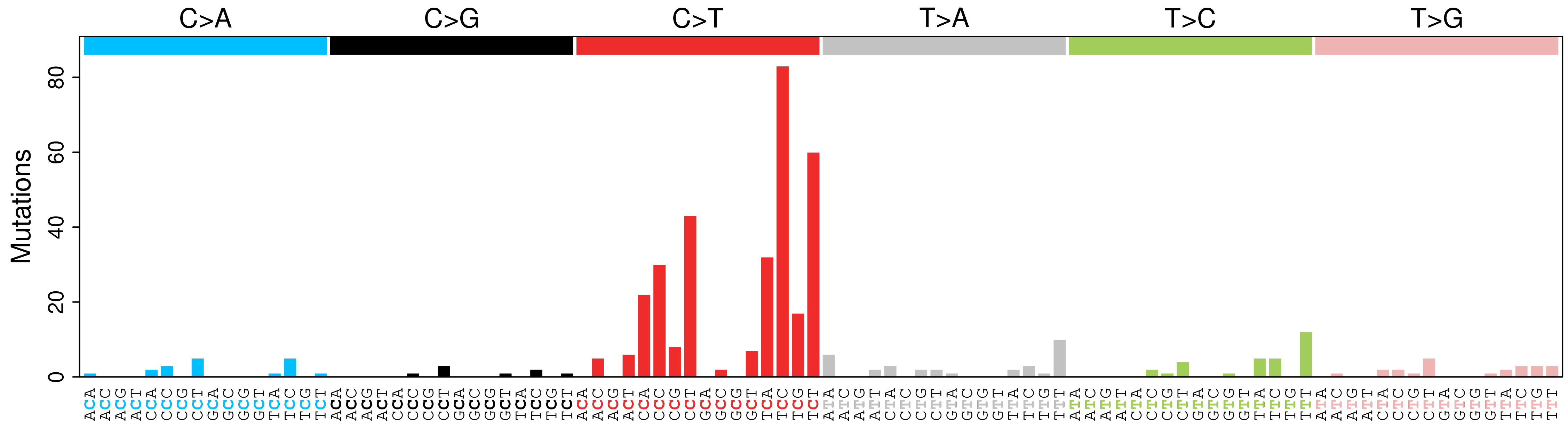


# CATD278a (426 mutations)



## Reconstructed spectrum (cosine similarity = 0.958)

