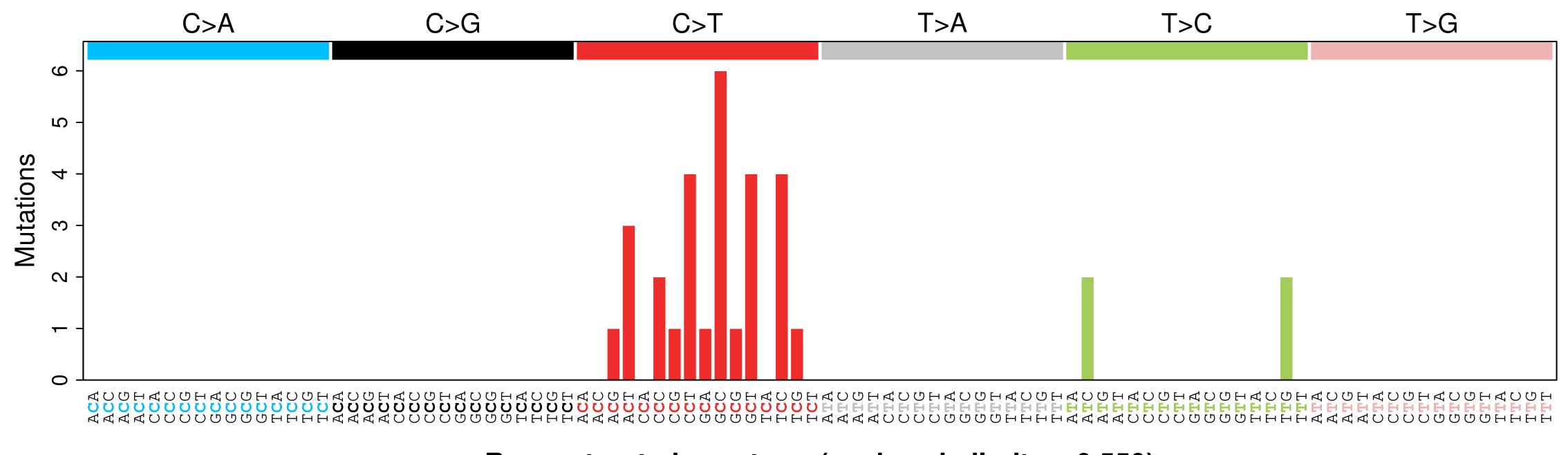
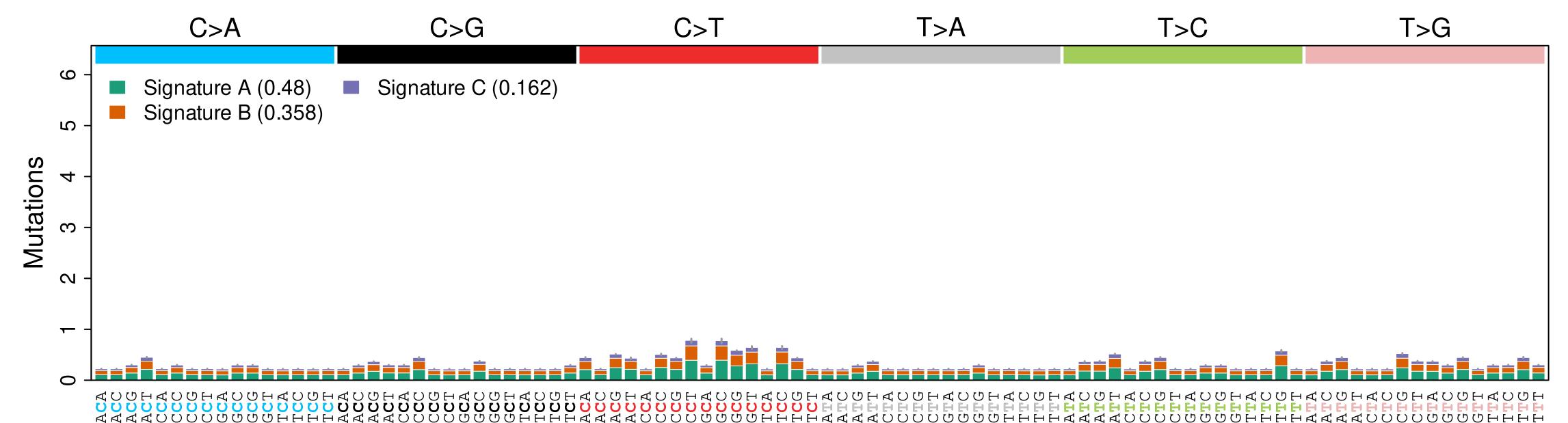
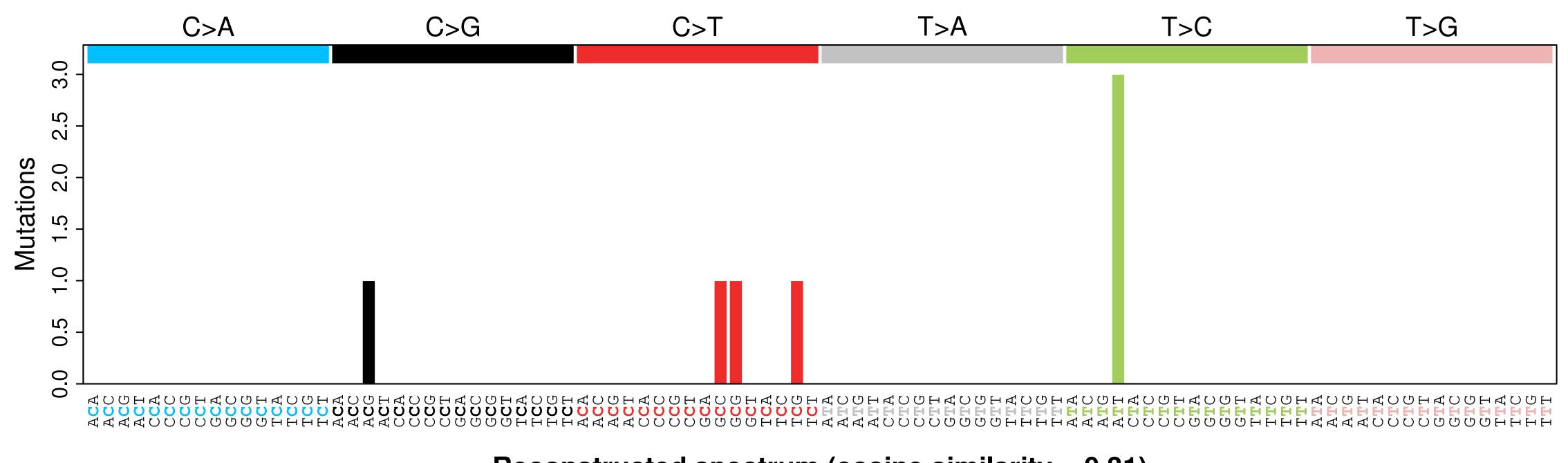
CATD0393a (32 mutations)



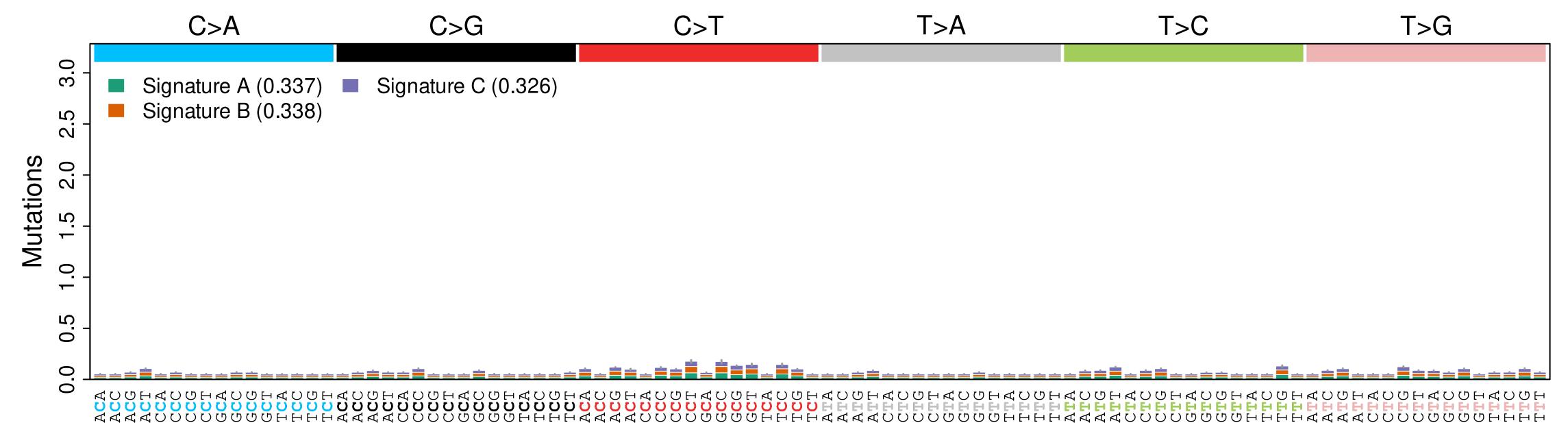
Reconstructed spectrum (cosine similarity = 0.559)

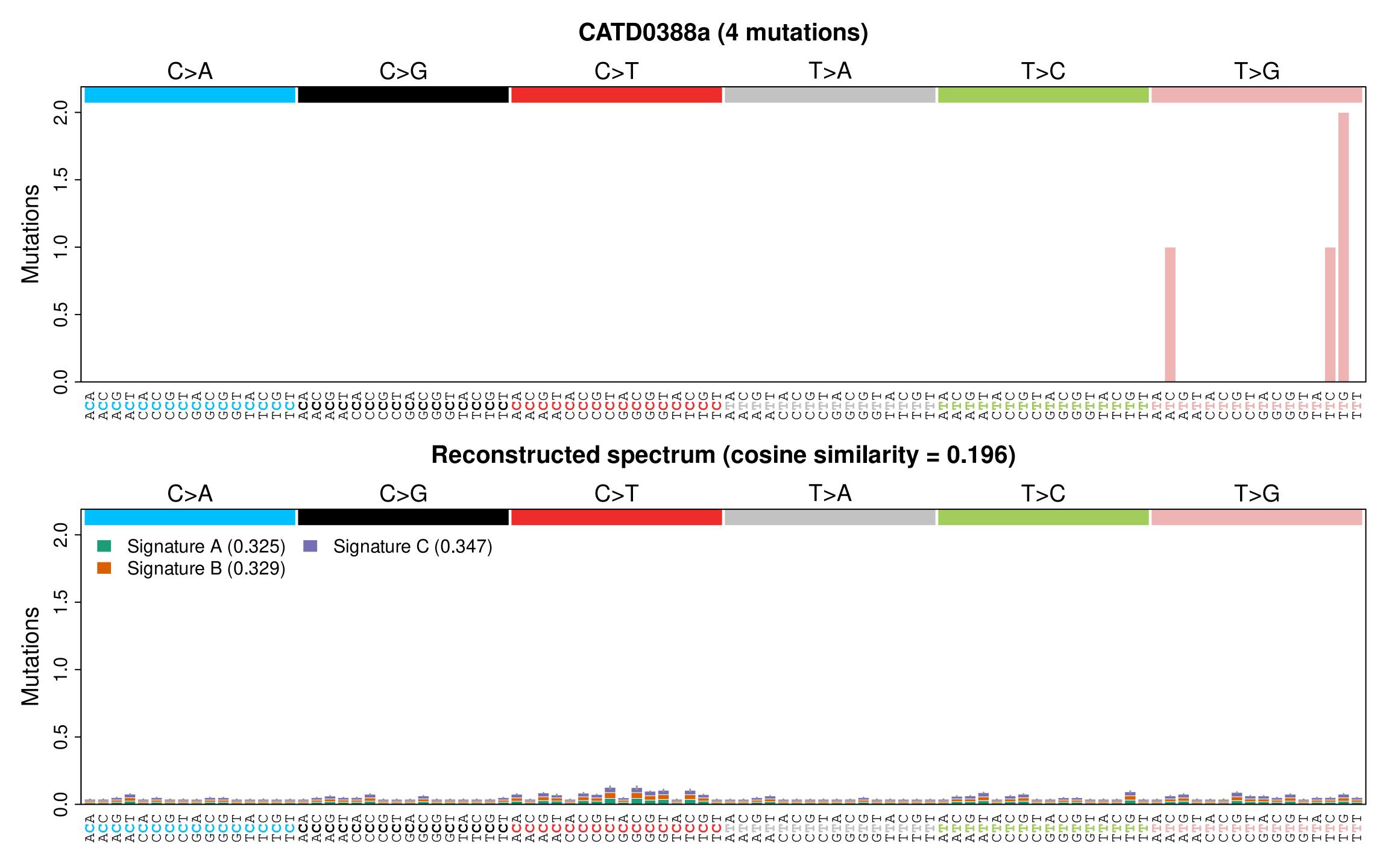


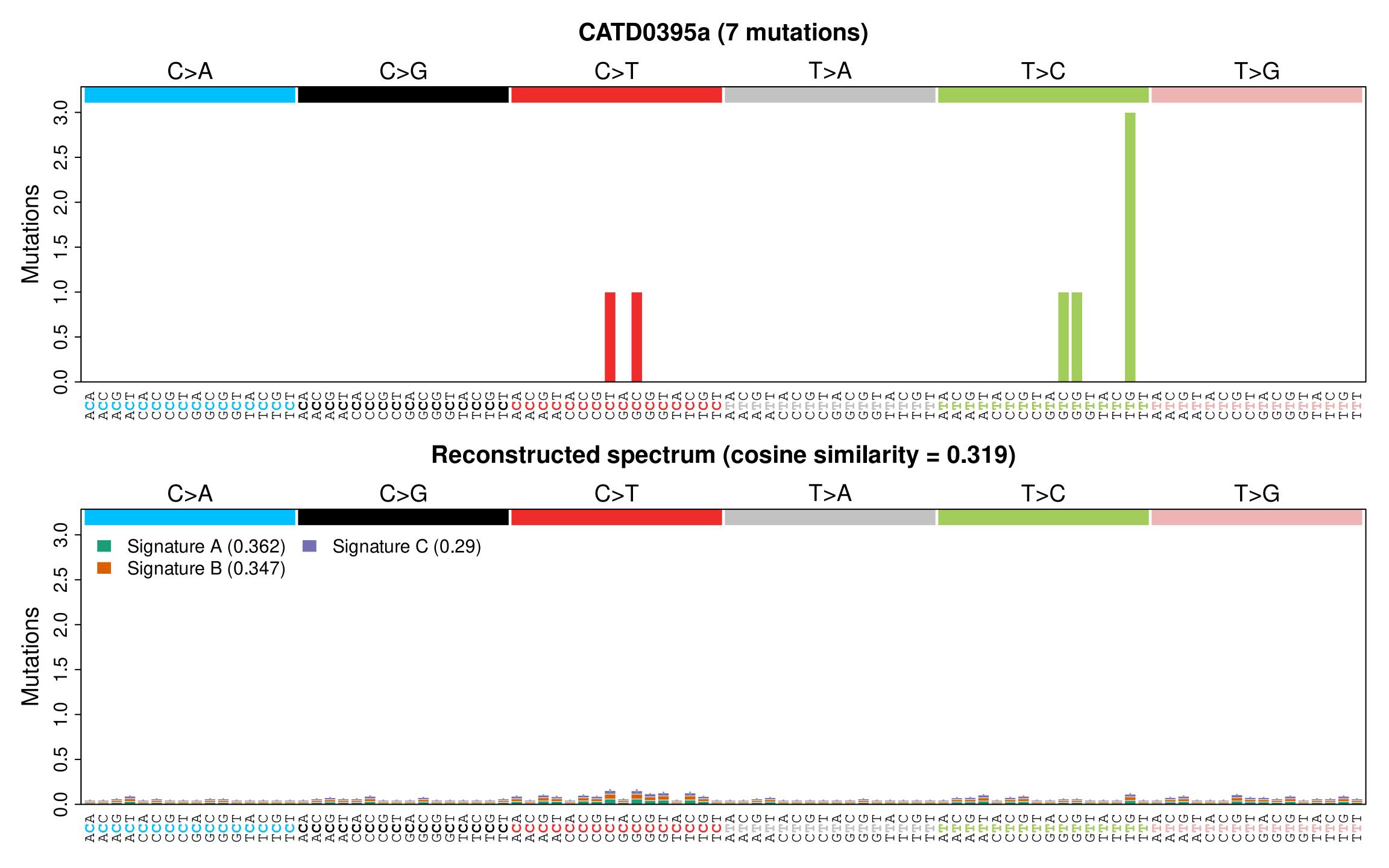
CATD0657a (7 mutations)



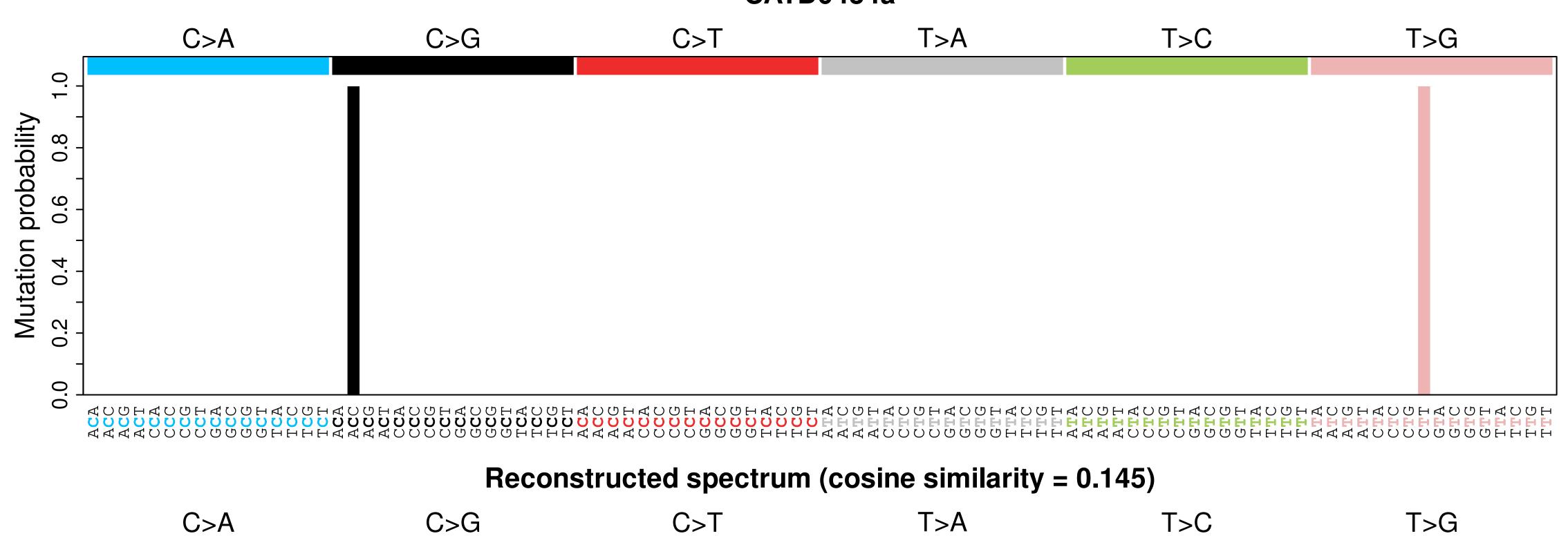
Reconstructed spectrum (cosine similarity = 0.31)

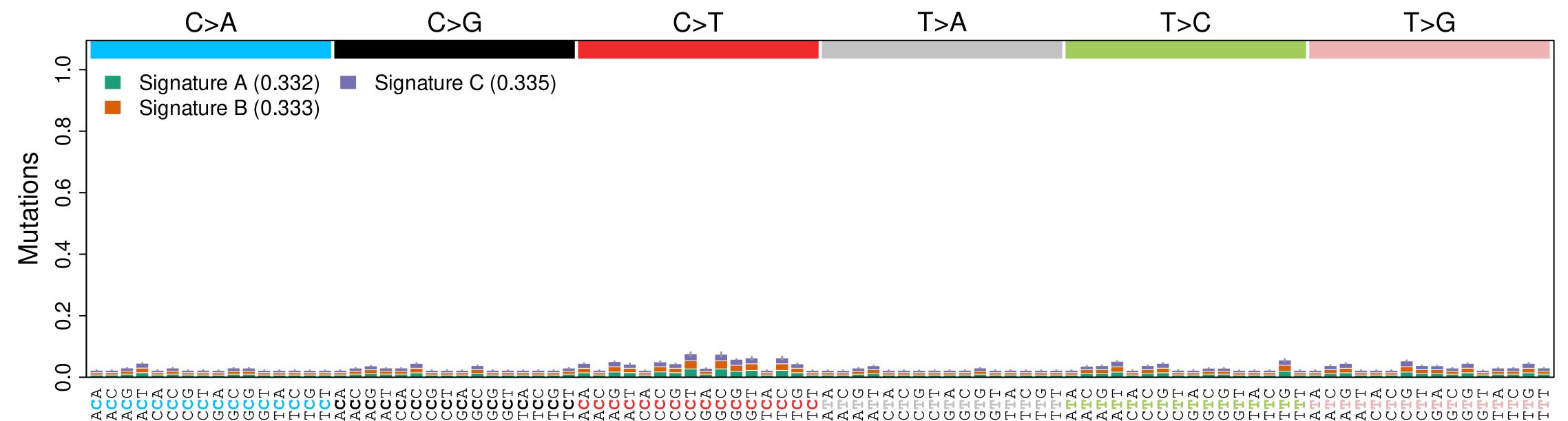




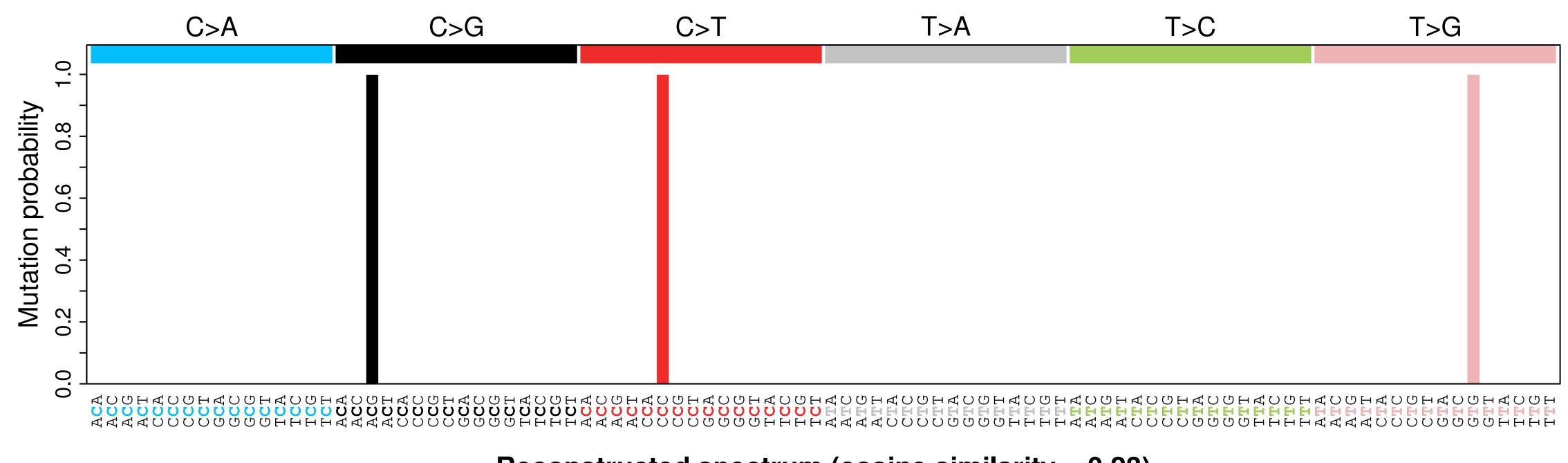


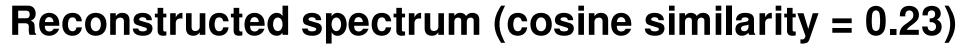


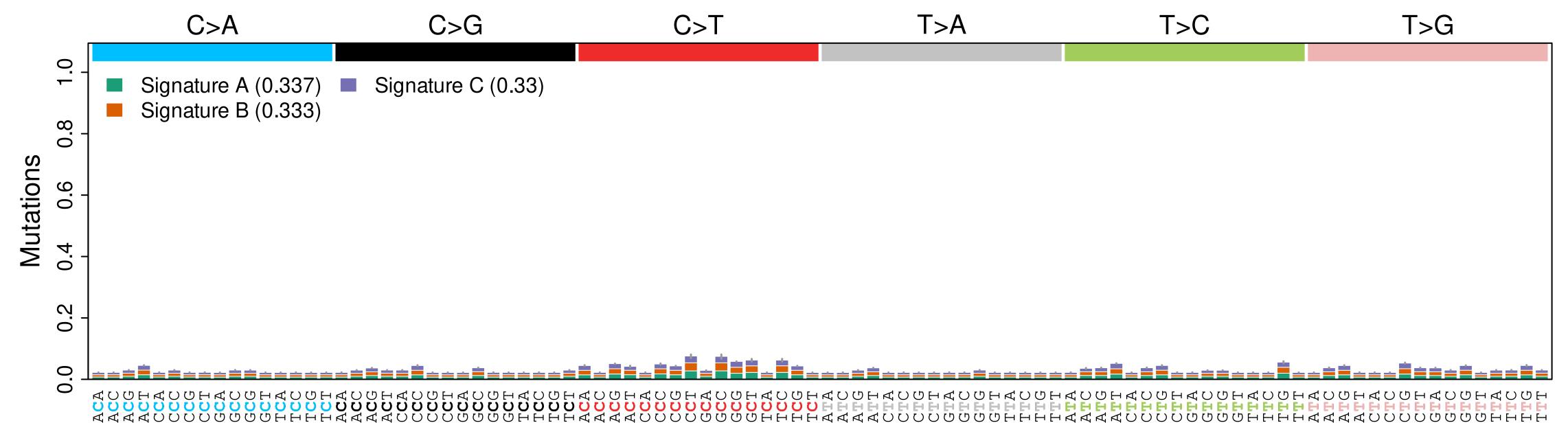






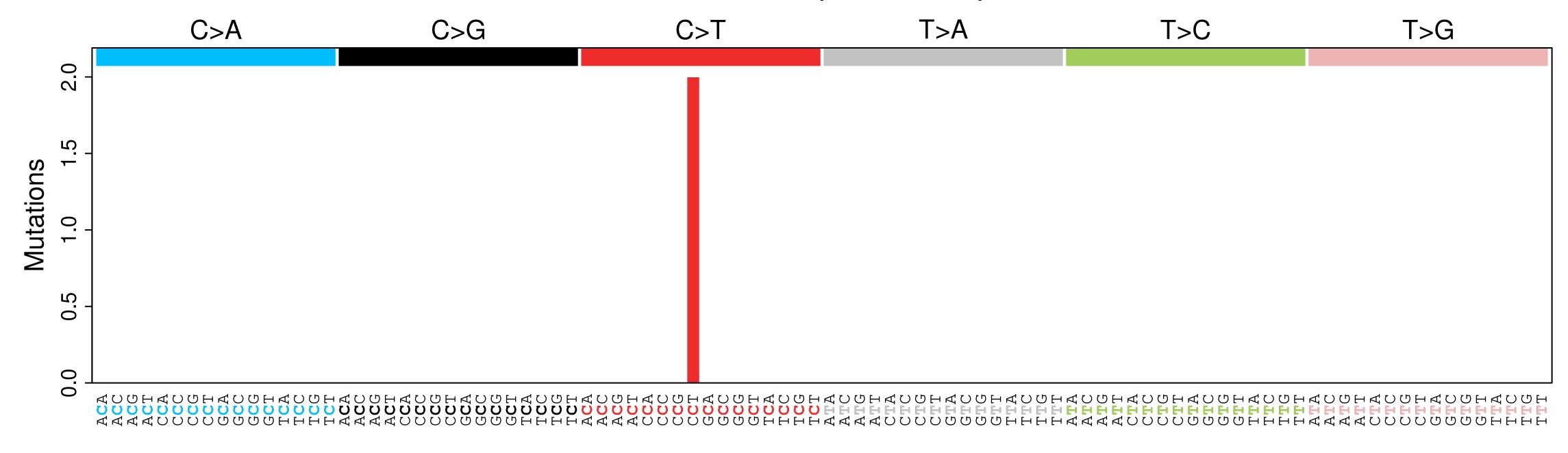


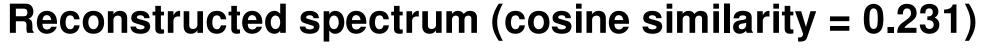


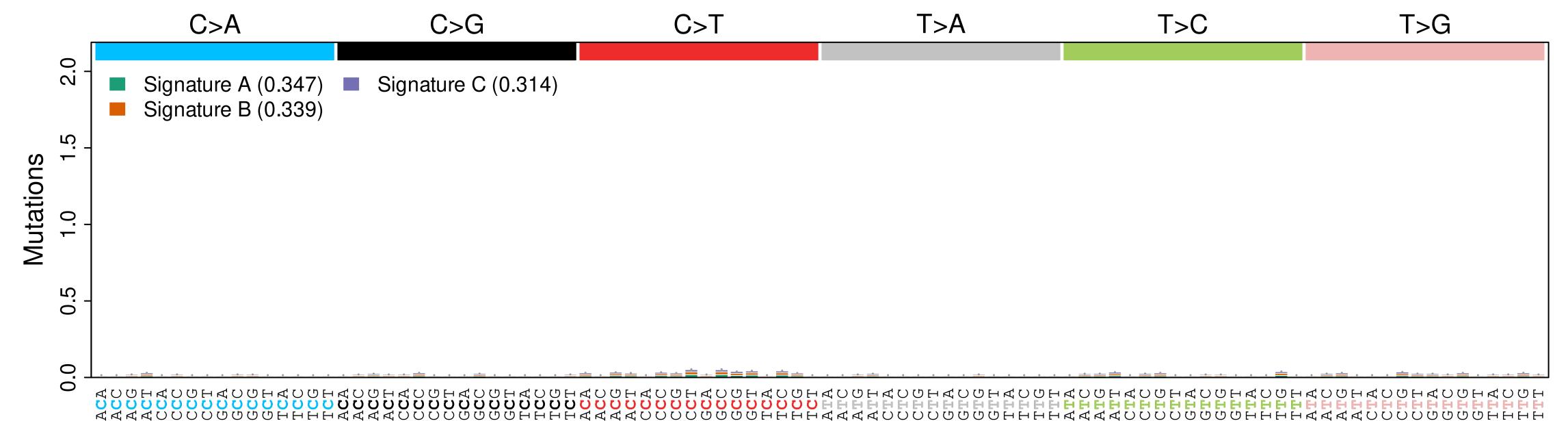


CATD0391a (2 mutations) T>G C>A C>G C>T T>A T>C 2.0 5 Mutations 0.5 0.0 **Reconstructed spectrum (cosine similarity = 0.161)** C>A C>G T>A T>C T>G C>T 2.0 Signature A (0.327) Signature C (0.342) Signature B (0.331) 5 Mutations 5 0

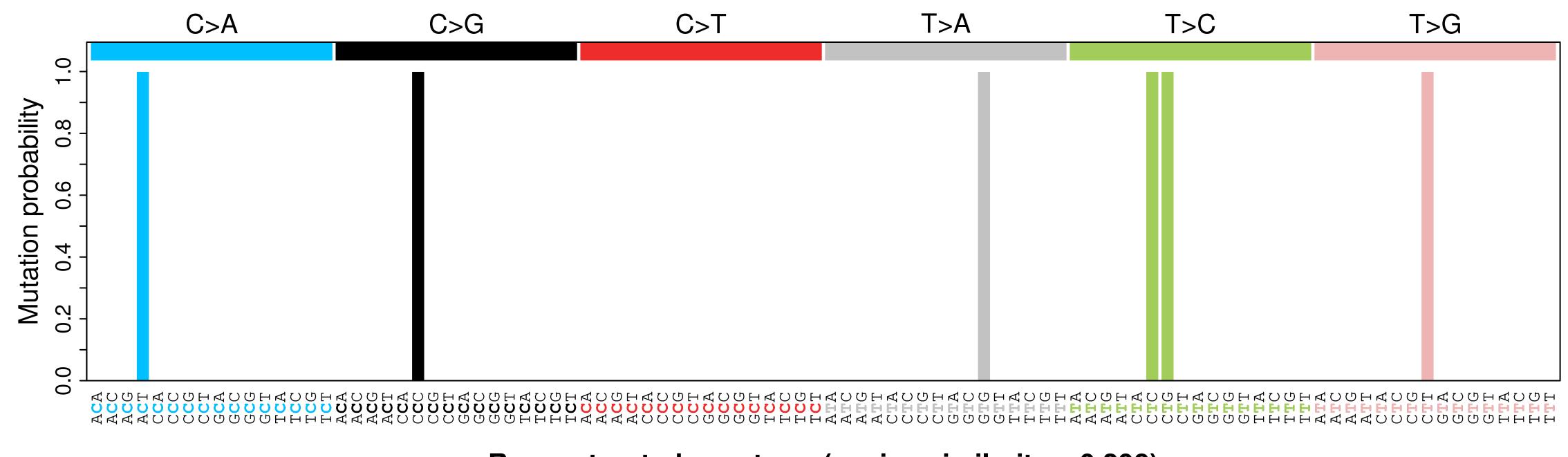
CATD0387a (2 mutations)

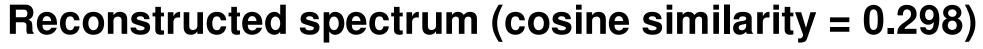


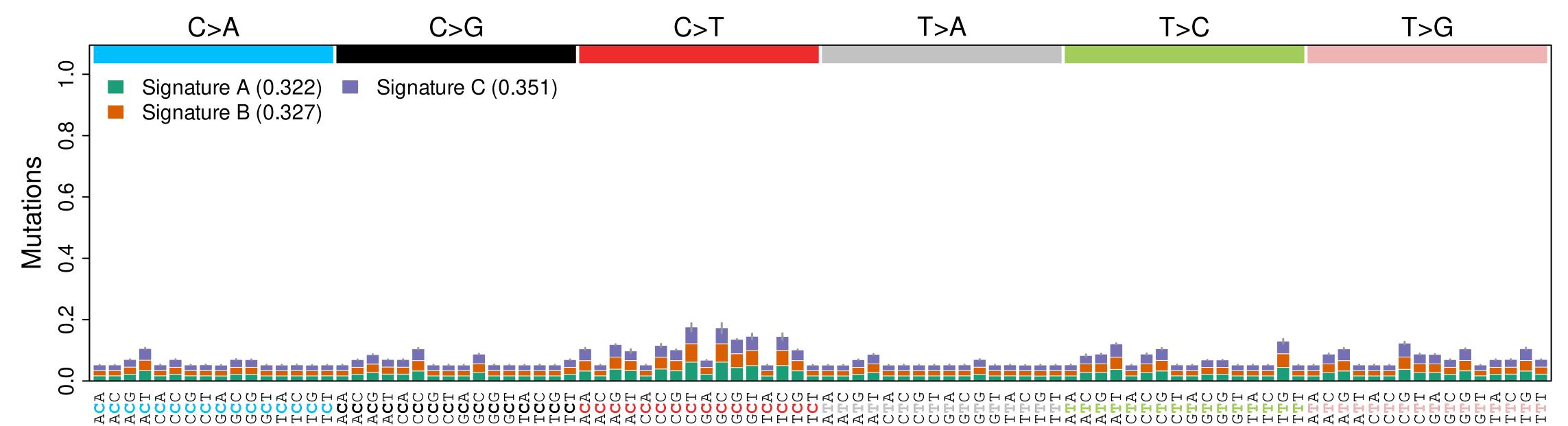




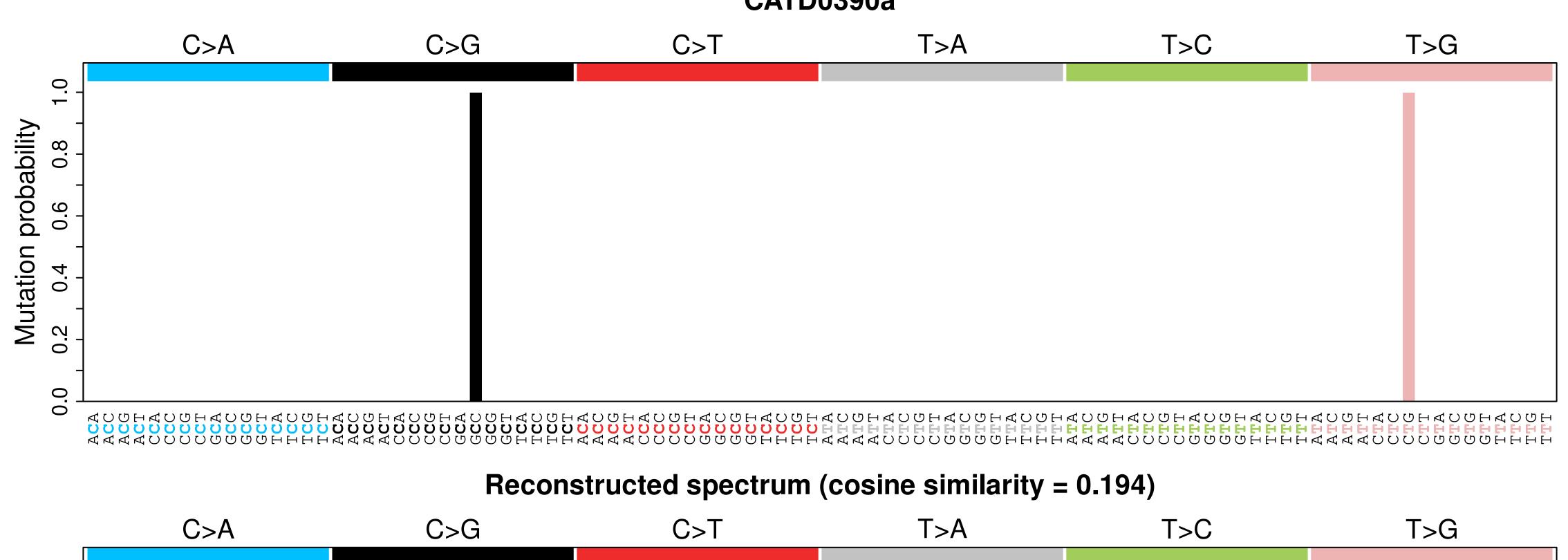
CATD0381a

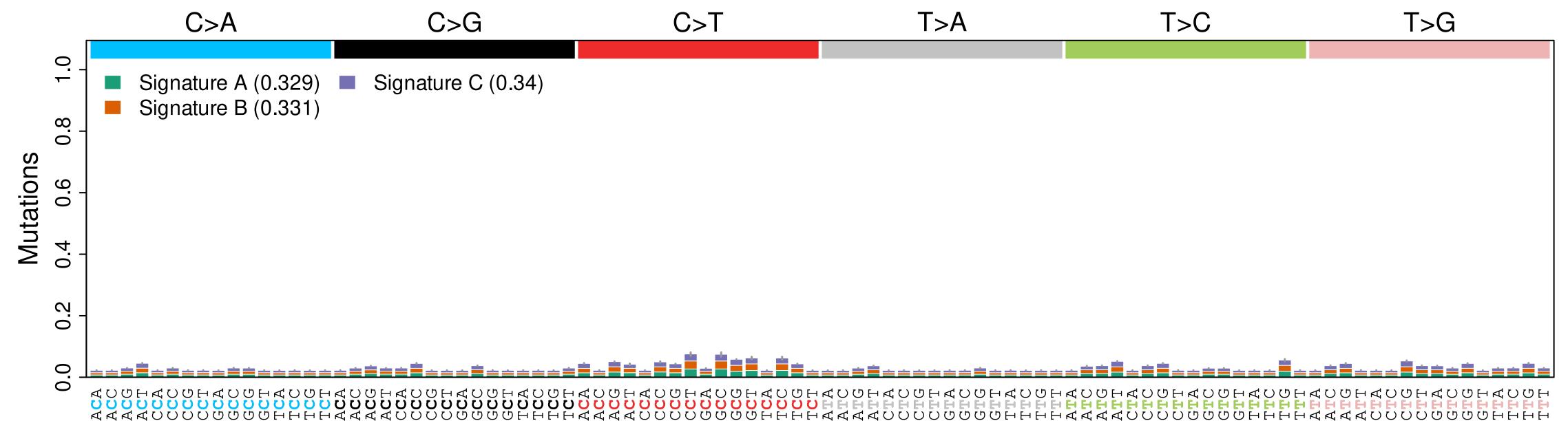




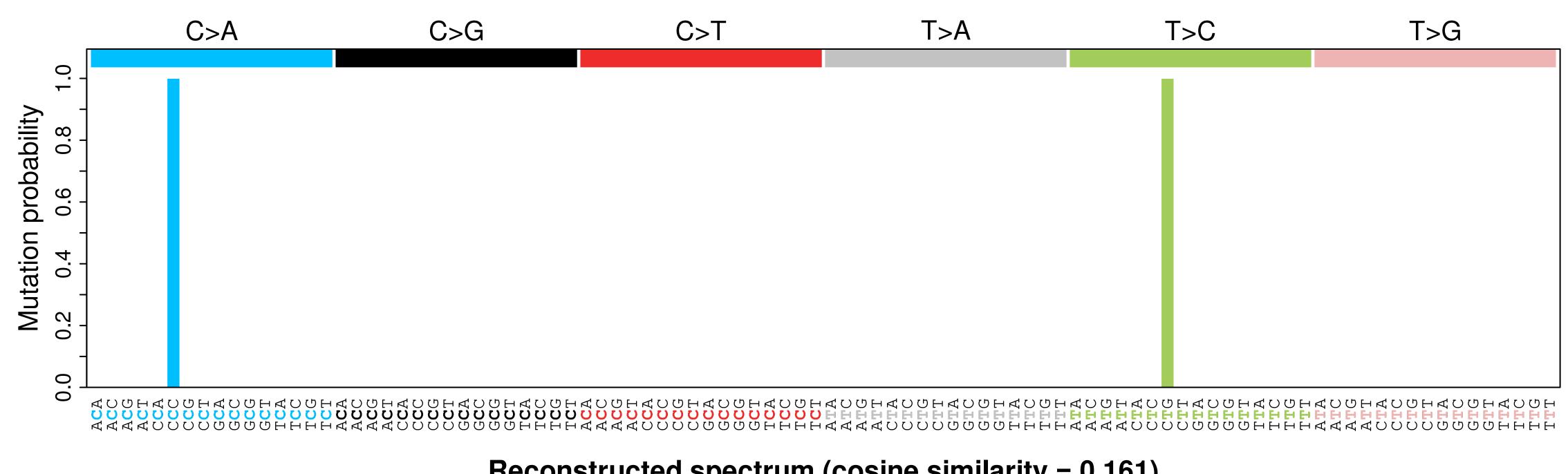


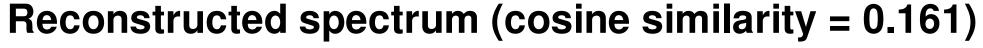


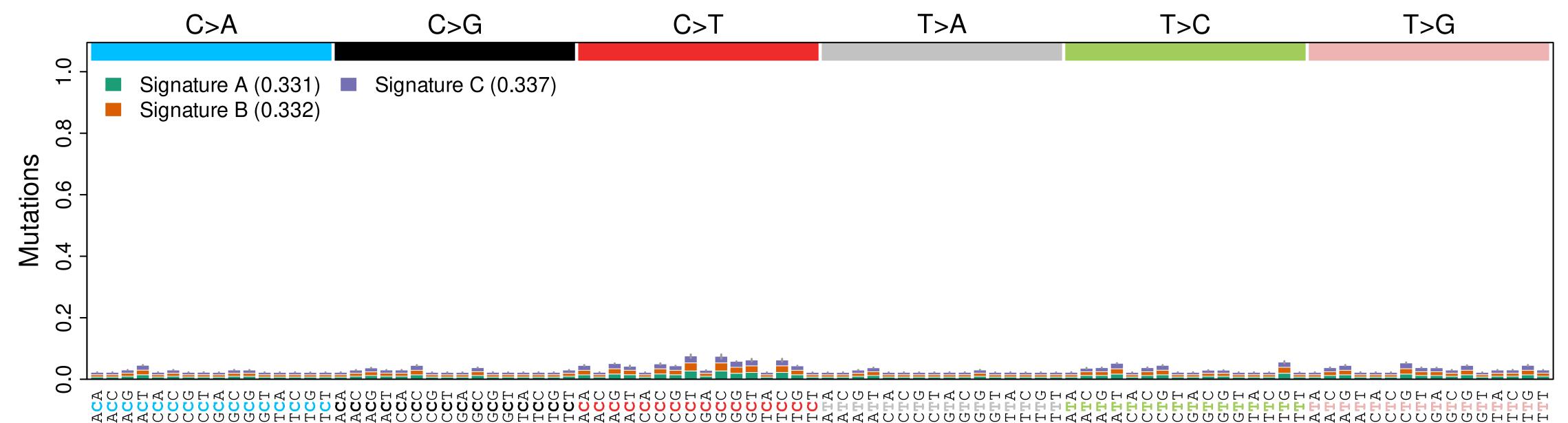




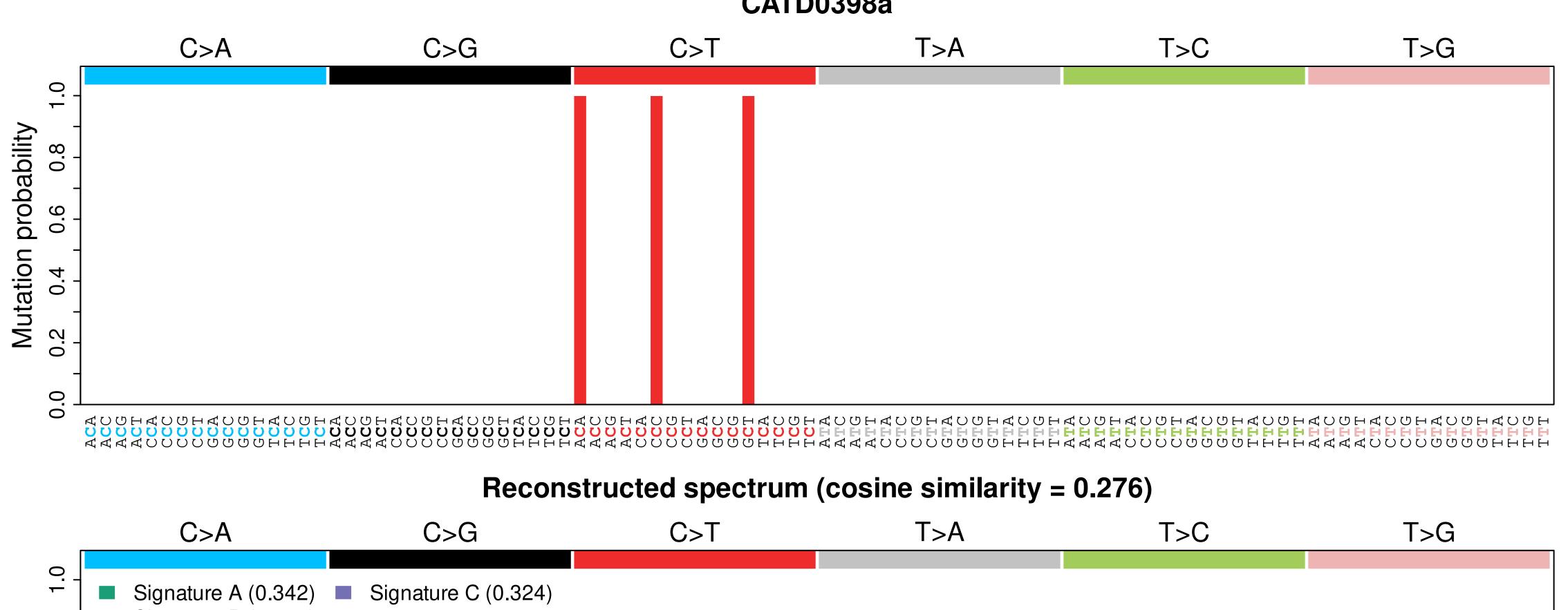


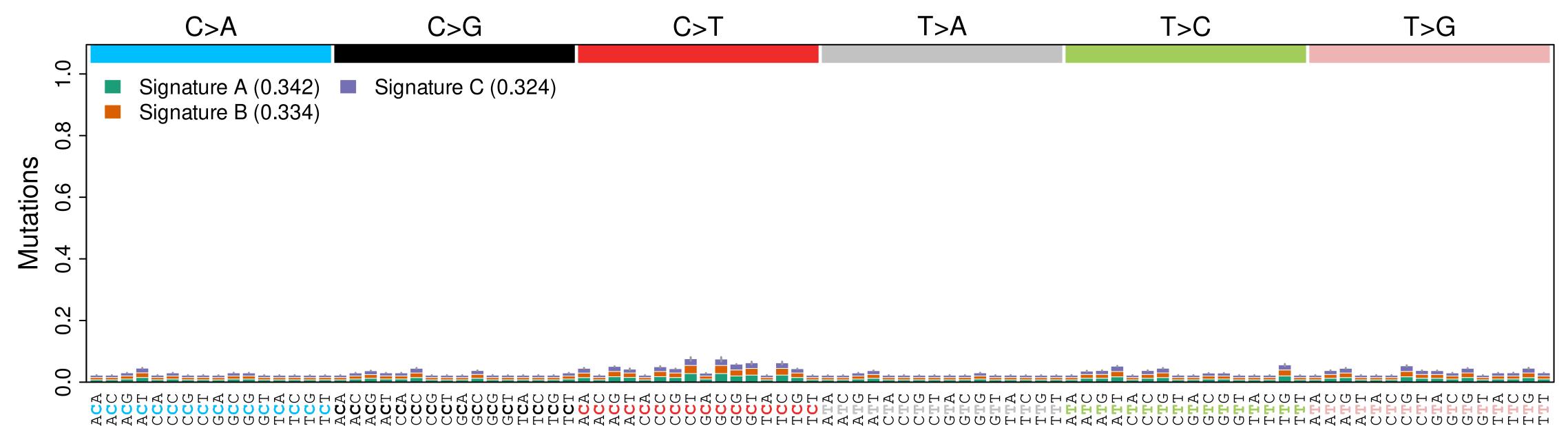




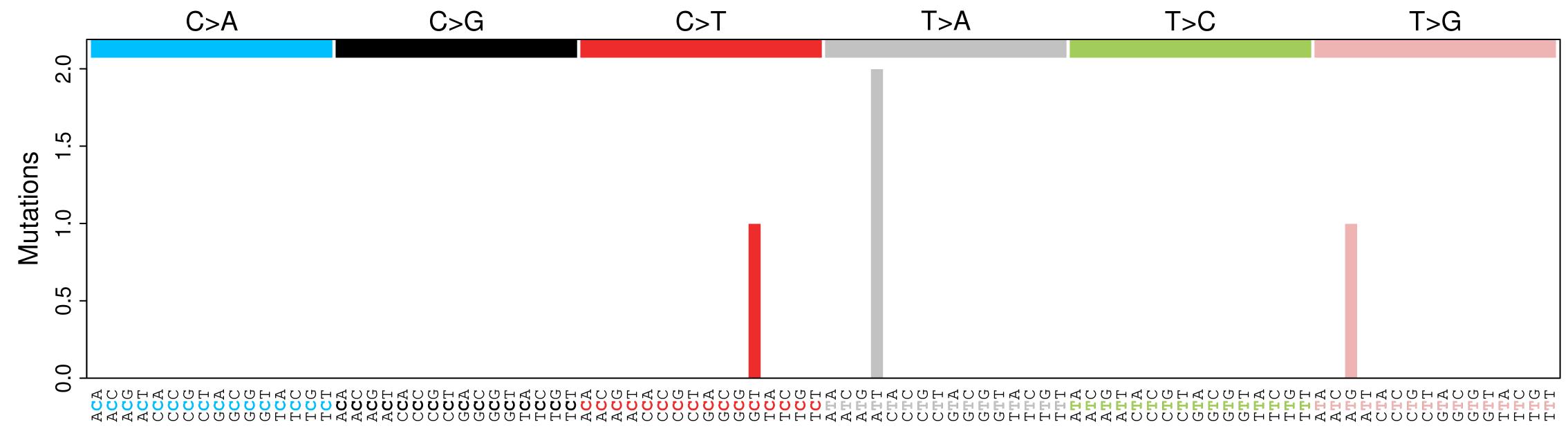




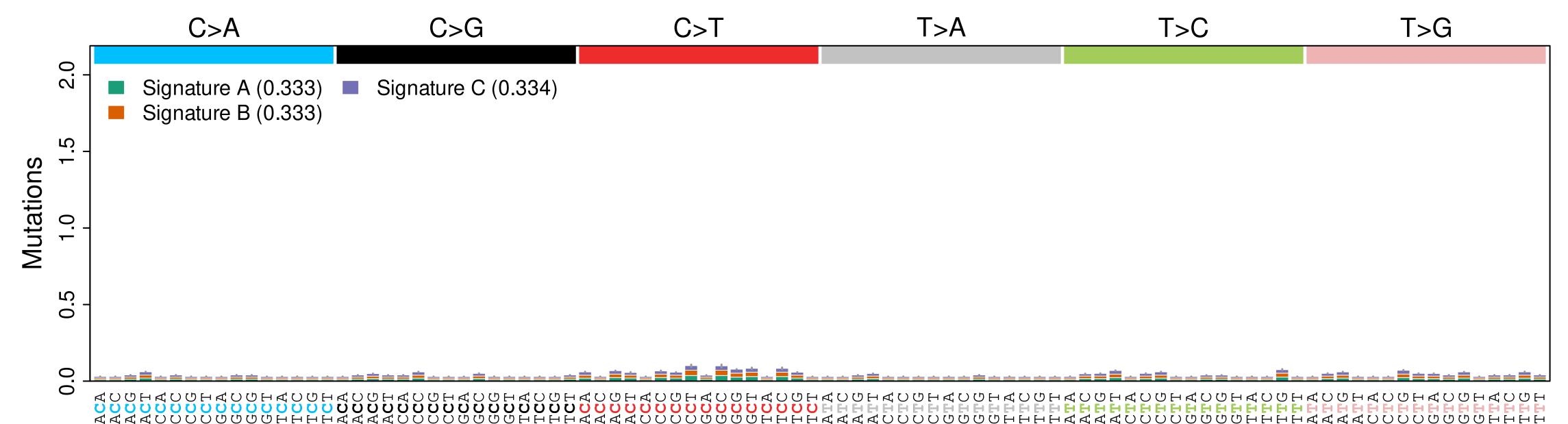




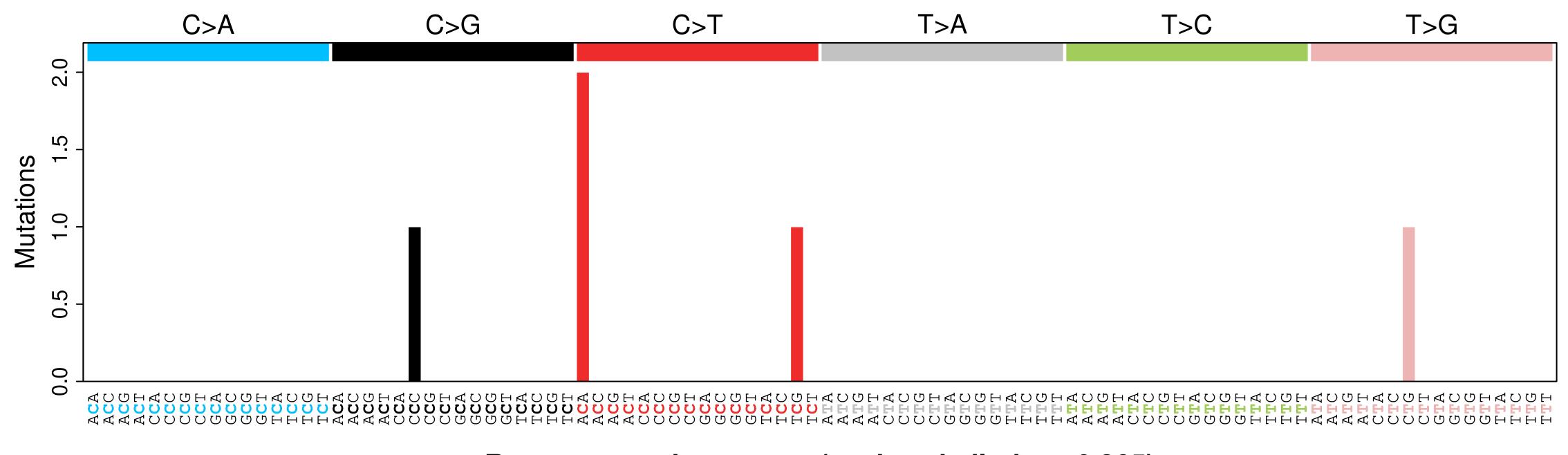
CATD0428a (4 mutations) C>A C>G C>T T>A



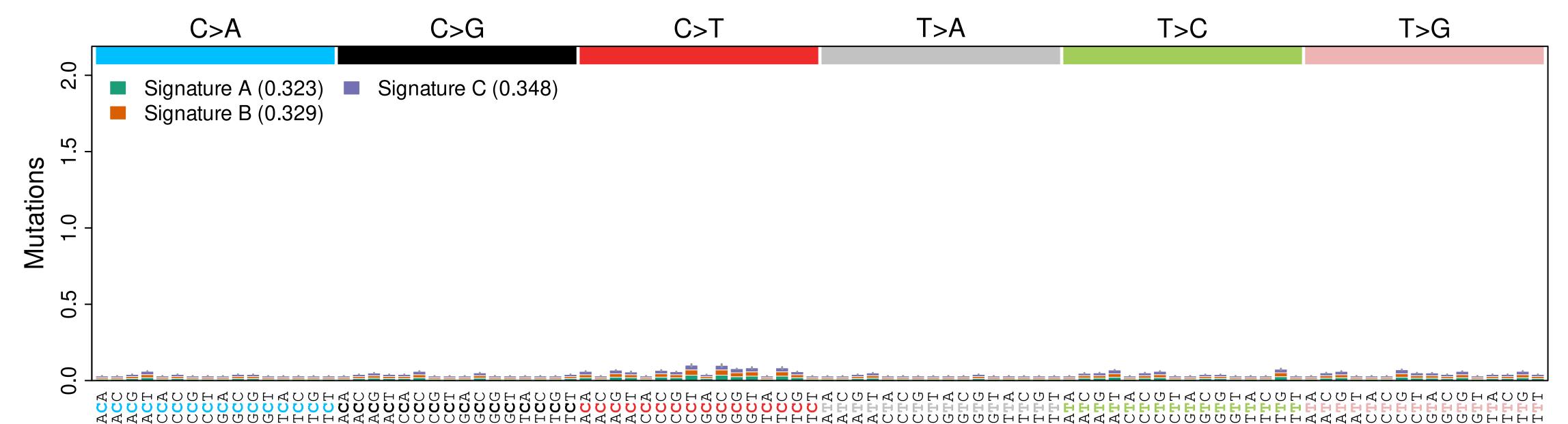
Reconstructed spectrum (cosine similarity = 0.225)



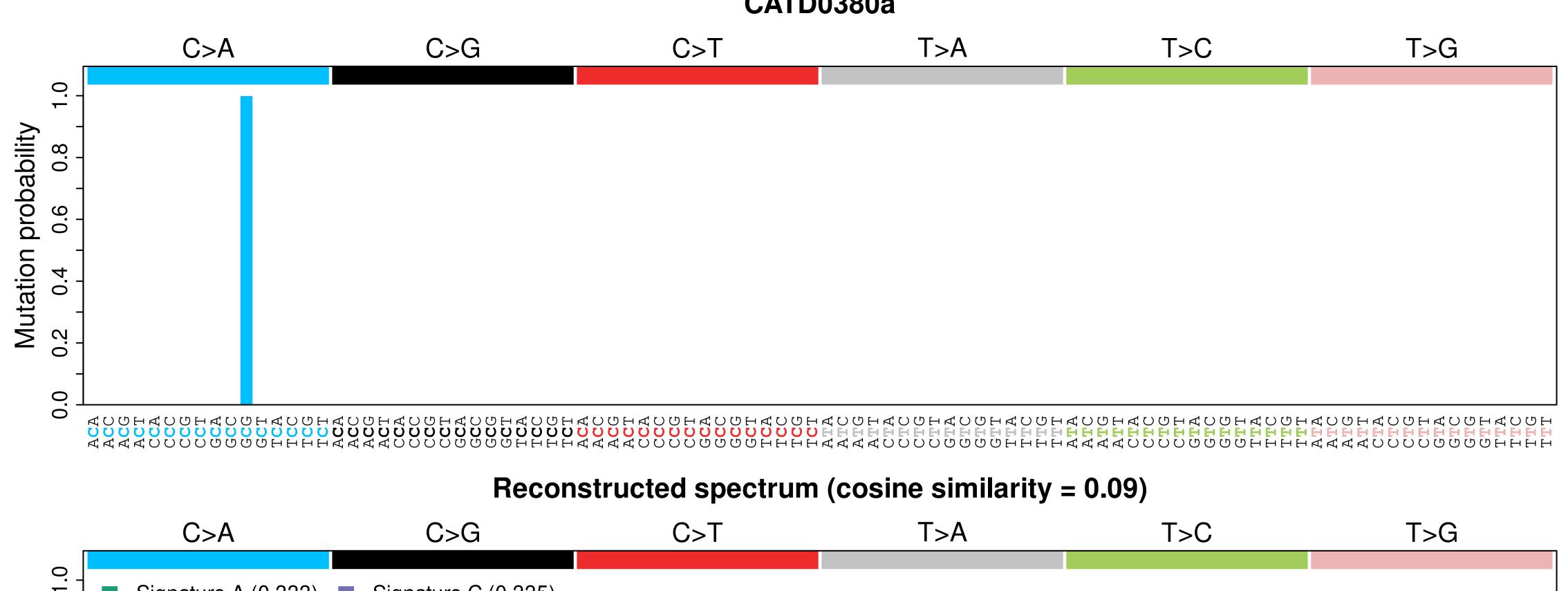
CATD0378a (5 mutations)

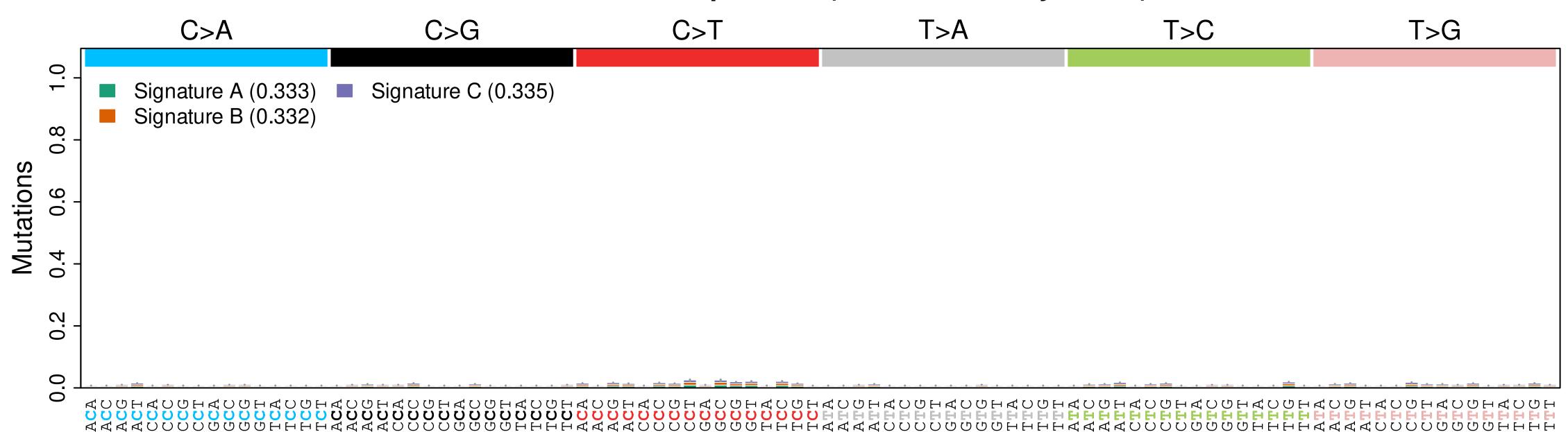




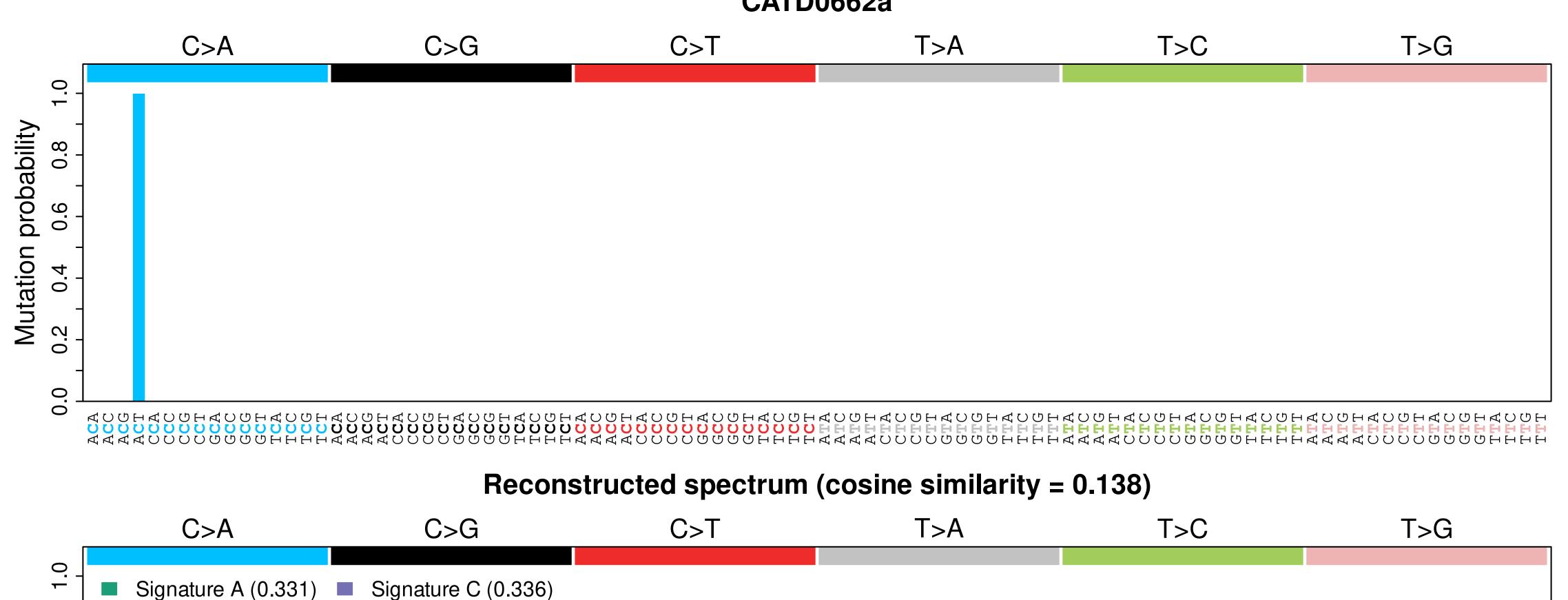


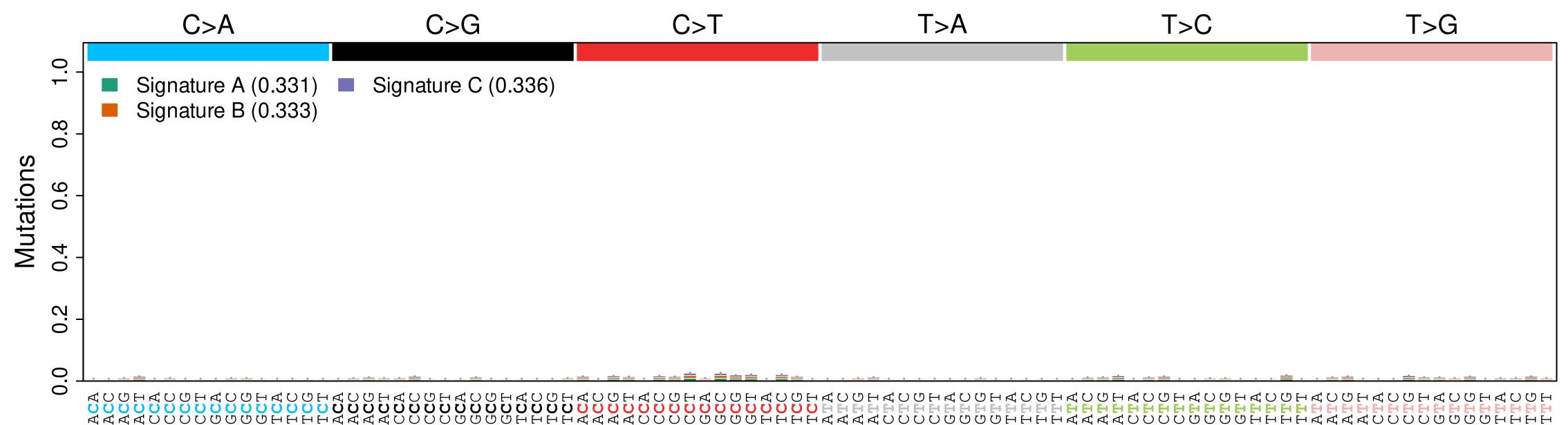




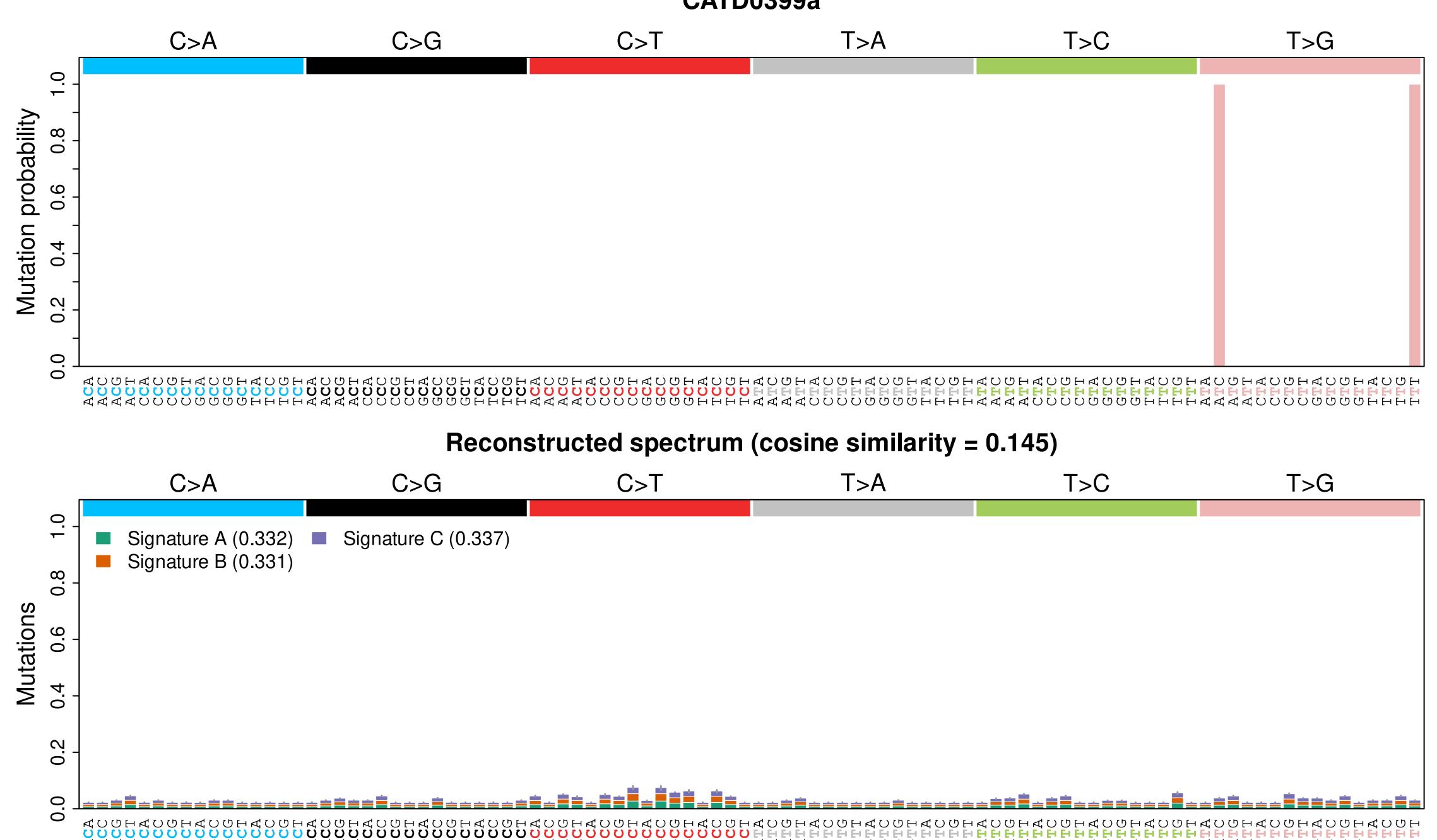




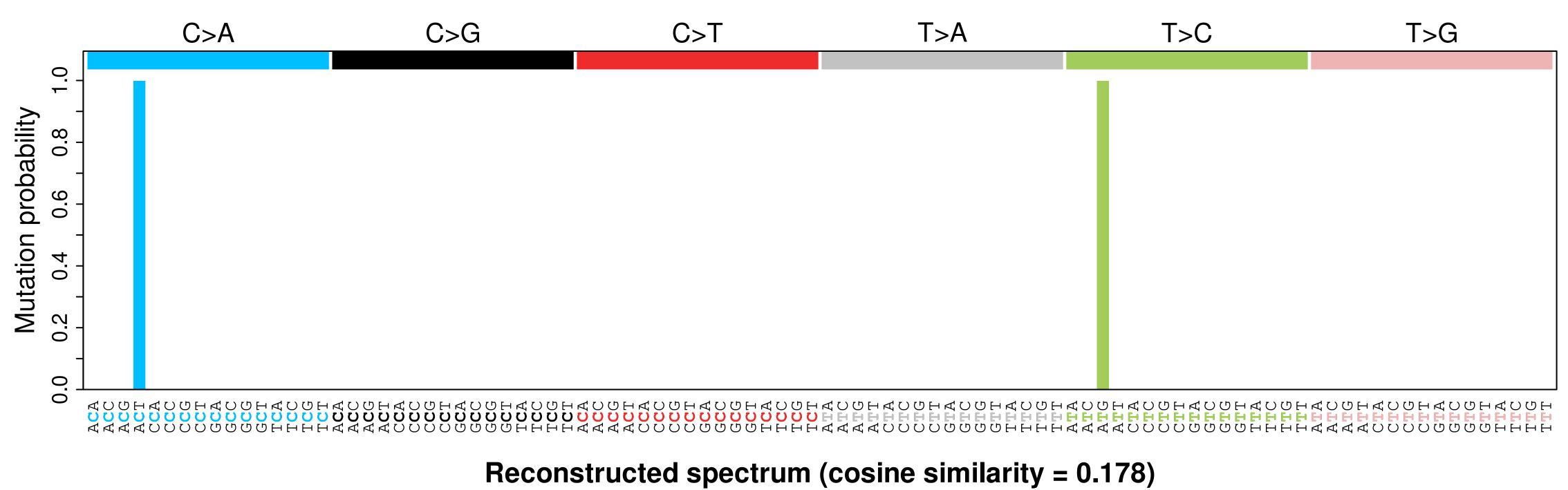


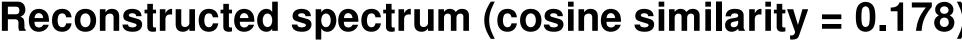


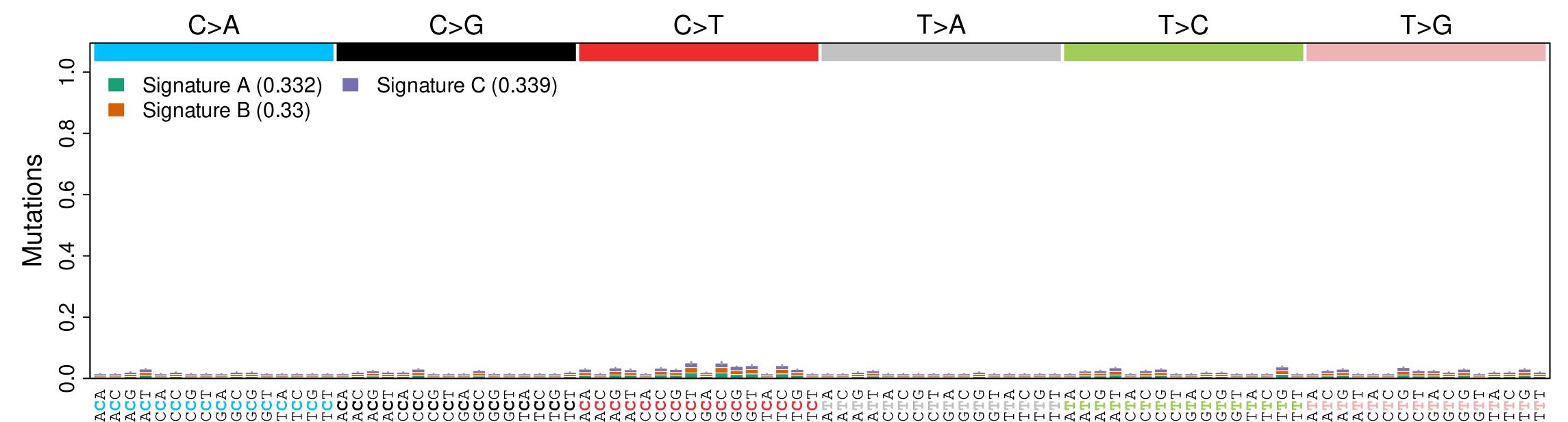




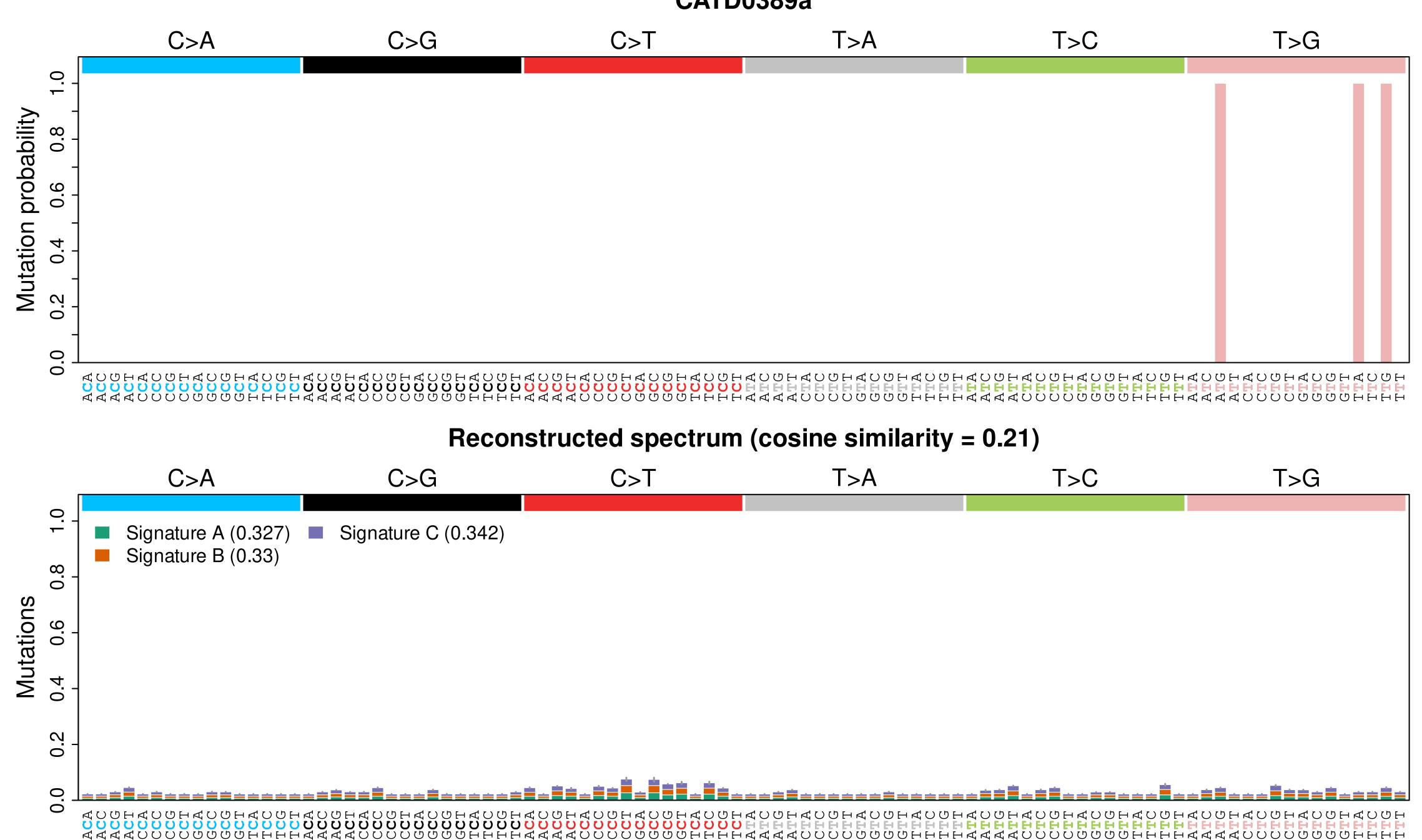




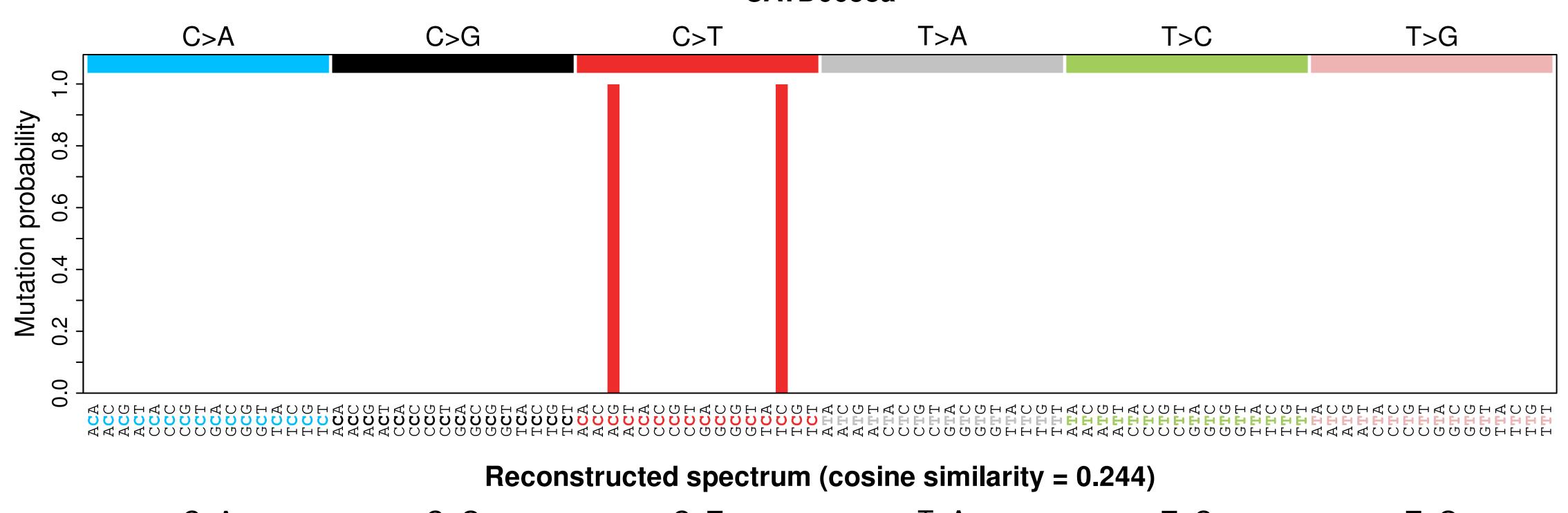


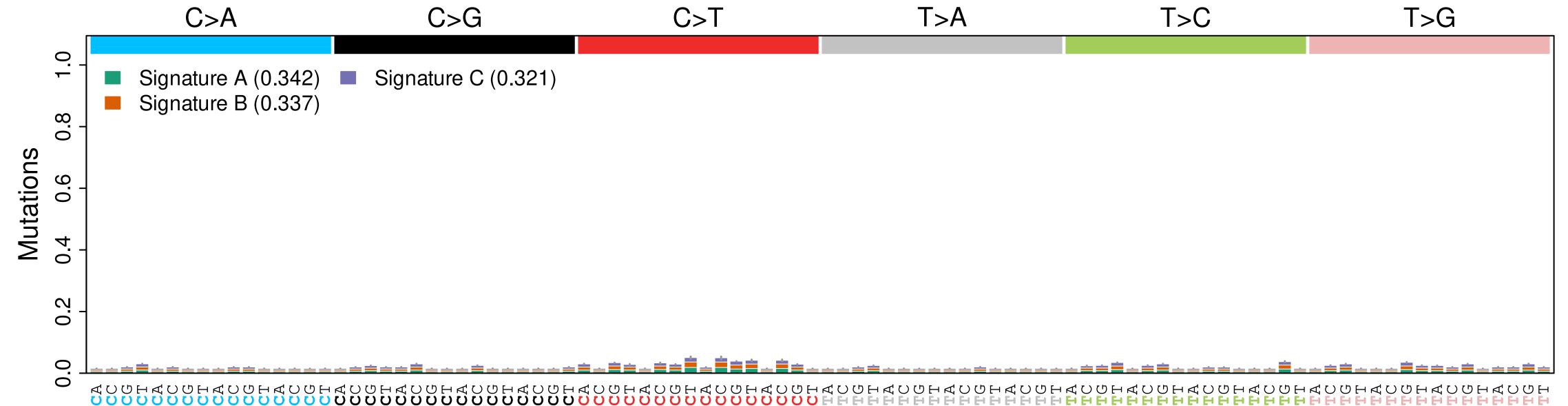






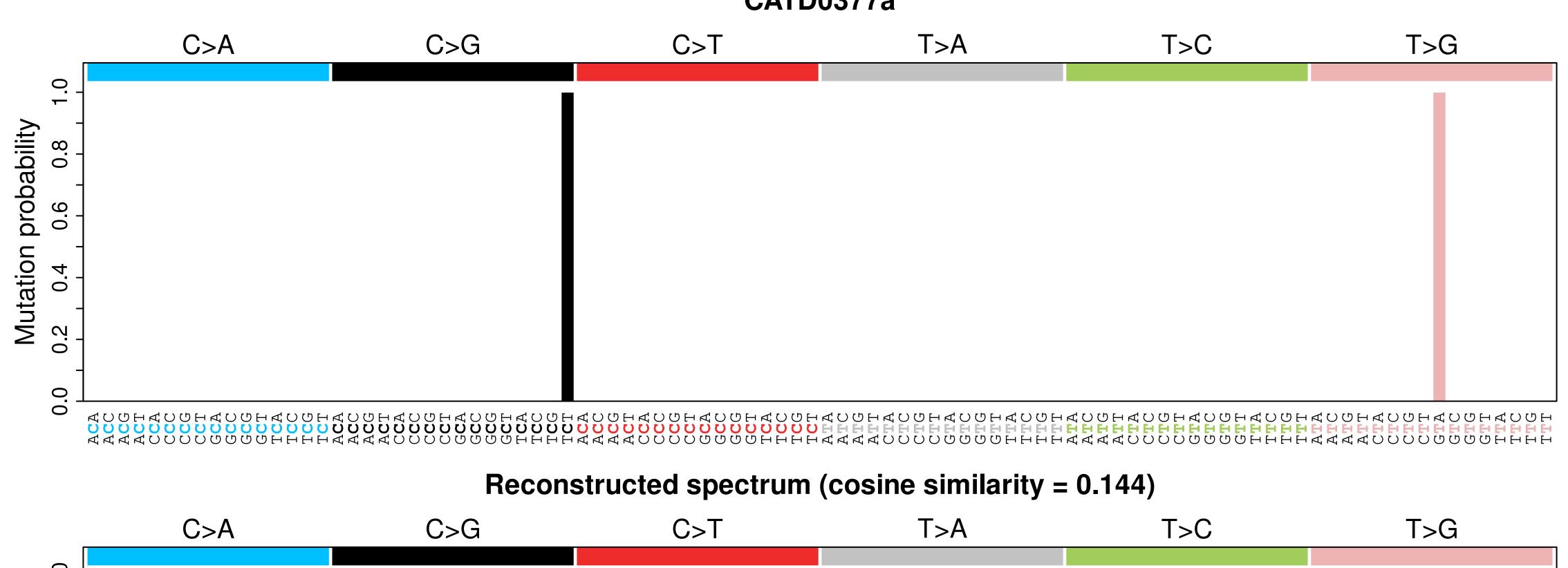


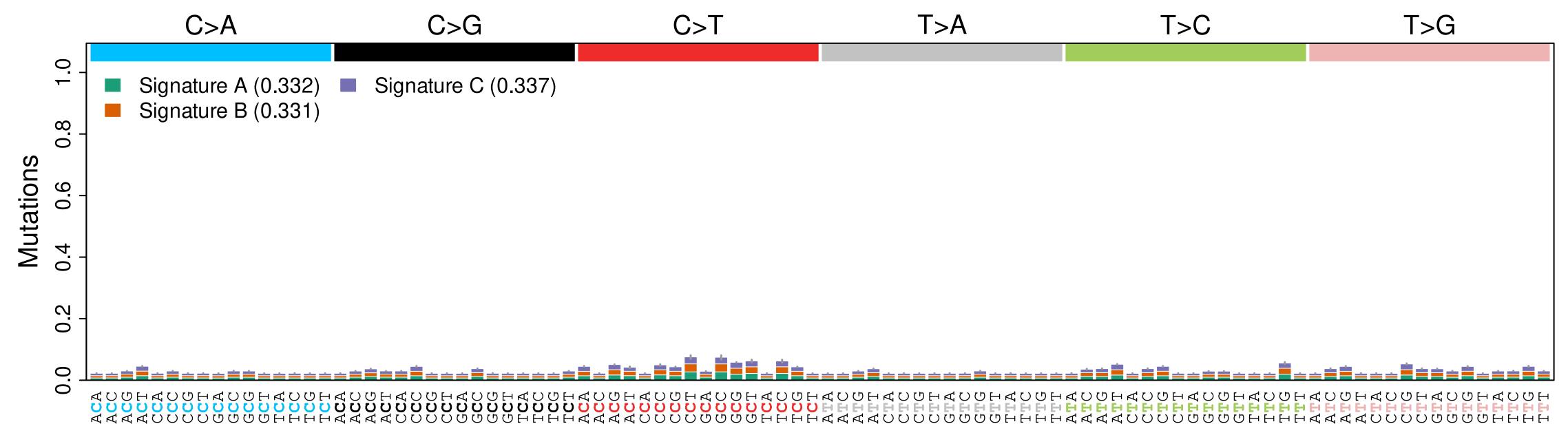




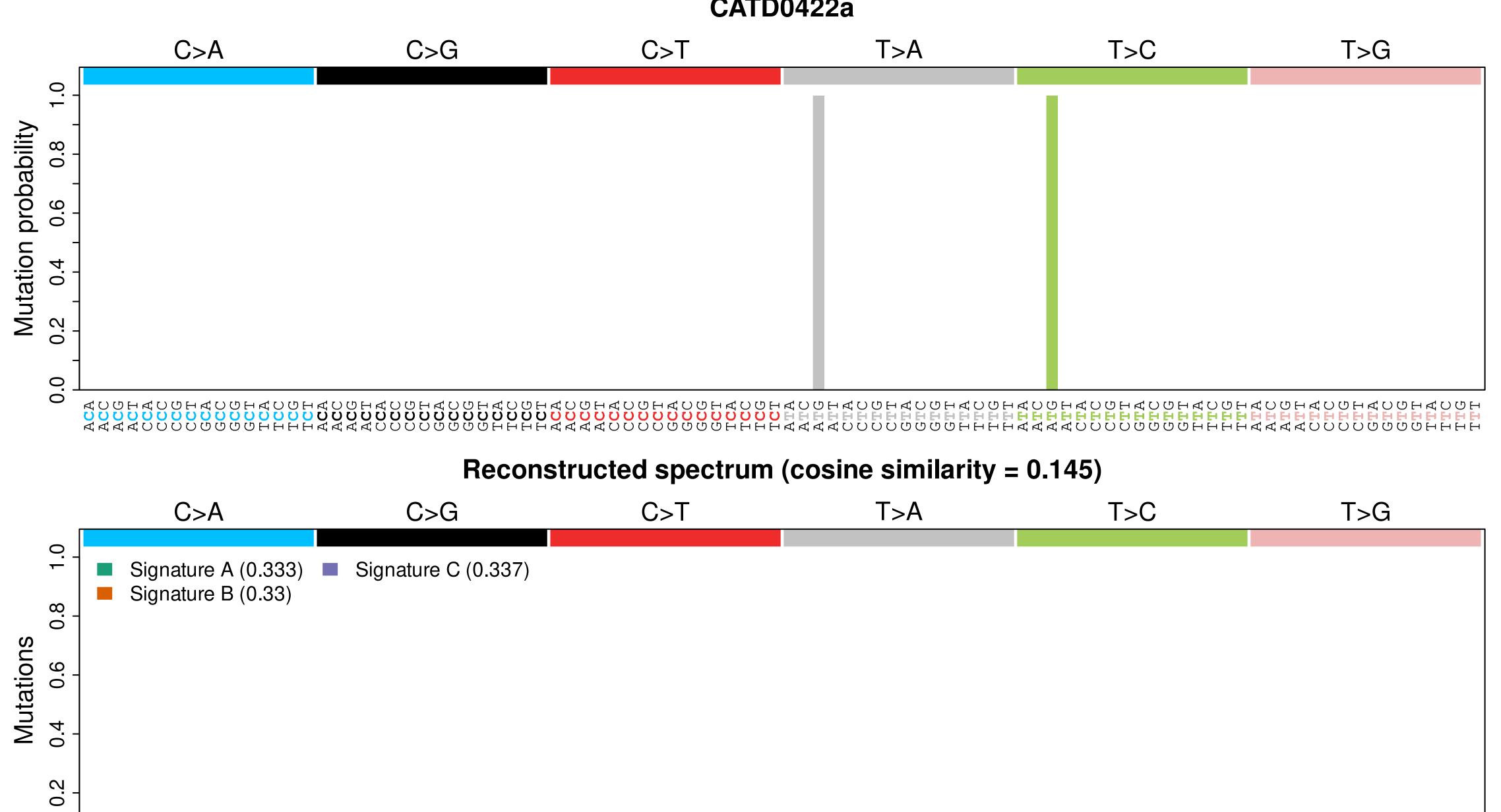
CATD0665a (4 mutations) T>G C>A C>G C>T T>A T>C 2.0 5 Mutations 0.5 0.0 **Reconstructed spectrum (cosine similarity = 0.237)** C>A C>G T>A T>C T>G C>T 2.0 Signature A (0.333) Signature C (0.335) Signature B (0.332) 5 Mutations 5 0



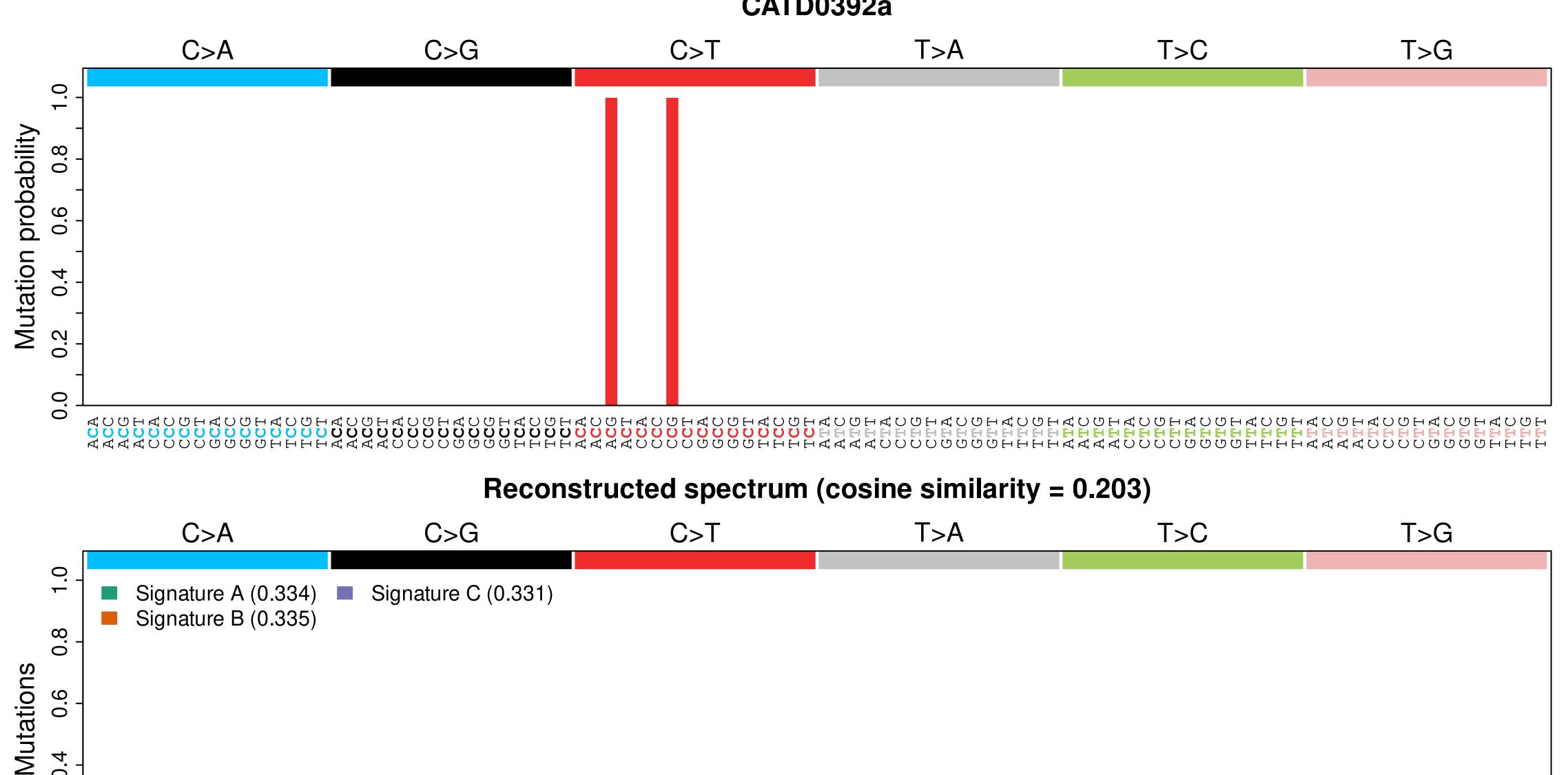






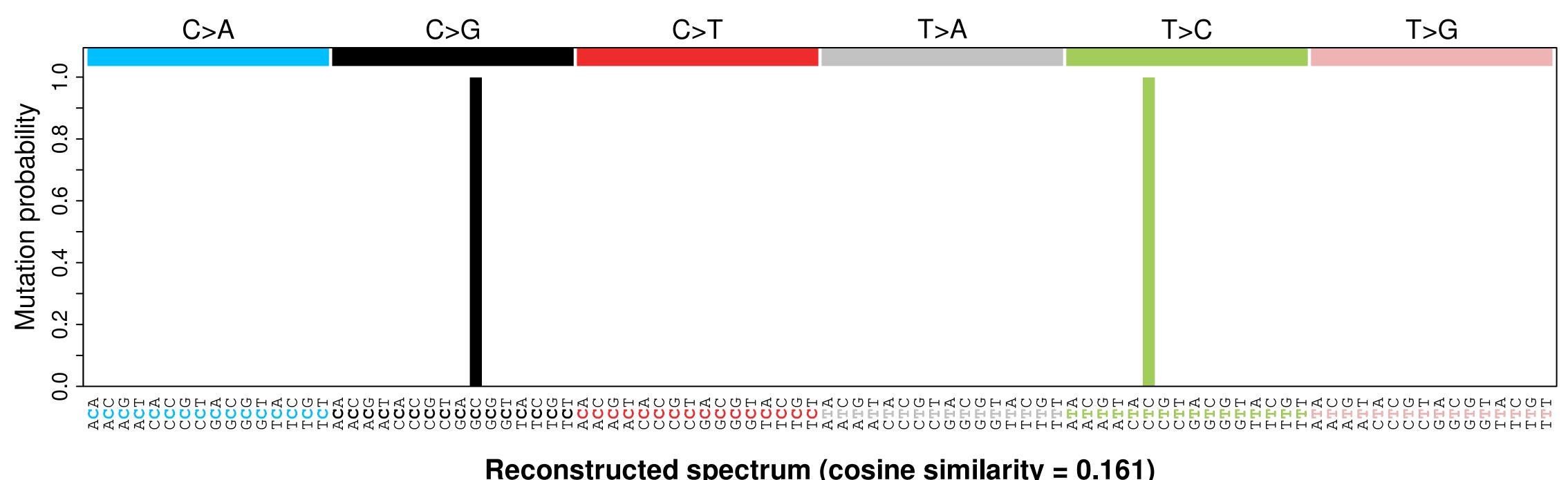


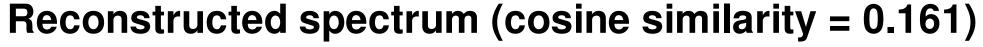


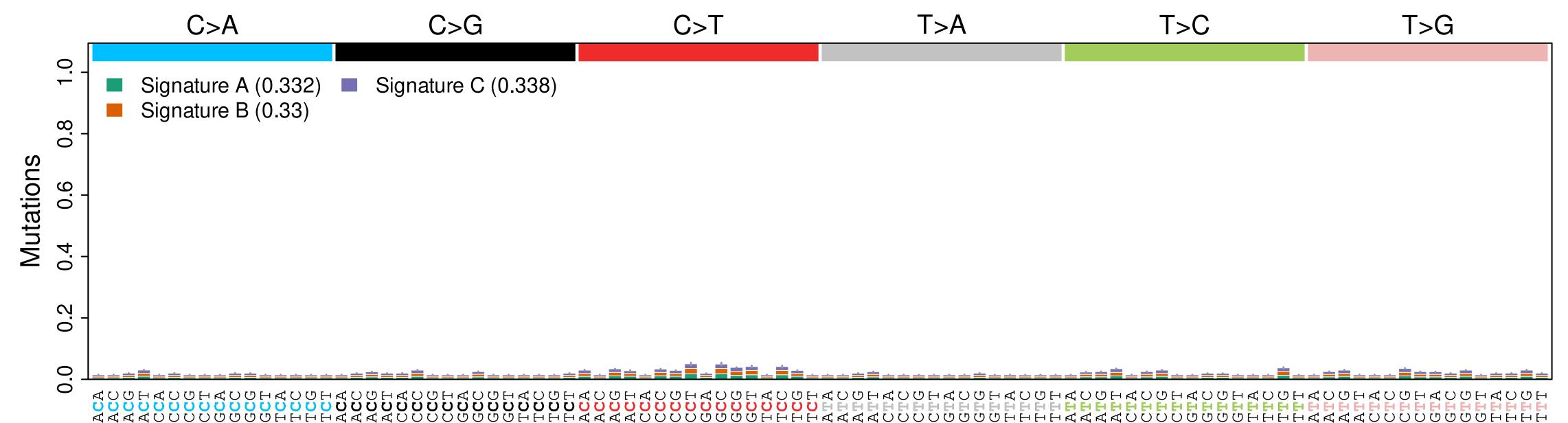


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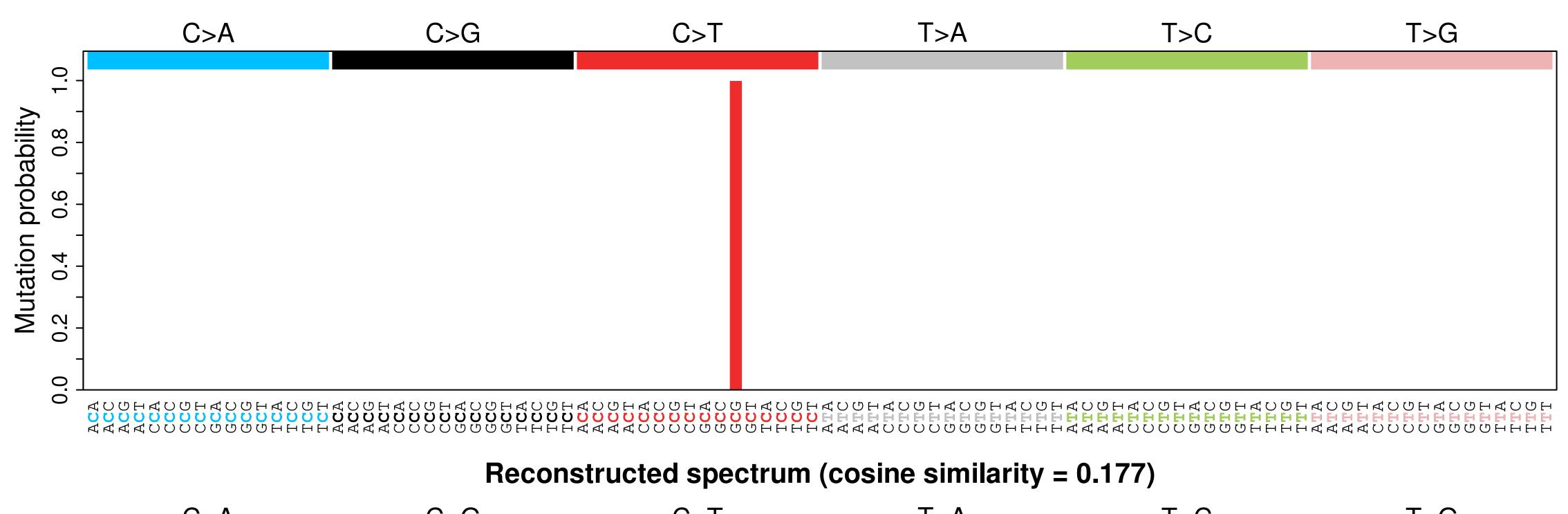


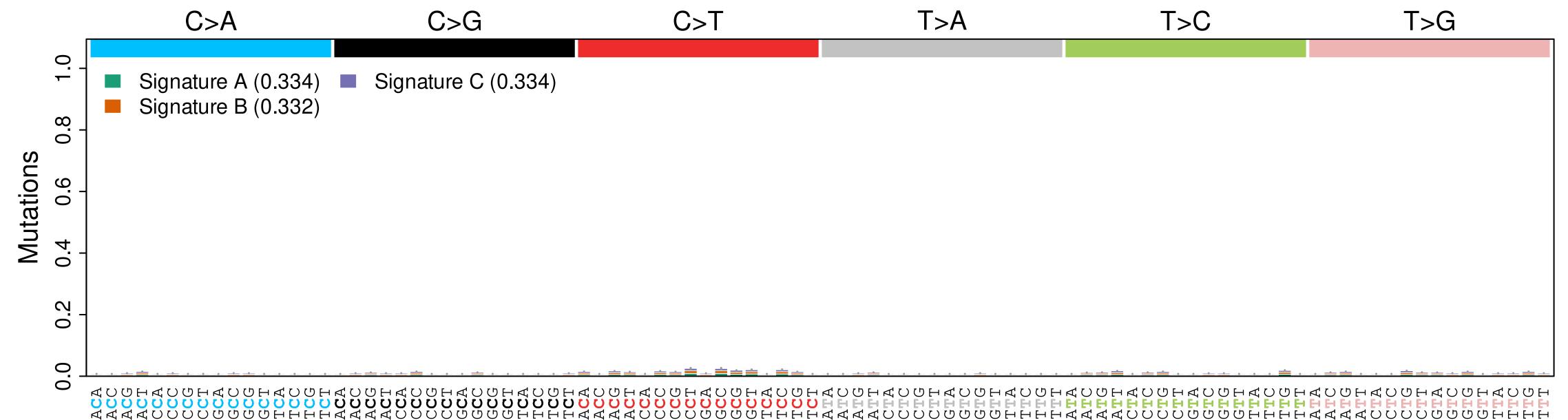




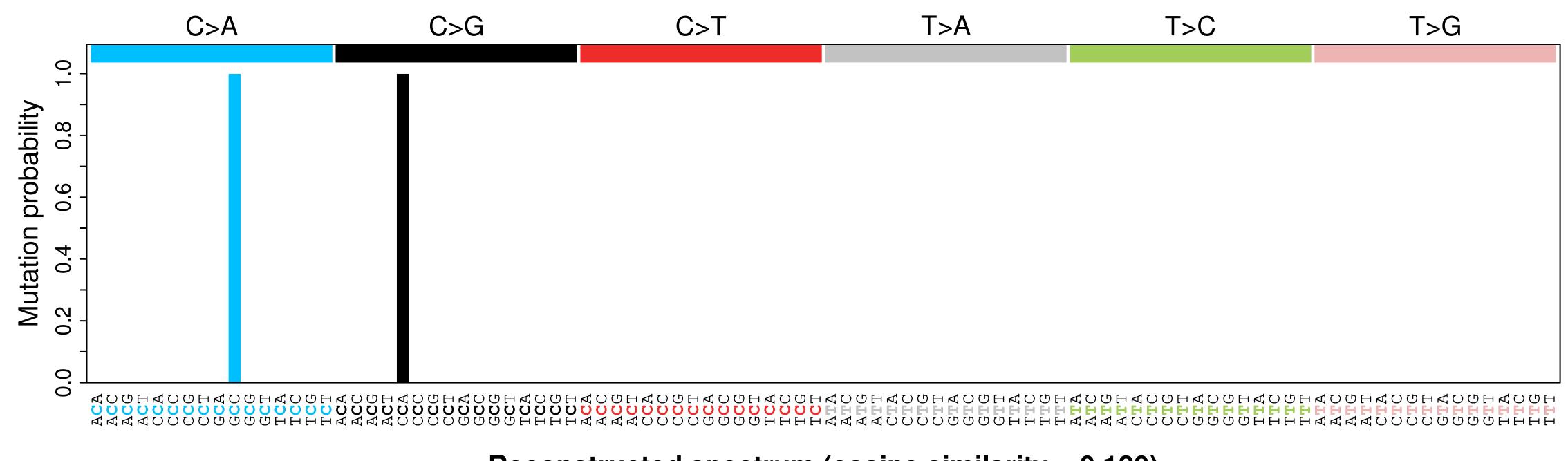


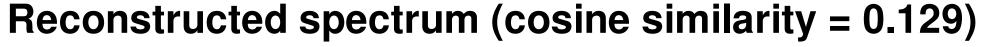


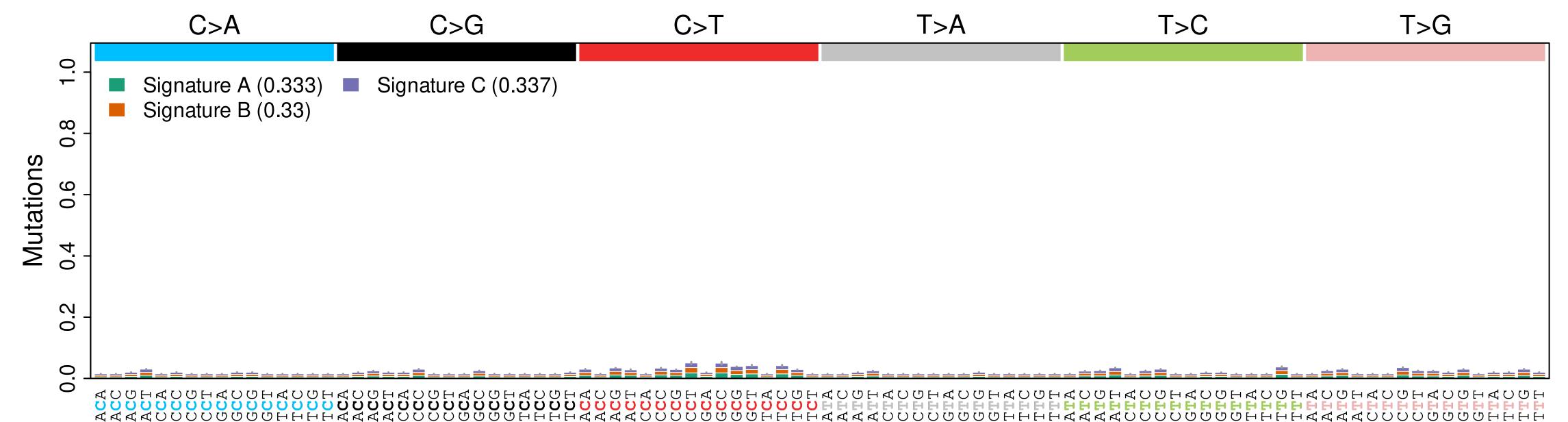




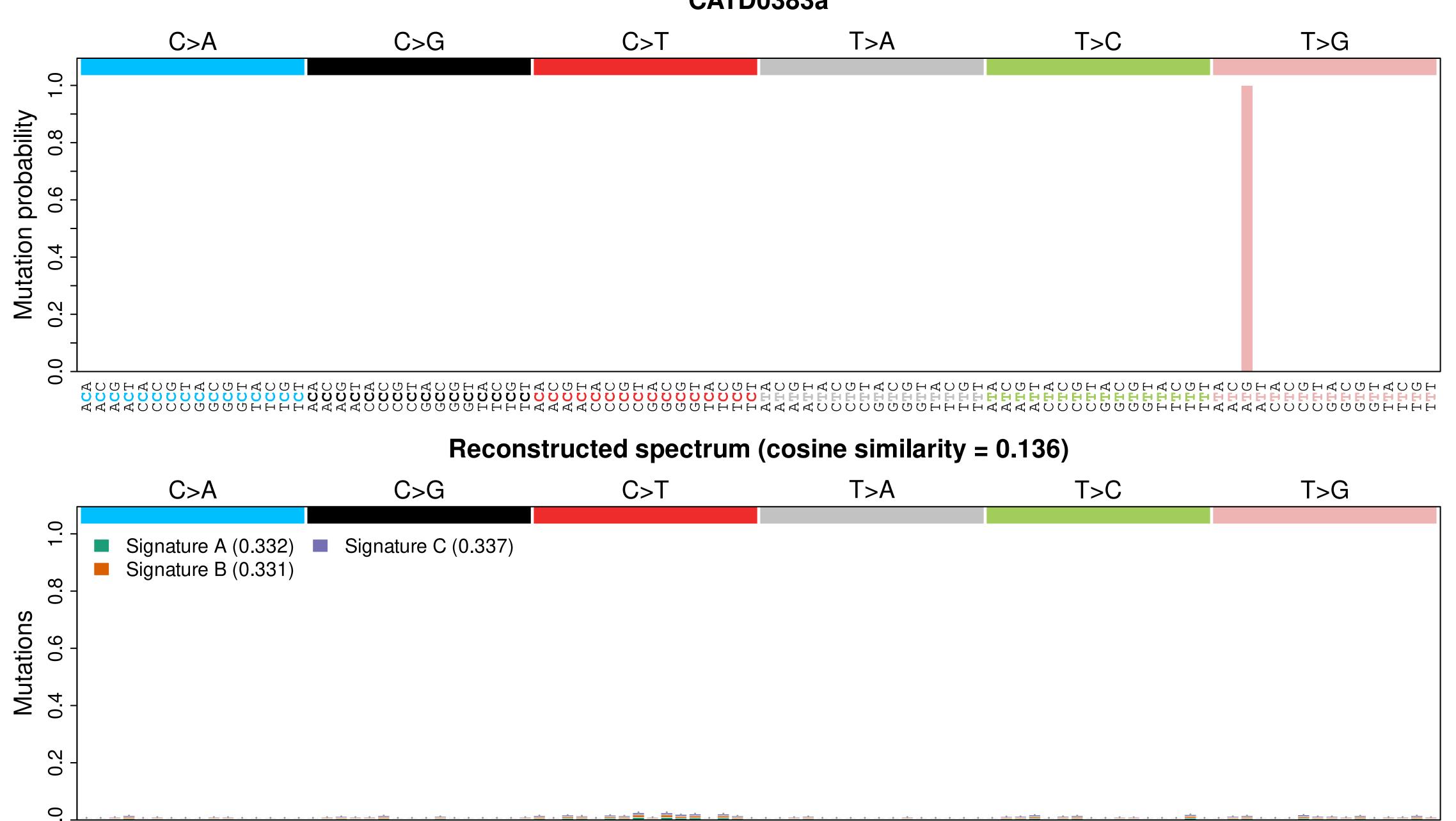




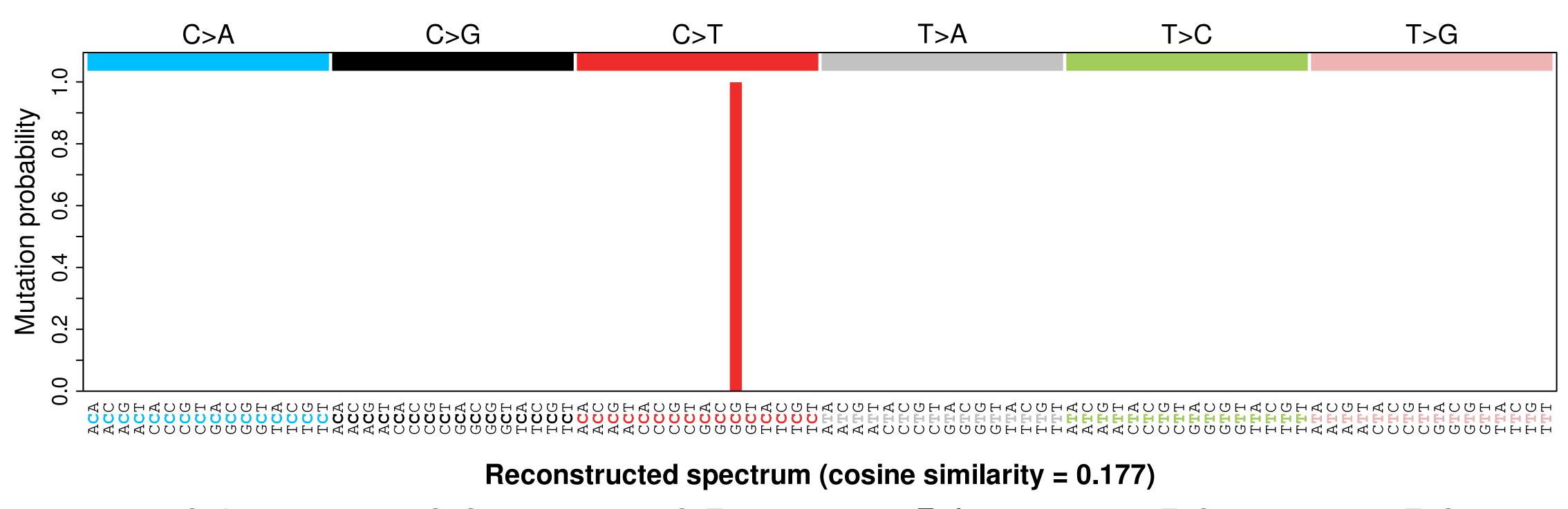


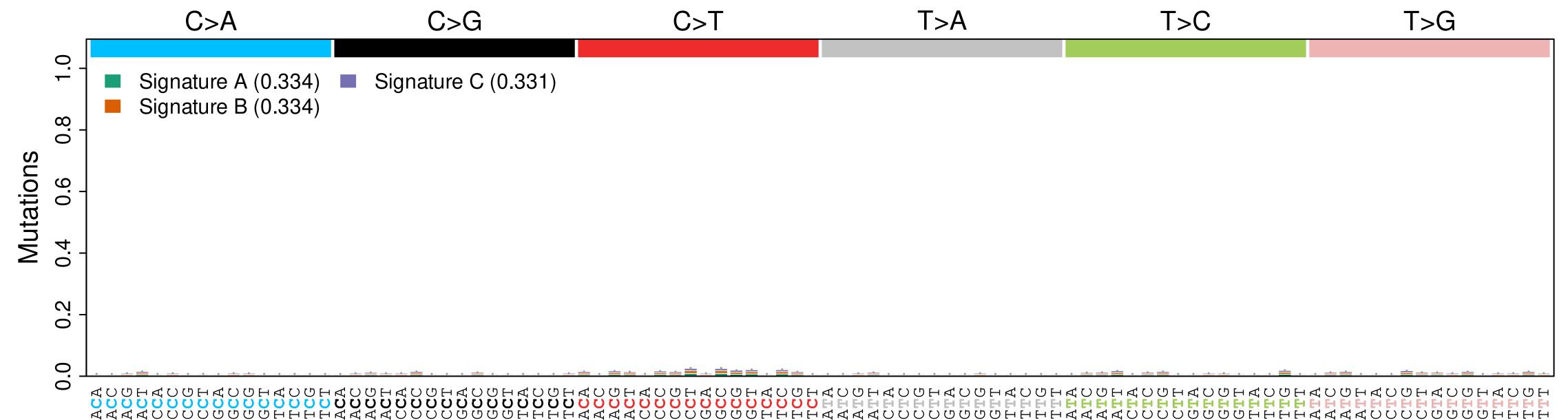




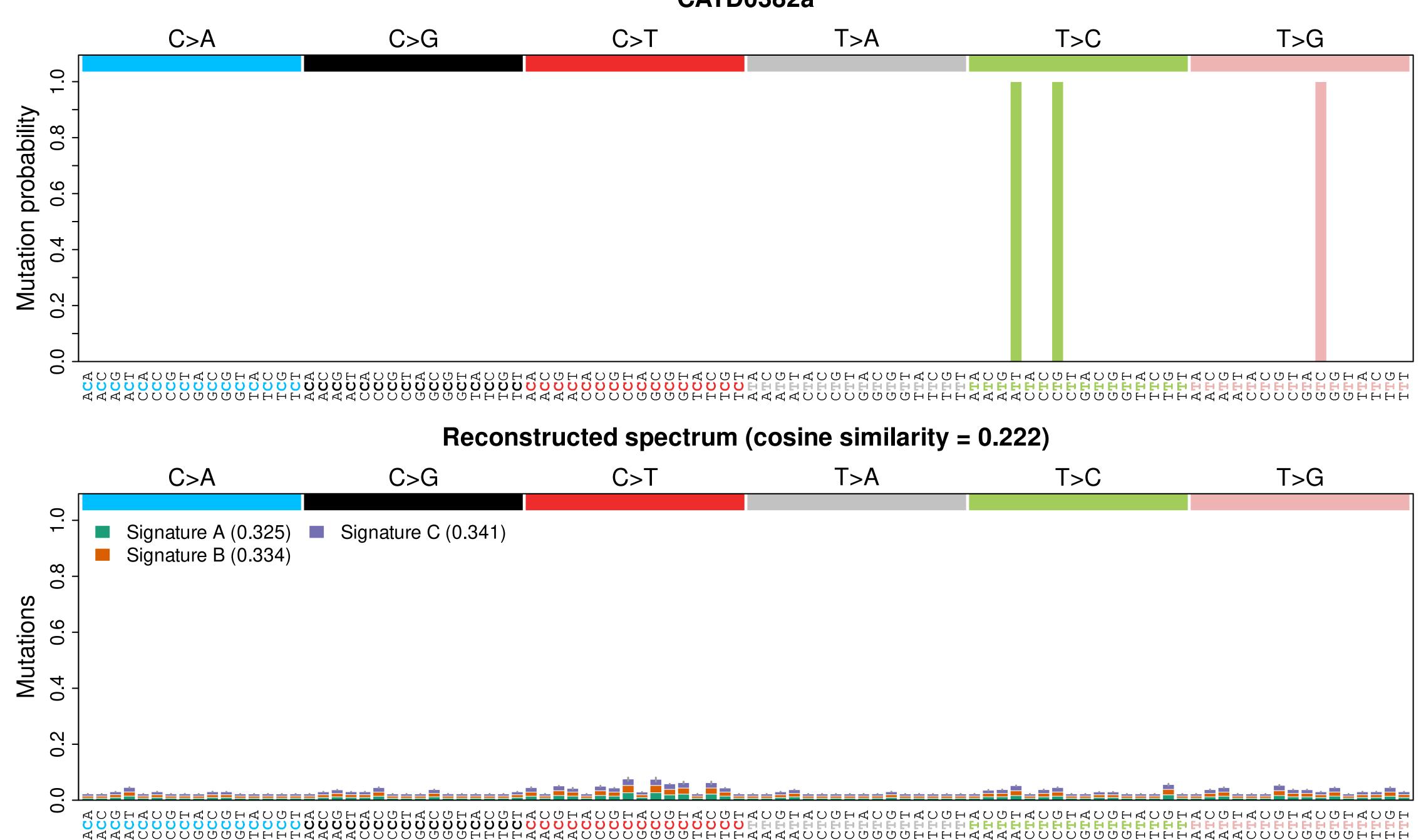




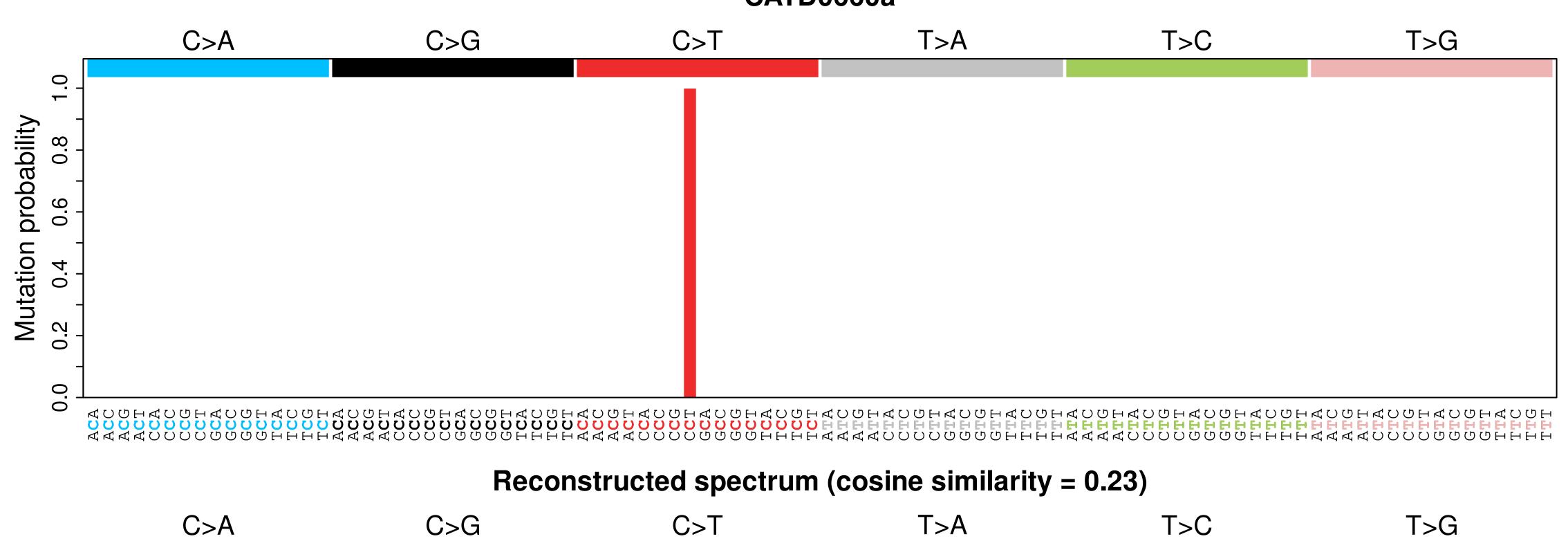


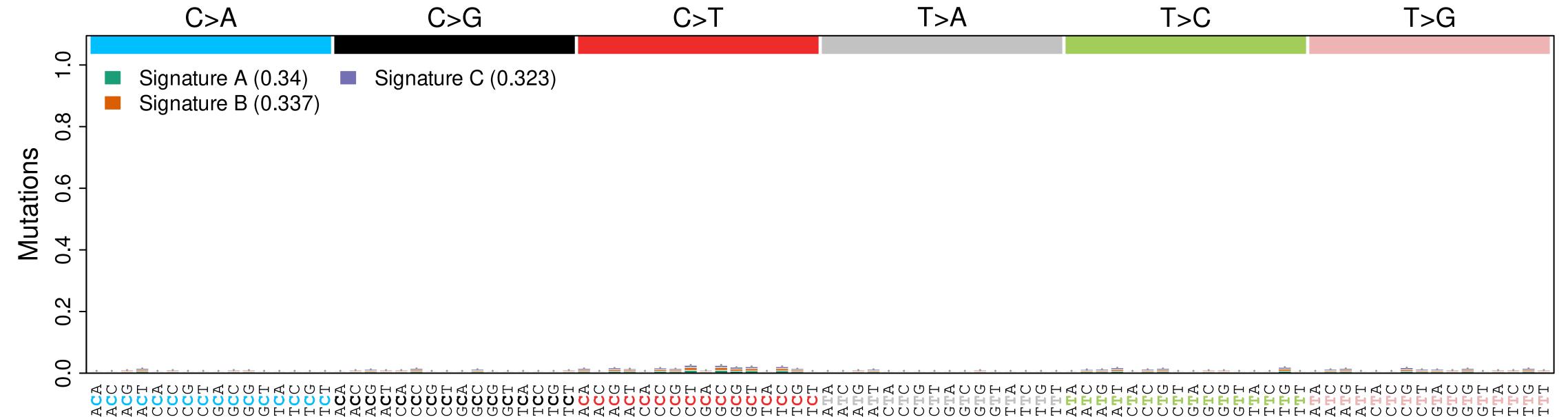




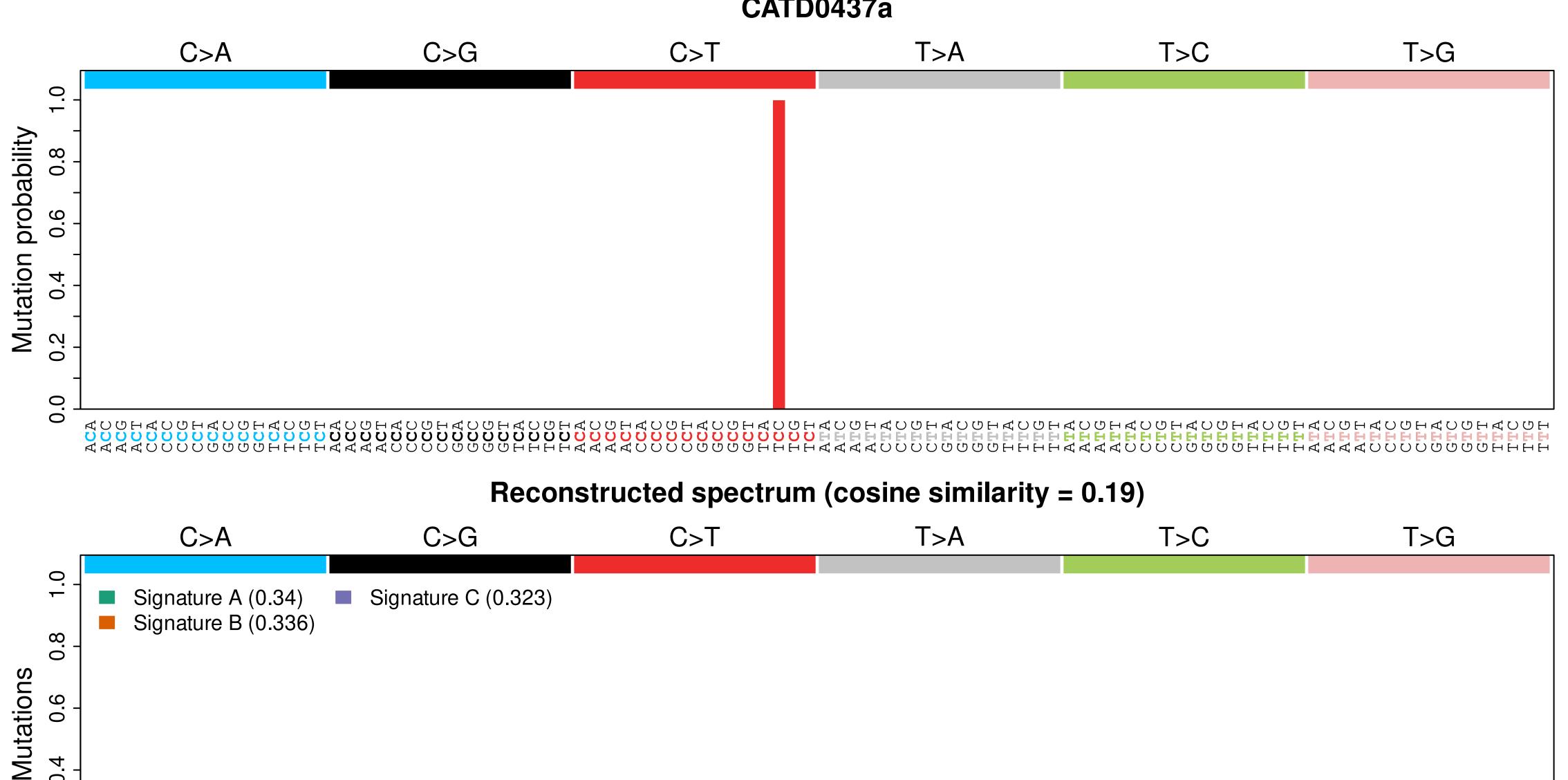












9.0

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