0.017 0.001 0.000

FDR, median

0.050

0.014

0.014

Hellmann

**TMB** 

all MHC binders

0.010 0.050

0.009

0.007 0.002

0.021 0.000 Sombined NSCLC

0.01 **CDNs ADNs** 

**IEDB** high

0.05

0.04

0.03

0.02

high dissimilarity

all dissimilar neopeptides