or a diet supplemented with PB (500 ppm) for 14 days. The blots were hybridized to (A) rST_{el} , (B) rST_{HA} , (C) rST_a , (D) mST_{pl} and (E) human β -actin. Lane 1, male control; lane 2, female control; lane 3, male TAM (11 mg/kg); lane 4, female TAM (2.8 mg/kg); lane 5, male TAM (2.8 mg/kg); lane 6, female TAM (11 mg/kg); lane 7, male TAM (45 mg/kg); lane 8, female TAM (45 mg/kg); lane 9, male PB (500 ppm); and lane 10, female PB (500 ppm).

PB (Table 1). In contrast to male rats, female rats lacked both basal and drug-induced expression of ST_{HA} mRNA.

Hybridization of the northern blot to an rST_a cDNA probe (Fig. 1C) showed a single transcript of about 1.2 kb in both male and female control rats (lanes 1 and 2). However, ST_a mRNA expression tended to be less abundant in the male control animals than in the female. The low basal level of expression of ST_a mRNA in male control rats increased following treatment with TAM. As shown in Table 1, mRNA levels for this particular ST isoform were 3.7, 7.6, and 3.0 times that of control in male rats administered low, medium, and high doses of TAM, respectively. While TAM dramatically increased the level of ST_a mRNA in male rats, PB appeared to have an opposite effect, slightly decreasing mRNA levels to 0.7 times that of control (Table 1). In contrast to male rats, female rats that displayed high basal levels of ST_a showed little response to any of the drug treatments. Figure 1D shows hybridization of the same blot to an mST_{pl} cDNA probe. Two dominant transcripts of 1.5 and 3.5 kb were observed in both male and female control

TABLE 1. Effect of TAM on hepatic mRNA levels of different phase II detoxifying enzymes in male and female F344/NCr rats

cDNA probe	mRNA (fold change vs control)							
	Male				Female			
	TAM (mg/kg)			PB (ppm)	TAM (mg/kg)			PB (ppm)
	2.8	11	45		2.8	11	45	
τST_{el}	0.7	1.6	1.9	2.4				
τST_{HA}	0.9	0.9	0.8	0.7				
τST_{α}	3.7	7.6	3.0	0.7	1.3	1.1	0.9	0.8
mST_{pl}	0.9*	1.8	3.8	1.4	1.0	0.9	1.9	1.7
	1.1†	1.0	1.3	1.2	0.9	0.8	1.3	1.0
$mUGT_{Br/p}$	2.0	2.9	1.8	1.9	2.1	1.9	1.7	2.2
$mUGT_{ml}$	1.4	1.7	2.0	1.6	1.3	1.1	1.5	1.4
$pEPH_{302}$	1.9	4.8	11.2	16.0	2.8	4.2	6.1	7.1

Total hepatic RNA was isolated from a tissue sample pooled from groups of three rats administered either corn oil, TAM (2.8 to 45 mg/kg body wt) dissolved in corn oil, or a diet supplemented with PB (500 ppm) for 14 days. The blots were hybridized to cDNA probes coding for various ST, UGT, and Eph mRNA transcripts. Values (β -actin normalized) were obtained by densitometry of the autoradiograms from northern blots.

* 3.5 kb mST_{pl} mRNA transcript.

† 1.5 kb mST_{pl} mRNA transcript.

rats (lanes 1 and 2). Whereas the smaller 1.5 kb mRNA transcript showed no substantial response to any of the drug treatments, the larger 3.5 kb transcript was increased with high doses of TAM (Table 1).

To normalize the amount of RNA loaded onto each lane, all blots used to probe the different cDNAs were hybridized to a human β -actin cDNA probe. A typical autoradiogram is pictured in Fig. 1E, and shows even loading for all lanes. Ethidium bromide staining of the 18S and 28S ribosomal RNA also indicated that all lanes were loaded evenly (data not shown).

Effect of Tamoxifen on Levels of UGT mRNA in Rat Liver

The effect of TAM on hepatic UGT mRNA is shown in Fig. 2. Hybridization of the northern blot to a family 1 UGT $mUGT_{Br/p}$ cDNA probe (Fig. 2A) showed a prominent RNA transcript of about 2.5 kb in male and female control animals (lanes 1 and 2). Densitometric analysis revealed that the expression of this mRNA transcript was consistently increased in both sexes of animals following

treatment with TAM. As shown in Table 1, $UGT_{Br/p}$ mRNA levels were 2.0, 2.9, and 1.8 times that of control in male rats, and 2.1, 1.9, and 1.7 times that of control in female rats treated with low, medium, and high doses of TAM, respectively. Similar to TAM, PB was also found to be a potent inducer of $UGT_{Br/p}$ expression, increasing mRNA levels 1.9 and 2.2 times that of control in male and female rats, respectively, as been shown by other investigators [36].

Figure 2B shows the northern blot hybridized to a family 2 UGT $mUGT_{ml}$ cDNA probe. A single transcript of approximately 2.5 kb was evident in both groups of control animals (lanes 1 and 2). This mRNA transcript was 1.4, 1.7, and 2.0 times that of control in male rats, and 1.3, 1.1, and 1.5 times that of control in female rats treated with low, medium, and high doses of TAM, respectively (Table 1). In addition, treatment of both male and female rats with PB increased UGT_{ml} mRNA levels 1.6- and 1.4-fold when compared with their respective control groups (Table 1).

Effect of Tamoxifen on Levels of EPH and GST mRNA in Rat Liver

The effect of TAM on hepatic microsomal EPH mRNA probed by $pEPH_{302}$ cDNA is shown in Fig. 3. A prominent mRNA transcript of about 2.0 kb was observed in both male and female control animals (lanes 1 and 2), and the level of expression of this mRNA transcript was increased markedly in both sexes of animals following treatment with TAM. Densitometric analysis shows that EPH mRNA levels increased 1.9, 4.8, and 11.2 times that of control in male rats, and 2.8, 4.2, and 6.1 times that of control in female rats treated with low, medium, and high doses of TAM, respectively (Table 1). Similar to TAM, PB was also found to elevate EPH mRNA to levels of 16.0 and 7.1 times that of control in male and female rats, respectively, consistent with that reported in the literature [32]. The elevation of

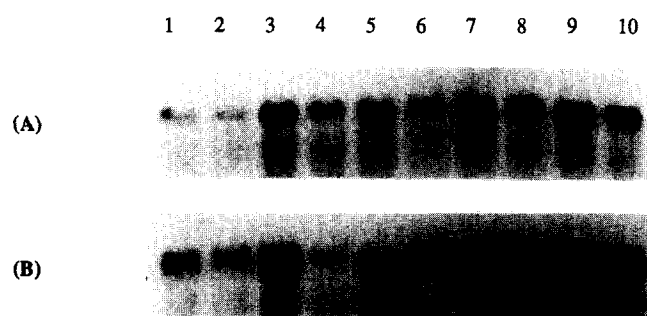


FIG. 2. Northern blot analysis of the effect of TAM on UGT mRNA levels in rat liver. The blots were identical to those described in the legend of Fig. 1 and were hybridized to (A) $mUGT_{Br/p}$ and (B) $mUGT_{ml}$.

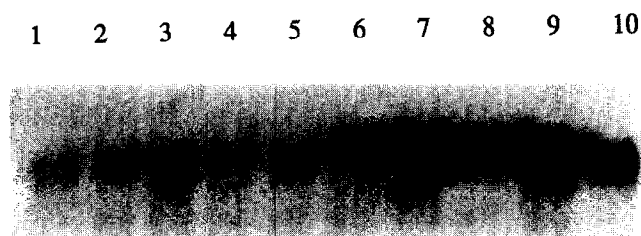


FIG. 3. Northern blot analysis of the effect of TAM on EPH mRNA levels in rat liver. The blot was identical to that of Fig. 1 and was hybridized to *pEPH*₃₀₂.

EPH mRNA level by TAM and PB was further confirmed in Fig. 4 using slot blot. Also in Fig. 4, slot-blot analysis of hepatic RNA showed relatively strong elevation of the *GST*_{ya/yc} family by PB and a somewhat weaker increase by the highest dose of TAM.

Effect of Tamoxifen on the Induction of Various CYP Genes**

The induction of *CYP2B1* and *CYP3A1* by the highest dose of TAM (45 mg/kg body wt) or PB is presented in Fig. 4. As can readily be seen, TAM had minimal effects on the mRNA level of *CYP2B1*, while PB highly increased this specific isozyme of CYP. We have confirmed these results (data not shown) employing both enzymatic methods, dealkylation of benzyloxyresorufin, and immunochemical methods employing slot-blot analysis. These studies showed that although TAM did increase the levels of *CYP2B1*, the maximal levels obtained were never greater than 8% of the levels obtained with PB. Interestingly, we found that *CYP3A1* was highly induced by both PB and the highest dose of TAM. We have similarly confirmed this finding employing immunochemical analysis (data not shown). In contrast, enzymatic methods examining dealkylation of methoxyresorufin (*CYP1A2*) or ethoxyresorufin (*CYP1A1*) showed minimal induction by any of the doses of TAM or PB (data not shown).

DISCUSSION

Chemopreventive agents, as proposed by Wattenberg [37] can potentially act by one of the following pathways: (1) compounds that prevent the formation of carcinogens from precursor substances; (2) compounds that act subsequent to exposure to carcinogenic agents [these have been called "suppressive agents" since they act by suppressing the expression of neoplasia in cells previously exposed to doses of a carcinogenic agent that will cause cancer, and examples of which include the retinoids, protease inhibitors, and non-steroidal anti-inflammatory drugs (NSAIDs) such as indomethacin]; and (3) compounds that inhibit carcino-

genesis by preventing carcinogenic agents from reaching or reacting with critical target sites, i.e. DNA in tissues (these have been termed "blocking agents"). This third group of agents "block" the action of carcinogens either via the induction of carcinogen-detoxifying enzymes in tissues such as the liver, leading to detoxification and subsequent excretion of the carcinogen in the urine or bile, before they have a chance to react with DNA molecules, or via blocking the metabolic activation process by enzymes such as the cytochrome P450s [38]. Hence, in theory, chemical carcinogenesis can be prevented, in most cases, either by blocking the metabolic activation process or by induction of enzymes that preferentially detoxify carcinogens. In fact, PB, which was examined in the present study, has proven to be a highly effective chemopreventive for aflatoxin-induced liver carcinogenesis in both rat and hamster [39] apparently by inducing a specific glutathione transferase gene that preferentially conjugates aflatoxin 8,9-epoxide [40].

As alluded to at the beginning of the paper, there are a variety of reasons for more detailed studies examining the widest range of potential biological effects of TAM. First, TAM has proven to be a highly effective chemopreventive and chemotherapeutic agent in the breast in a variety of species [1,2]. Although it has been proposed that most of the efficacy of TAM is due to its antiestrogenic/estrogenic properties, drugs that highly induce various phase I and phase II drug-metabolizing enzymes may alter levels of endogenous substrates including estrogens [15, 19]. Second, TAM is being employed extensively in both therapeutic trials and in chemoprevention trials in humans; thus, any further information relative to potential drug effects or drug-drug interactions is of great clinical relevance. Finally, TAM has proven to be a potent liver carcinogen in rats and

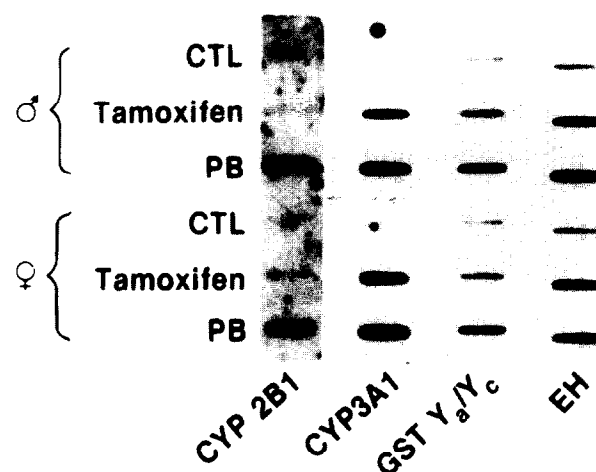


FIG. 4. Slot-blot analysis of the effect of a high dose of TAM (45 mg/kg body wt), or a diet supplemented with PB (500 ppm) for 14 days on the mRNA levels of EPH, GST, *CYP2B1* and *CYP3A1* in rat liver. The blots were hybridized to *rGST*_{ya/yc} or *pEPH*₃₀₂ cDNAs or to oligonucleotide probes for *CYP2B1* or *CYP3A1*.

** Prior experiments with the various phase II genes implied that the higher dose of TAM did appear to give results that were quantitatively similar to those of PB.

hamsters [17]. [The doses of TAM employed in the present studies (2.8 to 45 mg/kg body wt/day) were the same as those used in carcinogenesis assays with TAM and were similar to concentrations of TAM employed in therapy studies in rats.].

One of the most interesting aspects of the present studies is the striking sex-dependent differences in the expression of certain of the rat STs. Thus, ST_{el} and ST_{HA}, both of which are expressed at readily observable levels in male rats, are virtually unexpressed in either control or drug-treated female rats. In contrast ST_a mRNA is expressed at low level in male rats, is highly elevated by TAM, but not by PB, and is not affected by either drug in female rats (although female rats have much higher basal level transcript).

We employed PB as a control in these experiments for three reasons. First, PB induces a variety of phase I and phase II drug-metabolizing enzymes [41] and, therefore, can serve as a positive control for induction of the various UGT and CYP genes examined. Second, previous reports had contended that TAM could induce metabolism of certain substrates associated with CYP2B (pentoxyresorufin O-dealkylase), implying that TAM may be a PB-type inducer [19]. Third, we have shown previously that most strong PB-type inducers [42, 43] are effective liver tumor promoters in initiated rats, and such a mechanism could help to explain the carcinogenic properties of TAM in addition to its known abilities to form DNA adducts. The data with the STs, UGTs and EPH show a similar response of TAM and PB with respect to induction with the exception of ST_a. This ST mRNA transcript was highly elevated by TAM while being suppressed by PB treatment. The results with the CYP genes reinforce the clear dichotomy between these two inducers. Thus, CYP2B1 is highly induced by PB but is only very weakly induced by even the highest dose of TAM. The finding that TAM is an inducer of CYP3A and at best a weak inducer of CYP2B1 is in agreement with the recent results of Nuwaysir *et al.* [44]. This limited induction of CYP2B is in contrast to strong induction by classic promoters, e.g. PB-type inducers [42, 43]. These results indirectly argue that the "PB-type" effect is unlikely to be a primary contributor to the carcinogenic/promoting potential of TAM [42, 43]. Perhaps the most intriguing result is the finding that TAM appears to increase the levels of CYP3A1 as determined by RNA or immunochemical analysis (data not shown). This finding supports observations by White and coworkers [19] of induction of CYP3A1 by TAM. This finding is of particular interest since it is this form of P450 that has been implicated in the major metabolic routes for TAM and is apparently involved in its activation to reactive metabolites in rats and humans [45]. In view of certain of the known drug-drug interactions associated with CYP3A1, this is potentially of some interest in women exposed long term to TAM. In summary, TAM appears to induce a wide variety of phase I and phase II drug-metabolizing enzymes although this pleiotypic re-

sponse appears to be different from that observed with other prototype inducers such as PB, dioxin (TCDD), dexamethasone, or peroxisome proliferators.

Although we have examined the ability of TAM to alter the mRNA levels of phase I and phase II drug-metabolizing enzymes, it alters levels of other genes. Thus, TAM has been shown to stimulate the transcription of many estrogen-regulated genes in the uterus, including ovalbumin [46], insulin-like growth factor-1 (IGF-1) [47], alkaline phosphatase [48], complement C3 [49], progesterone receptor [50], and collagenase [51].*** Webb *et al.* [51] recently reported that TAM activates transcription of human collagenase promoter containing a consensus AP-1 site via the estrogen receptor and the AP-1 pathway. Examination of the xenobiotic detoxifying genes show that *gpST_e* [52], and *rGST_{ya}* [53] contain AP-1 and AP-1-like sites, respectively, at the 5'-flanking region. Future studies will elucidate whether TAM regulates phase I and phase II gene expression through transcription activation of the AP-1 complexes and/or other transcription factors, or through some other pathway such as mRNA stabilization.

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