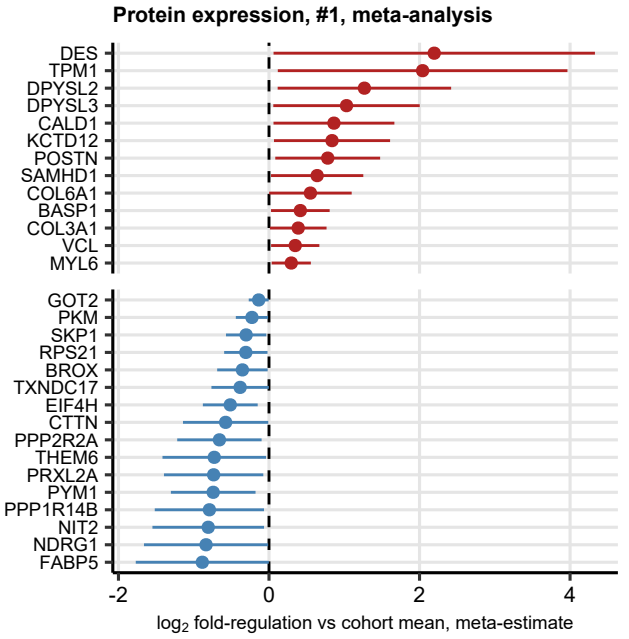
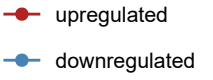
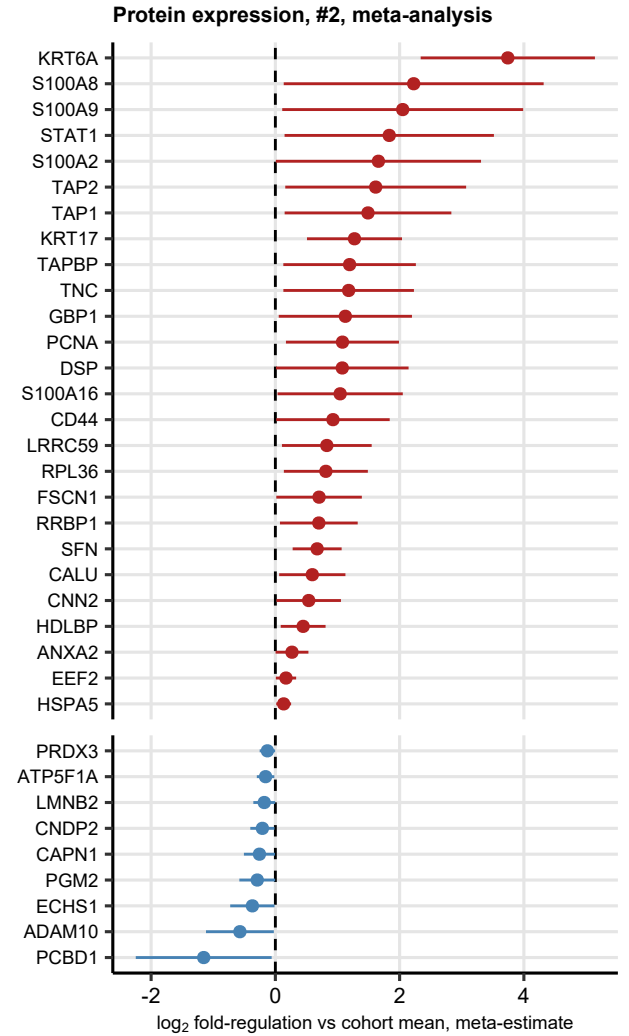
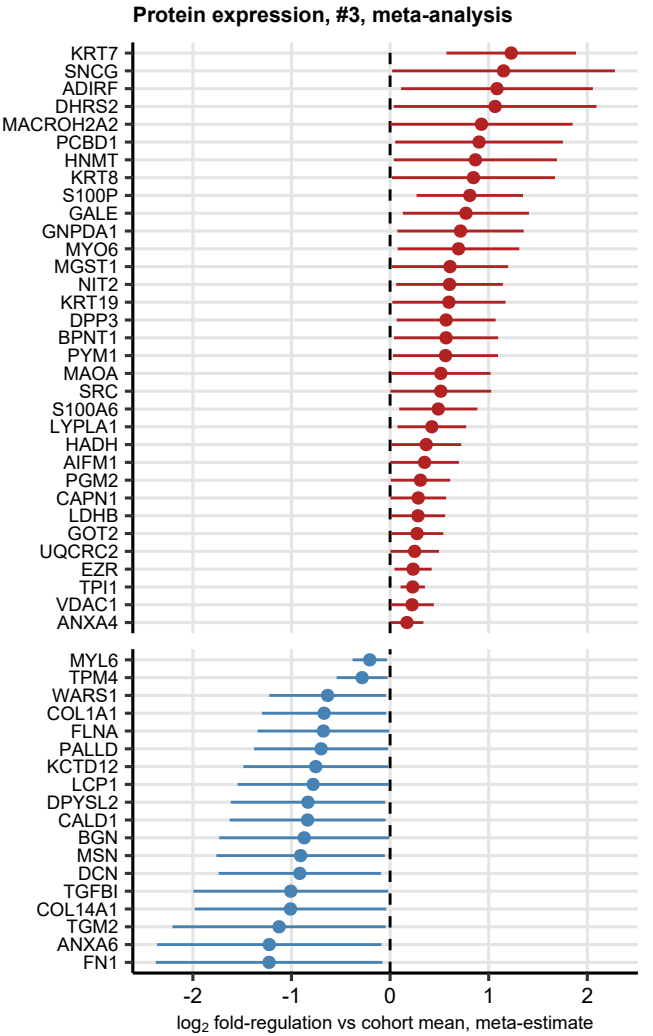

Protein expression, #1, meta-analysis
DES
TPM1
DPYSL2
DPYSL3
CALD1
KCTD12
POSTN
SAMHD1
COL6A1
BASP1
COL3A1
VCL
MYL6
GOT2
PKM
SKP1
RPS21
BROX
TXNDC17
EIF4H
CTTN
PPP2R2A
THEM6
PRXL2A
PYM1
PPP1R14B
NIT2
NDRG1
FABP5
-2
0
2
4
log2 fold-regulation vs cohort mean, meta-estimate


upregulated
downregulated


Protein expression, #2, meta-analysis
KRT6A
S100A8
S100A9
STAT1
S100A2
TAP2
TAP1
KRT17
TAPBP
TNC
GBP1
PCNA
DSP
S100A16
CD44
LRRC59
RPL36
FSCN1
RRBP1
SFN
CALU
CNN2
HDLBP
ANXA2
EEF2
HSPA5
PRDX3
ATP5F1A
LMNB2
CNDP2
CAPN1
PGM2
ECHS1
ADAM10
PCBD1
-2
0
2
4
log2 fold-regulation vs cohort mean, meta-estimate


Protein expression, #3, meta-analysis
KRT7
SNCG
ADIRF
DHRS2
MACROH2A2
PCBD1
HNMT
KRT8
S100P
GALE
GNPDA1
MYO6
MGST1
NIT2
KRT19
DPP3
BPNT1
PYM1
MAOA
SRC
S100A6
LYPLA1
HADH
AIFM1
PGM2
CAPN1
LDHB
GOT2
UQCRC2
EZR
TPI1
VDAC1
ANXA4
MYL6
TPM4
WARS1
COL1A1
FLNA
PALLD
KCTD12
LCP1
DPYSL2
CALD1
BGN
MSN
DCN
TGFBI
COL14A1
TGM2
ANXA6
FN1
-2
-1
0
1
2
log2 fold-regulation vs cohort mean, meta-estimate