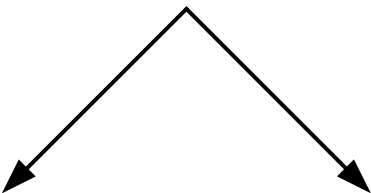
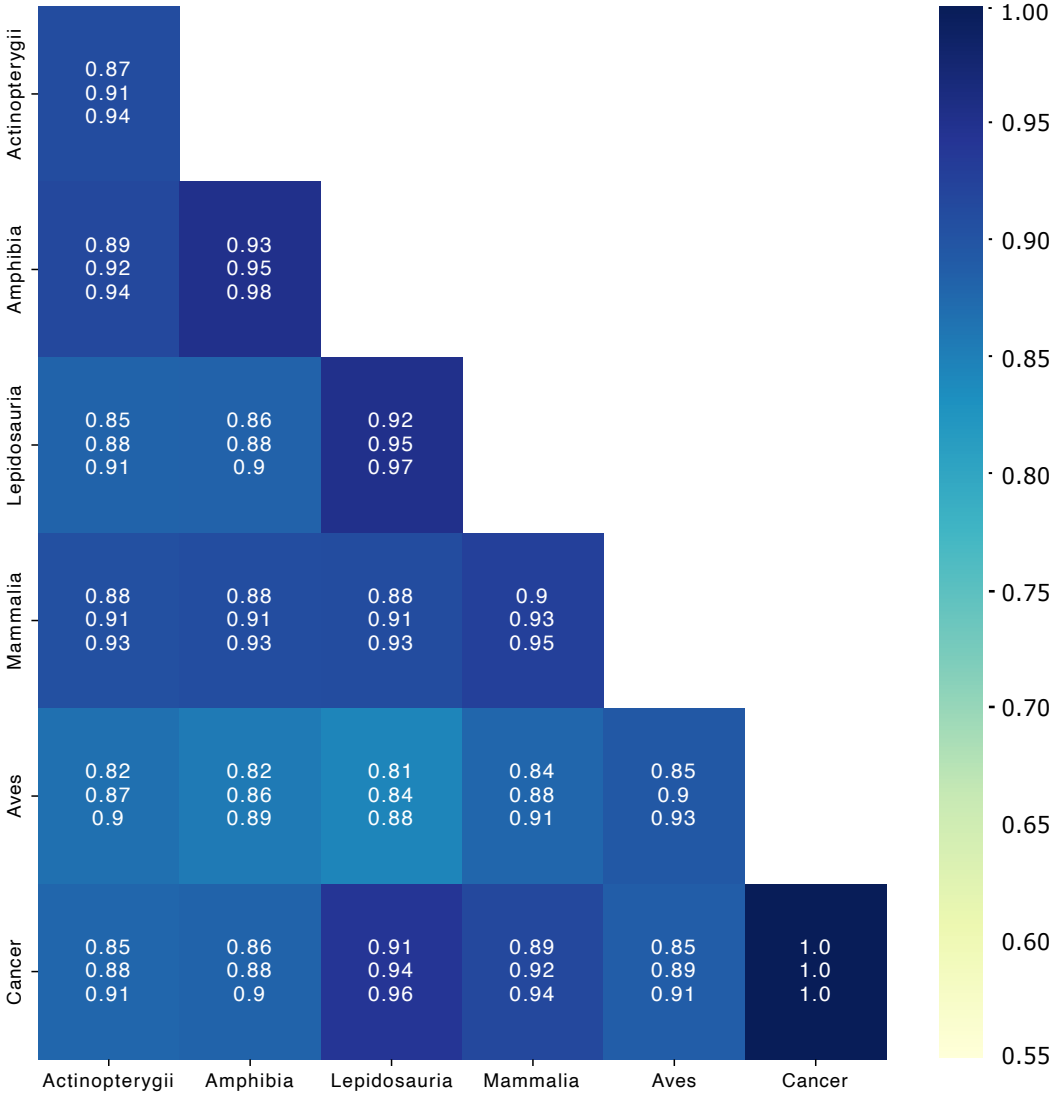


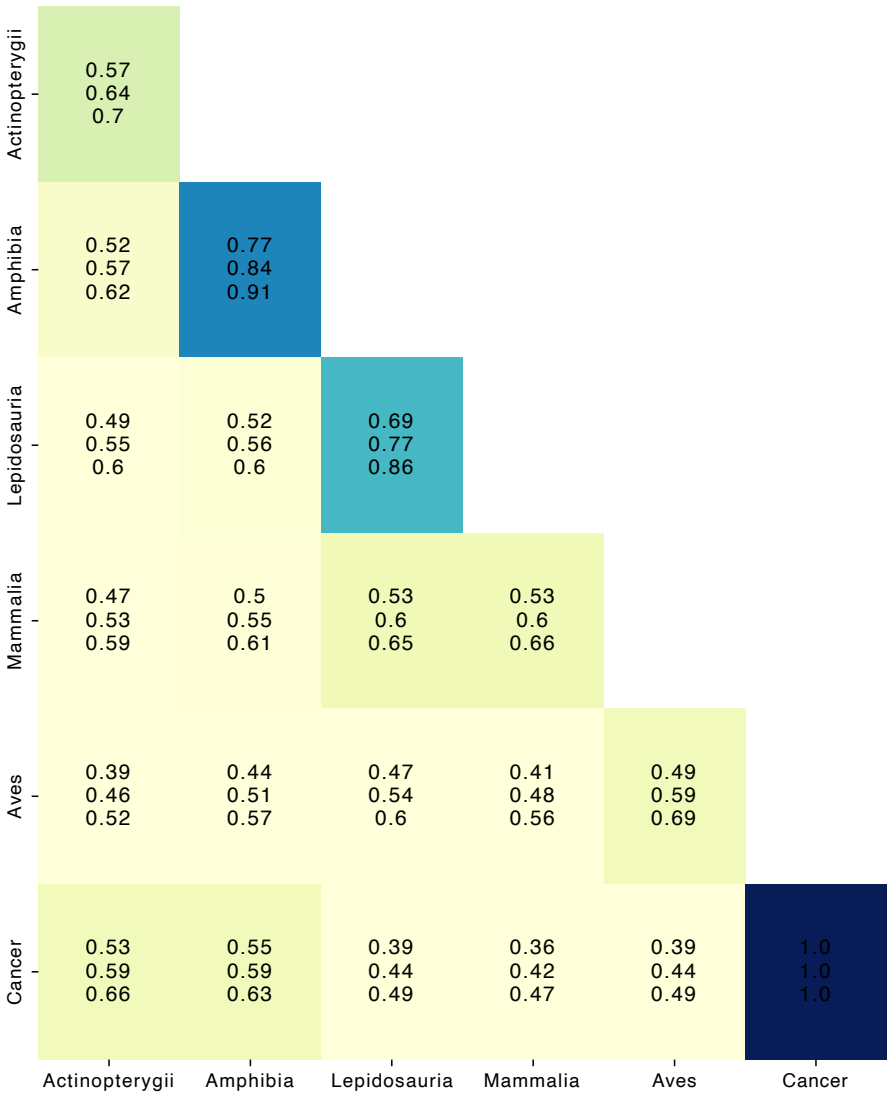
a

all mutations (n=192)



b

transitions (n=4 x 16)



c

transversions (n=8 x 16)

