all MHC binders
rho = 0.9488
p = 5.83e−159

CDNs
rho = 0.9282
p = 1.38e−136

ADNs
rho = 0.9259
p = 1.72e−134
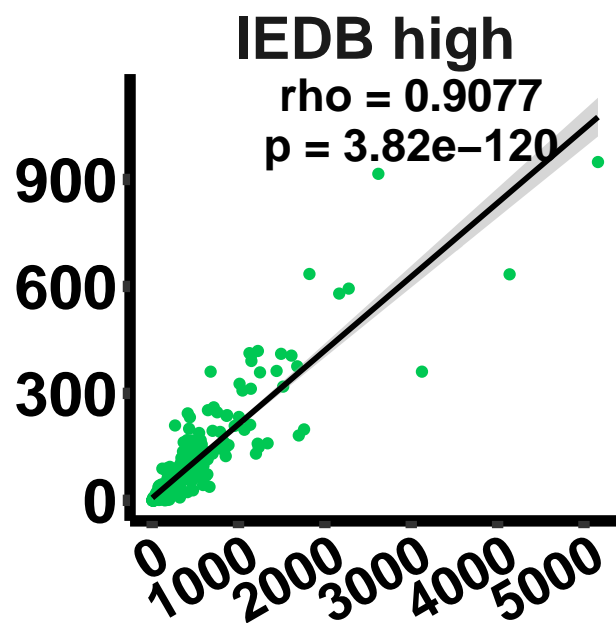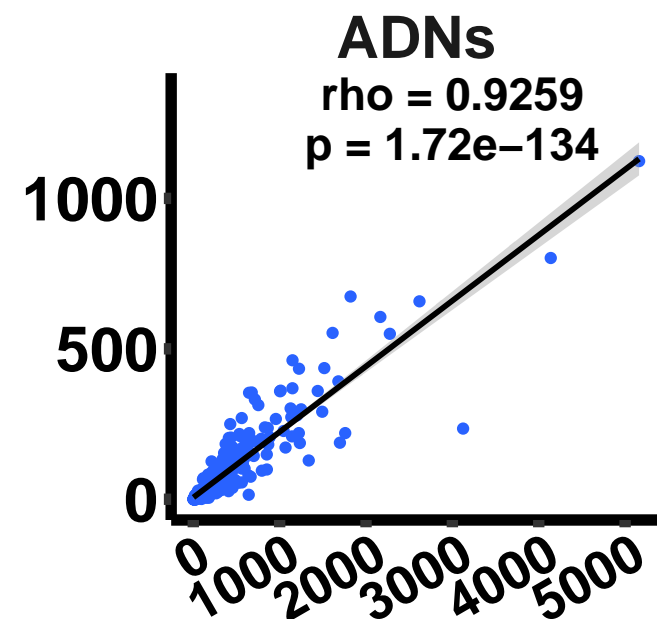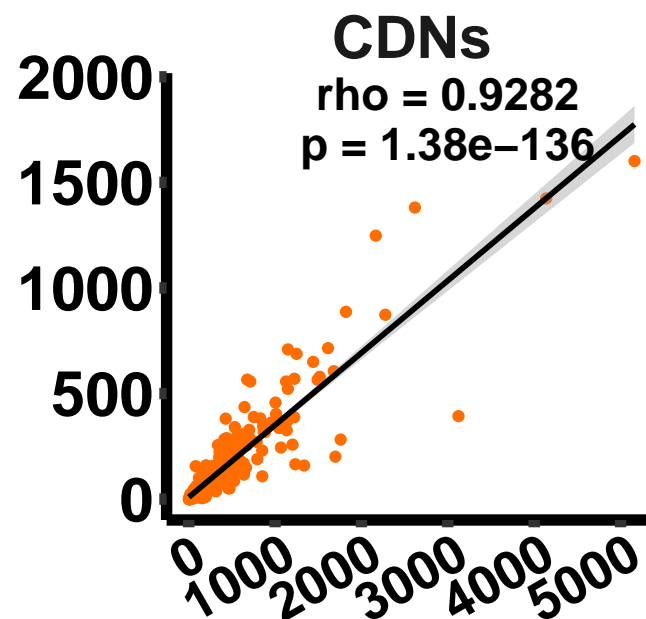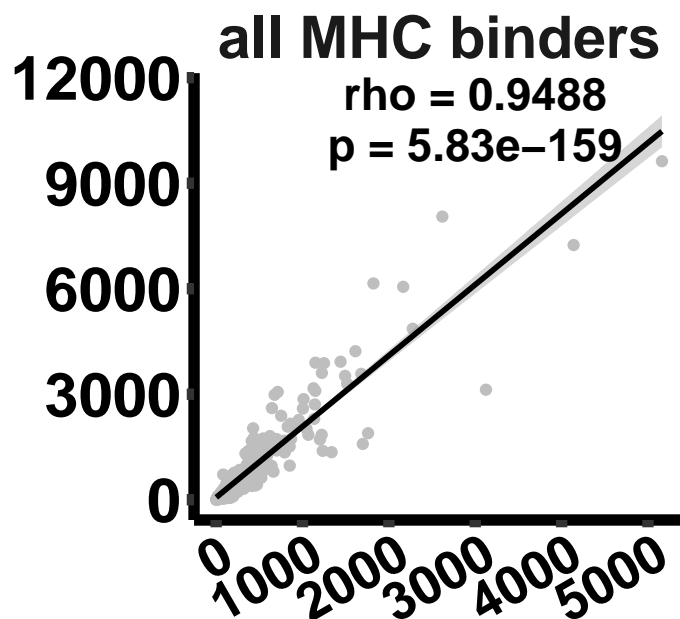
IEDB high
rho = 0.9077
p = 3.82e−120

high dissimilarity
rho = 0.8525
p = 4.87e−90

Peptides

Tumor mutational burden