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

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

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Abstract

Background /Aim: Intrahepatic cholestasis of pregnancy (ICP) is a liver disease which may occur in the third trimester of pregnancy. The etiology and pathogenesis of ICP is thought to be closely related to bile acid metabolism. The objective of this study was to examine the regulation of bile acid metabolism during normal pregnant and lactation in rats. **Materials and Methods:** Livers from timely 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 expression of small heterodimer partner (SHP) was increased at GD19 and lactation days, and farnesoid X receptor (FXR) increased on postpartum. 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 maintained during normal pregnancy in rats, probably through the regulation of SHP. The expression of bile acid synthesis genes and liver bile acids were increased during lactation, together with increased expression of bile acid transporters.

Keywords: Pregnant and lactating rats; Liver bile acids;Cyp7a1 and SHP; Ntcp and Bsep.

Introduction

Intrahepatic cholestasis of pregnancy (ICP) is a liver disease which can occur in the third trimester of pregnancy (Abu-Hayyeh, Papacleovoulou & Williamson, 2013a). The etiology and pathogenesis of ICP are still not clear, but many studies have related this disease to abnormal bile acid metabolism (Abu-Hayyeh, Papacleovoulou & Williamson, 2013a; Floreani, et al., 2013). ICP with elevated bile acids in serum and liver is a major cause for premature embryo development and embryonic death (Diken et al., 2013). Genetic variations or mutations of farnesoid X receptor (FXR) (Van Mil et al., 2007), bile salt export pump (BSEP/ABCB11)(Dixon et al., 2009), and ATP-binding cassette, sub-family B (MDR/TAP), member 4 (ABCB4/MDR3) and ABCB11 (Dixon et al., 2000; Anzivino et al., 2013) contribute to the etiology of ICP. To fully understand bile acid synthesis, transport, and regulation in normal pregnancy would help us to shed light on the pathology of ICP.

Significant physiological changes occur during pregnancy and lactation to support nutritional demand of the developing fetus and lactating pups (Carlin and Alfrevic, 2008; Athippozhy et al., 2011). Bile acids and cholesterol metabolism are important changes during pregnancy and lactation to support and to protect offspring development (Wooton-Kee, Cohen & Vore, 2008; Athippozhy et al., 2011; Abu-Hayyeh, Papacleovoulou & Williamson, 2013a). Such physiological changes would also affect hepatic drug processing genes of phase-1, phase-2 metabolism and transporters (Aleksunes et al., 2012; Shuster et al., 2013). The alteration of bile acid homeostasis during pregnancy could unmask cholestatic disease in genetically predisposed but otherwise asymptotic individuals (Milona et al., 2010). Recent

work suggests that in pregnant mice FXR-SHP (small heterodimer partner, NR0B2) regulation mechanism could be dysfunctional for its ability to down-regulate the rate-limiting bile acid synthetic enzyme Cyp7a1 and 8b1, resulting in bile acids accumulation in the liver of late pregnancy mice (Milona et al., 2010; Aleksunes et al., 2012). Furthermore, the disease usually develops in the third trimester of pregnancy when concentrations of estrogen are highest. Estradiol and/or its metabolites may interfere with FXR activity during pregnancy (Milona et al., 2010; Aleksunes et al., 2012), and estrogen signaling is associated with pregnancy-induced hepatotoxicity and cholestasis in mice (Arrese et al., 2008). Reduced hepatic PPAR-α function in the mouse also appears to be estrogen-dependent (Papacleovoulou, Abu-Hayyeh & Willamson, 2011).

   The above scenario has been studied extensively in mice (Milona et al., 2010; Aleksunes et al., 2012; Shuster et al., 2013), but little is known the FXR-SHP regulation of bile acid homeostasis in rats. Mice and rats are two most commonly used experimental animals, but some physiological responses are different. For example, in mice, Cyp7a1 and liver bile acid pool were not increased during lactation (Aleksunes et al., 2012), whereas the bile acid synthesis gene Cyp7a1 and hepatic bile acids are increased 2-3 fold in lactating rats (Wooton-Kee, Cohen & Vore, 2008; Wooton-Kee et al., 2010). In mice, pregnancy and lactation are associated with decreases in hepatic transporters, including bile acid transporters, but little is known whether such a phenomenon occurs in rats. This study was initiated to investigate bile acid metabolism and transport gene expressions in pregnant and lactating rats, and the results clearly demonstrate that under physiological conditions, PXR-SHP regulation plays important roles in bile acid homeostasis in pregnant rats.

Materials and Methods

Animals. Adult Sprague Dawley (SD) rats (250 g) were purchased from the Experimental

Animal Center of Third Military Medical College (Chongqing, China; certificate No CXK

2007-0005). Rats were kept in a SPF-grade animal facilities (certificate No SYXK 2011-004)

at Zunyi Medical College, with regulated environment (22 ± 1°C, 50 ± 2 % humidity and a 12

h: 12 h light: dark cycle) and free access to purified water and standard rodent chow. Rats

were acclimatized for 1 week, and subjected to timely mating overnight. A vaginal plug in the

next morning was designated as day 0 (GD 0) of gestation. Maternal livers were collected on

GD10, GD14 and GD19, as well as on the postnatal days (PND) 1, 7, 14 and 21. The

age-matched virgin rats were used as controls. Livers were weighed, snap frozen in liquid

nitrogen, and stored at -80°C until analysis. All animal procedures follow the NIH guide of

Humane Use and Care Animals, and approved by Institutional Animal Use and Care

Committee of Zunyi Medical College (2012-07).

Bile acid determination

Bile acids were extracted from the liver and measured with the total bile acid assay (TBA)

kit (Nanjing Jian-Cheng Bioengineering Co., China). Briefly, livers were homogenized in

physiological saline (1:9, wt :vol), followed by centrifugation at 2500 rpm/min for 10 min.

the supernatant (30 µl) was taken for determination of bile acids according to manufacturer's

protocol.

RNA Isolation and real-time RT-PCR analysis

Total RNA was isolated from frozen liver sample (50-100 mg) using 1 ml TRIzol (Takara,

Biotechnology, Dalian, China) and subsequently purified with Total RNA (Mini) Kit (Watson

Biotechnology, Shanghai, China). The quality of purified RNA was determined by

spectrophotometry with the 260/280 ratio >1.8. Purified RNA was reversed transcribed with

the High Capacity Reverse Transcriptase Kit (Applied Biosystems, Foster City, CA, USA).

The primer pairs were designed with the Primer3 software and listed in Supplementary Table

1. The Power SYBR Green Master Mix (Applied Biosystems, Foster City, CA, USA) was

used for real-time RT-PCR analysis. The cycle time for reaching threshold (Ct) of each target

gene was normalized to the housekeeping genes (G3PDH and β-actin), and expressed as % of

housekeeping genes.

Western Blot Analysis

Livers were homogenized in RIPA lysis buffer (Beyotime, P0013B, Shanghai, China)

containing freshly-prepared proteinase inhibitors. The supernatants were centrifuged at 12000

rpm 10 min at 4°C, and protein concentrations were quantified by the BCA assay (Beyotime,

P0012, Shanghai, China). Aliquot proteins were denatured with protein loading buffer

(Beyotime, P0015, Shanghai, China), and approximately 50 µg of protein/lane was separated

on 10% SDS-PAGE and transferred to PVDF membranes. Membranes were blocked in 5%

non-fat milk in TBST, followed by incubation overnight at 4 °C with 1:1000 CYP7A1

(ab65586) or β-actin (Ab8227) from Abcam (Cambridge, MA) in 1% BSA. After washing

with TBST three times, the membranes were incubated with HRP-conjugated anti-rabbit or

anti-mouse IgG (Beyotime, A0208 and A0216, Shanghai, China). Protein-antibody

complexes were visulized using an enhanced chemiluminescent reagent (ECL-Plus) (Beyotime, P0018, Shanghai, China), and exposed to Gel Imaging (Bio-Rad, ChemiDoc XRS, USA). The intensity of the band was semi-quantified with Quantity One software.

Statistical Analysis

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

Results

Liver bile acid levels in pregnant and lactating rat

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

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

Expression of the classic pathway bile acid synthetic enzyme genes (Cyp7a1 and 8b1) and alternative pathway (Cyp27a1 and 7b1), is shown in Fig.2. The expression of rate-limiting Cyp7a1 mRNA was unchanged during pregnancy, and increased on postpartum. Cyp8b1 mRNA decreased in GD10 and GD14, and increased about 2-fold in PND14. The expression of alternative pathway genes Cyp27a1 and Cyp7b1 were unchanged in gestation days and increased in postnatal days.

Hepatic expression of bile acid synthetic rate-limiting protein Cyp7A1 in pregnant and

postpartum rat.

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

Hepatic mRNA expression of nuclear receptors FXR, SHP, and ESR-1, PPAR-α in pregnant and postpartum rat

The expression of bile acid regulation nuclear receptor genes farnesoid X receptor (FXR, NR1H4) did not show significant increases during pregnancy, while FXR gradually increased on postpartum. The small heterodimer partner (SHP; NR0B2) significantly increased in the late gestational days, increased 3-fold on GD 19 as compared to controls. FXR plays an important role in bile acid homeostasis by inducing the transcription repressor SHP (Chiang, 2009). Estrogen receptor alpha (ESR-1) decreased to 64.7%, 57.7% on GD10 and GD14. In postnatal days, ESR-1 increased 2.33-fold in PND1 and then decreased to 68% of control on PND21. Proliferator-activated receptor α (PPARα) increased 3.79-fold compared to controls during lactation.

Hepatic mRNA expression of bile acid transporters in pregnant and postpartum rats.

As illustrated in Fig 5, the expression of bile acid uptake transporter Na+-taurocholate co-transporting polypeptide (Ntcp) and the efflux transporter bile salt export pump (Bsep/ABCB11) were decreased during pregnancy but increased during lactation. The

multidrug resistance protein 3 (Mrp3) and Mrp4 showed the similar pattern, with slightly increase during lactation.

Hepatic mRNA expression of canalicular uptake OATP transporters and efflux transporter Abcg2 in livers of pregnant and postpartum rats.

Figure 6 demonstrates that the expression of canalicular uptake transporter solute carrier organic anion transporter (Oatp1/Slco1a1), solute carrier organic anion transporter (Oatp2/Slco1b2), and organic anion-transporting polypeptide 4 (Oatp4/Slc21a10) were all decreased in the gestation days, on PND1 Oatp1 increased 1.68-fold, then decreased in postnatal days. In comparison, Oatp2 and Oatp4 decreased in the both gestation days and lactation days. The ATP-binding cassette sub-family G member 2 (Abcg2/BCRP) also decreased in the gestation and postnatal days expect PND1.

Discussion

The present study demonstrates that in pregnant rats, hepatic bile acids were not elevated. Consistent with hepatic bile acid concentrations, bile acid synthesis enzymes, i.e., Cyp7a1, Cyp8b1, Cyp27a1 and Cyp7b1 were not increased during pregnancy. Although FXR was unchanged, dramatic increases in SHP would be responsible for bile acid homeostasis during pregnancy. In comparison, lactating rats had increased liver bile acid, increased bile acid synthetic enzymes, and increased bile acid transporters. In general, OATP transporters were down-regulated during pregnancy and lactation in rats. These results fill the gap to add our understanding of FXR-SHP regulation of bile acid homeostasis and transport in rats during pregnancy and lactation.

ICP is characterized by raised serum bile acid levels and abnormal liver function tests

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

ICP has a complex etiology including genetic factors, endocrine factors, and the impact of pregnancy on FXR function (Abu-Hayyeh, Papacleovoulou & Williamson, 2013a; Floreani, et, al., 2013). The present study focused on FXR function under physiological conditions. It is proposed that pregnancy in mice resembles a state of FXR inactivation (Milona et al., 2010; Aleksunes et al., 2012). Indeed, attenuated FXR function during mouse pregnancy has been reported (Papacleovoulou, Abu-Hayyeh & Williamson, 2011;Aleksunes et al., 2012) and the 3β-sulfated progesterone metabolite epiallopregnanolone sulfate was found to inhibit FXR, resulting in reduced FXR-mediated bile acid efflux (Abu-Hayyeh et al., 2013b). In the present study, the expression of FXR in rats during pregnancy was basically unchanged with a reduction trend during pregnancy (Fig. 4). However, the FXR-inducible negative target SHP was marked increased at GD14 and reached more than 3-fold higher at GD19. This marked

increased in SHP would be responsible for maintaining the bile acid synthesis homeostasis

and preventing the liver bile acids to accumulate to protect the fetus from the bile acid

toxicity. It should also be realized that estrogen receptor alpha (ERS-1) and the peroxisome

proliferator-activated receptorα (PPAR-α) during normal pregnancy were not altered in rats

(Fig. 4). Thus, the estrogen and FXR interactions may not be evident in rats as compared to

that in mice (Aleksunes et al., 2012).

Lactation is a time of a five-fold increase in energy demand, as suckling young requires a

proportional adjustment in the ability of the lactating dam to absorb nutrients (Cripps &

Williams, 1975; Vernon et al., 2002). Lactating rats have a two to three-fold increase in food

consumption to ensure lactating dams to absorb nutrients and to synthesize critical molecules

including bile acids to meet the dietary needs of the offspring and the dam (Vernon et al.,

2002). The size and hydrophobicity of the bile acid pool increase during lactation, implying

an increased absorption and disposition of lipid, sterols, nutrients, and xenobiotics

(Athippozhy et al., 2011). In essence, rats (Wooton-Kee, Cohen &Vore, 2008) are different

from mice (Aleksunes et al., 2012) in bile acid homeostasis during lactation. In the present

study, hepatic bile acid pool (Fig. 1), bile acid synthesis gene Cyp7a1, Cyp8b1, Cyp27a1 and

Cyp7b1 (Fig. 2 and Fig. 3) were all increased during lactation, consistent with this scenario.

The bile acid transporters Ntcp and Bsep followed the similar pattern. Ntcp is the major

bile acid transporter for conjugated bile acid (Csanaky et al., 2011) and Bsep is the major bile

acid efflux pump located at the bile canalicular apical domain of hepatocytes (Lam, Soroka &

Boyer, 2010). Down-regulation of Ntcp and Bsep was observed in pregnant rats (Arrese et

al., 2003; Cao et al., 2001), however, they are increased on postpartum, probably under the

influence of prolactin (Cao et al., 2001). Sulfated progesterone metabolite (P4-S) levels are raised in normal pregnancy and elevated further in ICP, which can cause a competitive inhibition of NTCP-mediated uptake of taurocholate in Xenopus oocytes (Abu-Hayyeh et al., 2010). In the present study, both Ntcp and Besp (Fig. 5) were lower during pregnancy, and Bsep was increased during lactation, consistent with liver bile acid homeostasis profile. Mrp3 and Mrp4 are two major bile acid efflux (Cui et al., 2009; Aleksunes et al., 2012), and their expression showed the similar pattern (Fig. 5), i.e., lower during the pregnancy and higher during lactation. The pattern of these transporter mRNA levels coincide with FXR-SHP regulation of bile acid homeostasis, and fortifying the concept that under physiological conditions, SHP-regulation of bile acid synthesis is essential for maintaining the bile acid homeostasis to avoid the occurrence of ICP, an unusual pathological condition.

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

similar to Oatps, and can also be envisioned as an adaptive mechanism during pregnancy and lactation.

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

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Competing Interests

Jie Liu is an Academic Editor for PeerJ. There are no other competing interests.

Author Contributions

_ Hongmei Xie, Qiongni Zhu and Dan Zhang performed the experiments, analyzed the data.

_ Yuanfu Lu, Jie Liu conceived and designed the experiments, contributed

reagents/materials/analysis tools, wrote the paper.

_ Qiongni Zhu, Hongmei Xie, Yuanfu Lu and Jie Liu wrote the paper.

Animal Ethics

The following information was supplied relating to ethical approvals (i.e. approving body and any reference numbers):
All animal procedures follow the NIH guide of Humane Use and Care Animals, and approved

by Institutional Animal Use and Care Committee of Zunyi Medical College

References

Abu-Hayyeh S, Martinez-Becerra P, Sheikh Abdul Kadir SH, Selden C, Romero MR, Rees M, Marschall HU, Marin JJ, Williamson C.2010. Inhibition of Na+-taurocholate Co-transporting polypeptide-mediated bile acid transport by cholestatic sulfated progesterone metabolites. The Journal of Biological Chemistry **285**:16504-16512.

Abu-Hayyeh S, Papacleovoulou G, Williamson C.2013a.Nuclear receptors, bile acids and cholesterol homeostasis series - Bile acids and pregnancy. Molecular and Cellular Endocrinology **368**:120-128.

Abu-Hayyeh S, Papacleovoulou G, Lövgren-Sandblom A, Tahir M, Oduwole O, Jamaludin NA, Ravat S, Nikolova V, Chambers J, Selden C, Rees M, Marschall HU, Parker MG, Williamson C. 2013b.Intrahepatic cholestasis of pregnancy levels of sulfated progesterone metabolites inhibit farnesoid X receptor resulting in a cholestatic phenotype. Hepatology **57**:716-726.

Aleksunes LM, Yeager RL, Wen X, Cui JY, Klaassen CD. 2012.Repression of hepatobiliary transporters and differential regulation of classic and alternative bile acid pathways in mice during pregnancy. Toxicological Sciences**130**:257-268.

Anzivino C, Odoardi MR, Meschiari E, Baldelli E, Facchinetti F, Neri I, Ruggiero G, Zampino R, Bertolotti M, Loria P, Carulli L. 2013. ABCB4 and ABCB11 mutations in intrahepatic cholestasis of pregnancy in an Italian population. Digestive and Liver Disease **45**:226-232.

Arrese M, Trauner M, Ananthanarayanan M, Pizarro M, Solís N, Accatino L, Soroka C, Boyer JL, Karpen SJ, Miquel JF, Suchy FJ. 2003. Down-regulation of the Na+/taurocholate cotransporting polypeptide during pregnancy in the rat. Journal of Hepatology **38**:148-155.

Arrese M, Macias RI, Briz O, Perez MJ, Marin JJ. 2008.Molecular pathogenesis of intrahepatic cholestasis of pregnancy. Expert Reviews in Molecular Medicine **28:**10:e9.

Athippozhy A, Huang L, Wooton-Kee CR, Zhao T, Jungsuwadee P, Stromberg AJ, Vore M. 2011. Differential gene expression in liver and small intestine from lactating rats compared to age-matched virgin controls detects increased mRNA of cholesterol biosynthetic genes. BMC Genomics **3:**12:95.

Barth A, Rost M, Kindt A, Peiker G. 2005.Serum bile acid profile in women during pregnancy and childbed. Experimental and Clinical Endocrinology & Diabetes **113:**372-375.

Blazquez AG, Briz O, Romero MR, Rosales R, Monte MJ, Vaquero J, Macias RI, Cassio D, Marin JJ. 2012. Characterization of the role of ABCG2 as a bile acid transporter in liver and placenta. Molecular Pharmacology **81**:273-283.

Cao J, Huang L, Liu Y, Hoffman T, Stieger B, Meier PJ, Vore M. 2001. Differential regulation of hepatic bile salt and organic anion transporters in pregnant and postpartum rats and the role of prolactin. Hepatology **33**:140-147.

Cao J, Stieger B, Meier PJ, Vore M. 2002.Expression of rat hepatic multidrug resistance-associated proteins and organic anion transporters in pregnancy. American Journal of Physiology Gastrointestinal and Liver Physiology **283**:G757-G766.

Carlin A, Alfirevic Z.2008.Physiological changes of pregnancy and monitoring. Best

Practice & research. Clinical Obstetrics &Gynaecology **22:**801-823.

Chiang JY. 2009. Bile acids: regulation of synthesis. Journal of Lipid Research **50:**1955-1966.

Cripps AW, Williams VJ. 1975. The effect of pregnancy and lactation on food intake, gastrointestinal anatomy and the absorptive capacity of the small intestine in the albino rat. The British Journal of Nutrition **33:**17-32.

Cui YJ, Aleksunes LM, Tanaka Y, Goedken MJ, Klaassen CD. 2009. Compensatory induction of liver efflux transporters in response to ANIT-induced liver injury is impaired in FXR-null mice. Toxicological Sciences **110:** 47-60.

Csanaky IL, Lu H, Zhang Y, Ogura K, Choudhuri S, Klaassen CD. 2011. Organic anion-transporting polypeptide 1b2 (Oatp1b2) is important for the hepatic uptake of unconjugated bile acids: Studies in Oatp1b2-null mice. Hepatology **53:**272-281.

Diken Z,Usta IM, Nassar AH. 2013. A Clinical Approach to Intrahepatic Cholestasis of Pregnancy. American Journal of Perinatology 2013 Jan 28 [Epub ahead of print].

Dixon PH, Weerasekera N, Linton KJ, Donaldson O, Chambers J, Egginton E, Weaver J, Nelson-Piercy C, de Swiet M, Warnes G, Elias E, Higgins CF, Johnston DG, McCarthy MI, Williamson C. 2000. Heterozygous MDR3 missense mutation associated with intrahepatic cholestasis of pregnancy: evidence for a defect in protein trafficking. Human Molecular Genetics **9:**1209-1217.

Dixon PH, van Mil SW, Chambers J, Strautnieks S, Thompson RJ, Lammert F, Kubitz R, Keitel V, Glantz A, Mattsson LA, Marschall HU, Molokhia M, Moore GE, Linton KJ, Williamson C. 2009. Contribution of variant alleles of ABCB11 to susceptibility to intrahepatic cholestasis of pregnancy. Gut **58:**537-544.

Egan N, Bartels A, Khashan AS, Broadhurst DI, Joyce C, O'Mullane J, O'Donoghue K. 2012. Reference standard for serum bile acids in pregnancy. BJOG **119**:493-8.

Floreani A, Caroli D, Lazzari R, Memmo A, Vidali E, Colavito D, D'Arrigo A, Leon A, Romero R, Gervasi M. 2013. Intrahepatic Cholestasis Of Pregnancy: New Insights Into Its Pathogenesis. The Journal of Maternal-Fetal& Neonatal Medicine. 2013 Mar 12 [Epub ahead of print]

Geenes V, Williamson C. 2009. Intrahepatic cholestasis of pregnancy. World Journal of Gastroenterology**15**:2049-2066.

Lam P, Soroka CJ, Boyer JL. 2010. The bile salt export pump: clinical and experimental aspects of genetic and acquired cholestatic liver disease. Seminars in Liver Diseases **30**:125-133.

Lu H, Choudhuri S, Ogura K, Csanaky IL, Lei X, Cheng X, Song PZ, Klaassen CD. 2008. Characterization of organic anion transporting polypeptide 1b2-null mice: essential role in hepatic uptake/toxicity of phalloidin and microcystin-LR. Toxicological Sciences **103**:35-45.

Milona A, Owen BM, Cobbold JF, Willemsen EC, Cox IJ, Boudjelal M, Cairns W, Schoonjans K, Taylor-Robinson SD, Klomp LW, Parker MG, White R, van Mil SW, Williamson C. 2010. Raised hepatic bile acid concentrations during pregnancy in mice are associated with reduced farnesoid X receptor function. Hepatology **52:**1341-1349.

Papacleovoulou G, Abu-Hayyeh S, Williamson C.2011. Nuclear receptor-driven alterations in bile acid and lipid metabolic pathways during gestation. Biochimica et Biophysica

Acta **1812:**879-887.

Shuster DL, Bammler TK, Beyer RP, Macdonald JW, Tsai JM, Farin FM, Hebert MF, Thummel KE, Mao Q. 2013. Gestational age-dependent changes in gene expression of metabolic enzymes and transporters in pregnant mice. Drug Metabolism and Disposition **41:**332-342.

Van Mil SW, Milona A, Dixon PH, Mullenbach R, Geenes VL, Chambers J, Shevchuk V, Moore GE, Lammert F, Glantz AG, Mattsson LA, Whittaker J, Parker MG, White R, Williamson C. 2007. Functional variants of the central bile acid sensor FXR identified in intrahepatic cholestasis of pregnancy. Gastroenterology **133:**:507-516.

Vernon RG, Denis RG, Sorensen A, Williams G. 2002.Leptin and the adaptations of lactation in rodents and ruminants. Hormone Metabolic Research **34:**678-685.

Wooton-Kee CR, Cohen DE, Vore M.2008. Increased cholesterol 7alpha-hydroxylase expression and size of the bile acid pool in the lactating rat. AmericanJournal of Physiology Gastrointestinal and Liver Physiology **294:**G1009-G1016.

Wooton-Kee CR, Coy DJ, Athippozhy AT, Zhao T, Jones BR, Vore M. 2010. Mechanisms for increased expression of cholesterol 7alpha-hydroxylase (Cyp7a1) in lactating rats. Hepatology **51:** 277-285.

Zhang Y, Csanaky IL, Cheng X, Lehman-McKeeman LD, Klaassen CD.2012. Organic anion transporting polypeptide 1a1 null mice are sensitive to cholestatic liver injury. Toxicological Sciences **127:**451-462.

Figure Legends:

Figure 1. Liver bile acid levels in pregnant and lactating rat. Bile acids were quantified in Livers from control and pregnant rat on GD10, 14, and 19 and PND 1, 7, 14, and 21. Dark gray bars represent pregnant rat, and black bars represent lactating rat. Data are presented as mean ± SEM. Asterisks *represent statistically significant difference (p< 0.05) compared with control.

Figure 2. Hepatic mRNA expression of bile acid synthetic pathway genes in pregnant and lactating rat. The expression of bile acid synthetic classic pathway genes Cyp7a1, Cyp8b1 and alternative pathway genes Cyp27a1 and Cyp7b1 was quantified from control and GD10,14 and 19 and PND 1,7,14 and 21, Data were normalized to controls (set to 100%) and presented as mean ± SEM, Dark gray bars represent pregnant rat, and black bars represent lactating rat. Asterisks *represent statistically significant difference (p< 0.05) compared with control.

Figure 3. Hepatic expression of bile acid synthesis rate-limiting protein CYP7A1 in pregnant and postpartum 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 ± SEM. Dark gray bars represent pregnant rat, and black bars represent lactating rat. Significantly difference was confirmed by two-tailed independent Samples test method (P<0.05)

Figure 4. Hepatic mRNA expression of Nuclear Receptors SHP, FXR and ESR-1 and

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

Figure 5. Hepatic mRNA expression of bile acid transporter in pregnant and postpartum rats. The expression of bile acid uptake transporter Ntcp and efflux transporter Bsep as well as Mrp3 and Mrp4 was quantified using total hepatic RNA from pregnant rats on GD 10, 14 and 19, and postpartum rats on PND 1, 7, 14 and 21. Data were normalized to controls and presented as mean ± SEM. Dark gray bars represent pregnant rat, and black bars represent lactating rat. Asterisks (*) represent statistically significant differences (p<0.05) compared control.

Figure 6. Hepatic mRNA expression of canalicular uptake transporter Oatps and efflux transporter Abcg2 in the liver of pregnant and postpartum rats. The expression of uptake Oatp transporter sand Abcg2 was quantified using total hepatic RNA from pregnant rats on GD10, 14 and 19, and postpartum rats on PND 1, 7, 14 and 21. Data were normalized to controls and presented as mean ± SEM. Dark gray bars represent pregnant rat, and black bars represent lactating rat. Asterisks (*) represent statistically significant differences (p < 0.05) compared control.

Figure 1

Fig. 1

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Fig.1
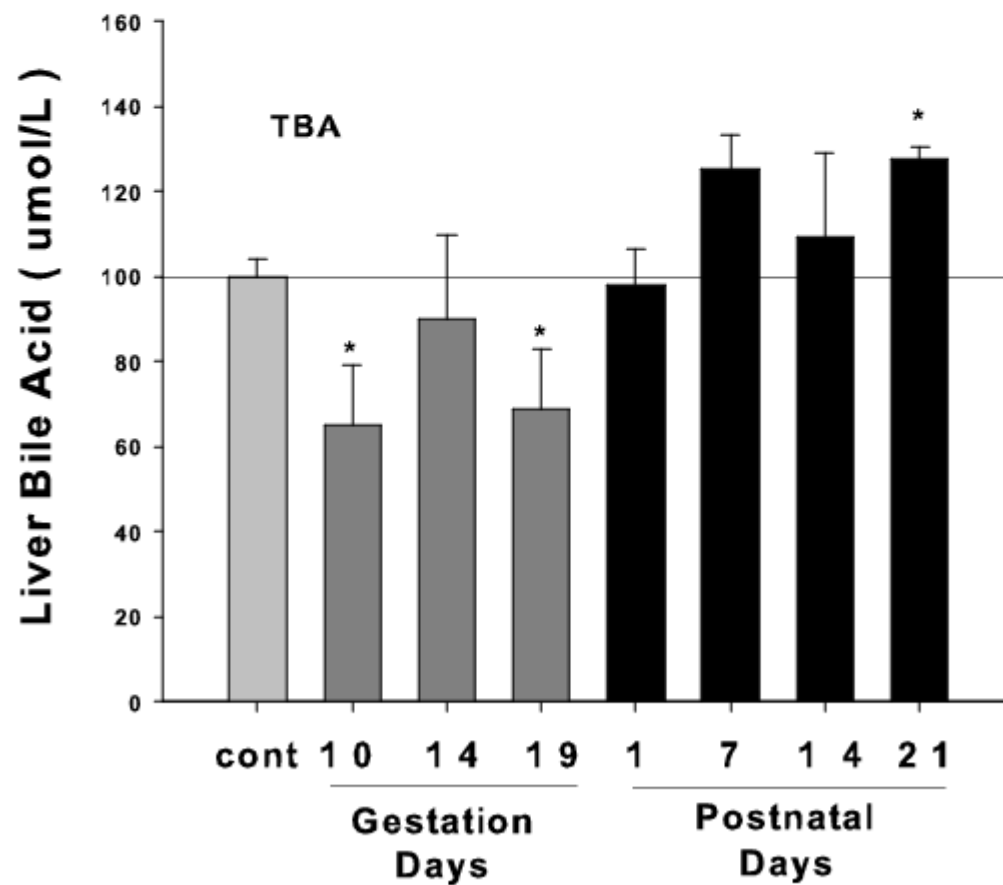
Figure 2

Fig. 2

Figure 2. Hepatic mRNA expression of bile acid synthetic pathway genes in pregnant and lactating rat. The expression of bile acid synthetic classic pathway genes Cyp7a1, Cyp8b1 and alternative pathway genes Cyp27a1 and Cyp7b1 was quantified from control and GD10,14 and 19 and PND 1,7,14 and 21, Data were normalized to controls (set to 100%) and presented as mean ± SEM, Dark gray bars represent pregnant rat, and black bars represent lactating rat. Asterisks 〔 * 〕 represent statistically significant difference ($p < 0.05$) compared with control.

Fig.2
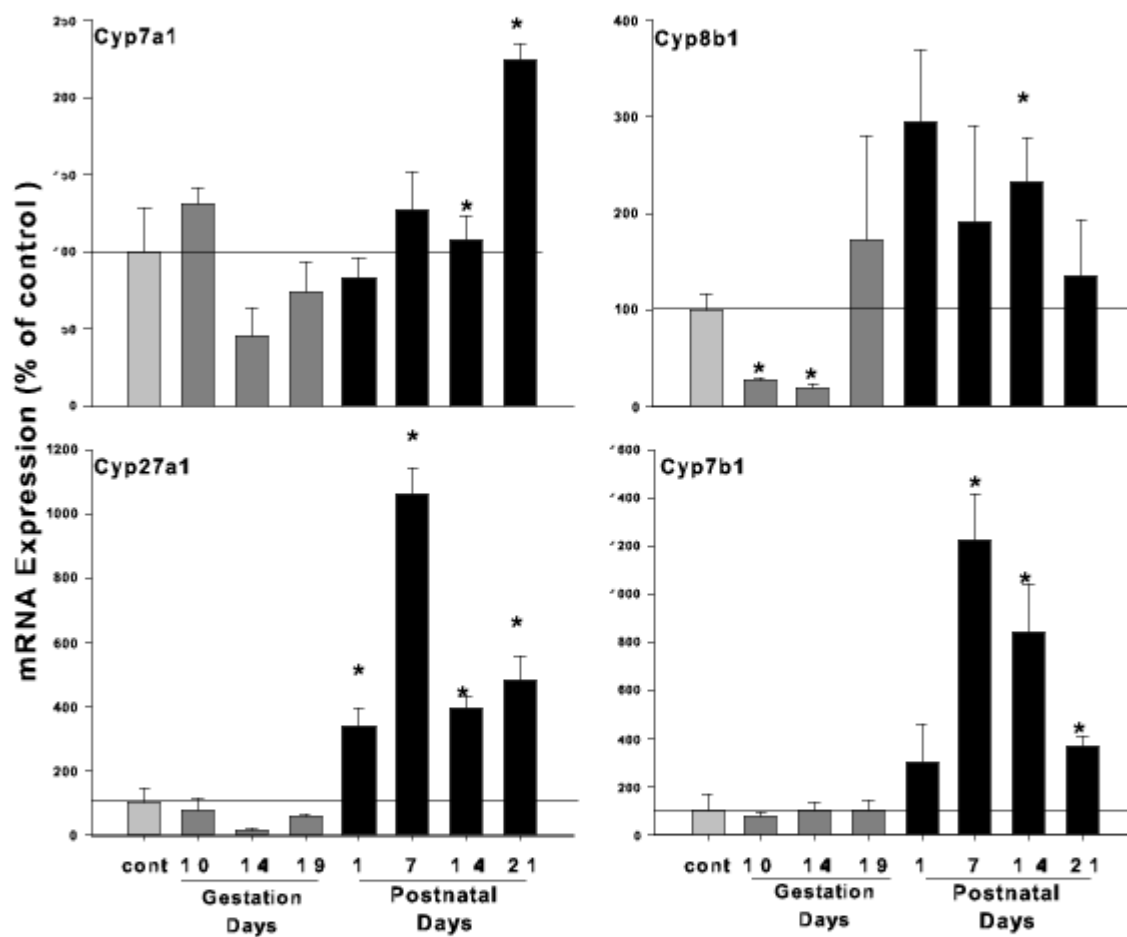
Figure 3

Fig. 3

Figure 3. Hepatic expression of bile acid synthesis rate-limiting protein CYP7A1 in pregnant and postpartum 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 ± SEM. Dark gray bars represent pregnant rat, and black bars represent lactating rat. Significantly difference was confirmed by two-tailed independent Samples test method (P<0.05)
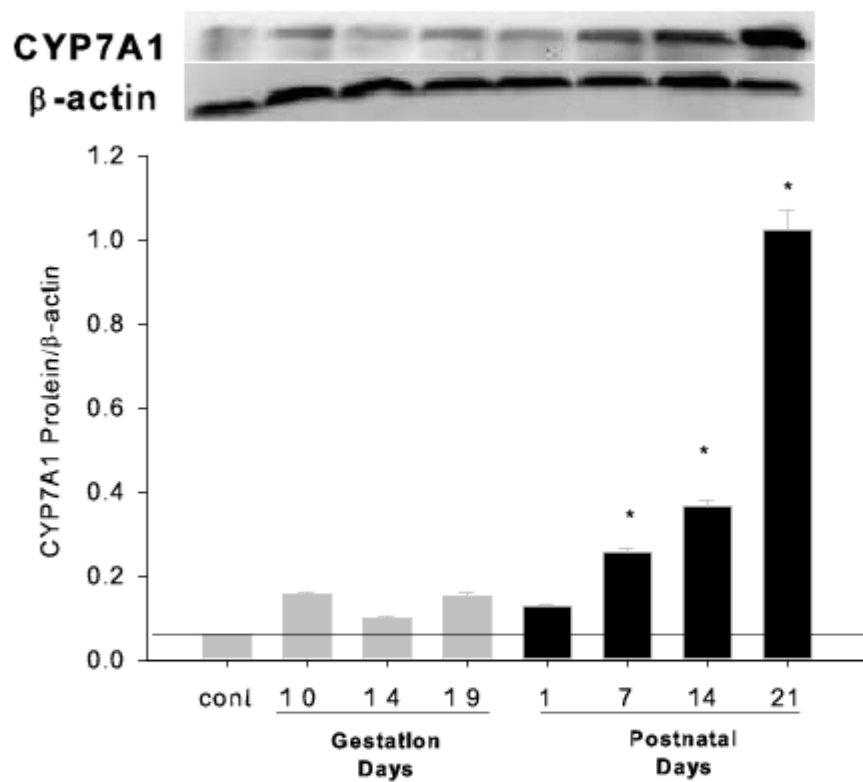
Figure 4

Fig. 4

Figure 4. Hepatic mRNA expression of Nuclear Receptors SHP, FXR and ESR-1 and PPAR-α in pregnant and postpartum rat. The expression of bile acid regulation Nuclear Receptors genes SHP, FXR and Esr-1,PPAR-α were quantified using total hepatic RNA from control and pregnant mice at gestational days 10, 14, 19 and postnatal days 1, 7, 14, 21. Data were normalized to controls and presented as mean ± SEM. Dark gray bars represent pregnant rat, and black bars represent lactating rat. Asterisks (*) represent statistically significant differences ($p < 0.05$) compared to control.
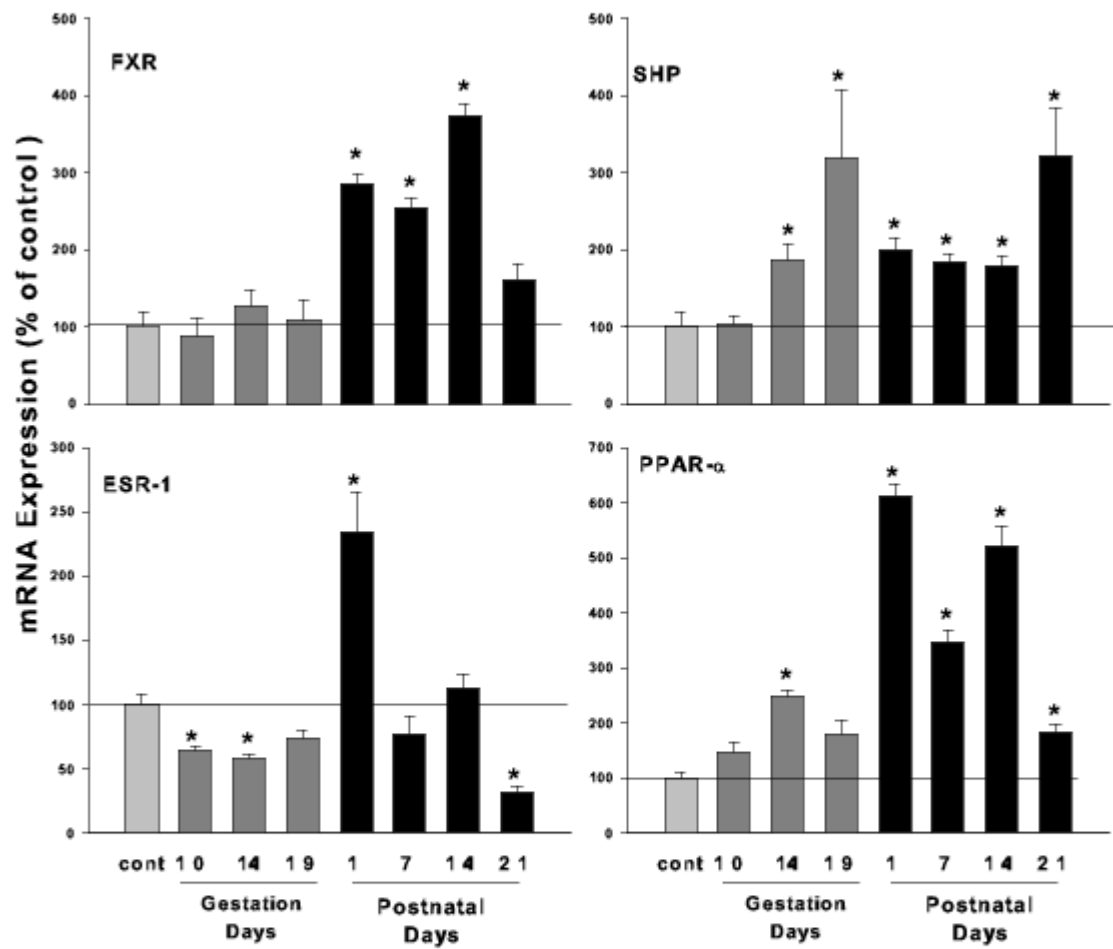
Fig.4

Figure 5

Fig. 5

Figure 5. Hepatic mRNA expression of bile acid transporter in pregnant and postpartum rats.
The expression of bile acid uptake transporter Ntcp and efflux transporter Bsep as well as Mrp3 and
Mrp4 was quantified using total hepatic RNA from pregnant rats on GD 10, 14 and 19, and postpartum
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Data were normalized to controls and presented as mean ± SEM. Dark gray bars represent pregnant
rat, and black bars represent lactating rat. Asterisks (*) represent statistically significant differences
($p<0.05$) compared control.

Fig.5

Figure 6

Fig. 6

Figure 6. Hepatic mRNA expression of canalicular uptake transporter Oatps and efflux transporter Abcg2 in the liver of pregnant and postpartum rats. The expression of uptake Oatp transporter sand Abcg2 was quantified using total hepatic RNA from pregnant rats on GD10, 14 and 19, and postpartum rats on PND 1, 7, 14 and 21. Data were normalized to controls and presented as mean ± SEM. Dark gray bars represent pregnant rat, and black bars represent lactating rat. Asterisks (*) represent statistically significant differences (p < 0.05) compared control.

Fig.6