

<b>VEGF vs Untreated</b>						
<b>ensembl</b>	<b>symbol</b>	<b>base mean</b>	<b>fold change</b>	<b>wald stat</b>	<b>p-value</b>	<b>q-value</b>
ENSMUSG00000020256	Aldh1l2	2986.08	3.27	7.36416347	1.78261E-13	3.2497E-09
ENSMUSG00000022122	Ednrb	549.08	4.16	7.08582612	1.38217E-12	1.25985E-08
ENSMUSG00000047139	Cd24a	278.41	0.11	-6.5636546	5.25047E-11	3.19054E-07
ENSMUSG00000039683	Sdk1	222.62	8.83	6.30887463	2.81072E-10	1.28098E-06
ENSMUSG00000010797	Wnt2	83.80	0.00	-6.1885979	6.07017E-10	2.0497E-06
ENSMUSG00000021806	Nid2	2494.46	4.41	6.17193025	6.74613E-10	2.0497E-06
ENSMUSG00000039377	Hlx	407.17	2.55	5.71242868	1.11375E-08	2.82651E-05
ENSMUSG00000079242	C730034F03Rik	9.46	8.11E+05	5.69408148	1.24038E-08	2.82651E-05
ENSMUSG00000048126	Col6a3	35562.32	2.01	5.62055847	1.90341E-08	3.85547E-05
ENSMUSG00000035566	Pcdh17	142.31	11.04	5.59240415	2.23947E-08	4.08255E-05

<b>VEGF+BAPN vs Untreated</b>						
<b>ensembl</b>	<b>symbol</b>	<b>base mean</b>	<b>fold change</b>	<b>wald stat</b>	<b>p-value</b>	<b>q-value</b>
ENSMUSG00000020256	Aldh1l2	2986.07549	3.60	7.97	1.5439E-15	2.5371E-11
ENSMUSG00000022122	Ednrb	549.083878	4.67	7.67	1.69436E-14	1.39217E-10
ENSMUSG00000056174	Col8a2	616.07749	16.98	6.94	4.06035E-12	2.22413E-08
ENSMUSG00000027460	Angpt4	105.647578	17.07	6.86	6.87924E-12	2.82616E-08
ENSMUSG00000001918	Slc1a5	3999.48366	2.25	6.67	2.64179E-11	8.68251E-08
ENSMUSG00000027832	Ptx3	2086.82172	8.11	6.52	7.2632E-11	1.98927E-07
ENSMUSG00000016024	Lbp	683.597346	8.06	6.39	1.6284E-10	3.82278E-07
ENSMUSG00000017493	Igfbp4	11604.7487	4.06	6.31	2.73147E-10	5.40072E-07
ENSMUSG00000055653	Gpc3	1069.56238	22.11	6.30	2.95786E-10	5.40072E-07
ENSMUSG00000000031	H19	3798.69503	48.13	6.25	4.22439E-10	6.94195E-07

<b>VEGF+BAPN+TGFB vs Untreated</b>						
<b>ensembl</b>	<b>symbol</b>	<b>base mean</b>	<b>fold change</b>	<b>wald stat</b>	<b>p-value</b>	<b>q-value</b>
ENSMUSG00000020256	Aldh1l2	2986.07549	7.43	12.50	7.81525E-36	1.49467E-31
ENSMUSG00000074971	Fibin	1357.27816	6.41	12.36	4.2306E-35	4.04551E-31
ENSMUSG00000028195	Ccn1	7725.09486	3.49	12.20	3.21455E-34	2.04927E-30
ENSMUSG00000029752	Asns	2609.19795	4.72	11.75	7.30209E-32	3.49131E-28
ENSMUSG00000030268	Bcat1	2324.58976	5.70	11.43	3.11569E-30	1.19175E-26
ENSMUSG00000005124	Ccn4	4891.40456	7.25	11.24	2.50571E-29	7.98694E-26
ENSMUSG00000028179	Cth	307.840501	16.65	10.50	9.00062E-26	2.4591E-22
ENSMUSG00000027422	Rrbp1	15581.2419	1.73	10.39	2.72236E-25	6.50814E-22
ENSMUSG00000024529	Lox	30556.1013	7.09	10.30	7.23804E-25	1.53808E-21
ENSMUSG00000041168	Lonp1	3138.68264	2.27	10.18	2.33299E-24	4.46183E-21

<b>VEGF+BAPN vs VEGF</b>						
<b>ensembl</b>	<b>symbol</b>	<b>base mean</b>	<b>fold change</b>	<b>wald stat</b>	<b>p-value</b>	<b>q-value</b>
ENSMUSG00000028039	Efna3	27.77	87.74	4.66	3.19E-06	7.70E-02
ENSMUSG00000089857	Zfp882	47.28	17.27	4.30	1.70E-05	1.70E-01
ENSMUSG00000020224	Llph	247.68	2.05	4.21	2.52E-05	1.70E-01
ENSMUSG00000051998	Lax1	14.03	170.05	4.17	3.08E-05	1.70E-01
ENSMUSG00000087600	Pmepa1os	20.26	78.15	4.13	3.62E-05	1.70E-01
ENSMUSG00000097284	NA	13.31	95.28	4.10	4.22E-05	1.70E-01

ENSMUSG00000026928	Card9	21.21	61.18	3.99	6.50E-05	2.24E-01
ENSMUSG00000040329	Il7	16.33	98.28	3.94	8.21E-05	2.48E-01
ENSMUSG00000040345	Arhgap9	26.85	14.39	3.89	1.01E-04	2.72E-01
ENSMUSG00000050423	Ppp1r3g	17.85	49.16	3.77	1.62E-04	3.52E-01

**VEGF+BAPN+TGFB vs VEGF**

ensembl	symbol	base mean	fold change	wald stat	p-value	q-value
ENSMUSG00000038400	Pmepa1	9183.25	2.82	10.92	9.50902E-28	1.8186E-23
ENSMUSG00000002997	Prkar2b	813.73	0.23	-8.71	3.04011E-18	2.30662E-14
ENSMUSG000000005124	Ccn4	4891.40	4.62	8.69	3.61823E-18	2.30662E-14
ENSMUSG000000040152	Thbs1	41041.82	7.37	8.46	2.68227E-17	1.28246E-13
ENSMUSG000000037408	Cnm4	500.49	3.38	8.29	1.12563E-16	4.30553E-13
ENSMUSG000000028195	Ccn1	7725.09	2.31	8.17	3.02132E-16	9.63045E-13
ENSMUSG000000020325	Fstl3	544.67	3.49	7.89	3.07214E-15	7.66558E-12
ENSMUSG000000024529	Lox	30556.10	4.48	7.88	3.20652E-15	7.66558E-12
ENSMUSG000000027955	Gask1b	4702.94	2.02	7.74	9.72453E-15	2.06646E-11
ENSMUSG000000031253	Srpx2	1515.84	3.06	7.72	1.15034E-14	2.20003E-11

**VEGF+BAPN+TGFB vs VEGF+BAPN**

ensembl	symbol	base mean	fold change	wald stat	p-value	q-value
ENSMUSG000000027955	Gask1b	4702.94	2.65	10.76	5.38259E-27	1.0052E-22
ENSMUSG000000038400	Pmepa1	9183.25	2.64	10.25	1.24088E-24	1.15868E-20
ENSMUSG000000037408	Cnm4	500.49	3.34	8.27	1.34364E-16	8.36416E-13
ENSMUSG000000005124	Ccn4	4891.40	4.22	8.18	2.77159E-16	1.29399E-12
ENSMUSG000000015568	Lpl	1657.61	0.11	-7.85	4.03381E-15	1.50663E-11
ENSMUSG000000000693	Loxl3	5243.93	3.11	7.46	8.42671E-14	2.62281E-10
ENSMUSG000000054588	NA	64.73	0.03	-7.42	1.18908E-13	3.1723E-10
ENSMUSG000000027820	Mme	1122.64	0.06	-7.40	1.40151E-13	3.27164E-10
ENSMUSG000000016024	Lbp	683.60	0.09	-7.29	3.08321E-13	6.39765E-10
ENSMUSG000000054196	Cthrc1	1628.98	5.75	7.15	8.42189E-13	1.57279E-09

Supp. Table 2