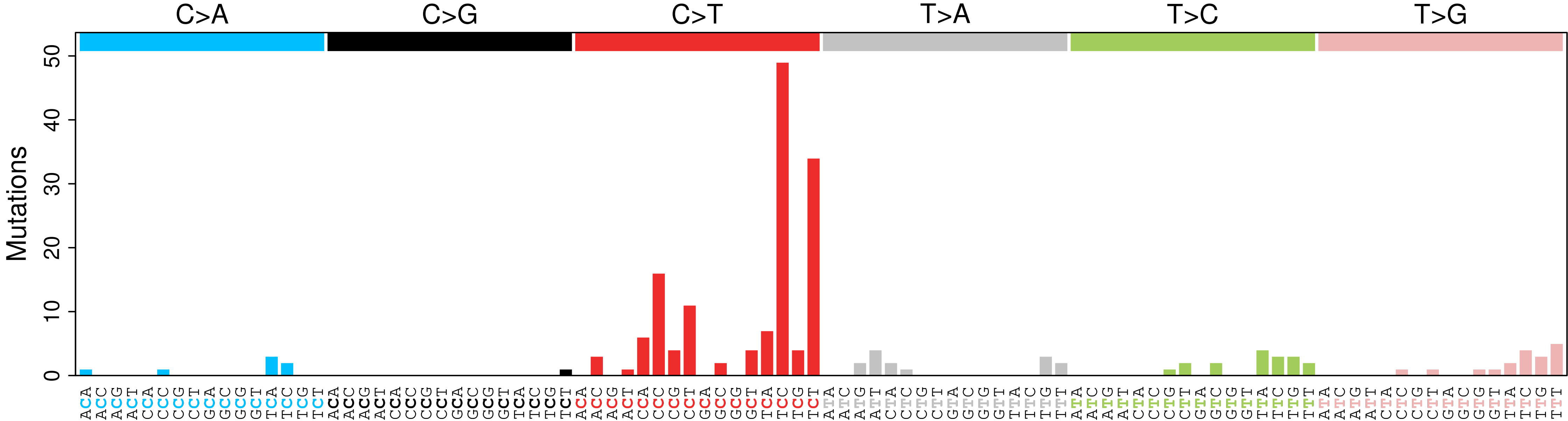


CATD245a (198 mutations)



Reconstructed spectrum (cosine similarity = 0.858)

