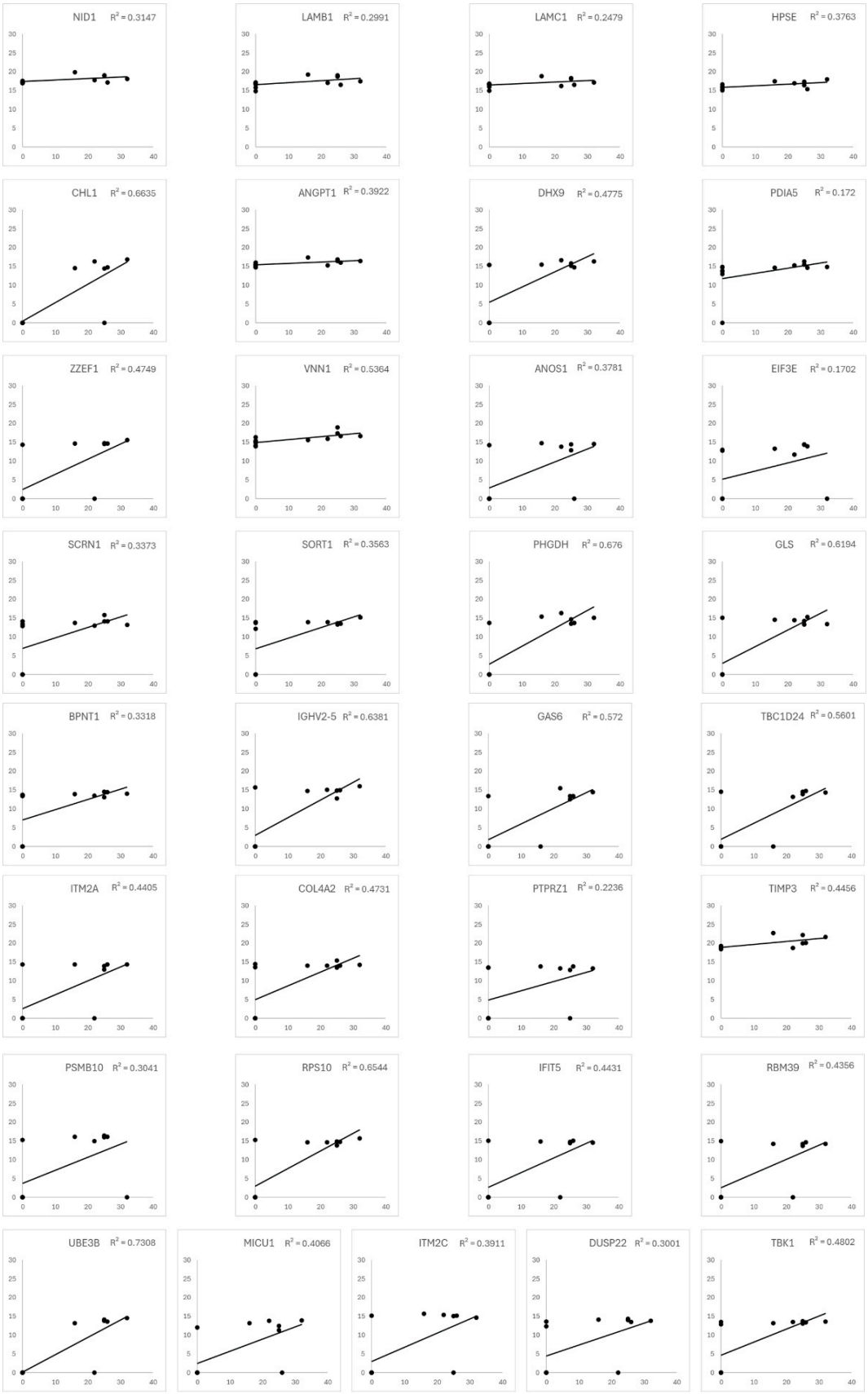
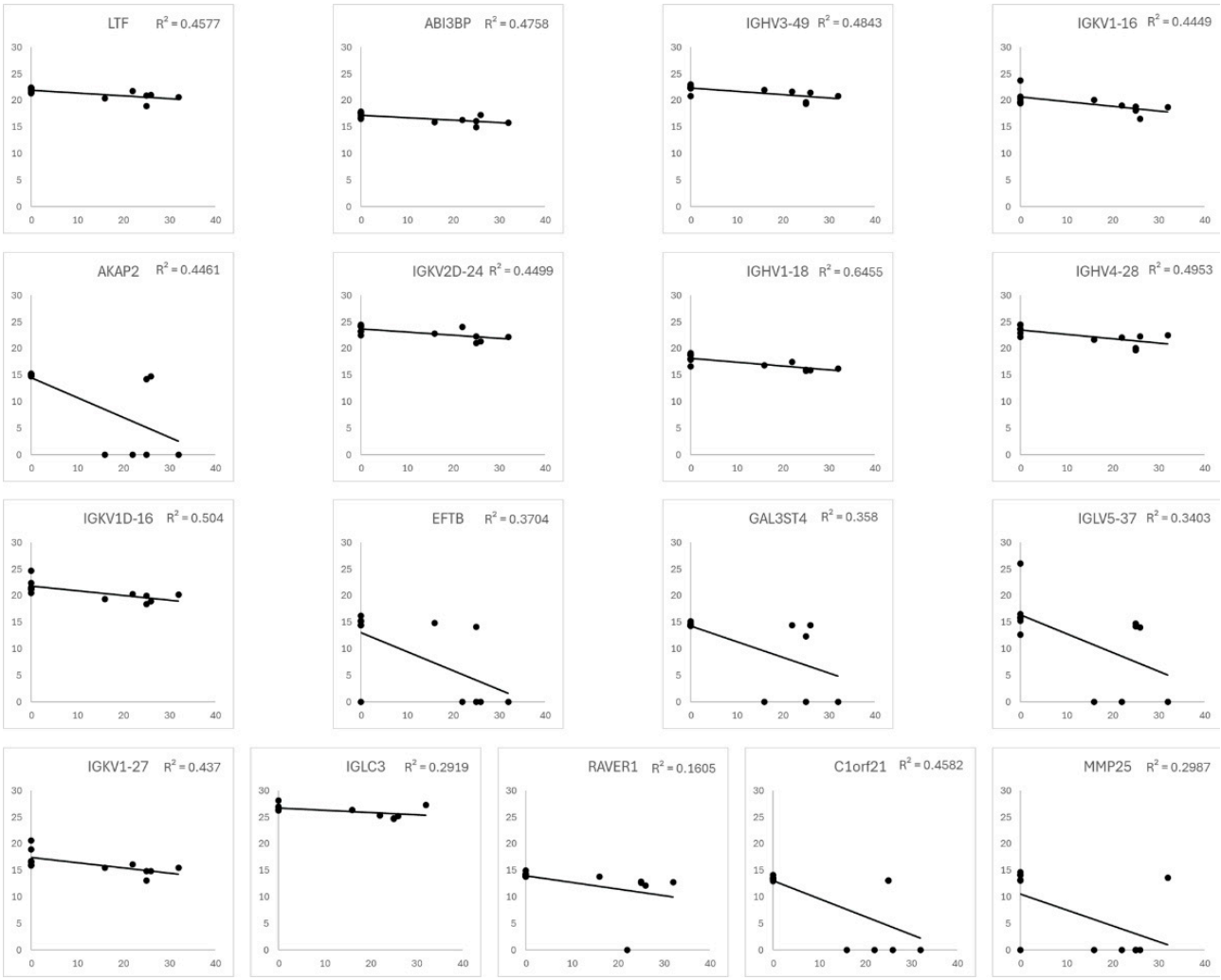

NID1 R² = 0.3147
LAMB1 R² = 0.2991
LAMC1 R² = 0.2479
HPSE R² = 0.3763
CHL1 R² = 0.6635
ANGPT1 R² = 0.3922
DHX9 R² = 0.4775
PDIA5 R² = 0.172
ZZEF1 R² = 0.4749
VNN1 R² = 0.5364
ANOS1 R² = 0.3781
EIF3E R² = 0.1702
SCRN1 R² = 0.3373
SORT1 R² = 0.3563
PHGDH R² = 0.676
GLS R² = 0.6194
BPNT1 R² = 0.3318
IGHV2-5 R² = 0.6381
GAS6 R² = 0.572
TBC1D24 R² = 0.5601
ITM2A R² = 0.4405
COL4A2 R² = 0.4731
PTPRZ1 R² = 0.2236
TIMP3 R² = 0.4456
PSMB10 R² = 0.3041
RPS10 R² = 0.6544
IFIT5 R² = 0.4431
RBM39 R² = 0.4356
UBE3B R² = 0.7308
MICU1 R² = 0.4066
ITM2C R² = 0.3911
DUSP22 R² = 0.3001
TBK1 R² = 0.4802

(a)



(b)

Figure S2. Correlation of up- and down-regulated proteins and CSS score

On the 33 up- (a) and 17 down-regulated (b) proteins, scatter plots of the log2-transformed protein masses versus the CSS scores in RTT group, with the regression line indicated. The vertical axis represents protein mass, and the horizontal axis represents the CSS score.