



Identification of Transmembrane Protease Serine 2 and Forkhead Box A1 As the Potential Bisphenol A Responsive Genes in the Neonatal Male Rat Brain

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Ubuka T, Moriya S, Soga T and Parhar I (2018) Identification of Transmembrane Protease Serine 2 and Forkhead Box A1 As the Potential Bisphenol A Responsive Genes in the Neonatal Male Rat Brain. Front. Endocrinol. 9:139. doi: 10.3389/fendo.2018.00139 Perinatal exposure of Bisphenol A (BPA) to rodents modifies their behavior in later life. To understand how BPA modifies their neurodevelopmental process, we first searched for BPA responsive genes from androgen and estrogen receptor signaling target genes by polymerase chain reaction array in the neonatal male rat brain. We used a transgenic strain of Wistar rats carrying enhanced green fluorescent protein tagged to gonadotropin-inhibitory hormone (GnIH) promoter to investigate the possible interaction of BPA responsive genes and GnIH neurons. We found upregulation of transmembrane protease serine 2 (Tmprss2), an androgen receptor signaling target gene, and downregulation of Forkhead box A1 (Foxa1), an ER signaling target gene, in the medial amygdala of male rats that were subcutaneously administered with BPA from day 1 to 3. Tmprss2immunoreactive (ir) cells were distributed in the olfactory bulb, cerebral cortex, hippocampus, amygdala, and hypothalamus in 3 days old but not in 1-month-old male rats. Density of Tmprss2-ir cells in the medial amygdala was increased by daily administration of BPA from day 1 to 3. Tmprss2 immunoreactivity was observed in 26.5% of GnIH neurons clustered from the ventral region of the ventromedial hypothalamic nucleus to the dorsal region of the arcuate nucleus of 3-day-old male rat hypothalamus. However, Tmprss2 mRNA expression significantly decreased in the amygdala and hypothalamus of 1-month-old male rats. Foxa1 mRNA expression was higher in the hypothalamus than the amygdala in 3 days old male rats. Intense Foxa1-ir cells were only found in the peduncular part of lateral hypothalamus of 3-day-old male rats. Density of Foxa1-ir cells in the hypothalamus was decreased by daily administration of BPA from day 1 to 3. Foxa1 mRNA expression in the hypothalamus also significantly decreased at 1 month. These results suggest that BPA disturbs the neurodevelopmental process and behavior of rats later in their life by modifying *Tmprss2* and *Foxa1* expressions in the brain.

Keywords: transmembrane protease serine 2, forkhead box A1, androgen receptor, estrogen receptor, polymerase chain reaction array, gonadotropin-inhibitory hormone neurons

INTRODUCTION

Bisphenol A (BPA) is an organic synthetic compound widely used to make polycarbonate plastics and epoxy resins utilized in reusable food and drink containers and inner lining of cans and bottles (1). Hydrolysis of the ester-bond linking BPA can occur at high temperature, and BPA can be leached out into food and beverages. Fetuses or young animals during their developmental stages are thought to be susceptible to BPA exposure than adults (2). Therefore, detectable level of BPA in 88% of human cord blood samples (3) generated social concerns about the effect of BPA on the fetuses.

Dodds and Lawson were the first to show the estrogenic property of BPA (4). However, the estrogenic effect of BPA is 10,000 times less potent than estradiol-17 β shown by uterine vascular permeability assay in ovariectomized mice (5). Similar level of agonistic activity of BPA to estrogen receptor (ER) was shown in yeast expressing human estrogen or androgen receptor (AR) (6). On the other hand, BPA is a potent ligand for the nonclassic membrane bound G protein-coupled receptor for estrogen (GPR30) (7, 8). BPA did not have an agonistic activity but had an antagonistic activity to AR in the same yeast-based assay (6). Stronger antagonistic activity of BPA to human AR was shown in an African monkey kidney cell line (9). It is also known that BPA can disturb steroidogenesis by interfering the activity of steroidogenic enzymes, such as CYP450scc, 36HSD, and CYP450arom (10). Perera et al. (11) examined the association between prenatal BPA exposure and child behavior. They found that high prenatal BPA exposure was associated with higher emotionally reactive and aggressive behavior syndromes in boys and lower anxious/ depressed and aggressive behavior in girls (11). There are numerous studies showing the effect of perinatally administered BPA on social behavior of rodents. BPA exposure during pregnancy increased display of nose-to-nose contacts, play solicitations, and approaches in both sexes (12). Administration of BPA during pregnancy or lactation increased defensive behavior in male and sexual behavior in female rats (13). In the other study, BPA exposed males during gestation and lactation showed persistent deficits in sexual behavior in adulthood (14).

To find BPA responsive genes in the brain mediating its effect on social behavior, we first focused on amygdala, because amygdala is the integrative center for the processing of emotion, which is pivotal for social behavior. It is thought that deficits in the development of amygdala may cause autism, a neurodevelopmental disorder that is characterized by impaired social interaction (the amygdala theory of autism) (15). It is also known that these brain nuclei express ARs and ERs and abnormally enlarge in autism infants (16). We subcutaneously administered a reference dose of BPA (50 µg/kg BW/day) determined by United States Environmental Protection Agency (EPA, www.epa.gov/ iris/subst/0356.htm) to neonatal male transgenic rats carrying enhanced green fluorescent protein tagged to gonadotropininhibitory hormone [EGFP-GnIH (17)] promoter from day 1 to 3 and differential expressions of AR and ER signaling target genes in the medial amygdala were analyzed by polymerase chain reaction (PCR) array. GnIH is a hypothalamic neuropeptide that decreases gonadotropin secretion directly acting on the pituitary

or by decreasing the activity of gonadotropin-releasing hormone (GnRH) neurons (18, 19). Recent studies have shown that GnIH neurons also decrease motivated behavior (20, 21). We found upregulation of transmembrane protease serine 2 (*Tmprss2*) and downregulation of Forkhead box A1 (*Foxa1*) mRNAs in the amygdala by BPA exposure. We further studied the location of Tmprss2 and Foxa1 immunoreactive (ir) cells, effect of BPA on the density of Tmprss2 and Foxa1 mRNA expressions in the male rat brain.

MATERIALS AND METHODS

Animals

EGFP-GnIH Wistar rats (17) were housed under a controlled 12 h light/dark cycle (light on at noon) and the temperature was maintained at 22°C in a specific pathogen free animal facility. All rats had free access to autoclaved food and water. The pups were counted after parturition, weighed, and remained with their biological mother. All procedures were approved by Monash University, Animal Ethics Committee (MARP/2016/037).

PCR Array

Bisphenol A at 50 µg/kg or vehicle (sesame oil) was subcutaneously injected to neonatal male rats after dawn daily from postnatal day 1 (P1) to P3. 2 h after the last injection the rats were deeply anesthetized by Zoletil/Ketamine/Xylazine (Z/K/X) at 13.5 mg/90 µl/kg and brains were collected and stored at -80°C. Medial amygdaloid tissue from both sides of the brain was dissected in a cryostat at -20° C by referring to a rat brain atlas (22). Medial amygdaloid tissues from two male siblings were combined in a tube for homogenization. Total medial amygdaloid tissue samples of six BPA treated rats from three different littermates and six vehicle treated rats from three different littermates were collected. Total RNA was extracted by using an RNA isolation kit (RNeasy mini kit; QIAGEN, Hilden, Germany) and reverse transcribed by using High-Capacity cDNA Reverse Transcription Kit (ThermoFisher SCIENTIFIC, Waltham, MA, USA). Differential gene expression was analyzed by PCR arrays, RT² ProfilerTM PCR Array Rat AR Signaling Targets, and RT² ProfilerTM PCR Array Rat ER Signaling Targets (QIAGEN). These PCR array kits analyze the expression levels of 84 AR signaling target genes (Table S1 in Supplementary Material) and 84 ER signaling target genes (Table S2 in Supplementary Material). Cbp/p300-interacting transactivator with Glu/Asp-rich carboxyterminal domain 2 (Cited2, NM_053698), insulin-like growth factor binding protein 5 (Igfbp5, NM_012817), kallikrein B plasma 1 (Klkb1, NM_012725), and myelocytomatosis oncogene (Myc, NM_012603) were analyzed in both assays. Real-time PCR was performed using a StepOnePlus[™] Real-Time PCR System (ThermoFisher SCIENTIFIC) with conditions of 95°C for 10 min, 40 cycles of 95°C for 15 s, and 60°C for 1 min, followed by a dissociation step according to the manufacturer's instruction. Expression levels of AR signaling target genes were normalized by the mean expression levels of housekeeping genes (HKGs), actin beta (Actb, NM_031144), Hypoxanthine

BPA Responsive Genes in Brain

TABLE 1 | Effect of Bisphenol A (BPA) on androgen receptor (AR) and estrogen receptor (ER) signaling target gene expression in the medial amygdala of neonatal male rats.

Gene symbol Fold changes P value symbol Gene changes Fold changes P value changes Abcc4 1.10 0.627 Adora1 1.09 0.735 Abhd2 1.12 0.508 Ahr 1.11 0.837 Acsl3 0.99 0.913 Akap1 1.13 0.634 Adamts1 0.91 0.751 Apbb1 1.11 0.6634 Apopb2 0.99 0.944 Bcl211 1.08 0.634 Ar 0.95 0.434 Bdnf 1.07 0.839 Camkk2 1.13 0.655 Bmp4 0.89 0.673 Camkk2 1.14 0.191 Bmp7 6.46 0.585 Cenpn 0.98 0.948 Brca1 0.62 0.571 Dhcr24 1.21 0.387 Cch11 0.98 0.944 Eaf2 0.56 0.0645 Cited2 1.60 0.839 Erla1 1.41 0.391 Ckb 0.97 0.	AR signaling target genes			ER signaling target genes			
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Lin 1.01 0.000 0.030 0.030 0.030 0.030 Ernf1 1.16 0.564 Cyp19a1 1.01 0.975 Fam105a 0.99 0.832 Cyp1a1 - - Fkbp5 1.08 0.571 Ebag9 15.05 0.266 Fos 0.80 0.252 Efna5 1.27 0.257 Fad5 0.99 0.795 Egr3 0.88 0.661 Gucy1a3 1.13 0.222 Erbb2 1.19 0.521 Herc3 1.01 0.974 Erbb3 1.52 0.992 Hpgd 1.17 0.482 Esr1 1.21 0.515 Igftr 1.08 0.733 Foxa1 0.32 0.0147* Jun 1.08 0.639 Fst 0.97 0.814 Kik1c2 2.43 0.135 G6pd 1.11 0.591 Kik4 0.90 0.354 Igftp5 0.98 0.867	FII2	1.01	0.935	Ctaf	0.76	0.359	
Int Int Otop Otop Otop Find 1.16 0.564 Cyp19a1 - - Fkbp5 1.08 0.571 Ebag9 15.05 0.266 Fos 0.80 0.252 Ena5 1.27 0.257 Fzd5 0.99 0.795 Egr3 0.88 0.651 Gucy1a3 1.13 0.222 Erbb2 1.19 0.521 Herc3 1.01 0.974 Erbb3 1.52 0.992 Hpgd 1.17 0.482 Esr1 1.21 0.515 Igftr 1.10 0.564 Esr2 0.86 0.606 Igfbp5 0.96 0.753 Fos 0.79 0.814 Klk1c2 2.43 0.135 G6pd 1.11 0.599 Klk4 0.95 0.994 Gper1 G 1.06 0.791 Klk4 0.90 0.304 Igfbp5 0.98 0.867 Lig1 1.00 0.9	Em1	1.01	0.392	Ctsd	1.06	0.899	
Link Link Gypen Link Gypen Fam105a 0.99 0.832 Cypen - - Fkbp5 1.08 0.571 Ebag9 15.05 0.266 Fos 0.80 0.252 Efna5 1.27 0.257 Fzd5 0.99 0.795 Egr3 0.88 0.661 Gucy1a3 1.13 0.222 Erbb2 1.19 0.521 Herc3 1.01 0.974 Erbb3 1.52 0.992 Hpgd 1.17 0.482 Esr1 1.21 0.515 Igf1r 1.10 0.564 Esr2 0.86 0.606 Igfbp5 0.96 0.753 Fost 0.97 0.814 Klk 1.08 0.733 Foxa1 0.32 0.0147* Jun 1.08 0.639 Fst 0.97 0.814 Klk1c2 2.43 0.135 G6pd 1.11 0.599 Klk4 0.95 0	Errfi1	1.21	0.564	Cvp19a1	1.00	0.000	
Tail Oda 0.33 0.357 0.351 0.357 0.351 0.557 0.351 0.557 0.351 1.21 0.515 0.366 0.606 0.357 Fos 0.79 0.351 Irs2 1.08 0.733 Foxa1 0.32 0.0147* Jun 1.08 0.639 Fst 0.97 0.814 Klk1c2 2.43 0.135 G6pd 1.11 0.599 Klk4 0.95 0.994 Gper1 G 1.06 0.791 Klkb1 - - Hsp00aa1 0.96 0.731 Krt8 1.06 0.503 Lama1 0.90 0.5451 Igftpp5 0.98	Eam105a	0.99	0.832	Cypital	-	0.070	
Hob Hob Hob Hob Hob Hob Hob Fos 0.80 0.252 Efna5 1.27 0.257 Fzd5 0.99 0.795 Egr3 0.88 0.661 Gucy1a3 1.13 0.222 Erbb2 1.19 0.521 Herc3 1.01 0.974 Erbb3 1.52 0.992 Hpgd 1.17 0.482 Esr1 1.21 0.515 Igftp 0.96 0.753 Fos 0.79 0.351 Irs2 1.08 0.733 Foxa1 0.32 0.0147* Jun 1.08 0.639 Fst 0.97 0.814 Klk1c2 2.43 0.135 G6pd 1.11 0.599 Klk4 0.95 0.994 Gper1 G 1.06 0.791 Klk4 0.90 0.545 Igfpp5 0.98 0.867 Lirg1 1.00 0.975 Irs1 1.12 0.629 Liff </td <td>Fkbn5</td> <td>1.08</td> <td>0.571</td> <td>EbadQ</td> <td>15.05</td> <td>0.266</td>	Fkbn5	1.08	0.571	EbadQ	15.05	0.266	
First 0.30 0.202 Endo 1.21 0.21 Fzd5 0.99 0.795 Egr3 0.88 0.661 Gucy1a3 1.13 0.222 Erbb2 1.19 0.521 Herc3 1.01 0.974 Erbb3 1.52 0.992 Hpgd 1.17 0.482 Esr1 1.21 0.515 Igftr 1.10 0.564 Esr2 0.86 0.6061 Igfbp5 0.96 0.753 Fos 0.79 0.814 Jun 1.08 0.639 Fst 0.97 0.814 Klk1c2 2.43 0.135 G6pd 1.11 0.599 Klk4 0.95 0.994 Gper1 G 1.06 0.731 Krt8 1.06 0.931 Igf1 0.86 0.503 Lama1 0.90 0.545 Igfbp5 0.98 0.867 Lifg1 1.00 0.975 Irs1 1.12 0.629 Liff	Fos	0.80	0.252	Ebag3 Efna5	1 27	0.200	
Fub 0.193 0.193 1.293 0.303 0.001 Gucy1a3 1.13 0.222 Erbb2 1.19 0.521 Herc3 1.01 0.974 Erbb3 1.52 0.992 Hpgd 1.17 0.482 Esr1 1.21 0.515 Igf1r 1.10 0.564 Esr2 0.86 0.606 Igfbp5 0.96 0.753 Fos 0.79 0.351 Irs2 1.08 0.733 Foxa1 0.32 0.0147* Jun 1.08 0.639 Fst 0.97 0.814 Klk1c2 2.43 0.135 G6pd1 1.11 0.599 Klk4 0.95 0.994 Gper1 G 1.06 0.731 Klk4 0.90 0.545 Igfbp4 1.51 0.164 Liff 0.90 0.545 Igfbp5 0.98 0.867 Lrig1 1.00 0.975 Irs1 1.12 0.629 Lrfip2	Fzd5	0.00	0.202	Enr2	0.89	0.207	
Cluby RD 1.13 0.222 Endb2 1.19 0.321 Herc3 1.01 0.974 Erbb3 1.52 0.992 Hpgd 1.17 0.482 Esr1 1.21 0.515 Igftr 1.10 0.564 Esr2 0.86 0.606 Igfbp5 0.96 0.753 Fos 0.79 0.351 Irs2 1.08 0.733 Foxa1 0.32 0.0147* Jun 1.08 0.639 Fst 0.97 0.814 Klk1c2 2.43 0.135 G6pd 1.11 0.599 Klk4 0.95 0.994 Gper1 G 1.06 0.731 Krt8 1.06 0.931 Igf1 0.86 0.503 Lama1 0.90 0.545 Igfbp5 0.98 0.867 Lift 0.90 0.545 Igfbp5 0.98 0.867 Lift 0.90 0.545 Igfbp5 0.98 0.867 Lift	Guov1o2	1 1 2	0.795	Erbh2	1 10	0.001	
Herds 1.01 0.974 Erobs 1.52 0.992 Hpgd 1.17 0.482 Esr1 1.21 0.515 Igftr 1.10 0.564 Esr2 0.86 0.606 Igftp5 0.96 0.753 Fos 0.79 0.351 Irs2 1.08 0.733 Foxa1 0.32 0.0147* Jun 1.08 0.639 Fst 0.97 0.814 Klk4 0.95 0.994 Gper1 G 1.06 0.791 Klk4 0.95 0.994 Gper1 G 1.06 0.731 Krt8 1.06 0.931 lgft 0.86 0.503 Lama1 0.90 0.545 lgfbp5 0.98 0.867 Liff 0.90 0.545 lgfbp5 0.98 0.867 Liff 0.90 0.545 lgfbp5 0.98 0.867 Liff 0.90 0.545 lgfbp4 1.51 0.164 Lifr 0.90 0.692 Klkb1 - - Map7d1 1.01<	Gucy 1a3	1.13	0.222	EIDD2 Erbb2	1.19	0.021	
Hygu 1.17 0.482 Esr1 1.21 0.515 Igf1r 1.10 0.564 Esr2 0.86 0.606 Igfbp5 0.96 0.753 Fos 0.79 0.351 Irs2 1.08 0.733 Foxa1 0.32 0.0147* Jun 1.08 0.639 Fst 0.97 0.814 Klk4 0.95 0.994 Gper1 G 1.06 0.791 Klk4 0.95 0.994 Gper1 G 1.06 0.731 Krt8 1.06 0.931 Igf1 0.86 0.503 Lama1 0.90 0.304 Igfbp5 0.98 0.867 Lrig1 1.00 0.975 Irs1 1.12 0.629 Lrig1 1.00 0.975 Irs1 1.12 0.629 Lrig1 1.00 0.975 Lgals1 1.01 0.883 Mt2A 1.15 0.563 Lpl 1.02 0.890 Myc <	Herco	1.01	0.974	EIDUS For1	1.02	0.992	
Ight 1.10 0.564 Esr2 0.86 0.506 Igfbp5 0.96 0.753 Fos 0.79 0.351 Irs2 1.08 0.733 Foxa1 0.32 0.0147* Jun 1.08 0.639 Fst 0.97 0.814 Klk1c2 2.43 0.135 G6pd 1.11 0.599 Klk4 0.95 0.994 Gper1 G 1.06 0.791 Klk4 0.95 0.994 Gper1 G 1.06 0.731 Krt8 1.06 0.931 Igftp5 0.98 0.867 Lirg1 0.90 0.545 Igfbp5 0.98 0.867 Lrig1 1.00 0.975 Irs1 1.12 0.629 Lrig1 1.00 0.975 Irs1 1.12 0.629 Lrig1 1.00 0.957 Lgals1 1.01 0.883 Mt2A 1.15 0.563 Lpl 1.02 0.890 Myc	npgu left#	1.17	0.462	ESI I	1.21	0.010	
Igilpp 0.90 0.733 Fos 0.73 0.32 0.0147* Jun 1.08 0.639 Fst 0.97 0.814 Klk1c2 2.43 0.135 G6pd 1.11 0.599 Klk4 0.95 0.994 Gper1 G 1.06 0.731 Klk4 0.95 0.994 Gper1 G 1.06 0.731 Klk4 0.90 0.304 Igfbp4 1.51 0.164 Lifr 0.90 0.545 Igfbp5 0.98 0.867 Lrig1 1.00 0.975 Irs1 1.12 0.629 Lrifp2 1.03 0.854 Junb 0.99 0.790 Maf 1.09 0.692 Klkb1 - - Map7d1 1.01 0.926 L1cam 1.02 0.890 Myc 1.23 0.286 Ltbp1 0.89 0.549 Ncapd3 1.00 0.990 Maff 0.71 0.473 <t< td=""><td>Igi II Iafha5</td><td>0.06</td><td>0.304</td><td>ESI2</td><td>0.80</td><td>0.000</td></t<>	Igi II Iafha5	0.06	0.304	ESI2	0.80	0.000	
Insz 1.08 0.733 Pokal 1 0.32 0.0147 Jun 1.08 0.639 Fst 0.97 0.814 Klk1c2 2.43 0.135 G6pd 1.11 0.599 Klk4 0.95 0.994 Gper1 G 1.06 0.731 Krt8 1.06 0.931 lgf1 0.86 0.503 Lama1 0.90 0.545 lgfbp5 0.98 0.867 Lrig1 1.00 0.975 Irs1 1.12 0.629 Lrfip2 1.03 0.854 Junb 0.99 0.790 Maf 1.09 0.692 Klkb1 - - Map7d1 1.01 0.926 L1cam 1.02 0.890 Mre 1.00 0.957 Lgals1 1.01 0.883 Mt2A 1.15 0.563 Lpl 1.02 0.890 Mrg1 1.27 0.127 Med1 1.07 0.844 Nfkb1 1.	Igippo	0.90	0.755	FUS	0.79	0.00147*	
Juli 1.08 0.039 Pst 0.97 0.814 Klk1c2 2.43 0.135 G6pd 1.11 0.599 Klk4 0.95 0.994 Gper1 G 1.06 0.791 Klk4 0.90 0.304 lgftp 0.86 0.503 Lama1 0.90 0.304 lgftp5 0.98 0.867 Lrig1 1.00 0.975 lrs1 1.12 0.629 Lrifp2 1.03 0.854 Junb 0.99 0.790 Maf 1.09 0.692 Klkb1 - - Map7d1 1.01 0.926 L1cam 1.02 0.890 Mre 1.00 0.957 Lgals1 1.01 0.883 Mt2A 1.15 0.563 Lpl 1.02 0.890 Mrg1 1.27 0.127 Med1 1.07 0.844 Nfkb1 1.05 0.866 Mmp9 2.81 0.357 Nfkb2 0.9	1152	1.00	0.733	Foxal	0.32	0.0147	
NKR22 2.43 0.133 Gdopd 1.11 0.399 Klk4 0.95 0.994 Gper1 G 1.06 0.791 Klkb1 - - Hsp90aa1 0.96 0.731 Krt8 1.06 0.931 lgf1 0.86 0.503 Lama1 0.90 0.304 lgfbp4 1.51 0.164 Lifr 0.90 0.545 lgfbp5 0.98 0.867 Lrig1 1.00 0.975 lrs1 1.12 0.629 Lrrfip2 1.03 0.854 Junb 0.99 0.790 Maf 1.09 0.692 Klkb1 - - Map7d1 1.01 0.926 L1cam 1.02 0.890 Mree 1.00 0.957 Lgals1 1.01 0.883 Mt2A 1.15 0.563 Lpl 1.02 0.890 Mrg1 1.27 0.127 Med1 1.07 0.844 Nfkb1 1.	Jun Kikto2	1.00	0.039	Cend	0.97	0.014	
NK4 0.95 0.994 Gper IG 1.06 0.731 Klkb1 - - Hsp90aa1 0.96 0.731 Krt8 1.06 0.931 lgf1 0.86 0.503 Lama1 0.90 0.304 lgfbp4 1.51 0.164 Lifr 0.90 0.545 lgfbp5 0.98 0.867 Lirg1 1.00 0.975 Irs1 1.12 0.629 Lrftp2 1.03 0.854 Junb 0.99 0.790 Maf 1.09 0.692 Klkb1 - - Map7d1 1.01 0.926 L1cam 1.02 0.890 Myc 1.23 0.286 Ltbp1 0.89 0.549 Ncapd3 1.00 0.990 Maff 0.71 0.473 Ndrg1 1.27 0.127 Med1 1.07 0.844 Nfkb1 1.05 0.866 Mmp9 2.81 0.357 Nfkb2 0.9	KIKTG2	2.43	0.135	Gopu Creart C	1.11	0.099	
Nk01 - - Hspstoaal 0.96 0.731 Krt8 1.06 0.931 lgf1 0.86 0.503 Lama1 0.90 0.304 lgfbp4 1.51 0.164 Lifr 0.90 0.545 lgfbp5 0.98 0.867 Lrig1 1.00 0.975 Irs1 1.12 0.629 Lrfip2 1.03 0.854 Junb 0.99 0.790 Maf 1.09 0.692 Klkb1 - - Map7d1 1.01 0.926 L1cam 1.02 0.994 Mme 1.00 0.957 Lgals1 1.01 0.883 Mt2A 1.15 0.563 Lpl 1.02 0.890 Myc 1.23 0.286 Ltbp1 0.89 0.549 Ncapd3 1.00 0.990 Maff 0.71 0.473 Nfkb1 1.05 0.866 Mmp9 2.81 0.357 Nfkb2 0.95 </td <td></td> <td>0.95</td> <td>0.994</td> <td>Gper I G</td> <td>1.06</td> <td>0.791</td>		0.95	0.994	Gper I G	1.06	0.791	
Nits 1.06 0.931 Ight 0.86 0.503 Lama1 0.90 0.304 Igfbp4 1.51 0.164 Lifr 0.90 0.545 Igfbp5 0.98 0.867 Lrig1 1.00 0.975 Irs1 1.12 0.629 Lrrfip2 1.03 0.854 Junb 0.99 0.790 Maf 1.09 0.692 Klkb1 - - Map7d1 1.01 0.926 L1cam 1.02 0.994 Mme 1.00 0.957 Lgals1 1.01 0.883 Mt2A 1.15 0.563 Lpl 1.02 0.890 Myc 1.23 0.286 Ltbp1 0.89 0.549 Ncapd3 1.00 0.990 Maff 0.71 0.473 Ndrg1 1.27 0.127 Med1 1.07 0.844 Nfkb1 1.05 0.866 Mmp9 2.81 0.357 Nfkb2 0.		1.06	-	HSp90aa i	0.96	0.731	
Lamai 0.90 0.304 Igrbp4 1.51 0.164 Lifr 0.90 0.545 Igfbp5 0.98 0.867 Lrig1 1.00 0.975 Irs1 1.12 0.629 Lrrfip2 1.03 0.854 Junb 0.99 0.790 Maf 1.09 0.692 Klkb1 - - Map7d1 1.01 0.926 L1cam 1.02 0.994 Mme 1.00 0.957 Lgals1 1.01 0.883 Mt2A 1.15 0.563 Lpl 1.02 0.890 Myc 1.23 0.286 Ltbp1 0.89 0.549 Ncapd3 1.00 0.990 Maff 0.71 0.473 Ndrg1 1.27 0.127 Med1 1.07 0.844 Nfkb1 1.05 0.866 Mmp9 2.81 0.357 Nfkb2 0.95 0.816 Mta1 1.06 0.678 Nfkb1 1	Krio	1.06	0.931	igi i	0.60	0.503	
Lirr 0.90 0.545 1grbps 0.98 0.567 Lrig1 1.00 0.975 1rs1 1.12 0.629 Lrrfip2 1.03 0.854 Junb 0.99 0.790 Maf 1.09 0.692 Klkb1 – – Map7d1 1.01 0.926 L1cam 1.02 0.994 Mme 1.00 0.957 Lgals1 1.01 0.883 Mt2A 1.15 0.563 Lpl 1.02 0.890 Myc 1.23 0.286 Ltbp1 0.89 0.549 Ncapd3 1.00 0.990 Maff 0.71 0.473 Ndrg1 1.27 0.127 Med1 1.07 0.844 Nfkb1 1.05 0.866 Mmp9 2.81 0.357 Nfkb2 0.95 0.816 Mta1 1.06 0.678 Nfkbia 1.01 0.773 Myc 1.33 0.348 Nkx3-1 0.64 0.102 Nab2 1.28 0.685 Orm1 – – Ncoa2 1.01 0.979 Pak1ip1 1.11 0.0870 Ncoa3 1.01 0.969 Pgc – – Ncor1 0.81 0.285 Pias1 1.10 0.789 Ncor2 0.98 0.888 Pik3r3 1.07 0.794 Nov 0.89 0.450 Pmepa1 1.14 0.489 Nr0b1 0.84 0.534 Ppap2a 0.92 0.341 Nr0b2 – – Rab4a 0.99 0.901 Nr2f6 1.07 0.787 Rel 0.80 0.131 Nr3c1 0.93 0.775	Lama	0.90	0.304	Igtop4	1.51	0.164	
Lrig1 1.00 0.975 Irs1 1.12 0.629 Lrrfip2 1.03 0.854 Junb 0.99 0.790 Maf 1.09 0.692 Klkb1 - - Map7d1 1.01 0.926 L1cam 1.02 0.994 Mme 1.00 0.957 Lgals1 1.01 0.883 Mt2A 1.15 0.563 Lpl 1.02 0.890 Myc 1.23 0.286 Ltbp1 0.89 0.549 Ncapd3 1.00 0.990 Maff 0.71 0.473 Ndrg1 1.27 0.127 Med1 1.07 0.844 Nfkb1 1.05 0.866 Mmp9 2.81 0.357 Nfkb2 0.95 0.816 Mta1 1.06 0.678 Nfkbia 1.01 0.773 Myc 1.33 0.348 Nkx3-1 0.64 0.102 Nab2 1.28 0.685 Orm1 - - Ncoa2 1.01 0.979 Pak1ip1 1.11	Littr	0.90	0.545	eqaigi	0.98	0.867	
Lmp2 1.03 0.854 Junb 0.99 0.790 Maf 1.09 0.692 Kikb1 - - Map7d1 1.01 0.926 L1cam 1.02 0.994 Mme 1.00 0.957 Lgals1 1.01 0.883 Mt2A 1.15 0.563 Lpl 1.02 0.890 Myc 1.23 0.286 Ltbp1 0.89 0.549 Ncapd3 1.00 0.990 Maff 0.71 0.473 Ndrg1 1.27 0.127 Med1 1.07 0.844 Nfkb1 1.05 0.866 Mmp9 2.81 0.357 Nfkb2 0.95 0.816 Mta1 1.06 0.678 Nfkbia 1.01 0.773 Myc 1.33 0.348 Nkx3-1 0.64 0.102 Nab2 1.28 0.685 Orm1 - - Ncoa2 1.01 0.979 Paktip1 1.11	Lrig i	1.00	0.975	Irsi	1.12	0.629	
Mat 1.09 0.692 Kikb1 - - Map7d1 1.01 0.926 L1cam 1.02 0.994 Mme 1.00 0.957 Lgals1 1.01 0.883 Mt2A 1.15 0.563 Lpl 1.02 0.890 Myc 1.23 0.286 Ltbp1 0.89 0.549 Ncapd3 1.00 0.990 Maff 0.71 0.473 Ndrg1 1.27 0.127 Med1 1.07 0.844 Nfkb1 1.05 0.866 Mmp9 2.81 0.357 Nfkb2 0.95 0.816 Mta1 1.06 0.678 Nfkb1a 1.01 0.773 Myc 1.33 0.348 Nkx3-1 0.64 0.102 Nab2 1.28 0.685 Orm1 - - Ncor1 0.979 9 Pak1ip1 1.11 0.0870 Ncoa3 1.01 0.969 Pgc -	Lmip2	1.03	0.854	Junb	0.99	0.790	
Map/a1 1.01 0.926 L1carn 1.02 0.994 Mme 1.00 0.957 Lgals1 1.01 0.883 Mt2A 1.15 0.563 Lpl 1.02 0.890 Myc 1.23 0.286 Ltbp1 0.89 0.549 Ncapd3 1.00 0.990 Maff 0.71 0.473 Ndrg1 1.27 0.127 Med1 1.07 0.844 Nfkb1 1.05 0.866 Mmp9 2.81 0.357 Nfkb2 0.95 0.816 Mta1 1.06 0.678 Nfkbia 1.01 0.773 Myc 1.33 0.348 Nkx3-1 0.64 0.102 Nab2 1.28 0.685 Orm1 - - Ncoa2 1.01 0.979 Pak1ip1 1.11 0.0870 Ncoa3 1.01 0.969 Pgc - - Ncor1 0.81 0.285 Pias1 1.10 <td>Mar Zala</td> <td>1.09</td> <td>0.692</td> <td>KIKD1</td> <td>-</td> <td>-</td>	Mar Zala	1.09	0.692	KIKD1	-	-	
Mme 1.00 0.957 Lgais I 1.01 0.883 Mt2A 1.15 0.563 Lpl 1.02 0.890 Myc 1.23 0.286 Ltbp1 0.89 0.549 Ncapd3 1.00 0.990 Maff 0.71 0.473 Ndrg1 1.27 0.127 Med1 1.07 0.844 Nfkb1 1.05 0.866 Mmp9 2.81 0.357 Nfkb2 0.95 0.816 Mta1 1.06 0.678 Nfkb1 1.01 0.773 Myc 1.33 0.348 Nkx3-1 0.64 0.102 Nab2 1.28 0.685 Orm1 - - Ncoa2 1.01 0.979 Pak1ip1 1.11 0.0870 Ncoa3 1.01 0.969 Pgc - - Ncor1 0.81 0.285 Pias1 1.10 0.789 Ncor2 0.98 0.450 Pmepa1 1.44 <td>Nap701</td> <td>1.01</td> <td>0.926</td> <td>Licam</td> <td>1.02</td> <td>0.994</td>	Nap701	1.01	0.926	Licam	1.02	0.994	
MtZA 1.15 0.563 Lpt 1.02 0.890 Myc 1.23 0.286 Ltbp1 0.89 0.549 Ncapd3 1.00 0.990 Maff 0.71 0.473 Ndrg1 1.27 0.127 Med1 1.07 0.844 Nfkb1 1.05 0.866 Mmp9 2.81 0.357 Nfkb2 0.95 0.816 Mta1 1.06 0.678 Nfkbia 1.01 0.773 Myc 1.33 0.348 Nkx3-1 0.64 0.102 Nab2 1.28 0.685 Orm1 - - Ncoa2 1.01 0.979 Pak1ip1 1.11 0.0870 Ncoa3 1.01 0.969 Pgc - - Ncor1 0.81 0.285 Pias1 1.10 0.789 Ncor2 0.98 0.888 Pik3r3 1.07 0.794 Nov 0.89 0.450 Pmepa1 1.14 <td>Nime</td> <td>1.00</td> <td>0.957</td> <td>Lgais1</td> <td>1.01</td> <td>0.883</td>	Nime	1.00	0.957	Lgais1	1.01	0.883	
Myc 1.23 0.286 Ltop1 0.89 0.349 Ncapd3 1.00 0.990 Maff 0.71 0.473 Ndrg1 1.27 0.127 Med1 1.07 0.844 Nfkb1 1.05 0.866 Mmp9 2.81 0.357 Nfkb2 0.95 0.816 Mta1 1.06 0.678 Nfkbia 1.01 0.773 Myc 1.33 0.348 Nkx3-1 0.64 0.102 Nab2 1.28 0.685 Orm1 - - Ncoa2 1.01 0.979 Pak1ip1 1.11 0.0870 Ncoa3 1.01 0.969 Pgc - - Ncor1 0.81 0.285 Pias1 1.10 0.789 Ncor2 0.98 0.888 Pik3r3 1.07 0.794 Nov 0.89 0.450 Pmepa1 1.14 0.489 Nr0b1 0.84 0.534 Ppap2a 0.92	Mt2A	1.15	0.563	Lpi	1.02	0.890	
Ncapos 1.00 0.990 Matr 0.71 0.473 Ndrg1 1.27 0.127 Med1 1.07 0.844 Nfkb1 1.05 0.866 Mmp9 2.81 0.357 Nfkb2 0.95 0.816 Mta1 1.06 0.678 Nfkbia 1.01 0.773 Myc 1.33 0.348 Nkx3-1 0.64 0.102 Nab2 1.28 0.685 Orm1 - - Ncora2 1.01 0.979 Pak1ip1 1.11 0.0870 Ncoa2 1.01 0.979 Pak1ip1 1.11 0.0870 Ncora3 1.01 0.969 Pgc - - Ncor1 0.81 0.285 Pias1 1.10 0.789 Ncor2 0.98 0.888 Pik3r3 1.07 0.794 Nov 0.89 0.450 Pmepa1 1.14 0.489 Nr0b1 0.84 0.534 Ppap2a <	IVIYC	1.23	0.286	Ltop I	0.89	0.549	
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Pmepa1 1.14 0.489 Nr0b1 0.84 0.534 Ppap2a 0.92 0.341 Nr0b2 - - Rab4a 0.99 0.901 Nr2f6 1.07 0.787 Rel 0.80 0.131 Nr3c1 0.93 0.775	Pik3r3	1.07	0.794	Nov	0.89	0.450	
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Rab4a0.990.901Nr2f61.070.787Rel0.800.131Nr3c10.930.775	Ppap2a	0.92	0.341	Nr0b2	-	-	
Rel 0.80 0.131 Nr3c1 0.93 0.775	Rab4a	0.99	0.901	Nr2f6	1.07	0.787	
	Kel	0.80	0.131	Nr3c1	0.93	0.775	

(Continued)

TABLE 1 | Continued

AR signaling target genes			ER signaling target genes		
Gene symbol	Fold changes	P value	Gene symbol	Fold changes	P value
Rela	0.97	0.788	Nr5a2	1.33	0.703
Ripk4	1.01	0.888	Nrip1	1.21	0.467
Sgk1	1.18	0.369	Nrp1	1.39	0.285
Slc26a2	1.04	0.801	Pdzk1	1.02	0.987
Slc45a3	0.75	0.509	Pelp1	0.95	0.650
Sms	0.98	0.685	Pgr	0.71	0.397
Snai2	0.79	0.328	Phb2	1.16	0.178
Sord	0.97	0.896	Ptch1	1.14	0.725
Sp1	1.06	0.692	Ptgs2	1.43	0.504
Spdef	1.15	0.475	Rala	1.06	0.587
Srf	1.12	0.564	Rara	1.27	0.548
Steap4	0.61	0.302	S100a6	0.87	0.539
Stk39	0.93	0.560	Safb	1.05	0.831
Tbc1d8	1.01	0.876	Snai1	1.15	0.725
Tiparp	1.56	0.988	Socs3	1.04	0.885
Tmprss2	2.17	0.0449*	Spp1	0.74	0.650
Tpd52	1.03	0.862	Tff1	-	-
Trib1	1.09	0.715	Tgfa	0.99	0.993
Tsc22d1	1.03	0.370	Tgfb3	1.16	0.466
Tsc22d3	1.12	0.587	Thbs1	1.05	0.861
Vapa	0.99	0.942	Vdr	1.15	0.686
Vipr1	0.96	0.821	Vegfa	1.14	0.719
Wipi1	1.08	0.655	Wisp2	1.01	0.974
Zbtb10	1.10	0.305	Wnt4	0.99	0.997
Zbtb16	0.99	0.889	Wnt5a	1.04	0.804
Zfp189	1.02	0.925	Xbp1	1.07	0.808

Fold changes: ratio of the average of BPA group (n = 3, two siblings in one sample) to control group (n = 3, two siblings in one sample).

–, under detectable.

*P < 0.05 by Student's t-test.

phosphoribosyltransferase 1 (Hprt1, NM_012583), lactate dehydrogenase A (Ldha, NM_017025), and ribosomal protein large P1 (Rplp1, NM_001007604) using the $\Delta\Delta$ Ct method. Expression levels of ER signaling target genes were normalized by the mean expression levels of Hprt1 and Rplp1, which were selected from the HKGs by the software based on their consistent expression levels within all samples. Ct cut-off was set to 35 in both assays and treated as undetectable.

Real-Time PCR

Five P3 male brains and five P35 male brains were collected under deep anesthesia by Z/K/X at 13.5 mg/90 µl/kg and stored at -80°C. Amygdala, hypothalamus, and telencephalon excluding amygdala and hippocampus were collected in a cryostat at -20°C. Tissues were homogenized in TRIzolTM Reagent (ThermoFisher SCIENTIFIC) and total RNA was extracted by chloroform. Total RNA was reverse transcribed by using High-Capacity cDNA Reverse Transcription Kit (ThermoFisher SCIENTIFIC). Expression levels of *Rattus norvegicus* Tmprss2 mRNA (NM_130424.3) and Foxa1 mRNA (NM_012742.1) were measured using Rplp1 mRNA (NM_001007604.2) as a reference HKG. Real-time PCR was performed using SensiFAST SYBR Master Mix (BioLine Reagent, London, United Kingdom) and primers (Tmprss2 forward: 5'-CACCTGCCATCCACATACAG-3', reverse: 5'-CCAGAACTTCCAAAGCAAGC-3'; Foxa1 forward:



5'-GGAGGCCTACTCCTCTGTCC-3', reverse: 5'-TTGGCGTA GGACATGTTGAA-3'; Rplp1 forward: 5'-GACGGTCACGGA GGATAAGA-3', reverse: 5'-GCAGATGAGGCTTCCAATGT-3'). Real-time PCR was performed using a StepOnePlus[™] Real-Time PCR System (ThermoFisher SCIENTIFIC) with conditions of 95°C for 2 min, 40 cycles of 95°C for 5 s, and 60°C for 30 s, followed by a dissociation step. The levels of each mRNA were normalized to Rplp1 mRNA using the $\Delta\Delta$ Ct method.

Immunohistochemistry and Fluorescent Microscopy

Immunohistochemistry was performed to investigate the location of Tmprss2 and Foxa1 protein in P3 male rat brains using rabbit monoclonal anti-TMPRSS2 antibody (EPR3861; abcam, Cambridge, United Kingdom) and anti-FOXA1 antibody (EPR10881: abcam). Briefly, five P3 male rats' brains were collected in 4% paraformaldehyde (PFA) under deep anesthetized by Z/K/X at 13.5 mg/90 µl/kg. After 5 days in 4% PFA at 4°C, brains were soaked in 30% sucrose in 0.1 M phosphate buffer at 4°C until they sank. Brains were kept at -80° C until sectioning at 20 μ m thickness on a cryostat at -20°C. Sections were incubated in 0.3% H₂O₂ in 20% methanol in 0.01 M phosphate buffered saline (PBS; pH 7.0) for 20 min to suppress endogenous peroxidase activity. Sections were then washed three times in PBS and incubated overnight at 4°C in the primary antibody at concentrations of 1:300 for anti-TMPRSS2 antibody and 1:200 for anti-FOXA1 antibody in blocking solution (0.5% Triton X, 2% normal goat serum in PBS). The next day, three subsequent washes in PBS were followed by incubation in biotinylated goat anti-rabbit IgG at 1:500 in blocking solution for 40 min. After the sections were washed in PBS three times, they were then incubated for 45 min in avidin-biotin complex (Vectastain Elite Kit, Vector, Burlingame, CA, USA) in blocking solution. The resulting complex was visualized by 3,3-diaminobenzidine after the sections were washed three times in PBS and rinsed in 0.05 M Tris-HCl buffer (pH 7.5). Immunohistochemistry without the primary antibodies served as control. The location of the immunoreactivities was identified by Nissl staining of the adjacent sections.

Number of Tmprss2-ir cells in the medial amygdala was counted in three 100 μ m grids and averaged for each P3 male rat that were subcutaneously injected with BPA at 50 μ g/kg/day (n = 5) or vehicle (n = 4) from P1 to P3. Number of Foxa1-ir cells in the hypothalamus was counted in three 100 μ m grids and averaged for each P3 male rat that were subcutaneously injected with BPA at 50 μ g/kg/day (n = 4) or vehicle (n = 4) from P1 to P3.

Co-localization of Tmprss2 or Foxa1 immunoreactivity and GnIH-EGFP was studied using a fluorescent microscope with bright field function. The secondary antibody (biotinylated goat anti-rabbit IgG) for immunohistochemistry was used at 1:200 to investigate the co-localization of Tmprss2 or Foxa1 and GnIH-EGFP.

Statistics

Differential expression of AR and ER signaling target genes in the medial amygdala, number of Tmprss2-ir cells in the medial amygdala, and number of Foxa1-ir cells in the hypothalamus between BPA administered male rats and control rats were analyzed by Student's *t*-test. Differential expression of Tmprss2 and Foxa1 in the amygdala, hypothalamus, and telencephalon excluding amygdala and hippocampus in P3 and P35 male rat brains was analyzed by one-way ANOVA followed by Fisher's protected least significant difference test.

RESULTS

Effect of BPA on AR and ER Signaling Target Gene Expression in the Medial Amygdala of Neonatal Male Rats

Differential expression of AR and ER signaling target genes in the medial amygdala by BPA administration was analyzed in neonatal male rats. Klkb1 (NM_012725), orosomucoid 1 (Orm1, NM_053288), and progastricsin (pepsinogen C) (Pgc, NM_133284) were undetectable within the 84 AR signaling target genes, and cytochrome P450 family 1 subfamily a polypeptide 1 (Cyp1a1, NM_012540), Klkb1 (NM_012725), nuclear receptor subfamily 0 group B member 2 (Nr0b2, NM_057133) were undetectable within the 84 ER signaling target genes (**Table 1**). Genes analyzed in both assays (Cited2, Igfbp5, Klkb1, Myc) produced similar results in terms of detectability, fold changes by BPA treatment, and *P* values by Student's *t*-test (**Table 1**).

An AR signaling target gene, Tmprss2 (NM_130424) was the only gene that was significantly increased in the medial

amygdala of neonatal male rats by BPA treatment (fold change 2.17, P = 0.0449; **Table 1**; **Figure 1A**). An ER signaling target gene, Foxa1 (NM_012742) was the only gene that decreased significantly in the medial amygdala of neonatal male rats by BPA treatment (fold change 0.32, P = 0.0147; **Table 1**; **Figure 1B**).



FIGURE 2 | Location of Tmprss2-immunoreactive cells in the neonatal male rat brain. (A) Tmprss2-immunoreactive (ir) cells in the olfactory bulb of Bisphenol A (BPA) treated P3 rat. AOL, anterior olfactory nucleus, lateral part. AOV, anterior olfactory nucleus, ventral part. (B) Tmprss2-ir cells in the secondary motor cortex (M2) of BPA treated P3 rat. (C) Tmprss2-ir cells in the hippocampus of BPA treated P3 rat. CA1, field CA1 of the hippocampus; CA2, field CA2 of the hippocampus; CA3, field CA3 of the hippocampus; MoDG, molecular layer of the dentate gyrus. (D) Tmprss2-ir cells in the hypothalamus of BPA treated P3 rat. DMH, dorsomedial hypothalamic area; VMH, ventromedial hypothalamic area; 3V, third ventricle. (E) Tmprss2-ir cells in the amygdala of vehicle treated P3 rat. MePD, medial amygdaloid nucleus, posteroventral part. (F) Tmprss2-ir cells in the amygdala of BPA treated P3 rat. (G) Higher magnification of MePV of BPA treated P3 rat (F). (H) Immunohistochemistry without the first antibody showed no immunoreactive cellular structure.









Distribution of Tmprss2 and Foxa1 Immunoreactive Cell Bodies in Neonatal Male Rat Brain

Location of Tmprss2 and Foxa1-ir cell bodies in P3 male rat brains was analyzed by immunohistochemistry. Abundant Tmprss2-ir cell bodies were found in the olfactory bulb (Figure 2A), cerebral cortex (Figure 2B), hippocampus (Figure 2C), hypothalamus (Figure 2D), and amygdala (Figures 2E–G). Spindle-like Tmprss2-ir cell bodies of about 10 µm in diameter (Figure 2G) were consistently found

in all brain areas. The weak blur staining of Tmprss2-ir cell bodies (Figure 2G) suggested that Tmprss2 exists in neuronal cellular membrane. Immunohistochemistry without the first antibody served as control (Figure 2H).

Foxa1-ir cell bodies were only found in the peduncular part of lateral hypothalamus (PLH) (**Figures 3A–C**). Three kinds of staining were observed. The first type of staining was even through the cytoplasm and the nucleus, and the cellular diameter was about $10 \,\mu m$ (**Figure 3C**, i). The second type of staining was a weak staining in the cytoplasm and darker staining in the nucleus of about 7 μm in



FIGURE 5 | Tmprss2-immunoreactive EGFP-GnIH neurons in the hypothalamus of neonatal male rat. (A,B) EGFP-GnIH neurons (A) and Tmprss2-immunoreactive (ir) cells (B) in the identical field and focus of hypothalamus. VMH, ventromedial hypothalamic area; Arc, arcuate hypothalamic nucleus; ME, median eminence. (C–E) Higher magnification of highlighted area in panel (A,B) with identical focus. Panel (D) is the merged image of (C,E). Arrows show Tmprss2-ir EGFP-GnIH neurons. [(C) i] An example of ventricle contacting rod-like EGFP-GnIH cell. [(C) ii] An example of EGFP-GnIH cell projecting its process to the ventricle. (F–H) Higher magnification of highlighted area in panel (A,B) with identical focus. Panel (G) is the merged image of (F,H). Arrows show Tmprss2-ir EGFP-GnIH neurons. (I–K) Higher magnification of highlighted area in panel (A,B) with identical focus. Panel (G) is the merged image of (F,H). Arrows show Tmprss2-ir EGFP-GnIH neurons. 3V, third ventricle. (L–N) Higher magnification of the central area in panel (F–H) with identical focus. Arrows show Tmprss2-ir EGFP-GnIH neurons.



FIGURE 6 | Developmental changes in Tmprss2 and Foxa1 gene expression in the amygdala, hypothalamus, and telencephalon excluding amygdala and hippocampus of male rat. (A) The ratio of Tmprss2 gene expression level relative to a housekeeping gene, Rplp1. Each column and the vertical line represent the mean \pm SEM (n = 5 samples, one sample from one rat). *******P* < 0.000001; ******P* < 0.00001, ****P* < 0.001 vs. amygdala of P3 male rat (A3) by one-way ANOVA followed by Fisher's protected least significant difference (PLSD) test. ⁺⁺⁺⁺P < 0.0001, hypothalamus of P3 male rat (H3) vs. hypothalamus of P35 male rat (H35) by Student's t-test. A35, amygdala of P35 male rat; T3, telencephalon excluding amygdala and hippocampus of P3 male rat; T35, telencephalon excluding amygdala and hippocampus of P35 male rat. (B) The ratio of Foxa1 gene expression level relative to Rplp1. Each column and the vertical line represent the mean \pm SEM (*n* = 5 samples, one sample from one rat). ****P* < 0.001; *P < 0.05 vs. A3 by one-way ANOVA followed by Fisher's PLSD test. [†]P < 0.05, H3 vs. H35 by Student's *t*-test.

diameter (**Figure 3C**, ii). The third type of staining was an intense staining only in the nucleus (**Figure 3C**, iii). Immunohistochemistry without the first antibody served as control (**Figure 3D**).

Effect of BPA Administration on the Number of Tmprss2-ir Cells in the Medial Amygdala and Foxa1-ir Cells in the Hypothalamus

To investigate the effect of BPA on Tmprss2 and Foxa1 protein expressions, number of Tmprss2-ir cells and Foxa1-ir cells were counted in the medial amygdala and hypothalamus, respectively, of P3 male rats that were daily administered with BPA or vehicle from P1 to P3. The number of Tmprss2-ir cells in the medial amygdala was significantly increased by BPA administration (P = 0.032; **Figures 2E,F** and **4A**). On the other hand, the number of Foxa1-ir cells in the hypothalamus was significantly decreased by BPA administration (P = 0.042; **Figures 3A,B** and **4B**).

Co-Localization of Tmprss2 and EGFP-GnIH Cells in the Hypothalamus

Because Tmprss2 was highly expressed in the hypothalamus of P3 male rat (**Figure 2D**), where GnIH neuronal cell bodies are located, we investigated if some Tmprss2-ir cells are also EGFP-GnIH positive (17). Abundant EGFP-GnIH cells were observed from the ventral region of ventromedial hypothalamic nucleus (VMH) to the dorsal region of the arcuate nucleus (Arc) as well as along the third ventricle in P3 male rat hypothalamus (**Figures 5A,C,F,I,L**). The diameter of EGFP-GnIH cell bodies in the VMH and Arc was 6–11 µm and had typical GnIH neuronal structure with dendrite or axon-like processes (**Figures 5A,F,L**). 26.5 \pm 6.8 (mean \pm SD from four different brains) percent of EGFP-GnIH cells in the VMH and Arc were immunoreactive to Tmprss2 antibody (**Figures 5A-N**).

EGFP-GnIH cells along the third ventricle were different between the dorsal population (Figure 5C) and the ventral population in the median eminence (ME) (Figure 5I). The dorsal population of EGFP-GnIH cells along the third ventricle had two cell types. The first type had a rod-like structure with a diameter of 4-7 µm and they were directly attached to the ventricle (Figure 5C,i). The second type was round shape with a diameter of 4–7 μ m situated in the sub-ventricular zone and had a protrusion to the ventricle (Figure 5C,ii). The ventral population of EGFP-GnIH cells in the ME had an oval shape with a minor axis of 4-7 µm directly contacted the ventricle and sent long processes to the external zone of the ME (Figure 5I). No clear cellular co-localization of Tmprss2 immunoreactivity was observed in EGFP-GnIH cells along the third ventricle either in the dorsal population (Figures 5A-E) or ventral population in the ME (Figures 5I–K).

Developmental Change in Tmprss2 and Foxa1 Gene Expression in the Amygdala, Hypothalamus, and Telencephalon in the Male Rat Brain

Tmprss2 was equally highly expressed in the amygdala and hypothalamus compared with the telencephalon excluding amygdala and hippocampus in P3 male rat brain (**Figure 6A**). However, Tmprss2 expression in the amygdala and hypothalamus decreased significantly in P35 male rat equivalent to Tmprss2 expression levels in the telencephalon (**Figure 6A**).

Foxal was expressed significantly higher in the hypothalamus compared with the amygdala of P3 male rats (**Figure 6B**). Although Foxal expression in the hypothalamus decreased significantly in P35, it was significantly higher than its expression level in the amygdala of P3 male rats (**Figure 6B**). Foxal expression in the telencephalon excluding amygdala and hippocampus was almost under detectable level (**Figure 6B**).

DISCUSSION

To understand how perinatal exposure of BPA can modify the neurodevelopmental process and behavior, we first searched for BPA responsive genes from AR and ER signaling target genes by PCR array in the neonatal male rat medial amygdala that were subcutaneously administered with a reference dose of BPA (50 μ g/kg BW/day) determined by United States EPA. We found upregulation of *Tmprss2*, an AR signaling target gene, and down-regulation of *Foxa1*, an ER signaling target gene. Number of Tmprss2-ir cells in the medial amygdala was also increased and Foxa1-ir cells in the hypothalamus was also decreased in the hypothalamus by BPA administration to neonatal male rats.

Tmprss2 encodes a multimeric protein that is translated into an N-terminal short cytoplasmic domain, a transmembrane domain, extracellular low-density lipoprotein receptor A domain, a scavenger receptor cysteine-rich domain, and a serine protease domain at the C-terminal (23). In human tissue, *TMPRSS2* is highly expressed in the small intestine, followed by heart, lung, liver, and small amounts in thymus and prostate. *TMPRSS2* is very highly expressed in the fetal brain but its expression level in the adult brain is minimum (23). *TMPRSS2* was found to be induced by androgen exposure to prostate cancer cells by microarray containing 1,500 cDNAs (24). It was shown that TMPRSS2 mediates androgen-induced prostate cancer cell invasion, tumor growth, and metastasis by stimulating a proteolytic cascade (25).

There is no study investigating the function of Tmprss2 in the brain besides only one study showing its possibility to mediate cancer pain by acting on trigeminal neurons (26). Significant decrease of *Tmprss2* expression in the amygdala and hypothalamus at 1 month of age suggests that Tmprss2 is related to neurogenesis or neuronal differentiation. In humans, TMPRSS2 is highly expressed in the fetus brain but its expression decreases to undetectable level in the adult brain (23). Tmprss2-ir cells were consistently distributed in olfactory bulb, cerebral cortex, hippocampus, amygdala, and hypothalamus in 3 days old male rats. We attempted Tmprss2 immunohistochemistry in 1-month-old and adult rat brains but we could not observe positive staining possibly due to its low expression level. The site of Tmprss2 expression corresponds well with that of AR expression in the rat brain (27, 28), suggesting that Tmprss2 expression is regulated by androgen.

In 3-day-old male rat hypothalamus, Tmprss2 was expressed in 26.5% of EGFP-GnIH neurons in the hypothalamus. EGFP-GnIH neurons were clustered from the ventral region of VMH to the dorsal region of Arc at this age, which was similar to fetal rat brain (29), but unlike GnIH neuronal population in the dorsomedial hypothalamic area in the adult (17, 30, 31). Abundant EGFP-GnIH cells were also found along the third ventricle, but clear expression of Tmprss2 was not observed in this region. EGFP-GnIH cells along the third ventricle can be determined as tanicytes from their morphology. Recent lineage-tracing experiments have shown that tanicytes along the third ventricle are hypothalamic stem cells (32) and postnatally differentiate into neurons and astrocytes in the hypothalamus (33). These results suggest that Tmprs2 is involved in the construction of neural and glial cellular architecture as a transmembrane protease in neonatal rat brain. Previous study has found that daily subcutaneous injection of a reference dose of BPA (50 µg/kg BW/day) to female Wister rats whose GnRH neurons express EGFP from P0 to P3 decreases numbers of GnIH-ir cell bodies, fiber density, and contacts on GnRH neurons and advances puberty at 1 month of age (34).

FOXA1 belongs to the FOXA family of transcriptional regulators (FOXA1, FOXA2, and FOXA3) that play pivotal roles in mammalian development (35). FOXA proteins serve as pioneering factors in the early sequence of the transcriptional regulatory program. FOXA induces nucleosomal rearrangement that facilitates binding of other transcriptional regulators such as sex steroid hormone nuclear receptors (36). FOXA1 serves as a pioneering factor that is required for ER α to bind estrogen-response-element after estrogen stimulation (37). FOXA1 also facilitates AR/chromatin interactions at the regulatory loci near androgen responsive genes (38). It was shown that BPA promotes epithelial mesenchymal transition of ER-negative breast cancer cells through downregulation of FOXA1 by PI3K/Akt activation (39). It is possible that BPA also downregulates Foxa1 by activating PI3K/Akt in the brain.

Foxa1 was highly expressed in the hypothalamus of P3 male rat but significantly decreased in the 1-month-old rat brain, suggesting that Foxa1 is also involved in neuronal differentiation. It was shown that Foxa1 is one of the regulators of neuronal differentiation of midbrain dopamine cells (40). A recent study showed that ablation of Foxa1 causes impaired development of subthalamic nucleus in mice (41), a brain region that is next to PLH where intense Foxa1-ir cells were found in P3 male rats in this study. Foxa1 immunoreactivity in the nucleus of the cells in PLH where AR and ER are also expressed (27) suggests that Foxa1 is involved in transcriptional activities of AR and ER in PLH.

Finally, it was shown that BPA upregulates Tmprss2 and downregulates Foxa1 in the neonatal male rat brain. Location of Tmprss2 and Foxa1-ir cells as well as significant decrease in Tmprss2 and Foxa1 expression in 1-month-old male rat brain suggest that BPA disturbs the neurodevelopmental process and behavior by modifying the expression of Tmprss2 and Foxa1 in the brain.

ETHICS STATEMENT

All procedures were approved by Monash University, Animal Ethics Committee (MARP/2016/037).

AUTHOR CONTRIBUTIONS

TU conceived the project. TU, SM, and TS performed the experiments. TU wrote and IP edited the paper.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at https://www.frontiersin.org/articles/10.3389/fendo.2018.00139/ full#supplementary-material.

REFERENCES

- Miyagawa S, Sato T, Iguchi T. Bisphenol A. In: Takei Y, Ando H, Tsutsui K, editors. *Handbook of Hormones*. Tokyo: Academic Press (2016). p. 577-8.
- Gore AC, Martien KM, Gagnidze K, Pfaff D. Implications of prenatal steroid perturbations for neurodevelopment, behavior, and autism. *Endocr Rev* (2014) 35:961–91. doi:10.1210/er.2013-1122
- Tan BL, Ali Mohd M. Analysis of selected pesticides and alkylphenols in human cord blood by gas chromatograph-mass spectrometer. *Talanta* (2003) 61:385–91. doi:10.1016/S0039-9140(03)00281-9
- Dodds EC, Lawson W. Synthetic oestrogenic agents without the phenanthrene nucleus. *Nature* (1936) 137:996. doi:10.1038/137996a0
- Milligan SR, Balasubramanian AV, Kalita JC. Relative potency of xenobiotic estrogens in an acute in vivo mammalian assay. *Environ Health Perspect* (1998) 106:23–6. doi:10.1289/ehp.9810623
- Sohoni P, Sumpter JP. Several environmental oestrogens are also anti-androgens. J Endocrinol (1998) 158:327–39. doi:10.1677/joe.0.1580327
- Dong S, Terasaka S, Kiyama R. Bisphenol A induces a rapid activation of Erk1/2 through GPR30 in human breast cancer cells. *Environ Pollut* (2011) 159:212–8. doi:10.1016/j.envpol.2010.09.004
- Wang C, Zhang J, Li Q, Zhang T, Deng Z, Lian J, et al. Low concentration of BPA induces mice spermatocytes apoptosis via GPR30. *Oncotarget* (2017) 8:49005–15. doi:10.18632/oncotarget.16923
- Xu LC, Sun H, Chen JF, Bian Q, Qian J, Song L, et al. Evaluation of androgen receptor transcriptional activities of bisphenol A, octylphenol and nonylphenol in vitro. *Toxicology* (2005) 216:197–203. doi:10.1016/j. tox.2005.08.006
- Bloom MS, Mok-Lin E, Fujimoto VY. Bisphenol A and ovarian steroidogenesis. Fertil Steril (2016) 106:857–63. doi:10.1016/j.fertnstert.2016.08.021
- Perera F, Vishnevetsky J, Herbstman JB, Calafat AM, Xiong W, Rauh V, et al. Prenatal bisphenol A exposure and child behavior in an inner-city cohort. *Environ Health Perspect* (2012) 120:1190–4. doi:10.1289/ehp.1104492
- Wolstenholme JT, Taylor JA, Shetty SR, Edwards M, Connelly JJ, Rissman EF. Gestational exposure to low dose bisphenol A alters social behavior in juvenile mice. *PLoS One* (2011) 6:e25448. doi:10.1371/journal.pone.0025448
- Farabollini F, Porrini S, Della Seta D, Bianchi F, Dessi-Fulgheri F. Effects of perinatal exposure to bisphenol A on sociosexual behavior of female and male rats. *Environ Health Perspect* (2002) 110(Suppl 3):409–14. doi:10.1289/ ehp.02110s3409
- Jones BA, Shimell JJ, Watson NV. Pre- and postnatal bisphenol A treatment results in persistent deficits in the sexual behavior of male rats, but not female rats, in adulthood. *Horm Behav* (2011) 59:246–51. doi:10.1016/j.yhbeh.2010.12.006
- Baron-Cohen S, Ring HA, Bullmore ET, Wheelwright S, Ashwin C, Williams SC. The amygdala theory of autism. *Neurosci Biobehav Rev* (2000) 24:355–64. doi:10.1016/S0149-7634(00)00011-7
- Nordahl CW, Scholz R, Yang X, Buonocore MH, Simon T, Rogers S, et al. Increased rate of amygdala growth in children aged 2 to 4 years with autism spectrum disorders: a longitudinal study. *Arch Gen Psychiatry* (2012) 69:53–61. doi:10.1001/archgenpsychiatry.2011.145
- Soga T, Kitahashi T, Clarke IJ, Parhar IS. Gonadotropin-inhibitory hormone promoter-driven enhanced green fluorescent protein expression decreases during aging in female rats. *Endocrinology* (2014) 155:1944–55. doi:10.1210/ en.2013-1786
- Tsutsui K, Ubuka T. Gonadotropin-inhibitory hormone. In: Kastin A, editor. Handbook of Biologically Active Peptides. London: Academic Press (2013). p. 802–11.
- Ubuka T, Son YL, Tsutsui K. Molecular, cellular, morphological, physiological and behavioral aspects of gonadotropin-inhibitory hormone. *Gen Comp Endocrinol* (2016) 227:27–50. doi:10.1016/j.ygcen.2015.09.009
- Ubuka T, Mukai M, Wolfe J, Beverly R, Clegg S, Wang A, et al. RNA interference of gonadotropin-inhibitory hormone gene induces arousal in songbirds. *PLoS One* (2012) 7:e30202. doi:10.1371/journal.pone.0030202
- Ubuka T, Haraguchi S, Tobari Y, Narihiro M, Ishikawa K, Hayashi T, et al. Hypothalamic inhibition of socio-sexual behaviour by increasing neuroestrogen synthesis. *Nat Commun* (2014) 5:3061. doi:10.1038/ncomms4061
- 22. Paxinos G, Watson C. *The Rat Brain in Stereotaxic Coordinates*. London: Academic Press (2007).
- Paoloni-Giacobino A, Chen H, Peitsch MC, Rossier C, Antonarakis SE. Cloning of the TMPRSS2 gene, which encodes a novel serine protease with transmembrane, LDLRA, and SRCR domains and maps to 21q22.3. *Genomics* (1997) 44:309–20. doi:10.1006/geno.1997.4845

- Lin B, Ferguson C, White JT, Wang S, Vessella R, True LD, et al. Prostatelocalized and androgen-regulated expression of the membrane-bound serine protease TMPRSS2. *Cancer Res* (1999) 59:4180–4.
- Ko CJ, Huang CC, Lin HY, Juan CP, Lan SW, Shyu HY, et al. Androgeninduced TMPRSS2 activates matriptase and promotes extracellular matrix degradation, prostate cancer cell invasion, tumor growth, and metastasis. *Cancer Res* (2015) 75:2949–60. doi:10.1158/0008-5472.CAN-14-3297
- Lam DK, Dang D, Flynn AN, Hardt M, Schmidt BL. TMPRSS2, a novel membrane-anchored mediator in cancer pain. *Pain* (2015) 156:923–30. doi:10.1097/j.pain.00000000000130
- Simerly RB, Chang C, Muramatsu M, Swanson LW. Distribution of androgen and estrogen receptor mRNA-containing cells in the rat brain: an in situ hybridization study. *J Comp Neurol* (1990) 294:76–95. doi:10.1002/cne. 902940107
- Xiao L, Jordan CL. Sex differences, laterality, and hormonal regulation of androgen receptor immunoreactivity in rat hippocampus. *Horm Behav* (2002) 42:327–36. doi:10.1006/hbeh.2002.1822
- Yano T, Iijima N, Hinuma S, Tanaka M, Ibata Y. Developmental expression of RFamide-related peptides in the rat central nervous system. *Brain Res Dev Brain Res* (2004) 152:109–20. doi:10.1016/j.devbrainres.2004.06.008
- Hinuma S, Shintani Y, Fukusumi S, Iijima N, Matsumoto Y, Hosoya M, et al. New neuropeptides containing carboxy-terminal RFamide and their receptor in mammals. *Nat Cell Biol* (2000) 2:703–8. doi:10.1038/35036326
- Ubuka T, Inoue K, Fukuda Y, Mizuno T, Ukena K, Kriegsfeld LJ, et al. Identification, expression, and physiological functions of Siberian hamster gonadotropin-inhibitory hormone. *Endocrinology* (2012) 153:373–85. doi:10.1210/en.2011-1110
- Rizzoti K, Lovell-Badge R. Pivotal role of median eminence tanycytes for hypothalamic function and neurogenesis. *Mol Cell Endocrinol* (2017) 445:7–13. doi:10.1016/j.mce.2016.08.020
- Haan N, Goodman T, Najdi-Samiei A, Stratford CM, Rice R, El Agha E, et al. Fgf10-expressing tanycytes add new neurons to the appetite/energy-balance regulating centers of the postnatal and adult hypothalamus. *J Neurosci* (2013) 33:6170–80. doi:10.1523/JNEUROSCI.2437-12.2013
- Losa-Ward SM, Todd KL, McCaffrey KA, Tsutsui K, Patisaul HB. Disrupted organization of RFamide pathways in the hypothalamus is associated with advanced puberty in female rats neonatally exposed to bisphenol A. *Biol Reprod* (2012) 87:28. doi:10.1095/biolreprod.112.100826
- Kaestner KH. The FoxA factors in organogenesis and differentiation. Curr Opin Genet Dev (2010) 20:527–32. doi:10.1016/j.gde.2010.06.005
- Augello MA, Hickey TE, Knudsen KE. FOXA1: master of steroid receptor function in cancer. *EMBO J* (2011) 30:3885–94. doi:10.1038/emboj.2011.340
- Carroll JS, Liu XS, Brodsky AS, Li W, Meyer CA, Szary AJ, et al. Chromosomewide mapping of estrogen receptor binding reveals long-range regulation requiring the forkhead protein FoxA1. *Cell* (2005) 122:33–43. doi:10.1016/j. cell.2005.05.008
- Wang Q, Li W, Zhang Y, Yuan X, Xu K, Yu J, et al. Androgen receptor regulates a distinct transcription program in androgen-independent prostate cancer. *Cell* (2009) 138:245–56. doi:10.1016/j.cell.2009.04.056
- Zhang XL, Wang HS, Liu N, Ge LC. Bisphenol A stimulates the epithelial mesenchymal transition of estrogen negative breast cancer cells via FOXA1 signals. Arch Biochem Biophys (2015) 585:10–6. doi:10.1016/j.abb.2015. 09.006
- 40. Metzakopian E, Bouhali K, Alvarez-Saavedra M, Whitsett JA, Picketts DJ, Ang SL. Genome-wide characterisation of Foxa1 binding sites reveals several mechanisms for regulating neuronal differentiation in midbrain dopamine cells. *Development* (2015) 142:1315–24. doi:10.1242/dev.115808
- Gasser E, Johannssen HC, Rülicke T, Zeilhofer HU, Stoffel M. Foxa1 is essential for development and functional integrity of the subthalamic nucleus. *Sci Rep* (2016) 6:38611. doi:10.1038/srep38611

Conflict of Interest Statement: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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