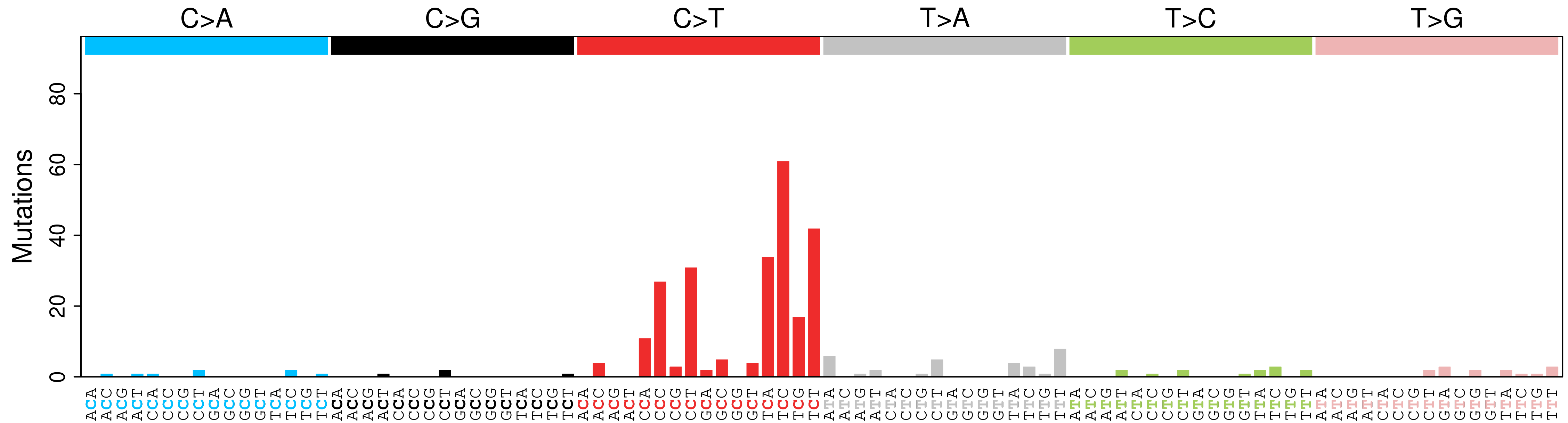


CATD0123a (311 mutations)



Reconstructed spectrum (cosine similarity = 0.962)

