



Supplementary information, Fig. S2 Impact of CNAs on the transcriptome, proteome and phosphoproteome of PitNETs, related to Fig. 2.

a Boxplot showing sample-wise mRNA-protein correlations in all PitNETs.

b Median of sample-wise mRNA-protein correlations in each clinicopathological subtype.

c The panel shows the correlation of CNAs to phosphoprotein abundance, with significant positive correlations in red and negative correlations in green (Spearman's correlation, FDR < 0.05). Genes were sorted by chromosomal location on the x and y axes.

d Significant *cis* and *trans* effects of selected 15 CAGs on the expression of cancer associated mRNA and protein.

e Significant *cis* and *trans* effects of selected 6 CAGs on the expression of cancer associated mRNA and phosphoprotein.