

Table S1. Genes that are differentially expressed in *Nkx2.9*-deficient mouse embryos

Fold change	Regulation	Gene description	GenBank
10.5	Up	inactive X specific transcripts	BC051222
4.6	Down	eukaryotic translation initiation factor 2, subunit 3, structural gene Y-linked	BC043656
4.3	Down	jumonji, AT rich interactive domain 1D (Rbp2 like)	AF127244
3.7	Down	DEAD (Asp-Glu-Ala-Asp) box polypeptide 3, Y-linked	BC021453
3.6	Down	ubiquitously transcribed tetratricopeptide repeat gene, Y chromosome	AF057367
2.9	Down	homeo box C8	X07646
2.9	Down	RIKEN cDNA 4922502B01 gene	AK013705
2.4	Down	homeo box C6	X16511
2.4	Down	calbindin-28K	M21531
2.3	Down	homeo box D8	NM_008276.2
2.2	Down	neuromedin S	AB164466
2.1	Up	forkhead box D1	L38607
2.0	Down	otoraplin	AJ243939
2.0	Down	diacylglycerol kinase kappa	BC075627
2.0	Down	homeo box B9	BC100743
1.9	Down	carbonic anhydrase 10	AB080741
1.9	Down	V-set and transmembrane domain containing 2A	BC027127
1.9	Up	Tex17	AF285579
1.9	Down	Riken cDNA C130021I20 gene	AK147796
1.9	Down	homeo box A6	BC119105
1.9	Down	protein tyrosine phosphatase, non-receptor type 3	NM_011207.2
1.9	Down	MACRO domain containing 2	NC_000068.6
1.8	Up	SLIT and NTRK-like family, member 6	BC145927
1.8	Up	solute carrier family 38, member 4	AY027919
1.8	Down	dehydrogenase/reductase (SDR family) member 3	BC010972
1.8	Down	neurotrophin 3	BC065785
1.8	Down	latexin	D88769
1.8	Down	RIKEN cDNA 6430550H21 gene	BC062956
1.8	Up	decorin	BC132521
1.8	Down	LIM domain binding 2	U89489
1.8	Down	MAM domain containing glycosylphosphatidylinositol anchor 2	AY371925
1.8	Down	protein tyrosine phosphatase, non-receptor type 3	NM_011207.2
1.8	Down	cadherin 7, type 2	BC083189
1.8	Up	wingless-related MMTV integration site 5A	BC018425
1.7	Down	hemicentin 1	NM_001024720.3

1.7	Down	CUB and Sushi multiple domains 3	NM_001081391.2
1.7	Down	visinin-like 1	AY101375
1.7	Down	leucine rich repeat containing 4C	BC094588
1.7	Down	cadherin 9	NM_009869.1
1.7	Down	MACRO domain containing 2	NC_000068.6
1.7	Down	Adams3	NM_177872.2
1.7	Down	SLIT and NTRK-like family, member 2	BC111888
1.7	Down	protein phosphatase 1, regulatory (inhibitor) subunit 1C	BC120706
1.7	Down	metallophosphoesterase domain containing 2	BC020182
1.7	Down	aldehyde dehydrogenase family 1, subfamily A2	BC075704
1.7	Down	CUB and Sushi multiple domains 3	NM_001081391.2
1.7	Up	metallothionein 2	BC031758
1.7	Down	tachykinin 1	D17584
1.7	Down	roundabout homolog 2 (Drosophila)	DQ533876
1.7	Up	gene model 784, (NCBI)	DQ192038 BC076625
1.6	Down	KH domain containing, RNA binding, signal transduction associated 2	BC132117
1.6	Up	meiotic nuclear divisions 1 homolog (S. cerevisiae)	BC027741
1.6	Down	zinc finger protein 804A	BC035535
1.6	Up	RIKEN cDNA 2610301F02 gene	AK011952.1
1.6	Up	vitronectin	BC012690
1.6	Up	RIKEN cDNA 3100002J23 gene	BC117741 BC126964 BC094630
1.6	Down	SLIT and NTRK-like family, member 2	AK141924
1.6	Down	neuritin 1	BC035531
1.6	Down	PNMA-like 2	NM_001099636.2
1.6	Up	small nucleolar RNA, C/D box 16A	AF357363
1.6	Down	solute carrier family 17 (Na-dependent inorganic phosphate cotransporter), member 6	BC038375
1.6	Up	versican	D16263
1.6	Up	chemokine (C-X-C motif) ligand 12	D43804
1.6	Down	synaptic vesicle glycoprotein 2c	BC132111 U24700 AK173092
1.6	Down	glutamate receptor, ionotropic, AMPA1 (alpha 1)	BC056397
1.6	Up	sulfatase 1	AY101178
1.6	Up	myeloblastosis oncogene-like 1	L35261
1.6	Up	RIKEN cDNA 1810011O10 gene	BC016562
1.6	Down	potassium voltage-gated channel, Shal-related family, member 2	AB045326
1.6	Down	transmembrane protein 108	BC052085
1.6	Down	ubiquitin specific peptidase 29	AF229257
1.6	Up	dopa decarboxylase	AF071068

Genes exhibiting a fold change greater than 1.5 in *Nkx2.9* null mice as compared with WT littermates.