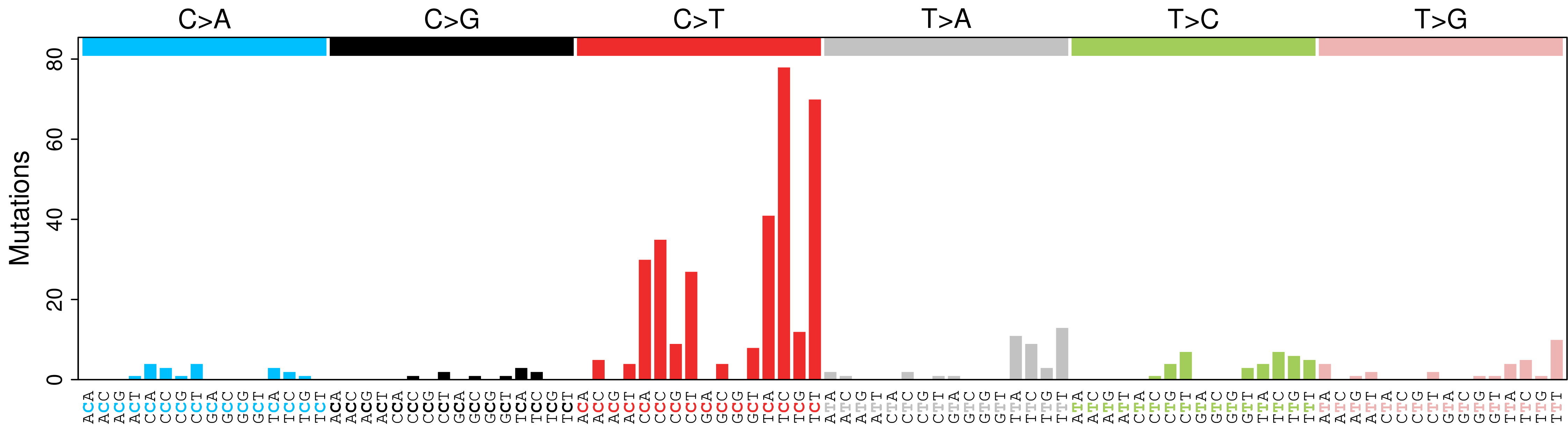


CATD248a (463 mutations)



Reconstructed spectrum (cosine similarity = 0.957)

