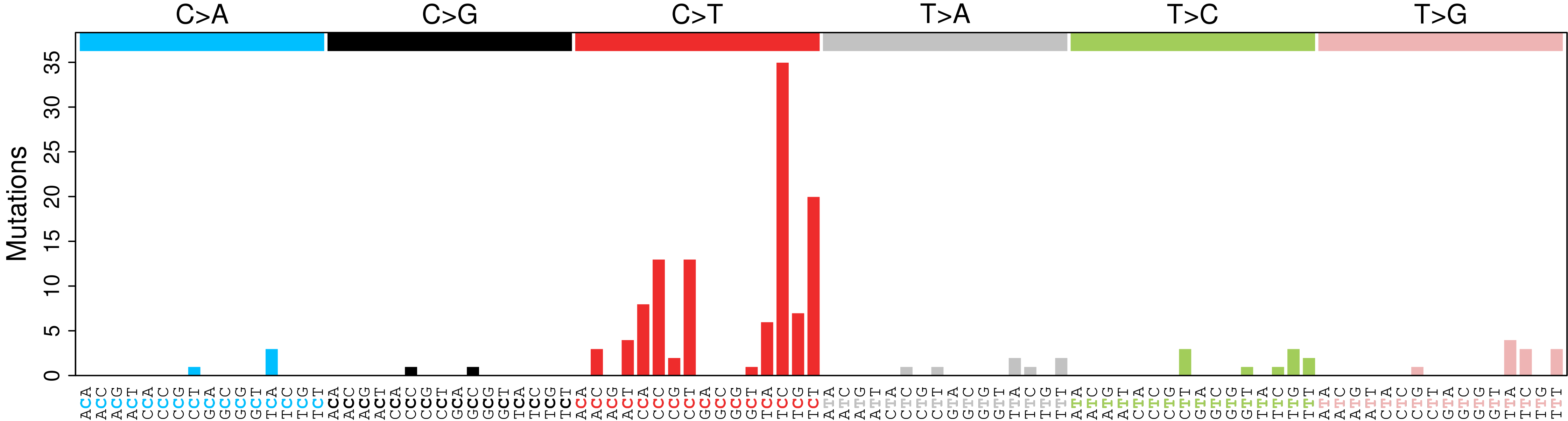CATD265a (146 mutations)

Reconstructed spectrum (cosine similarity = 0.906)

SBS1 (0.006)   SBS36 (0.006)
SBS2 (0.026)   SBS37 (0.007)
SBS3 (0.007)   SBS38 (0.005)
SBS4 (0.005)   SBS39 (0.006)
SBS5 (0.007)   SBS40a (0.008)
SBS6 (0.006)   SBS40b (0.008)
SBS7a (0.147) SBS40c (0.008)
SBS7b (0.205) SBS41 (0.011)
SBS7c (0.014) SBS42 (0.006)
SBS7d (0.008) SBS43 (0.005)
SBS8 (0.006)   SBS44 (0.006)
SBS9 (0.01)   SBS45 (0.005)