

Supplementary table 1. Sybr green primer pairs for RT-PCR

Gene	Forward/Reverse Primer	Primer
<i>Gapdh</i>	Forward Primer	AGGTCGGTGTGAACGGATTTG
	Reverse Primer	TGTAGACCATGTAGTTGAGGTCA
<i>Acta2</i>	Forward Primer	GTCCCAGACATCAGGGAGTAA
	Reverse Primer	TCGGATACTTCAGCGTCAGGA
<i>Cyp1b1</i>	Forward Primer	CACCAGCCTTAGTGCAGACAG
	Reverse Primer	GAGGACCACGGTTTCCGTTG
<i>Cldn1</i>	Forward Primer	GGGGACAACATCGTGACCG
	Reverse Primer	AGGAGTCGAAGACTTTGCACT
<i>P2rx2</i>	Forward Primer	GCGTTCCTGGGACTACGAGAC
	Reverse Primer	ACGTACCACACGAAGTAAAGC
<i>Fabp7</i>	Forward Primer	GGACACAATGCACATTCAGAAC
	Reverse Primer	CCGAACCACAGACTTACAGTTT
<i>Slitrk1</i>	Forward Primer	GAAGGGGACTTACACGTAGACT
	Reverse Primer	AGTGAGGGGAATTGCCATGCAG
<i>Ajap1</i>	Forward Primer	ACAGAGTTCAACGATTTGACTT
	Reverse Primer	GGGAAGGTGTTGGACTCCAG
<i>Itga9</i>	Forward Primer	AAGTGTCGTGTCCATACCAAC
	Reverse Primer	GGTCTGCTTCGTAGTAGATGTTT
<i>Hsd11b2</i>	Forward Primer	GGTTGTGACACTGGTTTTGGC
	Reverse Primer	AGAACACGGCTGATGTCCTCT
<i>Gfra2</i>	Forward Primer	CGCCTCTCGGACATCTTCAG
	Reverse Primer	TTGCAGTTGTCGTTCAAGTTG
<i>Syn3</i>	Forward Primer	CAATGGCTACATGCCGGAC
	Reverse Primer	CCTGGGTCTTTGAACCACGG
<i>Trpc6</i>	Forward Primer	AGCCAGGACTATTTGCTGATGG
	Reverse Primer	AACCTTCTTCCCTTCTCACGA
<i>Astn2</i>	Forward Primer	GGTGGACTCATTGCACTTCTC
	Reverse Primer	TTGCGCTCTTCTGTGGGATG
<i>Xkr4</i>	Forward Primer	ACTTCGCGGATGTGGGAAC
	Reverse Primer	CCAGCGGAAGCTGAACACTT
<i>Fgf5</i>	Forward Primer	TGTGTCTCAGGGGATTGTAGG
	Reverse Primer	AGCTGTTTTCTTGAATCTCTCC
<i>Tgfb1</i>	Forward Primer	CAGCACGGCCCCAATGTAT
	Reverse Primer	GGGACCTTTTCATATCCAGGACA
<i>Defa-rs4</i>	Forward Primer	AACGGCCTGCTTACCTTACT
	Reverse Primer	ACGTGTTCAAGTTCAACCCC
<i>Dpt</i>	Forward Primer	TGGATGGGTGAATCTTAACCGC
	Reverse Primer	TCAGAGCCTTCCTTCTTGCTA
<i>Sytl5</i>	Forward Primer	GGAAGACAAGAGGATAAGGAAGC

	<i>Reverse Primer</i>	TTGTGACAGTGAACGCAGACT
<i>Zcchc5</i>	<i>Forward Primer</i>	TGCCCCTAGAGTACCCTTTGG
	<i>Reverse Primer</i>	TCAGTGGGATACCCTCTGCTT
<i>Pdlim3</i>	<i>Forward Primer</i>	TGGGGGCATAGACTTCAATCA
	<i>Reverse Primer</i>	CTCCGTACCAAAGCCATCAATAG
<i>slc38a5</i>	<i>Forward Primer</i>	CTACAGGCAGGAACGCGAAG
	<i>Reverse Primer</i>	GGTTGAACACTGACATTCCGA
<i>Enpp2</i>	<i>Forward Primer</i>	TTTGC ACTATGCCAACAATCGG
	<i>Reverse Primer</i>	GGAGGCACTTTAGTCCTGTA CTT
<i>Dubr-202</i>	<i>Forward Primer</i>	ATAGCAACGGCCTGCTTACC
	<i>Reverse Primer</i>	GGTTCAACCCCTCGGTCTTT
<i>Ampd3</i>	<i>Forward Primer</i>	GTTGGCGGAGAAGGTGTTTG
	<i>Reverse Primer</i>	CTGCGACCGGATCATCTTGAA
<i>unc13c</i>	<i>Forward Primer</i>	AAAGCGAACTGCAAAGTGATGA
	<i>Reverse Primer</i>	TGCTCTCTGAAGAGTAGATCCAA
<i>Aspn</i>	<i>Forward Primer</i>	AAGGAGTATGTGATGCTACTGCT
	<i>Reverse Primer</i>	ACATTGGCACCCAAATGGACA
<i>Col6a6</i>	<i>Forward Primer</i>	GTGGACAGCTCCGATCACCTA
	<i>Reverse Primer</i>	GGGCGTTCCCTATCTTCAGG
<i>Col6a4</i>	<i>Forward Primer</i>	CTTCTGACAGCCGTA CTTGCC
	<i>Reverse Primer</i>	CGAGCACGAATAATGCGATGC
<i>Nrxn3</i>	<i>Forward Primer</i>	TGGGCTGCCTTAAAGAGGTTG
	<i>Reverse Primer</i>	GGGGTCTCAAAGTTGATGGGA
<i>Gpr50</i>	<i>Forward Primer</i>	AGAGCAACATGGGACCTACAA
	<i>Reverse Primer</i>	GCCAGAATTTCCGAGCTTCTTG
<i>Col14a1</i>	<i>Forward Primer</i>	TTTGGCGGCTGCTTGTTTC
	<i>Reverse Primer</i>	CGCTTTTGTTCAGTGTCTG
<i>Pgm5</i>	<i>Forward Primer</i>	TGCACCAGATGTTGTCTCAGA
	<i>Reverse Primer</i>	CTTCCCAGTCGAGATAGGTCA
<i>Nrk</i>	<i>Forward Primer</i>	GACCTGGGAGTTGGAGGGA
	<i>Reverse Primer</i>	CATAAGTACCAAGACCAATGGCT
<i>Ano1</i>	<i>Forward Primer</i>	CCCGTGCCAGTCACCTTTTT
	<i>Reverse Primer</i>	TCATCTGCTTCCGTTTCCAGT
<i>Col23a1</i>	<i>Forward Primer</i>	CCCCATCTGAGTGCATCTGTC
	<i>Reverse Primer</i>	CTTGCCGTCCAGACCTAGAG
<i>Mfap4</i>	<i>Forward Primer</i>	GCAACCCCTGGACTGTGATG
	<i>Reverse Primer</i>	TTGTCATGTCCGAGAAGACGG
<i>Enc1</i>	<i>Forward Primer</i>	CTGTTTCATAAGTCCTCCTACGC
	<i>Reverse Primer</i>	CACCACTGAACATGGCTTCG

Supplementary table 2. The top 20 downregulated genes in the absence of *Coup-TFII* in the mesonephros.

Gene Symbol	p-value (KO vs. WT)	Fold change (KO vs WT)
<i>Nr2f2</i>	1.32E-06	-2.51884
<i>Cyp1b1</i>	0.000187305	-2.24913
<i>Cldn1</i>	0.000964991	-1.87018
<i>Amhr2</i>	0.00027735	-1.84531
<i>P2rx2</i>	0.000170114	-1.76531
<i>Fabp7</i>	0.00126727	-1.76238
<i>Slitrk1</i>	0.000666716	-1.74521
<i>Ajap1</i>	3.20E-05	-1.69407
<i>Itga9</i>	0.00109656	-1.65097
<i>Zpld1</i>	0.00234644	-1.58798
<i>Tm4sf5</i>	0.0026883	-1.57505
<i>Hsd11b2</i>	0.00746169	-1.55323
<i>Gfra2</i>	0.00339666	-1.54512
<i>Syn3</i>	0.000105253	-1.53617
<i>Trpc6</i>	0.000126465	-1.51388
<i>Astn2</i>	0.000237335	-1.49525
<i>Xkr4</i>	0.00489398	-1.49496
<i>Fgf5</i>	0.00143297	-1.49425
<i>Tgfb1</i>	0.000120038	-1.49423
<i>Defa-rs4</i>	0.00750525	-1.48824

Supplementary table 3. The top 20 upregulated genes in the absence of *Coup-TFII* in the mesonephros.

Gene Symbol	p-value(KO vs. WT)	Fold change KO vs WT
<i>Dpt</i>	0.00279629	3.53294
<i>Sytl5</i>	0.00336484	2.20115
<i>Ret</i>	0.00086905	2.13712
<i>Zcchc5</i>	0.000359838	2.13014
<i>Pdlim3</i>	1.46E-05	2.10404
<i>Slc38a5</i>	1.89E-06	2.07221
<i>Enpp2</i>	0.00408385	2.0472
<i>Dubr</i>	0.0045779	2.04561
<i>Ampd3</i>	0.000446763	2.01873
<i>Unc13c</i>	7.46E-05	1.95979
<i>Aspn</i>	0.0473918	1.91231
<i>Col6a6</i>	0.00629587	1.87608
<i>Col6a4</i>	0.0305903	1.83411
<i>Ano4</i>	0.000156451	1.79544
<i>Acta2</i>	0.0013504	1.77138
<i>Nrxn3</i>	0.00786912	1.77108
<i>Gpr50</i>	0.0143848	1.76813
<i>Col14a1</i>	0.00350225	1.76605
<i>Pgm5</i>	0.000823237	1.76501
<i>Nrk</i>	0.00384636	1.73988

Supplementary Table 4: Target molecules of top upstream regulators in the list of differentially expressed genes.

Upstream regulators	Notes
TGFB1	ACTA2,ADAM12,ADM,ADORA2B,AGTR2,AIM2,ANGPT1,ANPEP,ANXA2,AQP1,ASPN,ATXN1,BMP2,BMP6,C1R,CADM1,CCL11,CD40,Cdkn1c,CELF2,CLDN1,Clec2d (includes others),COL6A3,CXCL13,CXCR6,DCN,DES,EGFR,ELN,ENPP1,ETS1,F3,FAS,FERMT1,FGF5,FGF7,FGFR2,FN1,FXD6,GDF6,GEM,GMPR,GNA14,GPR12,GRM3,GRM7,GUCY2F,HEXB,HOXA2,HTRA1,ID4,IGFBP5,IGFBP7,INHBB,ITGA11,KITLG,KLK3,MAPK13,MFAP4,MGP,MYH11,MYOCD,P2RY1,PLAGL1,PLCB1,PLCL1,PPFIBP2,PTGER4,RHOB,RUNX1,S1PR3,SCD,SERPINF1,SLC16A9,SLC7A5,SLIT3,SPP1,SVEP1,TAGLN,TBX2,TGFBI,TGM2,THBD,THBS1,THBS2,THBS4,THY1,TLL1,TNFAIP6,TRIM9,TRPC1,VA T1L,VCAM1,VDR,WNT11,WT1
progesterone	ACTA2,ADAMTS1,ADM,AEBP1,ALPL,AMPD3,AQP1,AR,BMP2,CA12,CADM1,CCNG1,CD40,Cdkn1c,CNR1,CYP1B1,EGFR,ENC1,ENPP1,ENPP2,F3,FGFR2,FN1,GREB1,HS D11B2,IGFBP5,IRS2,ITGA11,JAM2,KCNMA1,KLK3,LEPR,LGALS1,LGR5,LMO7,MME, NPY1R,PARM1,PLCL1,PRSS35,PTGER4,SCD,SPP1,ST3GAL1,TBX2,TGFBI,THBD,TN XB,TRPC1,VDR
beta-estradiol	ACTA2,ADAMTS1,ADAMTS18,ADAMTS5,ADM,AEBP1,AGTR2,ALX4,ANGPT1,ANPEP, ANXA1,AQP1,AR,BMP2,BMP5,CA12,CACNA1D,CCNG1,CD40,CELF2,CHRM3,CLDN1, CLDN11,Clec2d (includes others),CNR1,COL6A6,COLEC12,CXCL13,CYP1B1,DCLK1,DCN,DUSP10,DUSP6,EGF R,ENC1,ENPP2,ETS1,F3,FAS,FGF7,FGFR2,FHL2,FN1,FRK,GEM,GPR12,GREB1,HSD 11B2,HTRA1,HTRA3,ID4,IGFBP5,IGFBP7,IL6ST,INHBB,IRS2,ITGA11,JAM2,KCNN3,KI TLG,KLK3,LEPR,LGR5,LMO7,MACC1,MEDAG,MGP,MME,MPPED2,NPY1R,NRIP1,NR P2,NTN1,NTRK3,PLAC1,PRKCB,PRSS35,PTGER4,PTGFR,RET,RND3,RUNX1,RYR2, SFRP2,SLC7A5,SLC8A1,SLIT2,SMOC1,SNED1,SRD5A2,SSTR4,ST8SIA1,SYNE1,THB D,THBS1,THBS4,TNFAIP6,TP53INP1,TRIB2,TRPC1,VCAM1,VDR,WNT11
TNF	ABCD2,ACTA2,ADAMTS5,ADM,ADORA2B,AEBP1,AKR1B10,AMPD3,ANGPT1,ANPEP ,ANXA1,APCS,AQP1,AR,BMP2,BMP6,BMPER,CA12,CCL11,CD40,CDO1,CHSY3,CLD N11,CRISPLD2,CXCL13,CYP1B1,CYP27A1,DCN,DUSP10,DUSP6,EGFR,EMCN,ENPP 2,ENPP3,ETS1,EXOC3L4,F3,FAS,FERMT1,FGF5,FGFR2,FN1,FRZB,GABRA5,GEM,G FRA2,GRM3,GSDMD,HEXB,IGFBP5,IL1RL2,IRS2,KITLG,KLK3,LAMA4,LGR5,MFSD2A ,MGP,NRIP1,NTN1,OSMR,P2RX2,PDE7B,PDZD2,PTGFR,RGS5,RHOB,RND3,S1PR3, SCD,SCUBE1,SERPINB9,SERPINF1,SLC16A2,SLC8A1,SP7,SPP1,TAGLN,TGM2,THB D,THBS1,THBS2,THY1,TNFAIP6,TP53INP1,TRPC1,TRPC6,USP2,VCAM1,VDR
TWIST1	ACTA2,ADAMTS1,AR,CCDC3,CD40,DCN,EGFL6,ENPP3,ETS1,FAS,FAT4,FGFR2,FIBI N,FN1,FREM2,IGFBP5,MFGE8,MME,OLFML3,PDZD2,SPP1,ST8SIA2,TGFBI,THBS2

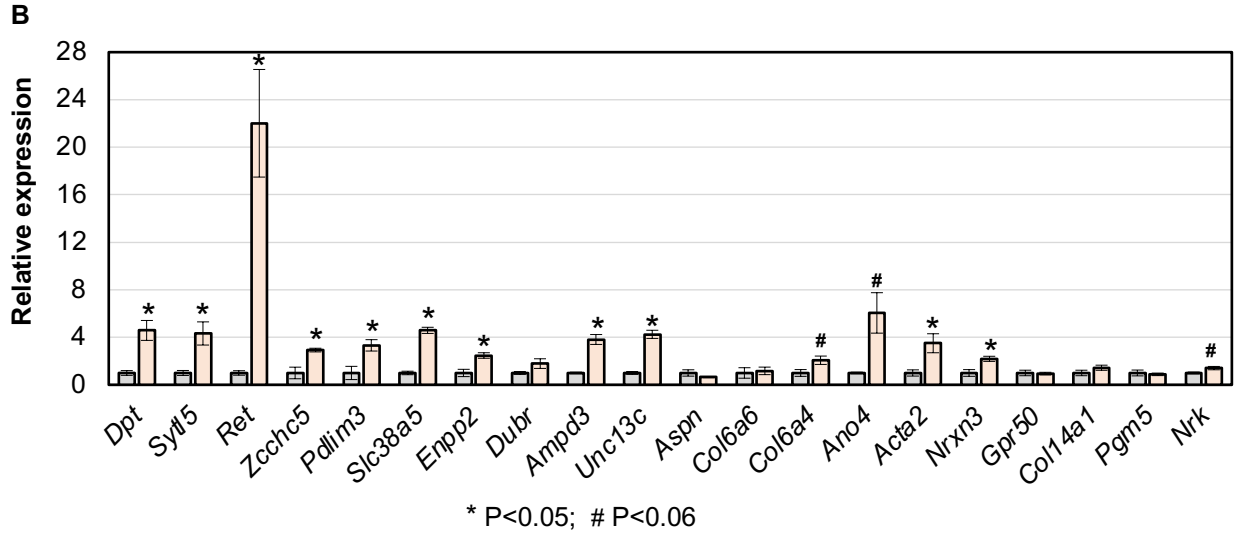
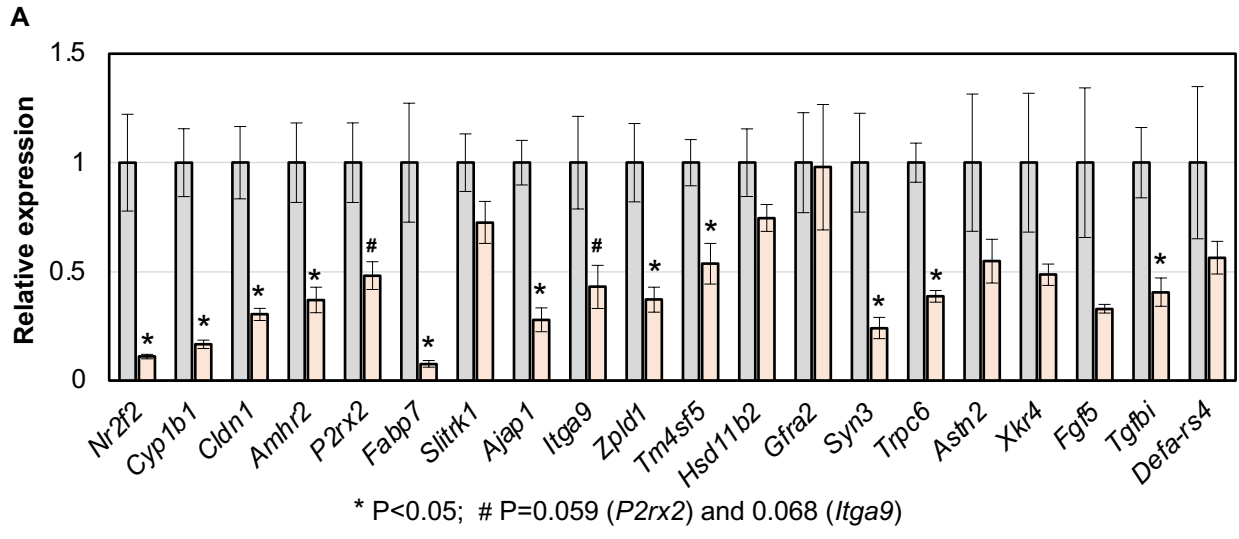
Supplementary Table 5: Mouses phenotype of putative COUP-TFII target genes with no expression information in the MGI database

Gene	Expression in fetal mesonephros	Mouse phenotype in reproduction	Notes
<i>Ampd3</i>	Unknown	No	
<i>Atrnl1</i>	Unknown	No	
<i>As3mt</i>	Unknown	No	
<i>Camk2d</i>	Unknown	No	
<i>Chsy3</i>	Unknown	Yes	Abnormal seminal vesicle; male and female infertility
<i>Epb41l2</i>	Unknown	Yes	Small testis and abnormal spermatogenesis
<i>Epha6</i>	Unknown	Yes	abnormal seminal vesicle and epididymis
<i>F3</i>	Unknown	No	
<i>Grk5</i>	Unknown	No	
<i>Ilf6st</i>	Unknown	Yes	Implantation failure
<i>Mme</i>	Unknown	No	
<i>Palld</i>	Unknown	Unknown	
<i>Rnu11</i>	Unknown	Unknown	
<i>Sned1</i>	Unknown	Unknown	
<i>Sulf1</i>	Unknown	No	
<i>Syn3</i>	Unknown	No	
<i>Tbc1d9</i>	Unknown	Unknown	
<i>Them7</i>	Unknown	Unknown	
<i>Trpc6</i>	Unknown	No	
<i>Usp2</i>	Unknown	Yes	Abnormal testis and reduced male fertility.
<i>Xkr4</i>	Unknown	Unknown	

Supplementary Table 6: Expression pattern and mouse phenotype of putative COUP-TFII target genes expressed in fetal mesonephros

Gene	Expression in fetal mesonephros	Mouse phenotype in reproduction	Notes (There will be no notes if expression localization in the mesonephros cannot be defined)
<i>Abcd2</i>	Yes	No	
<i>Ano1</i>	Yes	No	Wolffian duct mesenchyme and coelomic epithelium
<i>C1galt1</i>	Yes	No	
<i>Cacna1c</i>	Yes	Unknown	
<i>Cadm1</i>	Yes	No	Epithelium
<i>Cd24a</i>	Yes	Yes	
<i>Col23a1</i>	Yes	Unknown	Wolffian duct mesenchyme
<i>Cyp27a1</i>	Yes	No	
<i>Enc1</i>	Yes	No	Wolffian duct mesenchyme
<i>Fgfr2</i>	Yes	Yes	
<i>Gem</i>	Yes	No	
<i>Hmga2</i>	Yes	Yes	
<i>Kcnj8</i>	Yes	No	
<i>Lama1</i>	Yes	unknown	Expressed in Wolffian duct epithelium and mesenchyme
<i>Lama4</i>	Yes	No	
<i>Lsp1</i>	Yes	No	Wolffian duct mesenchyme?
<i>Man1a</i>	Yes	Unknown	Wolffian duct epithelium?
<i>Mfap4</i>	Yes	No	Wolffian duct mesenchyme
<i>Mybpc1</i>	Yes	unknown	Epithelium of Wolffian and Müllerian ducts
<i>Nr2f2</i>	Yes	Yes	Mesenchyme
<i>Slc7a5</i>	Yes	Unknown	Epithelium?
<i>Nrip1</i>	Yes	Yes	Epithelium
<i>Nrk</i>	Yes	Yes	

Supplementary figure 1



Supplementary figure 2

