

Appendix 3. All probes that changed only at P13/N13 but remained stable at N9/N30.

Gene symbol	Gene name	P13/N13	N9/N30
<i>1700027L20Rik</i>	RIKEN cDNA 1700027L20 gene	2.9097	1.1889
<i>Acta2</i>	actin, alpha 2, smooth muscle, aorta	2.4665	0.8237
<i>Adm</i>	adrenomedullin	2.2952	1.2184
<i>Ak311</i>	adenylate kinase 3-like 1	1.7040	0.7079
<i>Ankrd11</i>	ankyrin repeat domain 11	0.6465	1.2766
<i>Ankrd37</i>	ankyrin repeat domain 37	1.5206	1.0405
<i>Anp32a</i>	acidic (leucine-rich) nuclear phosphoprotein 32 family, member A	1.6889	0.7095
<i>Anxa2</i>	annexin A2	2.1625	1.2005
<i>B2m</i>	beta-2 microglobulin	1.5357	0.9678
<i>Bnip3</i>	BCL2/adenovirus E1B interacting protein 3	1.7988	0.7879
<i>Ccl12</i>	chemokine (C-C motif) ligand 12	1.6336	1.0920
<i>Ccl4</i>	chemokine (C-C motif) ligand 4	1.6343	1.0484
<i>Chd4</i>	chromodomain helicase DNA binding protein 4	0.5983	1.4473
<i>Dars</i>	aspartyl-tRNA synthetase	1.5999	1.3845
<i>Ddx5*</i>	DEAD (Asp-Glu-Ala-Asp) box polypeptide 5	1.8907, 1.7185, 1.5464	1.1269, 1.0889, 0.8610
<i>Dnttip2</i>	deoxynucleotidyltransferase, interacting protein 2	terminal, 1.5878	1.0796
<i>Edn2</i>	endothelin 2	2.8626	0.8801
<i>Eef2*</i>	eukaryotic translation elongation factor 2	1.6314, 1.5485	0.9331, 0.9661
<i>Egln1*</i>	EGL nine homolog 1 ( <i>C. elegans</i> )	1.8848, 2.0157, 1.6188	0.9181, 0.9040, 0.9575
<i>Ero1l*</i>	ERO1-like ( <i>S. cerevisiae</i> )	1.8372,	0.8408,

		1.9127	0.7613
<i>Gpi1</i>	glucose phosphate isomerase 1	1.6661	0.8129
<i>Higd1a</i>	HIG1 domain family, member 1A	1.9103	0.7834
<i>Hist1h1c</i>	histone cluster 1, H1c	1.6016	0.6822
<i>Hmox1</i>	heme oxygenase (decycling) 1	1.8236	1.1132
<i>Hsd17b2</i>	hydroxysteroid (17-beta) dehydrogenase 2	2.3535	1.0329
<i>Lgals3</i>	lectin, galactose binding, soluble 3	1.7933	0.9829
<i>Loxl2</i>	lysyl oxidase-like 2	1.7291	1.0689
<i>MGI:107378</i>		1.7439	0.9516
<i>Muc2</i>	mucin 2	2.0306	0.9220
<i>Mxi1</i>	Max interacting protein 1	1.7752	1.0872
<i>P4ha2</i>	procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha II polypeptide	1.6242	0.7561
<i>Pdk1</i>	pyruvate dehydrogenase kinase, isoenzyme 1	1.8361	1.1600
<i>Pdlim1</i>	PDZ and LIM domain 1 (elfin)	1.7719	1.1408
<i>Pdlim3</i>	PDZ and LIM domain 3	1.6370	1.1068
<i>Pgam1</i>	phosphoglycerate mutase 1	2.0699	0.8084
<i>Pgk1</i>	phosphoglycerate kinase 1	1.5450	0.7590
<i>Ppm1b</i>	protein phosphatase 1B, magnesium dependent, beta isoform	1.6208	1.0097
<i>Rmcs2*</i>	response to metastatic cancers 2 (synonym: histocompatibility 2, class II antigen A, beta 1)	2.2525, 1.6821	1.0778, 1.0434
<i>Rras</i>	Harvey rat sarcoma oncogene, subgroup R	1.7135	0.7321
<i>Slc14a1</i>	solute carrier family 14 (urea transporter), member 1	1.6253	0.9476
<i>Syt5</i>	synaptotagmin V	2.7145	0.7828

<i>Tpi1</i>	triosephosphate isomerase 1	1.6187	0.6997
<i>Triobp</i>	TRIO and F-actin binding protein	1.7222	1.0455
<i>Tubb6</i>	tubulin, beta 6	1.7014	1.3052
<i>Vegfa</i>	vascular endothelial growth factor A	2.4823	1.2186
<i>Vim</i>	vimentin	2.3312	1.4946

Note \*: These genes have two or three probes in this array setting, thus have more than one ratio in each condition.