

Supplementary Table 7. Genes commonly de-regulated in the transcriptomic and proteomic analyses. Up-regulated genes are in red and down-regulated genes in blue. The list of genes is ordered according to log2foldchange in the proteomic study.

Gene	log2Foldchange RNA	log2Foldchange PROT
RARRES2	8,84	5,97
FABP3	3,92	4,50
LMOD1	4,65	4,12
COL1A2	5,61	3,70
CSPG4	6,94	3,25
CAV1	4,08	3,24
ALDH1L2	1,64	3,14
TAGLN	6,08	3,04
PTN	1,77	2,89
TNC	2,94	2,89
PODXL	4,38	2,74
FLNC	4,64	2,50
CYSTM1	1,54	2,50
COL1A1	2,20	2,49
NCALD	2,31	2,47
CAV2	2,45	2,44
TPST2	1,75	2,32
ALDH1B1	2,60	2,23
LCP1	4,61	2,13
ENAM	3,90	2,07
SORBS1	4,86	2,07
GAS6	1,50	2,07
HSPA2	2,20	2,01
JPH2	3,63	1,95
ALCAM	1,60	1,92
FAS	1,62	1,82
MYOF	2,09	1,81
JAG1	2,51	1,72
COL4A1	1,82	1,72
GPC4	2,56	1,64
GPC1	1,53	1,64
CPA4	3,23	1,57
ITGA5	1,70	1,55
MYL9	3,41	1,52
NT5E	2,45	1,51
PTGIS	2,69	1,51
ASMTL	-1,78	-1,55
CADM1	-4,03	-1,62
VAT1L	-2,44	-1,74
SLC14A1	-4,18	-1,80
CYP11A1	-5,82	-1,89
LPXN	-1,66	-2,13
CALB2	-5,96	-2,44
ALDH1A3	-2,80	-2,44