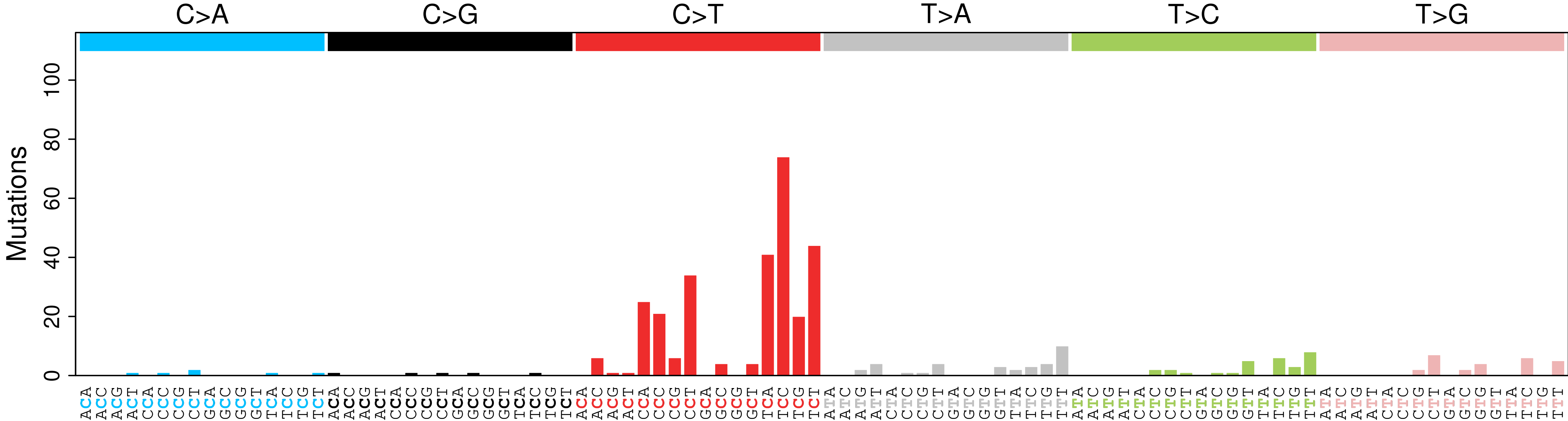


CATD0132a (381 mutations)



Reconstructed spectrum (cosine similarity = 0.961)

