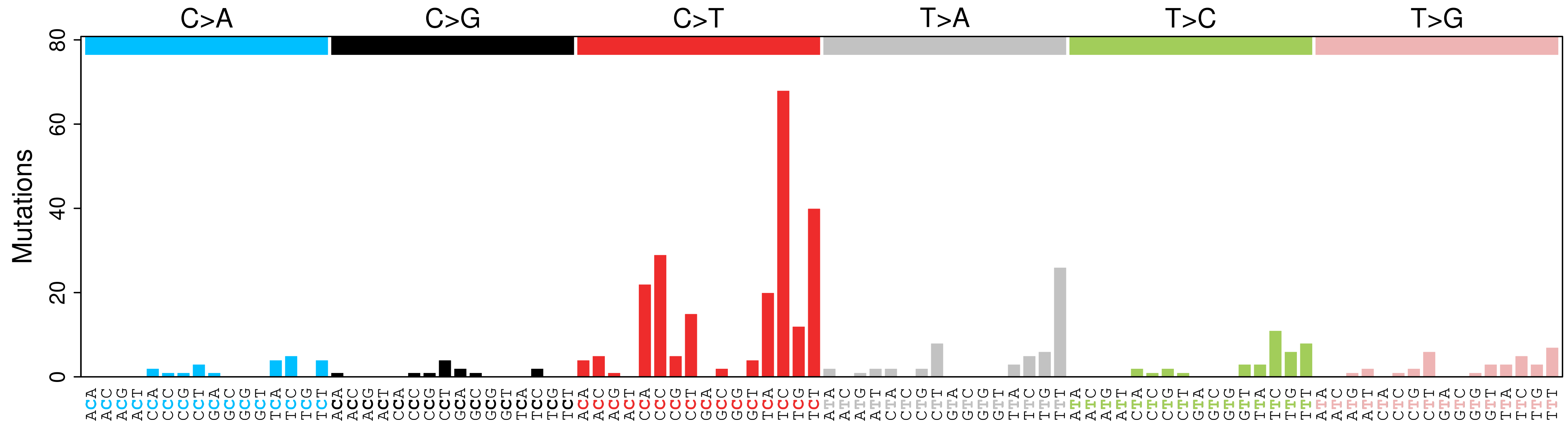


CATD260a (388 mutations)



Reconstructed spectrum (cosine similarity = 0.963)

