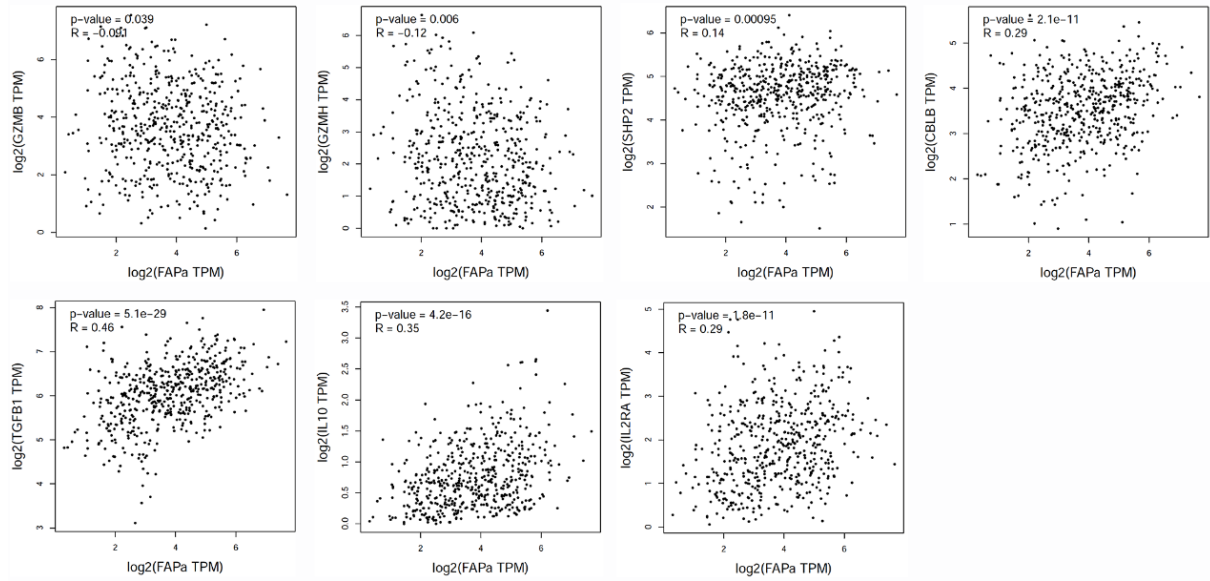


Supplementary Figure S8



Supplementary Figure S8. Correlation analysis between expression of FAP α and immune regulatory molecules. Expression data were extracted from the *TCGA head and neck squamous cell carcinoma RNA sequencing dataset*. Spearman's correlation test was used to calculate the correlation between mRNA expressions of two select genes.