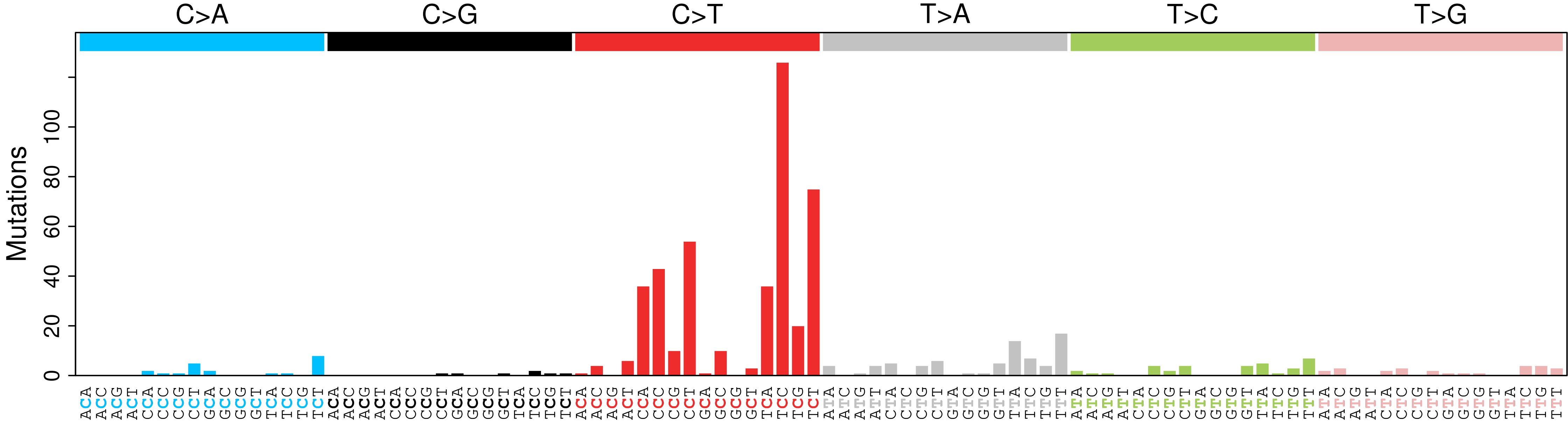**CATD0119a (586 mutations)**

Reconstructed spectrum (cosine similarity = 0.967)

Legend:
SBS1 (0.002)
SBS2 (0.036)
SBS3 (0.004)
SBS4 (0.003)
SBS5 (0.004)
SBS6 (0.003)
SBS7a (0.214)
SBS7b (0.337)
SBS7c (0.051)
SBS7d (0.005)
SBS8 (0.004)
SBS9 (0.005)
SBS36 (0.003)
SBS37 (0.004)
SBS38 (0.003)
SBS39 (0.003)
SBS40a (0.004)
SBS40b (0.006)
SBS40c (0.005)
SBS41 (0.006)
SBS42 (0.004)
SBS43 (0.004)
SBS44 (0.003)
SBS45 (0.003)