

Supplementary data: Table I. SAGE analysis, LIF-specific genes.

SAGE Tag	Control	LIF	NRG-1	AngII	Aldo	LL ID	Unigene	Gene Name
GTATTGAAAA	1	20	9	5	5	76745	Mm.154109	RIKEN cDNA 1810009F10 gene
TTTGTAACATA	3	46	17	23	5	68029	Mm.32247	RIKEN cDNA 2010203O03 gene
GTTATGTTTG	0	13	1	3	0	27428	Mm.46014	shroom
ATTGTAGATG	4	45	21	10	4	14694	Mm.5305	guanine nucleotide binding protein, beta 2, related sequence 1
TTTATCCTTG	0	10	2	0	0	16412	Mm.4712	integrin beta 1 (fibronectin receptor beta)
TTCTATGCCT	1	10	2	3	0		Mm.45149	ESTs
GTTTCAAGGT	4	39	13	18	7	67115	Mm.43749	RIKEN cDNA 3100001N19 gene
CTAAAAAAA	4	28	14	4	15	68275	Mm.180734	RIKEN cDNA 5031405K23 gene
CTGCTTCAAT	3	17	7	2	3		Mm.100622	ESTs
CTGCTTCAAT	3	17	7	2	3	76356	Mm.18249	RIKEN cDNA 2610027H02 gene
CTGCTTCAAT	3	17	7	2	3		Mm.210150	ESTs
ACCAGATGCT	5	25	8	4	1	20753	Mm.625	small proline-rich protein 1A
AACCATGAAG	0	5	0	0	0	19646	Mm.12145	retinoblastoma binding protein 4
CTGCCACTGC	0	5	0	0	0	56361	Mm.23653	pseudouridine synthase 1
CACAATCCAG	0	5	0	0	0	67185	Mm.193688	RIKEN cDNA 2700059D21 gene
GTAGAACTTG	0	5	0	0	0	11469	Mm.116989	actin-like
GTAGAACTTG	0	5	0	0	0	11465	Mm.196173	actin, gamma, cytoplasmic
TTTTTTATAC	4	14	5	2	2	12567	Mm.6839	cyclin-dependent kinase 4
CAGGAGTTCA	74	23	66	55	65	67268	Mm.29057	RIKEN cDNA 2900073G15 gene
CAGGAGTTCA	74	23	66	55	65	67938	Mm.196536	RIKEN cDNA 1500001M02 gene
AGGAGCAGGC	35	10	20	25	29	69669	Mm.21864	RIKEN cDNA 2310075E07 gene
CAGGACCCAG	39	11	25	26	39	14314	Mm.22763	folliculin-like
CAGGACCCAG	39	11	25	26	39		Mm.173253	ESTs
TGGTGTCCCT	13	3	12	15	17	17096	Mm.1834	Yamaguchi sarcoma viral (v-yes-1) oncogene homolog
TGGTGAGAGG	9	1	8	11	7	76563	Mm.5904	RIKEN cDNA 2700038P16 gene
ACCAAACCT	35	2	16	26	55		Mm.214965	synaptosomal-associated protein, 25 kDa
GTCTAATCAC	32	1	12	19	10	18823	Mm.1268	proteolipid protein (myelin)
GTGCTTTCTG	16	0	5	9	27	233489	Mm.218602	MGC19382 Hypothetical protein
AGAGCTGGGA	12	0	5	12	9	65970	Mm.182607	epithelial protein lost in neoplasm
AGAGCTGGGA	12	0	5	12	9		Mm.87267	ESTs
AGGGCGACAC	16	0	8	13	19	11747	Mm.1620	annexin A5
CTGCAAGGCT	10	0	7	8	6	20597	Mm.4628	sphingomyelin phosphodiesterase 1, acid lysosomal
TCCTCTTTGG	7	0	11	9	6	17906	Mm.1529	myosin light chain, phosphorylatable, cardiac ventricles
AGTGTGAGTG	6	0	5	5	3	52693	Mm.196527	DNA segment, Chr 10, ERATO Doi 625, expressed

Note: Number of SAGE tags found in each of the 5 libraries is indicated.