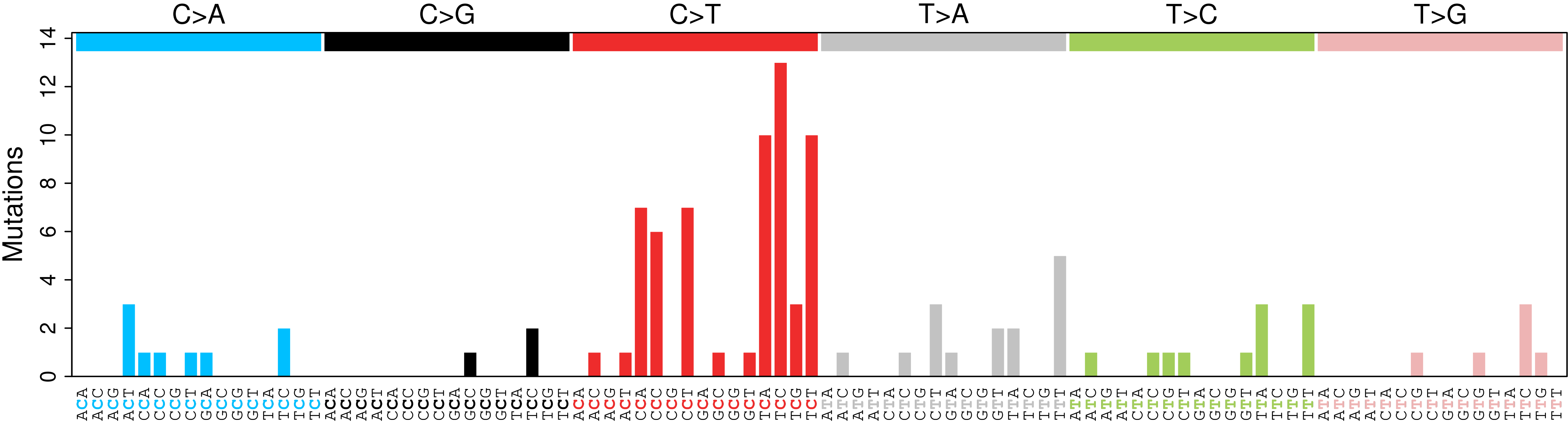


CATD0126a (104 mutations)



Reconstructed spectrum (cosine similarity = 0.787)

