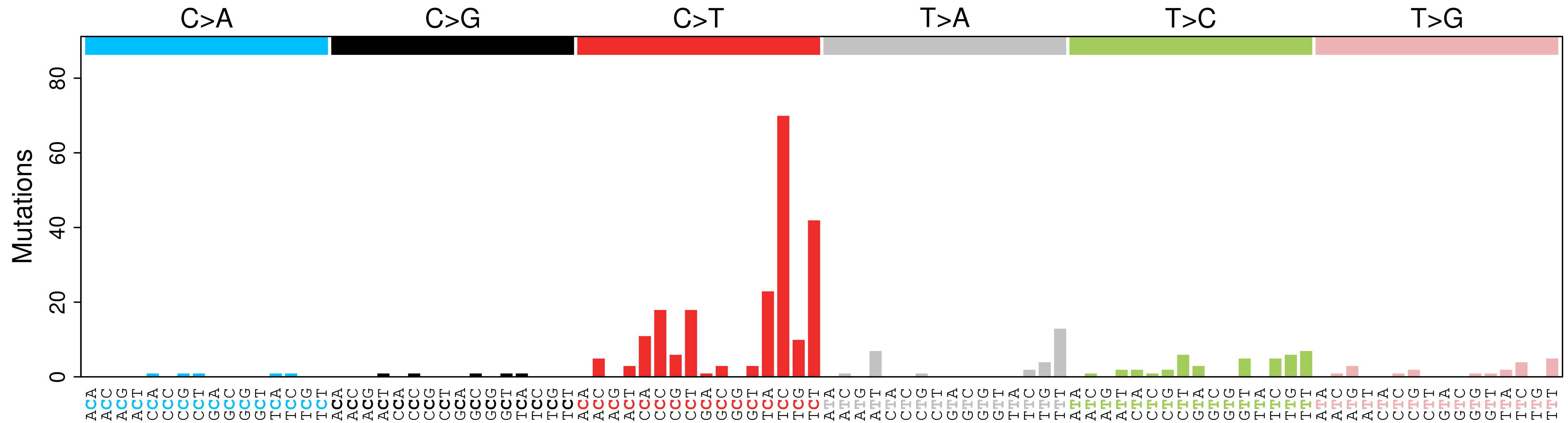


CATD273a (311 mutations)



Reconstructed spectrum (cosine similarity = 0.928)

