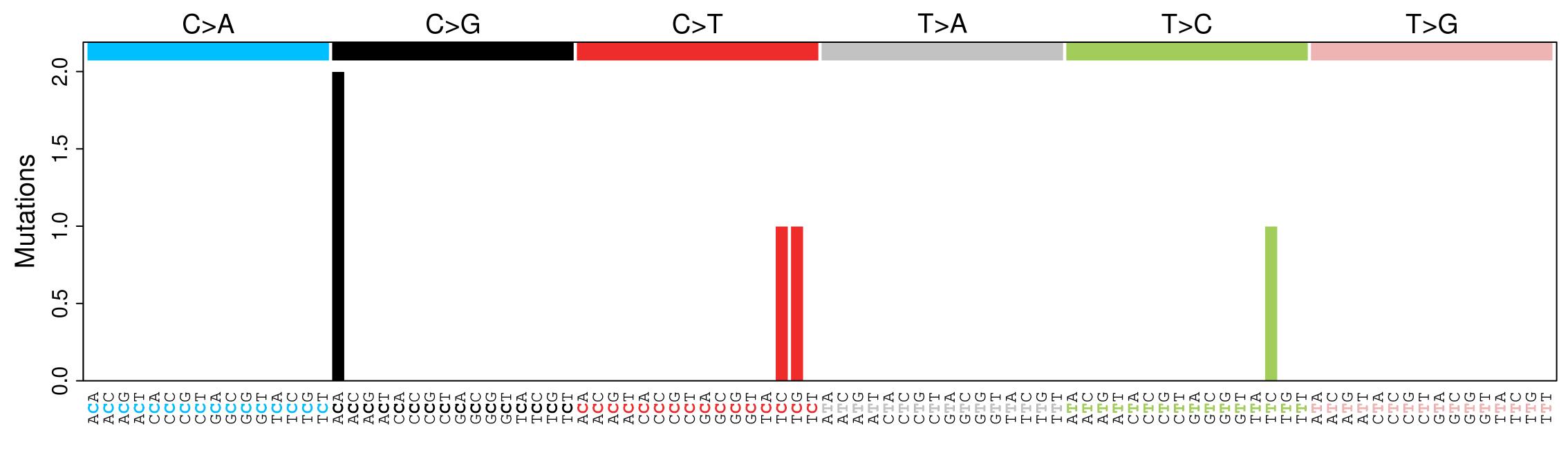
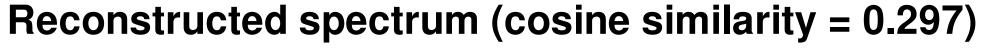


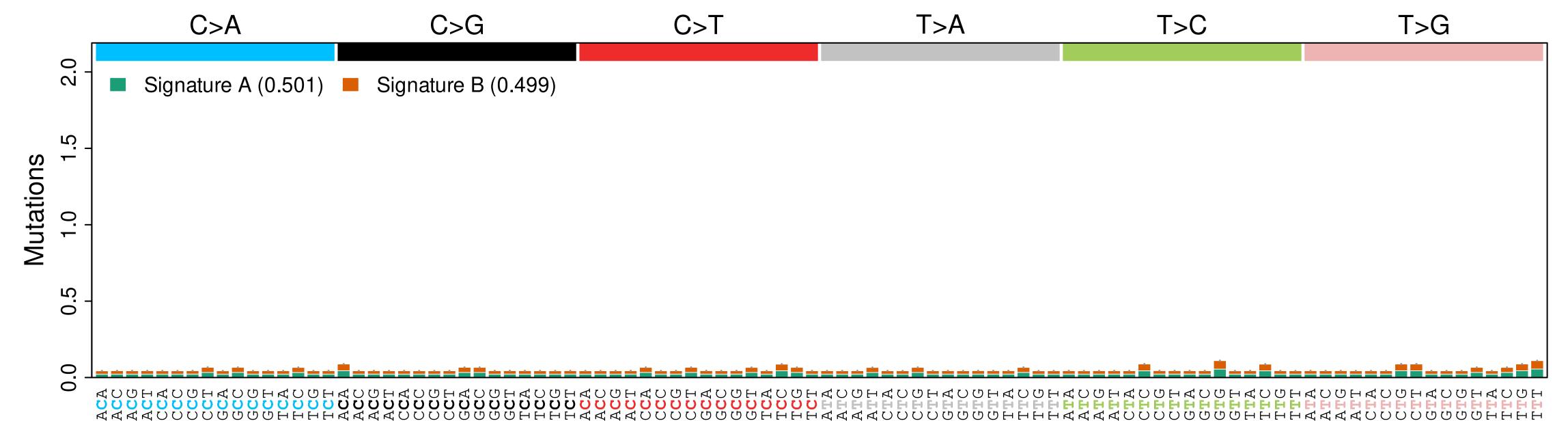
CATD0716a C>G T>C T>G C>A C>T T>A 0 Mutation probability 0.8 9.0 0.4 0.2 0.0 **Reconstructed spectrum (cosine similarity = 0.235)** C>A C>G T>A T>C T>G C>T 1.0 Signature A (0.501) Signature B (0.499) 0.8 Mutations 9.0 \mathcal{O}

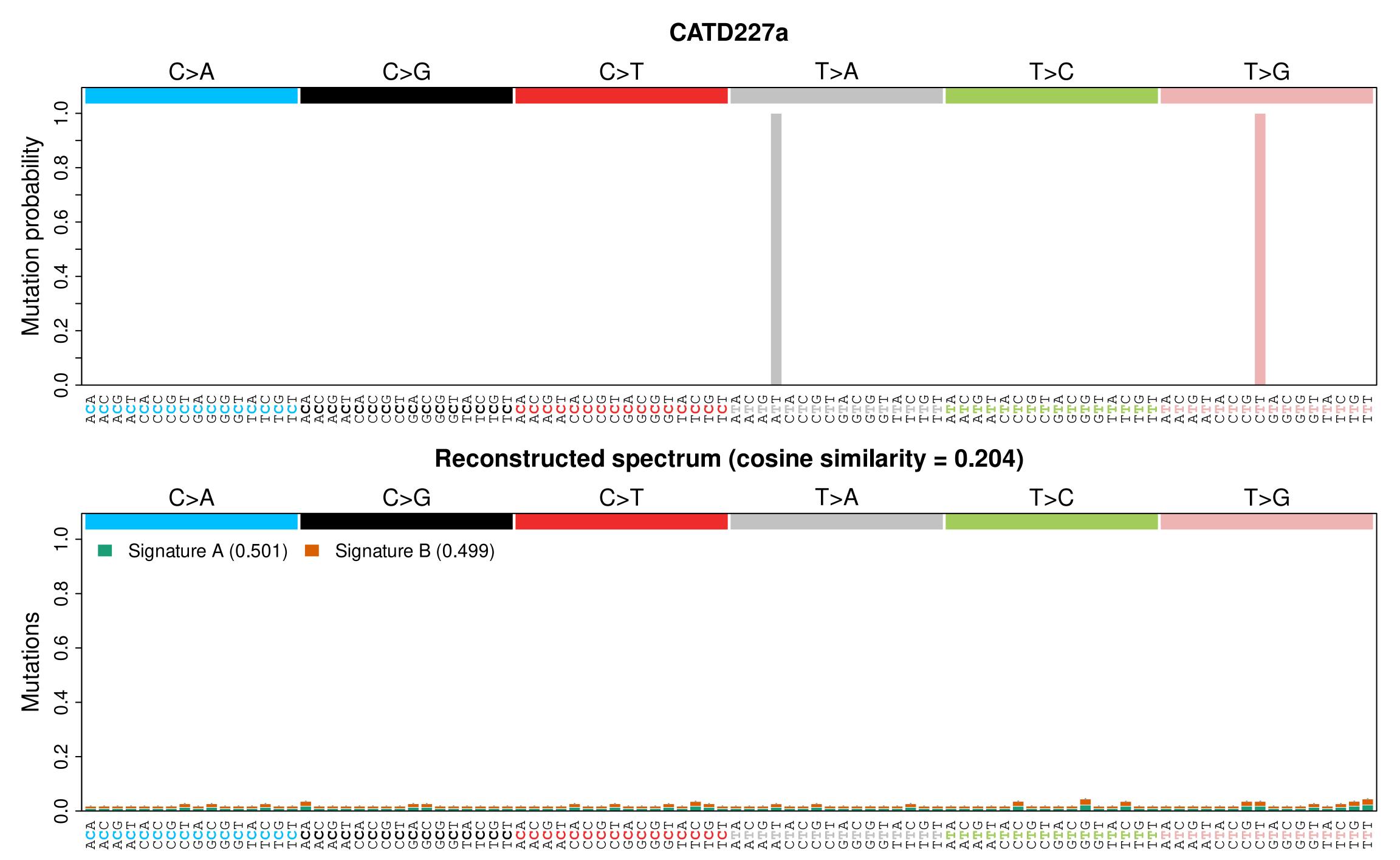
0

CATD217a (5 mutations)

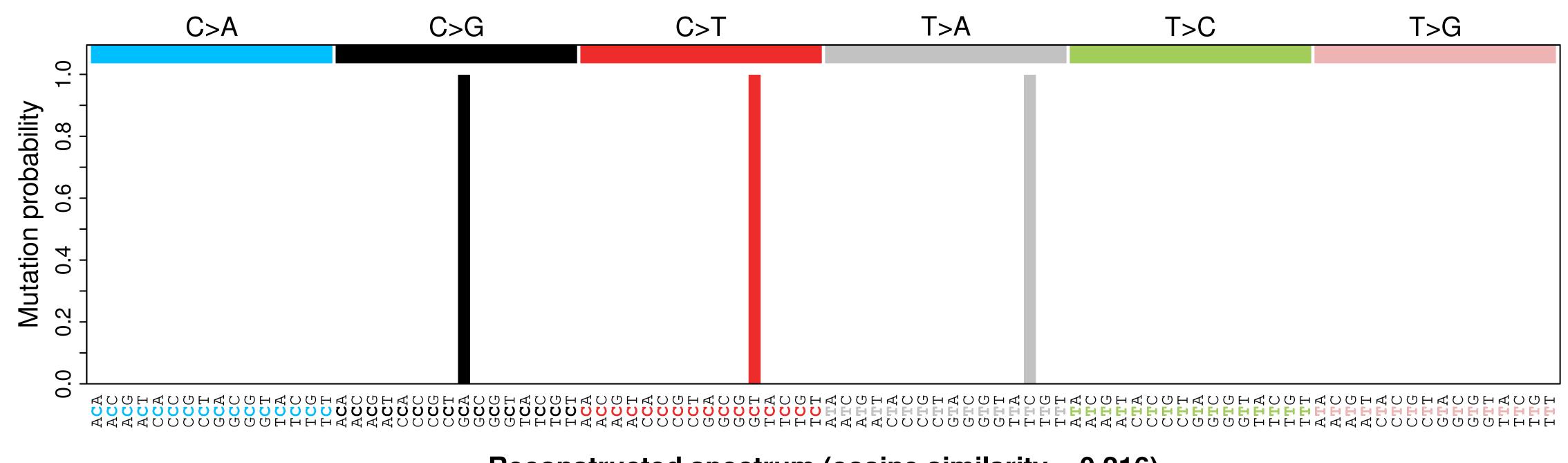


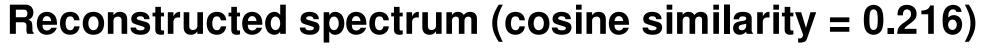


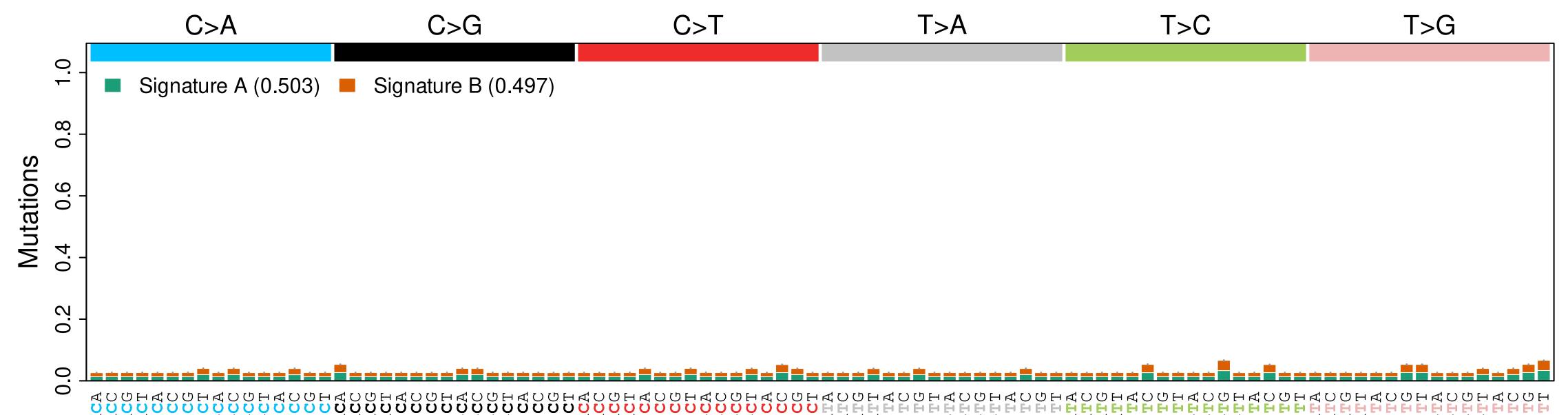




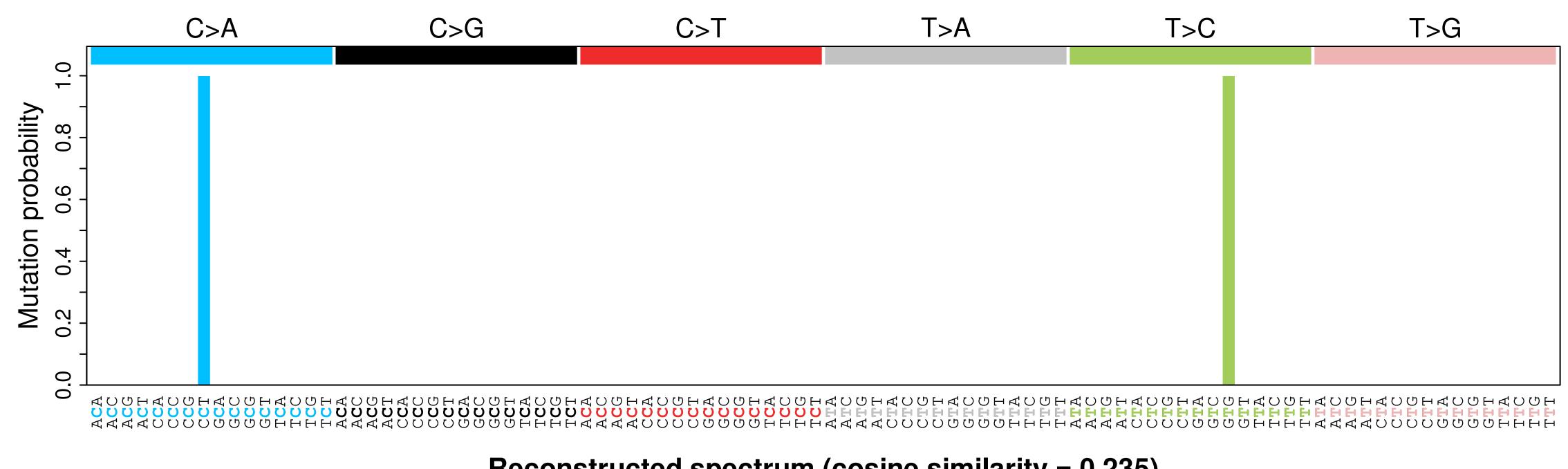




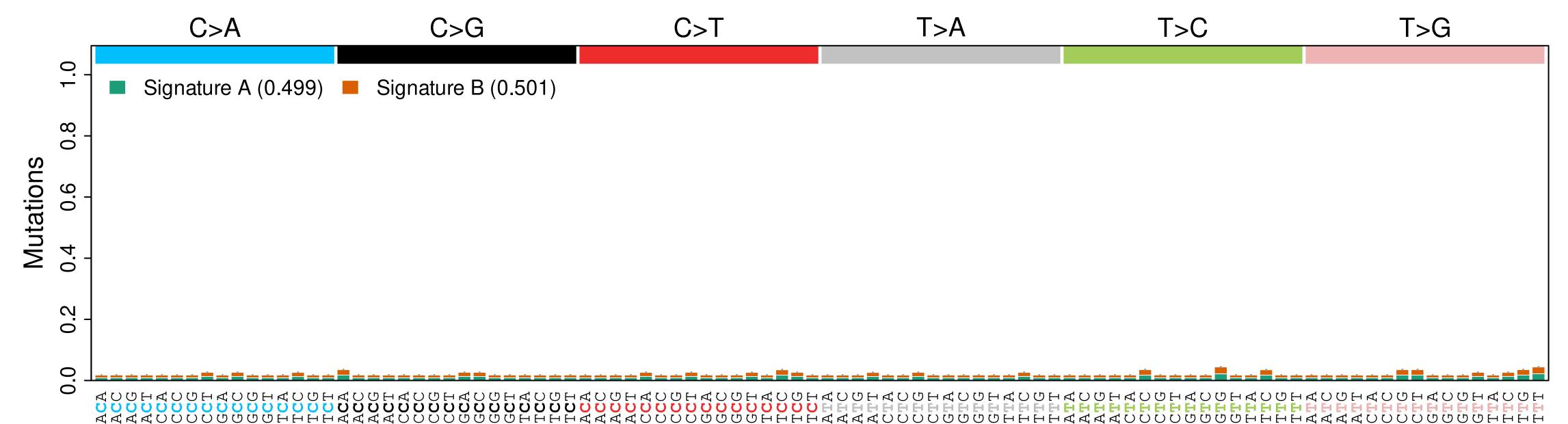


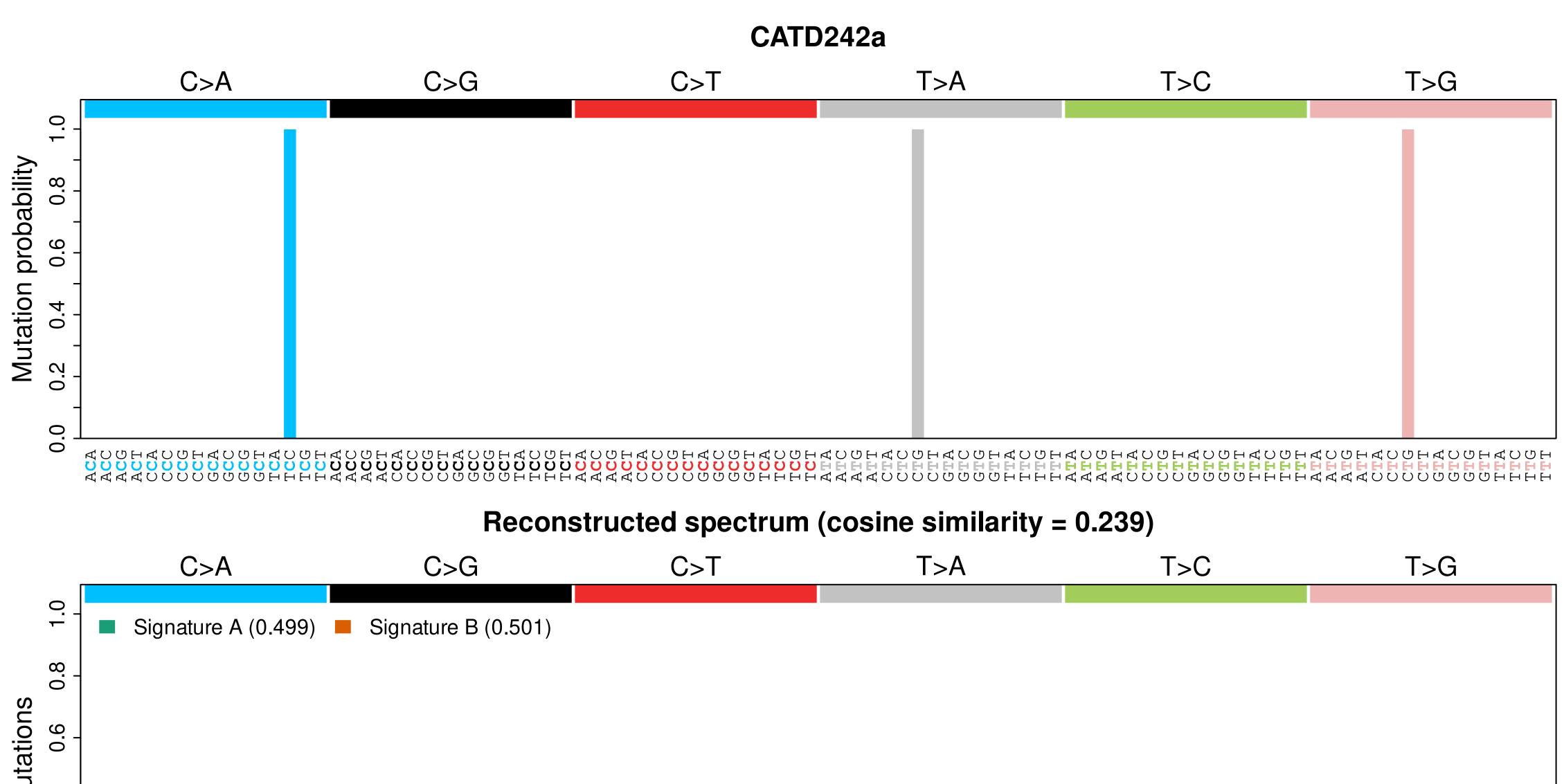


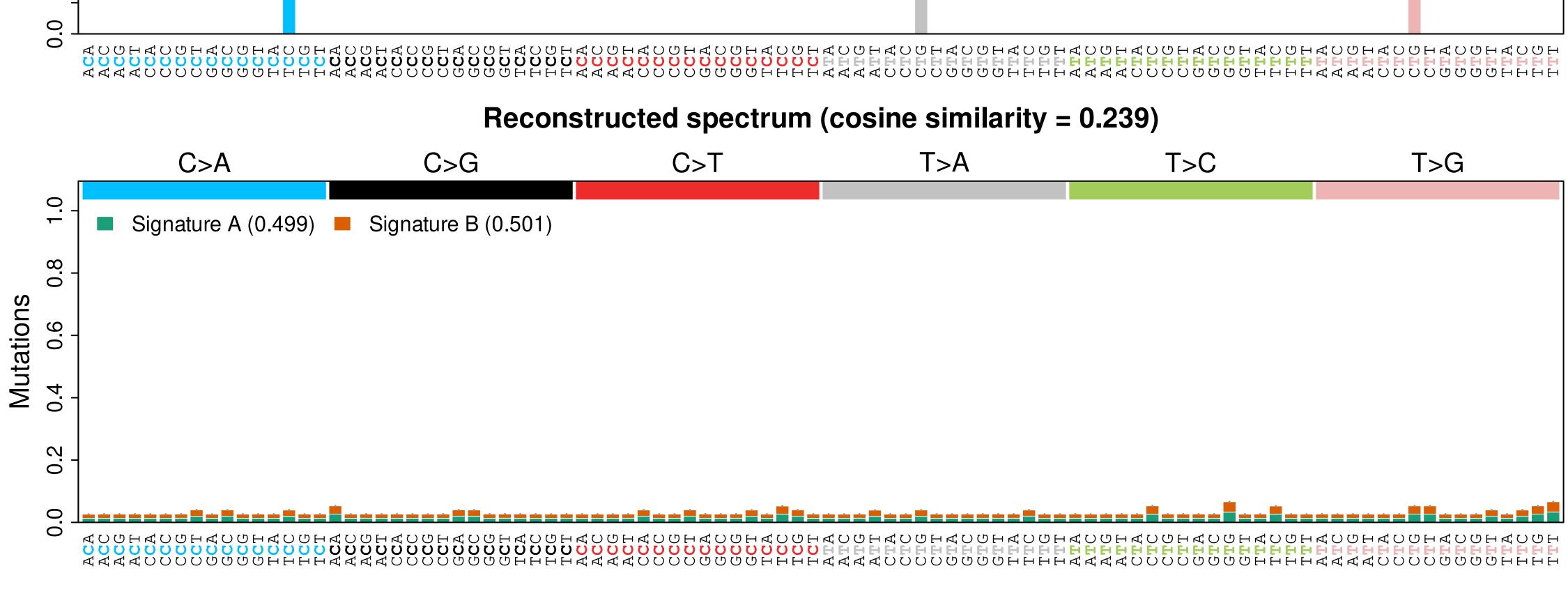




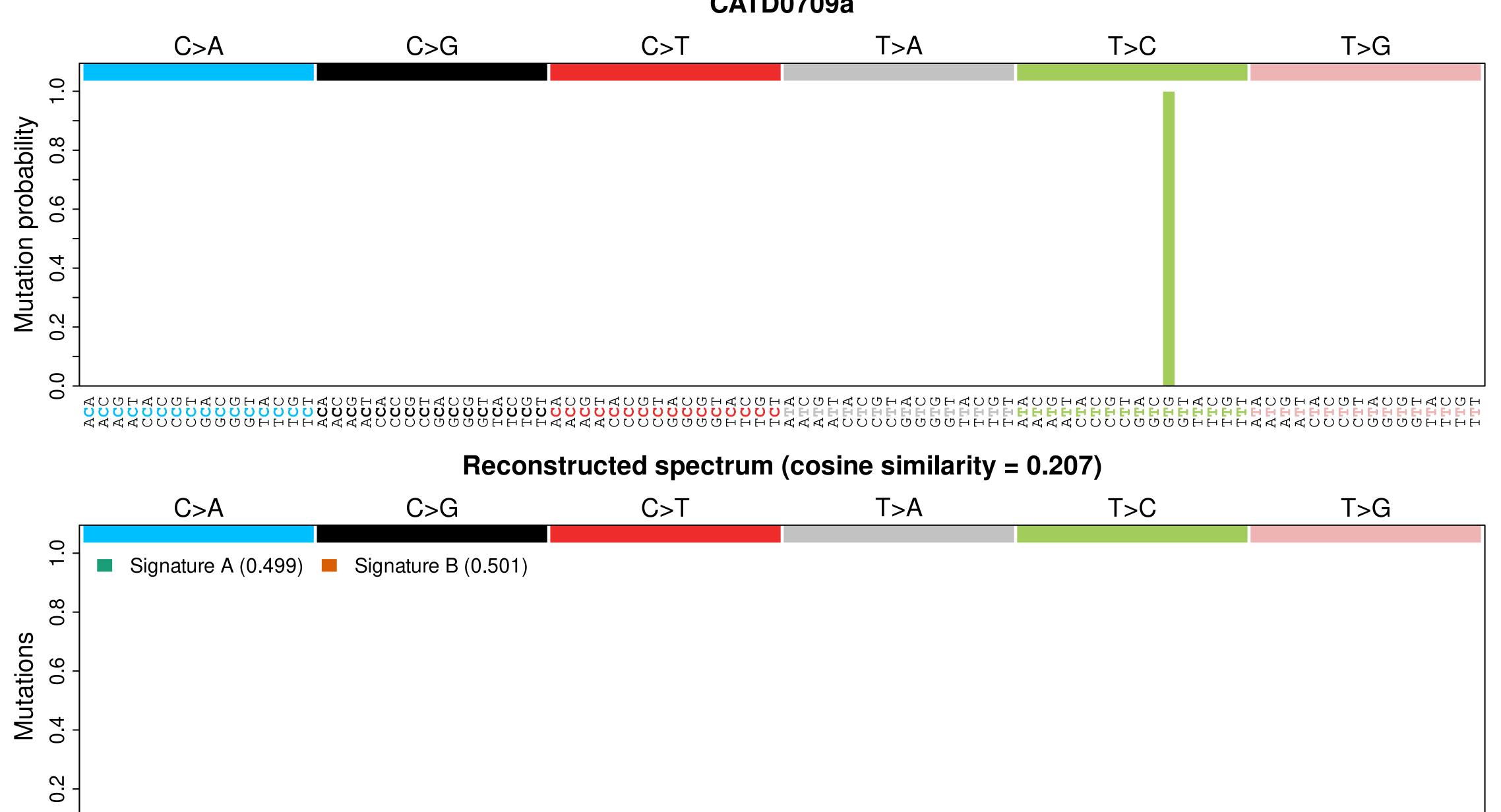
Reconstructed spectrum (cosine similarity = 0.235)

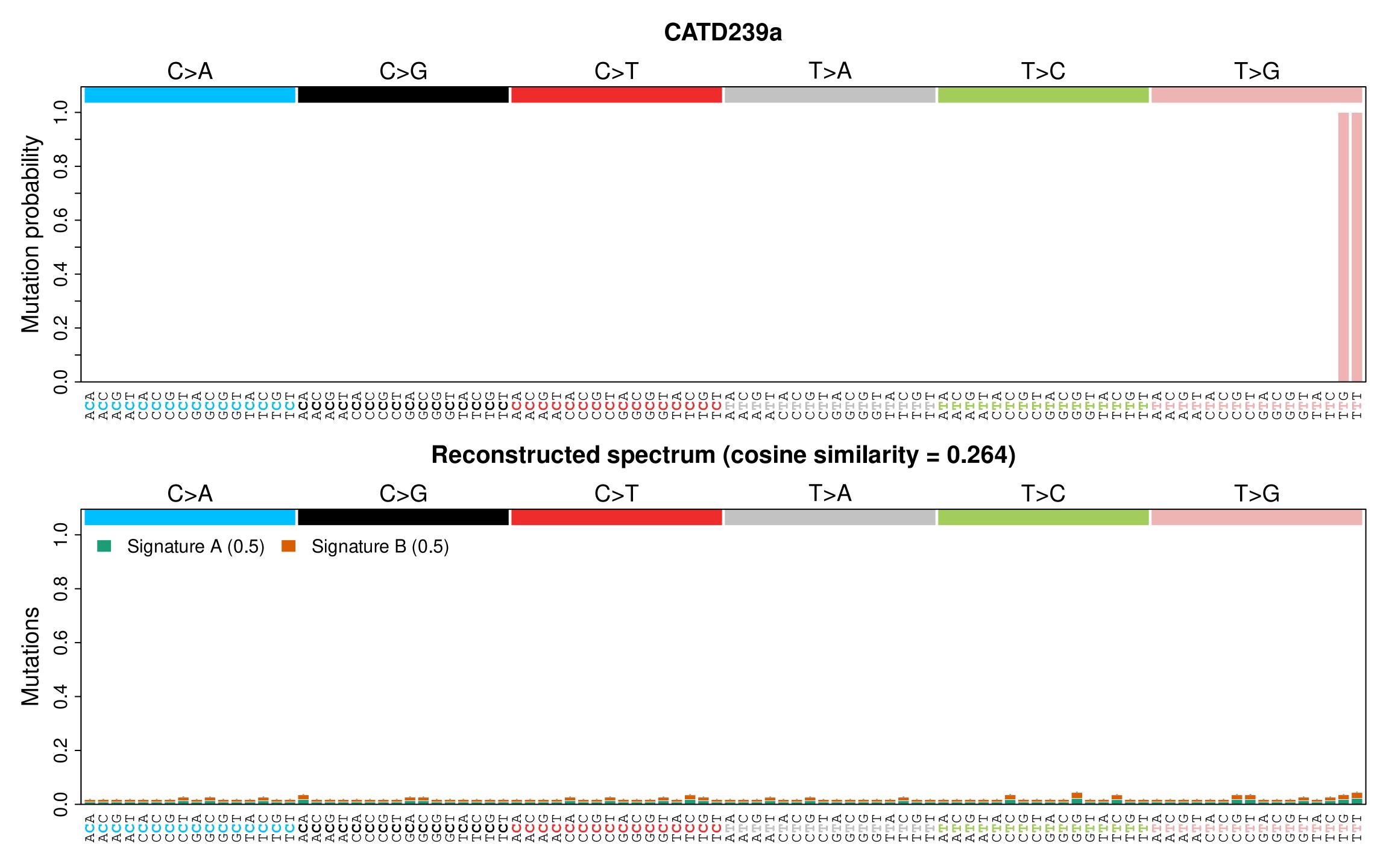


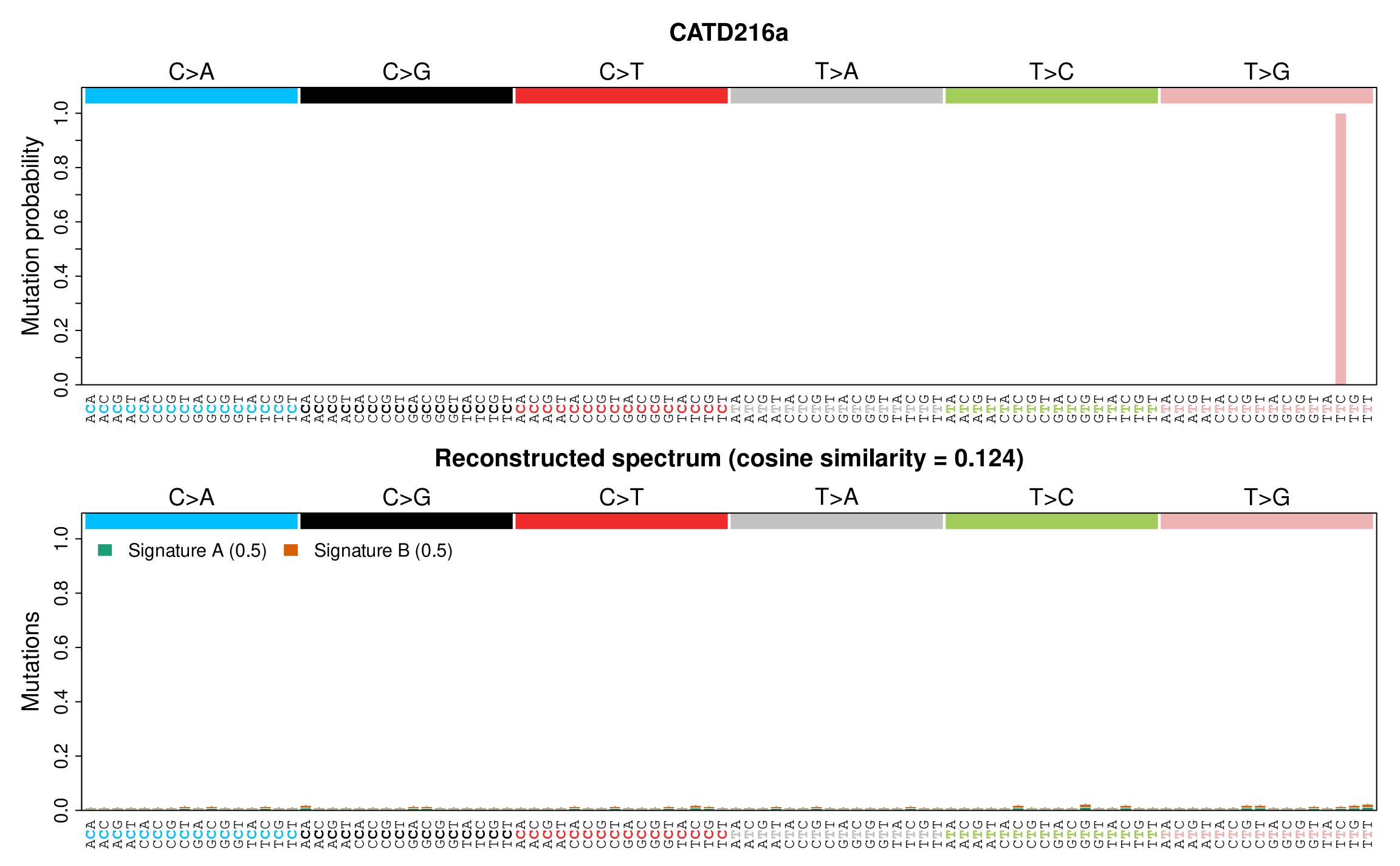


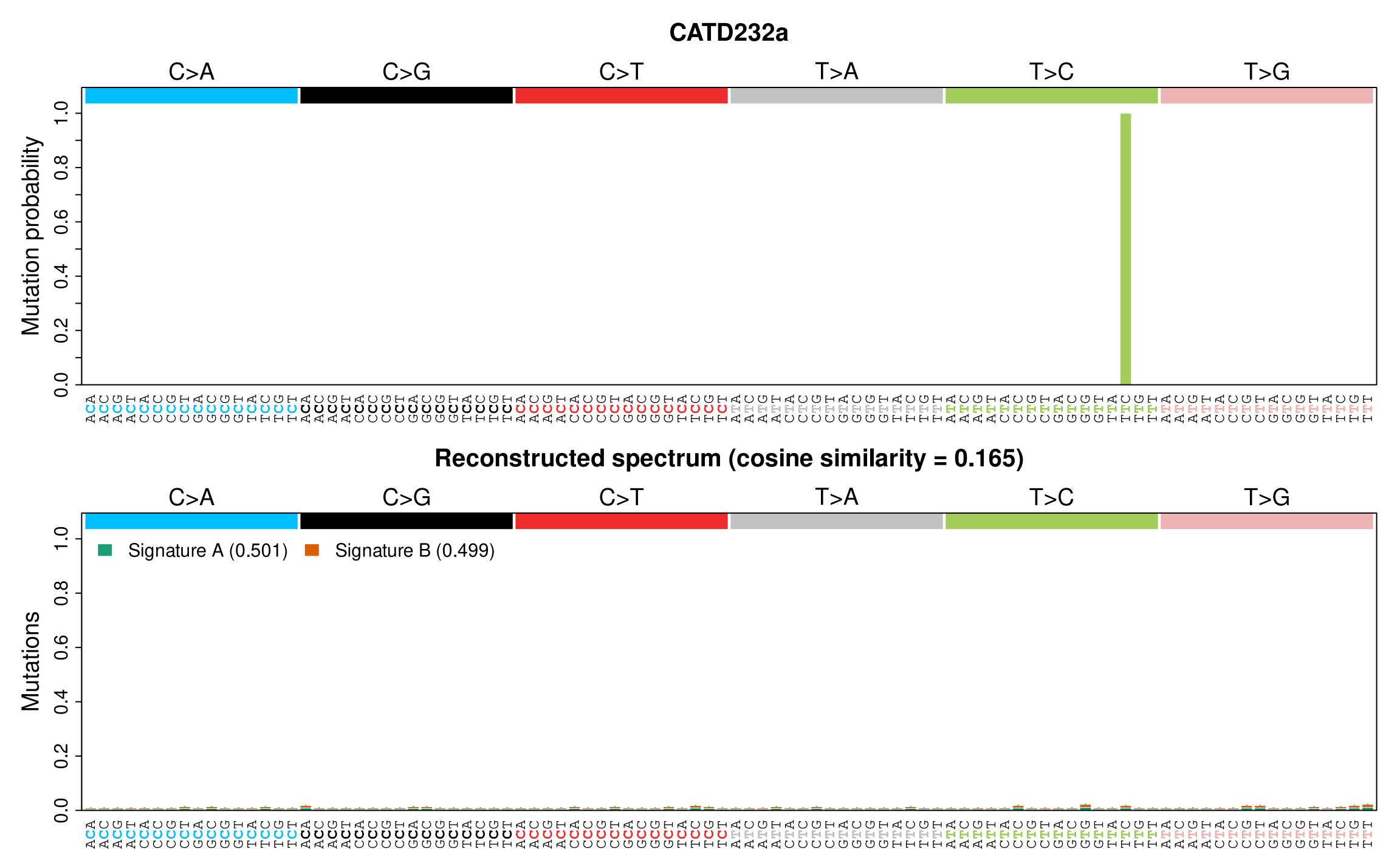






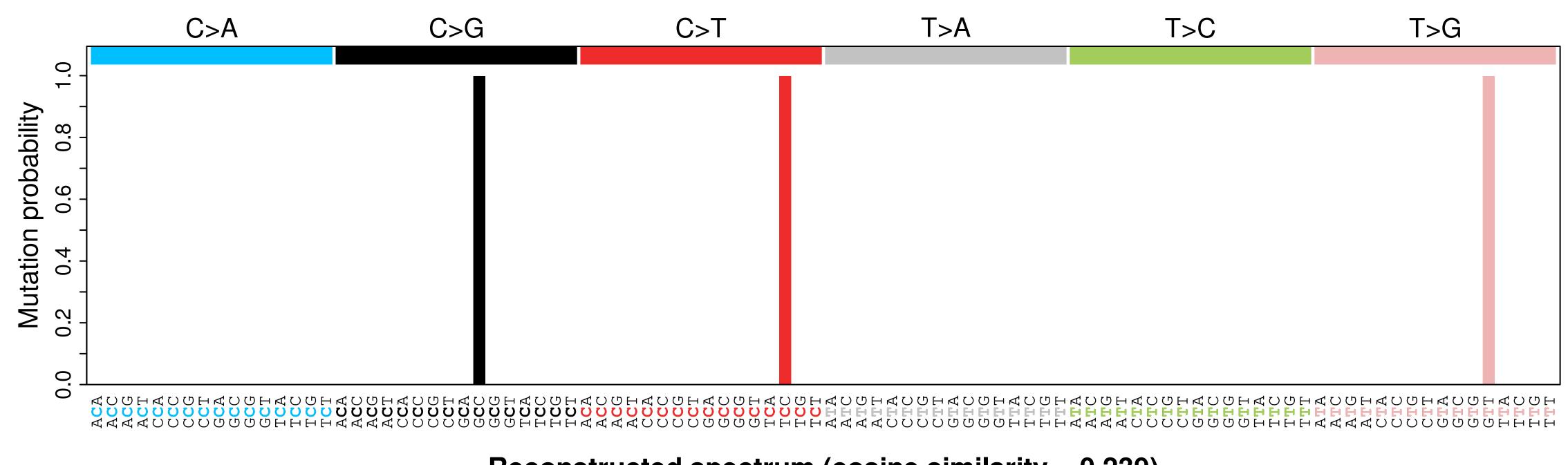


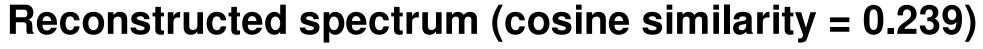


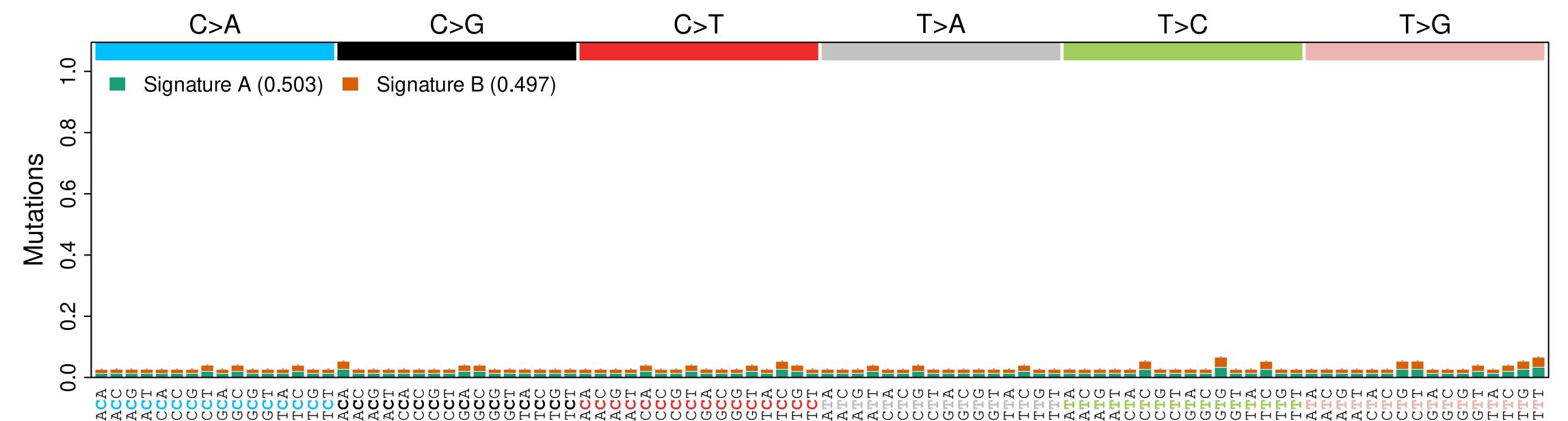


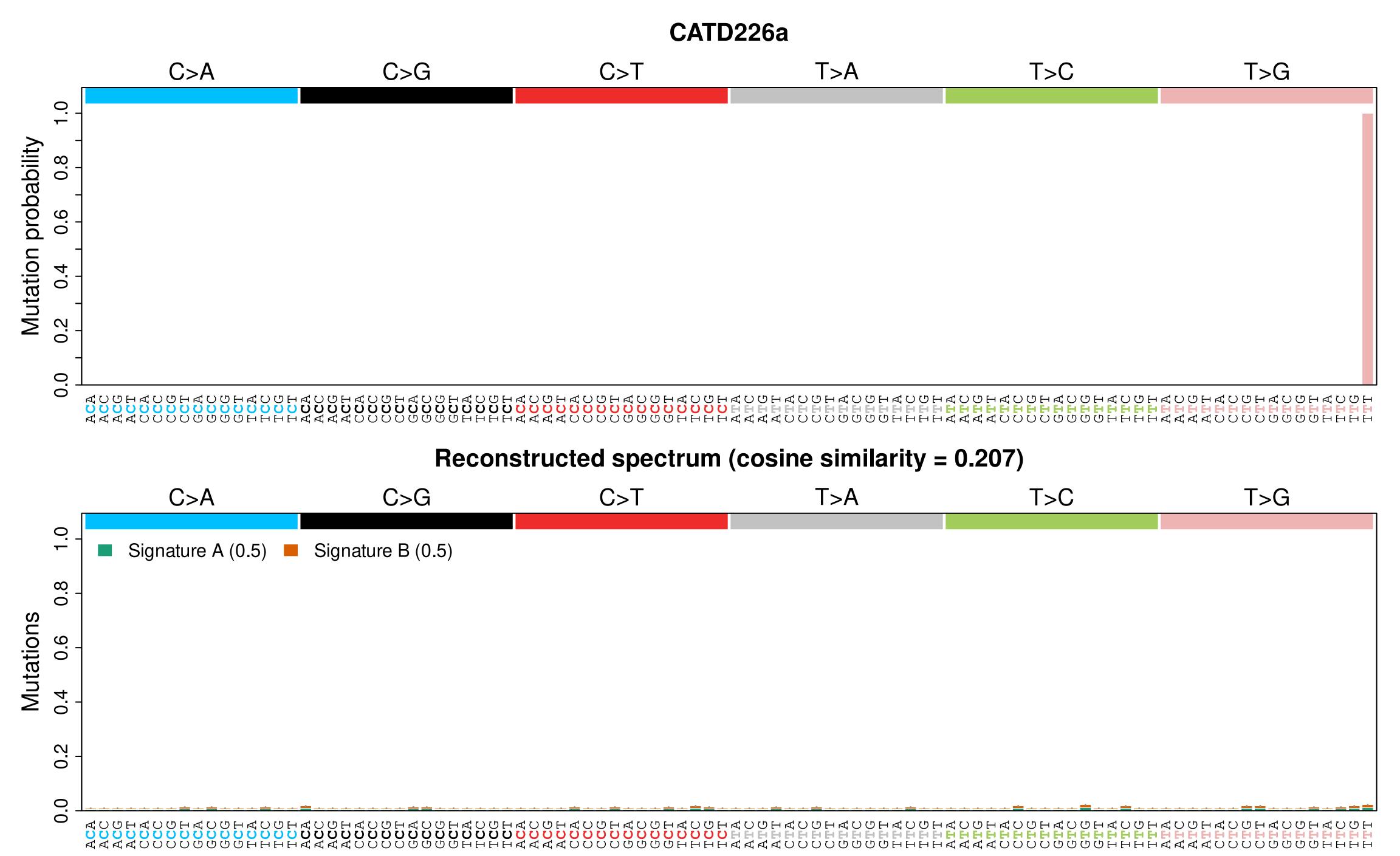
CATD229a T>C T>G C>A C>G C>T T>A 1.0 Mutation probability 0.8 9.0 0.4 0.0 **Reconstructed spectrum (cosine similarity = 0.311)** C>G T>A T>C T>G C>A C>T 1.0 Signature A (0.502) Signature B (0.498) 0.8 Mutations 9.0 \mathcal{O} 0



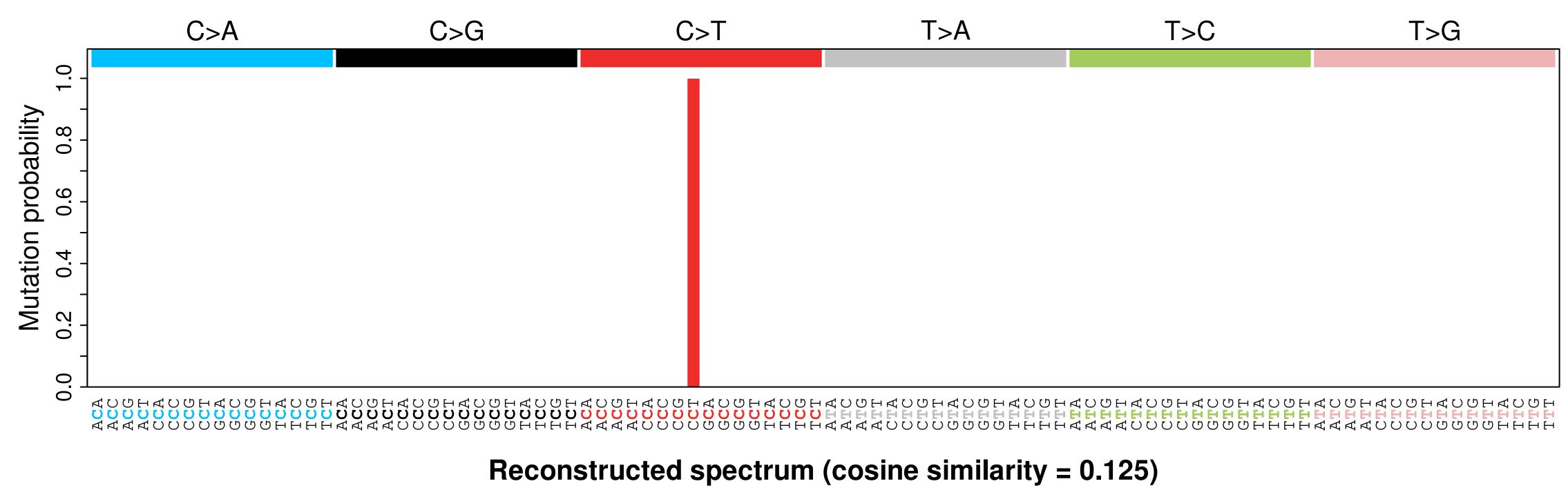


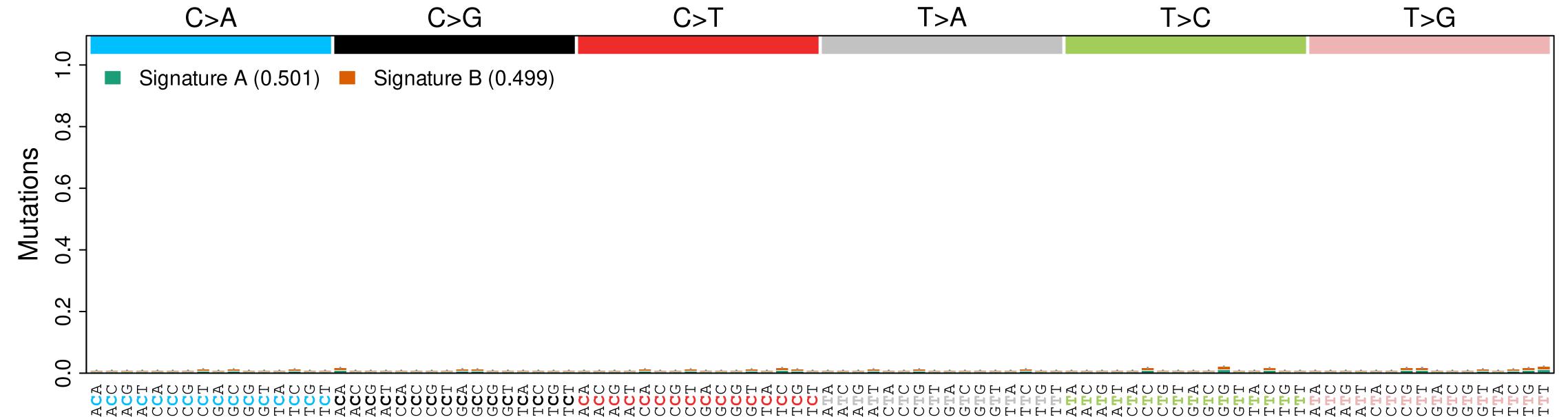












CATD224a (2 mutations) T>G C>A C>G C>T T>A T>C 2.0 5 Mutations 0.5 0.0 **Reconstructed spectrum (cosine similarity = 0.166)** C>G T>A T>C T>G C>A C>T 2.0 Signature A (0.496) Signature B (0.504) 5 Mutations 5 0

CATD0561e C>G T>C T>G C>A C>T T>A 1.0 Mutation probability 0.8 9.0 0.4 0.2 0.0 **Reconstructed spectrum (cosine similarity = 0.166)** C>A T>G C>G T>A T>C C>T 1.0 Signature A (0.5) Signature B (0.5) 0.8 Mutations 9.0

0.2

0.0