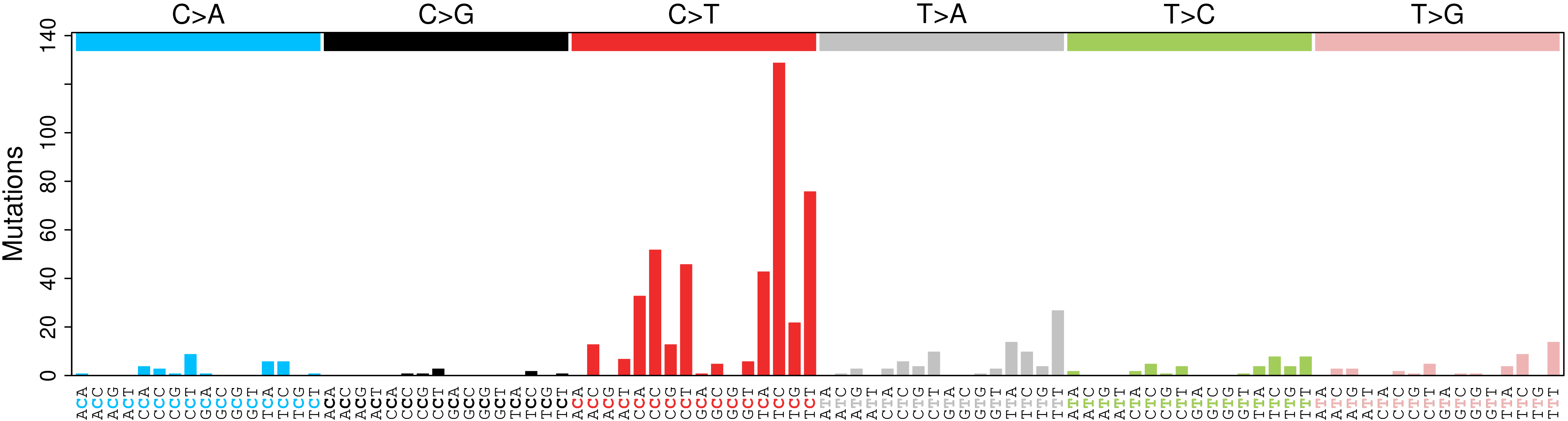


CATD0120a (654 mutations)



Reconstructed spectrum (cosine similarity = 0.972)

