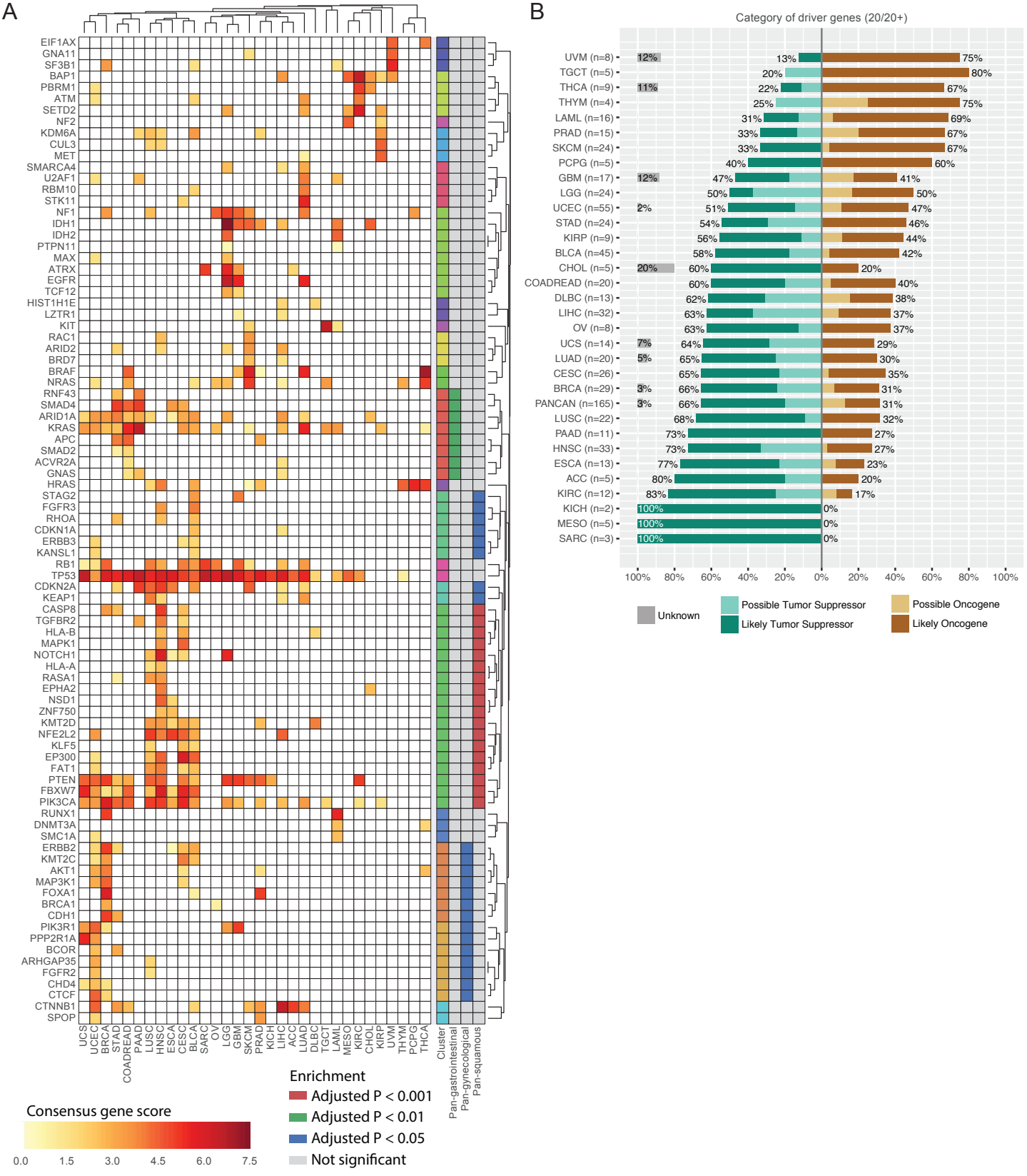
Figure S4


A
B
Category of driver genes (20/20+)
UVM (n=8)
TGCT (n=5)
THCA (n=9)
THYM (n=4)
LAML (n=16)
PRAD (n=15)
SKCM (n=24)
PCPG (n=5)
GBM (n=17)
LGG (n=24)
UCEC (n=55)
STAD (n=24)
KIRP (n=9)
BLCA (n=45)
CHOL (n=5)
COADREAD (n=20)
DLBC (n=13)
LIHC (n=32)
OV (n=8)
UCS (n=14)
LUAD (n=20)
CESC (n=26)
BRCA (n=29)
PANCAN (n=165)
LUSC (n=22)
PAAD (n=11)
HNSC (n=33)
ESCA (n=13)
ACC (n=5)
KIRC (n=12)
KICH (n=2)
MESO (n=5)
SARC (n=3)
Unknown
Possible Tumor Suppressor
Likely Tumor Suppressor
Possible Oncogene
Likely Oncogene
Consensus gene score
Enrichment
Adjusted P < 0.001
Adjusted P < 0.01
Adjusted P < 0.05
Not significant
Cluster
Pan-gastrointestinal
Pan-gynecological
Pan-squamous