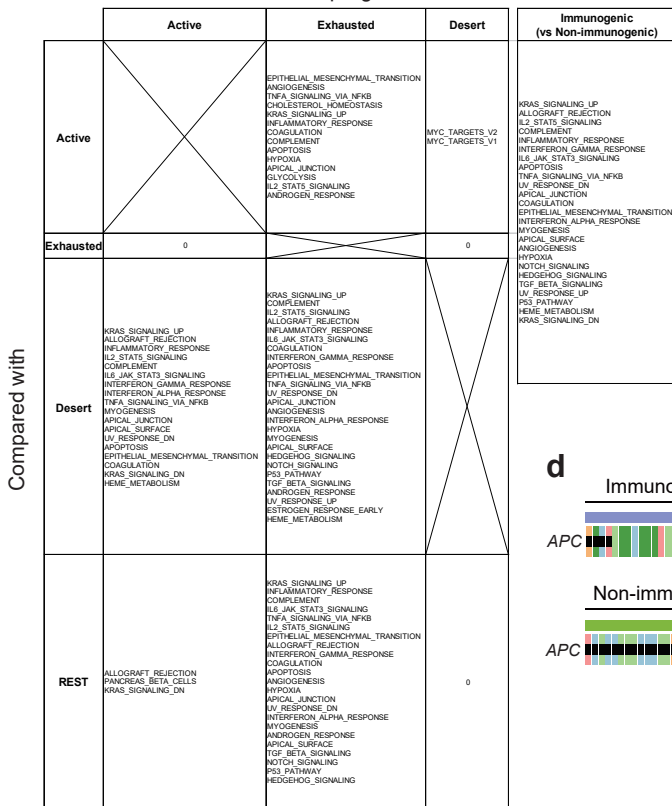


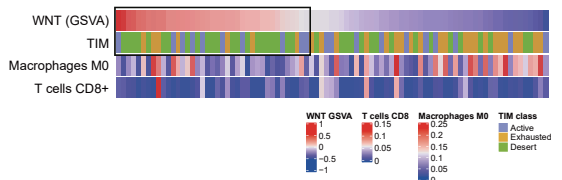
## a Pathway analysis using GSEA in primary tissues

Upregulated in

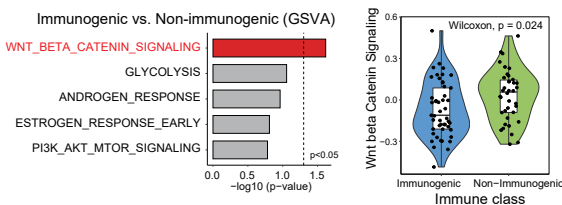


Compared with

b



c



d

Immunogenic (Active/Exhausted)

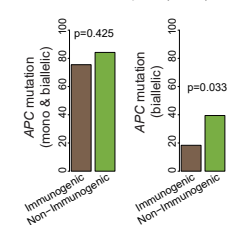


Non-immunogenic (Desert)



e

All samples (n=87)



f

