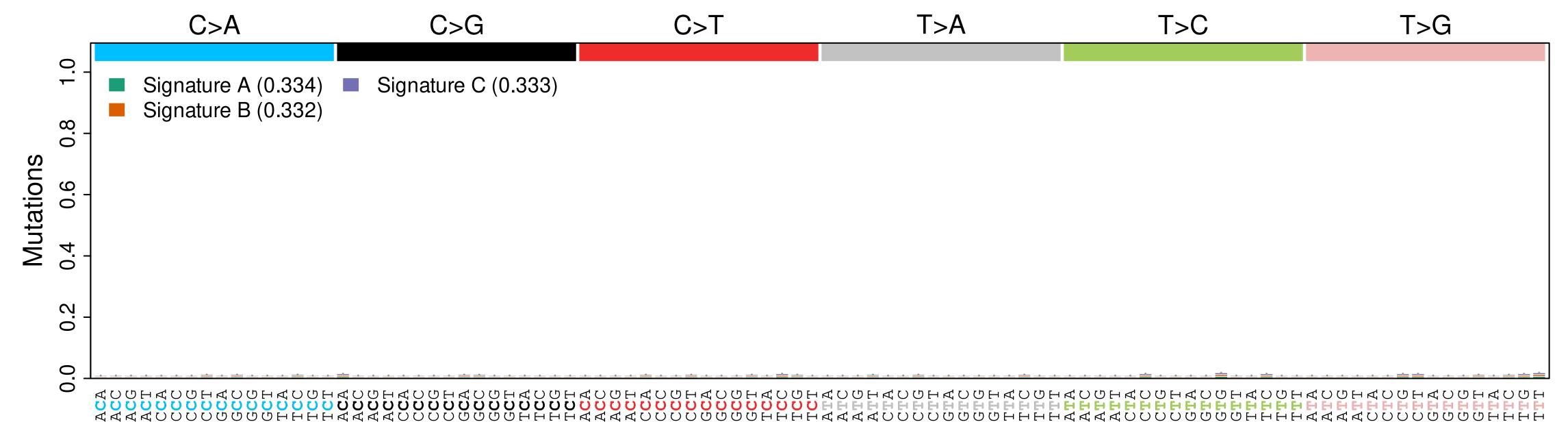
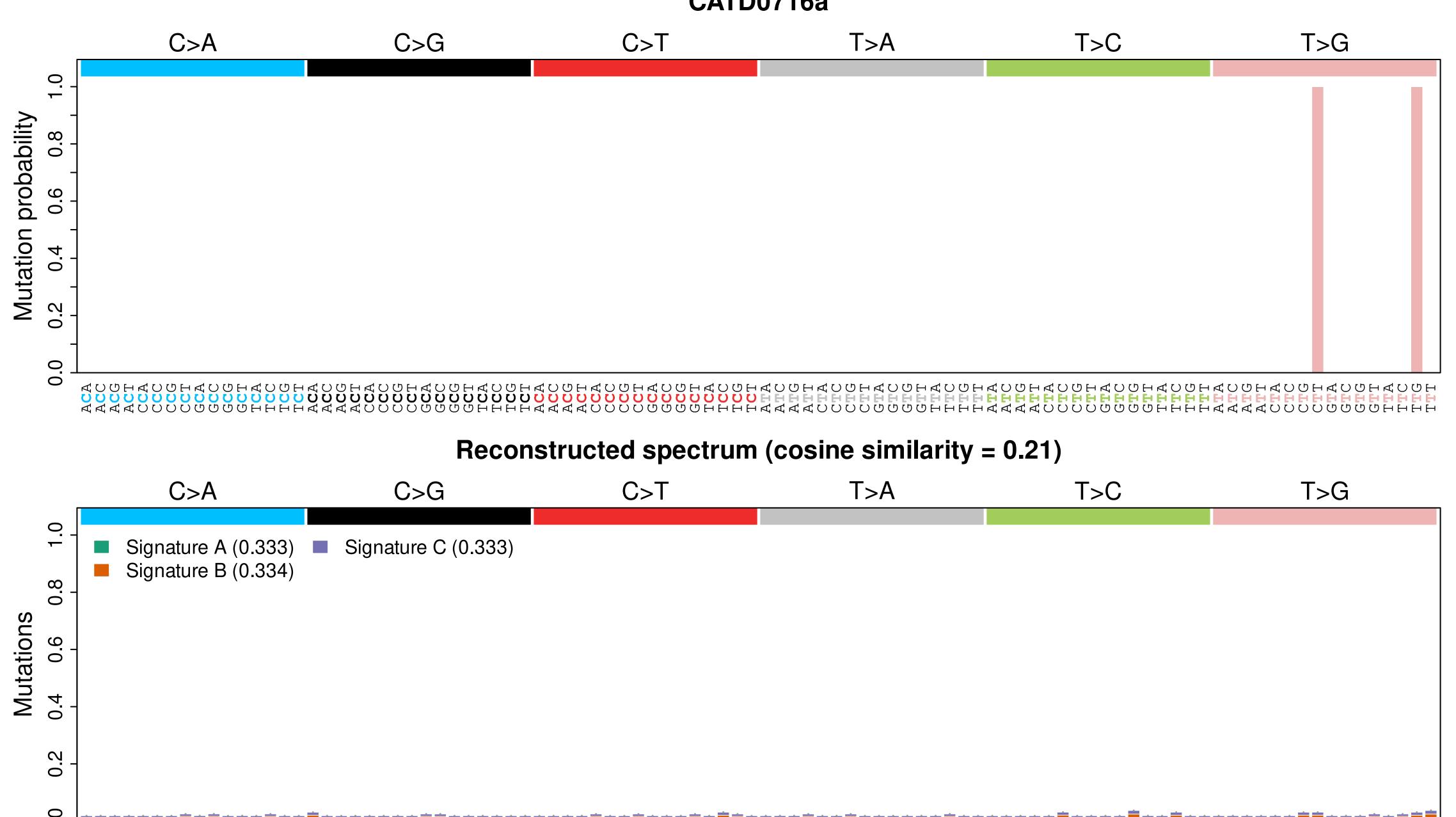


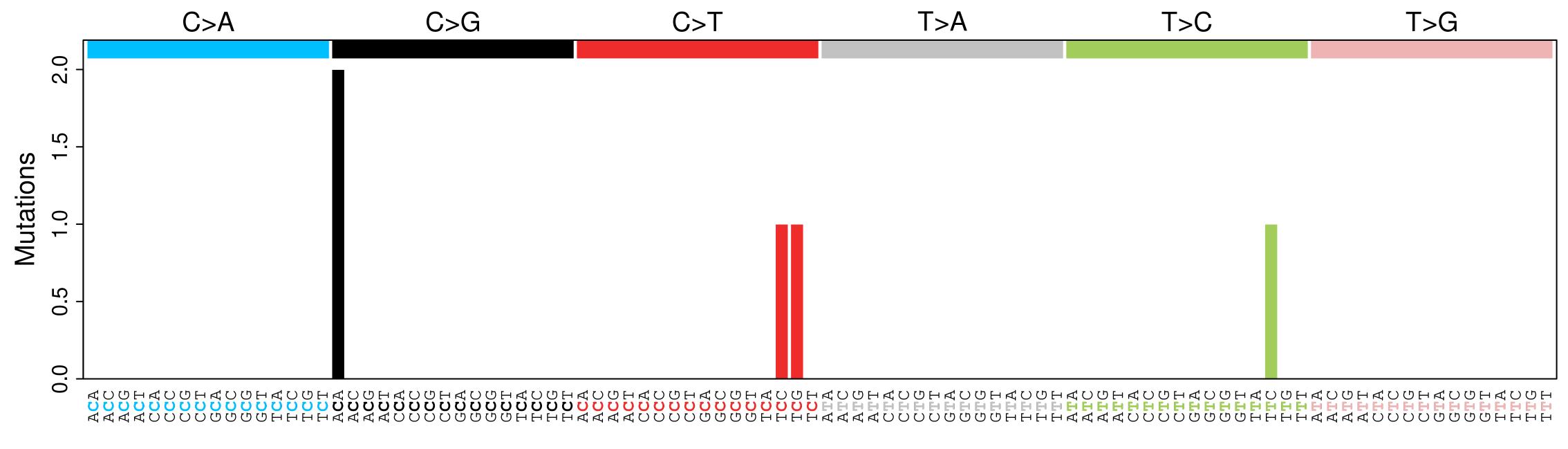
Reconstructed spectrum (cosine similarity = 0.12)



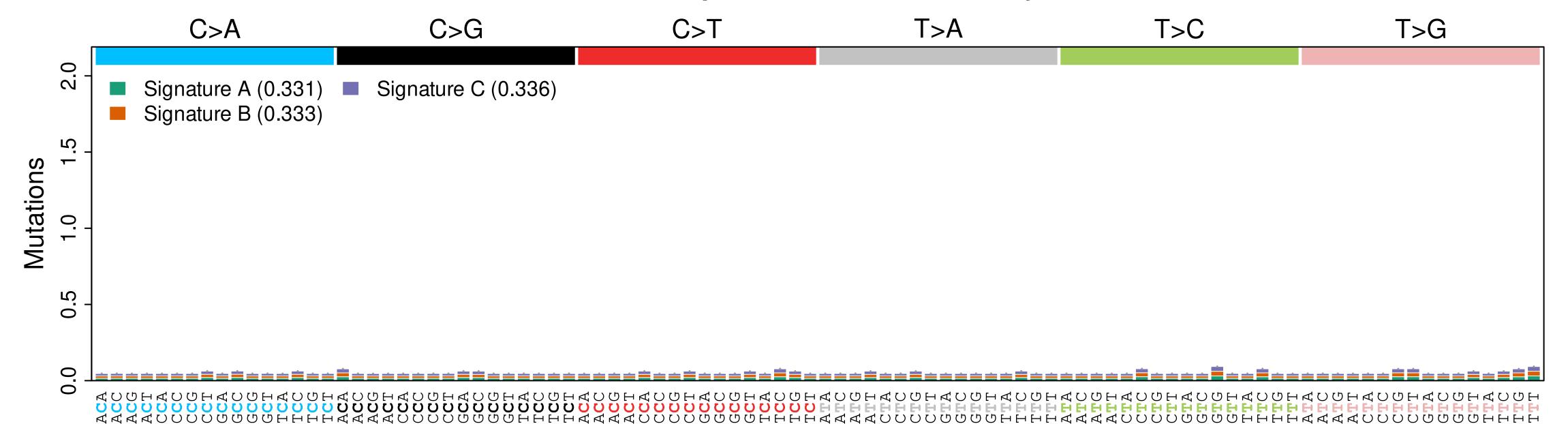




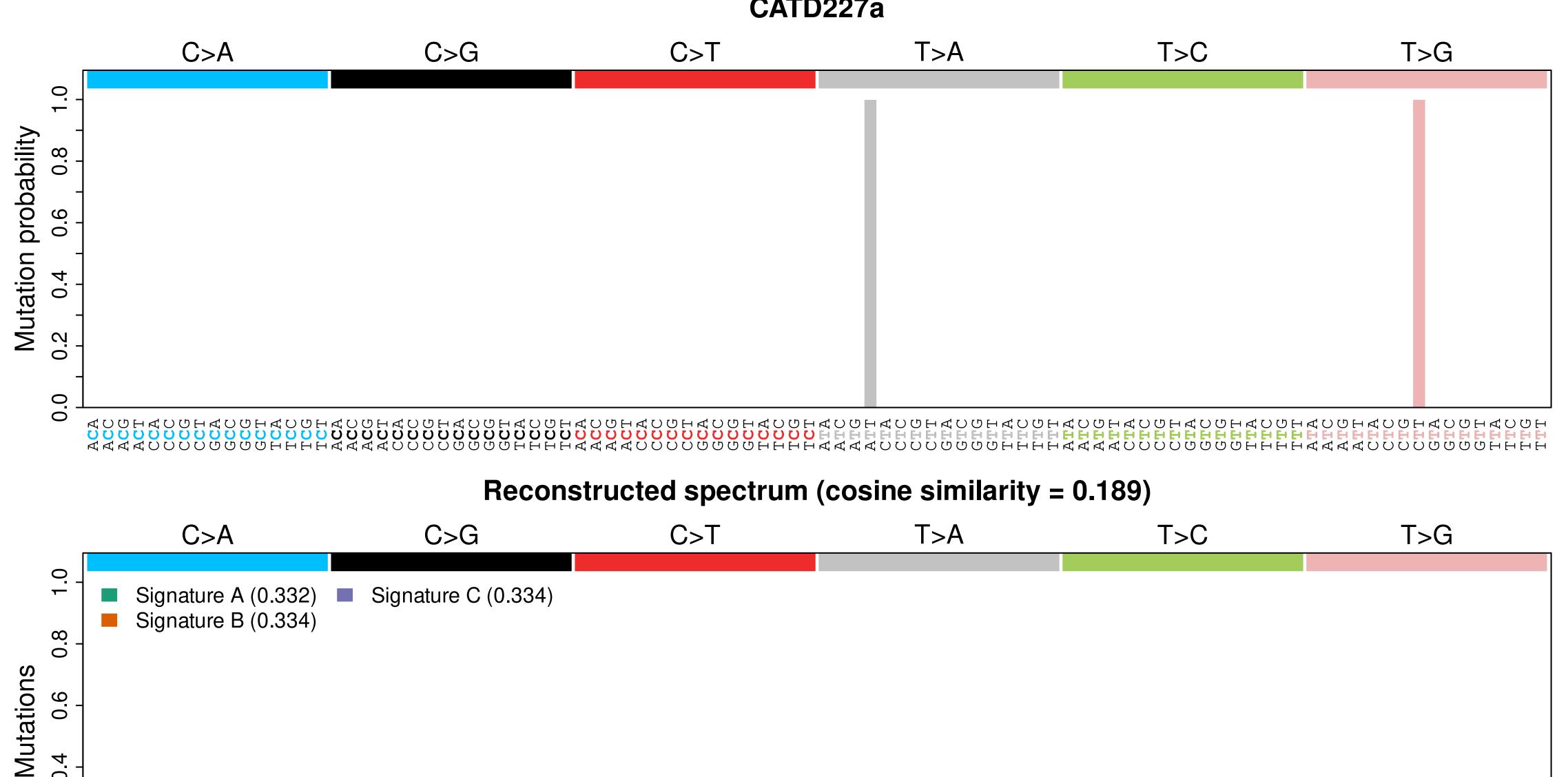
CATD217a (5 mutations)



Reconstructed spectrum (cosine similarity = 0.269)

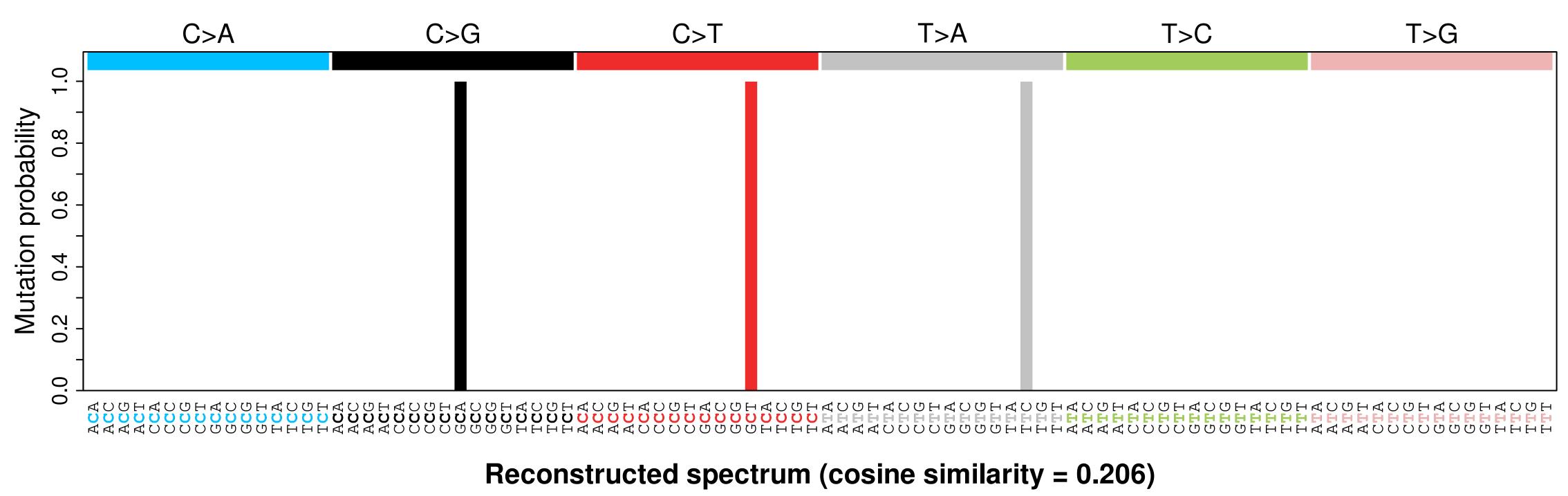


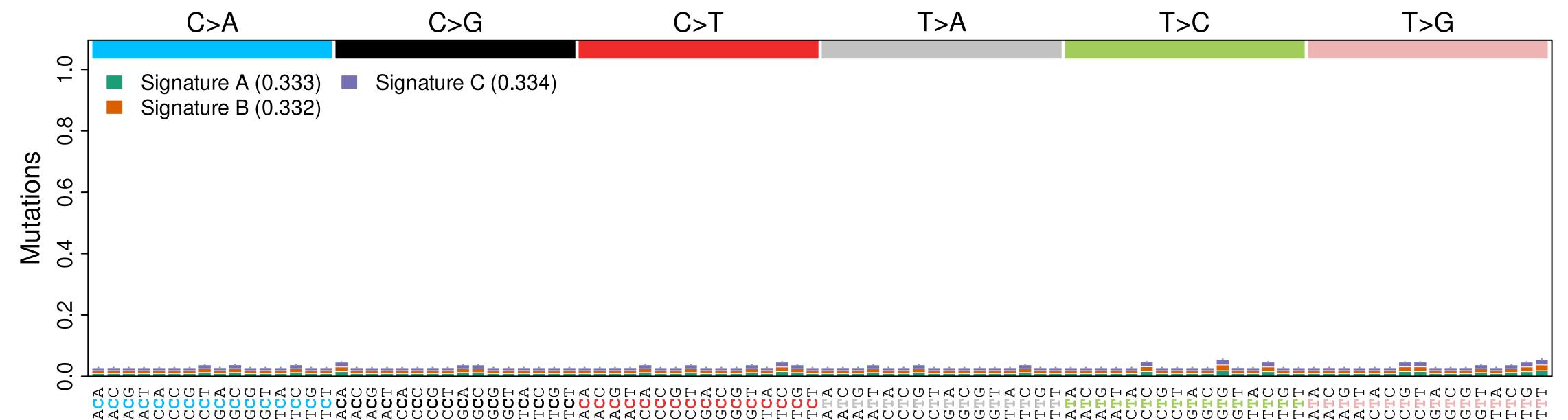




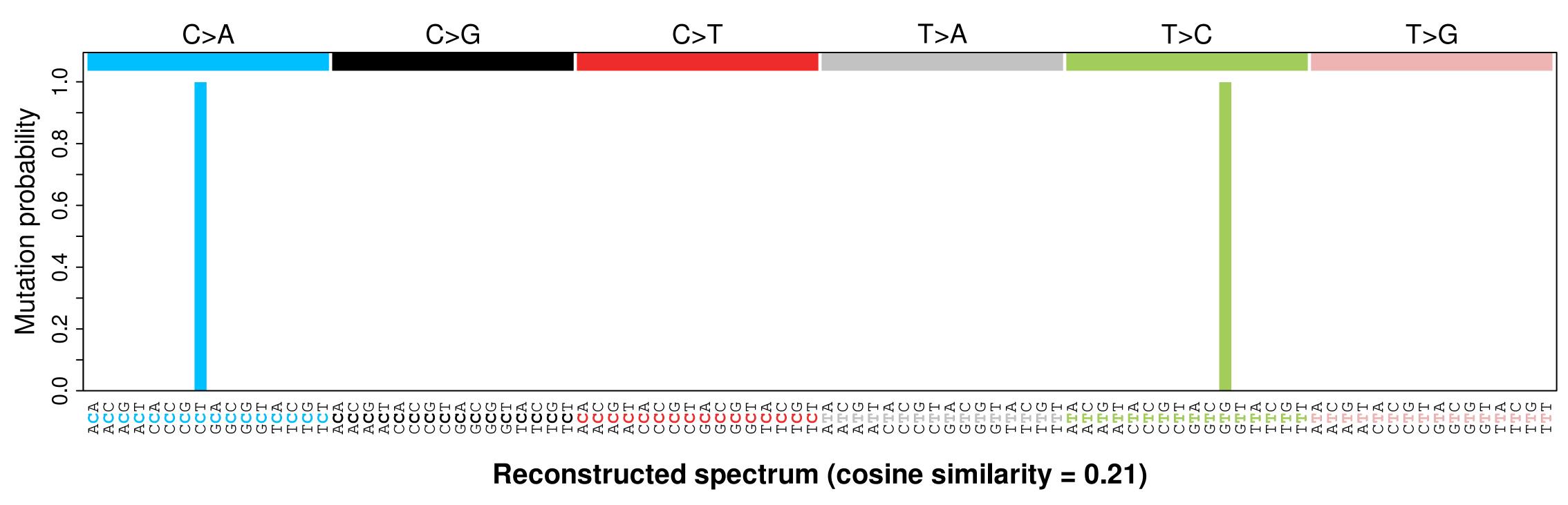
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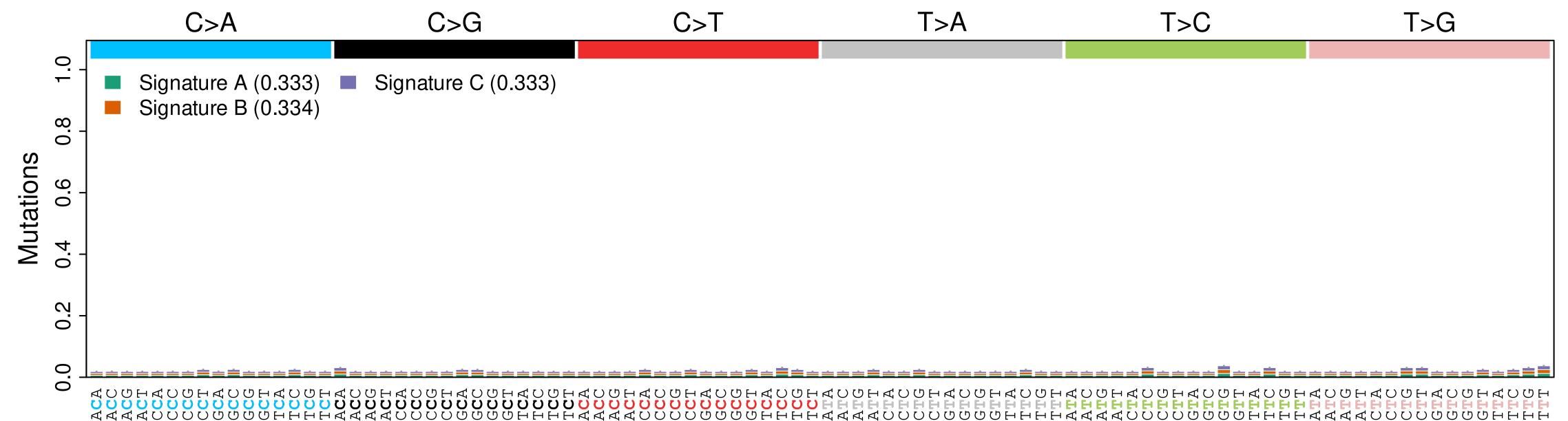




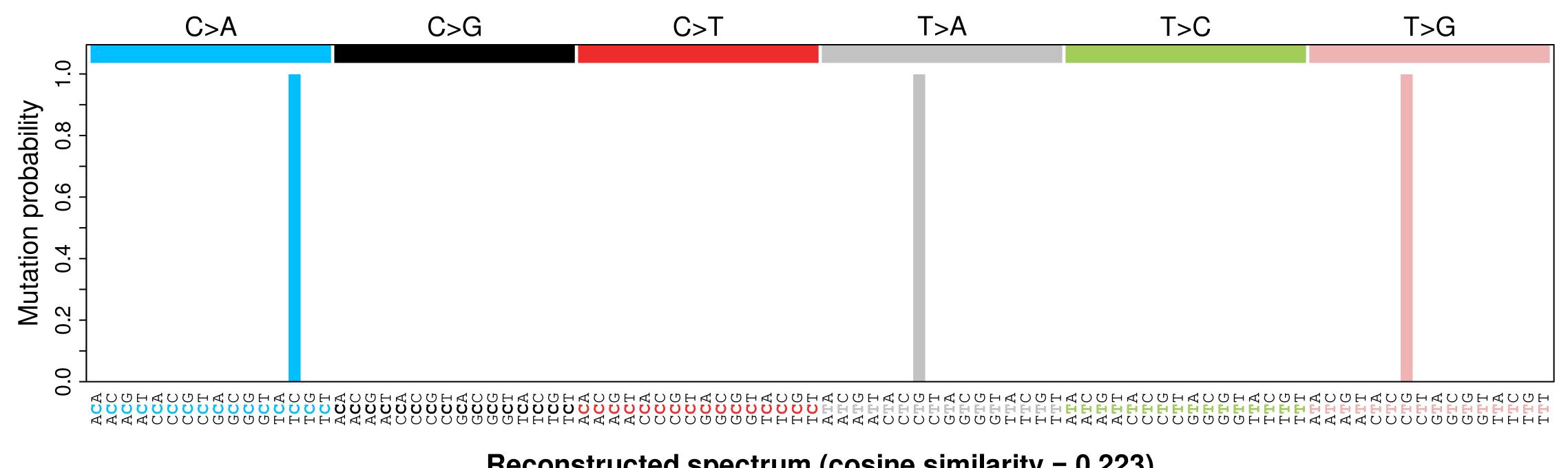


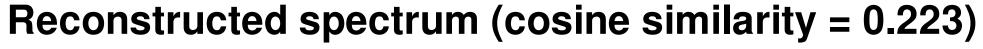


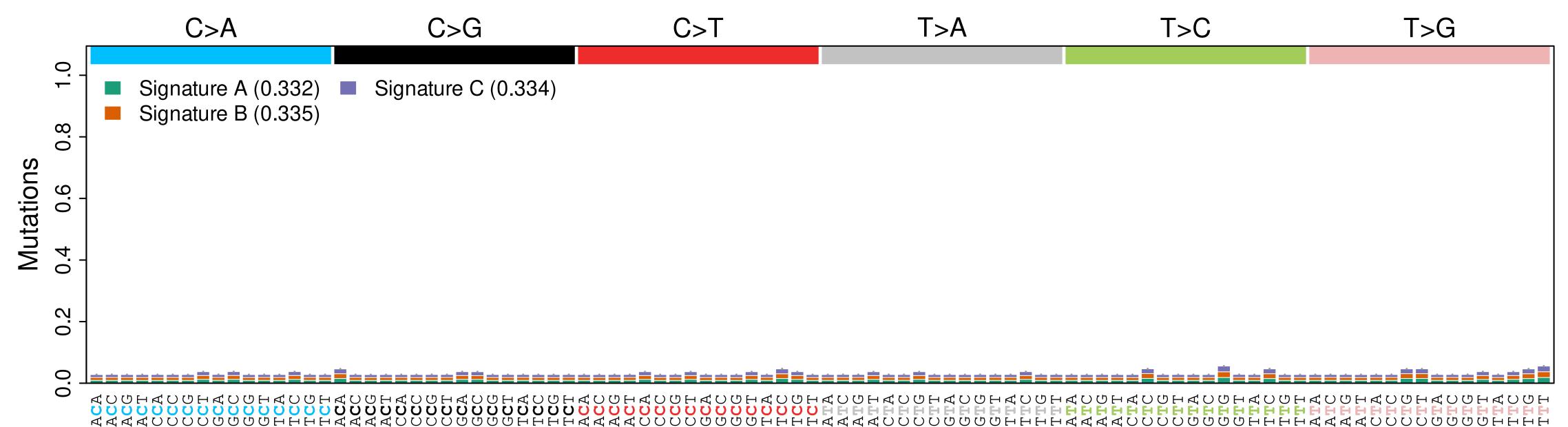




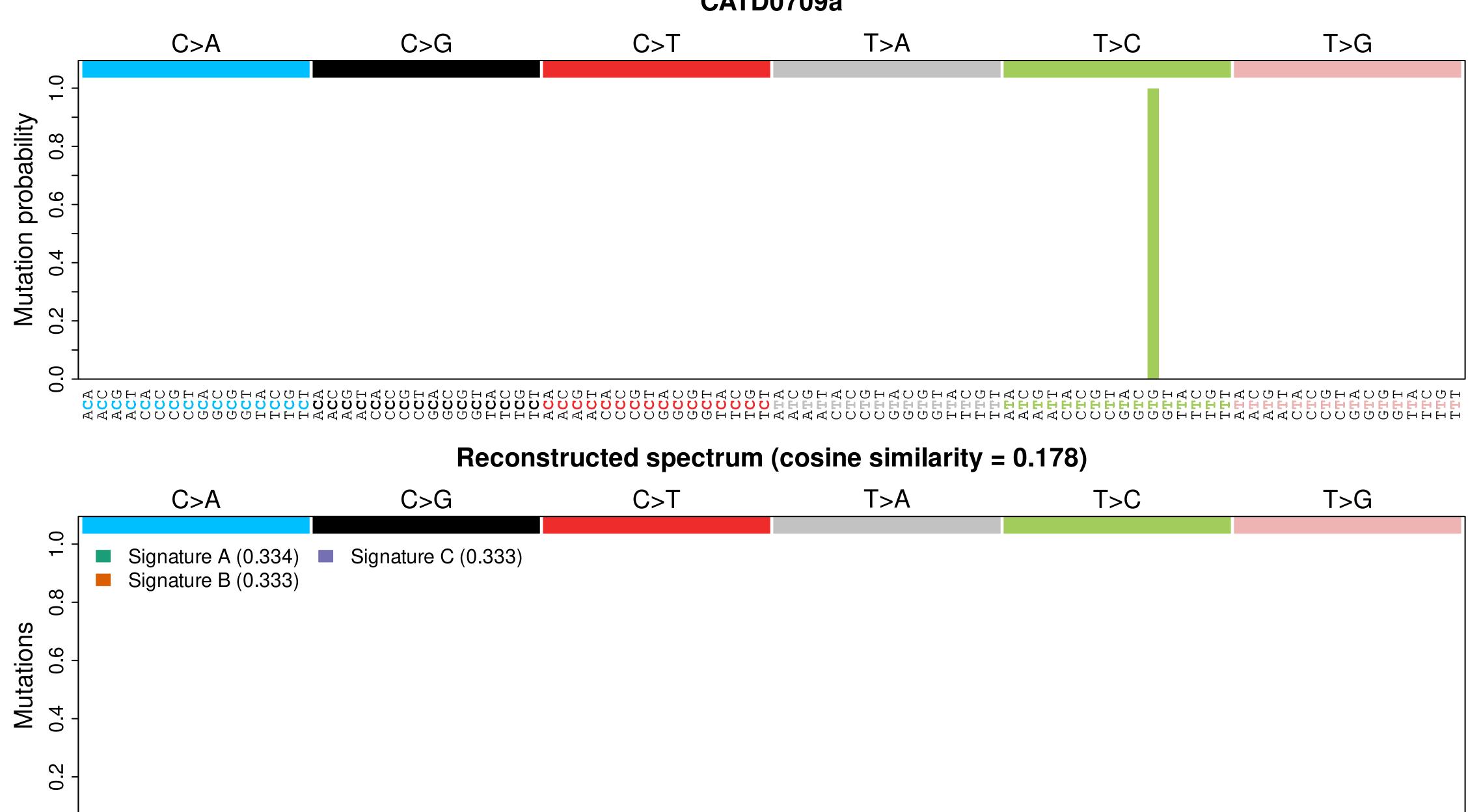


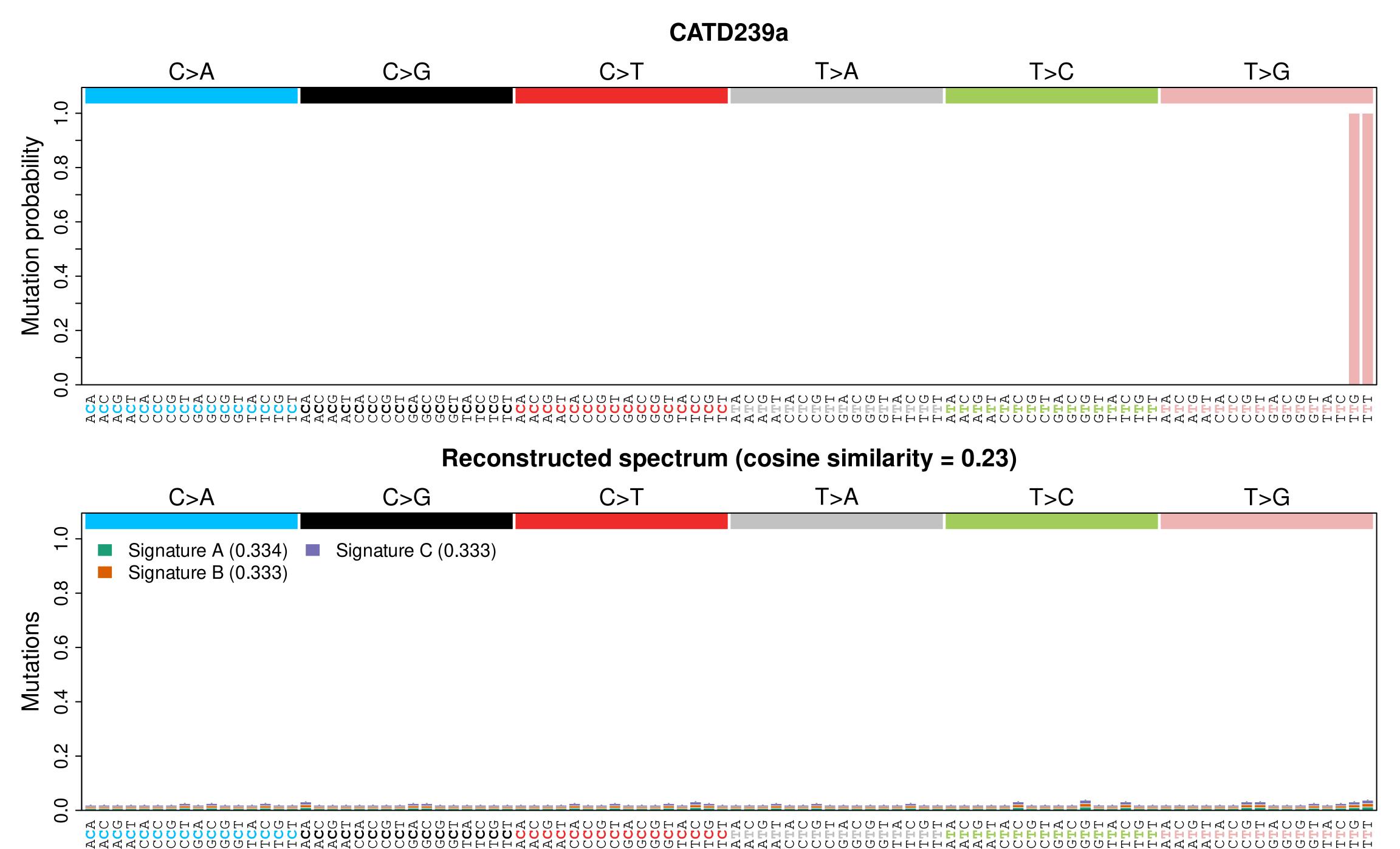


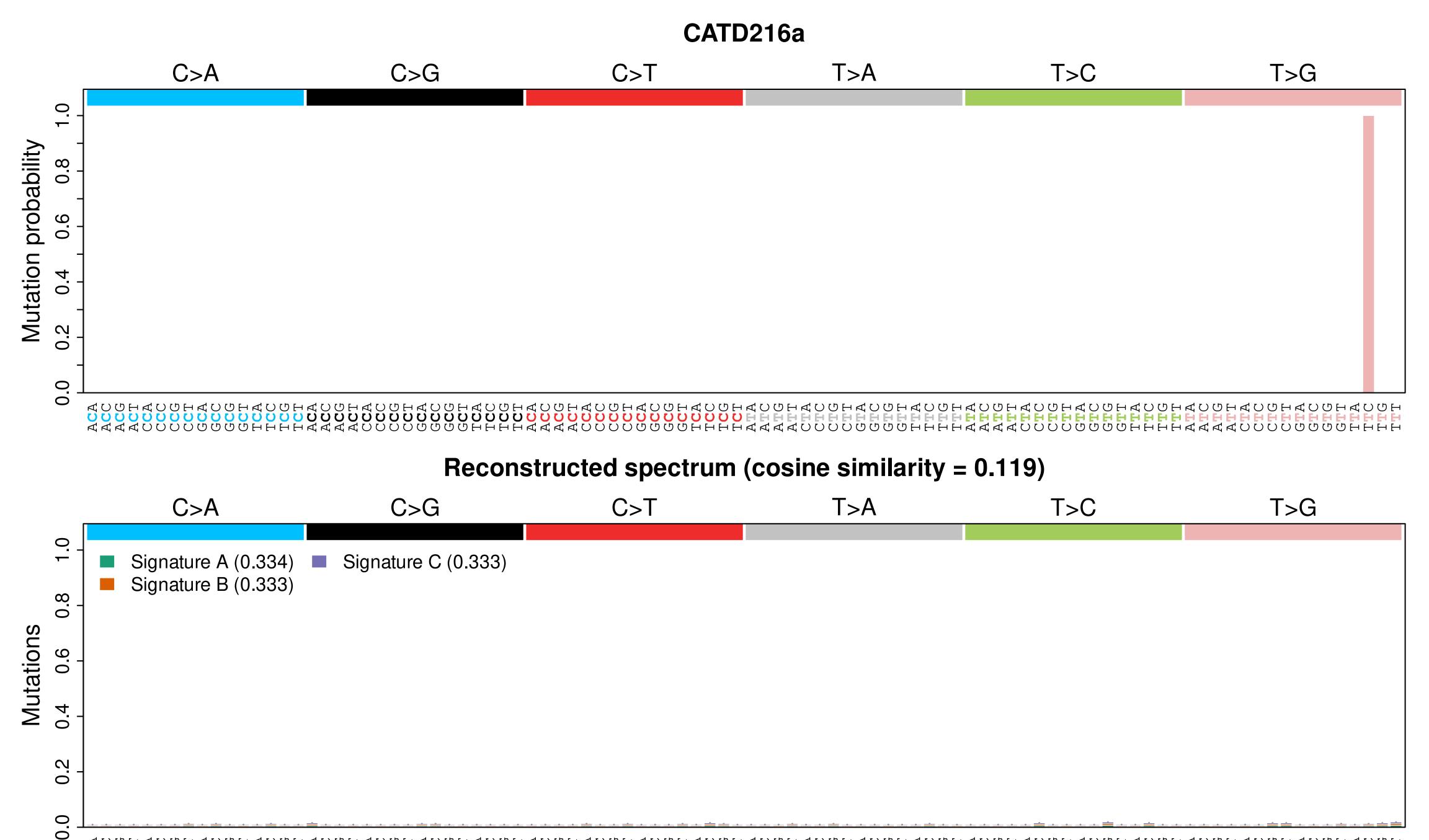




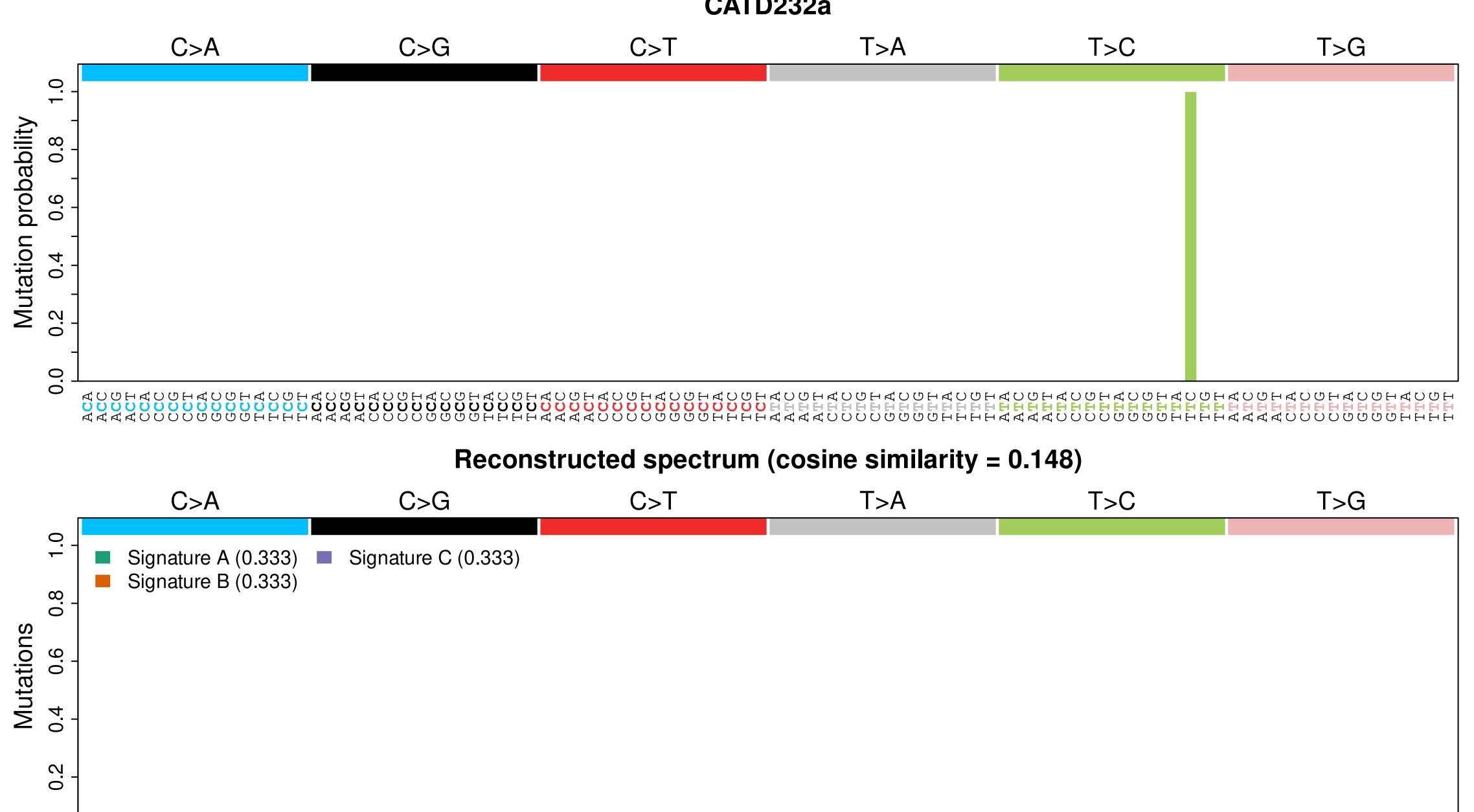




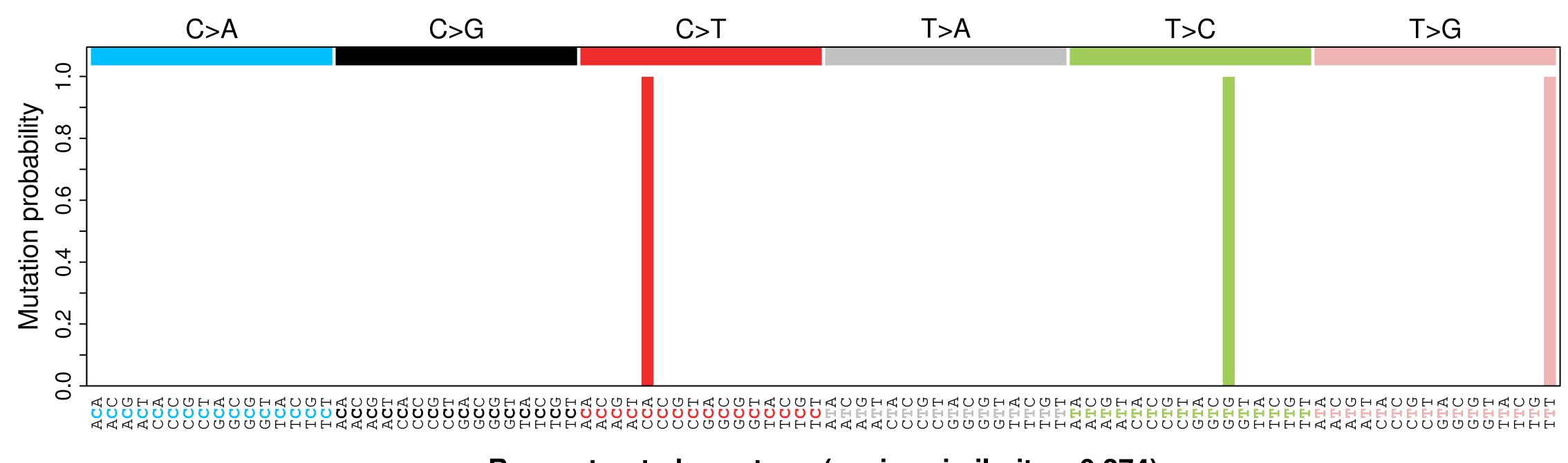




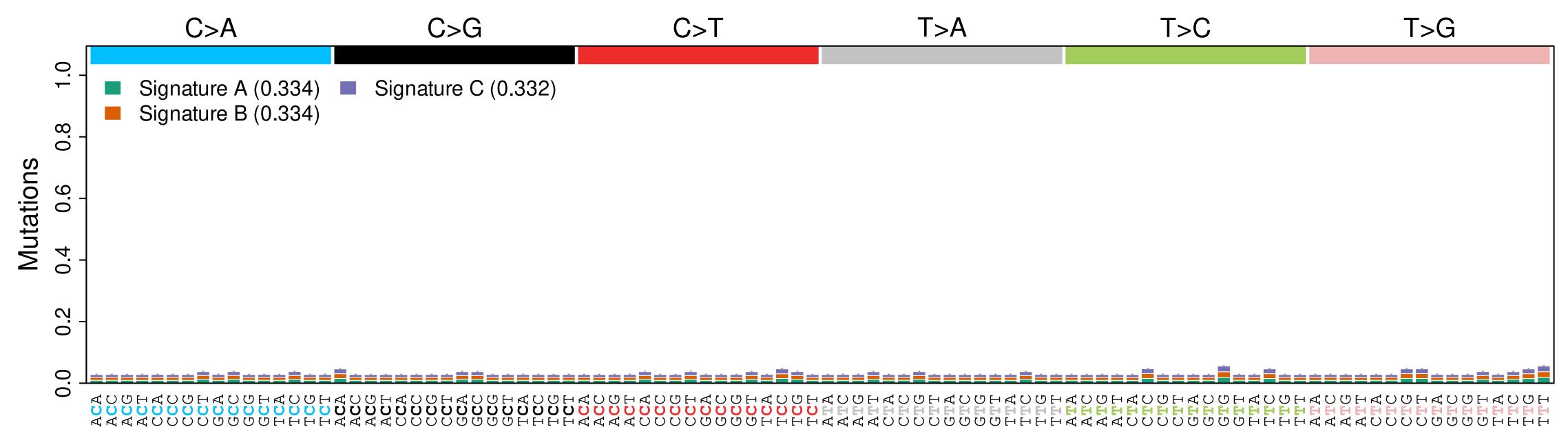




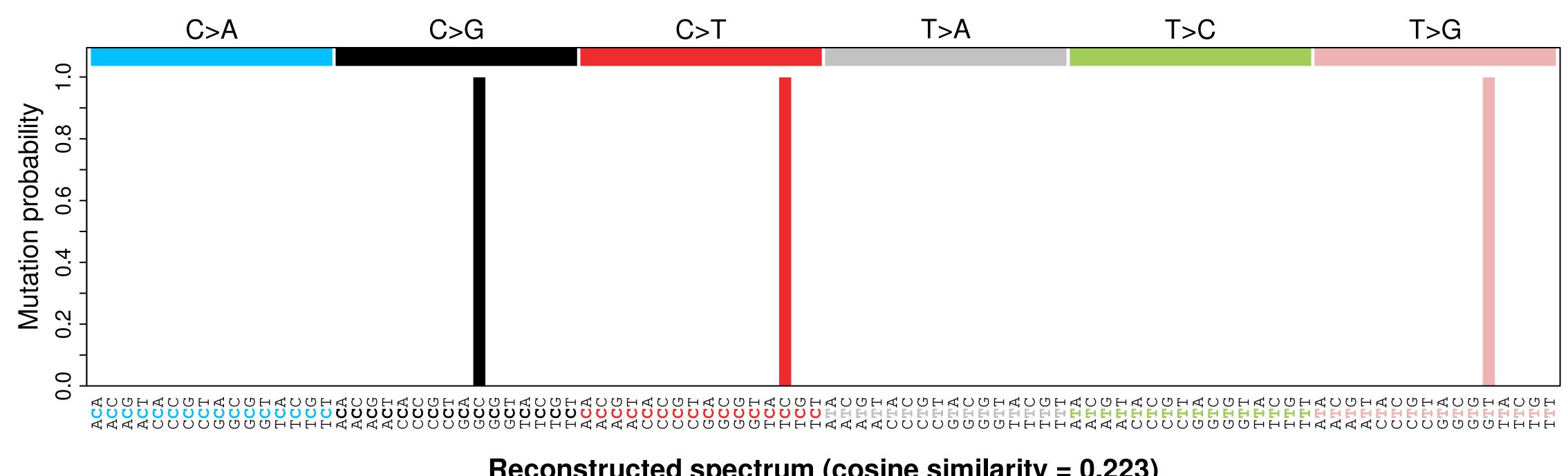


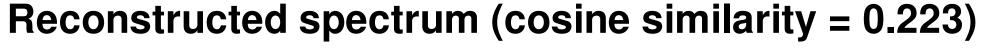


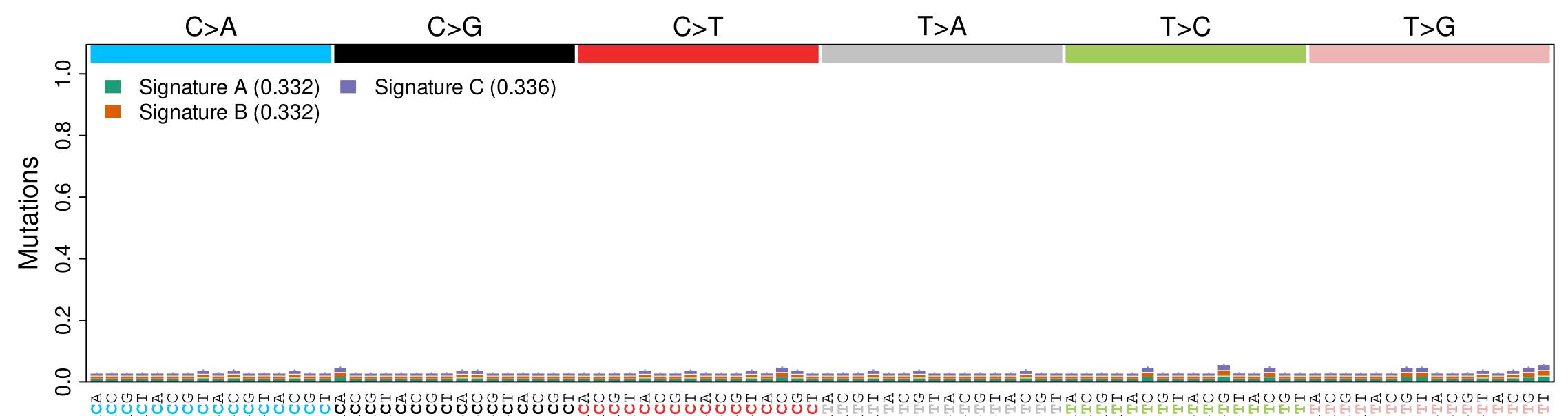




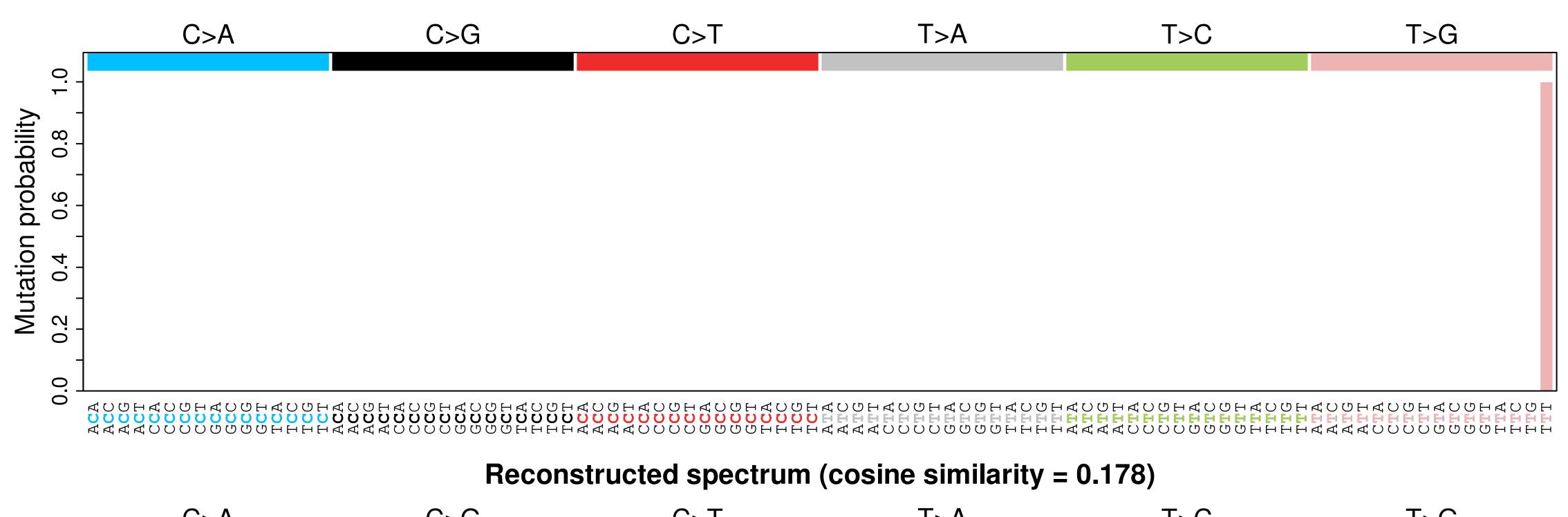


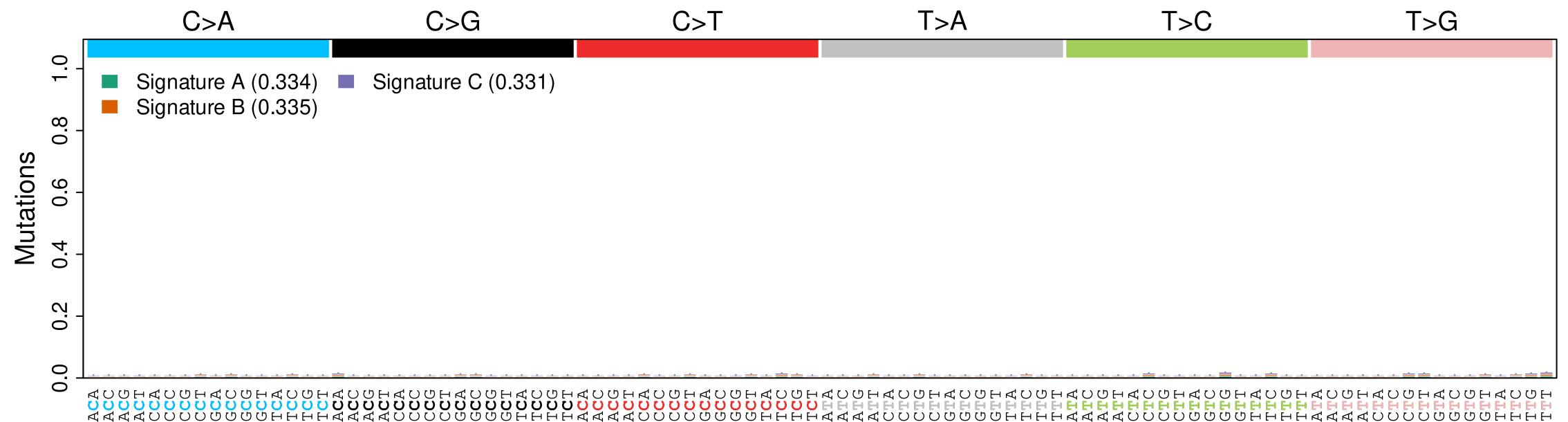




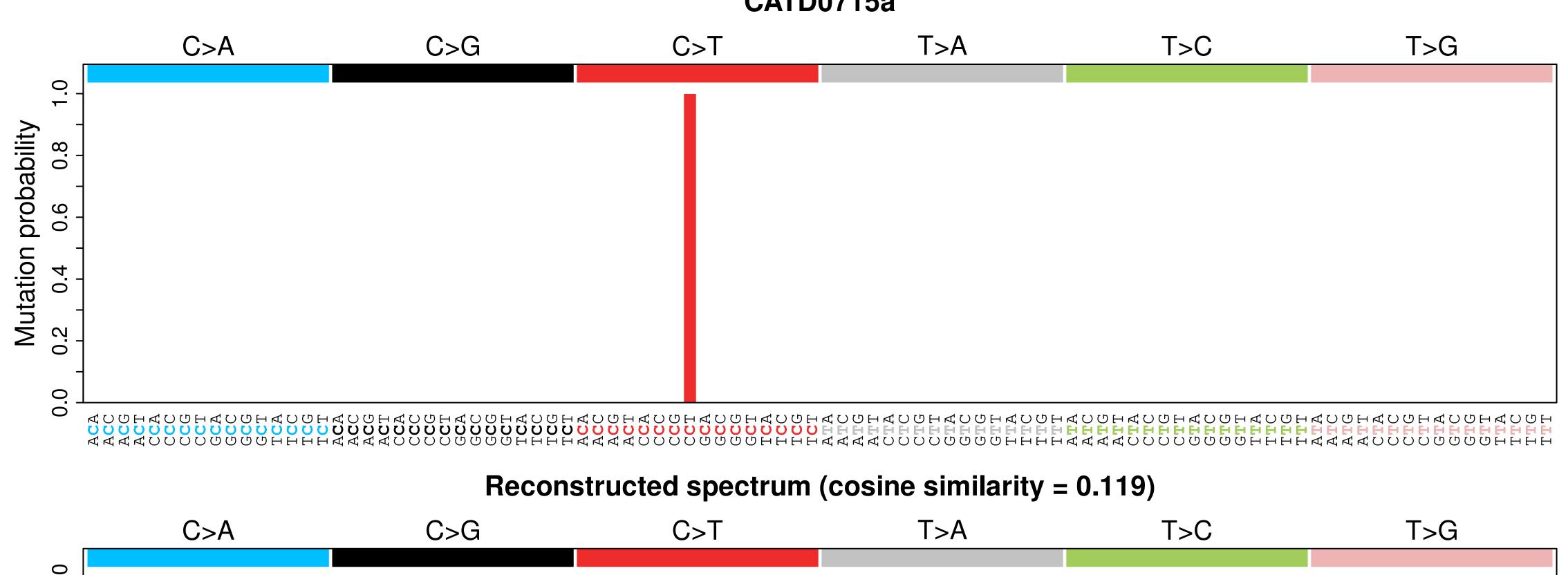


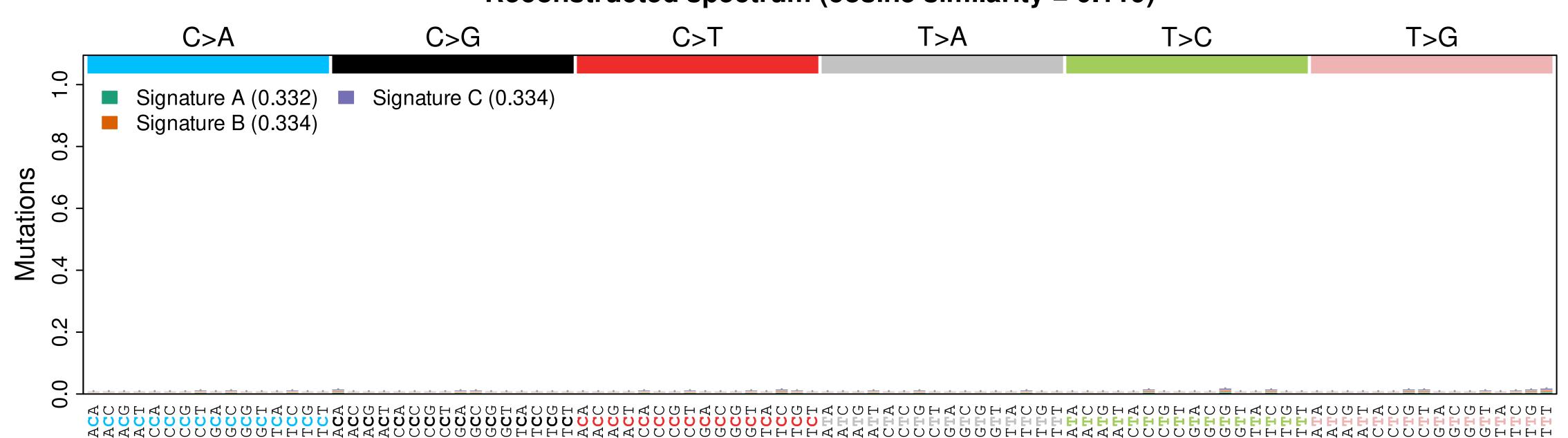












CATD224a (2 mutations)

