

Supplemental table S10. List of the 63 common factors (listed as Gene names) between microarray and proteomic datasets with relative fold change of expression.

GeneName	Microarray	Proteome
Acaa2	-1.34	-1.16
Ace	1.14	1.69
Acot1	-1.51	-2.28
Acta1	2.75	3.70
Aldh1a1	1.90	1.74
Aldh5a1	-1.05	-2.94
Anxa2	1.16	1.19
Anxa5	1.12	1.79
Aspn	3.23	4.32
Bdh1	1.12	2.20
Carhsp1	1.35	1.42
Ccbl1	-1.06	-1.21
Ces3	-4.60	-4.61
Ckap4	1.17	2.58
Ckb	1.08	0.69
Cst3	1.24	2.12
Ctgf	3.30	4.70
Cygb	1.51	4.32
Dci	-1.11	-0.83
Decr1	-1.15	-0.78
Dpysl3	1.35	1.58
Dynll1	1.63	1.54
Efhd2	1.16	3.46
Endod1	1.40	5.39
Eno3	-1.00	-0.75
Ephx1	1.20	2.81
Fabp4	-1.37	-0.63
Fah	-1.32	-5.00
Fbn1	1.40	0.83
Fhl1	1.71	2.82
Gpt	-1.49	-5.64
Gstk1	-1.43	-2.26
Hspa1a	1.59	1.03
Igfbp7	1.56	2.50
Lgals1	1.17	1.38
Lmcd1	1.67	2.85
Lrp1	1.42	2.86
Lum	1.43	0.91
M6prbp1	-1.19	-0.84
Maob	-1.66	-2.00
Msn	1.03	1.35
Myh10	1.91	3.33
Myh7	3.80	4.82
Myom2	-1.13	-1.99
Naalad2	1.33	2.74

Nampt	-1.13	-1.34
Nppa	1.86	3.09
Nt5e	2.07	3.00
P4ha1	1.48	4.00
Picalm	1.20	2.22
Prelp	1.49	2.70
Psme1	-1.12	-1.84
Ptgfrn	1.29	3.70
Rcn3	1.71	4.17
Rnase4	1.29	4.00
Rtn4	1.97	3.95
S100a11	1.37	1.81
Sod2	-1.01	-0.91
Sorbs2	1.00	1.69
Thbs4	3.35	4.46
Tuba1a	1.02	0.71
Tubb2a	2.20	0.52
Vim	1.26	0.83