

Lefevre et al., Supplemental Table S1. Genes selected for PCR array analysis and specific to ovarian functions.

Gene Symbol	Gene Name	RefSeq NB	Tagman Assay
Ar	androgen receptor	NM_012502	Rn00560747_m1
Arh	aryl hydrocarbon receptor	NM_013149	Rn00565750_m1
Arnt	aryl hydrocarbon receptor nuclear translocator	NM_012780	Rn00562847_m1
Bax	Bcl2-associated X protein	NM_017059	Rn02532082_g1
Bcl2	B-cell CLL/lymphoma 2	NM_016993	Rn99999125_m1
Bmp4	bone morphogenetic protein 4	NM_012827	Rn00432087_m1
Bmp7	bone morphogenetic protein 7	NM_001191856	Rn01528889_m1
Casp2	caspase 2	NM_022522	Rn00574684_m1
Ccnd2	cyclin D2	NM_022267	Rn03020897_m1
Cmklr1	chemokine-like receptor 1	NM_022218	Rn00573616_s1
Cx43	gap junction protein, alpha 1	NM_012567	Rn01433957_m1
Cyp11a1	cytochrome P450, family 11, subfamily a, polypeptide 1	NM_017286	Rn00568733_m1
Cyp17a1	cytochrome P450, family 17, subfamily a, polypeptide 1	NM_012753	Rn00562601_m1
Cyp19	aromatase	NM_017085	Rn01422546_m1
Esr1	estrogen receptor 1 (ER alpha)	NM_012689	Rn01640372_m1
Esr2	estrogen receptor 2 (ER beta)	NM_012754	Rn00562610_m1
Foxo3a	forkhead box O3	NM_001106395	Rn01441087_m1
Fshr	follicle stimulating hormone receptor	NM_199237	Rn01648507_m1
Gdf9	growth differentiation factor 9	NM_021672	Rn00572328_m1
Gzma	granzyme A	NM_153468	Rn00596401_m1
Gzmb	granzyme B	NM_138517	Rn00821752_g1
Hsd17b1	hydroxysteroid (17-beta) dehydrogenase 1	NM_012851	Rn00563388_g1
Hsd3b1	hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 1	NM_001007719	Rn01774741_m1
Igf1	insulin-like growth factor	NM_178866	Rn00710306_m1
Insl3	insulin-like 3	NM_053680	Rn00586632_m1
Kitlg	KIT ligand	NM_021843	Rn01502851_m1
Lfn3	O-fucosylpeptide 3-beta-N-acetylglucosaminyltransferase	NM_133393	Rn00591325_m1
Lhcgr	luteinizing hormone/choriogonadotropin receptor	NM_012978	Rn00564309_m1
Mis	anti-Mullerian Hormone	NM_012902	Rn00563731_g1
Ngf	nerve growth factor (beta polypeptide)	XM_227525	Rn01533872_m1
Notch2	notch 2	NM_024358	Rn01534371_m1
Nr3c1	nuclear receptor subfamily 3, group C, member 1	NM_012576	Rn00561369_m1
Ppia	peptidylprolyl isomerase A (cyclophilin A)	NM_017101	Rn00690933_m1
Prlr	prolactin receptor	NM_012630	Rn01525459_m1
Pten	phosphatase and tensin homolog deleted on chr ten	NM_031606	Rn00477208_m1
Ptgs2	prostaglandin-endoperoxide synthase 2	NM_017232	Rn01483828_m1

Lefevre et al., Supplemental Table S1. *Continued*: Genes selected for PCR array analysis and specific to ovarian functions.

Gene Symbol	Gene Name	RefSeq NB	Taqman Assay
Rarres2	retinoic acid receptor responder 2	NM_001013427	Rn01451853_m1
Rxfp2	relaxin/insulin-like family peptide receptor 2	NM_001012475	Rn01412901_m1
Sohlhl1	spermatogenesis and oogenesis specific basic helix-loop-helix 1	NM_001191852.1	Rn04280363_g1
Spp1	secreted phosphoprotein 1	NM_012881	Rn01449972_m1
Srd5a1	steroid-5-alpha-reductase, alpha polypeptide 1	NM_017070	Rn00567064_m1
Srd5a2	steroid-5-alpha-reductase, alpha polypeptide 2	NM_022711	Rn00575595_m1
Star	steroidogenic acute regulatory protein	NM_031558	Rn00580695_m1
Tspo	translocator protein	NM_012515	Rn00560892_m1

Lefevre et al., Supplemental Table S2. Fold change in gene expression relative to *Ppia* expression and the control group.

Gene symbol	Gene name	Fold change relative to control				p value
		0	0.06	20	60	
Rxfp2	relaxin/insulin-like family peptide receptor 2	1.00	0.76	0.92	0.46	0.11
Spp1	secreted phosphoprotein 1	1.00	0.70	1.25	1.39	0.12
Ptgs2	prostaglandin-endoperoxide synthase 2	1.00	0.80	0.62	0.63	0.15
Rarres2	retinoic acid receptor responder 2	1.00	0.88	0.89	0.82	0.19
Lhcgr	lutinizing hormone/choriogonadotropin receptor	1.00	0.68	1.07	0.92	0.20
Tspo	translocator protein	1.00	1.22	0.96	1.04	0.27
Esr1	estrogen receptor 1 (ER alpha)	1.00	0.83	0.94	0.86	0.27
Cmklr1	chemokine-like receptor 1	1.00	1.30	0.99	1.05	0.28
Bmp7	bone morphogenetic protein 7	1.00	0.74	1.07	0.78	0.32
Cyp11a1	cytochrome P450, family 11, subfamily a, polypeptide 1	1.00	1.20	1.05	1.22	0.34
Cat	catalase	1.00	1.30	1.02	1.15	0.38
Gzmb	granzyme B	1.00	0.62	0.73	0.84	0.41
Hsd17β1	hydroxysteroid (17-beta) dehydrogenase 1	1.00	1.14	1.07	1.16	0.43
Gdf9	growth differentiation factor 9	1.00	0.62	0.93	0.96	0.44
Sod1	superoxide dismutase 1	1.00	1.07	1.11	1.12	0.45
Ngf	nerve growth factor (beta polypeptide)	1.00	0.91	0.87	0.80	0.47
Casp2	caspase 2	1.00	0.91	1.00	0.98	0.48
Arh	aryl hydrocarbon receptor	1.00	1.15	0.96	1.04	0.59
Bmp4	bone morphogenetic protein 4	1.00	1.08	0.88	0.82	0.60
Bax	bcl2-associated X protein	1.00	0.91	0.95	0.88	0.60
Srd5a1	steroid-5-alpha-reductase, alpha polypeptide 1	1.00	0.82	0.76	0.80	0.62
Cx43	gap junction protein, alpha 1	1.00	1.07	1.00	0.87	0.65
Cyp19	aromatase	1.00	1.19	1.12	1.25	0.68
Fshr	follicle stimulating hormone receptor	1.00	0.91	1.09	0.86	0.73
Lfng	O-fucosylpeptide 3-beta-N-acetylglucosaminyltransferase	1.00	0.90	0.88	0.89	0.74
Igf1	insulin-like growth factor	1.00	0.91	1.08	0.94	0.76
Mis	anti-Mullerian Hormone	1.00	0.75	0.86	0.93	0.76
Hsd3β1	hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 1	1.00	1.15	1.10	1.16	0.77
Foxo3a	forkhead box O3	1.00	0.87	0.96	0.96	0.78
Esr2	estrogen receptor 2 (ER beta)	1.00	0.90	1.01	0.91	0.78
Ar	androgen receptor	1.00	0.92	0.93	0.97	0.80

Lefevre et al., Supplemental Table S2. *Continued*: Fold change in gene expression relative to *Ppia* expression and the control group.

Gene symbol	Gene name	Fold change relative to control				p value
		0	0.06	20	60	
<i>Bcl2</i>	B-cell CLL/lymphoma 2	1.00	0.88	0.93	0.90	0.81
<i>Arnt</i>	aryl hydrocarbon receptor nuclear translocator	1.00	0.92	0.94	0.94	0.82
<i>Notch2</i>	notch 2	1.00	0.91	0.98	0.90	0.84
<i>Ccnd2</i>	cyclin D2	1.00	1.03	0.98	1.03	0.88
<i>Star</i>	steroidogenic acute regulatory protein	1.00	1.00	1.12	1.06	0.90
<i>Nr3c1</i>	nuclear receptor subfamily 3, group C, member 1	1.00	0.96	0.97	0.96	0.90
<i>Gzma</i>	granzyme A	1.00	1.00	0.84	0.93	0.93
<i>Kitlg</i>	KIT ligand	1.00	0.94	0.91	0.91	0.93
<i>Pten</i>	phosphatase and tensin homolog deleted on chr ten	1.00	1.03	1.00	0.99	1.00
<i>Prlr</i>	prolactin receptor	1.00	1.02	0.99	1.00	1.00
<i>Sohlh1</i>	spermatogenesis and oogenesis specific basic helix-loop-helix 1	1.00	1.00	1.00	1.01	1.00

Lefevre et al., Supplemental Table S3. Reproductive Parameters (adapted from [29]).

Parameter	Nominal BFR Mixture Dose (mg/kg/day)			
	Control (0)	0.06	20	60
Estrous cycle length - days	4.8 ± 0.4 (31)	4.6 ± 0.3 (29)	4.7 ± 0.2 (29)	5.6 ± 0.6 (29)
Mating Index ^a	70.4 (27)	85.2 (27)	75.0 (24)	70.8 (24)
Fecundity index ^b	79.0 (15)	91.3 (21)	94.4 (17)	100.0 (17)
Number of live fetuses per litter	17.0 ± 0.3 (15)	17.4 ± 0.4 (21)	16.8 ± 0.80 (17)	16.7 ± 0.9 (17)
Mean placental weights (mg)	477.0 ± 10.5	470.4 ± 9.6	464.6 ± 14.2	494.4 ± 21.7

Values are expressed as means ± SEM with the total number of observations in parentheses.

^a*Mating index = (number of sperm positive females / number of mated females) x 100.*

^b*Fecundity index = (number of pregnant females / number of sperm positive females) x 100.*