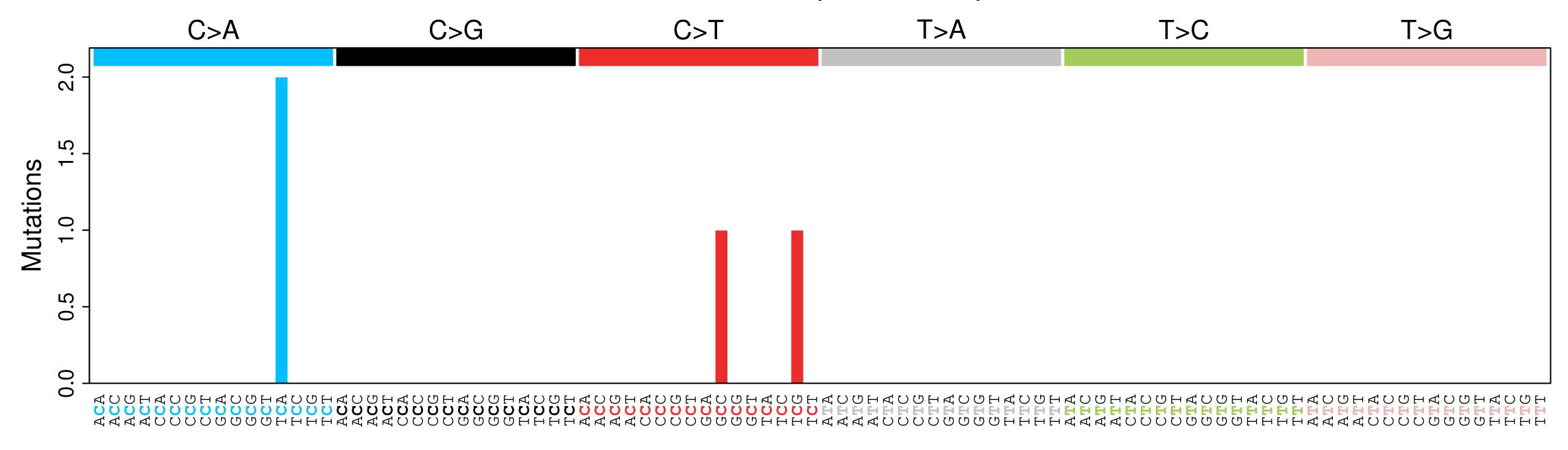
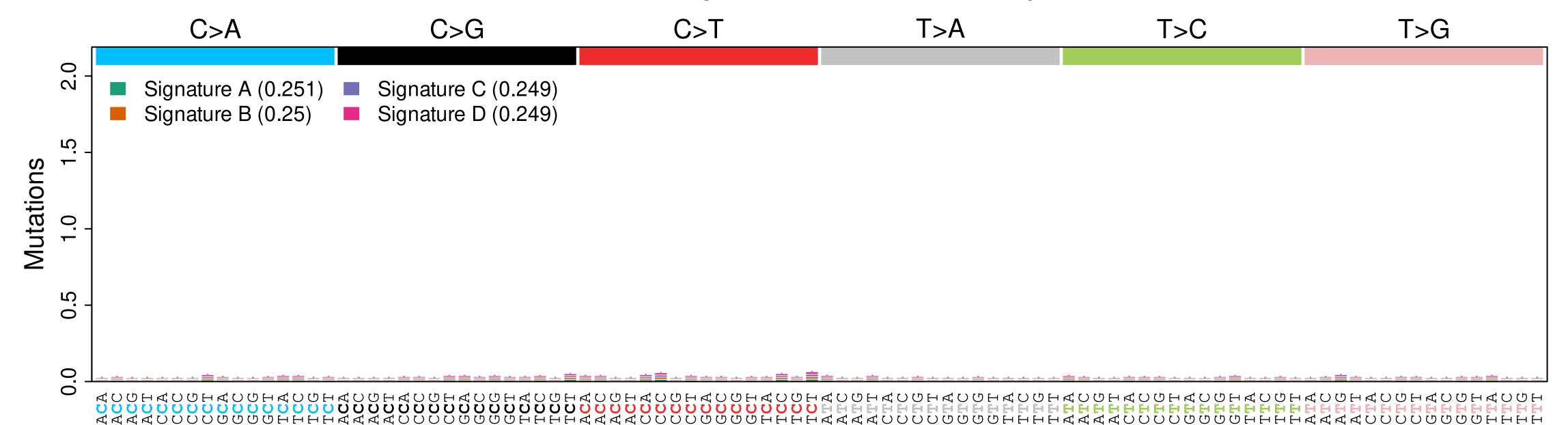
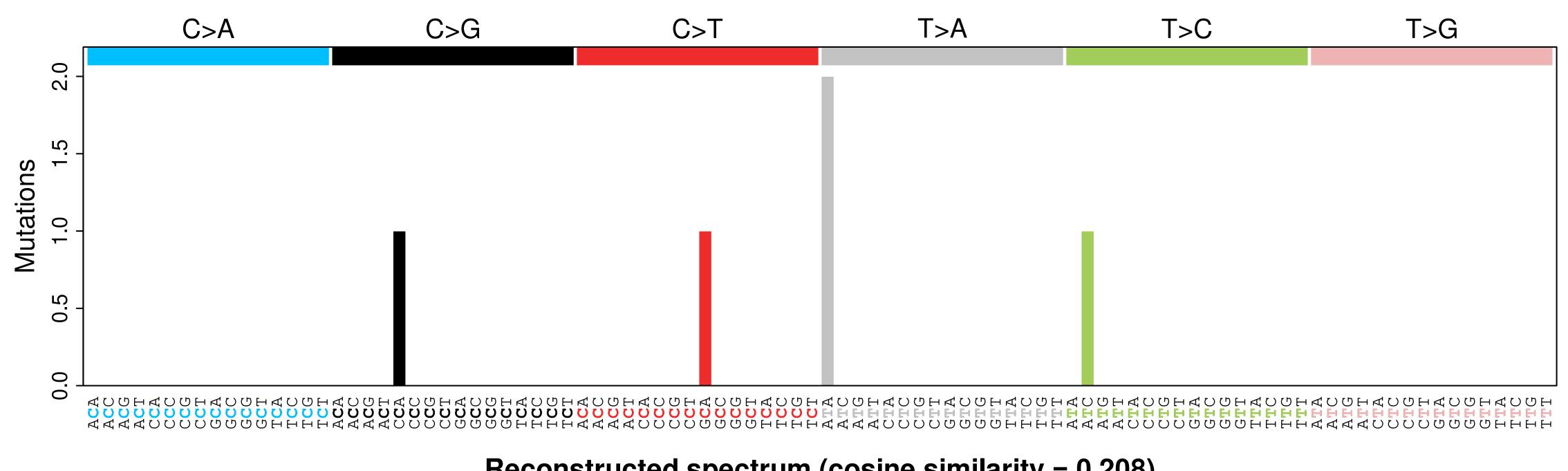
CATD0511a (4 mutations)



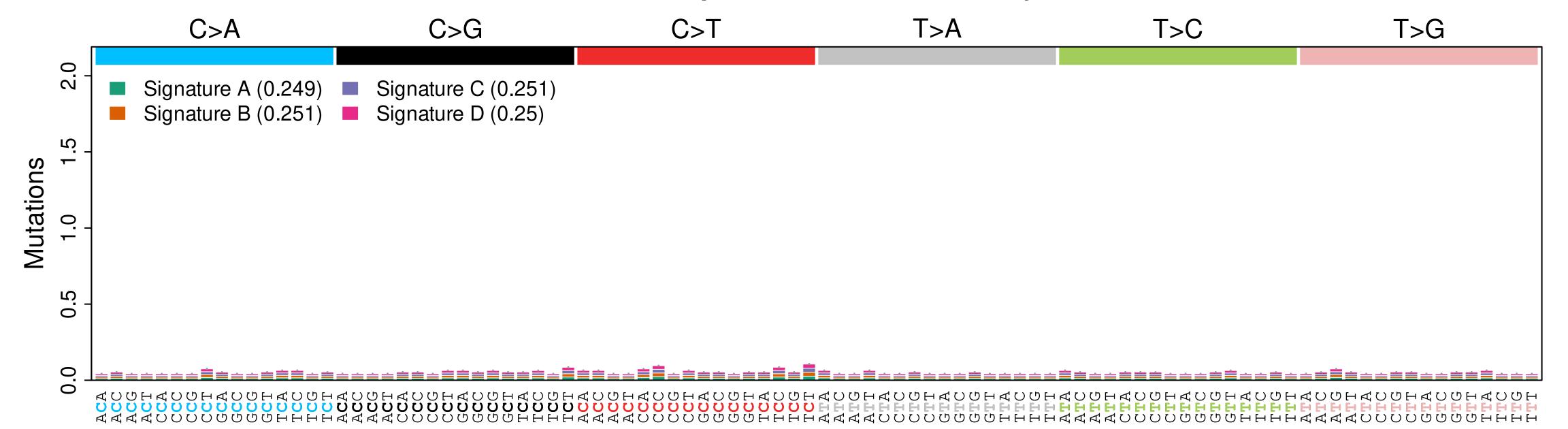
Reconstructed spectrum (cosine similarity = 0.183)



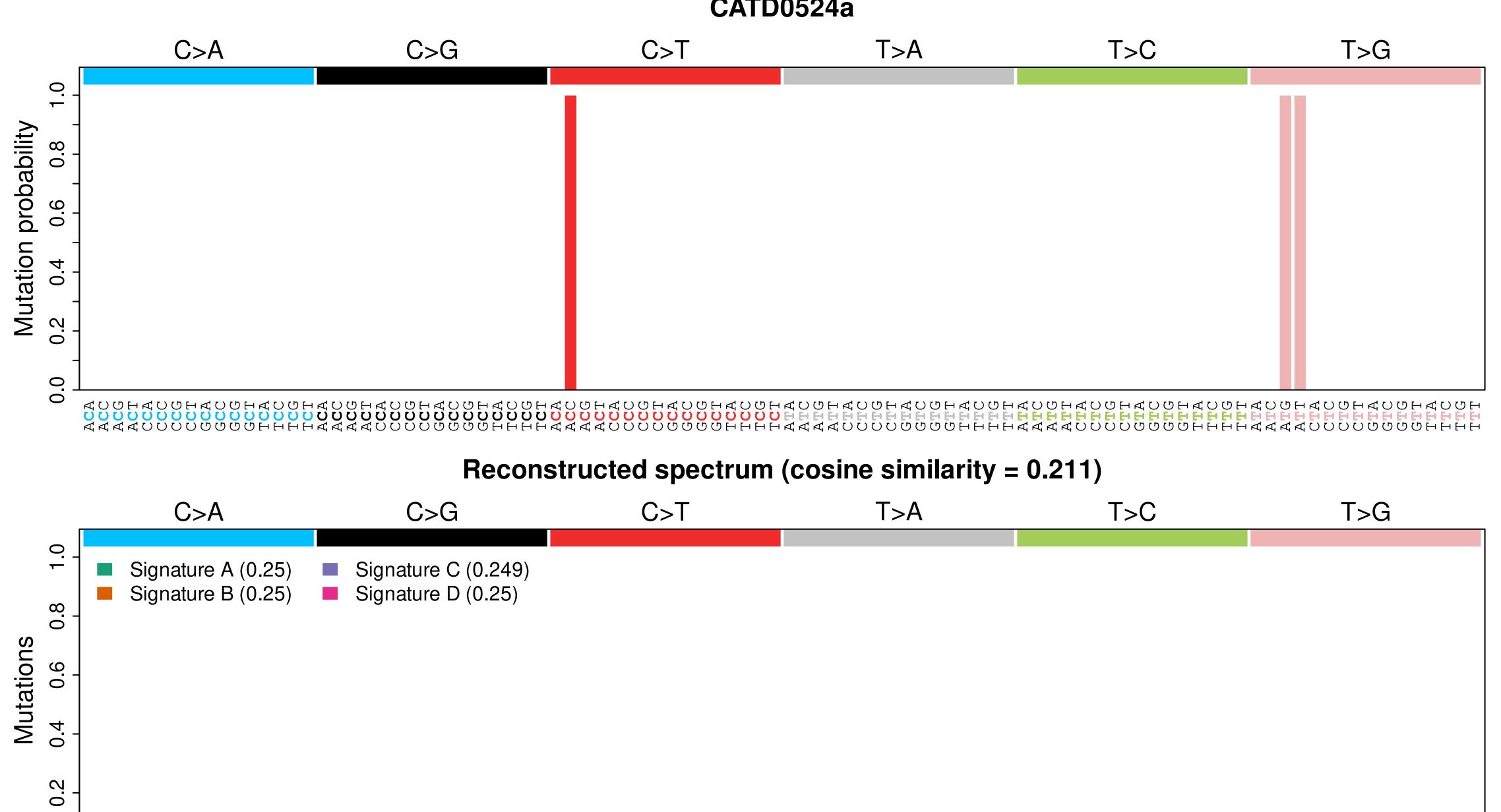
CATD0518a (5 mutations)



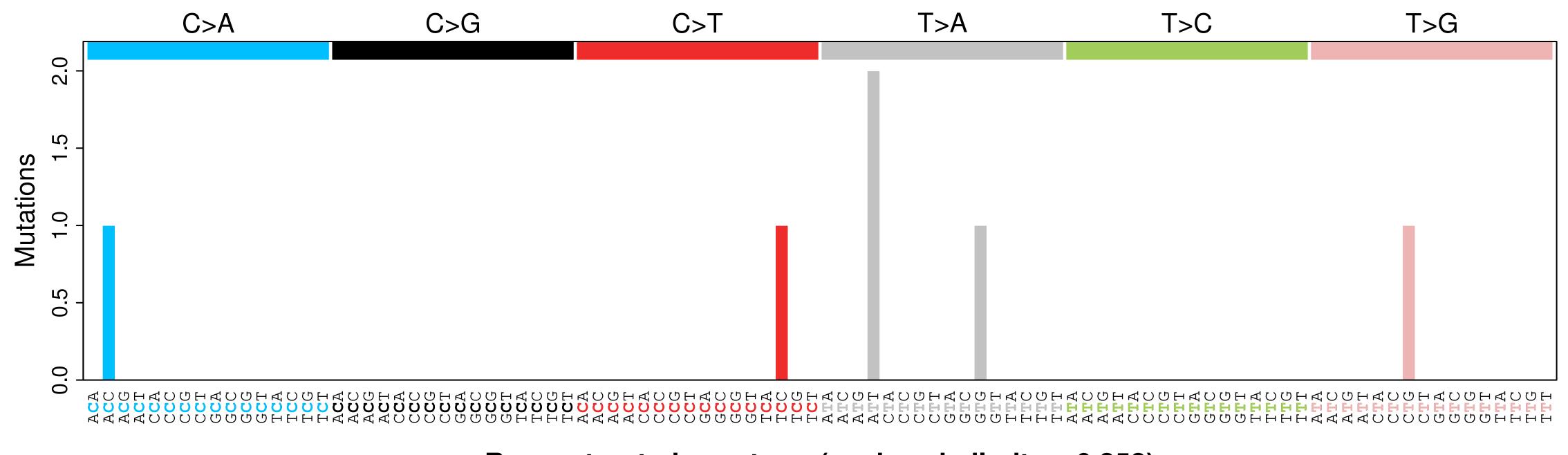




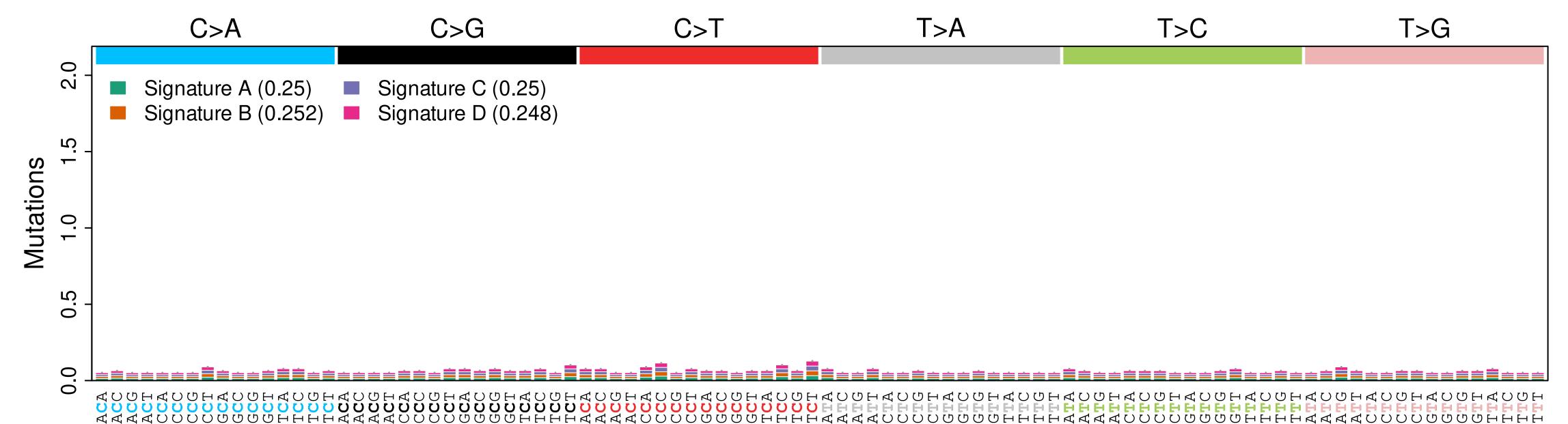




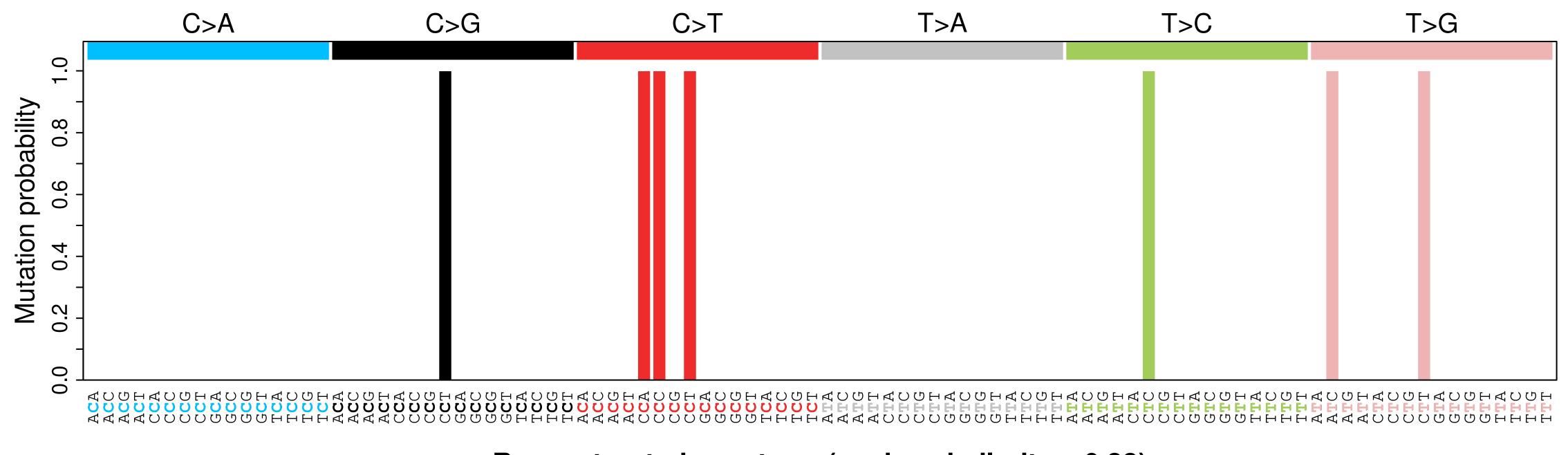
CATD0515a (6 mutations)



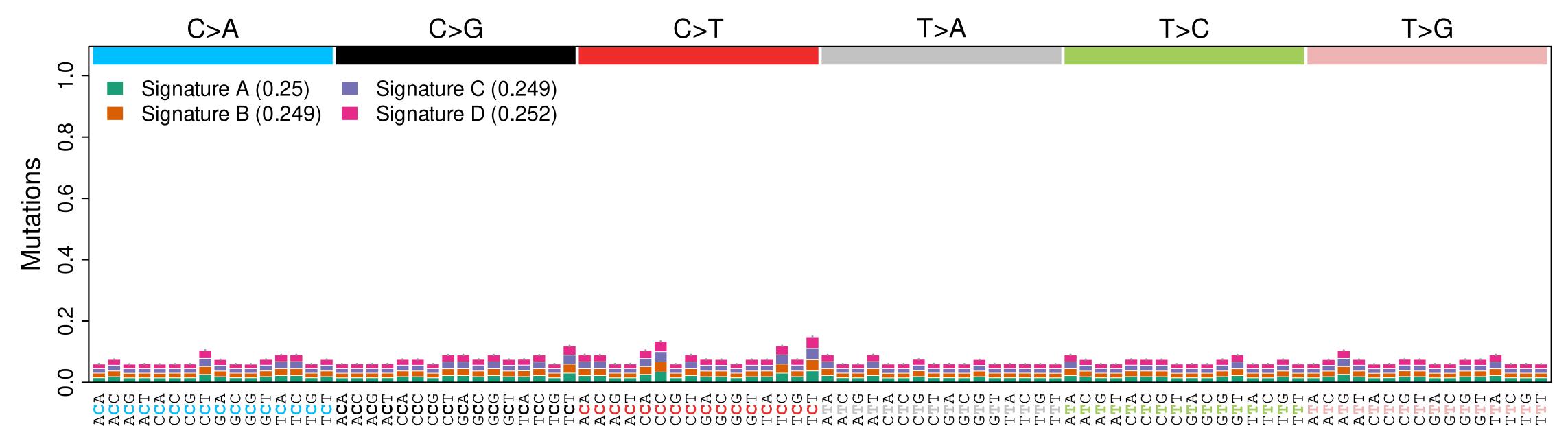
Reconstructed spectrum (cosine similarity = 0.252)



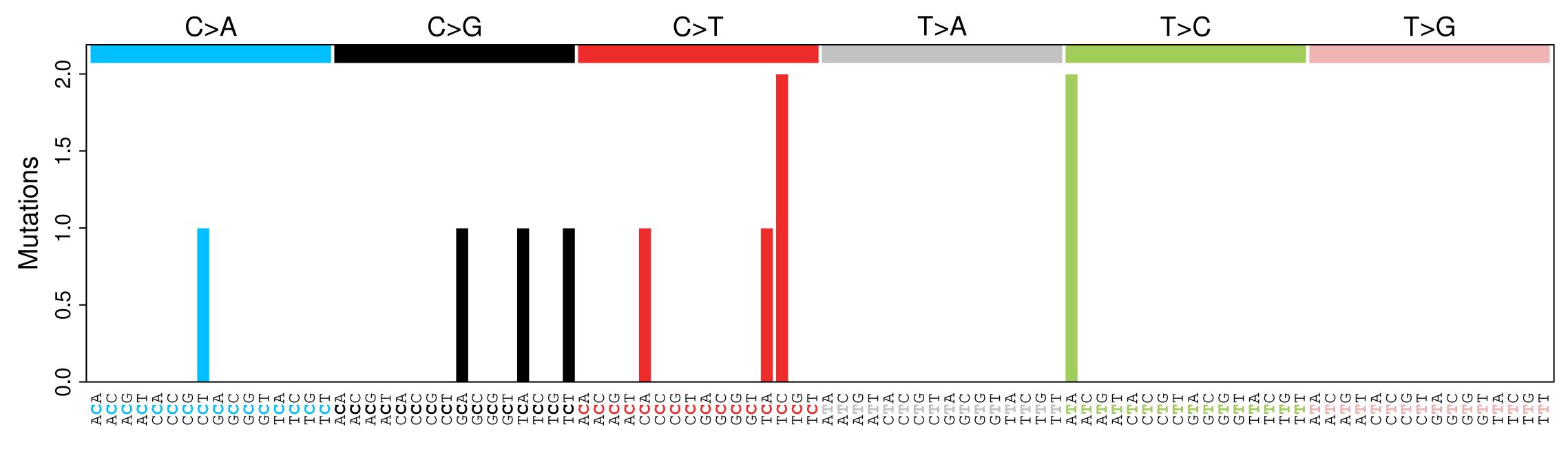
CATD196a



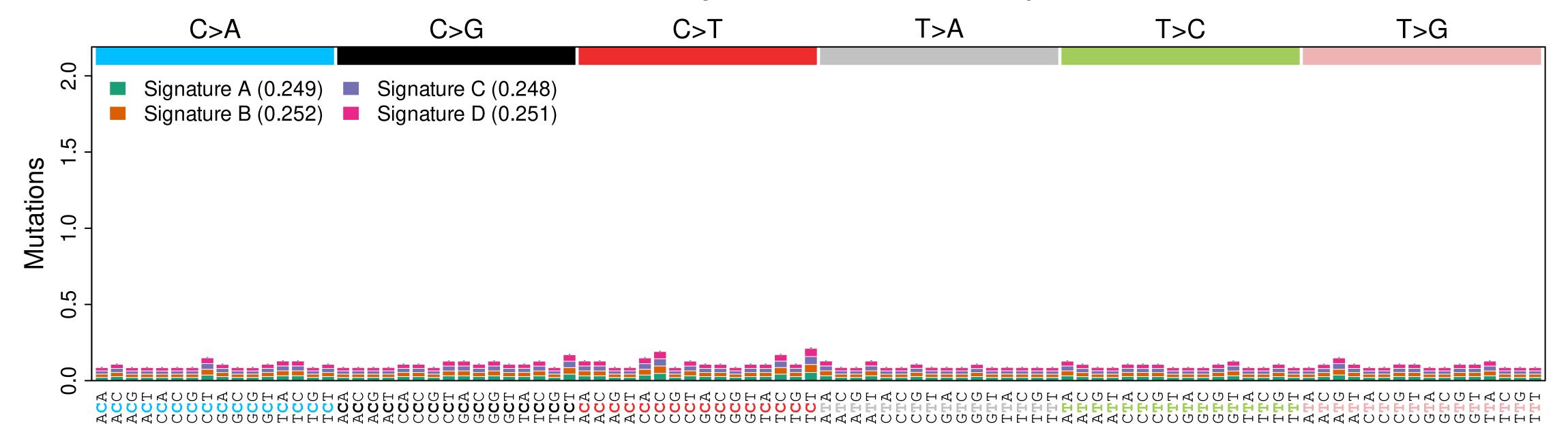
Reconstructed spectrum (cosine similarity = 0.33)



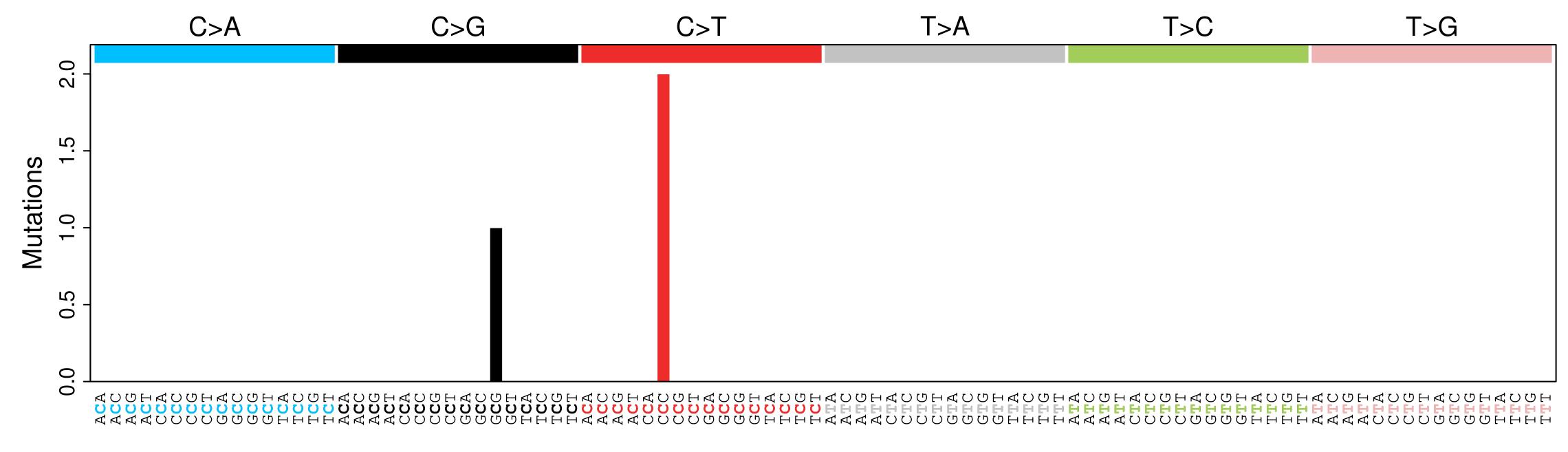
CATD205a (10 mutations)



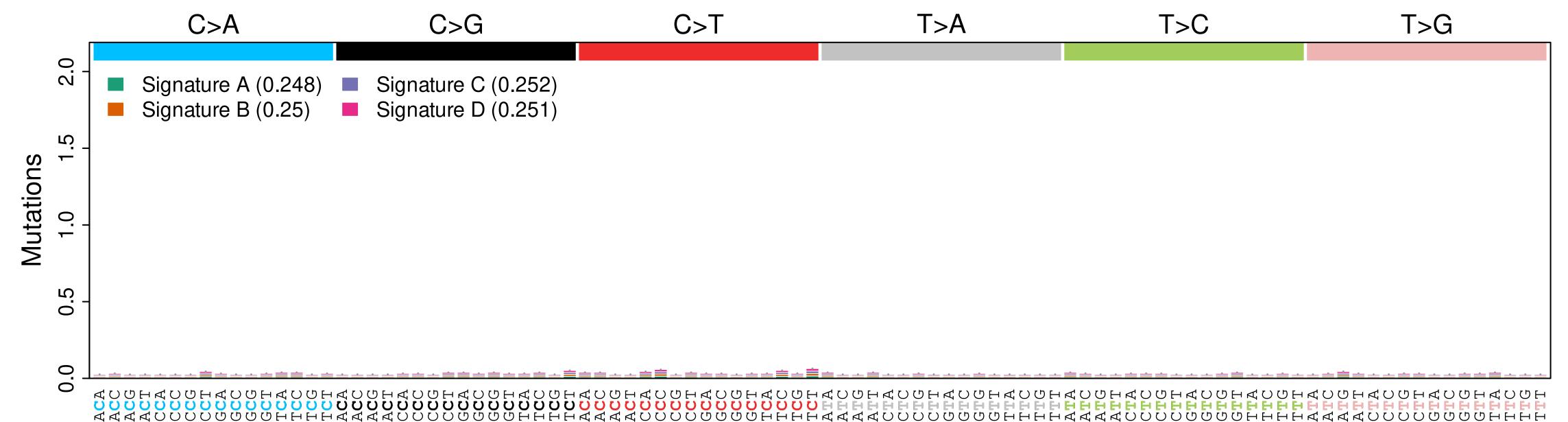
Reconstructed spectrum (cosine similarity = 0.358)



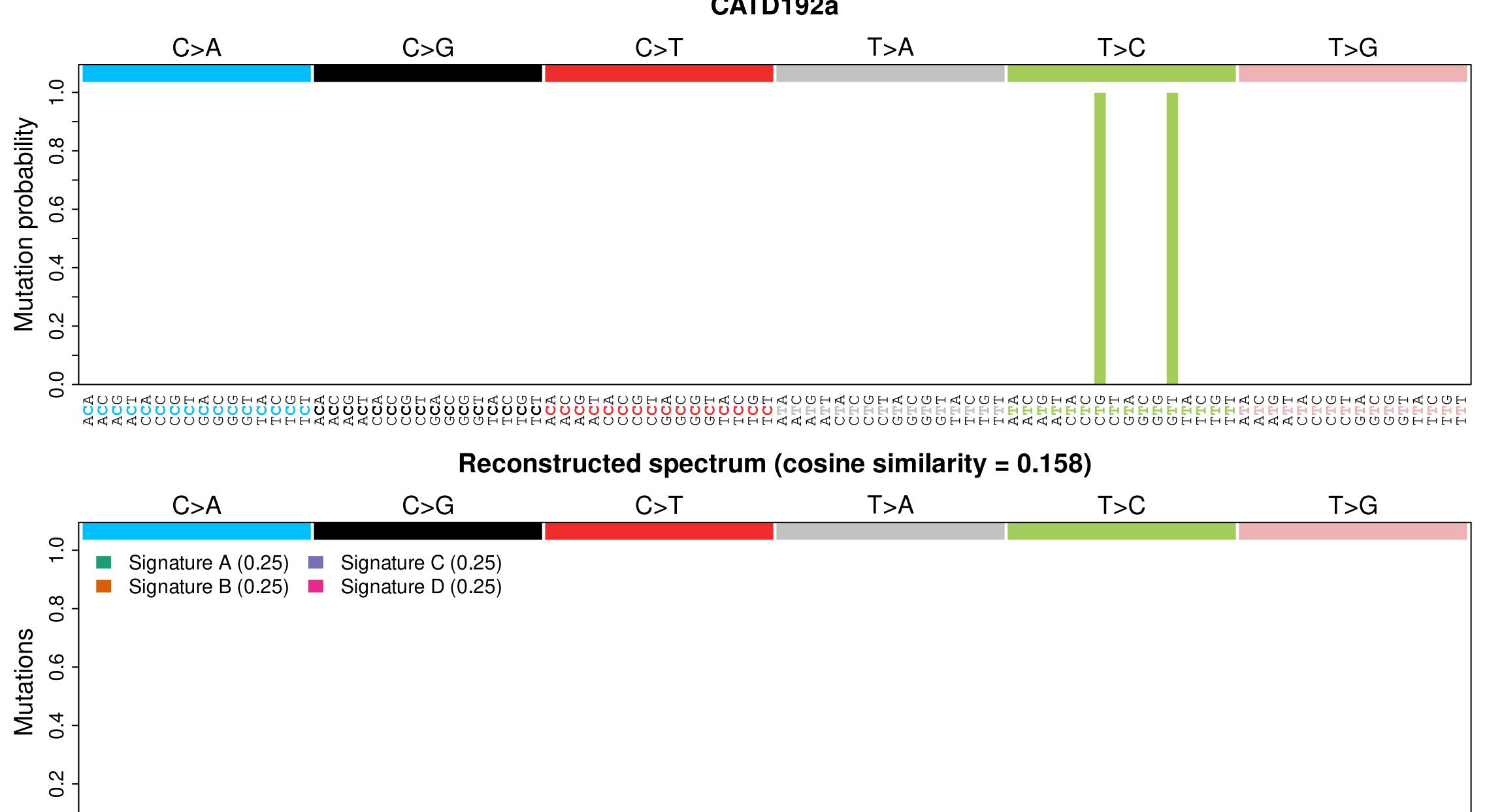
CATD193a (3 mutations)



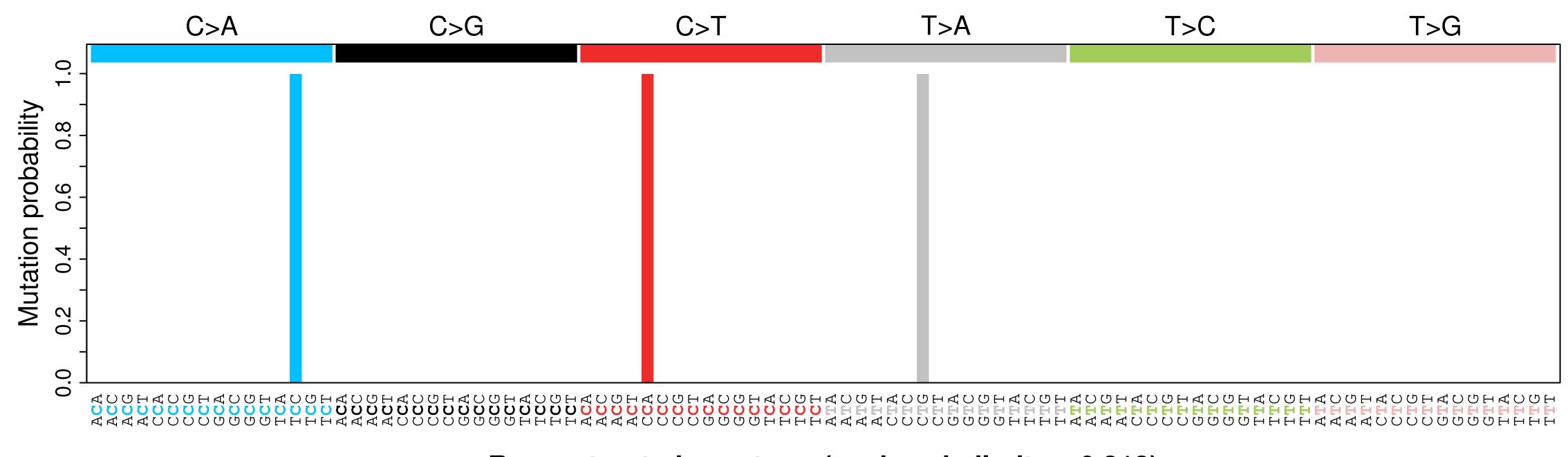
Reconstructed spectrum (cosine similarity = 0.217)

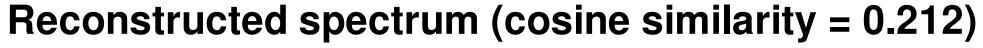


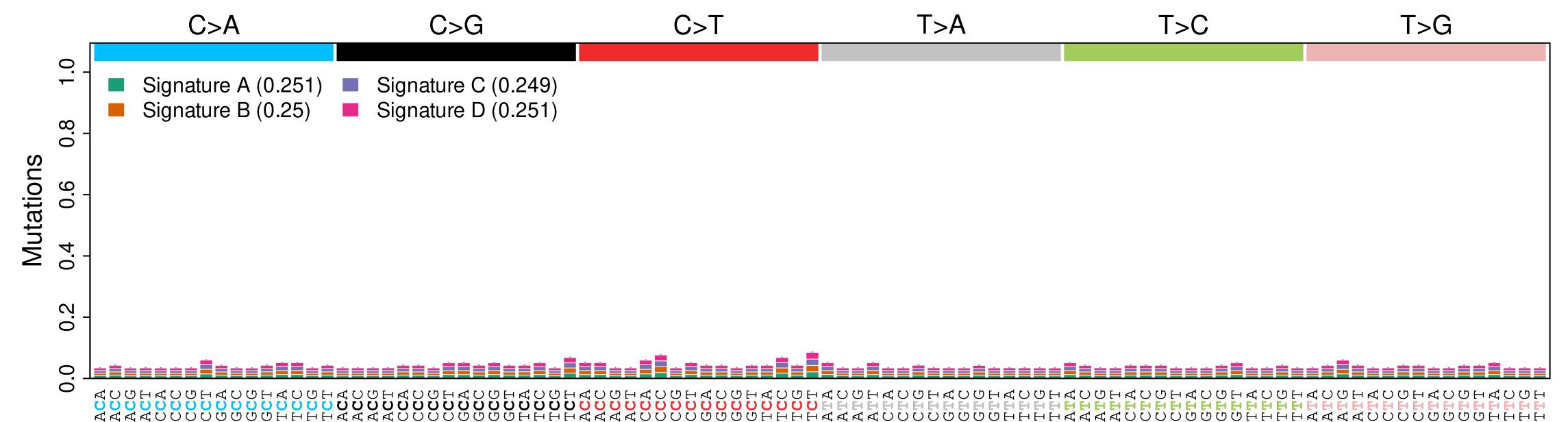




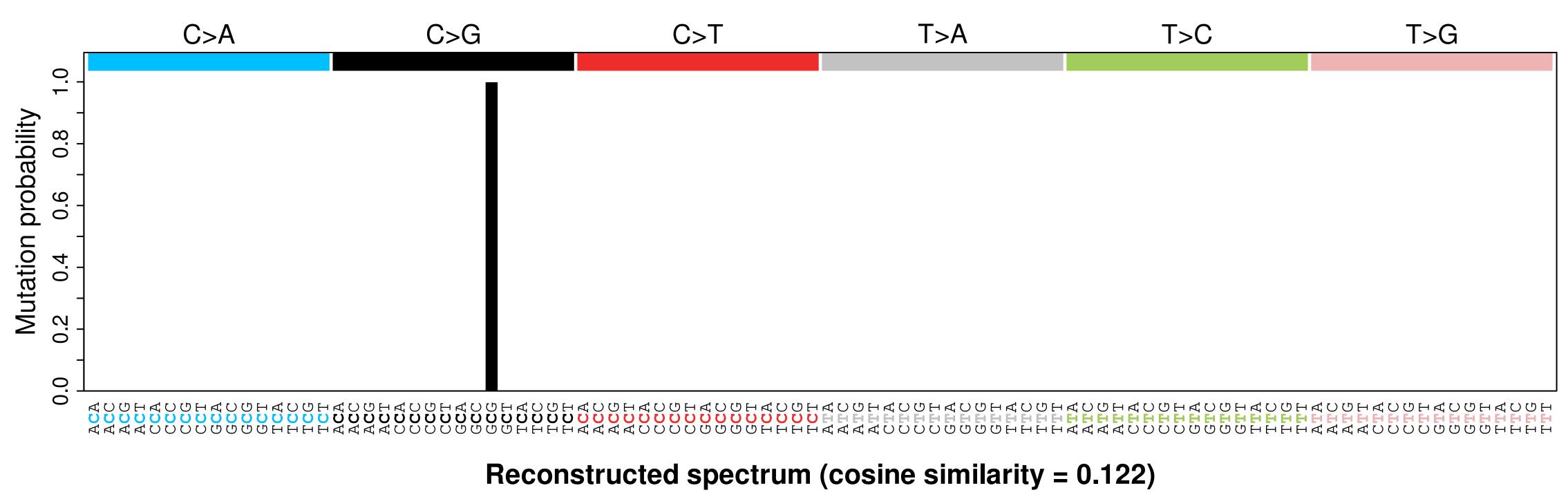


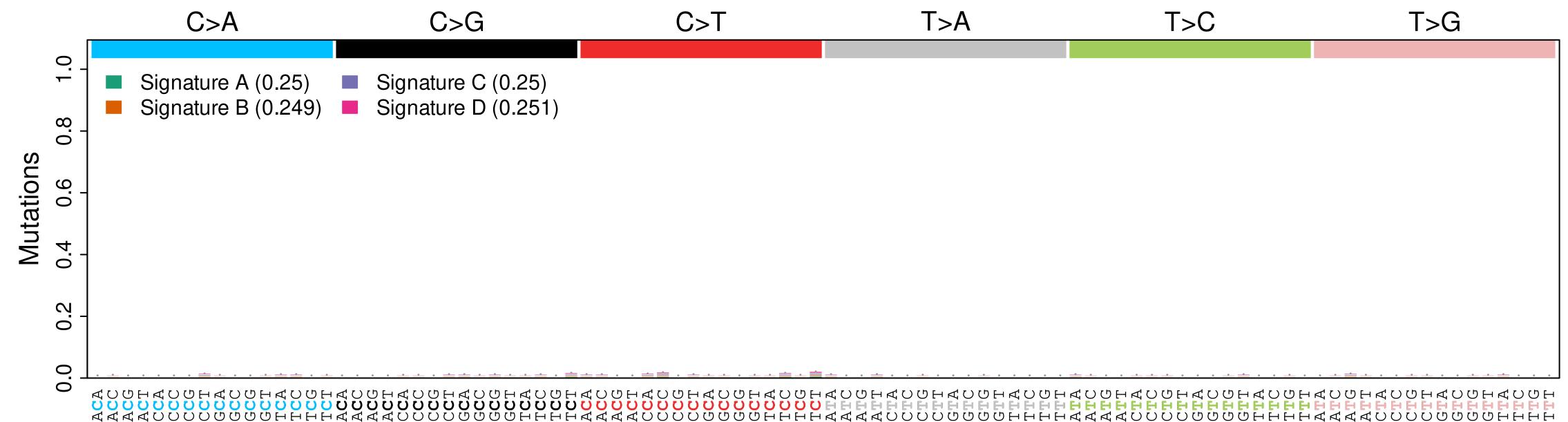




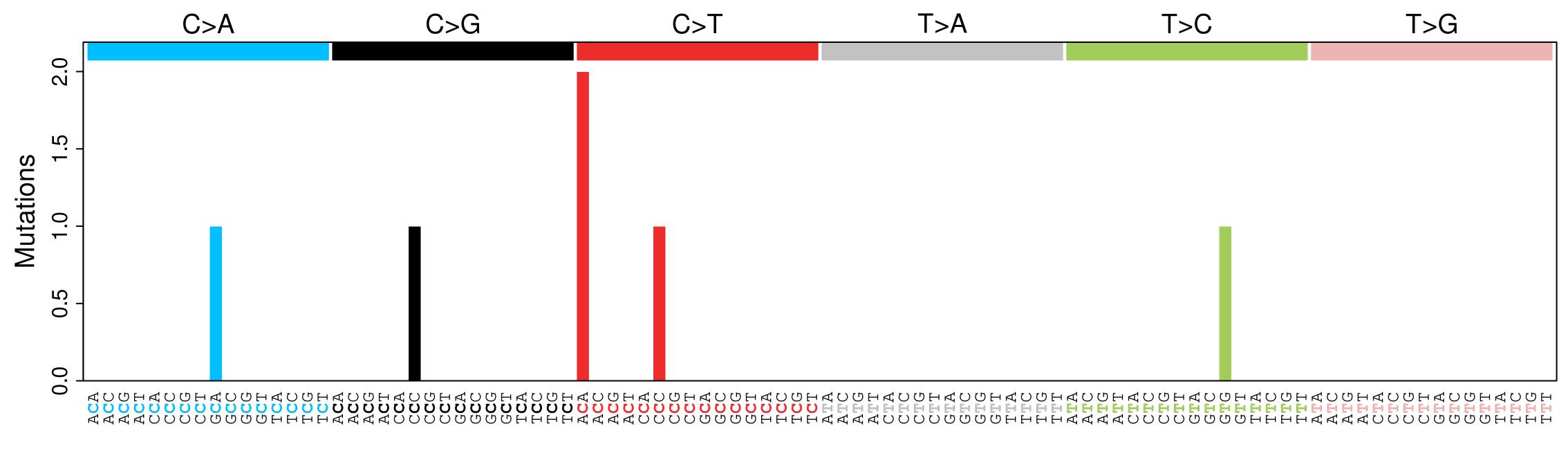




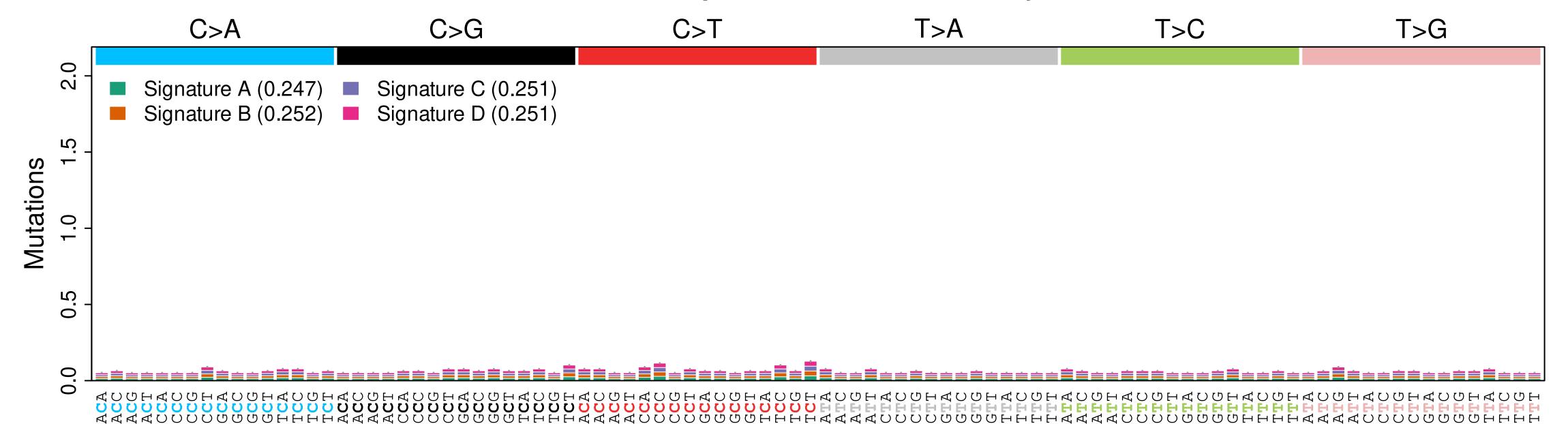




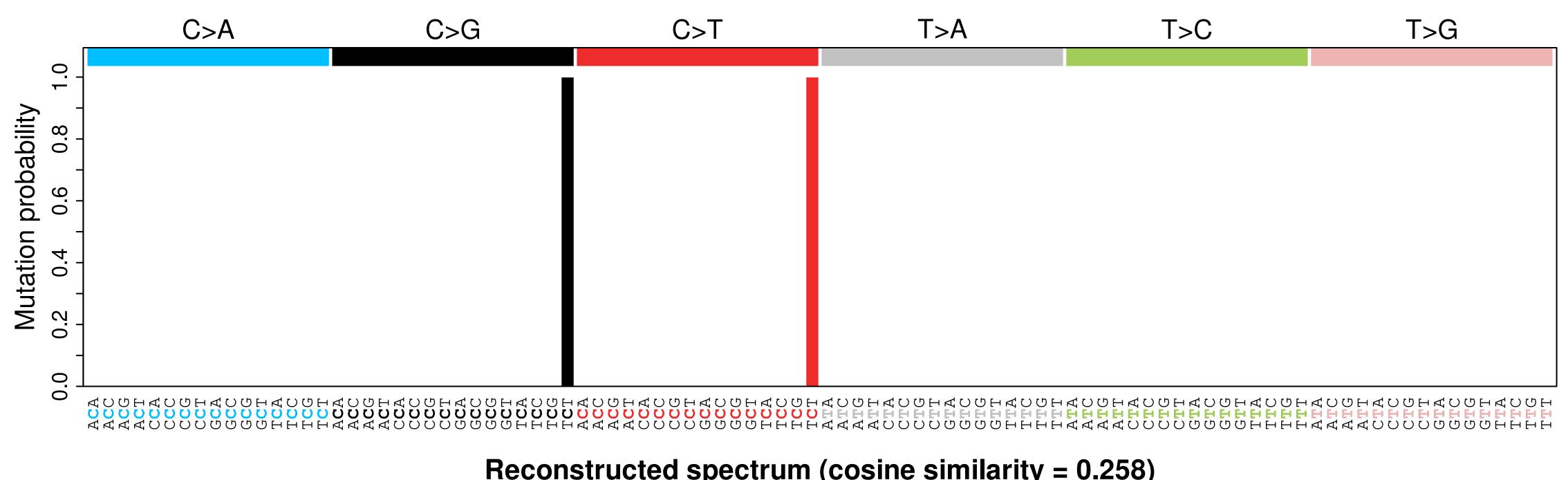
CATD0517a (6 mutations)

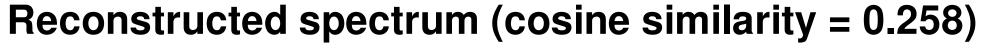


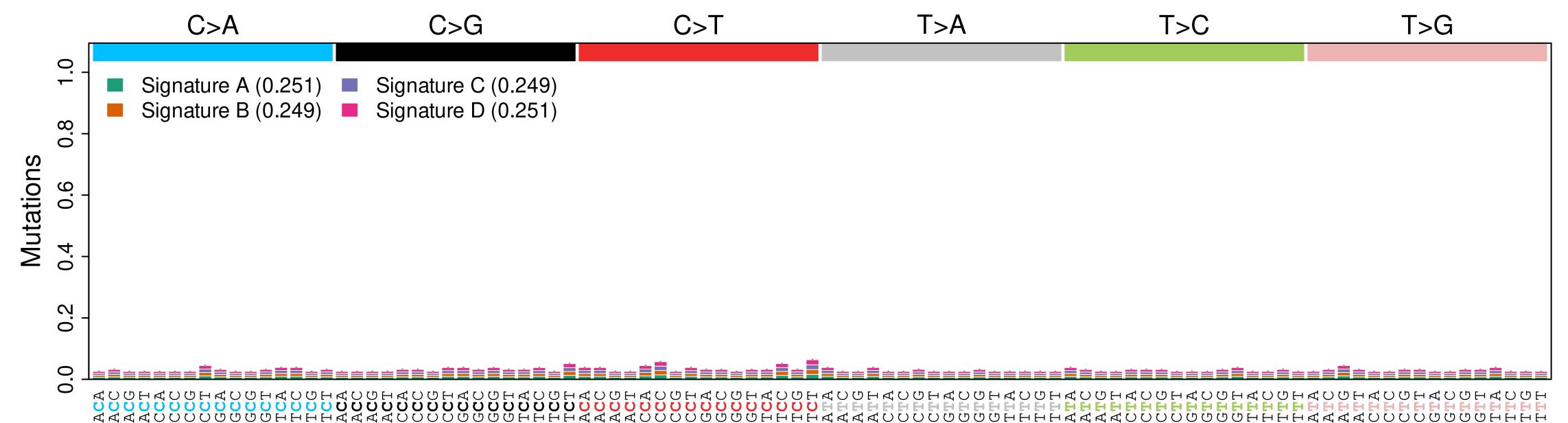
Reconstructed spectrum (cosine similarity = 0.259)



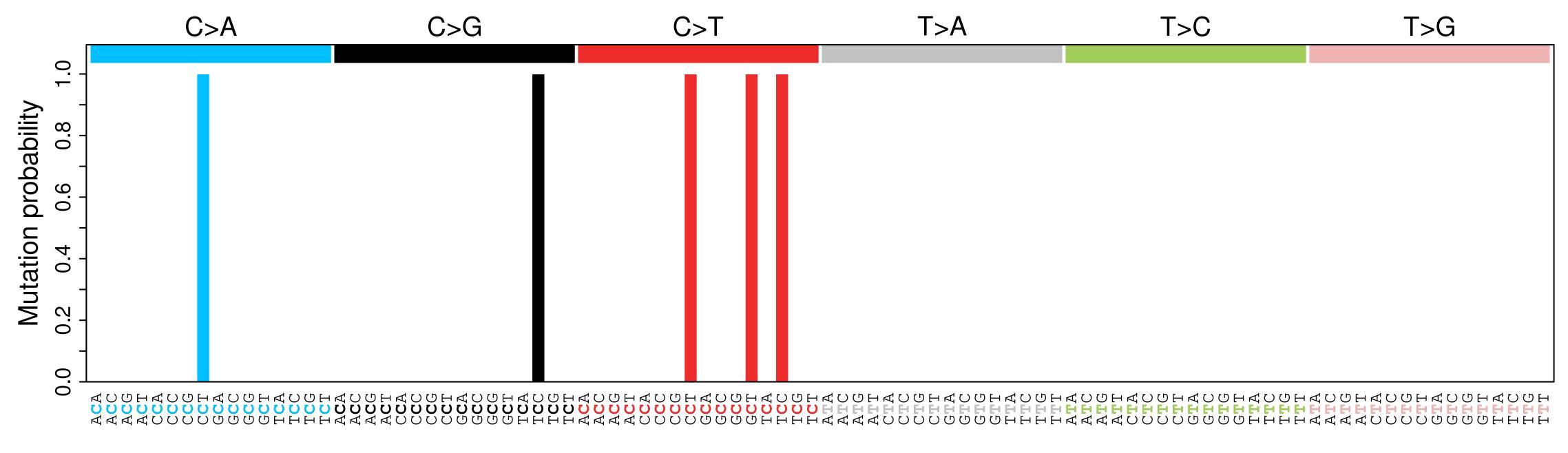




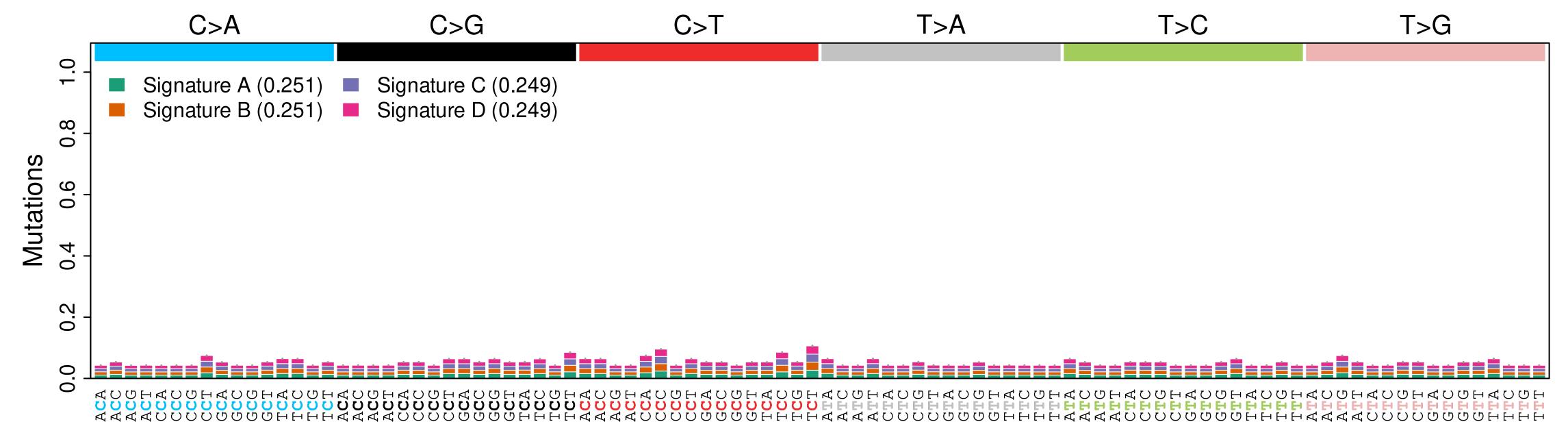




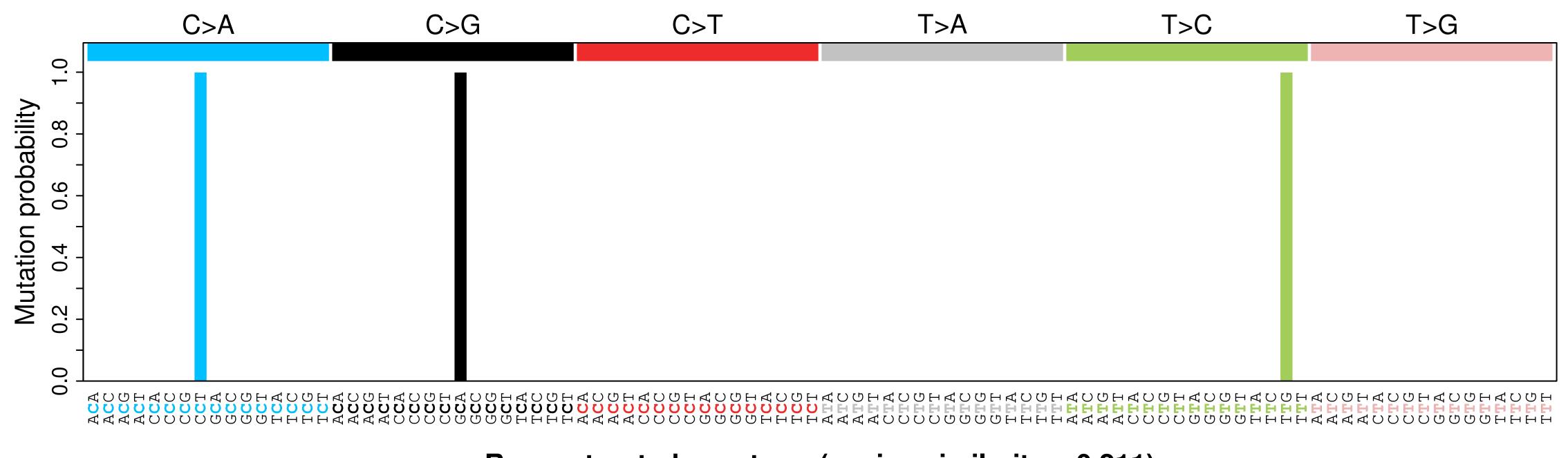




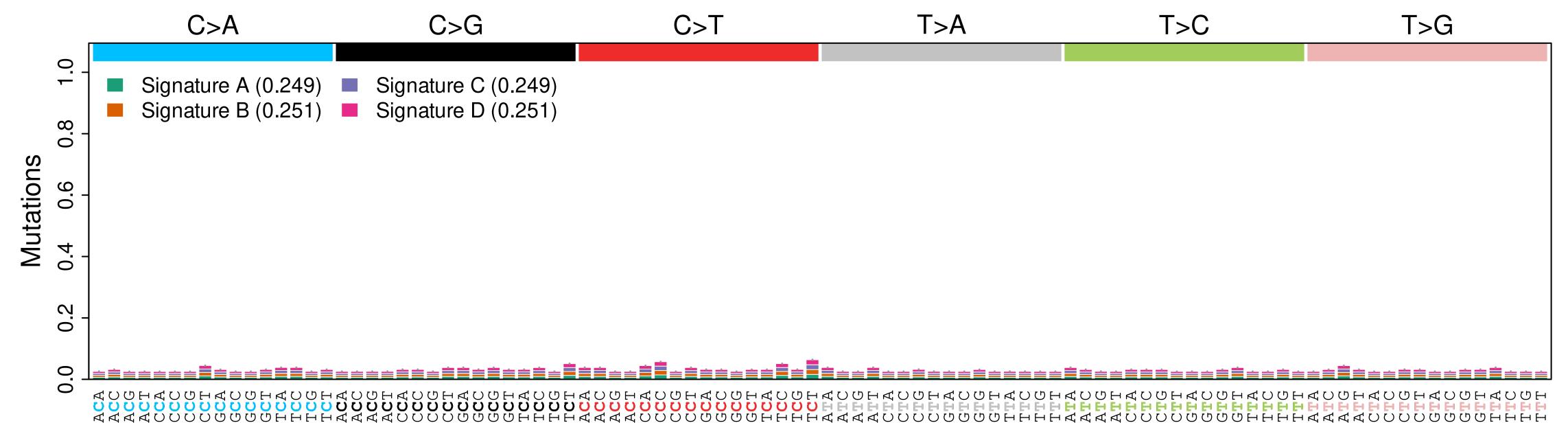
Reconstructed spectrum (cosine similarity = 0.291)



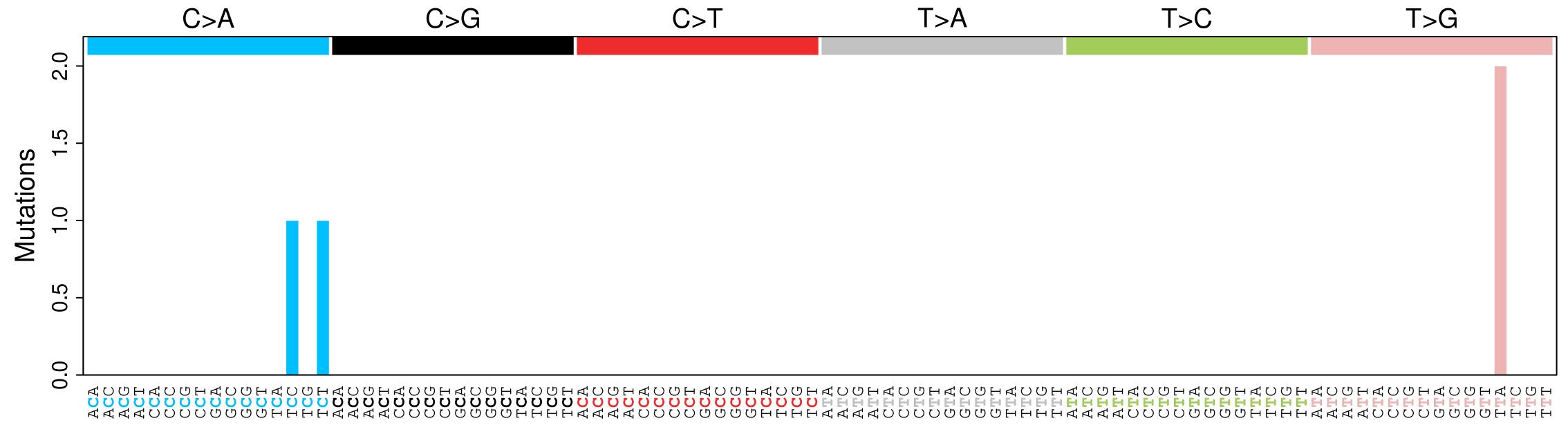


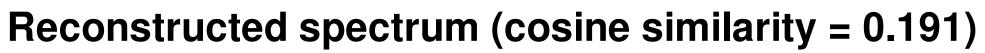


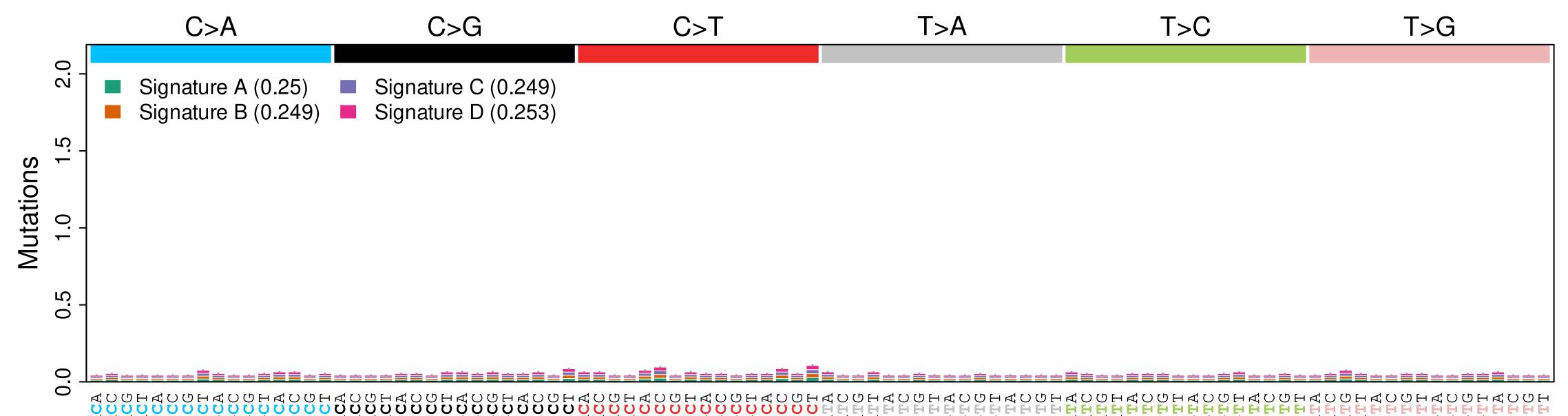
Reconstructed spectrum (cosine similarity = 0.211)



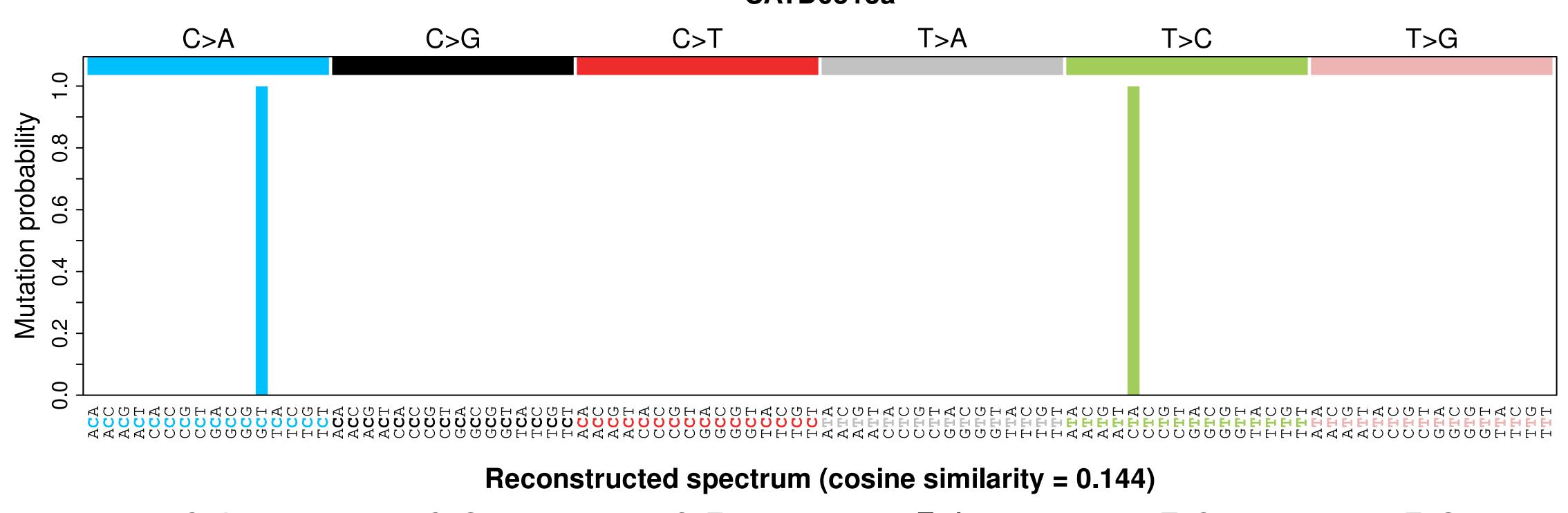
CATD206a (4 mutations) C>A C>G C>T T>A

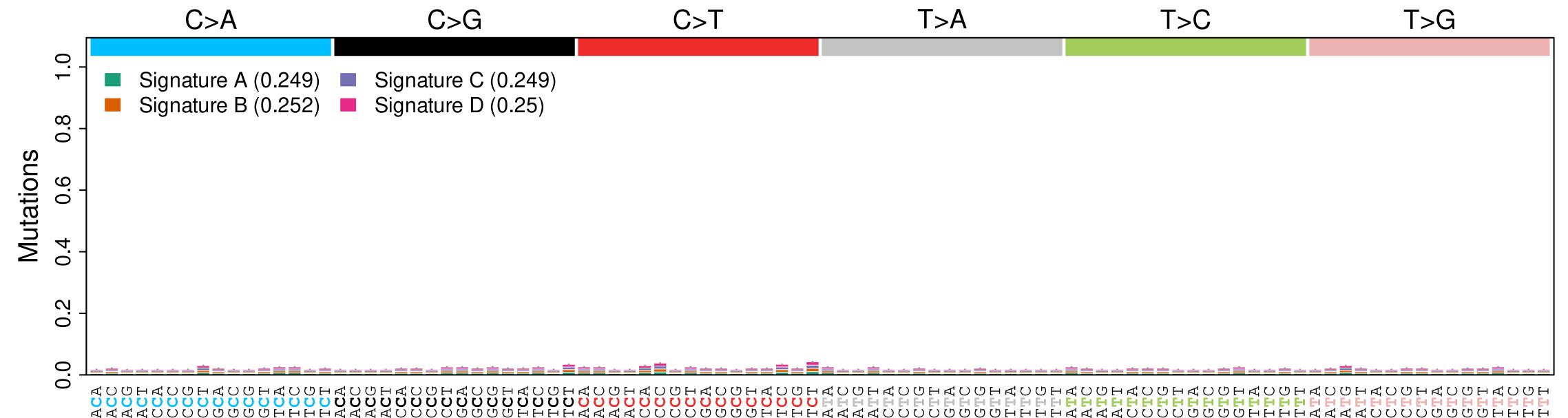






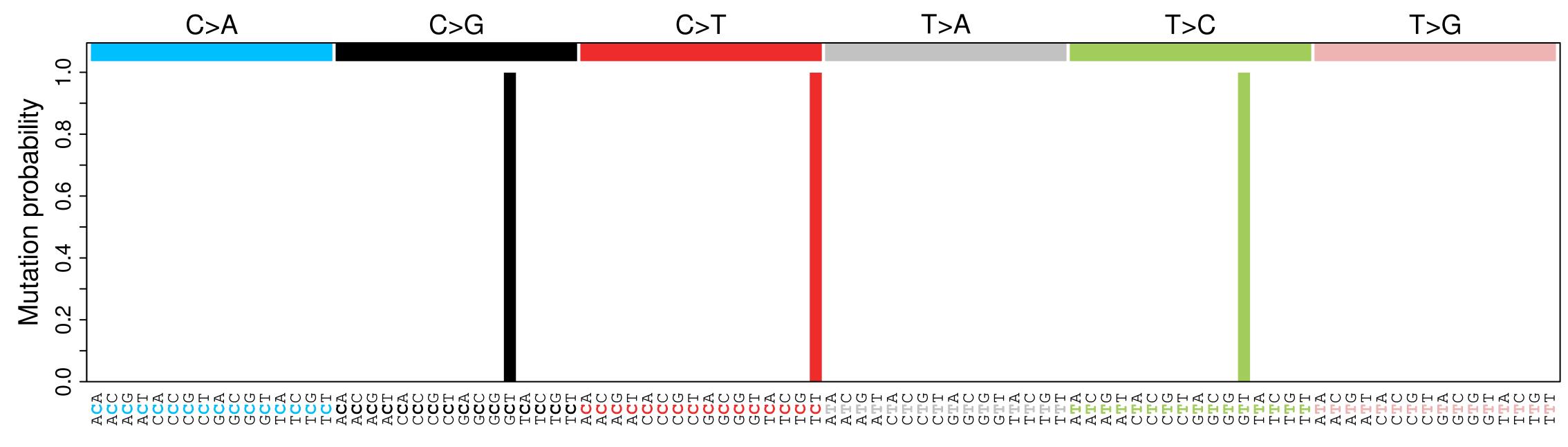




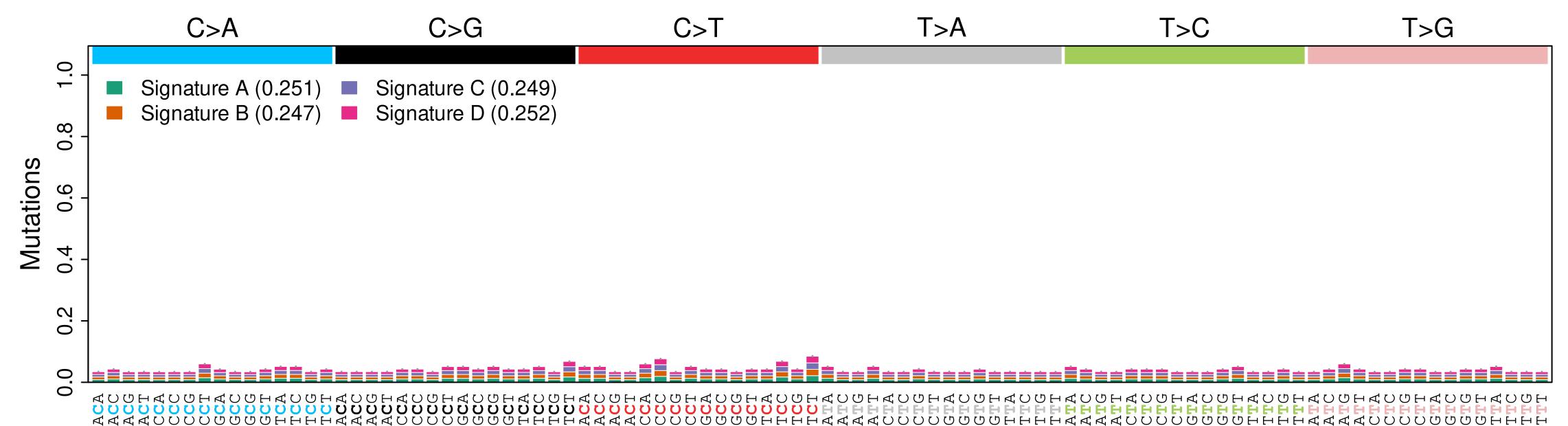


CATD0509a (3 mutations) T>G C>A C>G C>T T>A T>C 2.0 5 Mutations 0.5 0.0 **Reconstructed spectrum (cosine similarity = 0.172)** T>C T>G C>A C>G T>A C>T 2.0 Signature A (0.252) Signature C (0.252) Signature B (0.249) Signature D (0.248) 5 Mutations 5 0

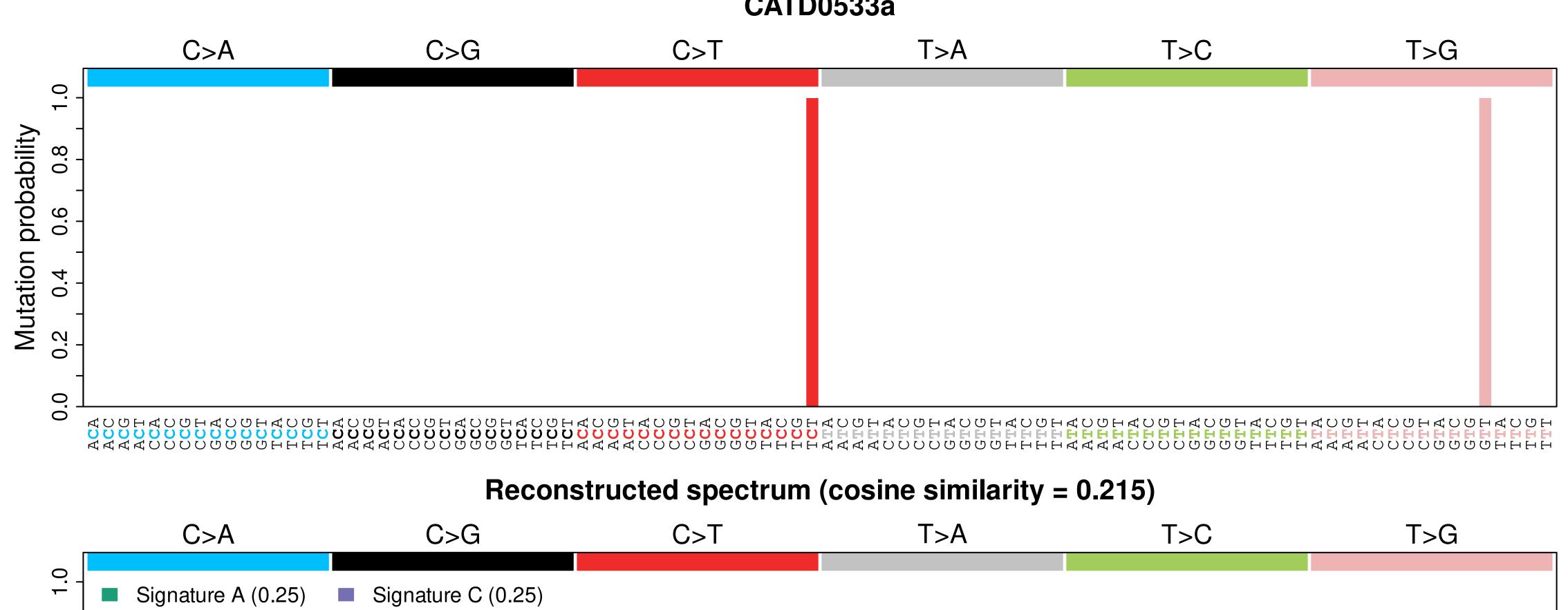


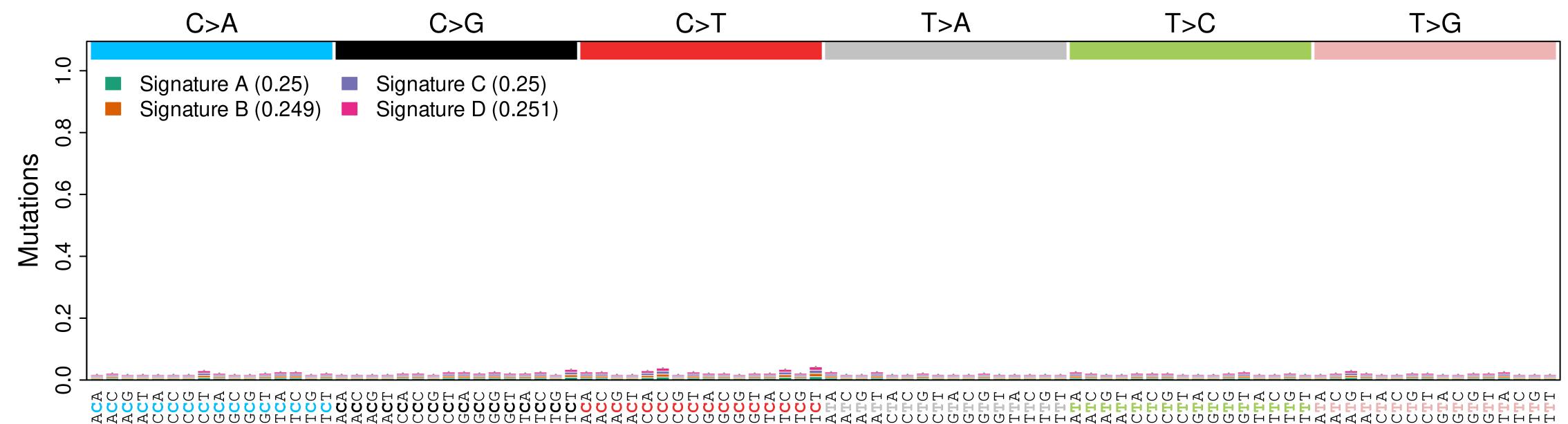


Reconstructed spectrum (cosine similarity = 0.246)

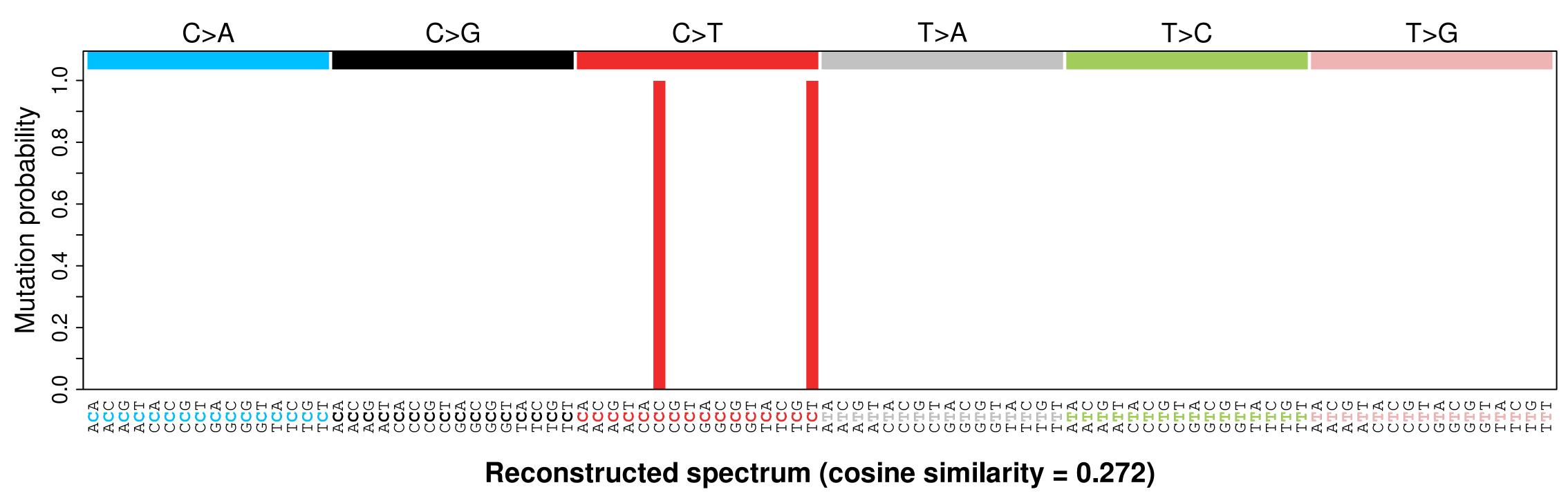


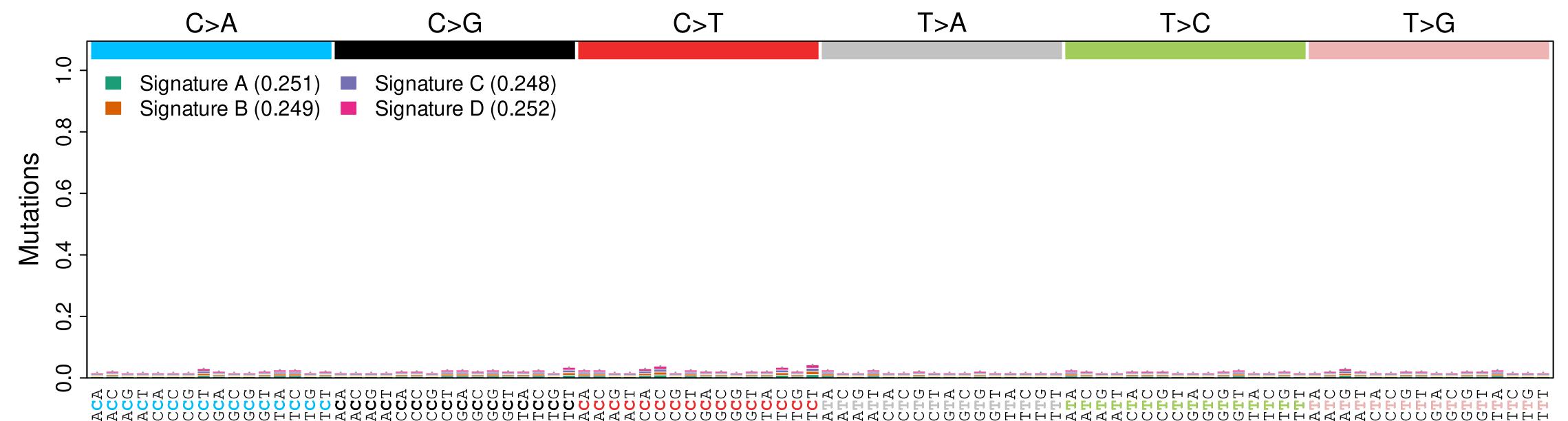


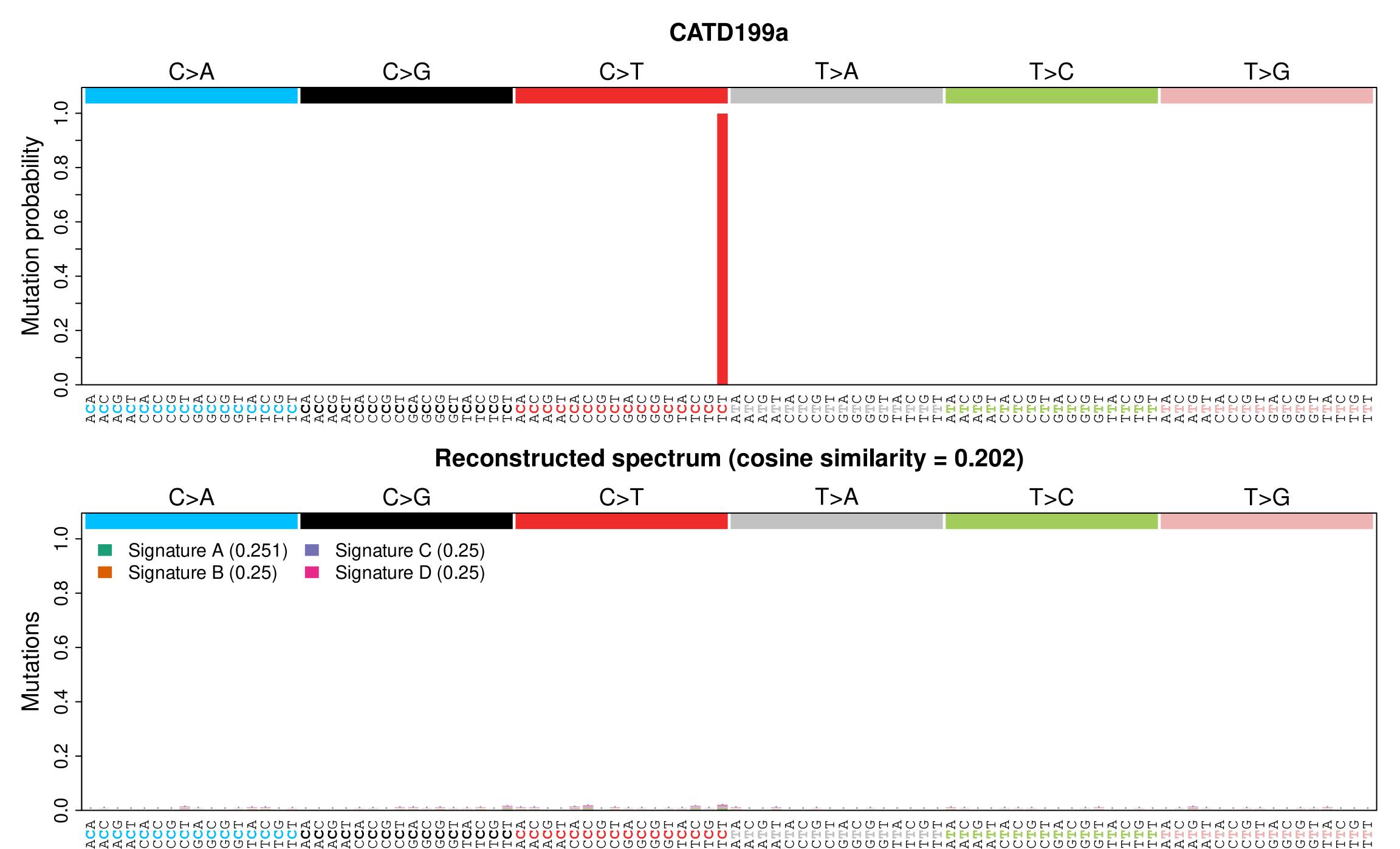


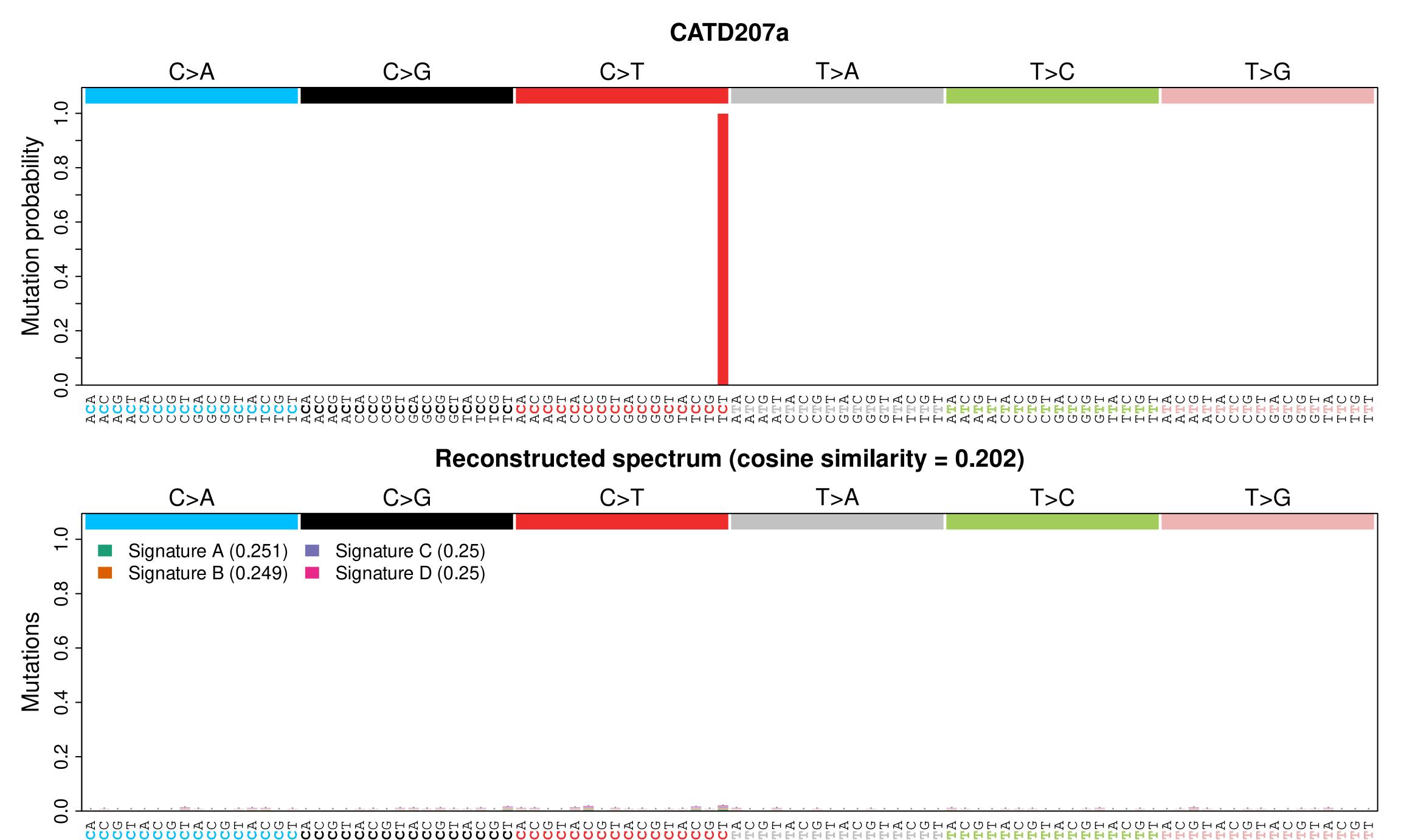




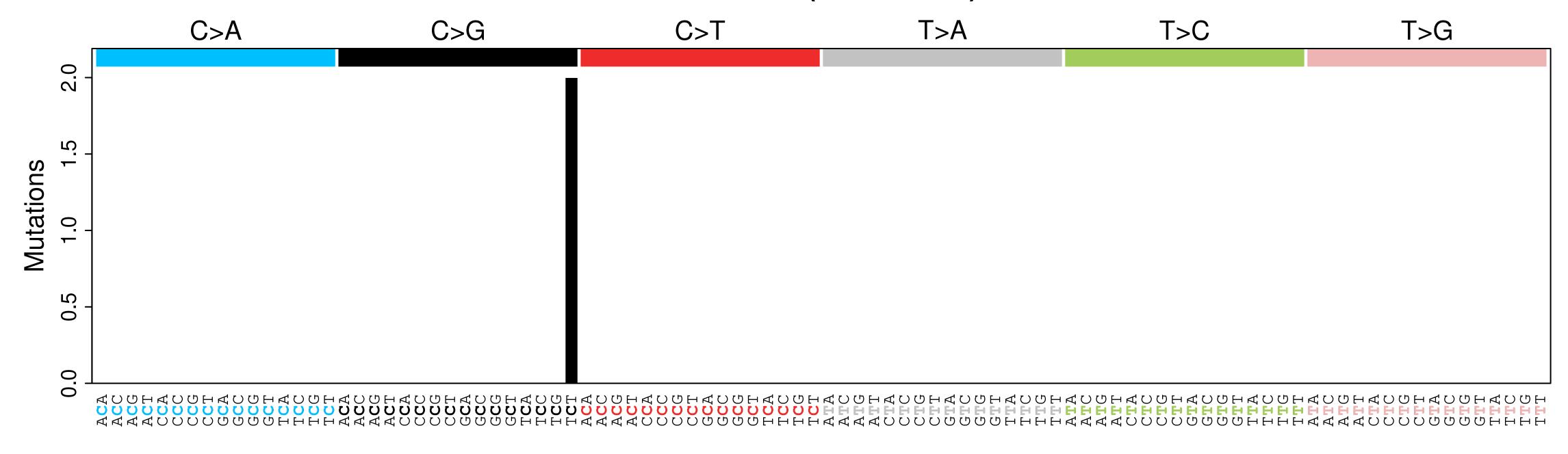








CATD197a (2 mutations)



Reconstructed spectrum (cosine similarity = 0.162)

