

Hepatic bile acids and bile acid-related gene expression in pregnant and lactating rat

Background /Aim: Significant physiological changes occur during pregnancy and lactation.

Intrahepatic cholestasis of pregnancy (ICP) is a liver disease closely related to disruption of bile acid homeostasis. The objective of this study was to examine the regulation of bile acid metabolism and transport in normal pregnant and lactating rats.

Materials and Methods: Livers from timed pregnant SD rats were collected on gestational days (GD) 10, 14 and 19, and postnatal days (PND) 1,7,14 and 21. Total bile acids were determined by the enzymatic method, total RNA was isolated and subjected to real time RT-PCR analysis. Liver protein was extracted for western-blot analysis.

Results: Under physiological conditions hepatic bile acids were not elevated during pregnancy but increased during lactation in rats. Bile acid synthesis rate-limiting enzyme Cyp7a1 was unchanged in gestations days, but increased on PND14 and21 at mRNA and protein levels. Expression of Cyp8b1, Cyp27a1 and Cyp7b1 was also higher during lactation. The mRNA levels of small heterodimer partner (SHP) and protein levels of farnesoid X receptor (FXR) were increased during pregnancy and lactation. Bile acid transporters Ntcp, Bsep, Mrp3 and Mrp4 were lower at gestation, but increased during lactation.

Hepatic Oatp transporters were decreased during pregnancy and lactation. **Conclusion:** Hepatic bile acid homeostasis is maintained during normal pregnancy in rats , probably through the FXR-SHP regulation. The expression of bile acid synthesis genes and liver bile acid accumulation were increased during lactation, together with increased bile acid efflux transporter Bsep, Mrp3 and Mrp4 .

1 **Hepatic bile acids and bile acid-related gene expression in pregnant and**
2 **lactating rats**

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11 **Abstract**

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14 acid homeostasis. The objective of this study was to examine the regulation of bile acid synthesis
15 and transport in normal pregnant and lactating rats. **Materials and Methods:** Livers from timed
16 pregnant SD rats were collected on gestational days (GD) 10, 14 and 19, and postnatal days
17 (PND) 1,7,14 and 21. Total bile acids were determined by the enzymatic method, total RNA was
18 isolated and subjected to real time RT-PCR analysis. Liver protein was extracted for western-blot
19 analysis. **Results:** Under physiological conditions hepatic bile acids were not elevated during
20 pregnancy but increased during lactation in rats. Bile acid synthesis rate-limiting enzyme Cyp7a1
21 was unchanged in gestations days, but increased on PND14 and21 at mRNA and protein levels.
22 Expression of Cyp8b1, Cyp27a1 and Cyp7b1 was also higher during lactation. The mRNA levels
23 of small heterodimer partner (SHP) and protein levels of farnesoid X receptor (FXR) were
24 increased during pregnancy and lactation. Bile acid transporters Ntcp, Bsep, Mrp3 and Mrp4
25 were lower at gestation, but increased during lactation. Hepatic Oatp transporters were decreased
26 during pregnancy and lactation. **Conclusion:** Hepatic bile acid homeostasis is maintained during
27 normal pregnancy in rats, probably through the FXR-SHP regulation. The expression of bile acid
28 synthesis genes and liver bile acid accumulation were increased during lactation, together with
29 increased expression of bile acid efflux transporter Bsep, Mrp3 and Mrp4.

30 **Keywords:** Pregnant and lactating rats; Liver bile acids; Cyp7a1; FXR-SHP; Ntcp and Bsep.

31 Introduction

32 Significant physiological changes occur during pregnancy and lactation to support nutritional
33 demand of the developing fetus and lactating pups (Carlin and Alfrevic, 2008; Athipozhy et al.,
34 2011). Bile acids and cholesterol metabolism are important changes during pregnancy and
35 lactation to support and to protect offspring development (Wooton-Kee, Cohen & Vore, 2008;
36 Athipozhy et al., 2011; Abu-Hayyeh, Papacleovoulou & Williamson, 2013a). Such
37 physiological changes would also affect hepatic drug processing genes of phase-1, phase-2
38 metabolism and transporters (Aleksunes et al., 2012; Shuster et al., 2013). The alteration of bile
39 acid homeostasis during pregnancy could unmask cholestatic disease in genetically predisposed
40 but otherwise asymptomatic individuals (Milona et al., 2010). Recent work suggests that in
41 pregnant mice farnesoid X receptor (FXR)-SHP (small heterodimer partner, NR0B2) regulation
42 could be dysfunctional in its ability to down-regulate the rate-limiting bile acid synthetic enzyme
43 Cyp7a1 and 8b1, resulting in bile acids accumulation in the liver of late pregnancy mice (Milona
44 et al., 2010; Aleksunes et al., 2012).

45 Intrahepatic cholestasis of pregnancy (ICP) is a liver disease which can occur in the third
46 trimester of pregnancy (Abu-Hayyeh, Papacleovoulou & Williamson, 2013a). The etiology and
47 pathogenesis of ICP are still not clear, but many studies have related this disease to abnormal bile
48 acid metabolism (Abu-Hayyeh, Papacleovoulou & Williamson, 2013a; Floreani, et al., 2013).
49 ICP with elevated bile acids in serum and liver is a major cause for premature embryo
50 development and embryonic death (Diken et al., 2013). Genetic variations or mutations of
51 farnesoid X receptor (FXR) (Van Mil et al., 2007), bile salt export pump (BSEP/ABCB11) (Dixon

52 [et al., 2009](#)), and ATP-binding cassette, sub-family B (MDR/TAP), member 4 (ABCB4/MDR3)
53 and ABCB11 ([Dixon et al., 2000](#); [Anzivino et al., 2013](#)) contribute to the etiology of ICP. To
54 fully understand bile acid synthesis, transport, and regulation in normal pregnancy would help us
55 to shed light on the pathology of ICP.

56 Estradiol and/or its metabolites may interfere with FXR activity during pregnancy ([Milona et](#)
57 [al., 2010](#); [Aleksunes et al., 2012](#)), and a defect in progesterone metabolism is also implicated in
58 the etiology of ICP ([Pascual et al., 2002](#)). Estrogen signaling is associated with
59 pregnancy-induced hepatotoxicity and cholestasis in mice ([Arrese et al., 2008](#)), and reduced
60 hepatic PPAR- α function in the mouse also appears to be estrogen-dependent ([Papacleovoulou,](#)
61 [Abu-Hayyeh & Williamson, 2011](#)).

62 The above scenario has been studied extensively in mice ([Milona et al., 2010](#); [Aleksunes et al.,](#)
63 [2012](#); [Shuster et al., 2013](#)). Mice and rats are two most commonly used experimental animals, but
64 some physiological responses are different. For example, in mice, Cyp7a1 and liver bile acid pool
65 were not increased during lactation ([Aleksunes et al., 2012](#)), whereas the bile acid synthesis gene
66 Cyp7a1 and hepatic bile acids are increased 2-3 fold in lactating rats ([Wooton-Kee, Cohen &](#)
67 [Vore, 2008](#); [Wooton-Kee et al., 2010](#)). In mice, pregnancy and lactation are associated with
68 decreases in hepatic transporters, including bile acid transporters ([Aleksunes et al., 2012](#)), and
69 such a phenomenon should also be characterized in rats. To fully understand bile acid synthesis,
70 transport, and regulation in normal pregnancy would help us to shed light on the pathology of
71 ICP. This study was initiated to investigate bile acid metabolism and transport gene expressions
72 in pregnant and lactating rats, and the results suggest that under physiological conditions,

73 FXR-SHP regulation might play roles in bile acid homeostasis in pregnant and lactating rats.

74 **Materials and Methods**

75 **Animals.** Adult Sprague Dawley (SD) rats (250 g) were purchased from the Experimental
76 Animal Center of Third Military Medical College (Chongqing, China; certificate No CXK
77 2007-0005). Rats were kept in a SPF-grade animal facilities (certificate No SYXK 2011-004) at
78 Zunyi Medical College, with regulated environment ($22 \pm 1^{\circ}\text{C}$, $50 \pm 2\%$ humidity and a 12 h: 12
79 h light: dark cycle) and free access to purified water and standard rodent chow. Rats were
80 acclimatized for 1 week, and subjected to timely mating overnight. A vaginal plug in the next
81 morning was designated as day 0 (GD 0) of gestation. Maternal livers were collected without
82 fasting on GD10, GD14 and GD19, as well as on the postnatal days (PND) 1, 7, 14 and 21. The
83 age-matched virgin rats were used as controls. The experimental design followed similar time
84 points in mice ([Aleksunes et al., 2012](#)). Livers were weighed, snap frozen in liquid nitrogen, and
85 stored at -80°C until analysis. All animal procedures follow the NIH guide of Humane Use and
86 Care Animals, and approved by Institutional Animal Use and Care Committee of Zunyi Medical
87 College

88 **Bile acid determination**

89 Bile acids were extracted from the liver and measured with the “Total” Bile Acid assay (TBA)
90 kit (Nanjing Jian-Cheng Bioengineering Co., China). Briefly, livers were homogenized in
91 physiological saline (1:9, wt :vol), followed by centrifugation at 2500 rpm/min for 10 min. the

92 supernatant (30 μ l) was taken for determination of bile acids according to manufacturer's
93 protocol.

94 **RNA Isolation and real-time RT-PCR analysis**

95 Total RNA was isolated from frozen liver sample (50-100 mg) using 1 ml TRIzol (Takara,
96 Biotechnology, Dalian, China) and subsequently purified with Total RNA (Mini) Kit (Watson
97 Biotechnology, Shanghai, China). The quality of purified RNA was determined by
98 spectrophotometry with the 260/280 ratio >1.8 . Purified RNA was reversed transcribed with the
99 High Capacity Reverse Transcriptase Kit (Applied Biosystems, Foster City, CA, USA). The
100 primer pairs were designed with the Primer3 software and listed in Supplementary Table 1. The
101 Power SYBR Green Master Mix (Applied Biosystems, Foster City, CA, USA) was used for
102 real-time RT-PCR analysis. The cycle time for reaching threshold (Ct) of each target gene was
103 normalized to the housekeeping genes (G3PDH and β -actin), and expressed as % of
104 housekeeping genes.

105 **Western Blot Analysis**

106 Livers were homogenized in RIPA lysis buffer (Beyotime, P0013B, Shanghai, China)
107 containing freshly-prepared proteinase inhibitors. The supernatants were centrifuged at 12000
108 rpm 10 min at 4°C, and protein concentrations were quantified by the BCA assay (Beyotime,
109 P0012, Shanghai, China). Aliquot proteins were denatured with protein loading buffer (Beyotime,
110 P0015, Shanghai, China), and approximately 50 μ g of protein/lane was separated on 10%
111 SDS-PAGE and transferred to PVDF membranes. Membranes were blocked in 5% non-fat milk

112 in TBST, followed by incubation overnight at 4 °C with 1:1000 CYP7A1 (ab65586) and β -actin
113 (Ab8227) from Abcam (Cambridge, MA), or FXR (sc-13063) from Santa Cruz Biotechnology
114 (Santa Cruz, CA) in 1% BSA. After washing with TBST three times, the membranes were
115 incubated with HRP-conjugated anti-rabbit or anti-mouse IgG (Beyotime, A0208 and A0216,
116 Shanghai, China). Protein-antibody complexes were visualized using an enhanced
117 chemiluminescent reagent (ECL-Plus) (Beyotime, P0018, Shanghai, China), and exposed to Gel
118 Imaging (Bio-Rad, ChemiDoc XRS, USA). The intensity of the band was semi-quantified with
119 Quantity One software.

120 **Statistical Analysis**

121 The software SPSS17.0 was used for statistical analysis. Results were described using mean \pm
122 SEM. Difference between virgin and pregnant rats was determined by two-tailed independent
123 samples test, $P < 0.05$ was considered statistically significant.

124 **Results**

125 **Liver bile acid levels in pregnant and lactating rat**

126 Bile acids were quantified in livers from control and pregnant rats at GD10, 14, and 19 and
127 PND 1, 7, 14, and 21. Liver bile acid levels slightly decreased in late pregnancy, especially on
128 GD 10, and 19. After birth, liver bile acid concentrations tended to increase, and there is a
129 significant increase in PND 21 (30% over control) (Fig.1).

130 **Hepatic mRNA expression of bile acid synthesis genes in pregnant and lactating rat.**

131 Expression of the classic pathway bile acid synthetic enzyme genes (Cyp7a1 and 8b1) and

132 alternative pathway (Cyp27a1 and 7b1), is shown in Fig.2. The expression of rate-limiting
133 Cyp7a1 mRNA was unchanged during pregnancy, and increased on postpartum. Cyp8b1 mRNA
134 decreased in GD10 and GD14, and increased about 2-fold in PND14. The expression of
135 alternative pathway genes Cyp27a1 and Cyp7b1 were unchanged in gestation days and increased
136 in postnatal days.

137 **Hepatic expression of bile acid synthetic rate-limiting protein Cyp7A1 in pregnant and** 138 **lactating rat.**

139 Western blots were performed using liver homogenates from control rats, pregnant rats at GD
140 10, 14, 19 and lactating rats at PND 1, 7, 14 and 21. The expressions of CYP7A1 protein were
141 semi-quantified by band intensity. CYP7A1 protein was basically unchanged during pregnancy, a
142 result similar to Cyp7a1 mRNA expression, but increased on lactation days PND7, 14 and 21
143 (Fig. 3).

144 **Hepatic mRNA expression of nuclear receptors FXR, SHP, and ESR-1, PPAR- α in pregnant** 145 **and lactating rat**

146 The expression of bile acid regulation nuclear receptor genes farnesoid X receptor (FXR,
147 NR1H4) did not show significant increases during pregnancy, while FXR gradually increased on
148 postpartum. The small heterodimer partner (SHP; NR0B2) significantly increased in the late
149 gestational days, increased 3-fold on GD 19 as compared to controls. FXR plays an important
150 role in bile acid homeostasis by inducing the transcription repressor SHP ([Chiang, 2009](#)).
151 Estrogen receptor alpha (ESR-1) decreased to 64.7% and 57.7% on GD10 and GD14. In
152 postnatal days, ESR-1 increased 2.33-fold in PND1 and then decreased to 68% of control on

153 PND21. Proliferator-activated receptor α (PPAR α) increased 3.79-fold compared to controls
154 during lactation.

155 **Hepatic expression of FXR protein in pregnant and lactating rat.**

156 Western blots were performed using liver homogenates from control rats, pregnant rats at GD
157 10, 14, 19 and lactating rats at PND 1, 7, 14 and 21. The expression of FXR protein was
158 semi-quantified by band intensity. FXR protein was increased during late pregnancy (GD10 to
159 GD19) and early lactation (PND1 to PND7) (Fig. 5).

160 **Hepatic mRNA expression of bile acid transporters in pregnant and lactating rats.**

161 As illustrated in Fig 6, the expression of bile acid efflux transporter bile salt export pump
162 (Bsep/ABCB11) was decreased during pregnancy but increased during lactation. The multidrug
163 resistance protein 3 (Mrp3) and Mrp4 showed the similar pattern, with slightly increases during
164 lactation. The ATP-binding cassette sub-family G member 2 (Abcg2/BCRP) was also decreased
165 in the gestation and lactation days expect PND1.

166 **Hepatic mRNA expression of uptake OATP transporters and Ntcp in pregnant and** 167 **lactating rats.**

168 Figure 7 demonstrates that the expression of canalicular uptake transporter solute carrier
169 organic anion transporter (Oatp1/Slco1a1), solute carrier organic anion transporter
170 (Oatp2/Slco1b2), and organic anion-transporting polypeptide 4 (Oatp4/Slc21a10) were all
171 decreased in the gestation days, on PND1 Oatp1 increased 1.68-fold, then decreased in postnatal
172 days. In comparison, Oatp2 and Oatp4 decreased in the both gestation days and lactation days.

173 The uptake transporter Na⁺-taurocholate co-transporting polypeptide (Ntcp) was also decreased
174 during pregnancy, increased on PND1, but decreased again thereafter during lactation.

175 **Discussion**

176 The present study demonstrates that in pregnant rats, hepatic bile acids were not elevated.
177 Consistent with hepatic bile acid concentrations, bile acid synthesis genes and/or enzymes, i.e.,
178 Cyp7a1, Cyp8b1, Cyp27a1 and Cyp7b1 were not increased during pregnancy. Increased FXR
179 protein and SHP mRNA are associated with bile acid homeostasis during pregnancy. In
180 comparison, lactating rats had increased liver bile acid, increased bile acid synthetic enzymes,
181 and increased expression of bile acid efflux transporters. In general, OATP transporters and bile
182 acid uptake transport Ntcp were down-regulated during pregnancy and lactation in rats.

183 ICP is characterized by raised serum bile acid levels and abnormal liver function tests ([Geenes](#)
184 & [Williamson, 2009](#); [Diken, Usta & Nassar, 2013](#)). However, in normal pregnant women, serum
185 bile acid levels are not necessarily increased during pregnancy, regardless of gestation days
186 ([Barth et al., 2005](#); [Egan et al., 2012](#)). In experimental animal studies, a mild increase in liver bile
187 acid levels during normal pregnancy in mice was reported in some studies ([Aleksunes et al.,](#)
188 [2012](#)), but not in others ([Abu-Hayyeh, Papacleovoulou & Williamson, 2013a](#)). In the majority
189 cases such mild increases do not reach pathological levels and remain below the upper end of the
190 reference range for serum bile acid levels ([Abu-Hayyeh, Papacleovoulou & Williamson, 2013a](#)).
191 Thus, it is not surprising that in the present study, liver bile acids were not elevated during
192 pregnancy in gestation days (Fig. 1). The expression of bile acid synthesis gene and proteins
193 during the gestation days (Fig. 2 and 3) is in agreement with hepatic bile acid profiles.

194 ICP has a complex etiology including genetic factors, endocrine factors, and the impact of

195 pregnancy on FXR function ([Abu-Hayyeh, Papacleovoulou & Williamson, 2013a](#); [Floreani, et,](#)
196 [al., 2013](#)). The present study focused on FXR-SHP regulation under physiological conditions. It
197 is proposed that pregnancy in mice resembles a state of FXR inactivation ([Milona et al., 2010](#);
198 [Aleksunes et al., 2012](#)), and attenuated FXR function during mouse pregnancy has been reported
199 ([Papacleovoulou, Abu-Hayyeh & Williamson, 2011](#);[Aleksunes et al., 2012](#)) and the 3 β -sulfated
200 progesterone metabolite epiallopregnanolone sulfate was found to inhibit FXR, resulting in
201 reduced FXR-mediated bile acid efflux ([Abu-Hayyeh et al., 2013b](#)). In the present study, the
202 expression of FXR mRNA in rats during pregnancy was basically unchanged. However, the FXR
203 protein and FXR-inducible negative target SHP were markedly increased at the late gestation
204 days and reached approximately 3-fold higher at GD14 and GD19, despite FXR mRNA was not
205 increased. The increases in FXR-SHP may play important role in maintaining the bile acid
206 homeostasis and preventing the liver bile acids to accumulate to protect the fetus from the bile
207 acid toxicity.

208 Lactation is a time of a five-fold increase in energy demand, as suckling young requires a
209 proportional adjustment in the ability of the lactating dam to absorb nutrients ([Cripps &](#)
210 [Williams, 1975](#); [Vernon et al., 2002](#)). Lactating rats have a two to three-fold increase in food
211 consumption to ensure lactating dams to absorb nutrients and to synthesize critical molecules
212 including bile acids to meet the dietary needs of the offspring and the dam ([Vernon et al., 2002](#)).
213 The size and hydrophobicity of the bile acid pool increase during lactation, implying an increased
214 absorption and disposition of lipid, sterols, nutrients, and xenobiotics ([Athipposhy et al., 2011](#)).
215 In essence, rats ([Wooton-Kee, Cohen & Vore, 2008](#)) are different from mice ([Aleksunes et al.,](#)
216 [2012](#)) in bile acid homeostasis during lactation. In the present study, hepatic bile acid pool (Fig.

217 1), mRNA levels of bile acid synthesis gene Cyp7a1, Cyp8b1, Cyp27a1 and Cyp7b1 (Fig. 2 and
218 Fig. 3) were all increased during lactation, consistent with this scenario.

219 The mRNA levels of bile acid transporters Ntcp and Bsep followed the similar pattern. Ntcp is
220 the major bile acid transporter for conjugated bile acid (Csanaky et al., 2011) and Bsep is the
221 major bile acid efflux pump located at the bile canalicular apical domain of hepatocytes (Lam,
222 Soroka & Boyer, 2010). Down-regulation of Ntcp and Bsep was observed in pregnant rats
223 (Arrese et al., 2003; Cao et al., 2001), however, they are increased on early postpartum, probably
224 under the influence of prolactin (Cao et al., 2001). In the present study, the changes in mRNA
225 levels of Ntcp and Bsep showed a similar pattern, i.e., lower expression during pregnancy but
226 returned to normal and even increased during lactation.

227 Sulfated progesterone metabolite (P4-S) levels are raised in normal pregnancy and elevated
228 further in ICP, which can cause a competitive inhibition of NTCP-mediated uptake of
229 taurocholate in *Xenopus* oocytes (Abu-Hayyeh et al., 2010), and also can cause inhibition of
230 BSEP (Vallejo et al., 2006). In the present study, mRNA levels of Ntcp and Bsep (Fig. 5) were
231 lower during pregnancy, and Bsep was increased during lactation, consistent with liver bile acid
232 homeostasis profile. Mrp3 and Mrp4 are two major bile acid efflux (Cui et al., 2009; Aleksunes
233 et al., 2012), and mRNA levels of these two genes expression showed the similar pattern (Fig. 5),
234 i.e., lower during the pregnancy and higher during lactation. The pattern of these transporter
235 mRNA levels coincide with FXR-SHP regulation of bile acid homeostasis, and fortifying the
236 concept that under physiological conditions, FXR-SHP regulation of bile acid synthesis could be
237 essential for maintaining the bile acid homeostasis and could prevent the occurrence of ICP, an
238 unusual pathological condition.

239 One of the major findings in the study is the decreased mRNA levels of Oatp transporters (Fig.
240 6), and this finding is consistent with that observed in mice (Aleksunes et al., 2012; Shuster et al.,
241 2013). Oatps are important not only for bile acid transport (Zhang et al., 2012), but also
242 important for drugs and xenobiotic transport (Lu et al., 2008). In pregnant rats, the expression of
243 Oatp2, but not Oatp1, was reported to decrease (Cao et al., 2001; 2002). The generalized
244 down-regulation of Oatp transporters could be an adaptive mechanism for dam to protect
245 developing fetus and nursing pups from toxicants. Abcg2 is involved in epithelial
246 transport/barrier functions, including bile acid transport (Blazquez et al., 2012). Abcg2 is
247 proposed to play a key role in bile acid transport in placenta, as Bsep does in liver (Blazquez et
248 al., 2012). In the present study, mRNA levels of Abcg 2 was depressed during pregnancy and
249 lactation except for a transient increase at PND1. The pattern of Abcg2 expression is similar to
250 Oatps, and can also be envisioned as an adaptive mechanism during pregnancy and lactation.

251 In summary, the present study suggests that in pregnant rats, FXR-SHP could regulate bile acid
252 synthesis enzyme genes to prevent the accumulation of bile acids in the liver, together with
253 down-regulation of bile acid transporters Ntcp and Bsep. Pregnancy and lactation is associated
254 with a general down-regulation of Oatp and Abcg2 in rats. These data would add to our
255 understanding of FXR-SHP regulation of bile acid homeostasis under physiological conditions.

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366 **Figure Legends:**

367 **Figure 1. Liver bile acid levels in pregnant and lactating rat**

368 Bile acids were quantified in Livers from control and pregnant rat on GD10, 14, and 19 and PND
369 1, 7, 14, and 21. Dark gray bars represent pregnant rat, and black bars represent lactating rat. Data
370 are presented as mean \pm SEM (n=3-6). Asterisks [*] represent statistically significant difference (
371 $p < 0.05$) compared with control.

372 **Figure 2. Hepatic mRNA expression of bile acid synthetic pathway genes in pregnant and**
373 **lactating rat.**

374 The expression of bile acid synthetic classic pathway genes Cyp7a1, Cyp8b1 and alternative
375 pathway genes Cyp27a1 and Cyp7b1 was quantified from control and GD10,14 and 19 and PND
376 1,7,14 and 21, Data were normalized to controls (set to 100%) and presented as mean \pm SEM
377 (n=3-6). Dark gray bars represent pregnant rat, and black bars represent lactating rat. Asterisks
378 [*] represent statistically significant difference ($p < 0.05$) compared with control.

379 **Figure 3. Hepatic expression of bile acid synthesis rate-limiting protein CYP7A1 in**
380 **pregnant and lactating rat.**

381 Western bolts were performed using liver homogenates from control, pregnant rats in GD 10, 14,
382 19 and PND 1, 7, 14 and 21. The expression of CYP7A1 was semi-quantified by band intensity.
383 Values are mean \pm SEM (n=5). Dark gray bars represent pregnant rat, and black bars represent
384 lactating rat. Significantly difference was confirmed by two-tailed independent Samples test
385 method ($P < 0.05$)

386 **Figure 4. Hepatic mRNA expression of Nuclear Receptors SHP, FXR and ESR-1 and**
387 **PPAR- α in pregnant and lactating rat**

388 The expression of bile acid regulation Nuclear Receptors genes SHP, FXR and Esr-1, PPAR- α
389 were quantified using total hepatic RNA from control and pregnant mice at gestational days 10,
390 14, 19 and postnatal days 1, 7, 14, 21. Data were normalized to controls and presented as mean \pm
391 SEM (n=3-6). Dark gray bars represent pregnant rat, and black bars represent lactating rat.
392 Asterisks (*) represent statistically significant differences (p < 0.05) compared to control.

393 **Figure 5. Hepatic expression of nuclear receptor FXR in pregnant and lactating rat.**

394 Western blots were performed using liver homogenates from control, pregnant rats in GD 10, 14,
395 19 and PND 1, 7, 14 and 21. The expression of FXR was semi-quantified by band intensity.
396 Values are mean \pm SEM (n=3). Dark gray bars represent pregnant rat, and black bars represent
397 lactating rat. Significant difference was confirmed by two-tailed independent Samples test
398 method (P < 0.05)

399 **Figure 6. Hepatic mRNA expression of bile acid transporter in pregnant and lactating rats.**

400 The expression of bile acid efflux transporter Bsep, Mrp3, Mrp4 and Abcg2 was quantified using
401 total hepatic RNA from pregnant rats on GD 10, 14 and 19, and postpartum rats on PND 1, 7, 14
402 and 21. Data were normalized to controls and presented as mean \pm SEM (n=3-6). Dark gray bars
403 represent pregnant rat, and black bars represent lactating rat. Asterisks (*) represent statistically
404 significant differences (p < 0.05) compared control.

405 **Figure 7. Hepatic mRNA expression of uptake transporter Oatps and Ntcp in pregnant and**

406 **lactating rats.**

407 The expression of hepatic canalicular uptake transporter solute carrier organic anion transporters
408 (Oatps) and Na⁺-taurocholate co-transporting polypeptide (Ntcp) transporters was quantified
409 using total hepatic RNA from pregnant rats on GD10, 14 and 19, and postpartum rats on PND 1,
410 7, 14 and 21. Data were normalized to controls and presented as mean ± SEM (n=3-6). Dark gray
411 bars represent pregnant rat, and black bars represent lactating rat. Asterisks (*) represent
412 statistically significant differences ($p < 0.05$) compared control.

413 **Supplementary Table 1.**

Gene	Access	Forward	Reverse
G3PDH	NM_017008.4	AACTTTGGCATTGTGGAAGG	GGATGCAGGGATGATGTTCT
<i>β-actin</i>	NM_031144	AGCCATGTACGTAGCCATCC	ACCCTCATAGATGGGCACAG
<i>Cyp7a1</i>	NM_012942	GAGAACGGGTTGATTCCGTA	AAAAACGTGACCATGCTTC
<i>Cyp8b1</i>	NM_031241	CACGTAGCCAGTACCAAGCA	GGTCCTAGCATCACCAAGGA
<i>Cyp27a1</i>	NM_178847	TCTGGCTACCTGCACTTCT	GTCTACCCAGCCAAGATCA
<i>Cyp7b1</i>	NM_019183	TCATCCGTGAAGTGCAAGAG	GGAGCATCGAAGACTTCTGG
FXR	NM_021745	CGAGATGCCTGTGACAAAGA	GCAGACCACACACAGCTCAT
<i>SHP</i>	NM_053908	TTATGTGTGAGGGTGGACGA	CCCGTCTTCTTGAAGTGCTC
<i>Esr-1</i>	NM_012689	TCCGGCACATGAGTAACAAA	TGAAGACGATGAGCATCCAG
<i>PPAR-α</i>	NM_013196	GAGACCCTCGGGGATCTTAG	TGTGTCCTGAGCTTGACCAG
<i>Ntcp</i>	NM_017047	CACAACGTATCAGCCCCTTT	ATGCTAAGCGCCTTGTCTGT
<i>Bsep</i>	NM_031760	CCACCAGAACATGACAAACG	CCCAGTGATGACCCATAACC
MRP3	NM_080581	CCAGACCTCACACCCTGTTT	CGTCTTGAGCCTGGATAAGC
MRP4	NM_133411	TGAAGCAACTGCAAATGTGG	AGTGCACTGGGCAAACCTTCT
<i>Oatp1</i>	NM_017111	GGATGTAGCTGAGGCAGAGG	CAGCTCCCAGTGGCATTAT
<i>Oatp2</i>	NM_0131906	CCTAGGCATAGGCATTTGGA	TCAACCAAAGCACAAAGCAG
<i>Oatp4</i>	NM_031650	AACATGCTTCGTGGGATAGG	CATGGAAGTGTGCCCTTCTT
<i>Abcg2</i>	NM_181381	GAAAGACCCACGGGGATTAT	CCCATCACAACGTCATCTTG

Figure 1

Fig.1

Liver bile acid concentrations

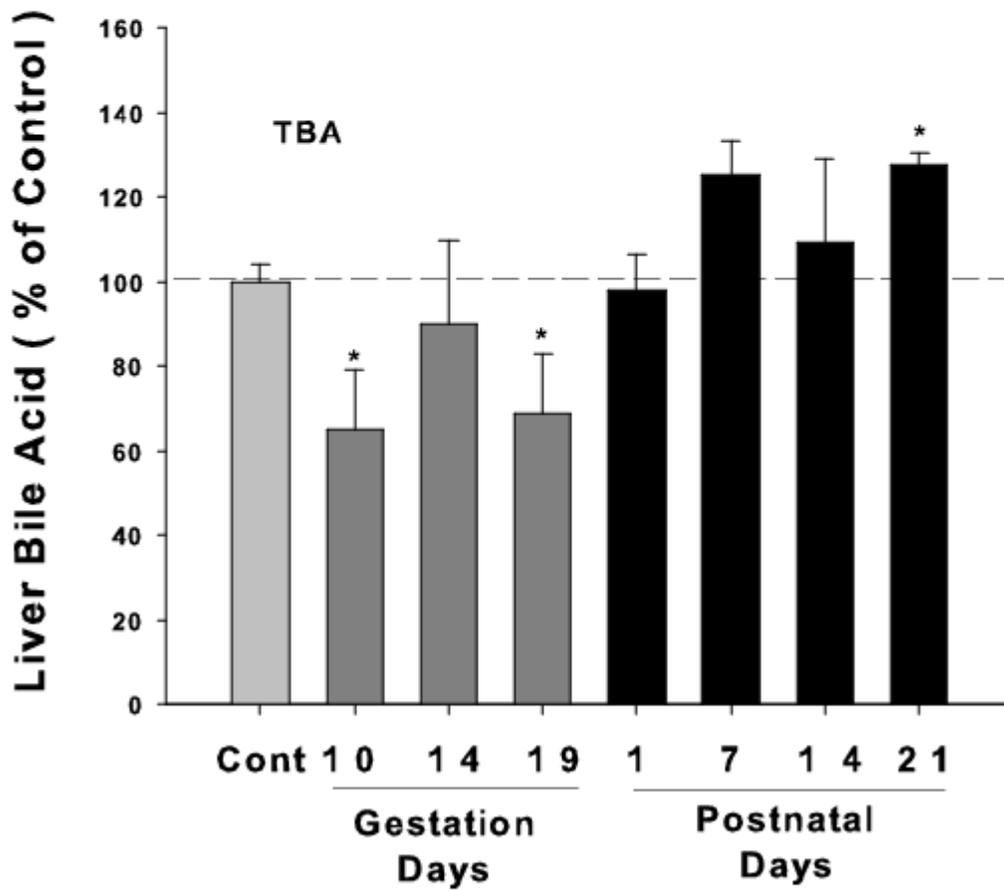


Fig.1

Figure 2

Fig. 2

Bile acid synthesis gene expression

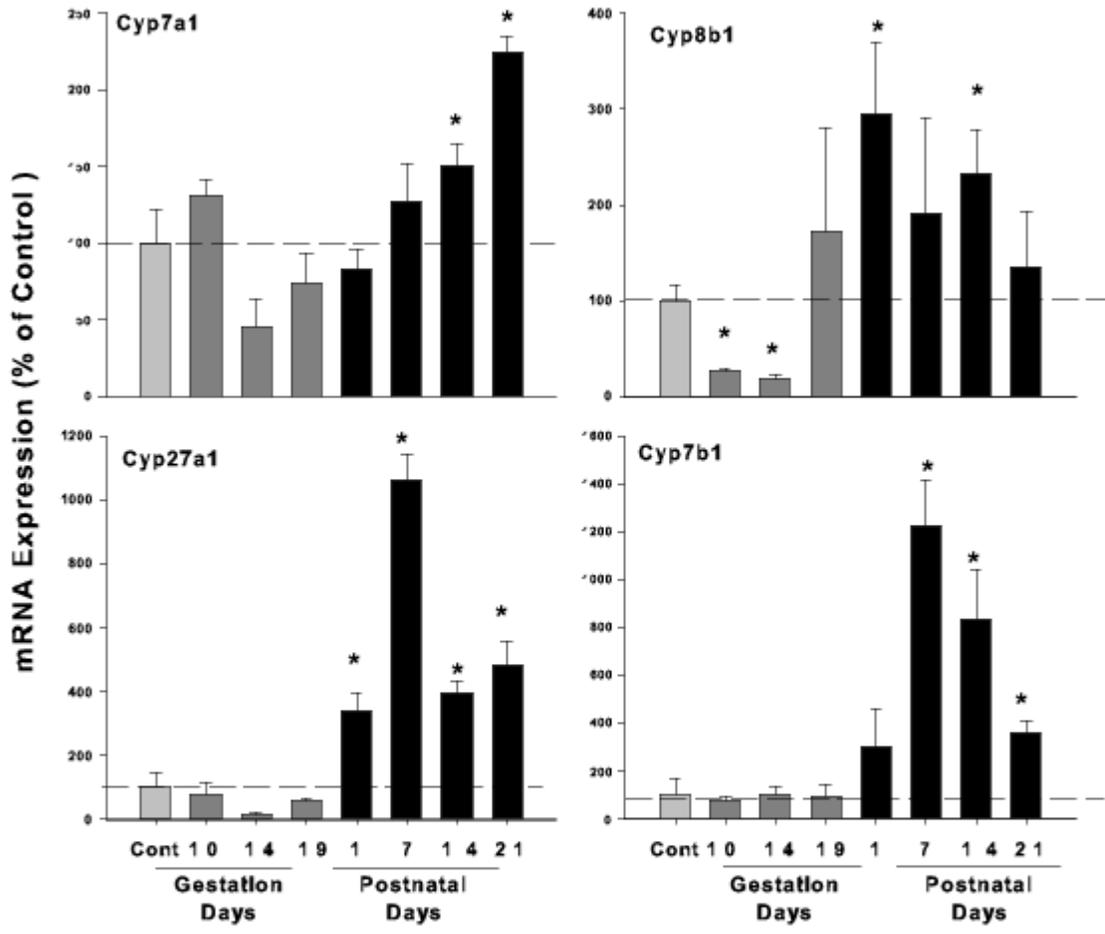


Fig.2

Figure 3

Hepatic expression of bile acid synthesis rate-limiting protein CYP7A1 in pregnant and lactating rat.

Western bolts were performed using liver homogenates from control, pregnant rats in GD 10, 14, 19 and PND 1, 7, 14 and 21. The expression of CYP7A1 was semi-quantified by band intensity. Values are mean \pm SEM. Dark gray bars represent pregnant rat, and black bars represent lactating rat.

Significant difference was confirmed by two-tailed independent Samples test method ($P < 0.05$).

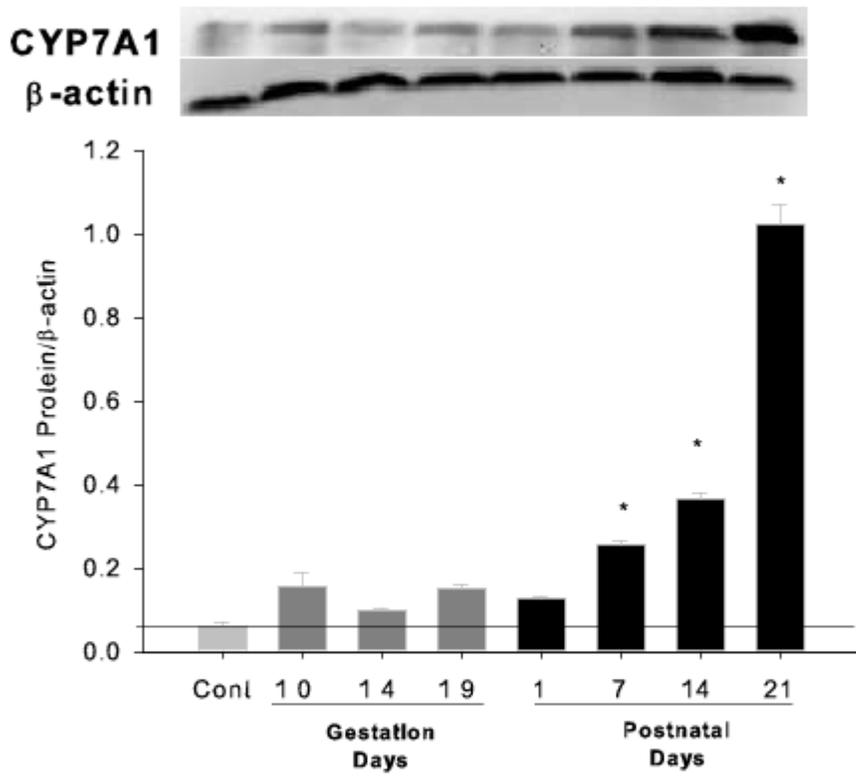


Figure 4

Fig. 4

Expression of nuclear receptors

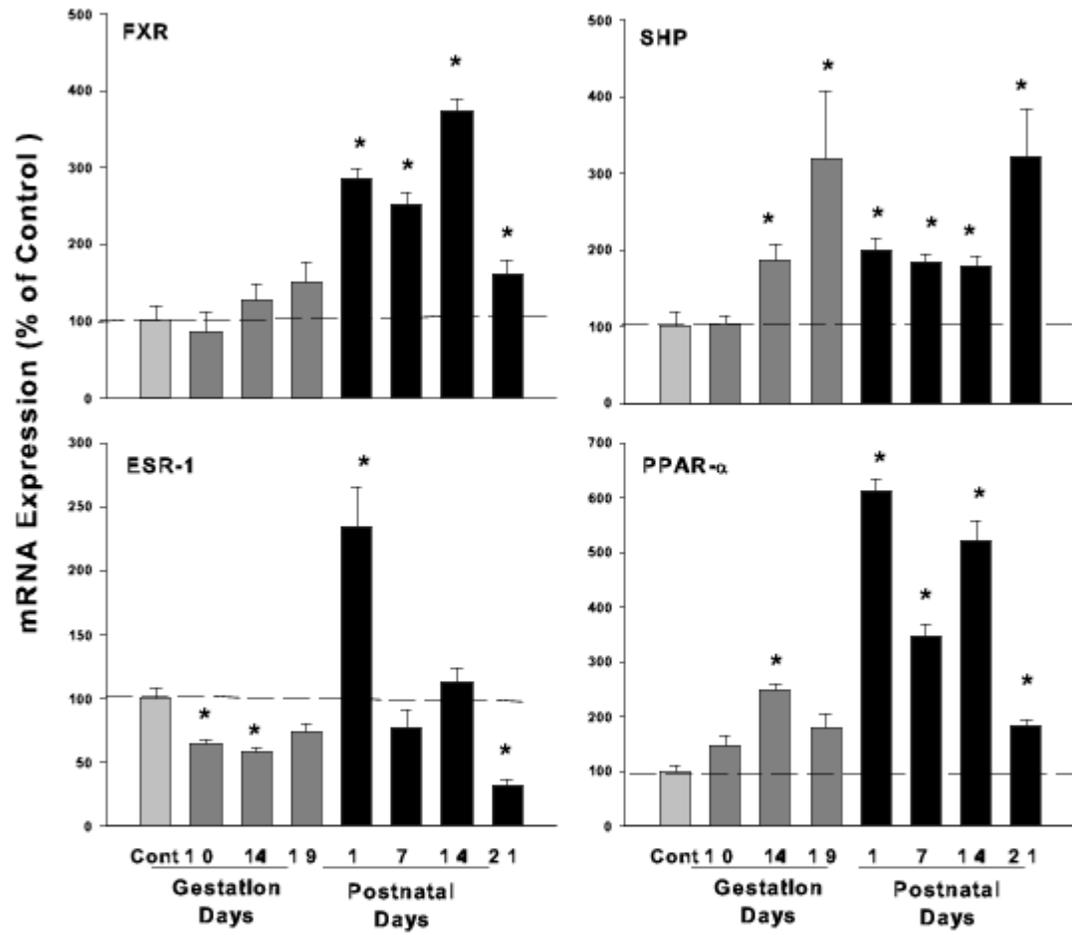


Fig.4

Figure 5

Fig. 5

FXR protein expression

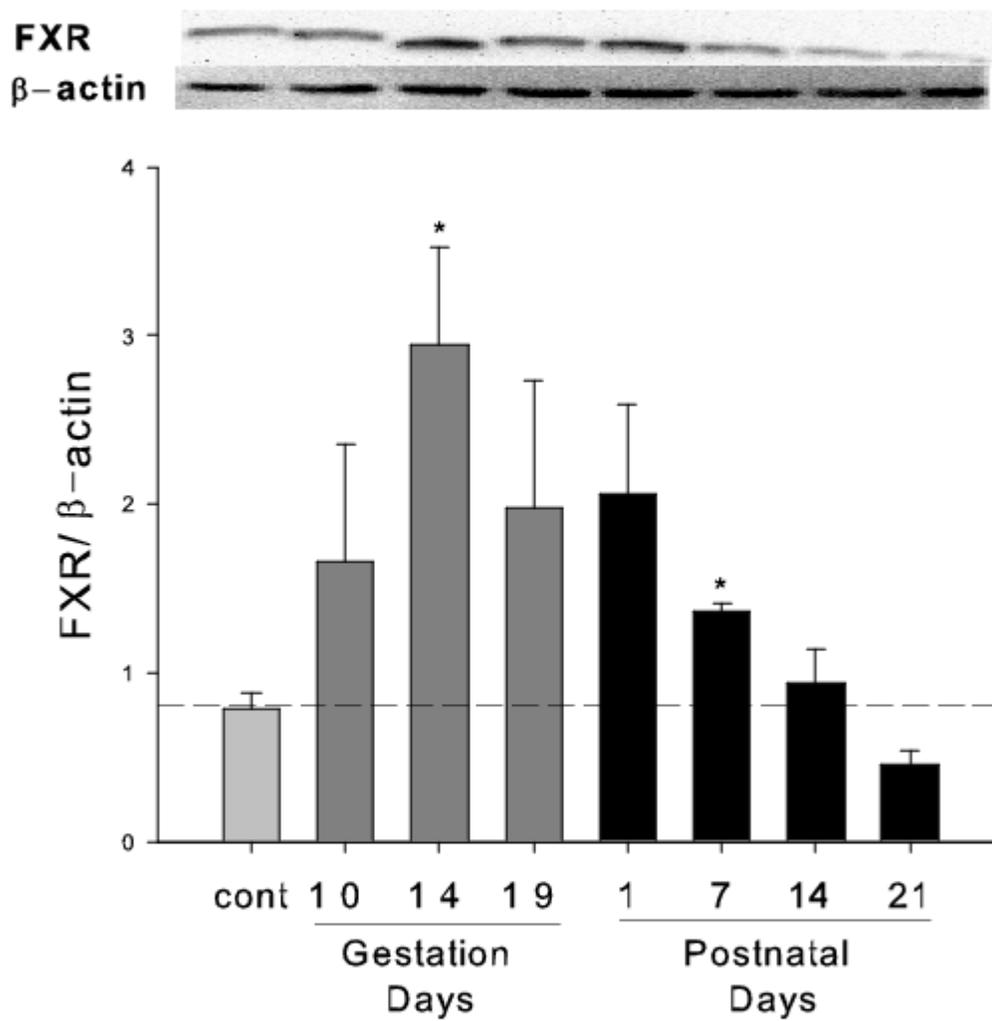


Fig.5

Figure 6

Fig. 6

Efflux transporters

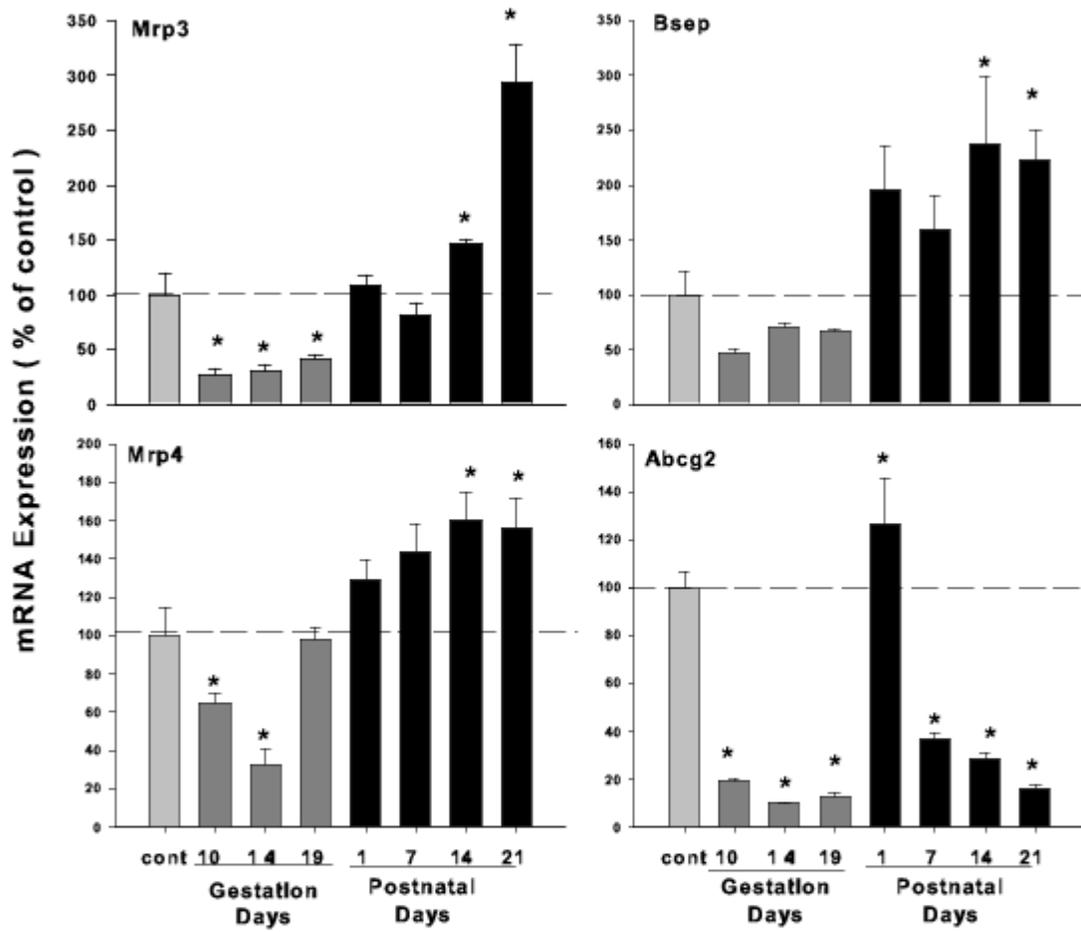


Fig.6

Figure 7

Fig. 7

Uptake transporter expression

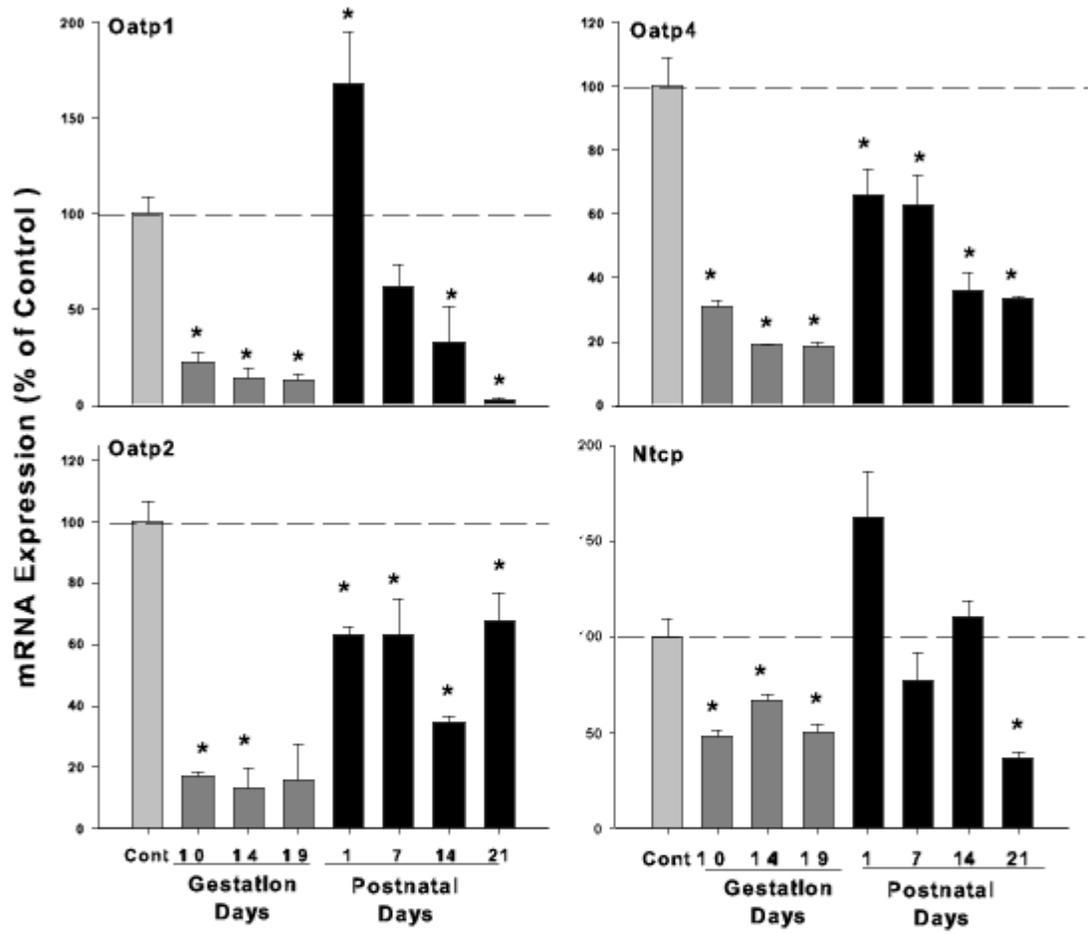


Fig.7