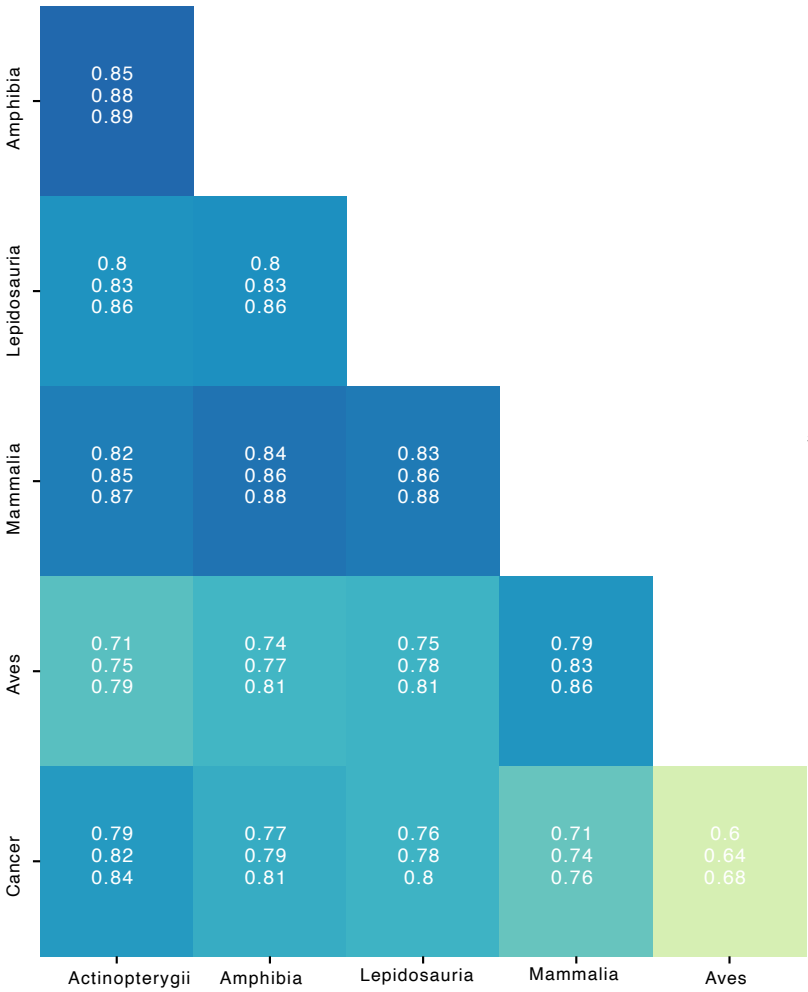
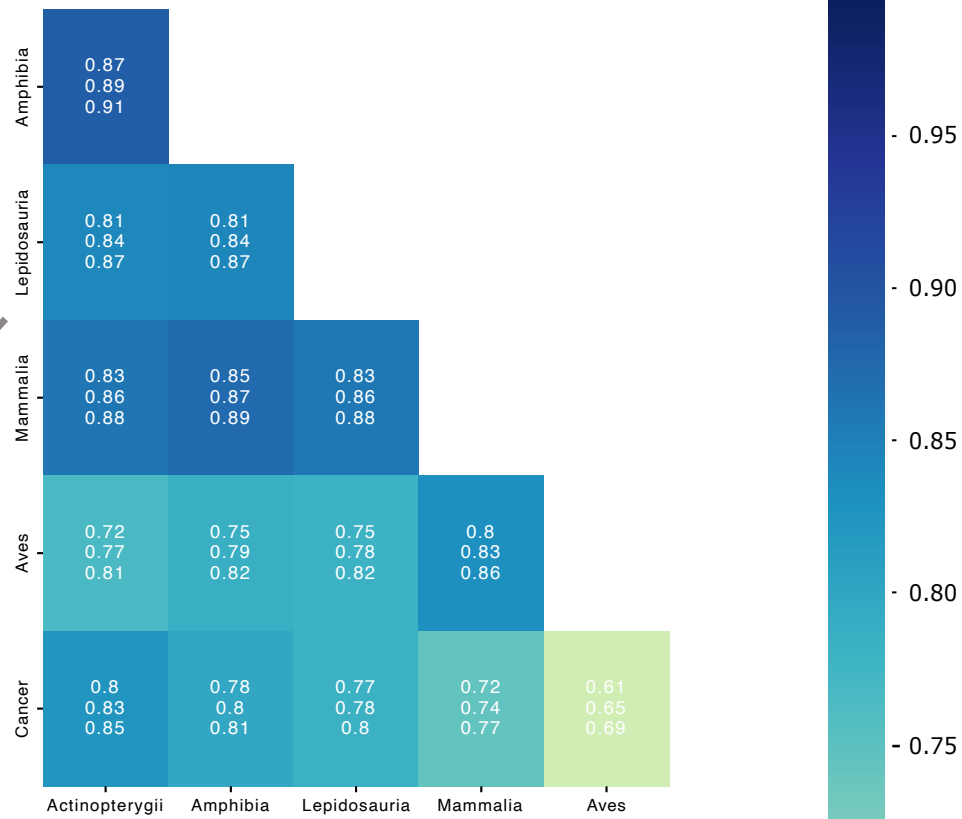


**a** all mutations (n=192)



**b** transitions (n=4 x 16)



**c** transversions (n=8 x 16)

