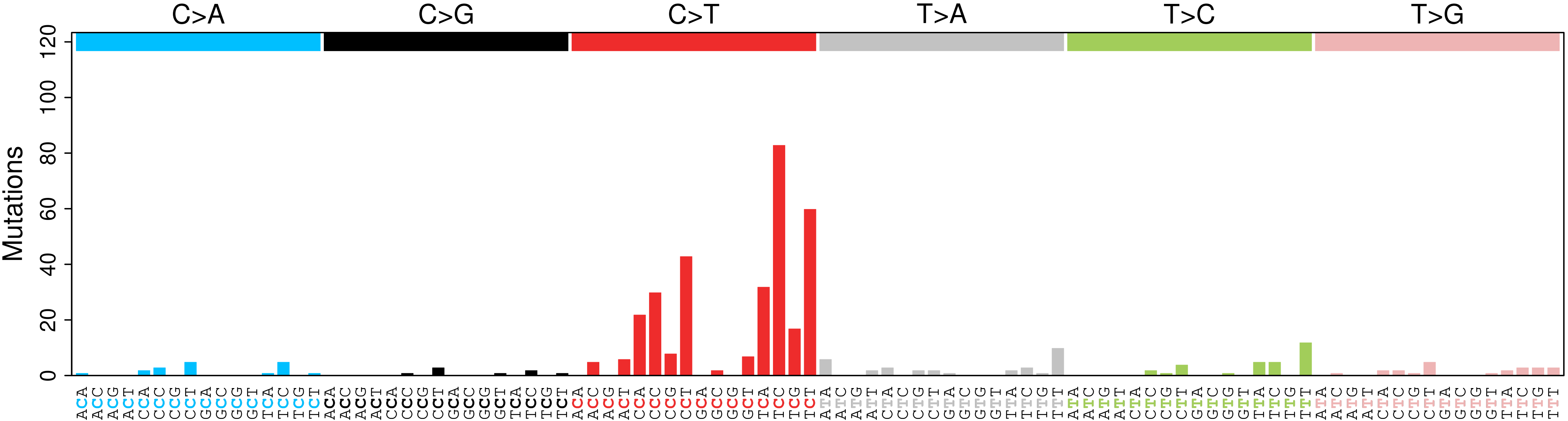


CATD278a (426 mutations)



Reconstructed spectrum (cosine similarity = 0.949)

