

Figure S1. Transcriptome analysis of somatotroph and lactotroph adenomas. A: Volcano map of differentially expressed proteins. B: The most enriched GO terms. CD: KEGG pathway analysis based on DEGs.

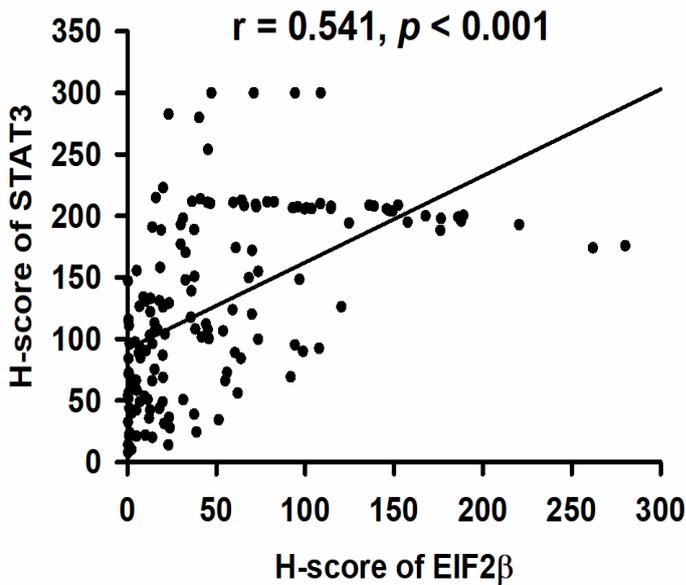


Figure S2. Correlation between EIF2 β H-score and STAT3 H-score in patients with pituitary adenomas.