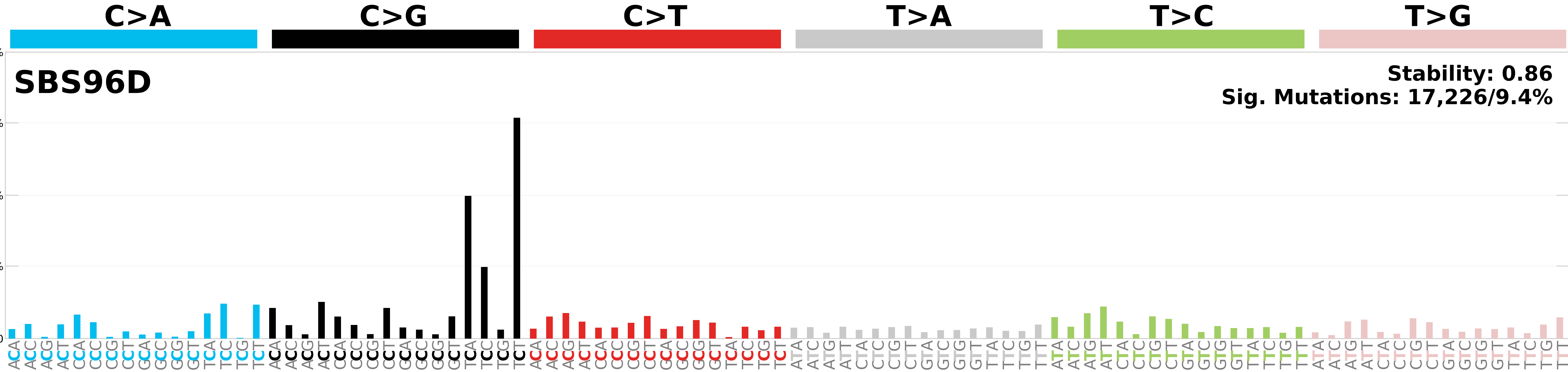
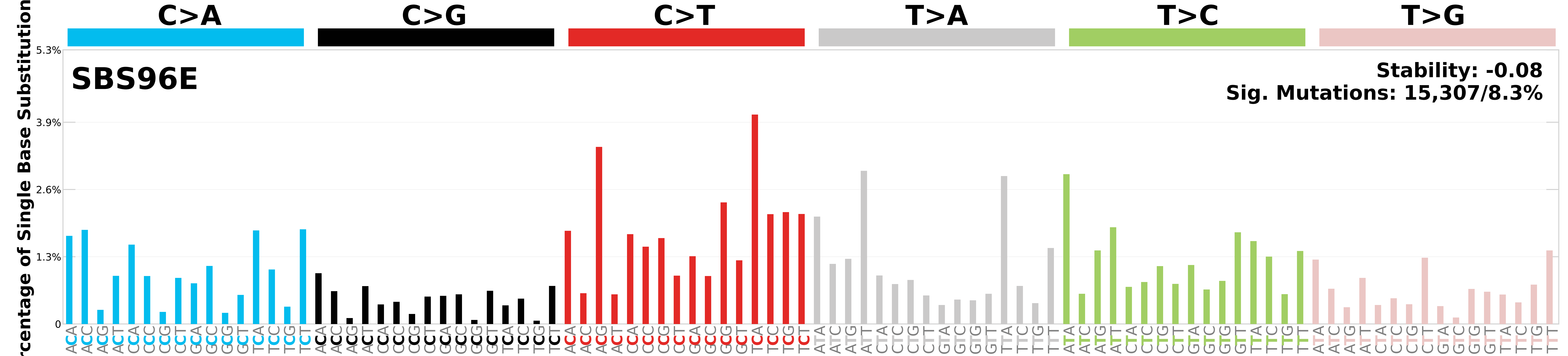


Percentage of Single Base Substitution





T > G

Stability: 0.16
Sig. Mutations: 14,914/8.1%

