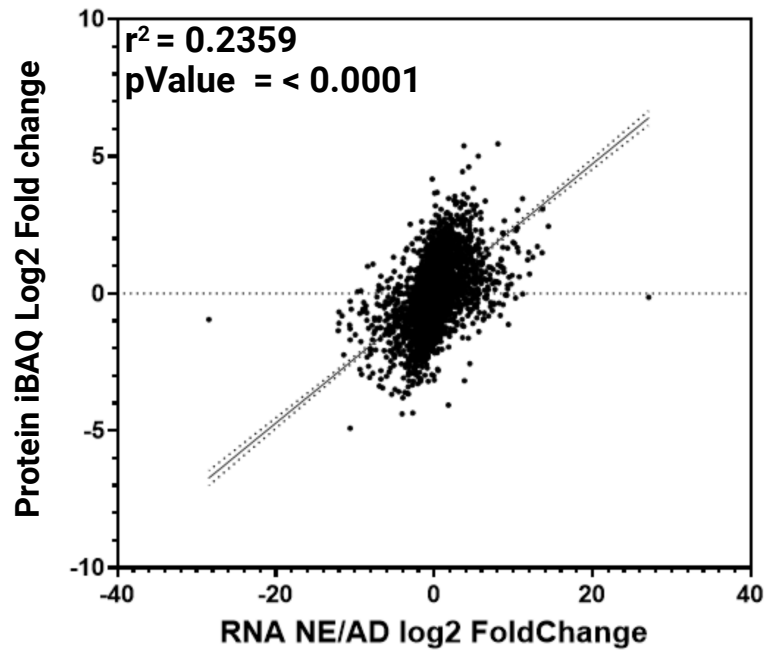**Supplemental Figure S3**

Overall linear regression plot between protein Log2 fold change relative intensity based average quantification (iBAQ) against the correspondent transcript FPKM Log2 fold change with a correlation of r^2=0.2359 and a p-value of < 0.0001