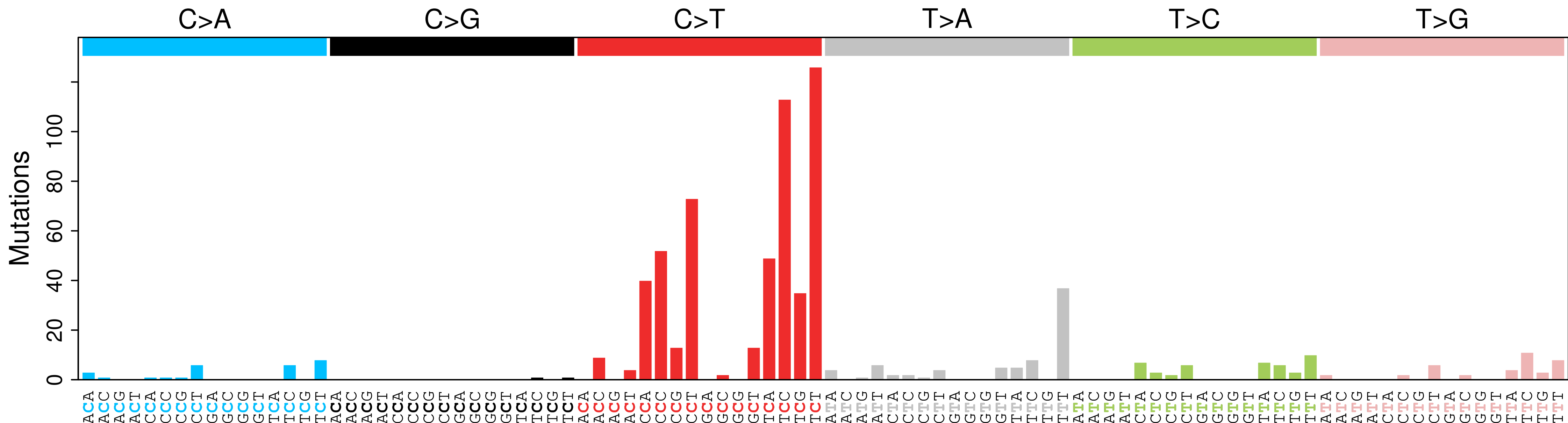


CATD268a (715 mutations)



Reconstructed spectrum (cosine similarity = 0.948)

