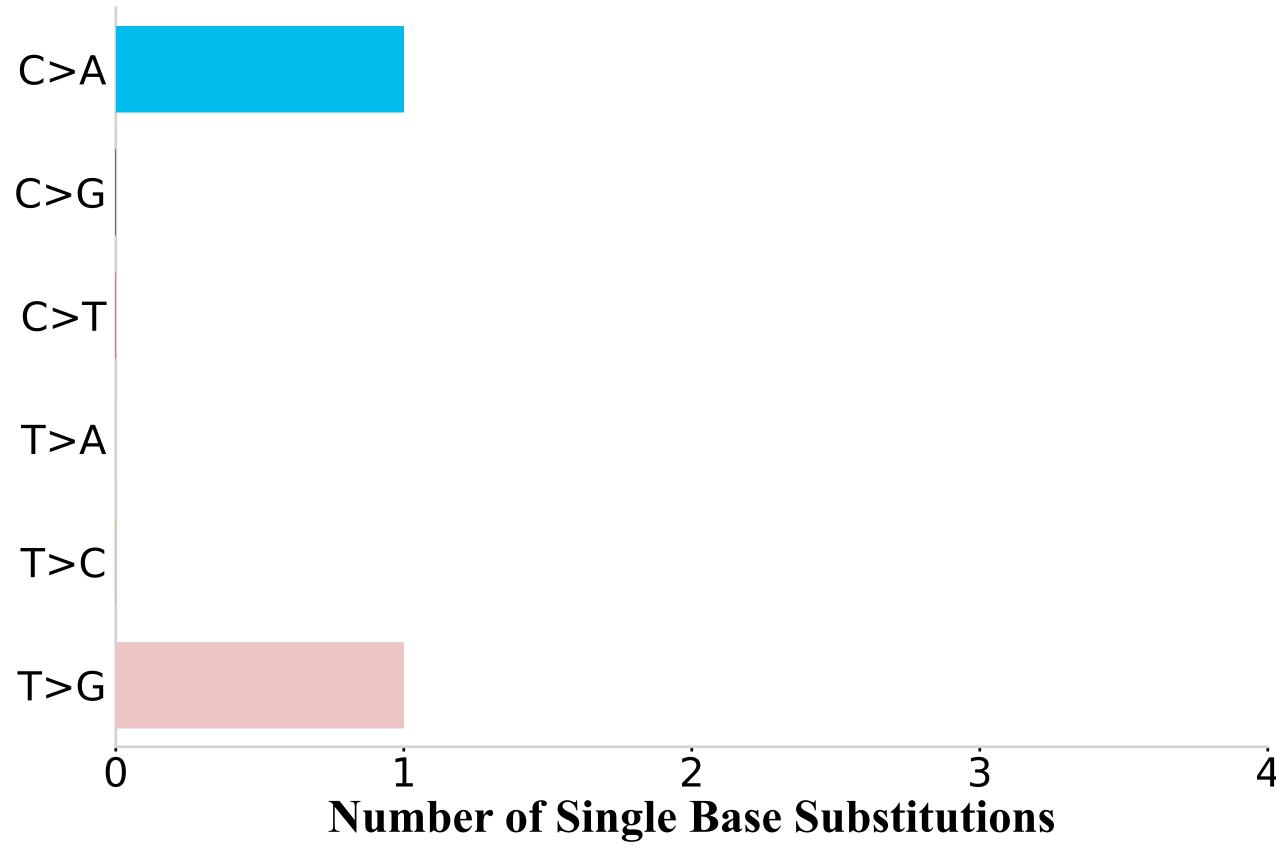
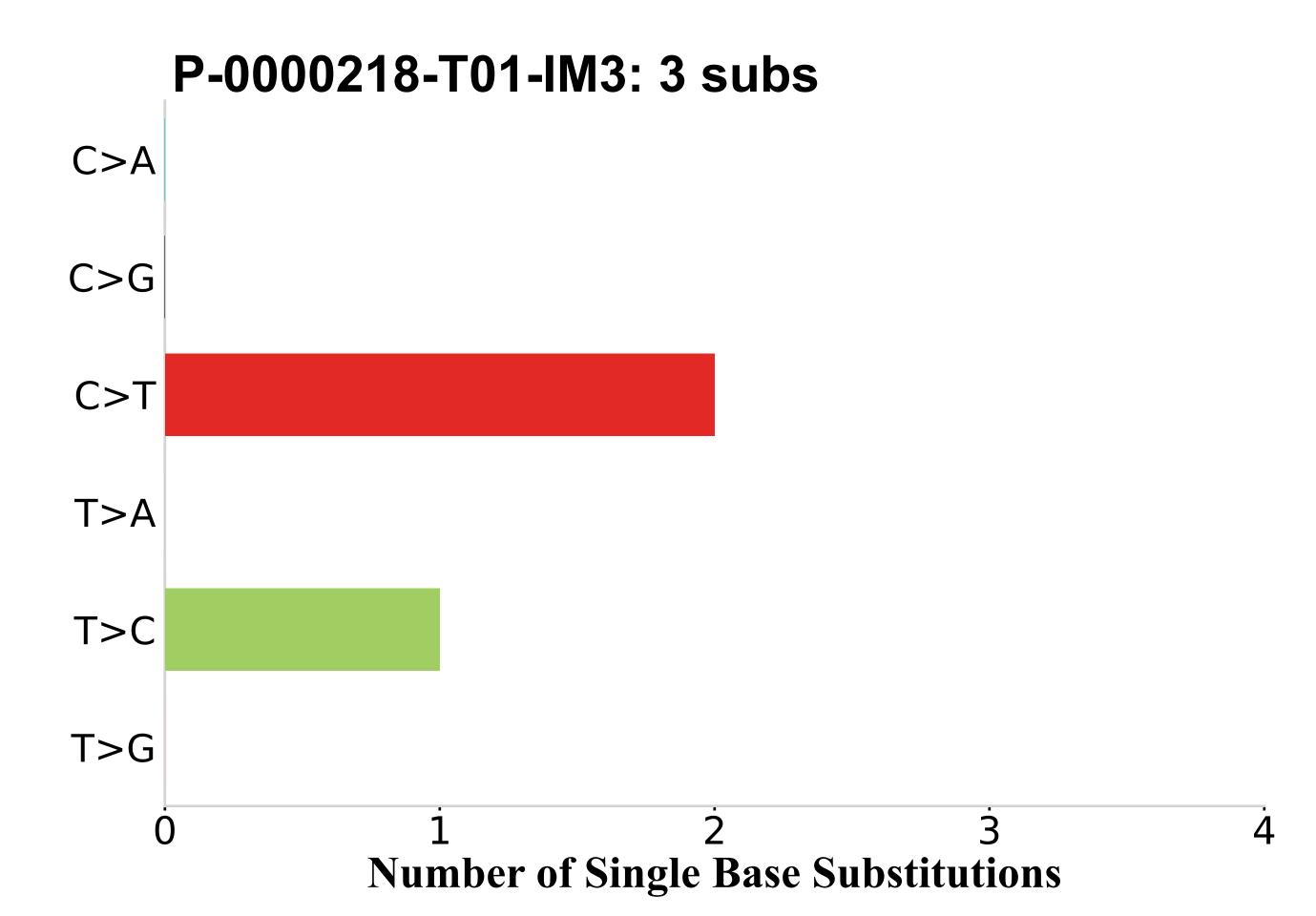
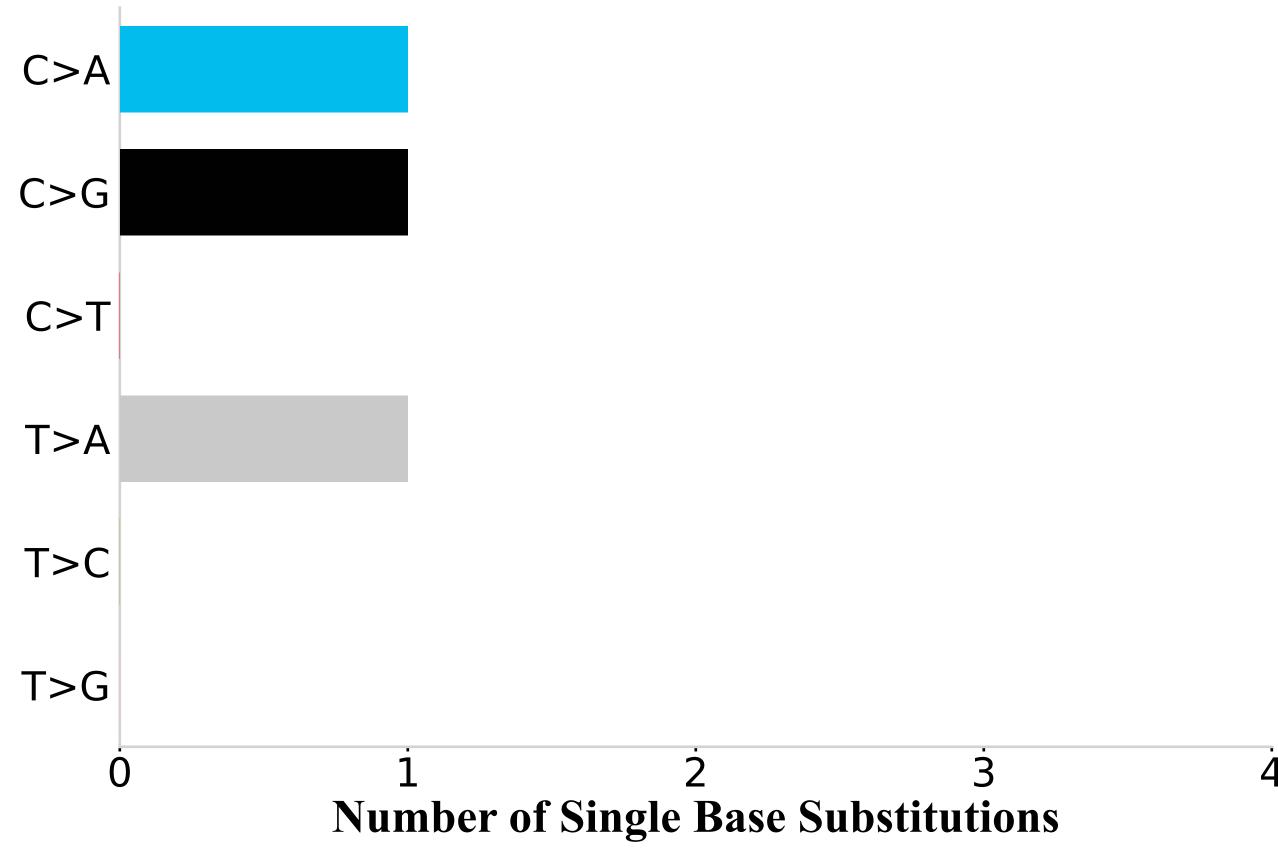


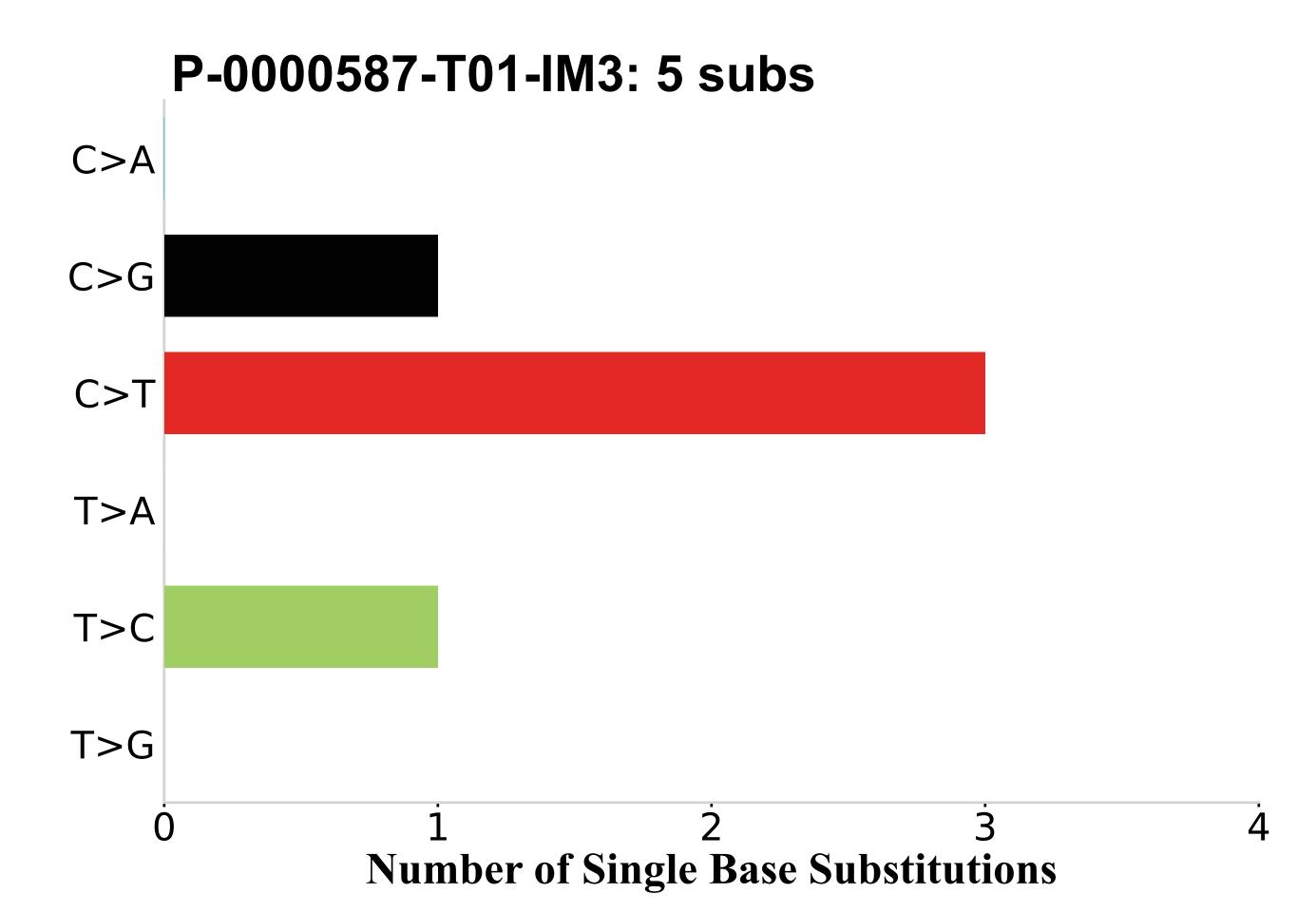
P-0000182-T01-IM3: 2 subs

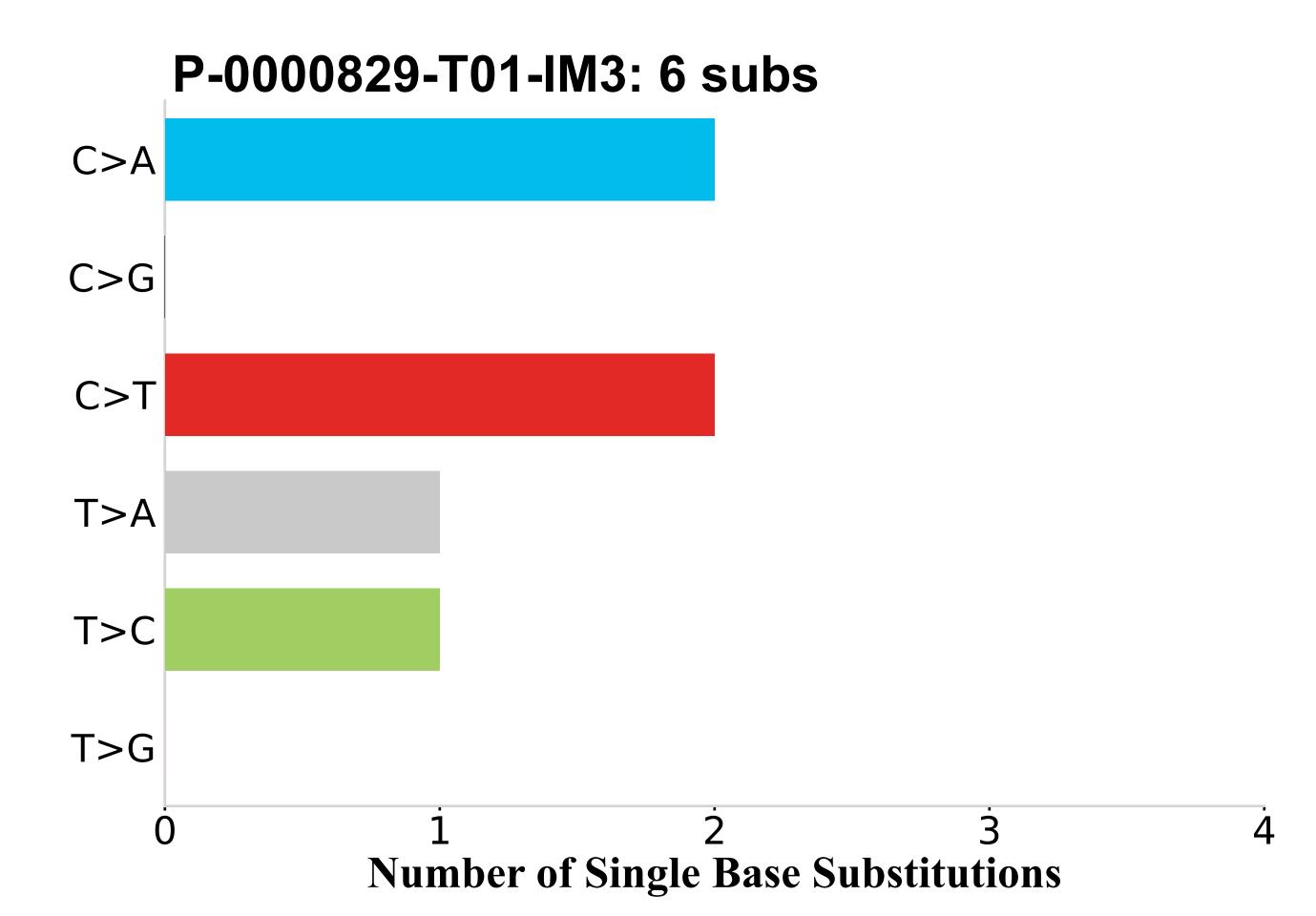


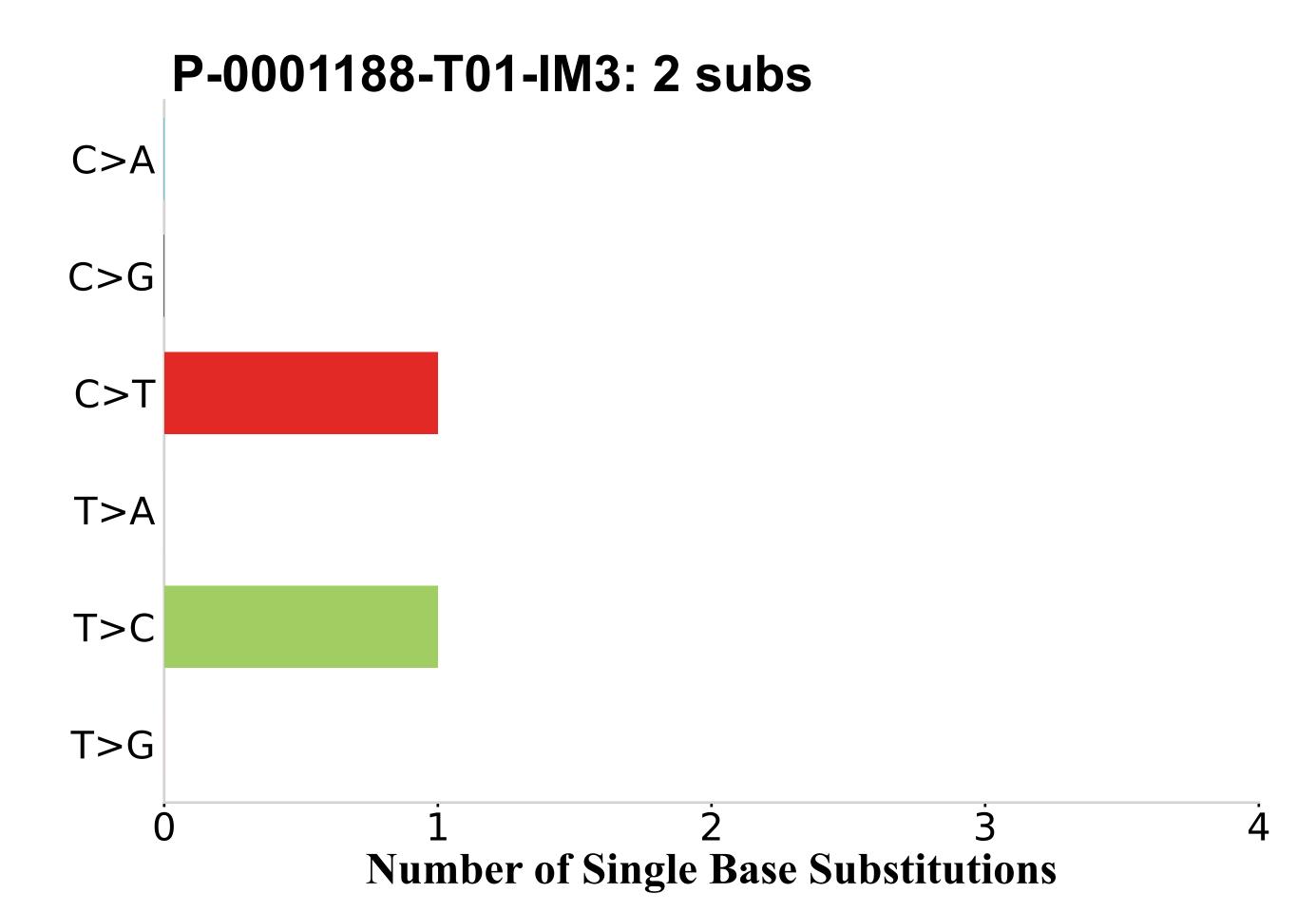


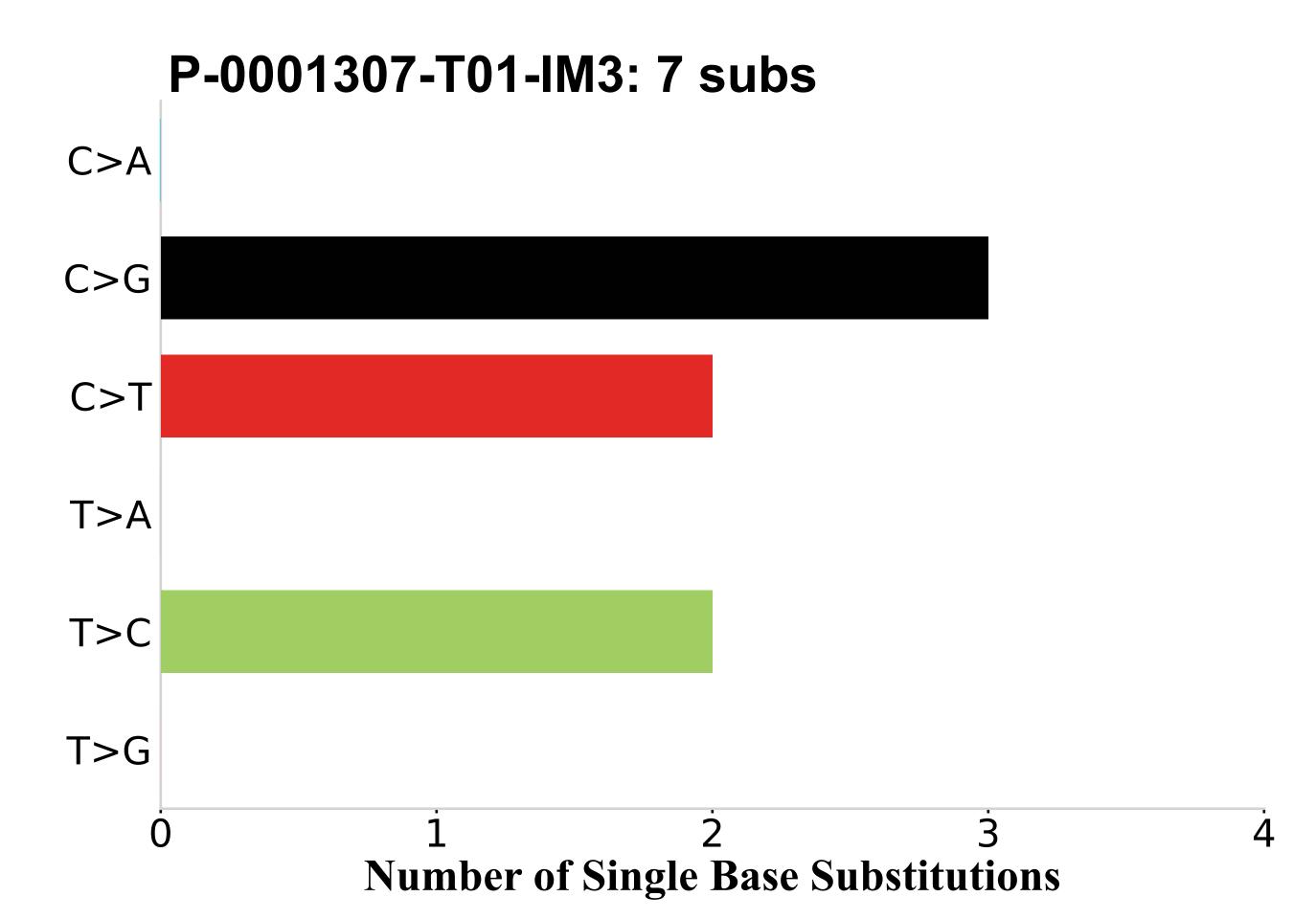
P-0000228-T01-IM3: 3 subs



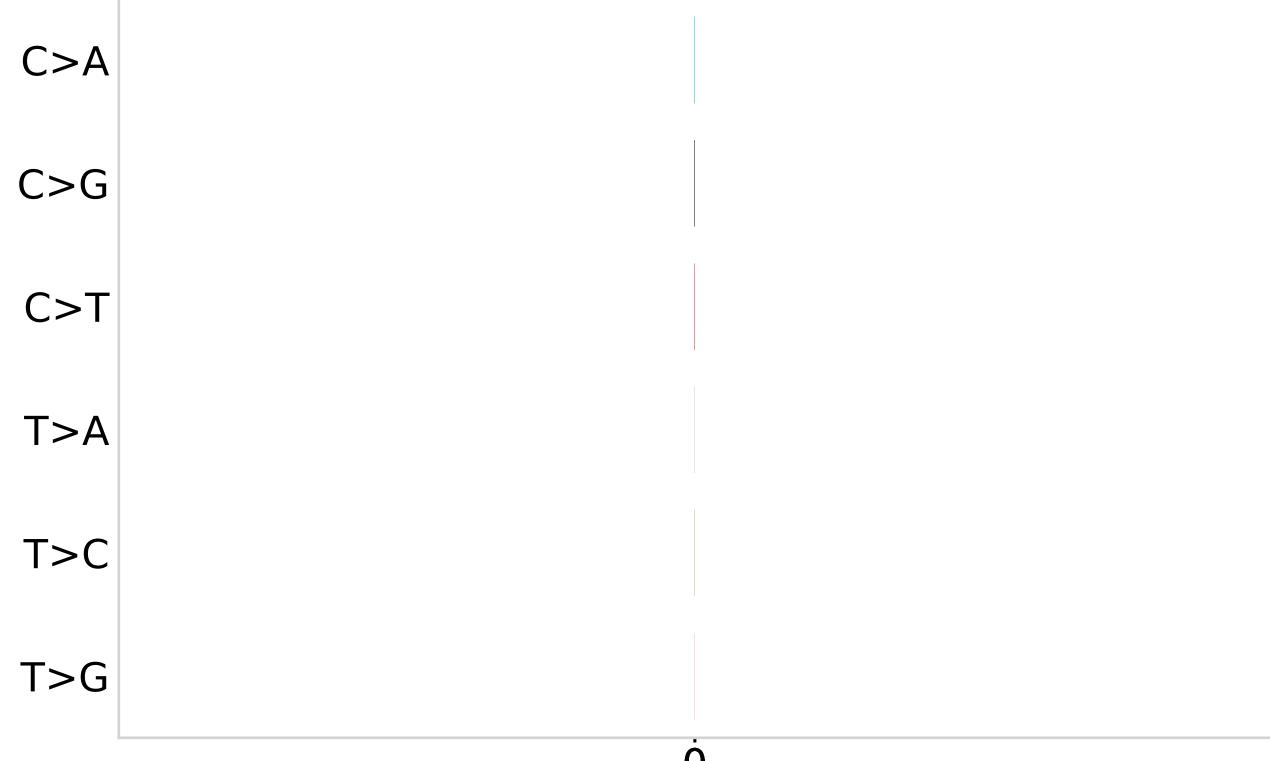




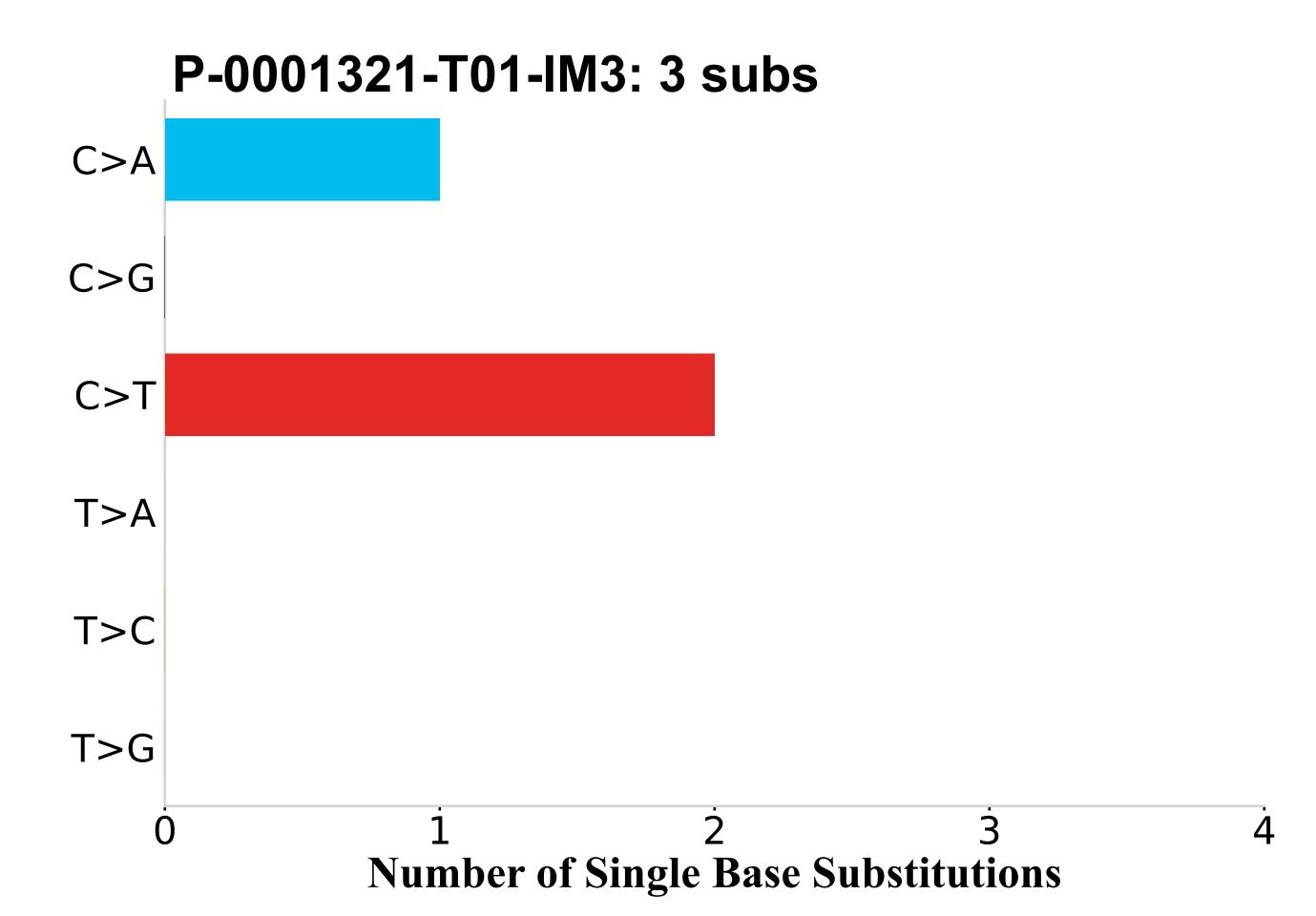


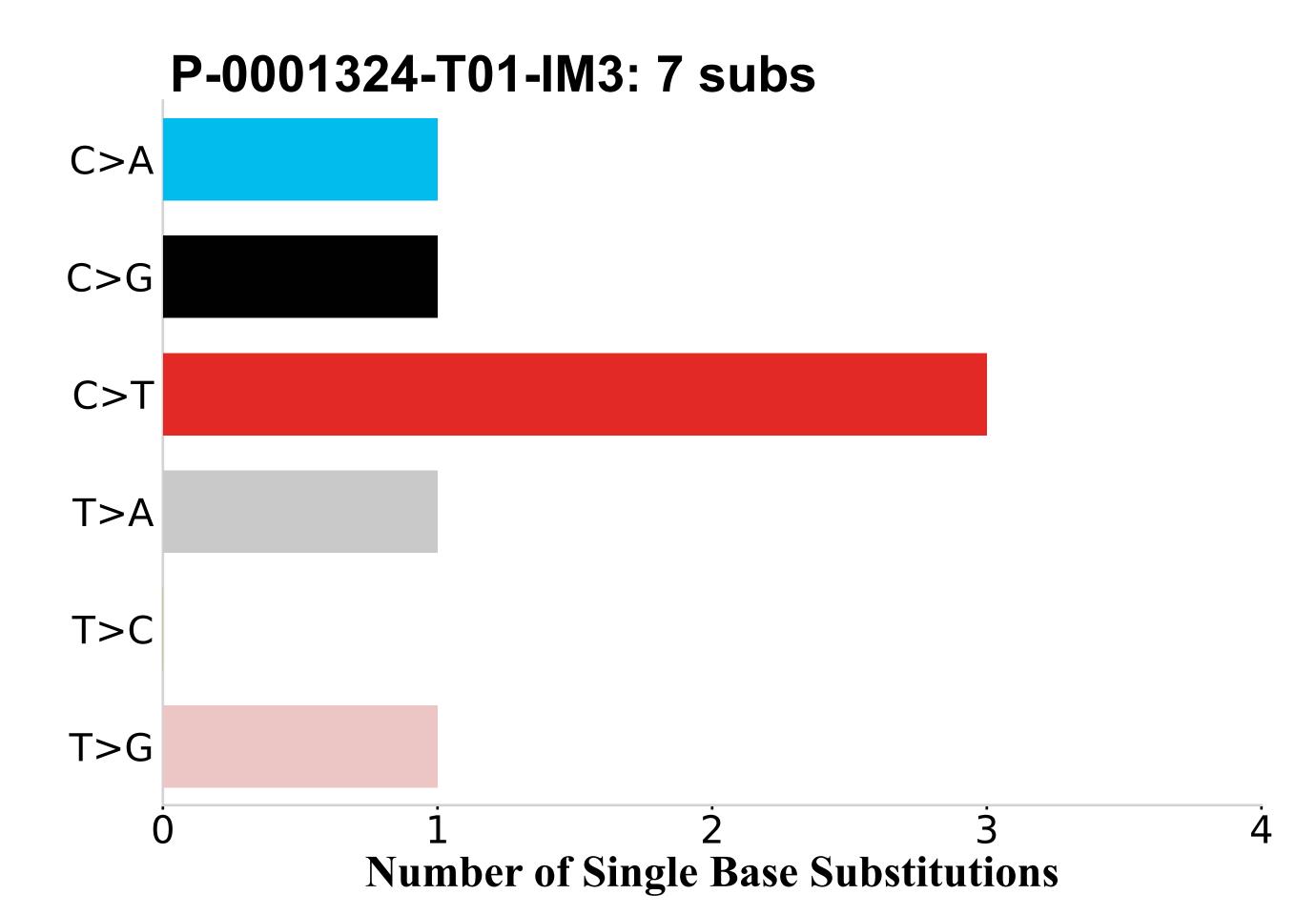


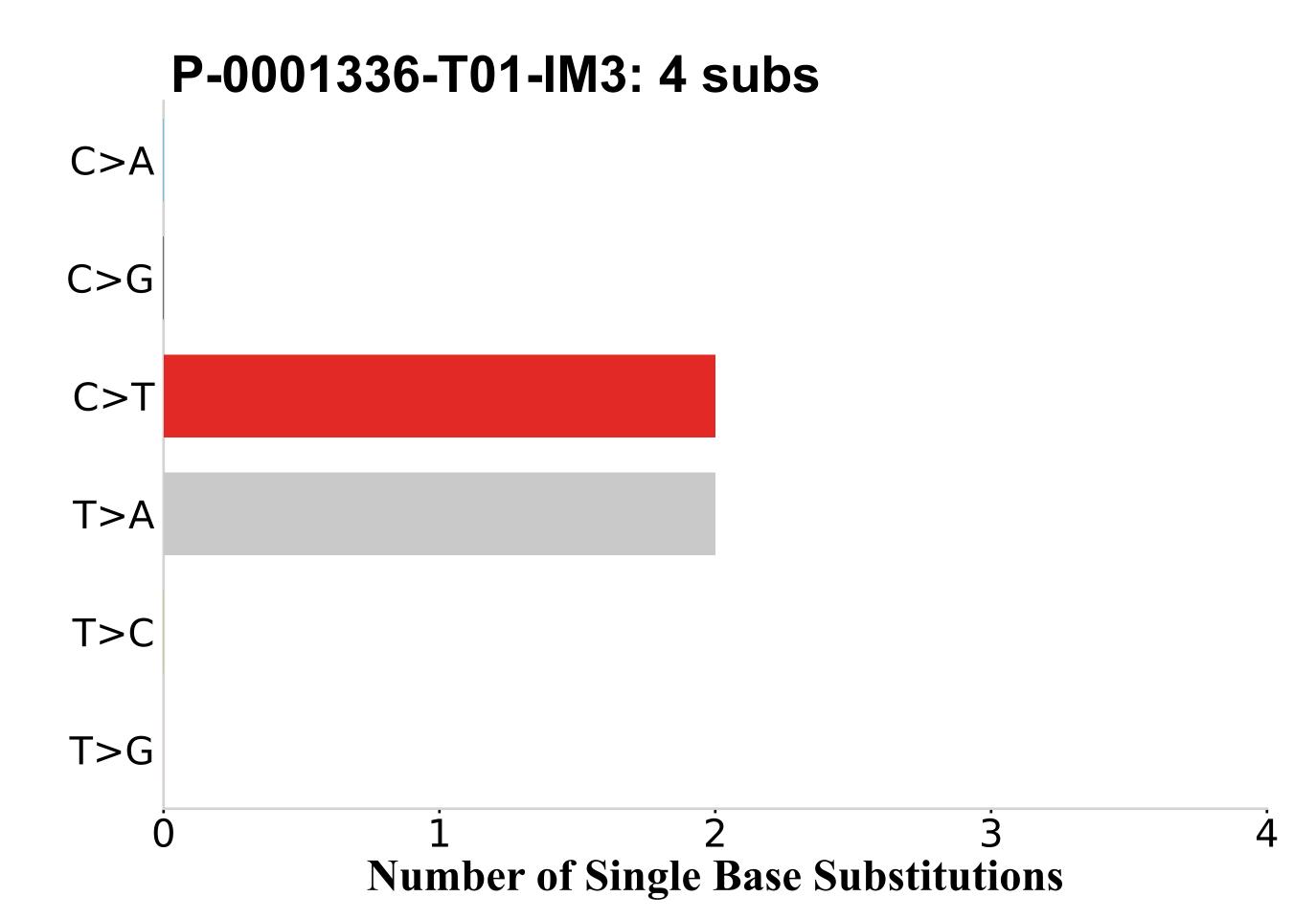
P-0001310-T01-IM3: 0 subs

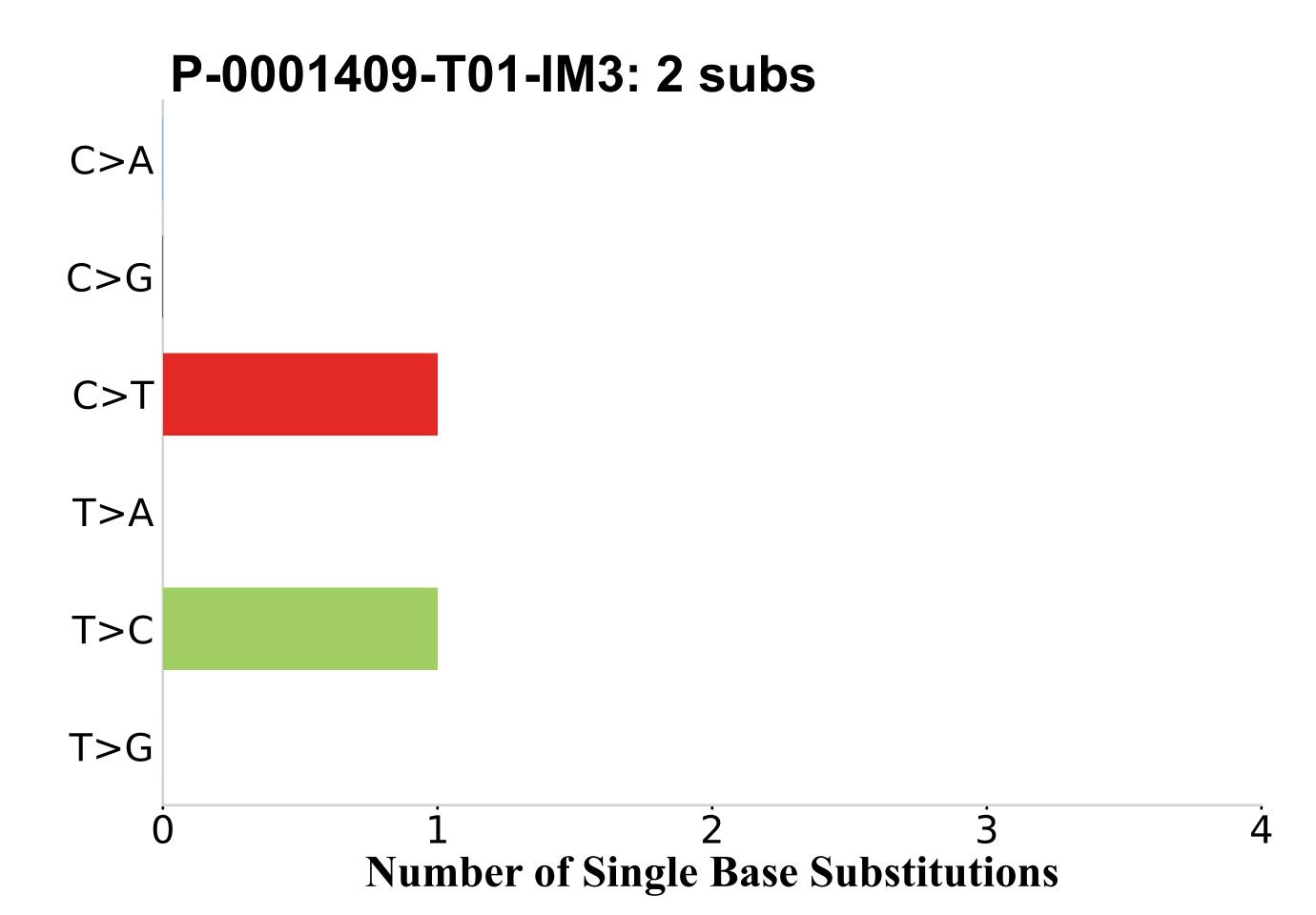


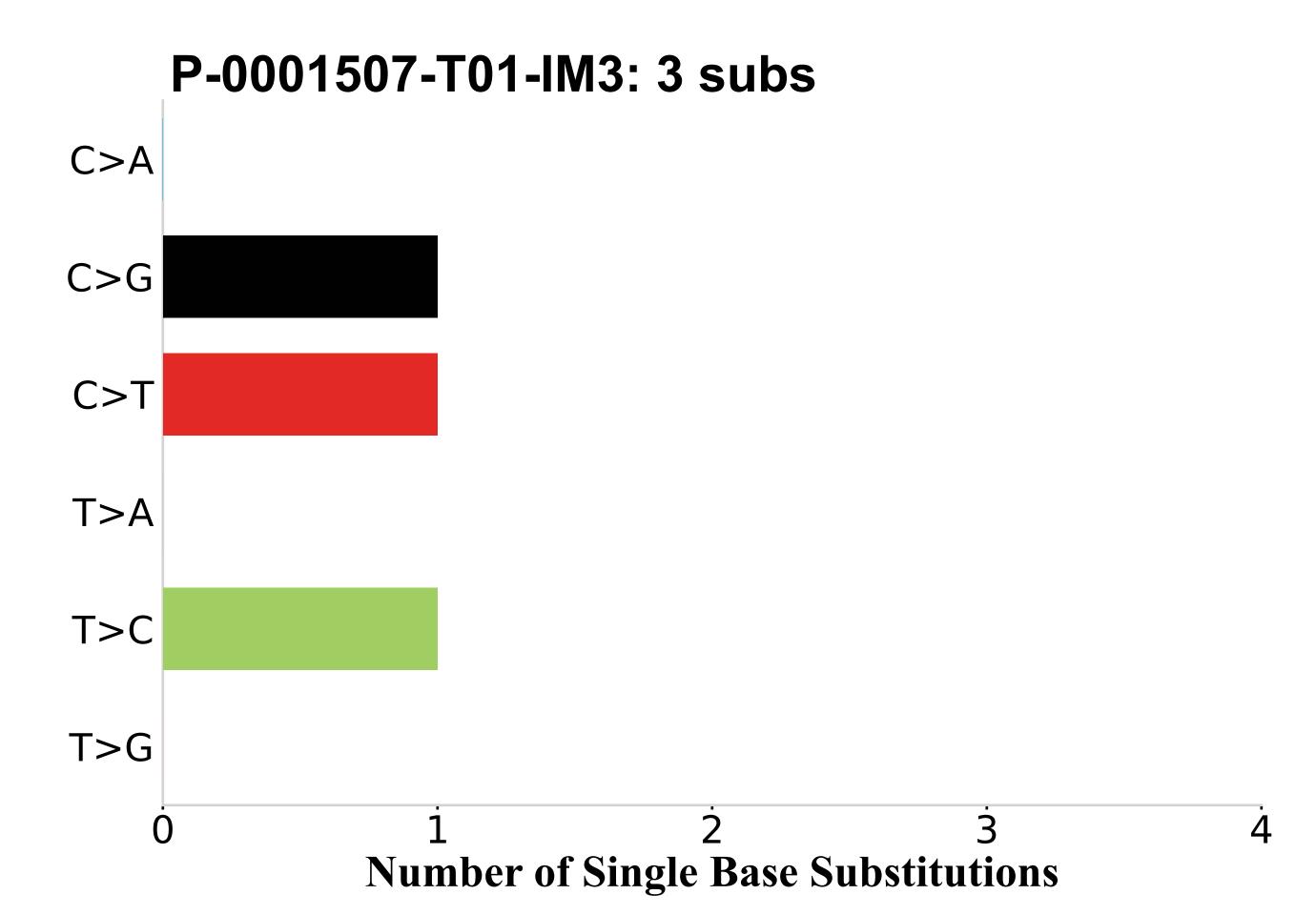
Number of Single Base Substitutions





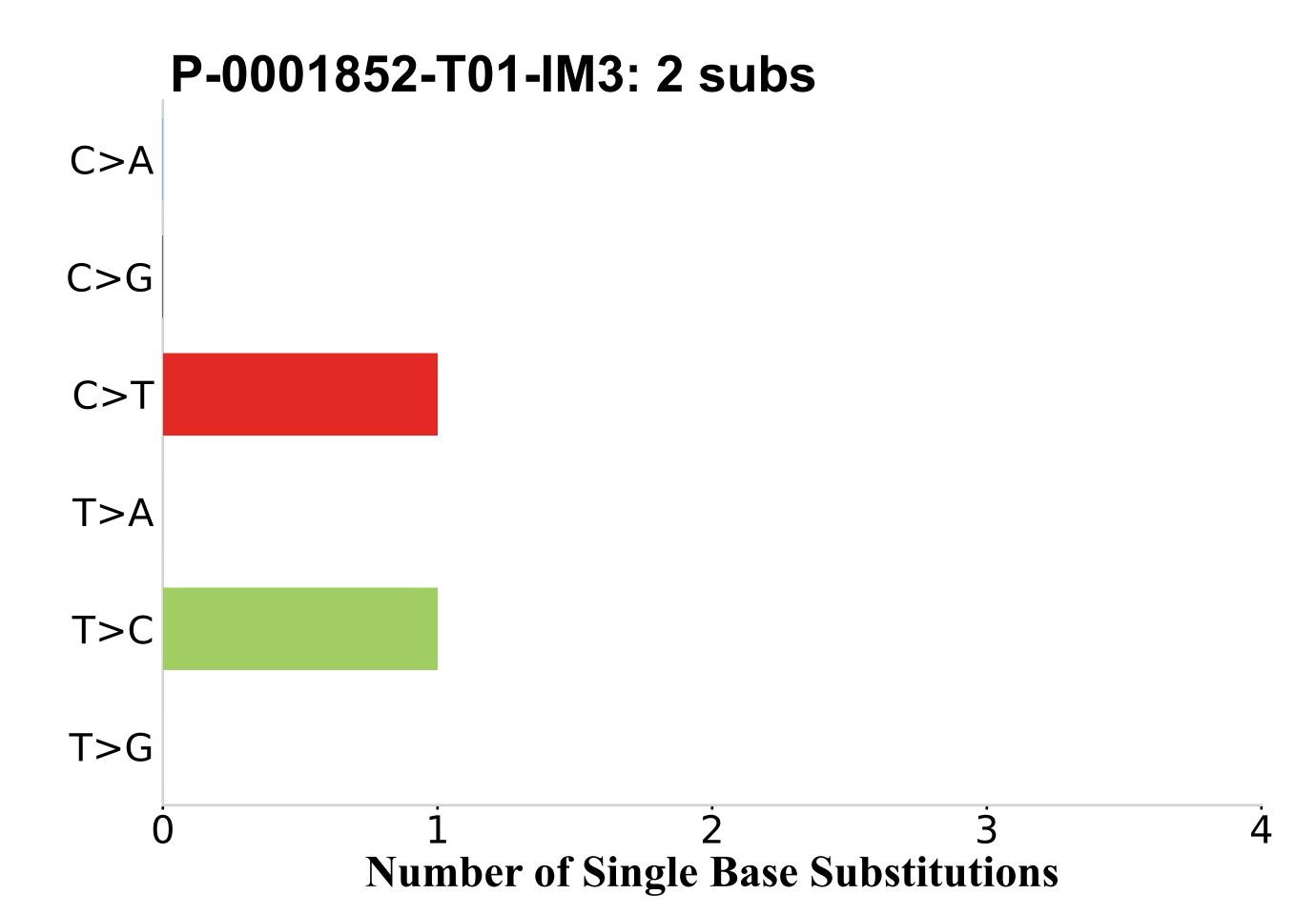


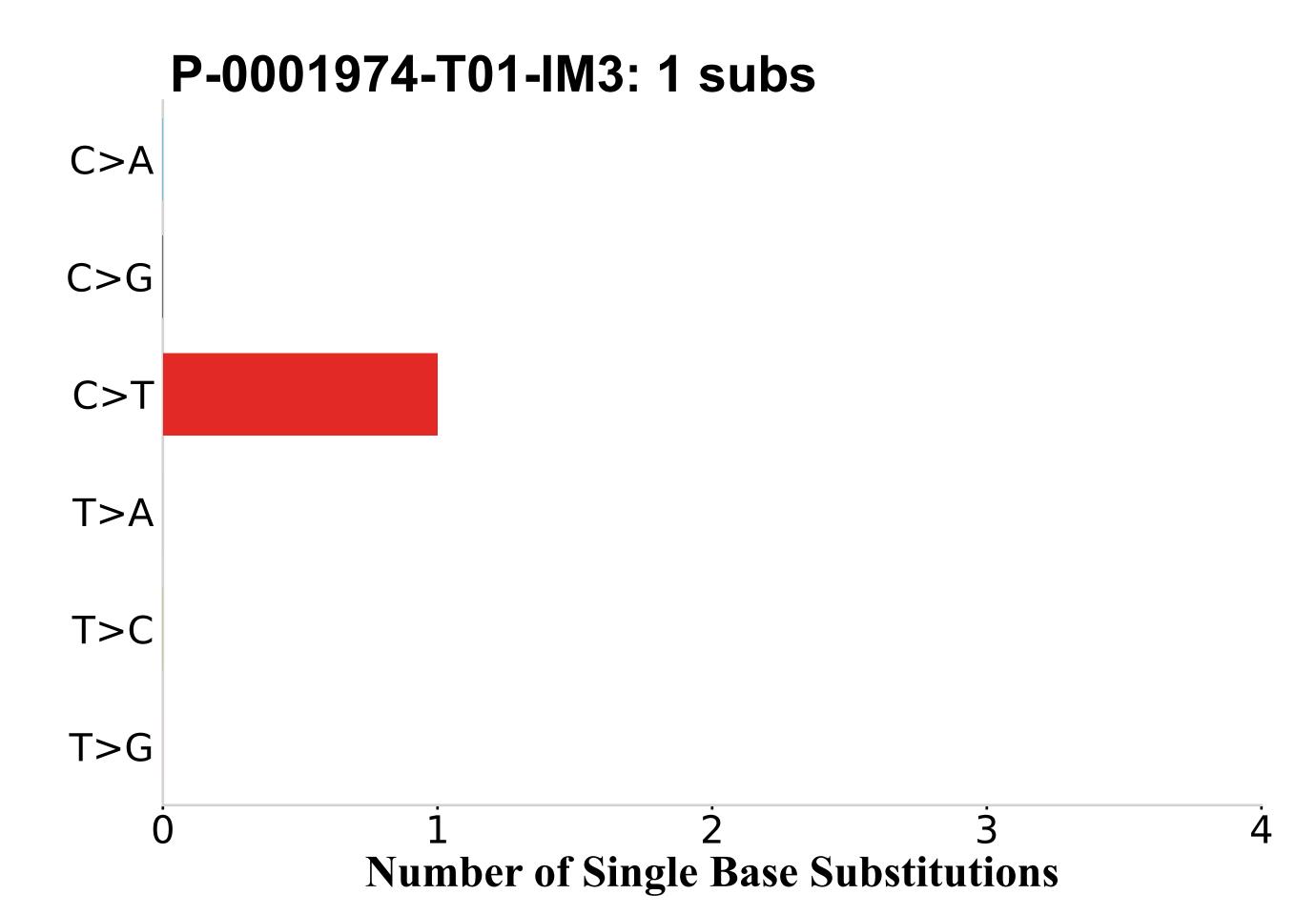


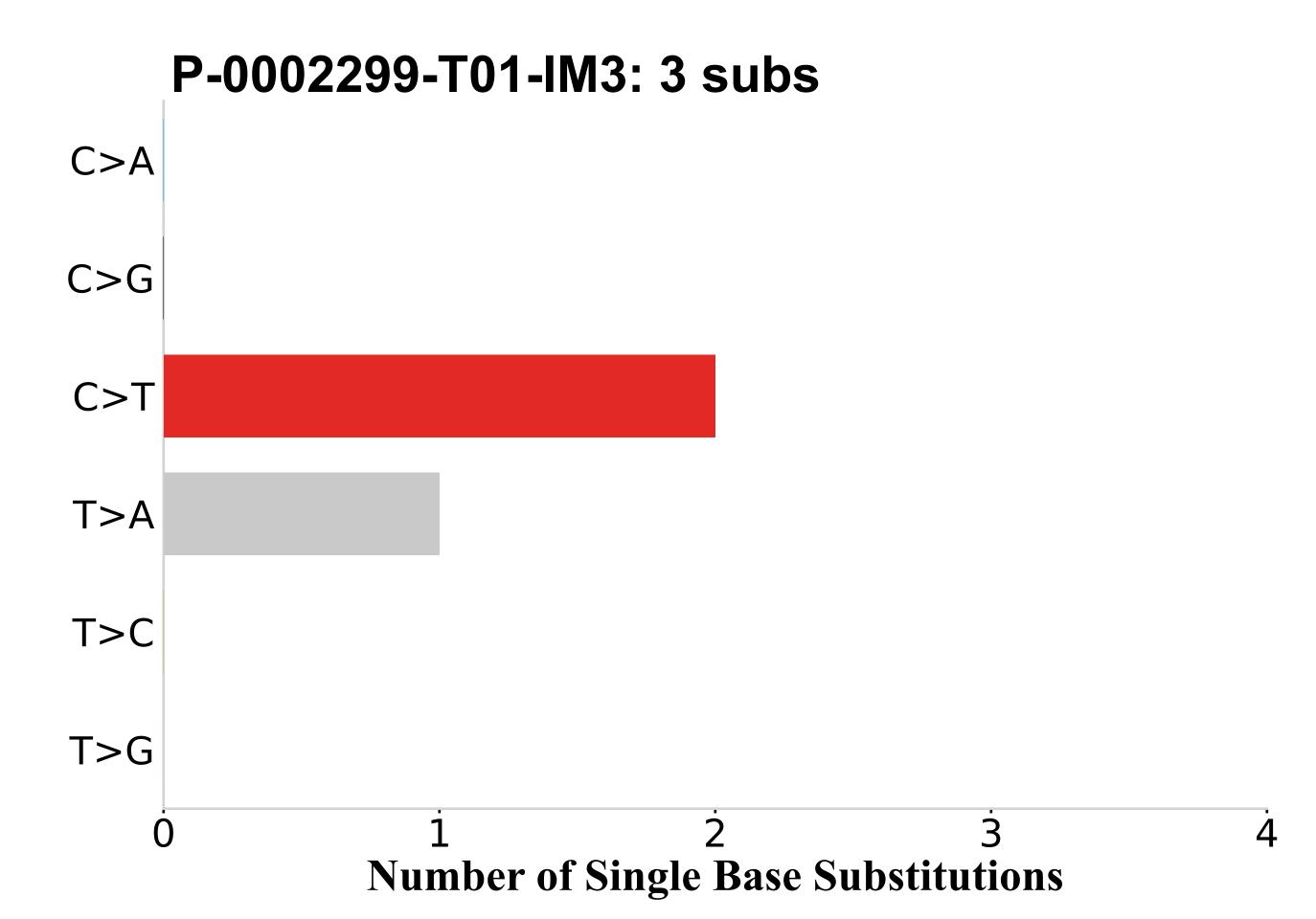


P-0001806-T01-IM3: 3 subs C>A C>G C>T T>A T>C T>G

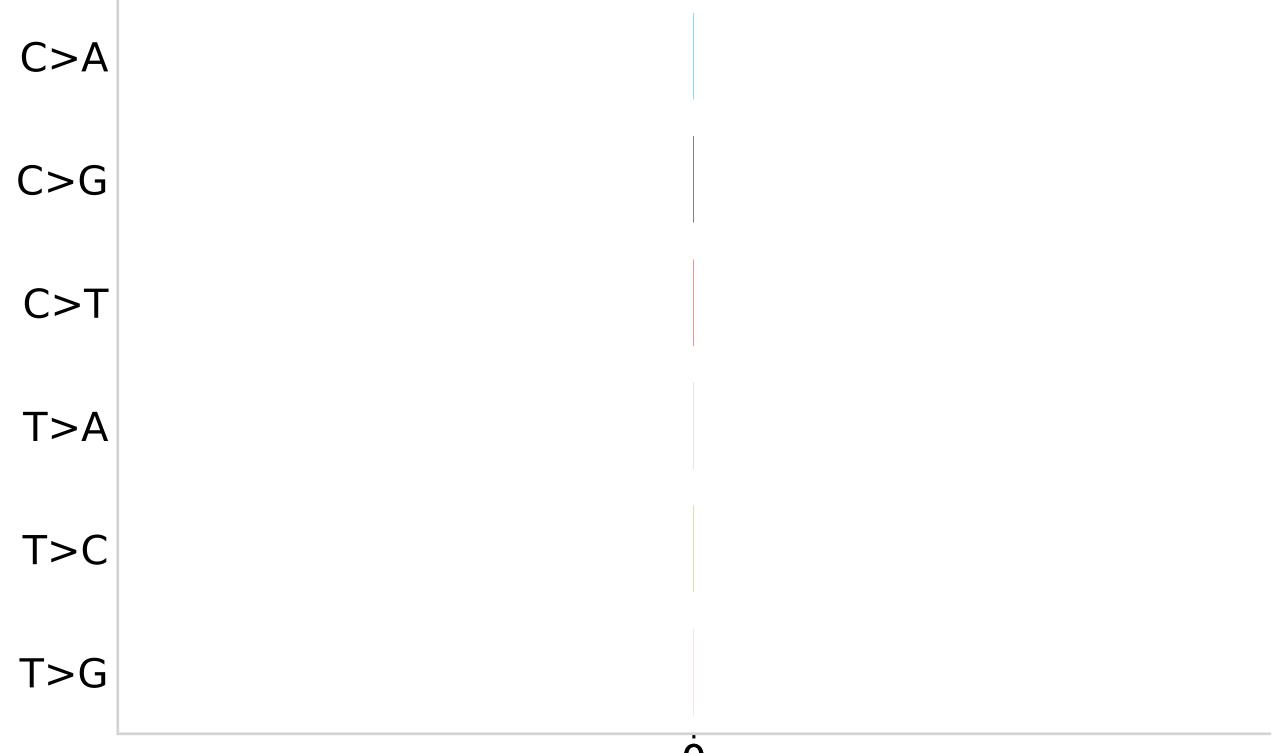
Number of Single Base Substitutions





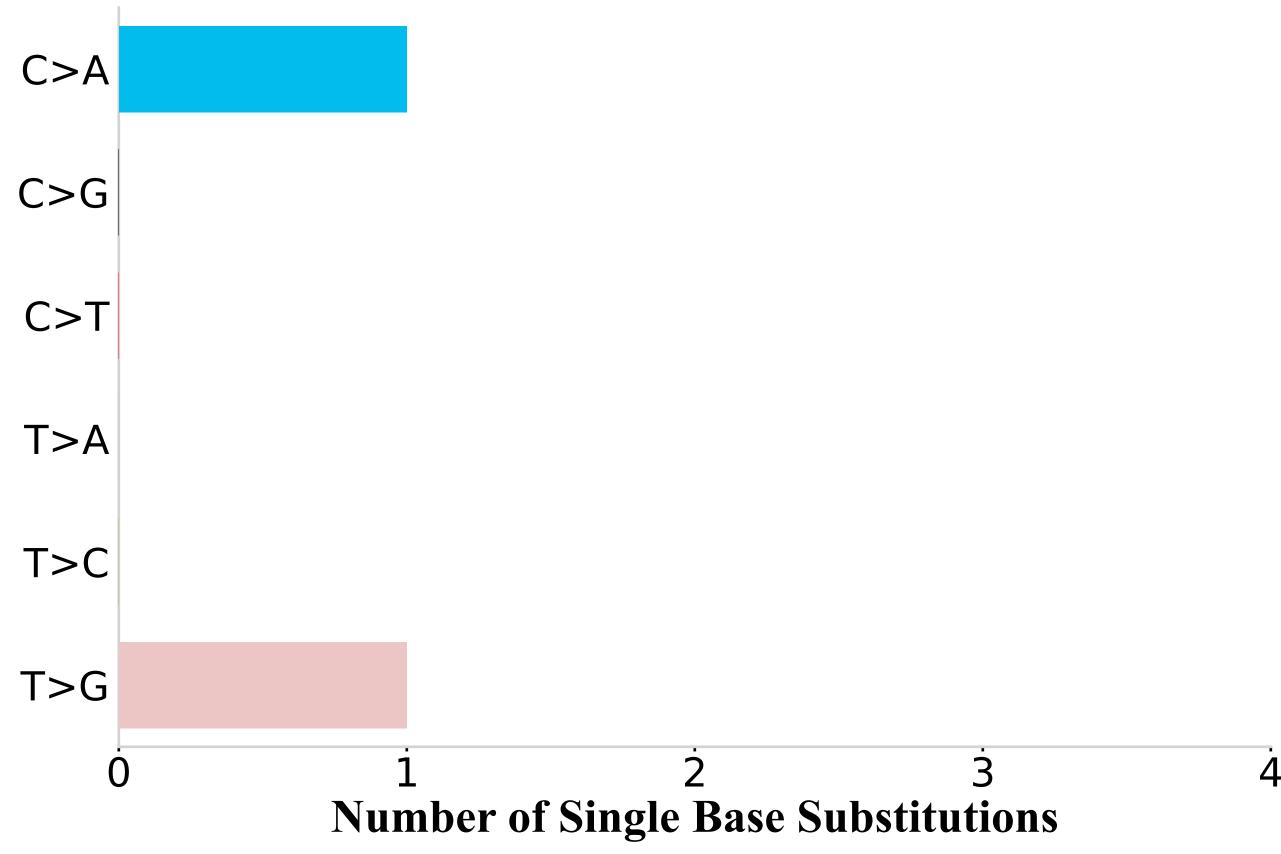


P-0002317-T01-IM3: 0 subs

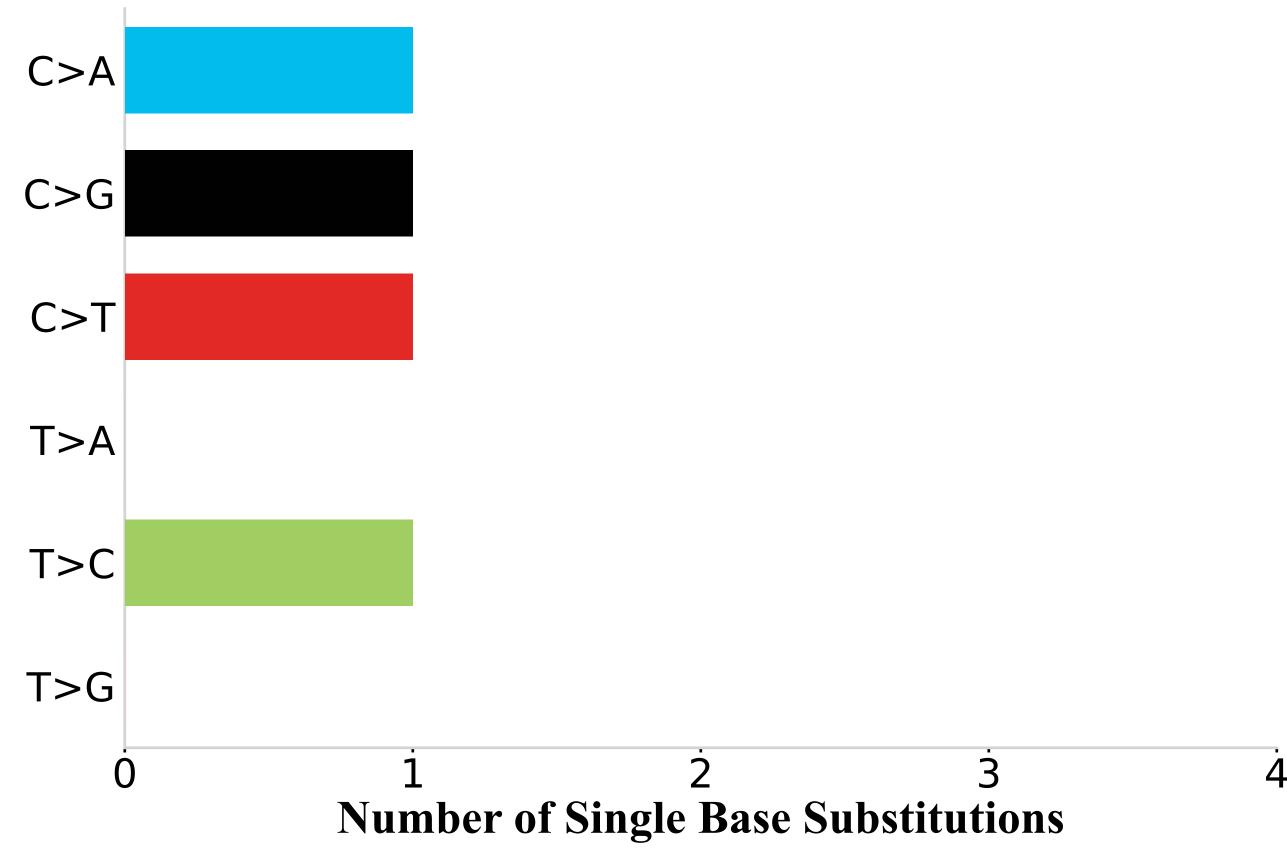


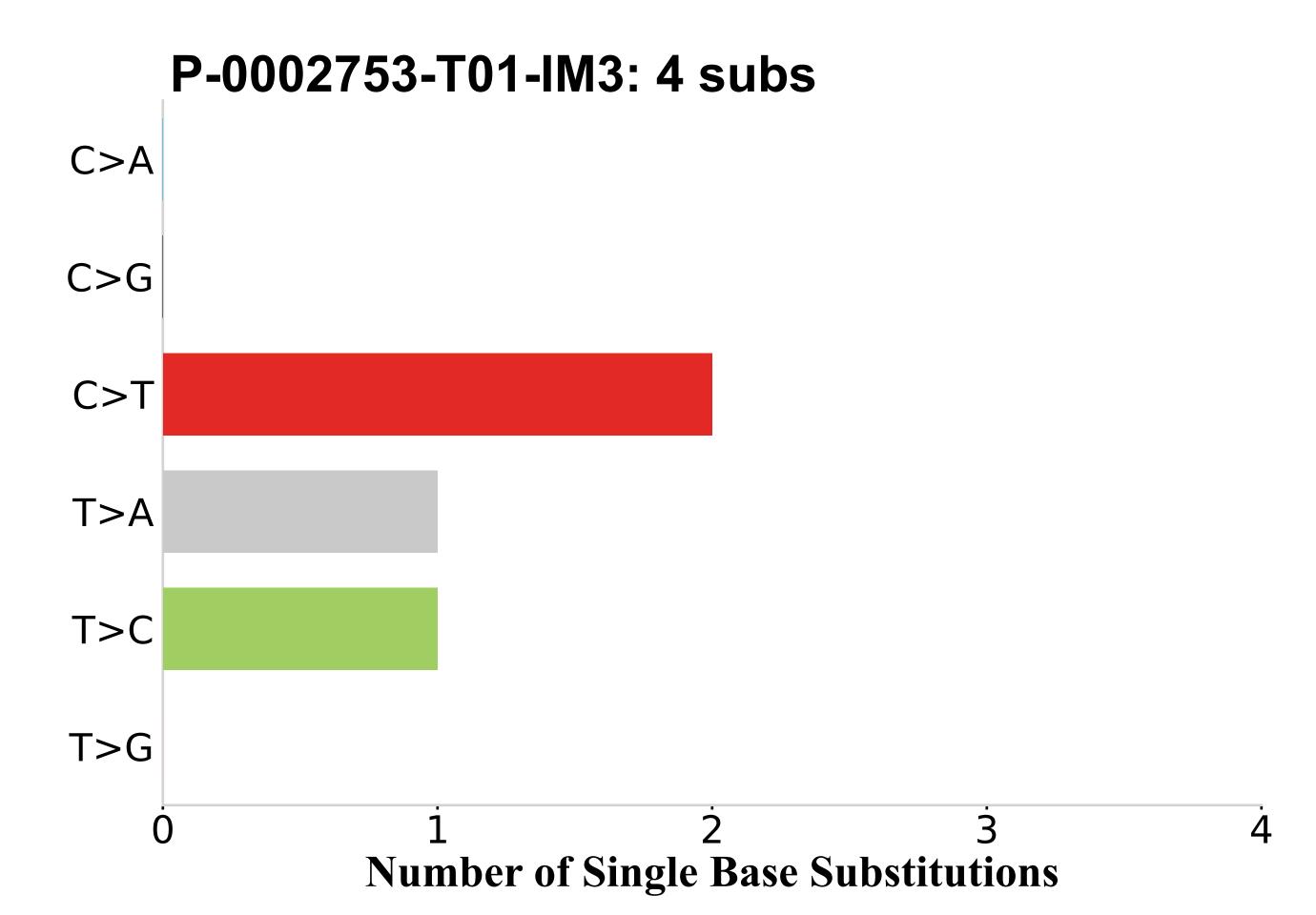
Number of Single Base Substitutions

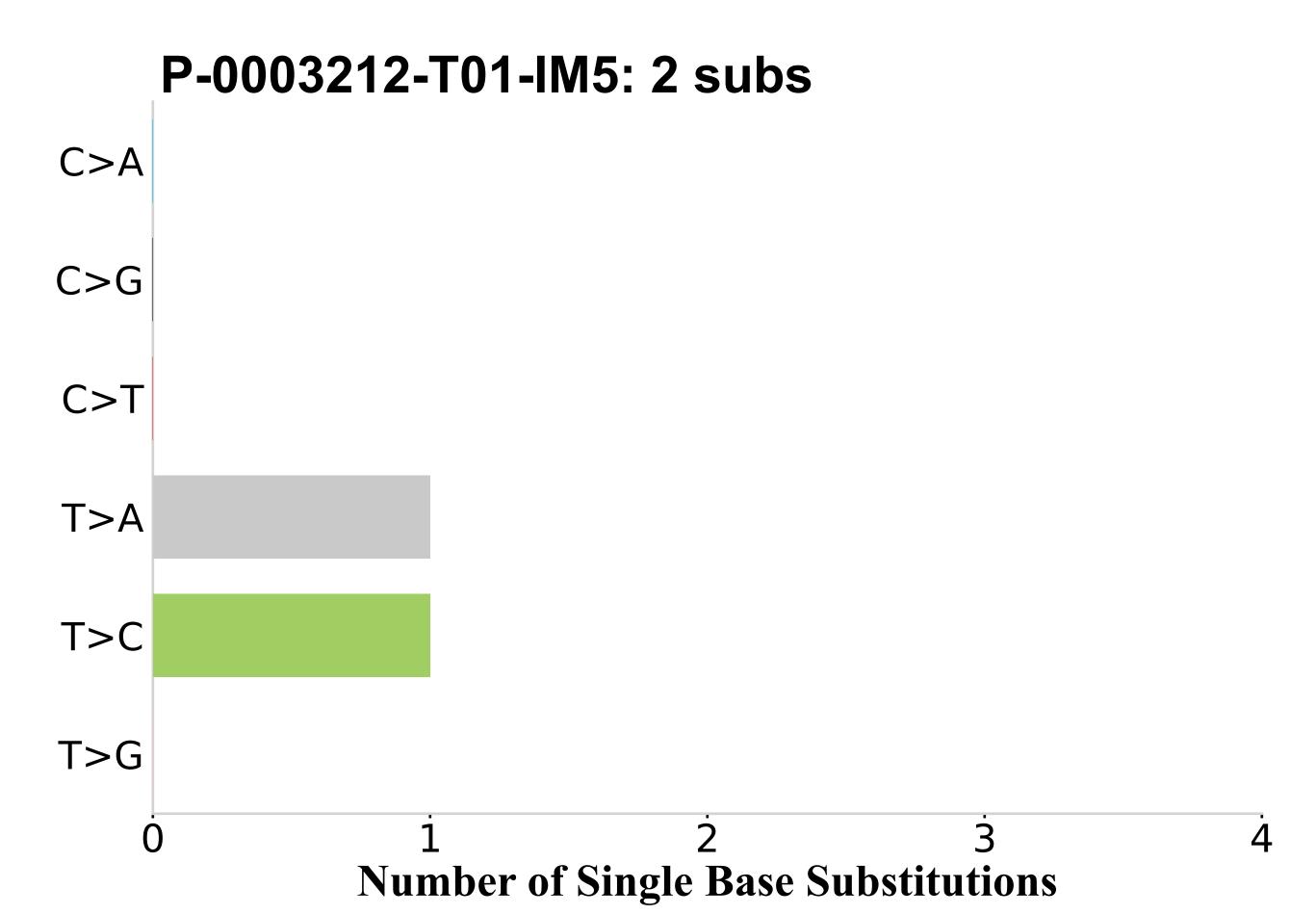
P-0002697-T01-IM3: 2 subs



P-0002719-T01-IM3: 4 subs

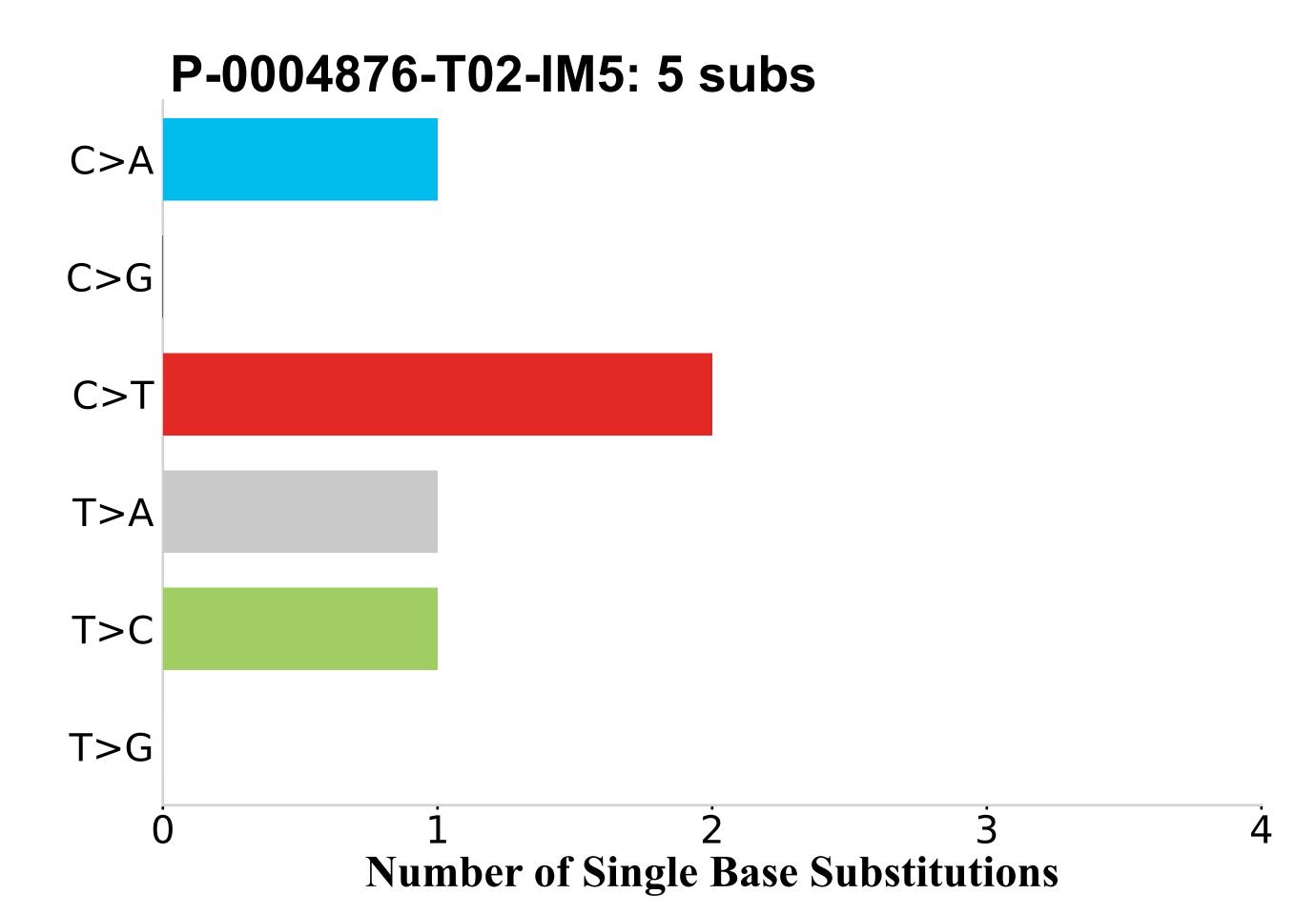


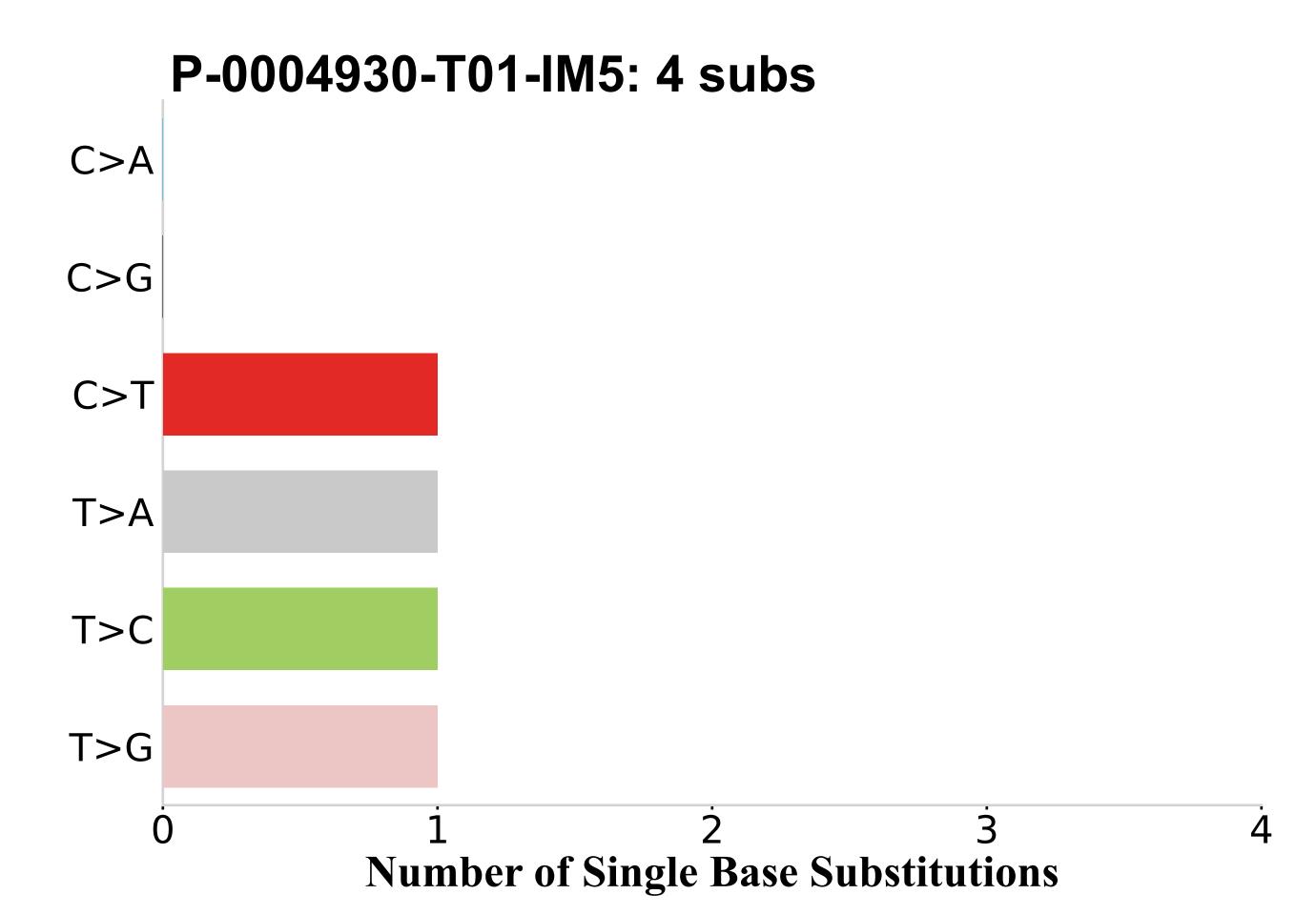


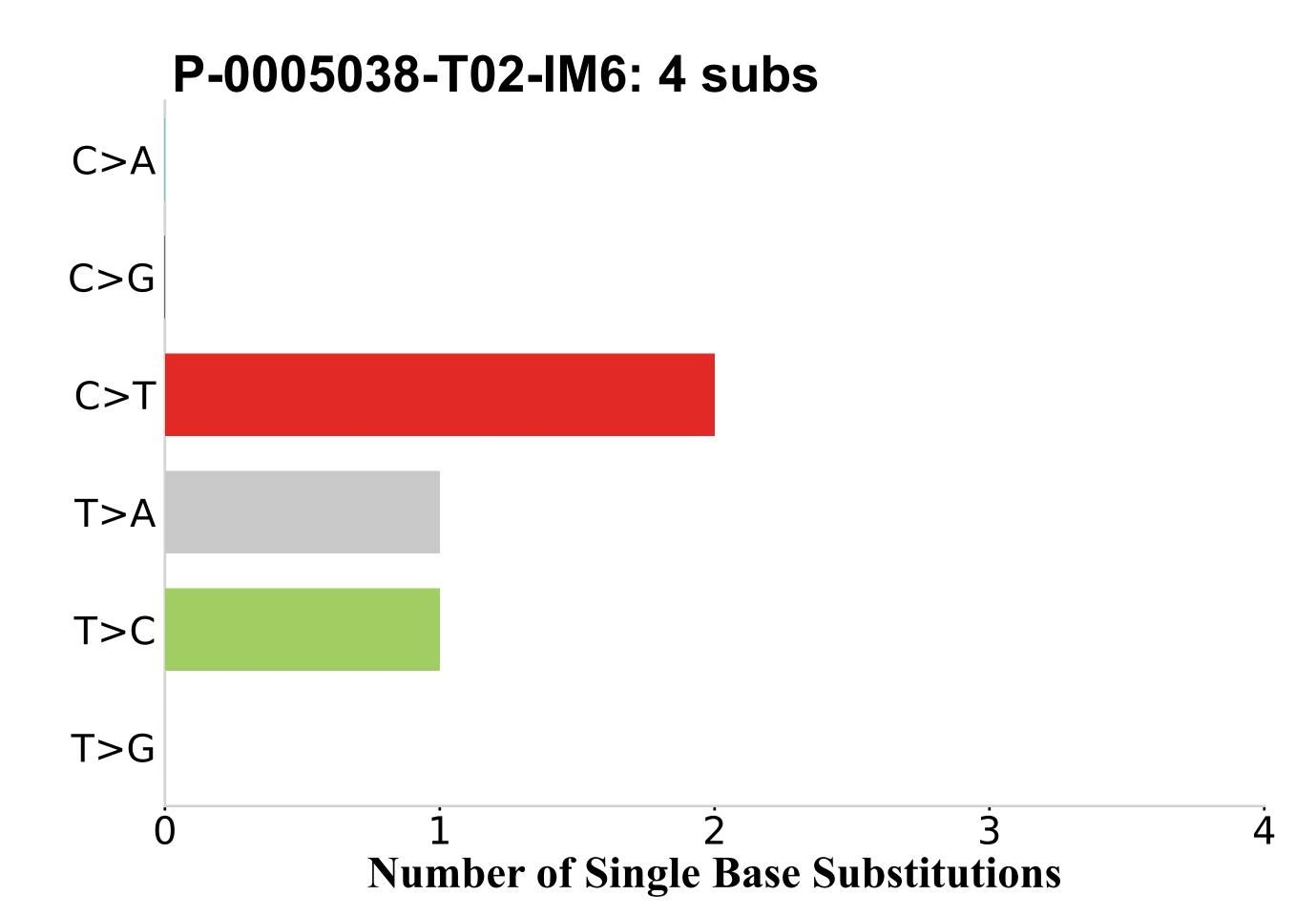


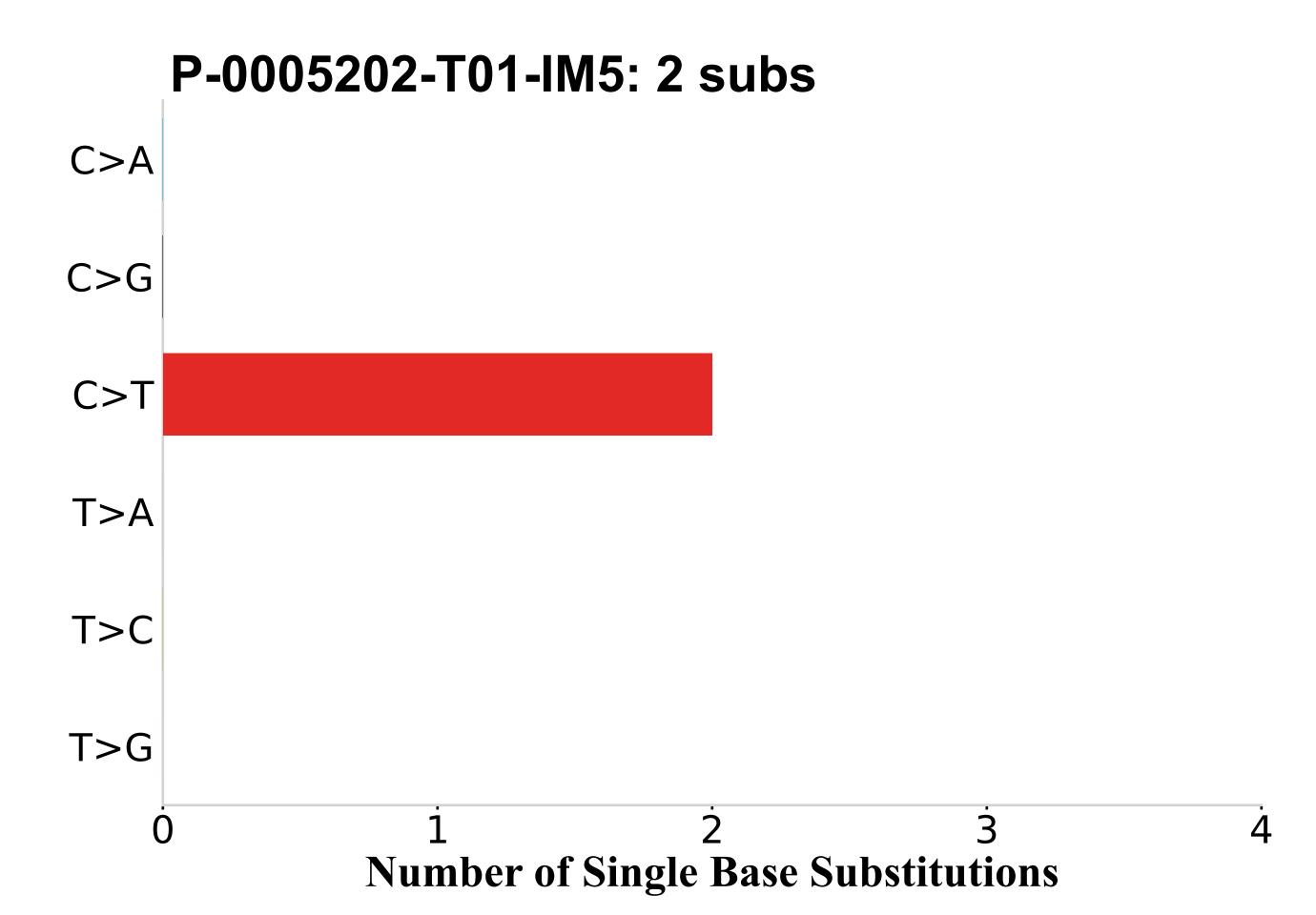
P-0004247-T01-IM5: 3 subs C>A C>G C>T T>A T>C T>G

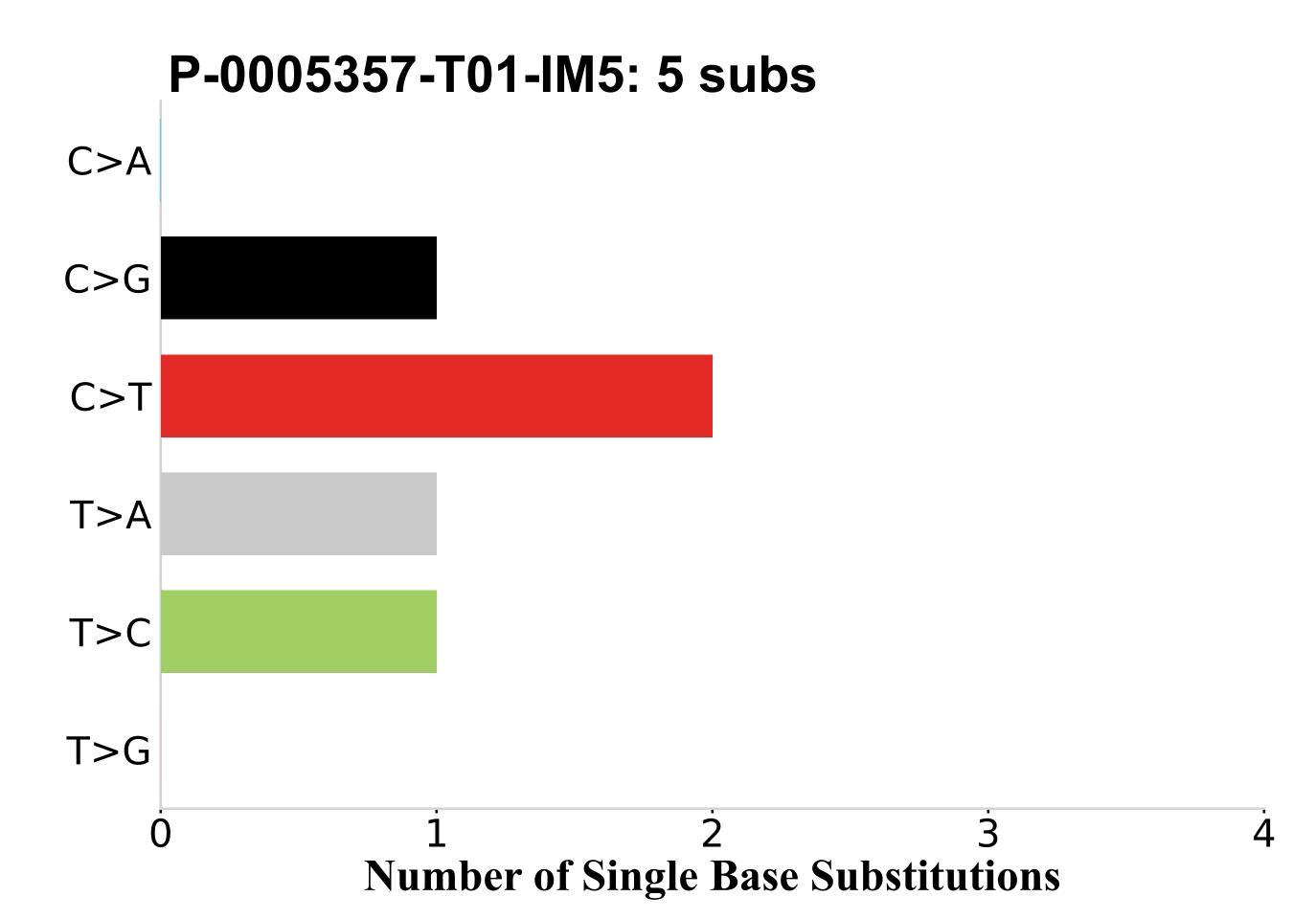
Number of Single Base Substitutions

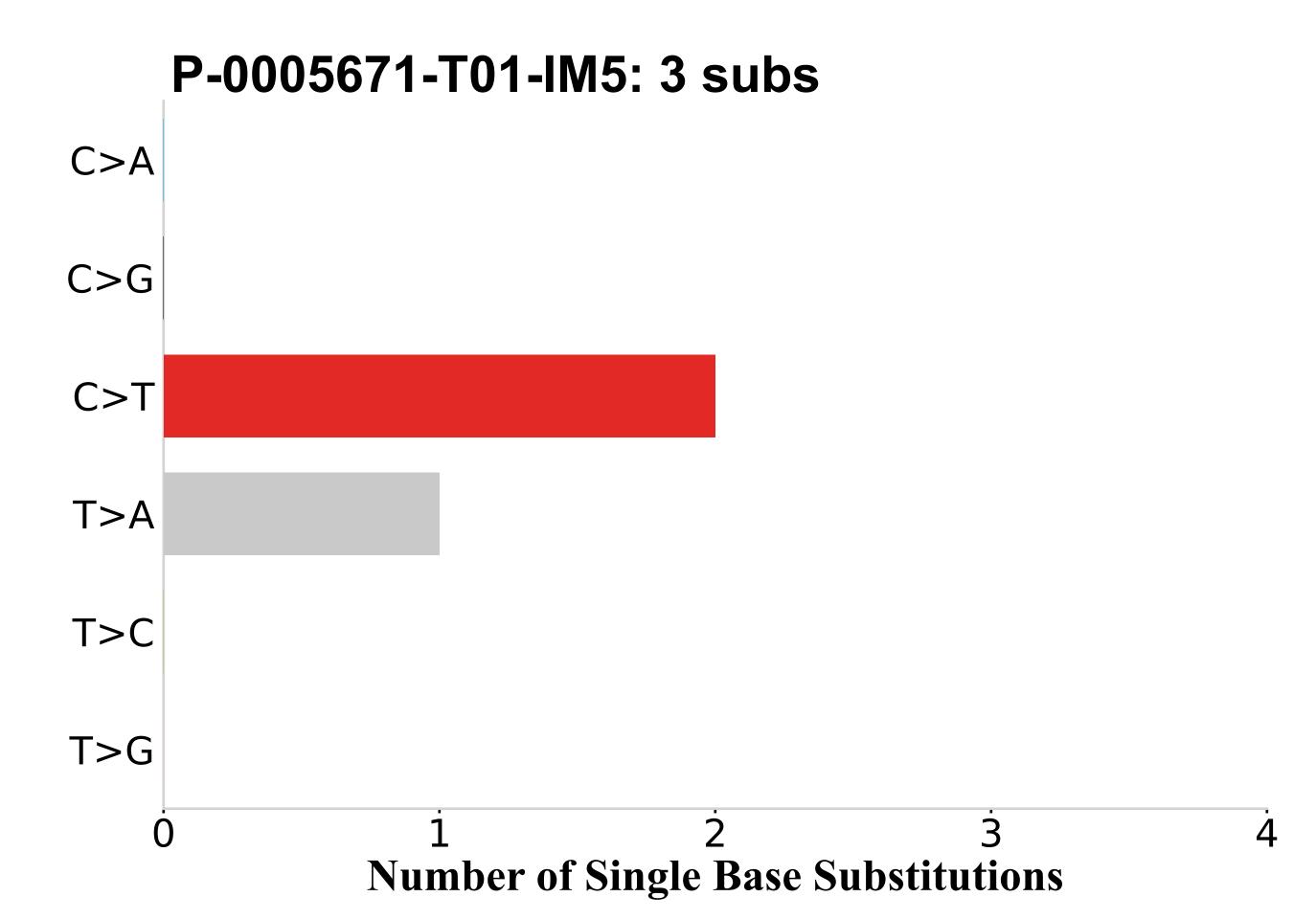


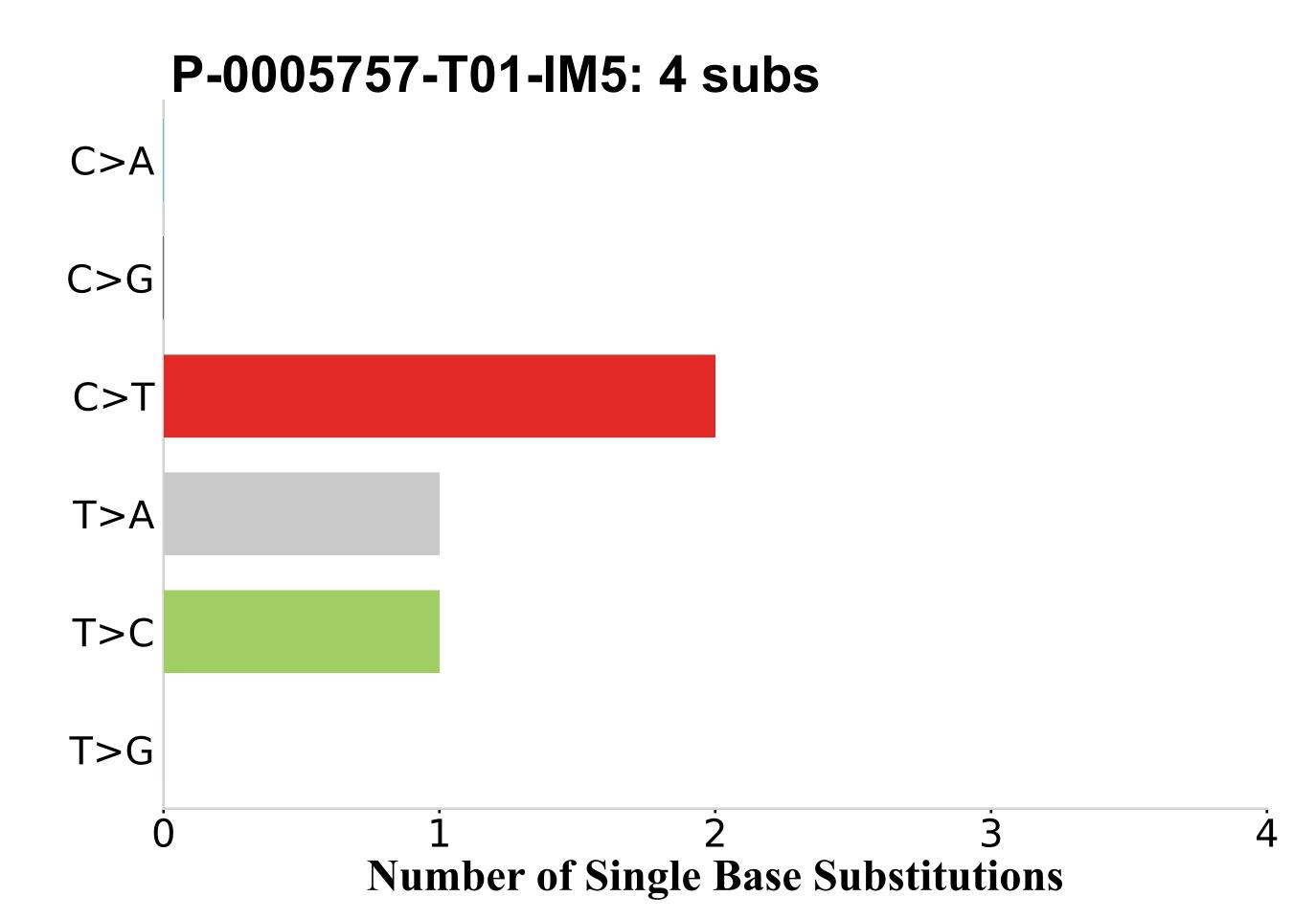


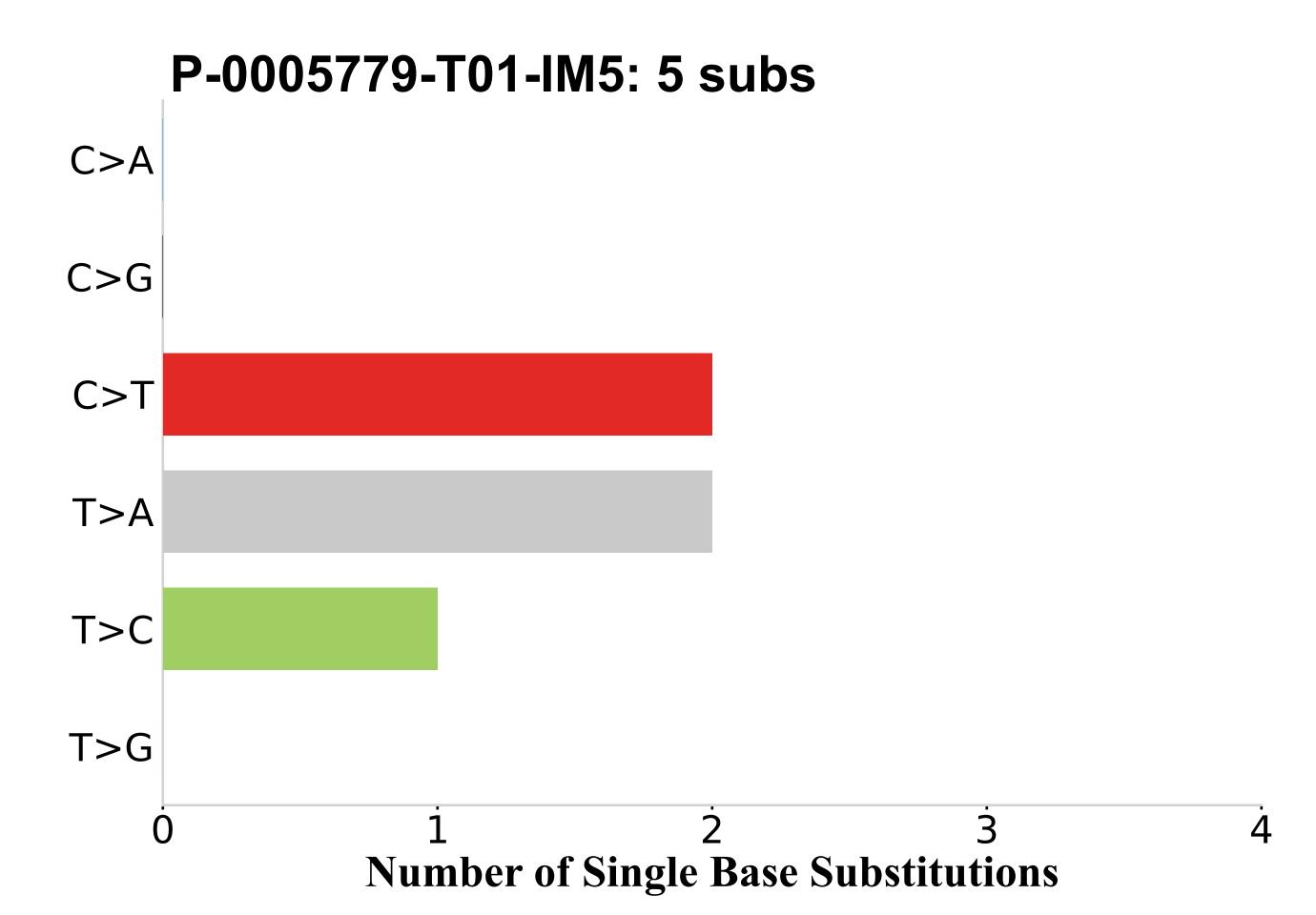


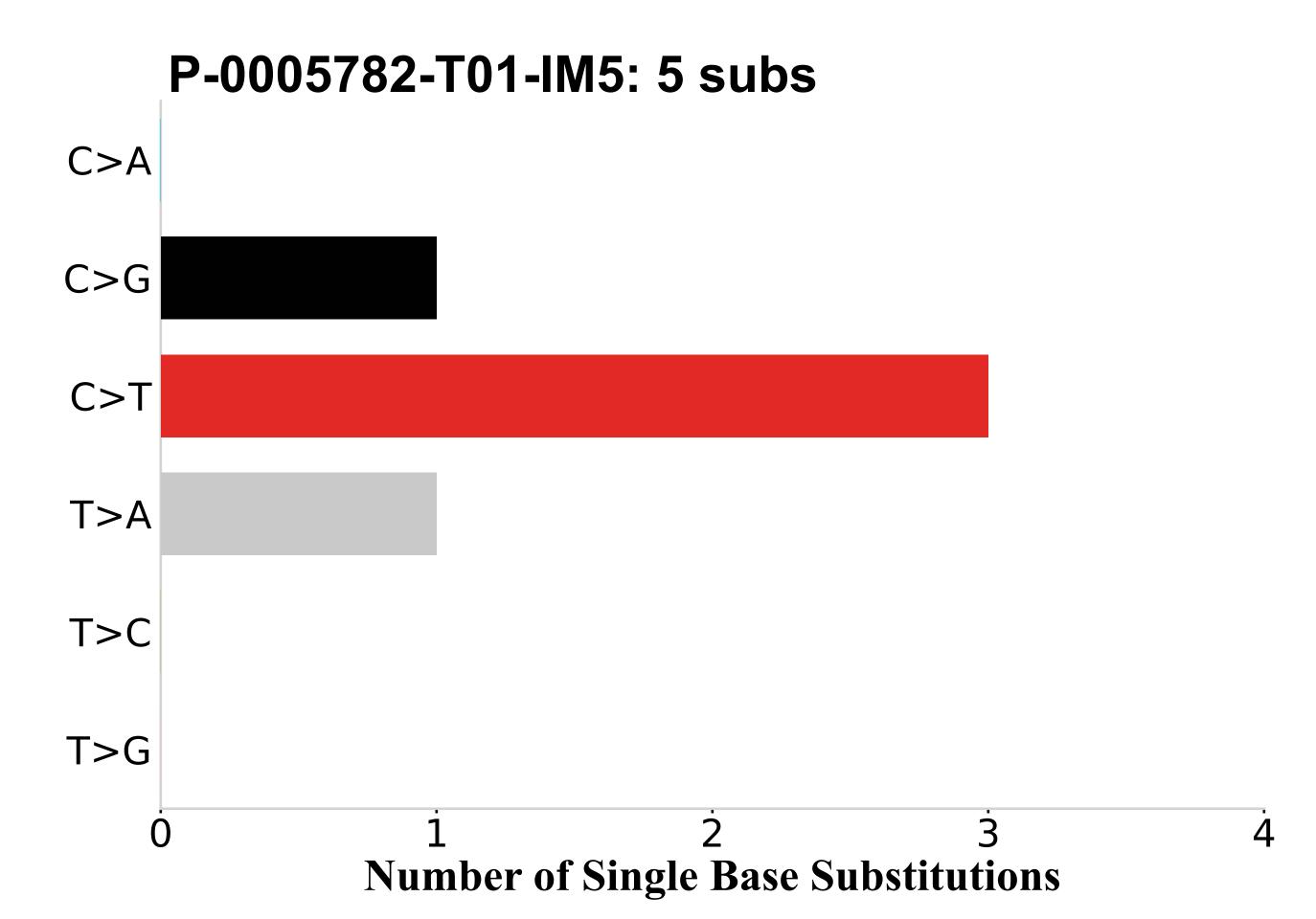


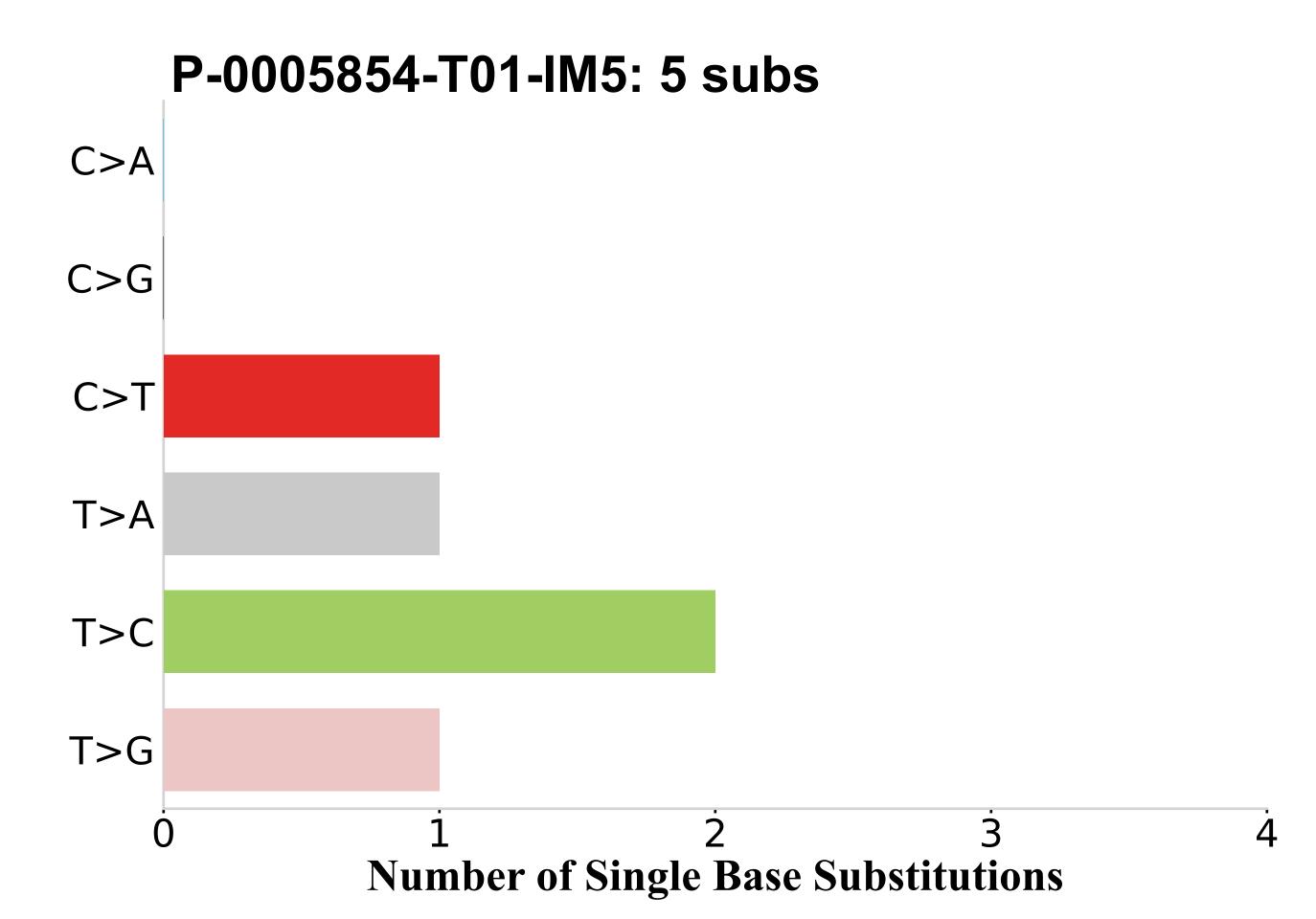


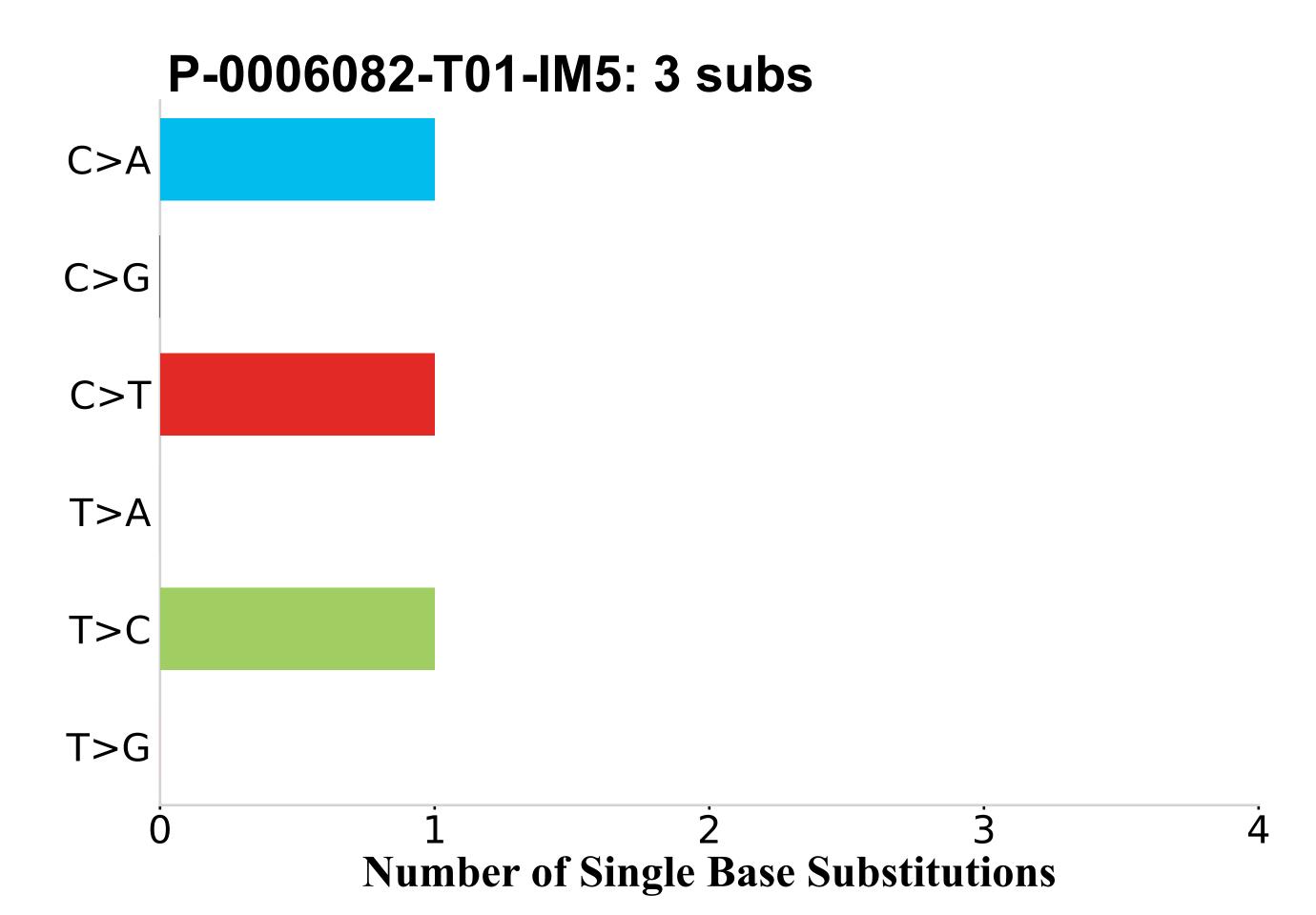




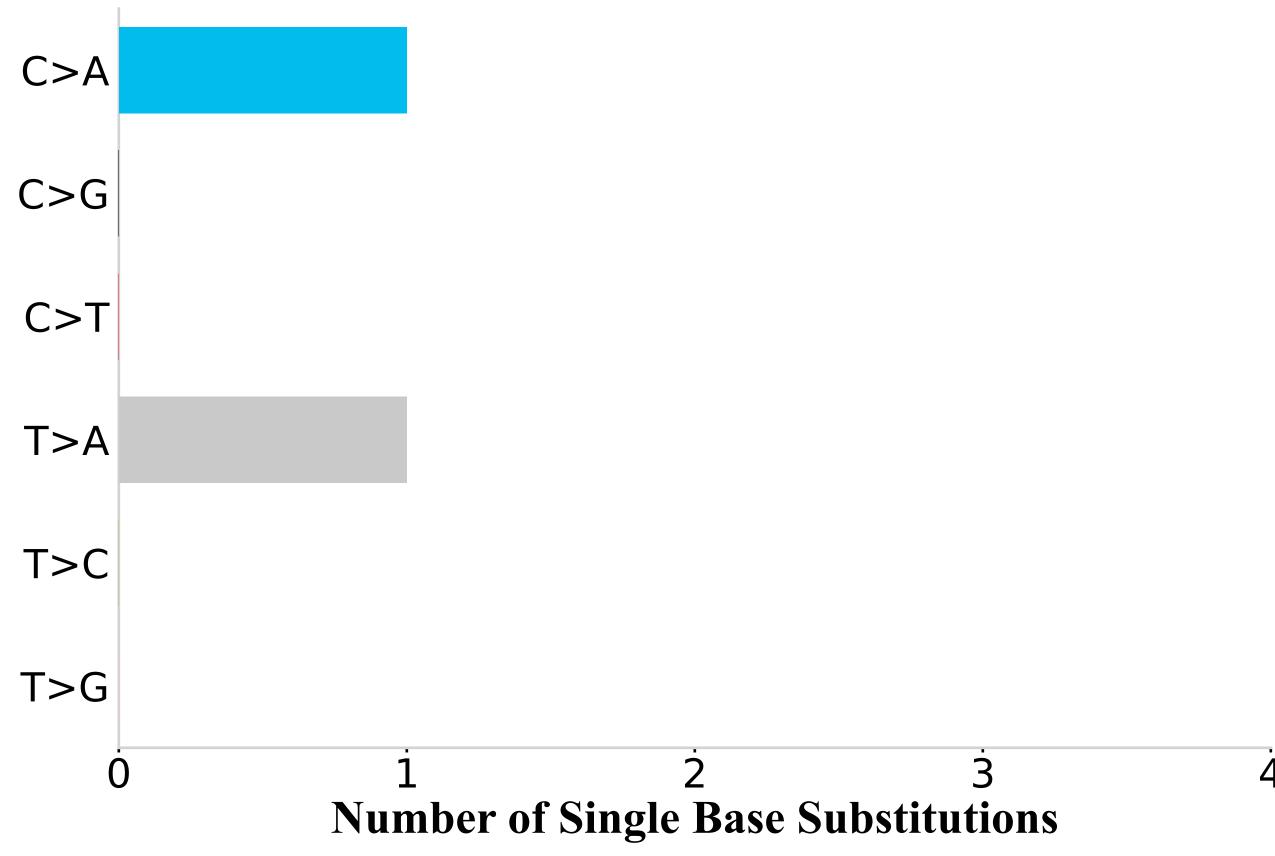


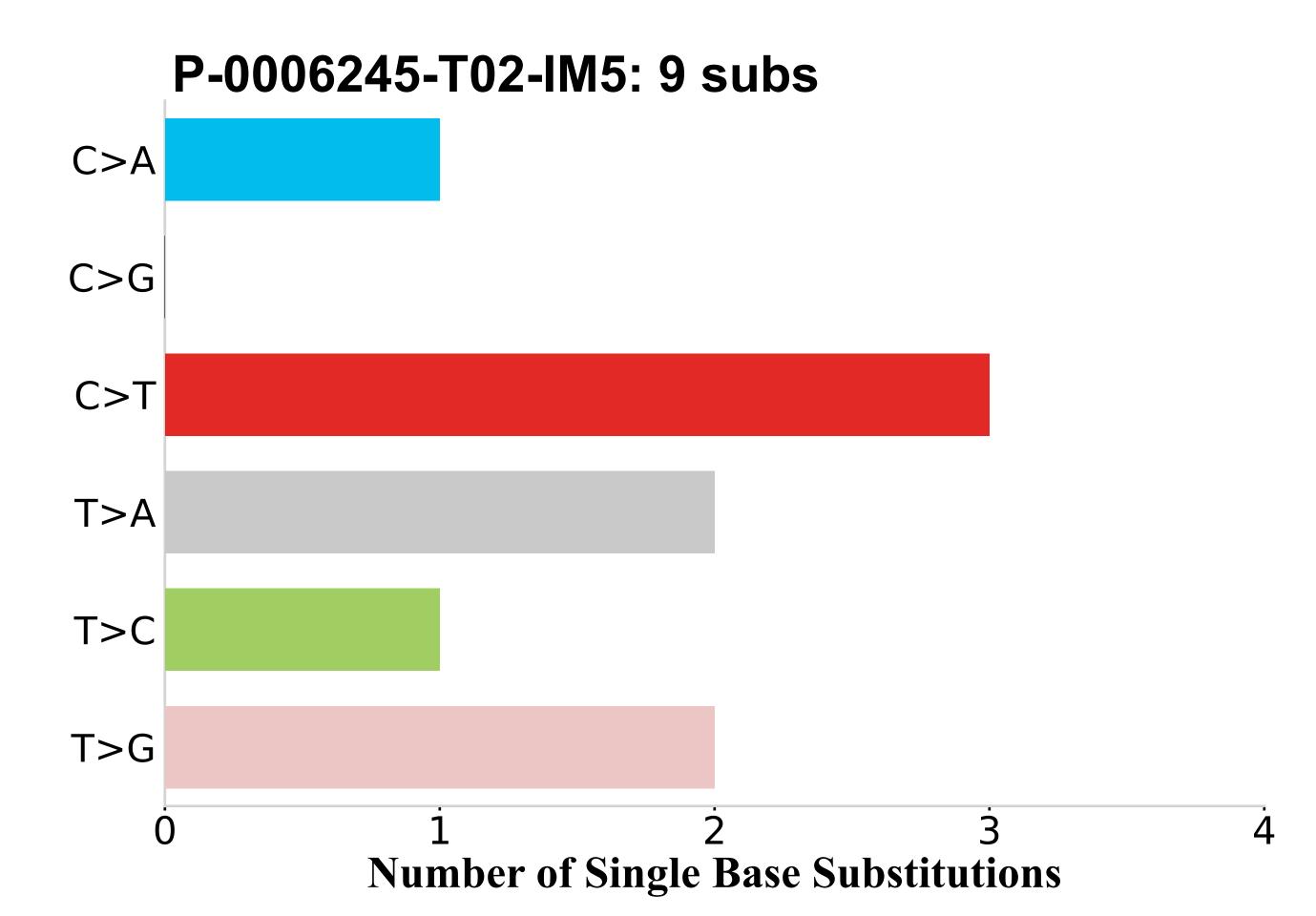




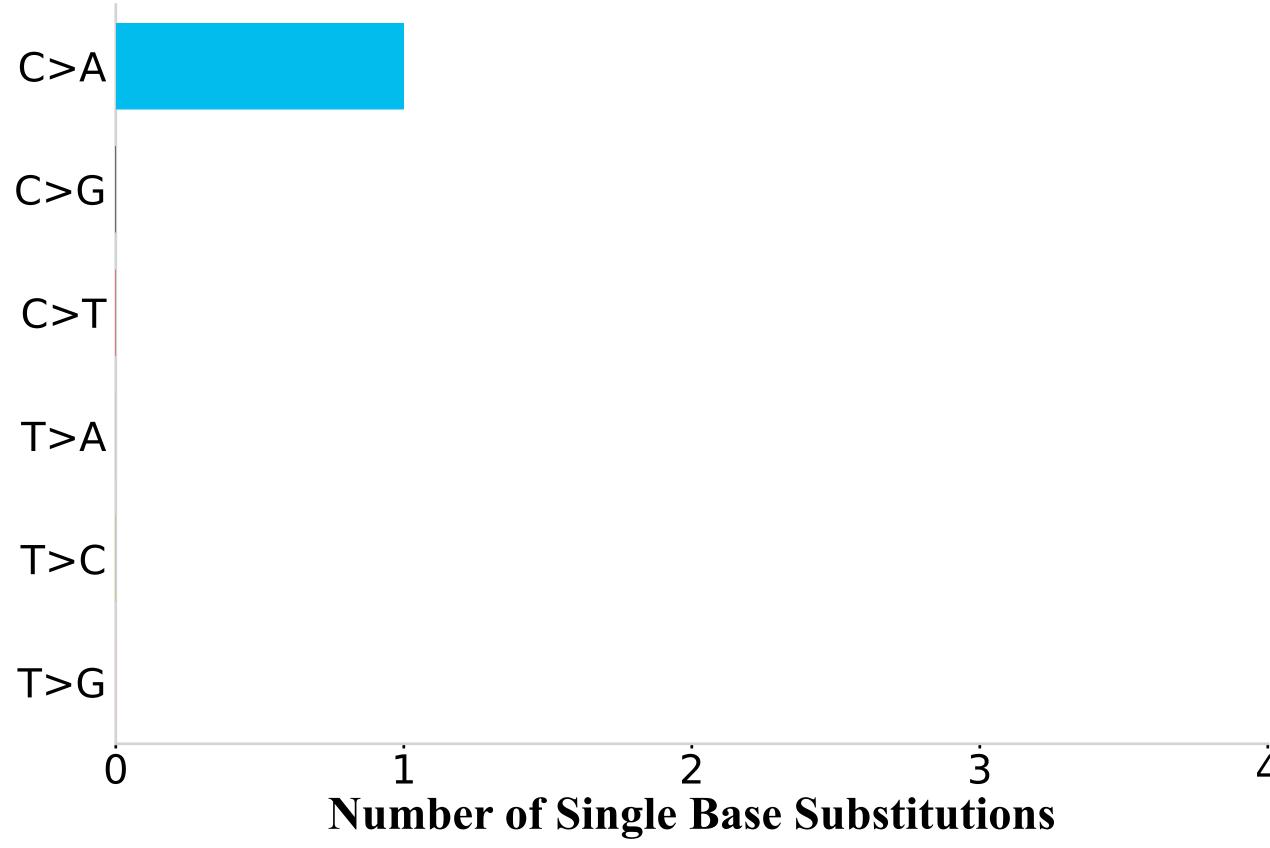


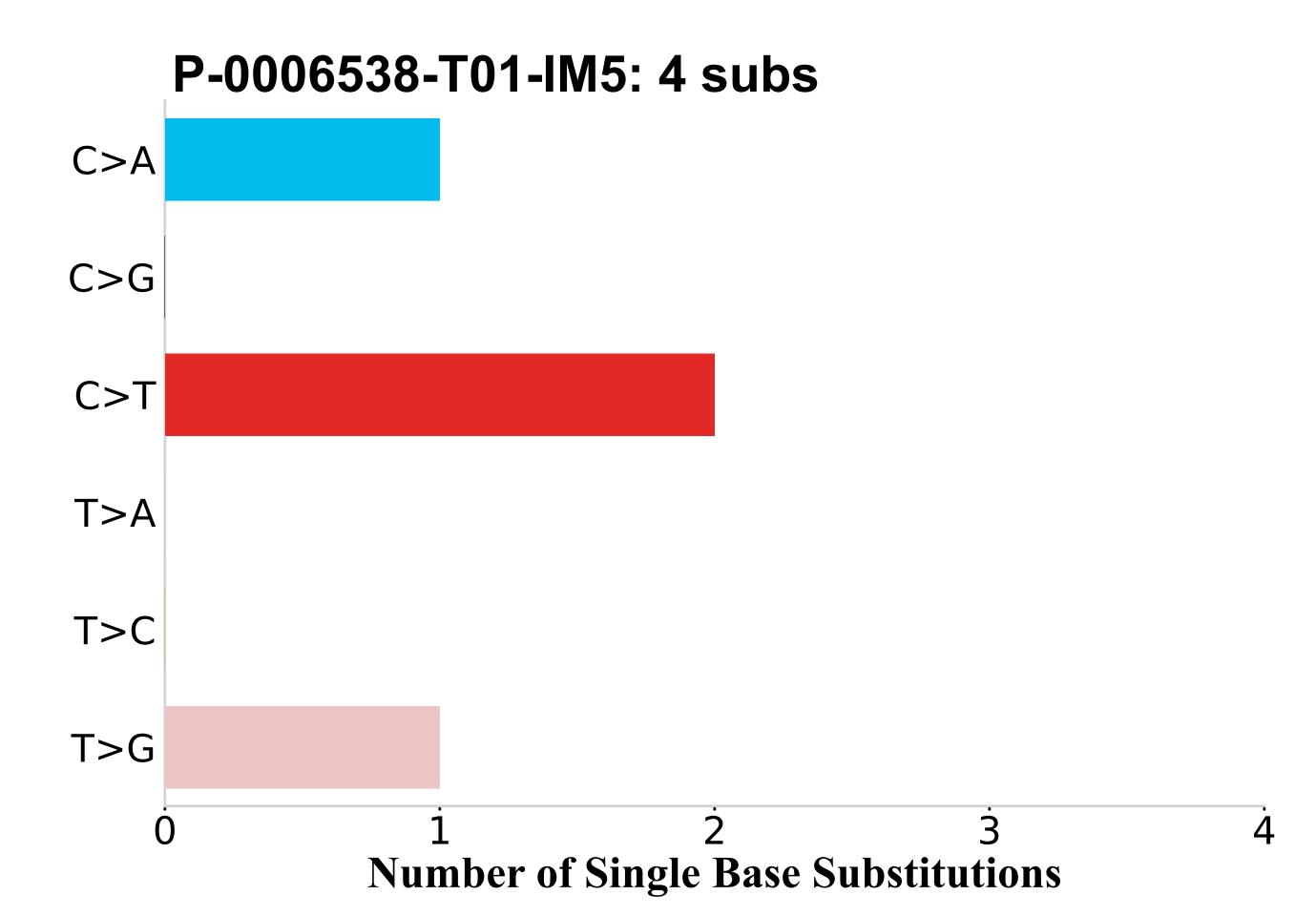
P-0006102-T01-IM5: 2 subs

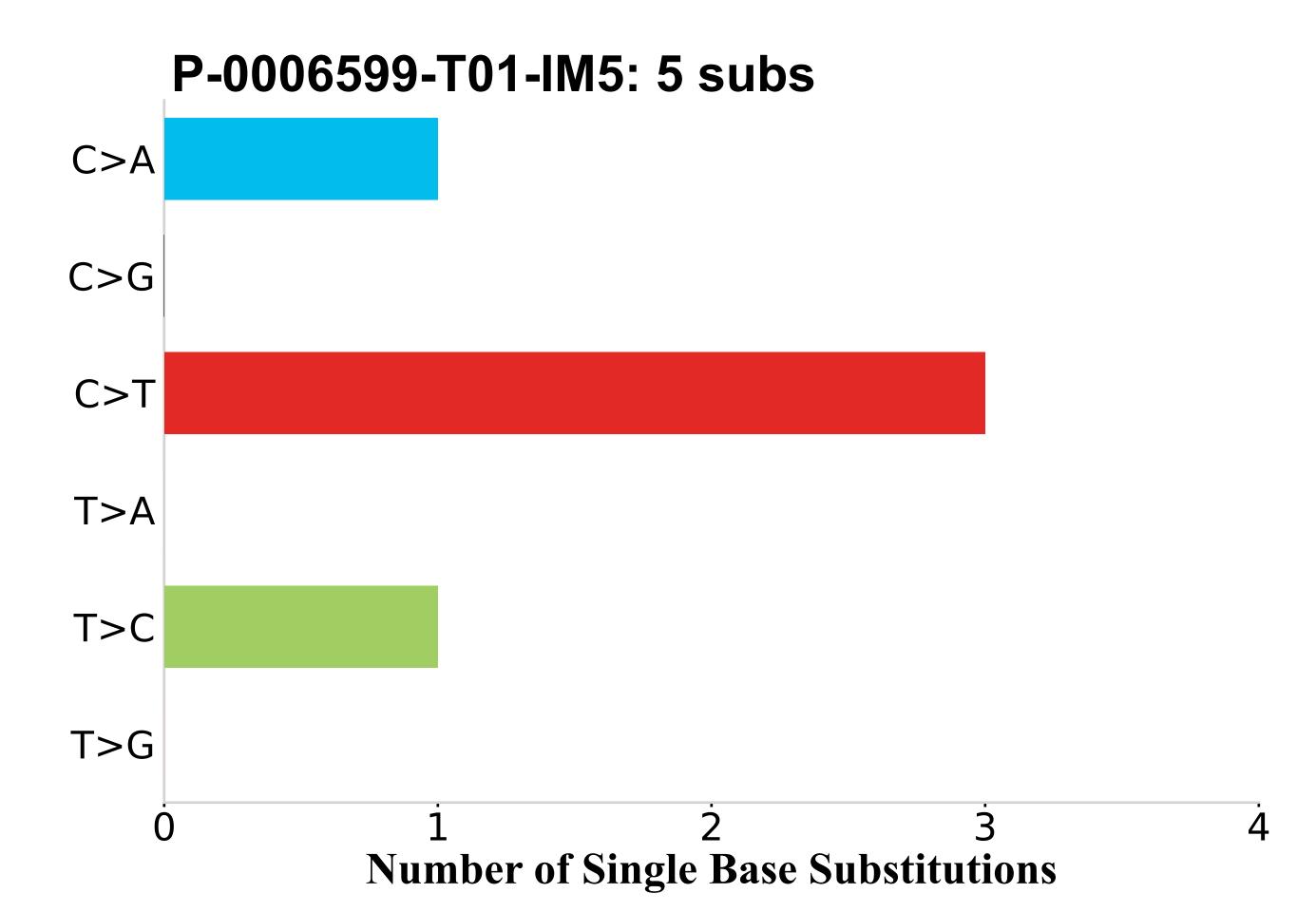


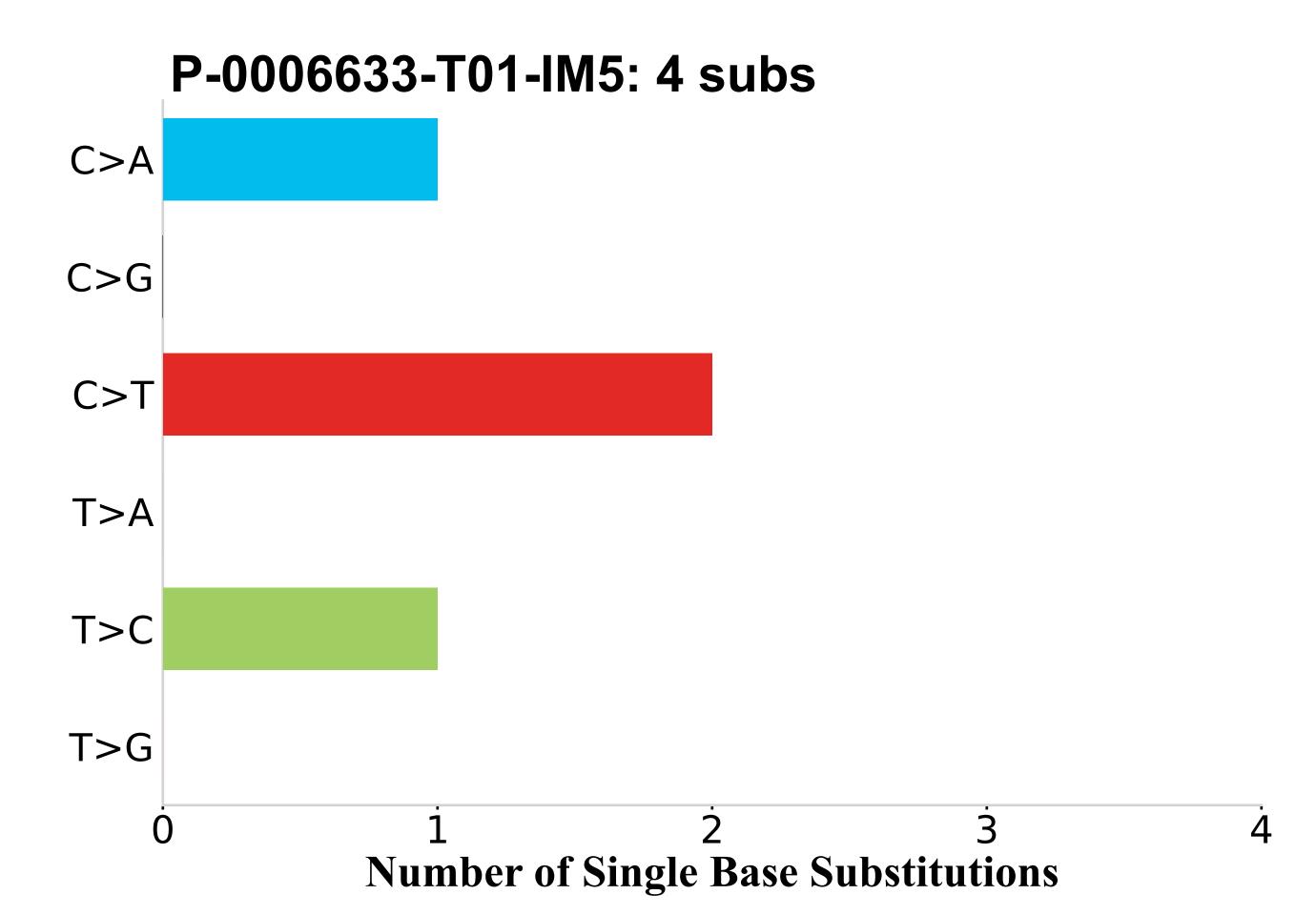


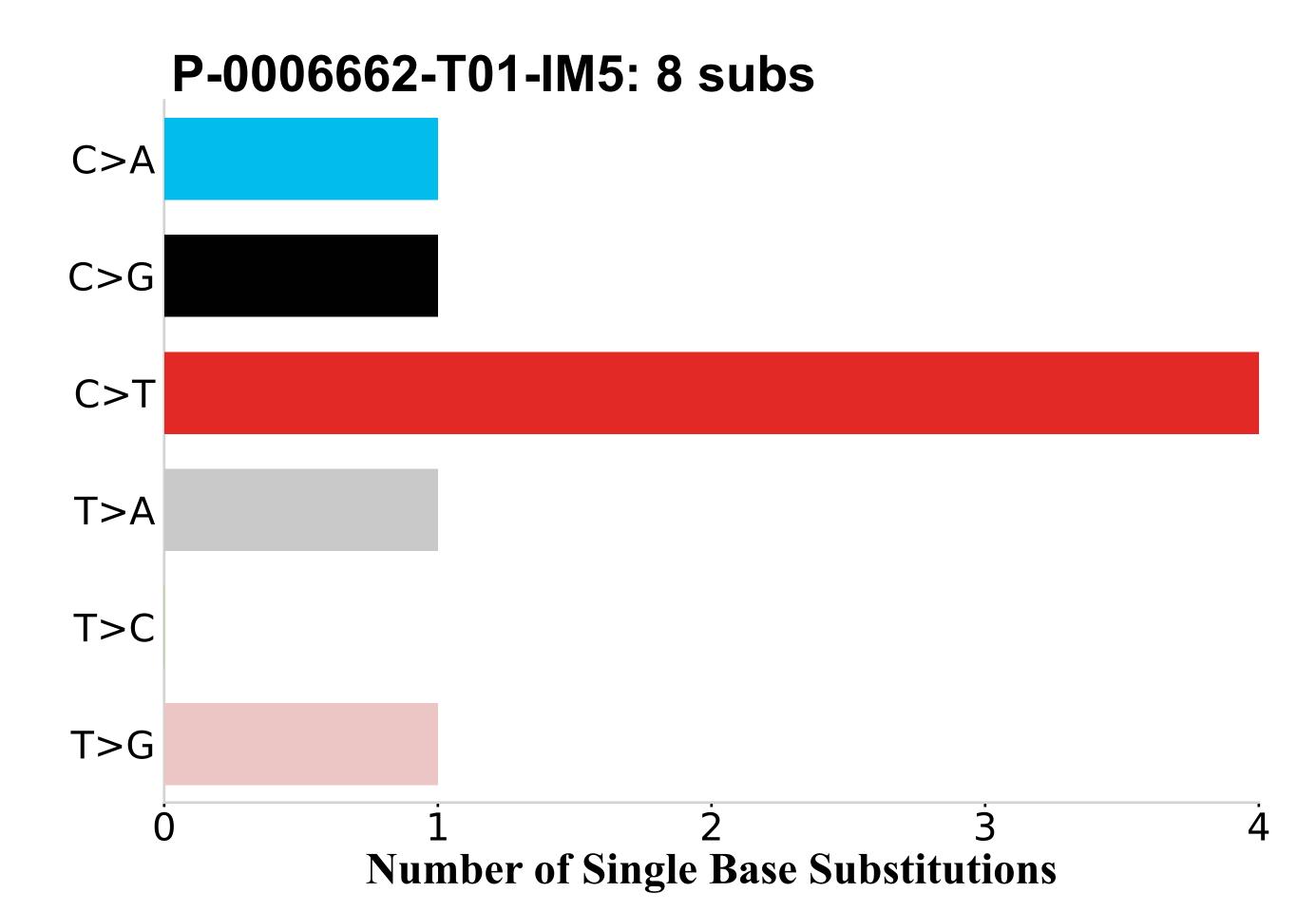


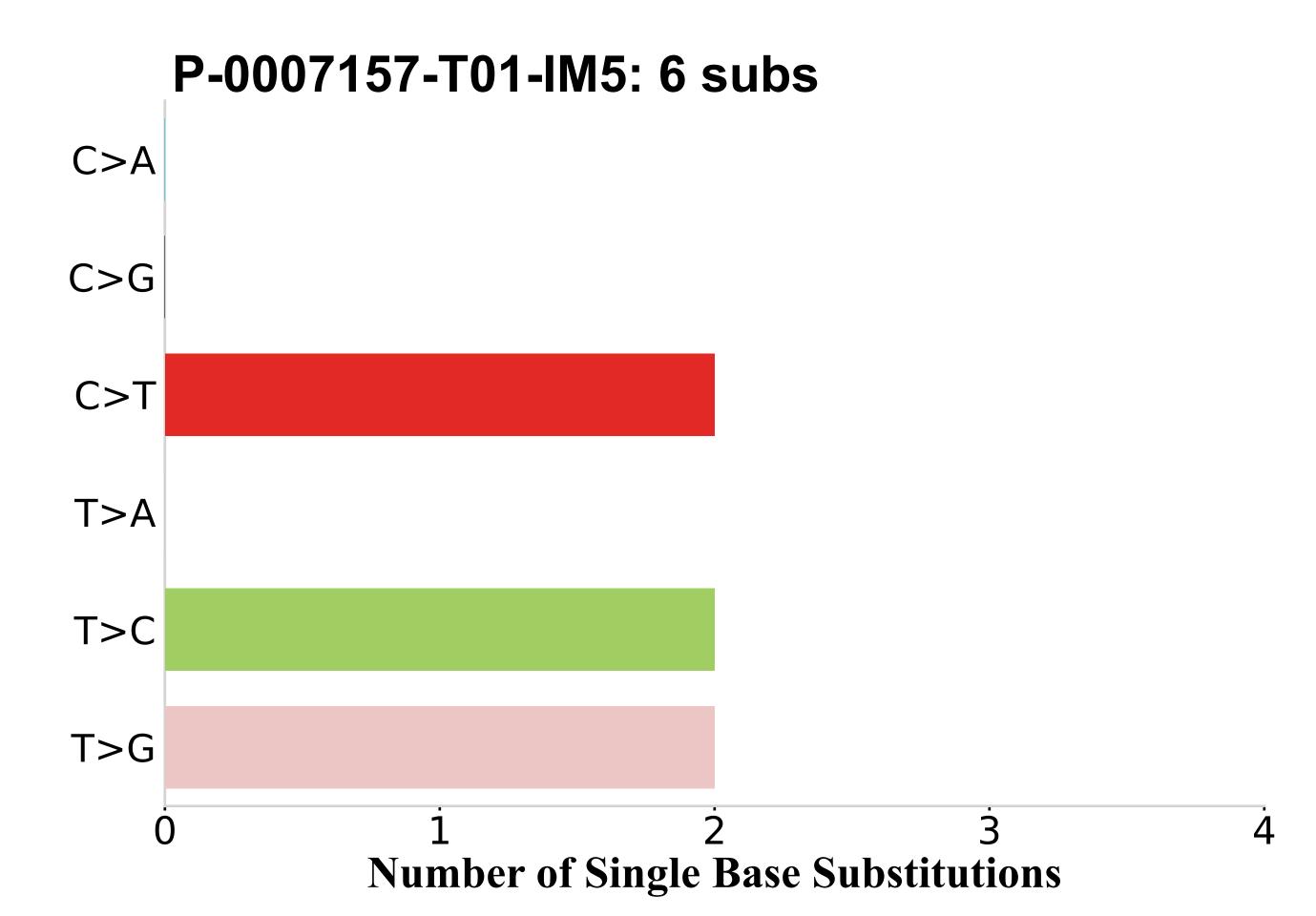


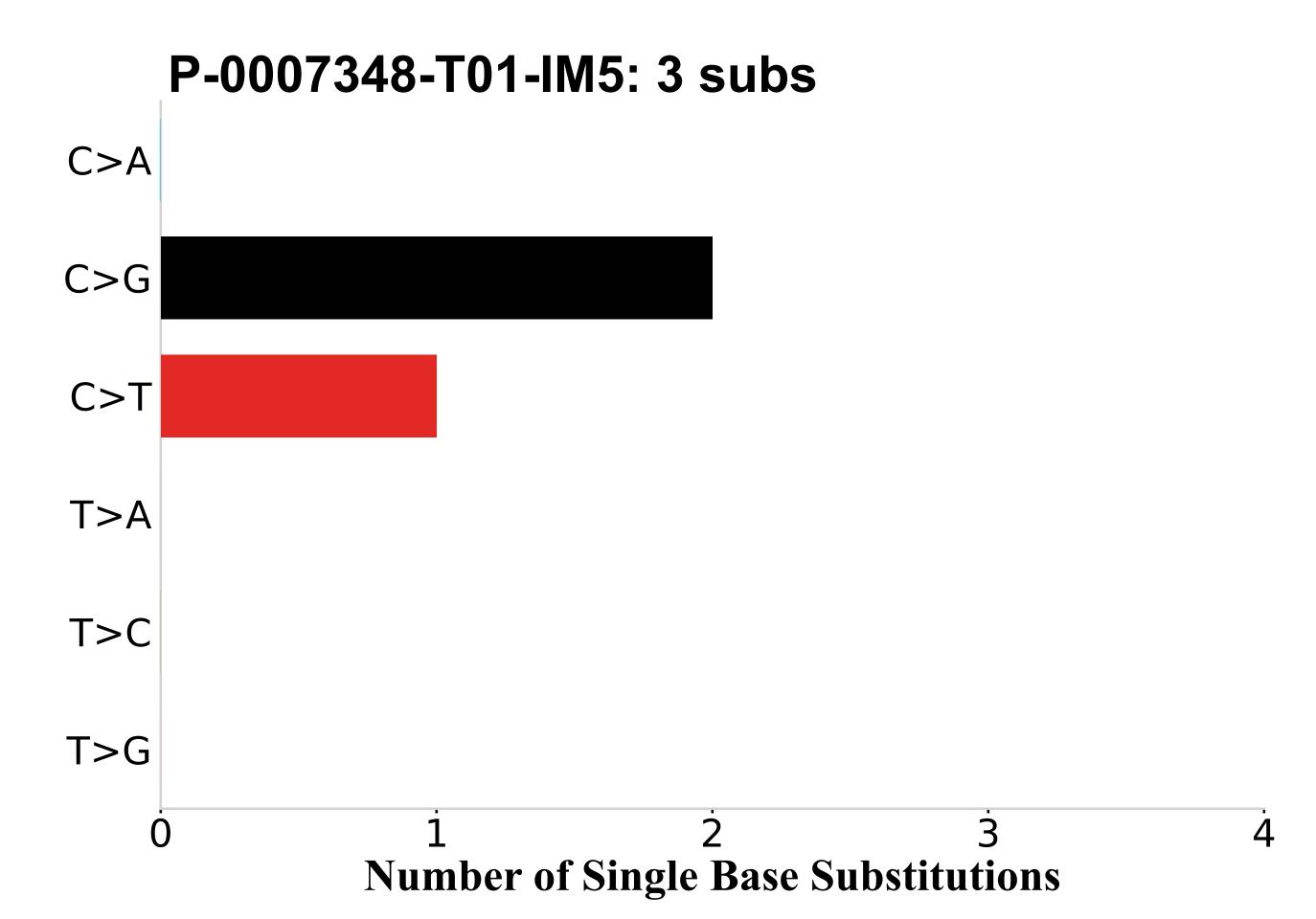


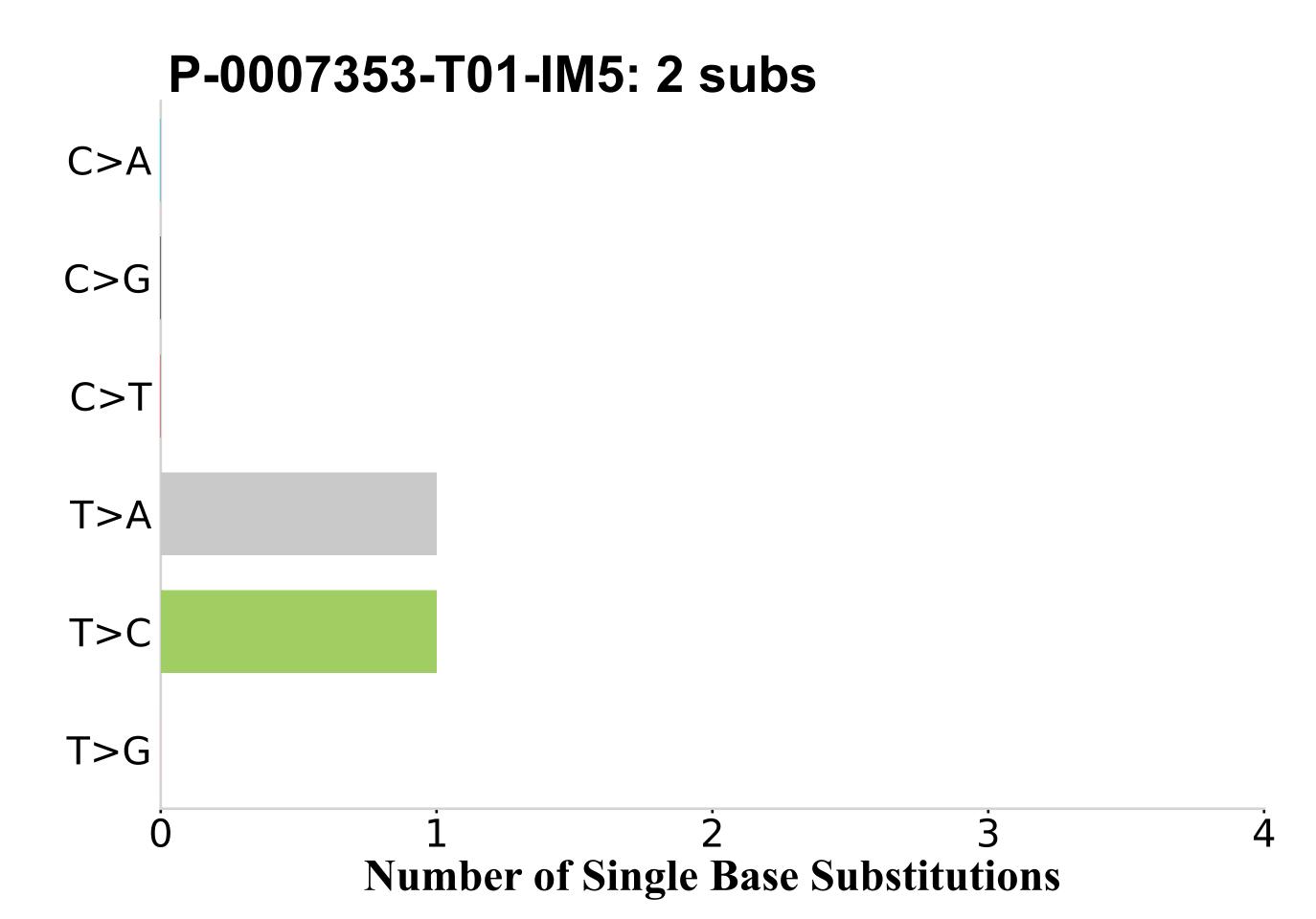


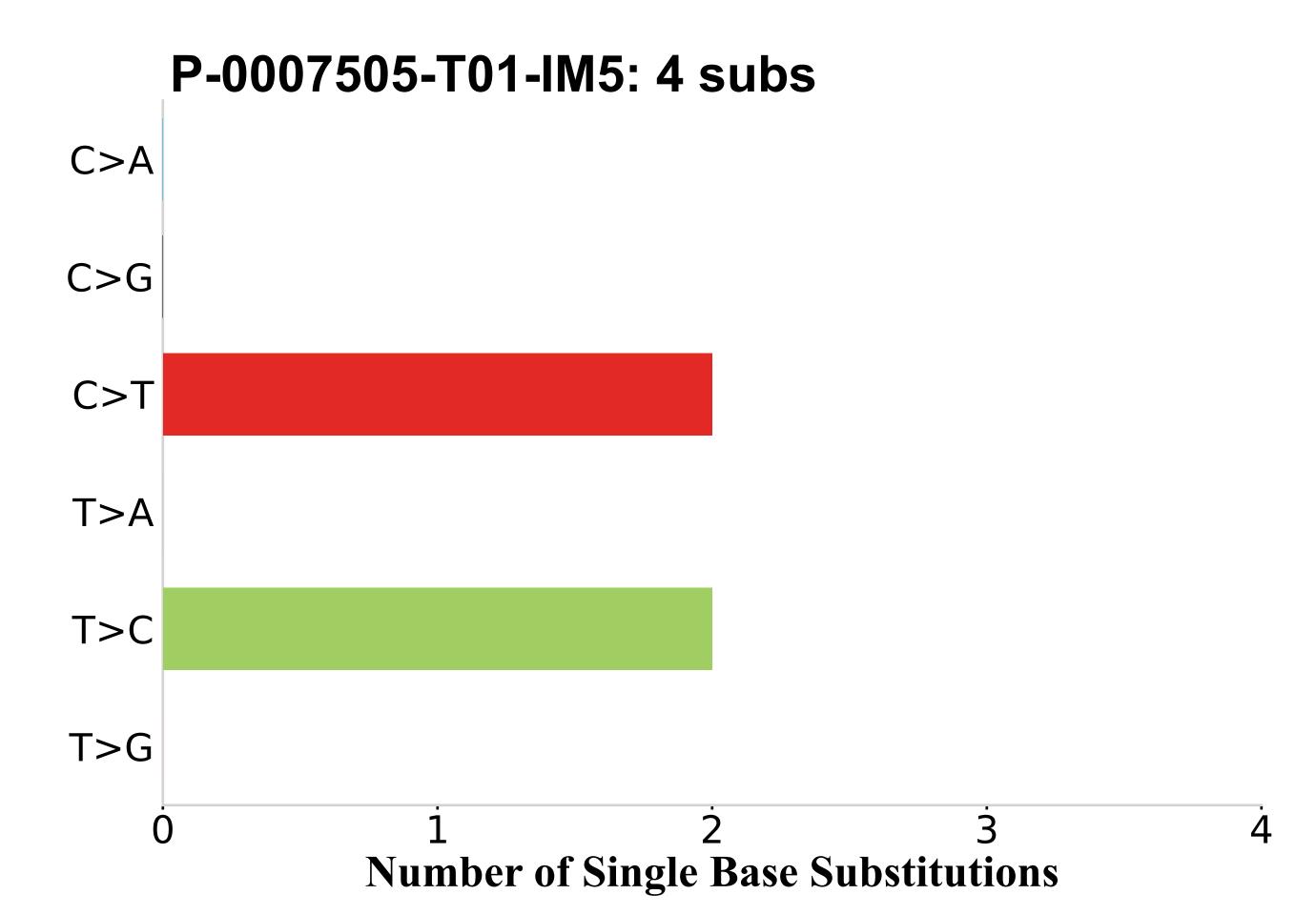




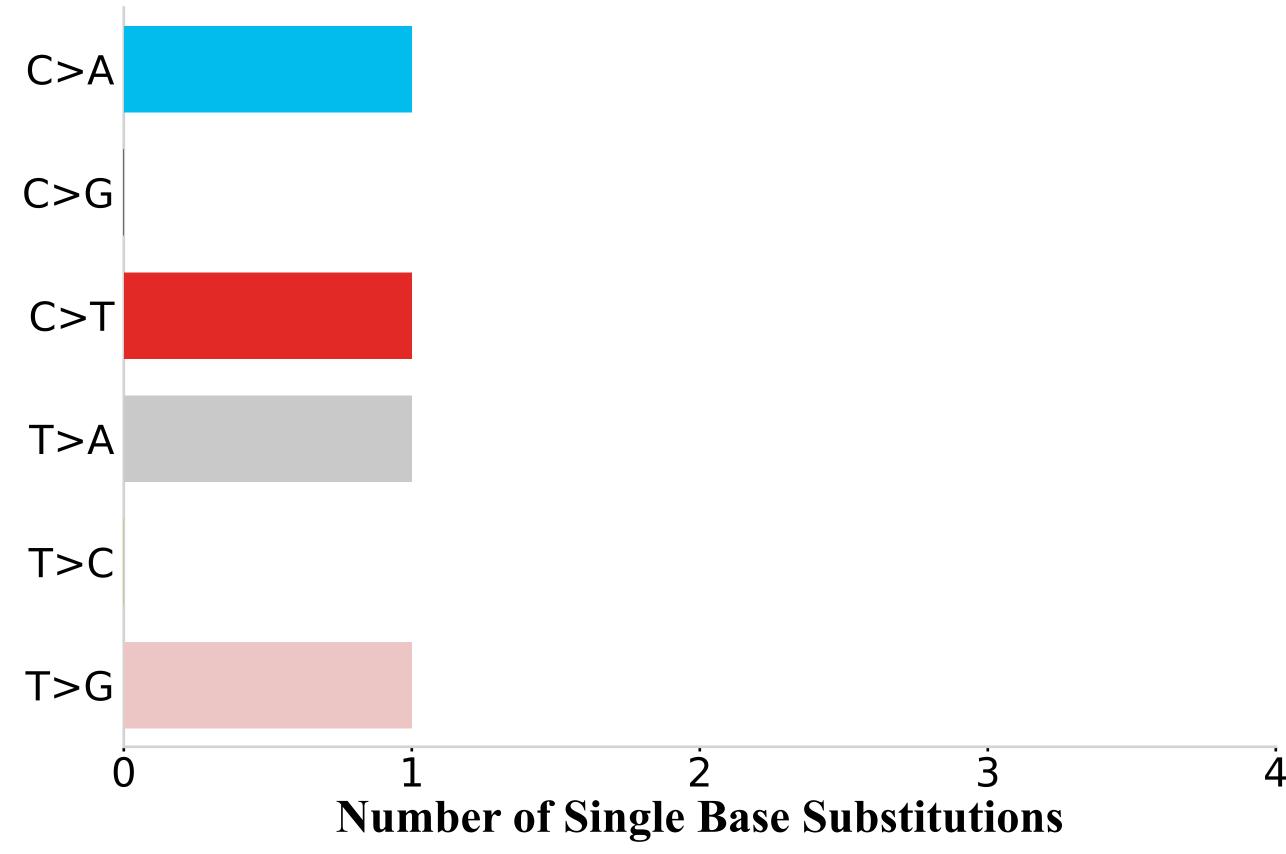


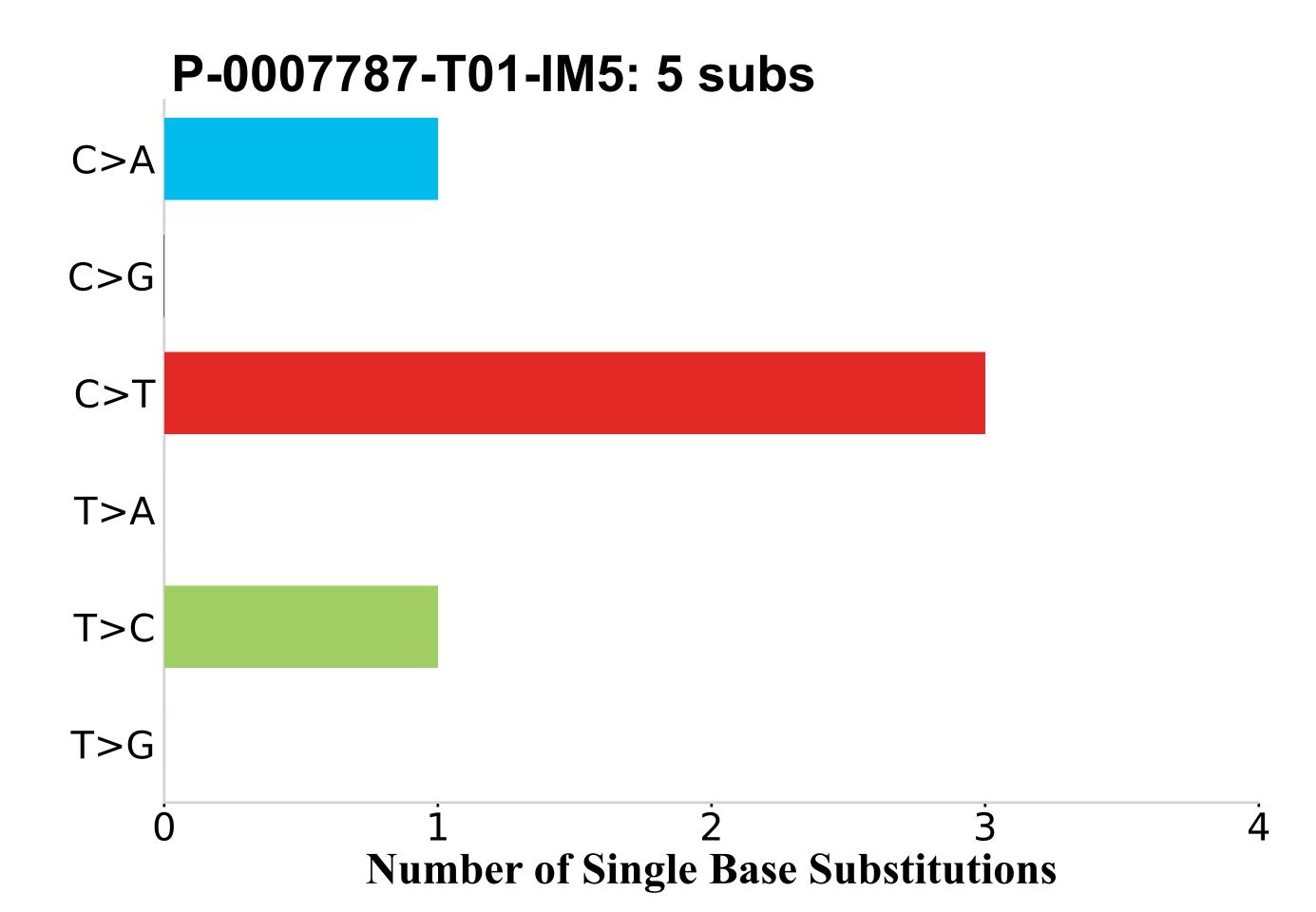


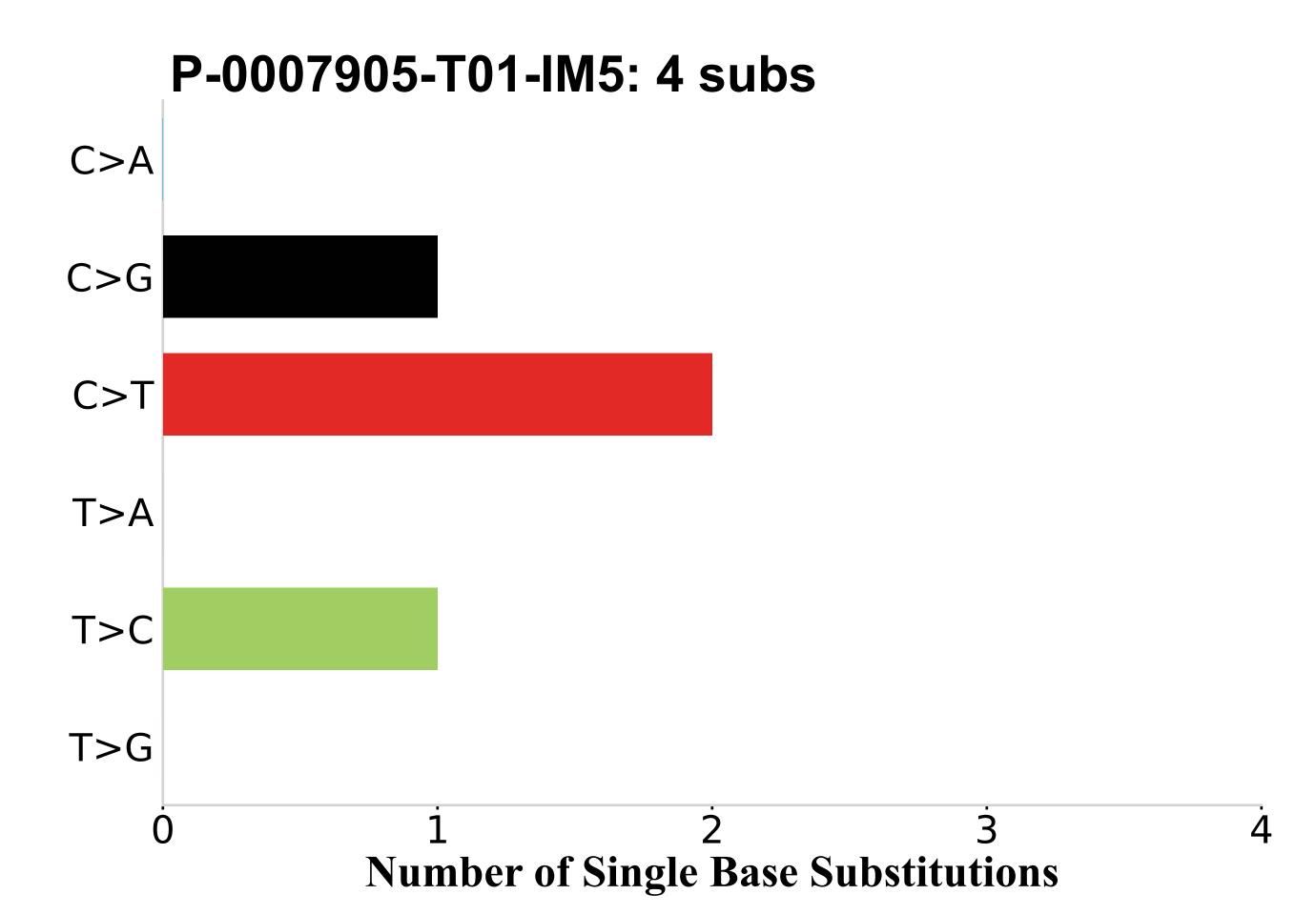


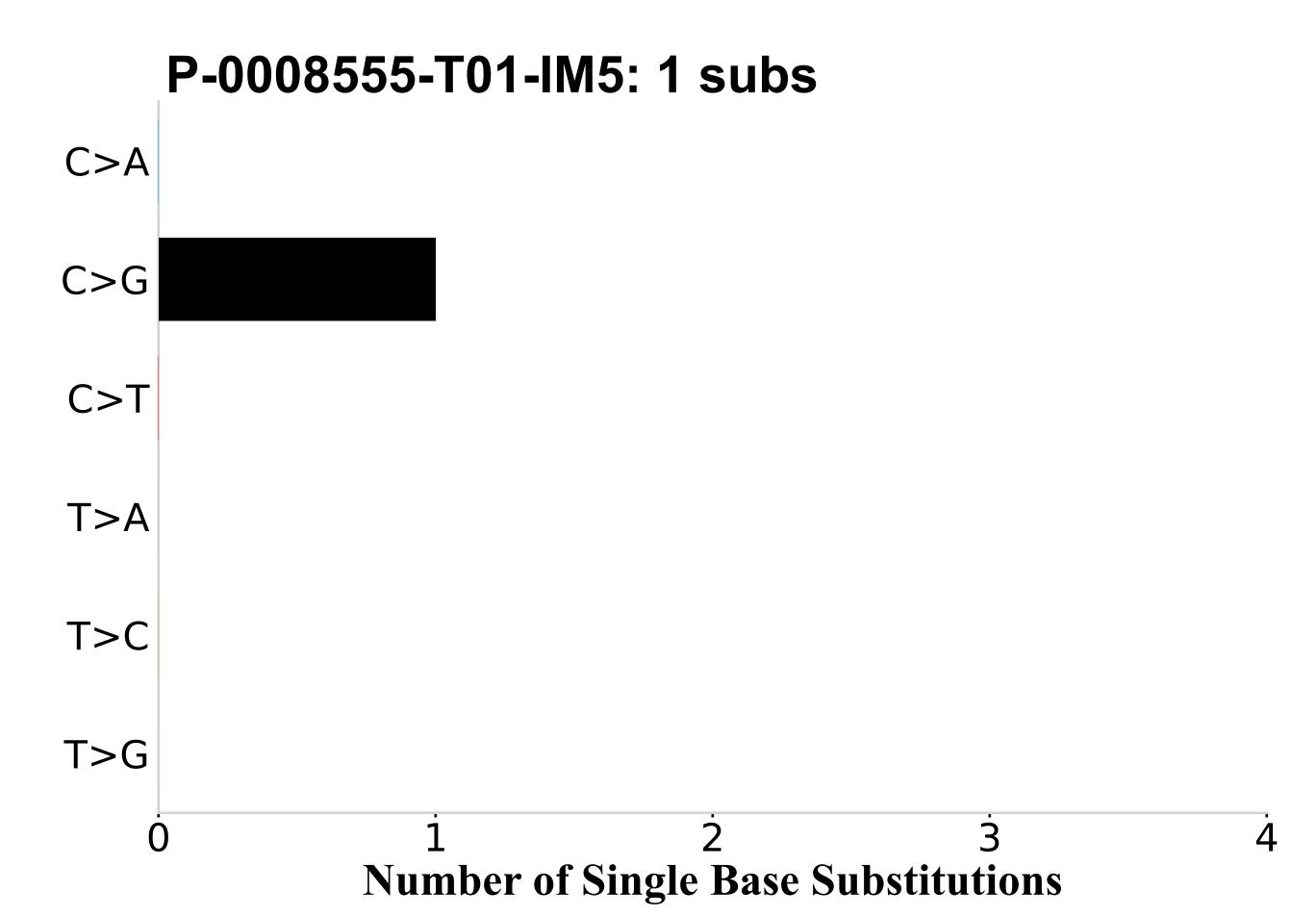


P-0007773-T01-IM5: 4 subs

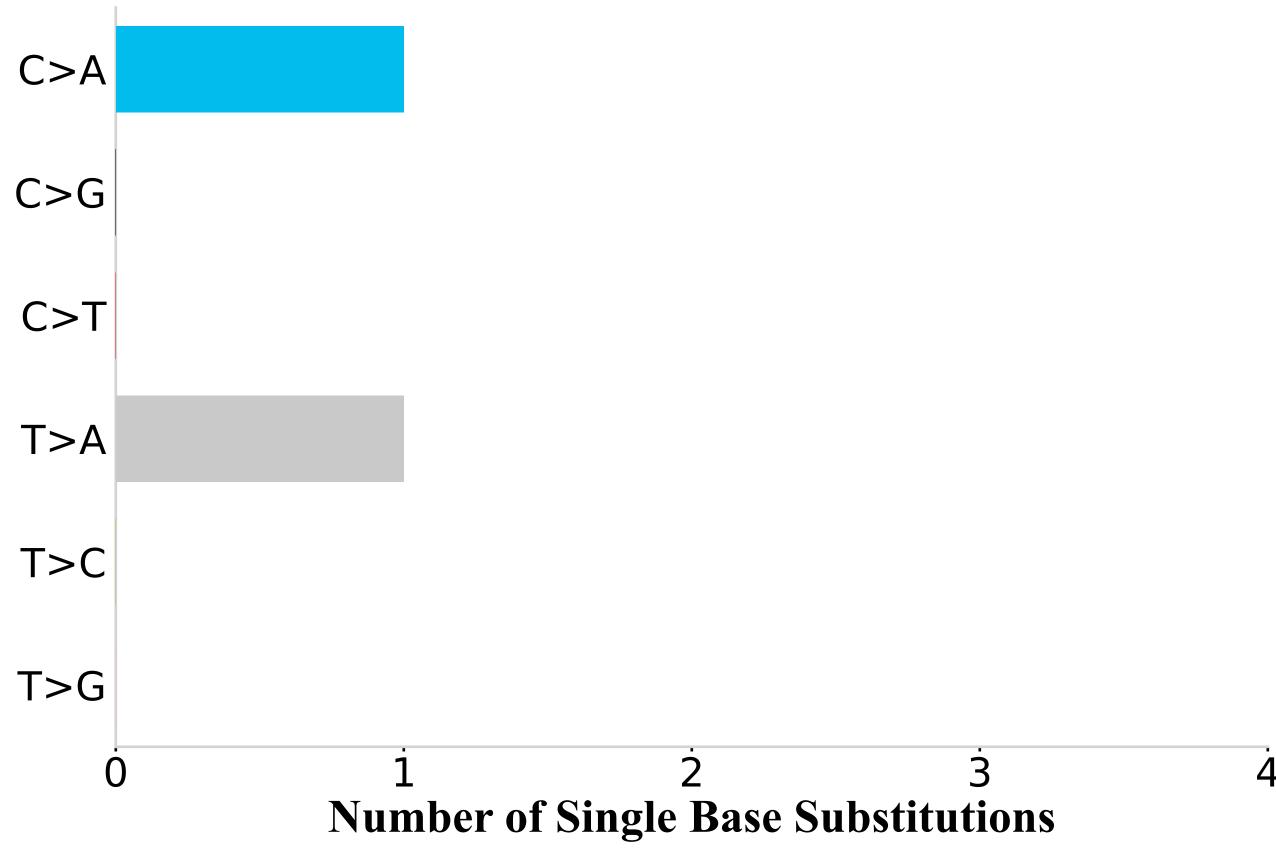




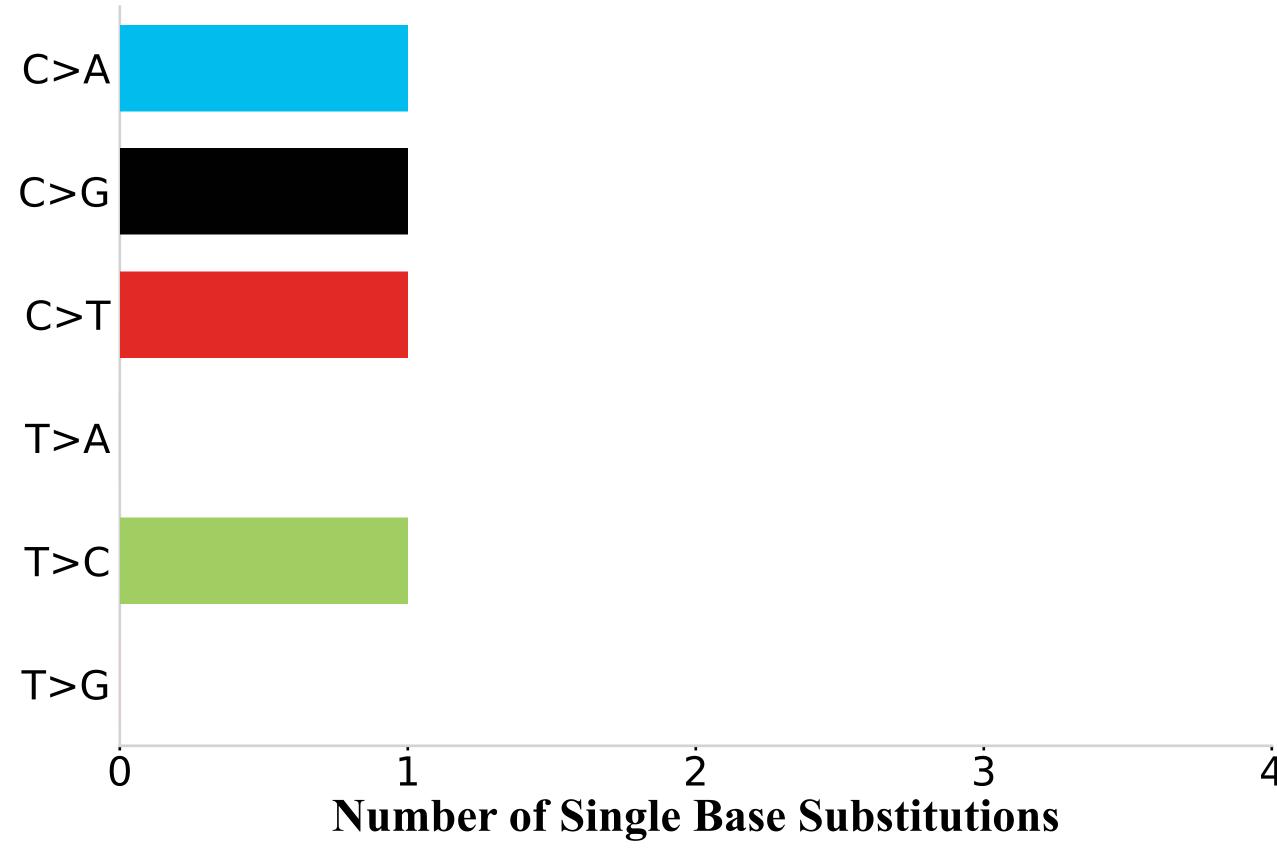


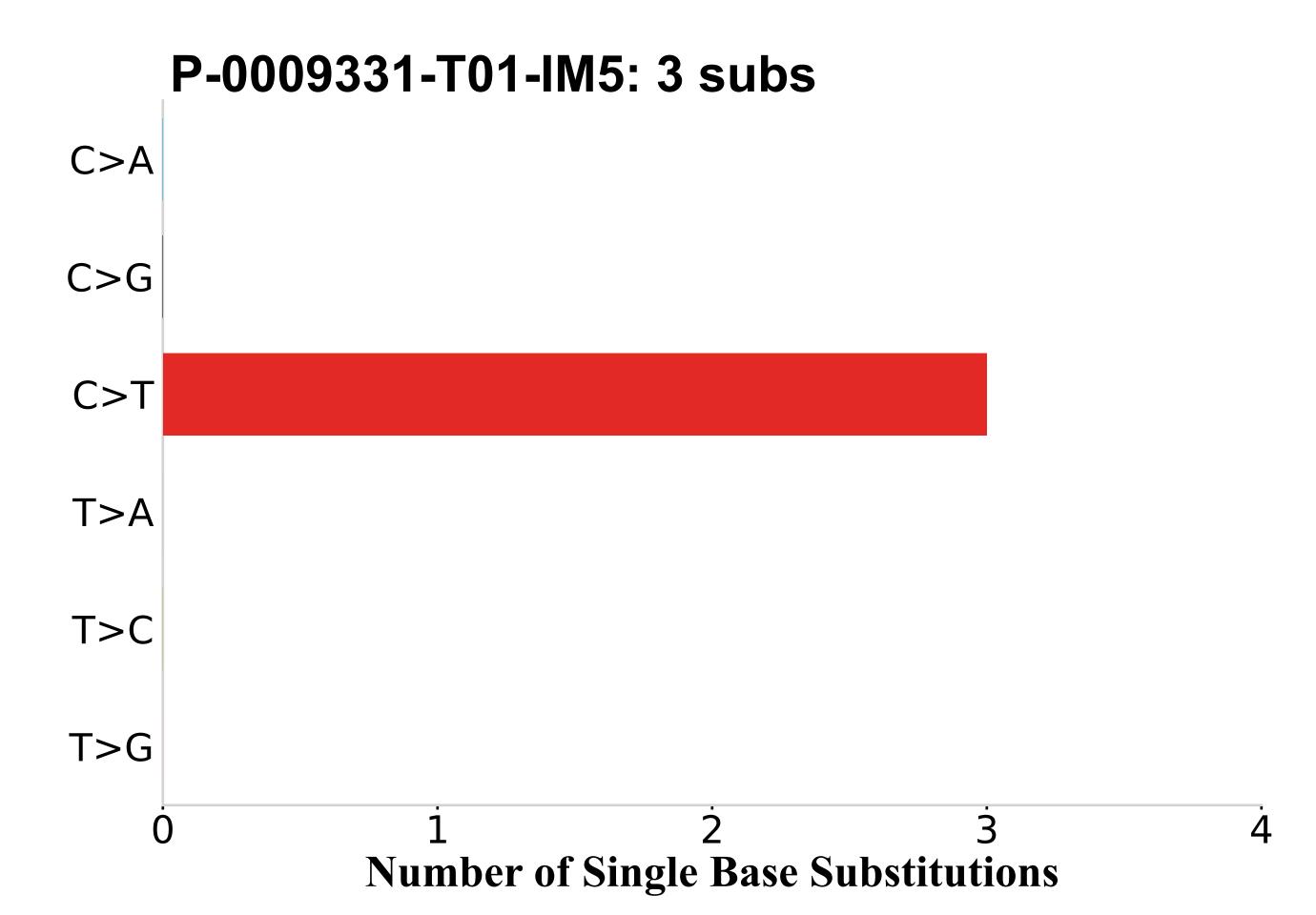


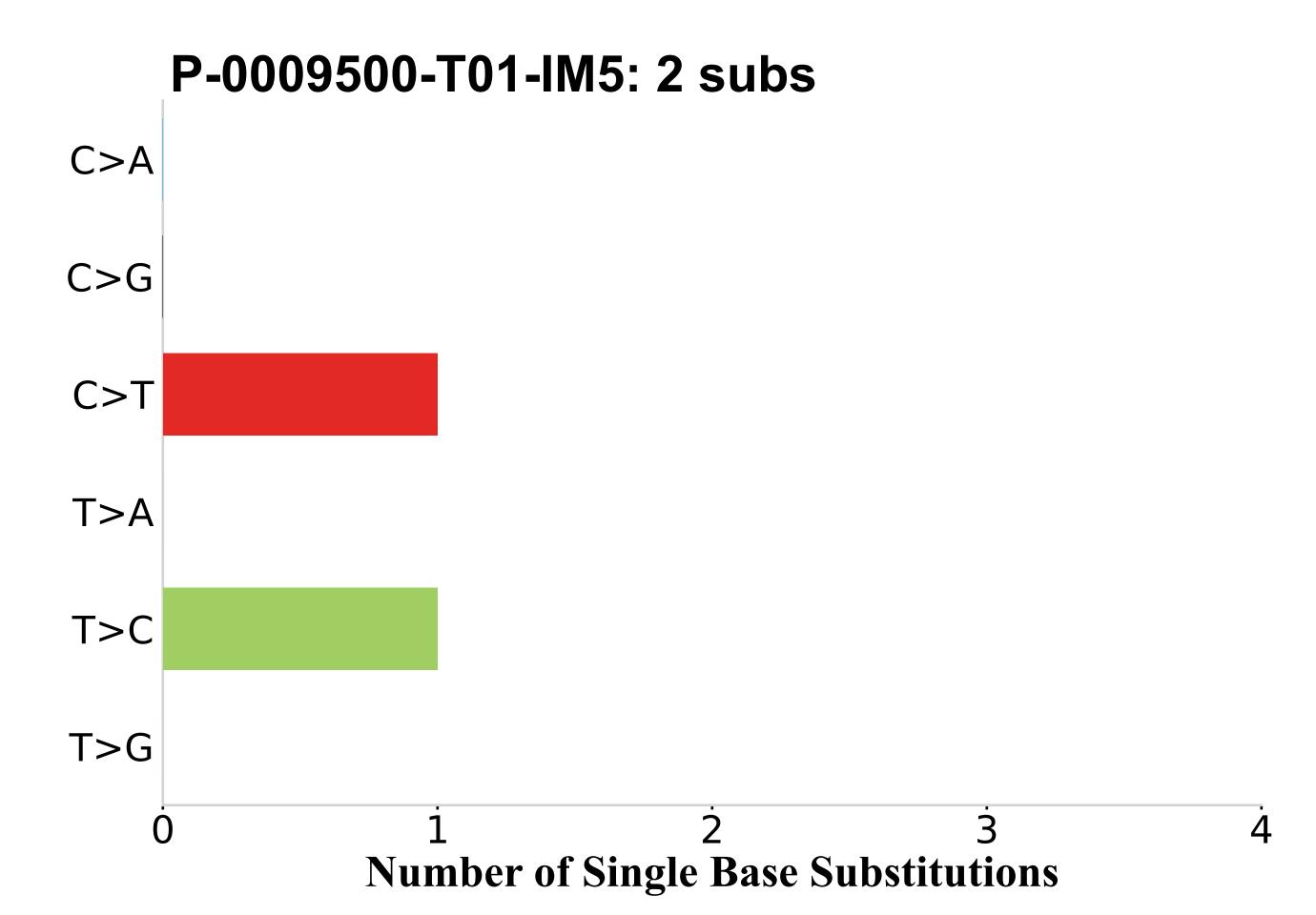
P-0008564-T01-IM5: 2 subs

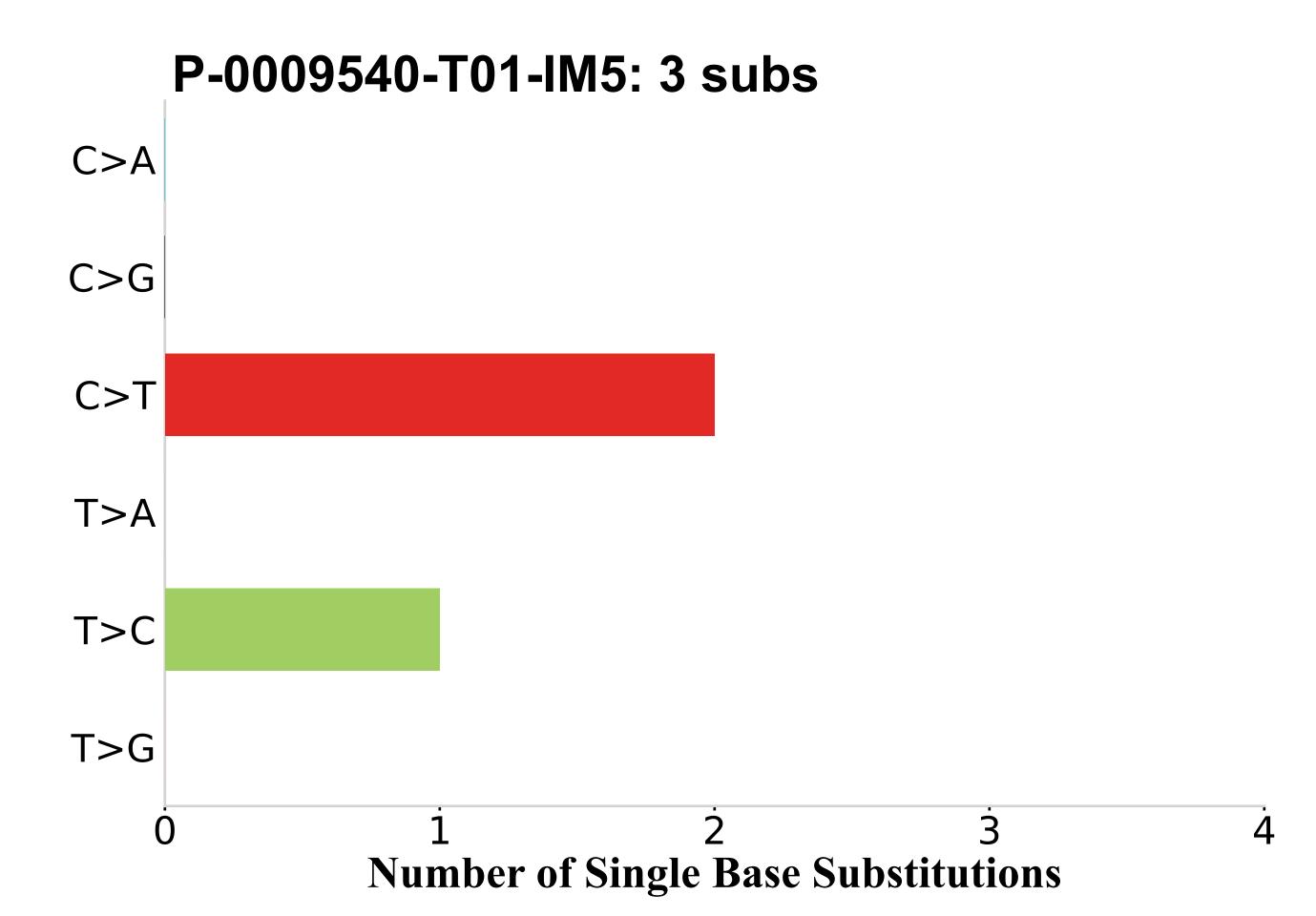


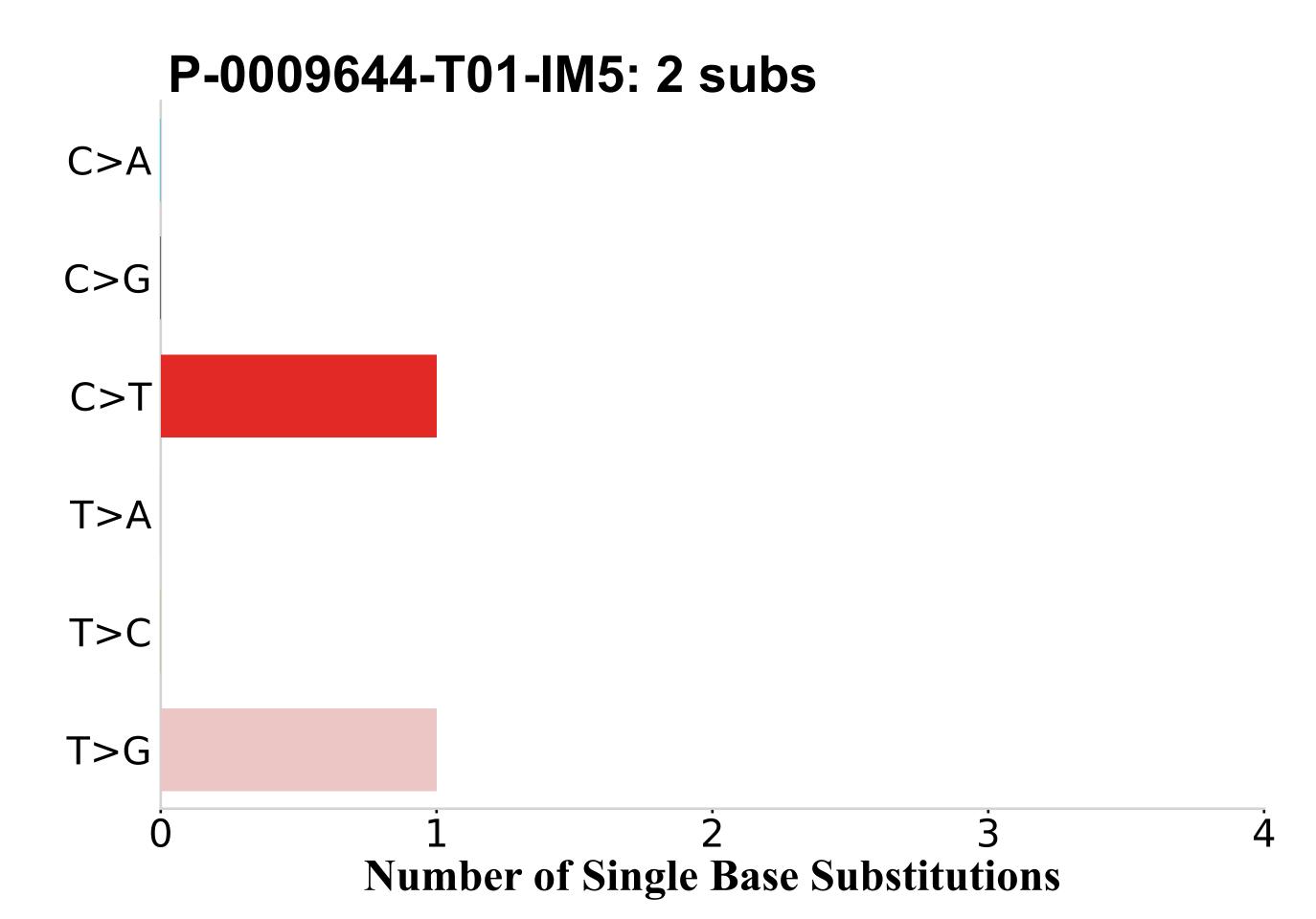


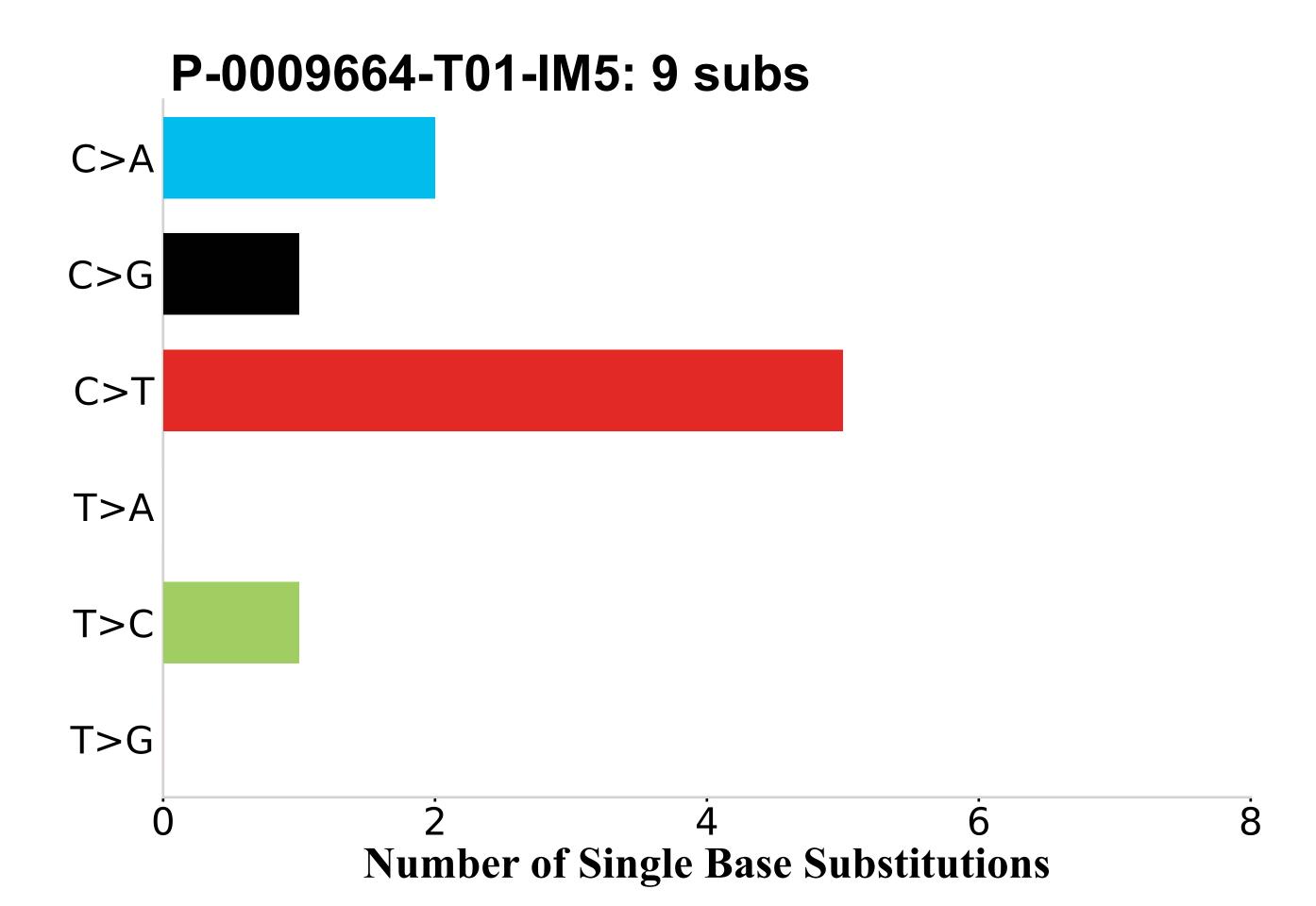


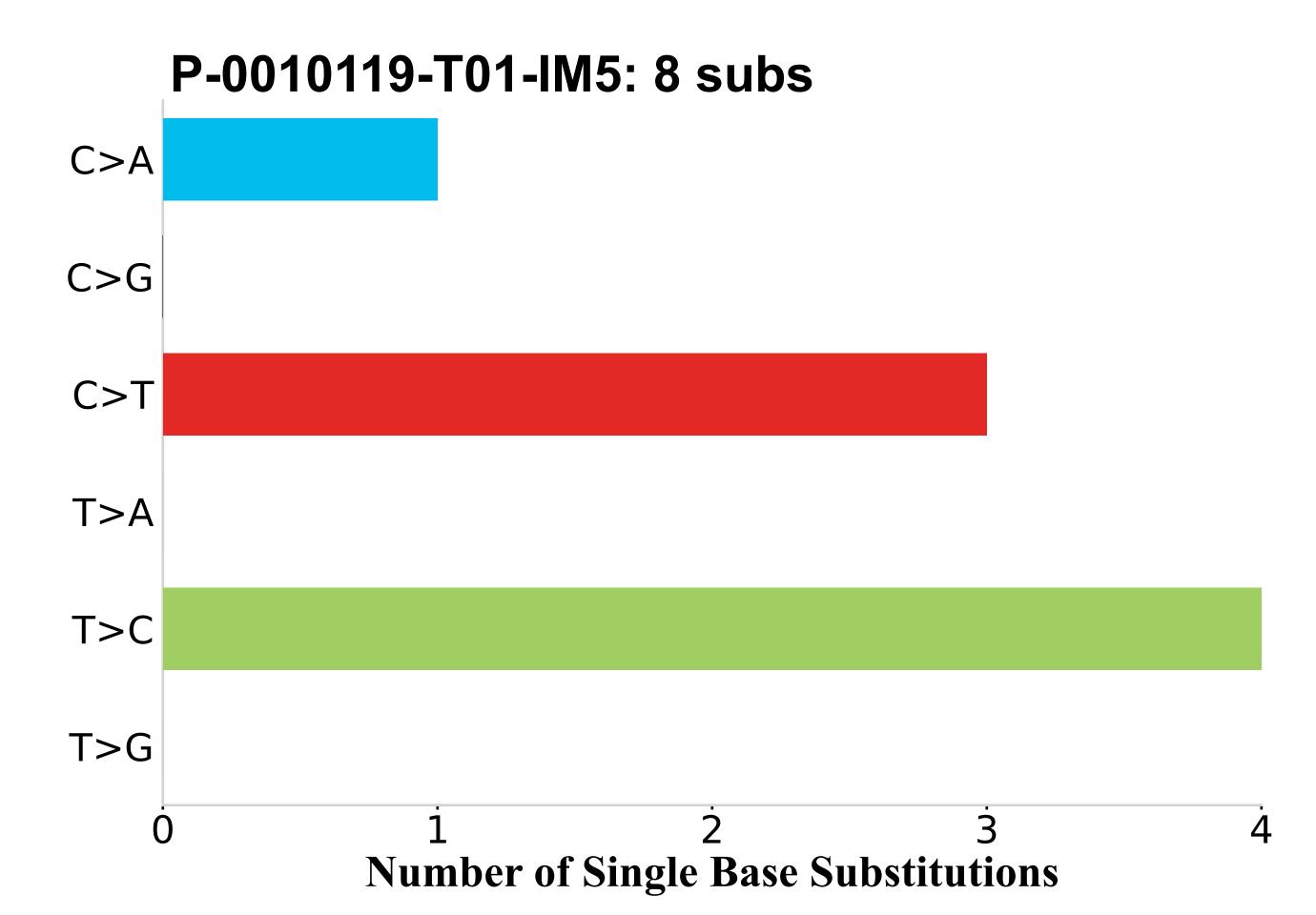


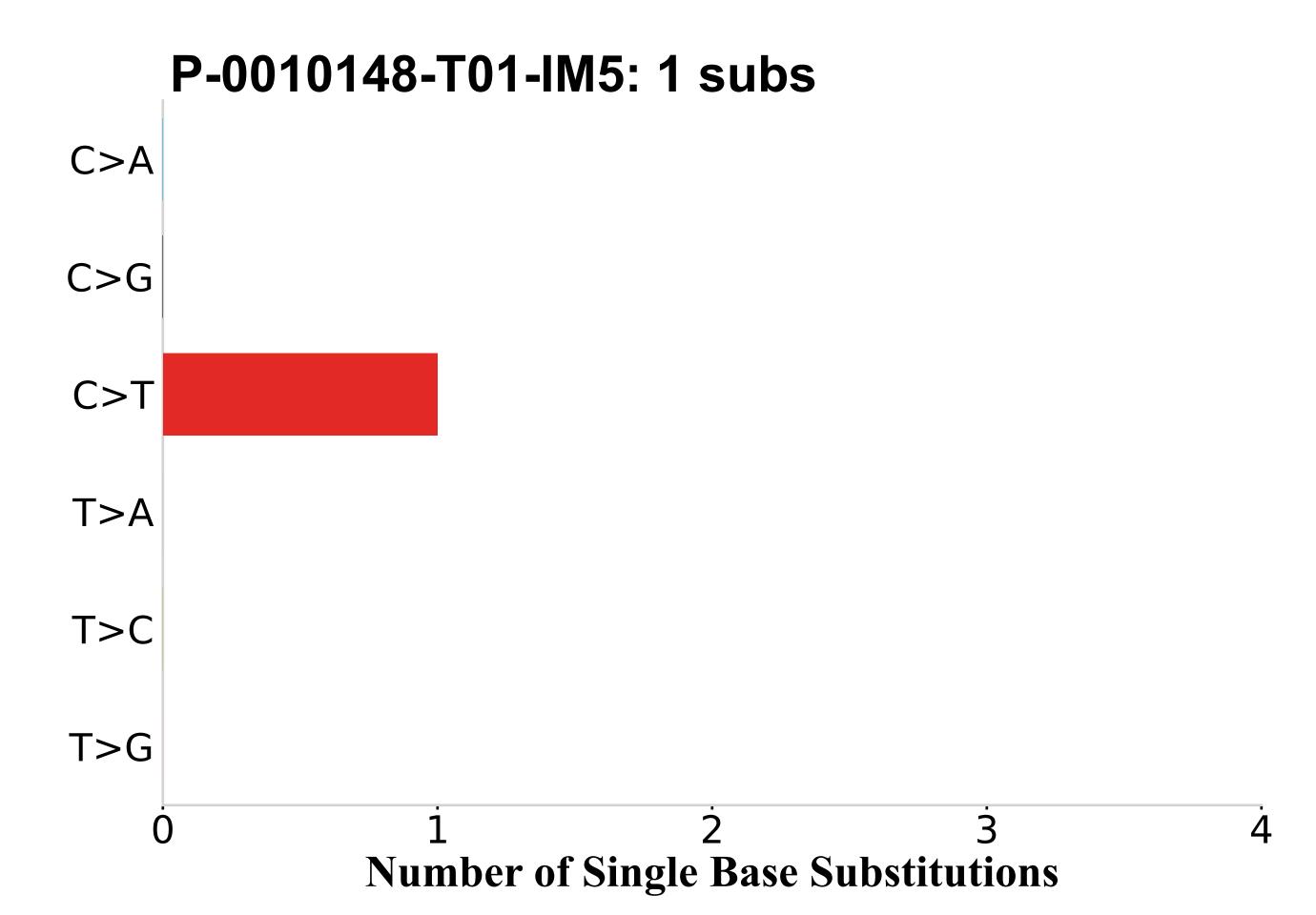




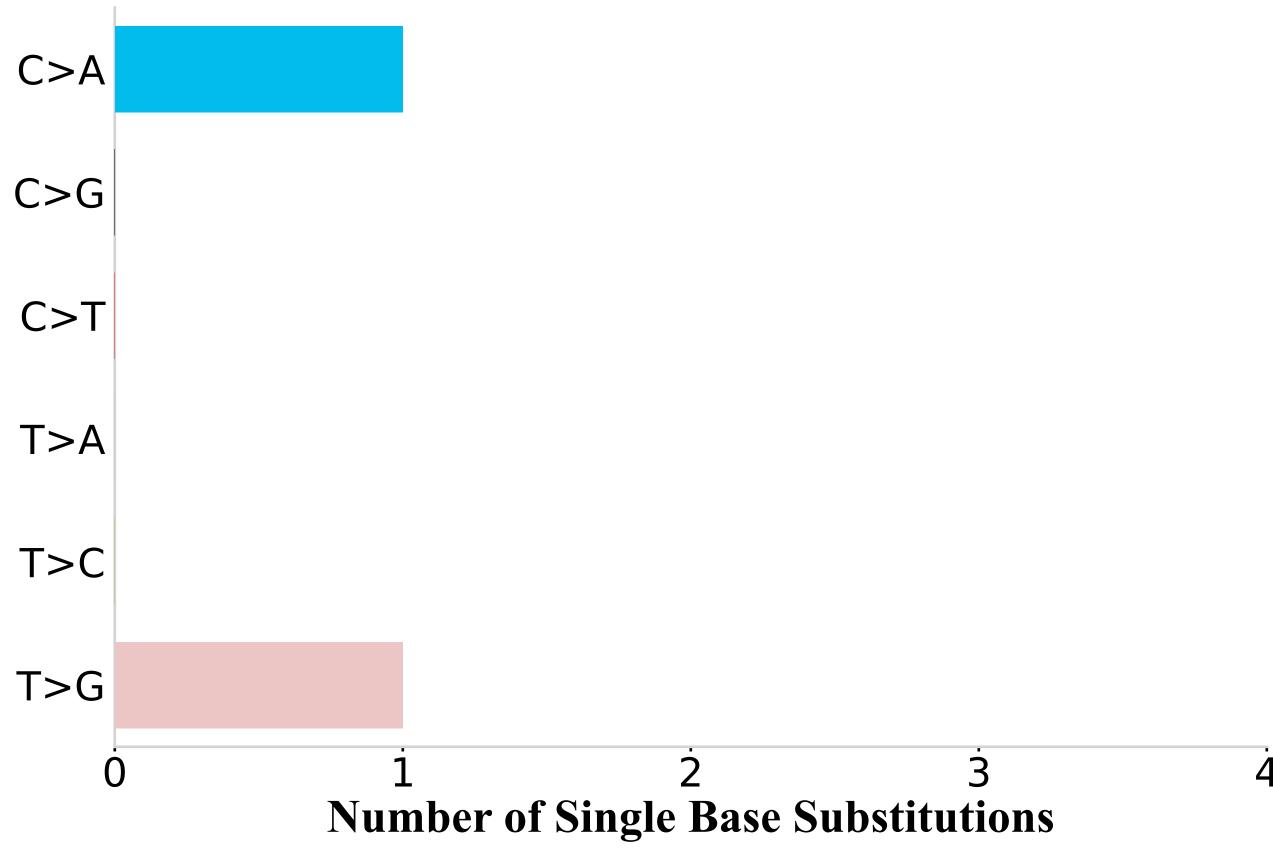


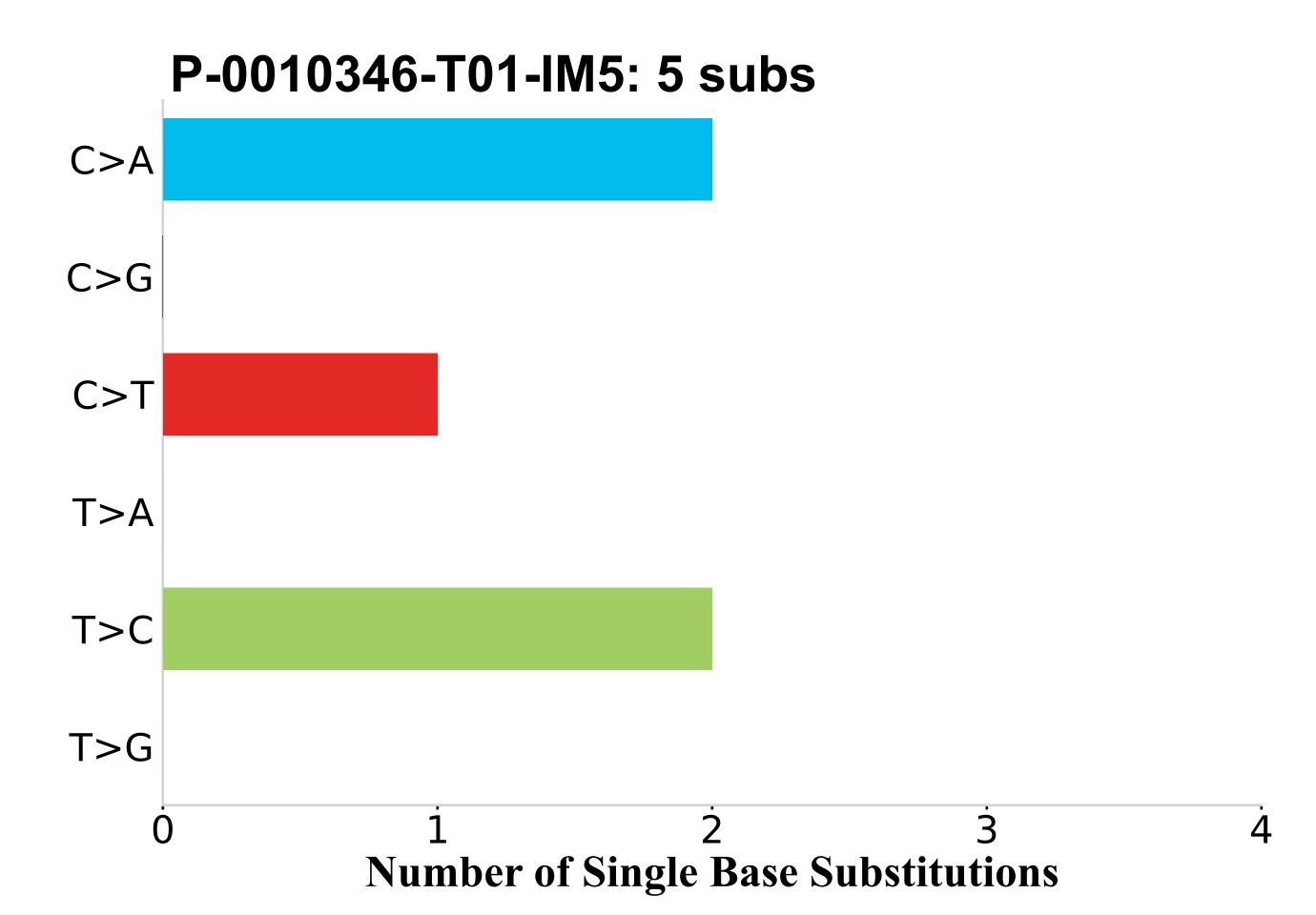


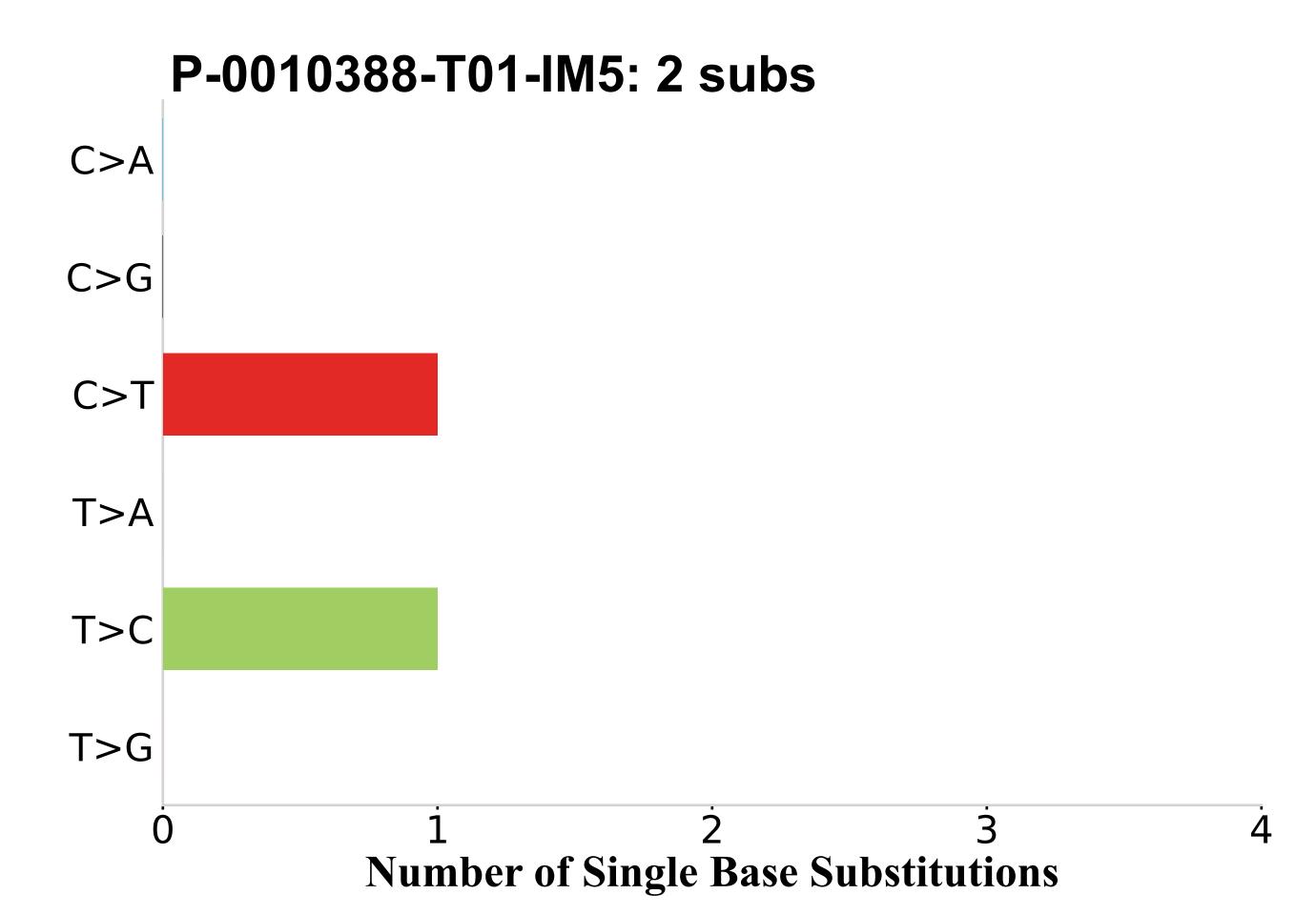


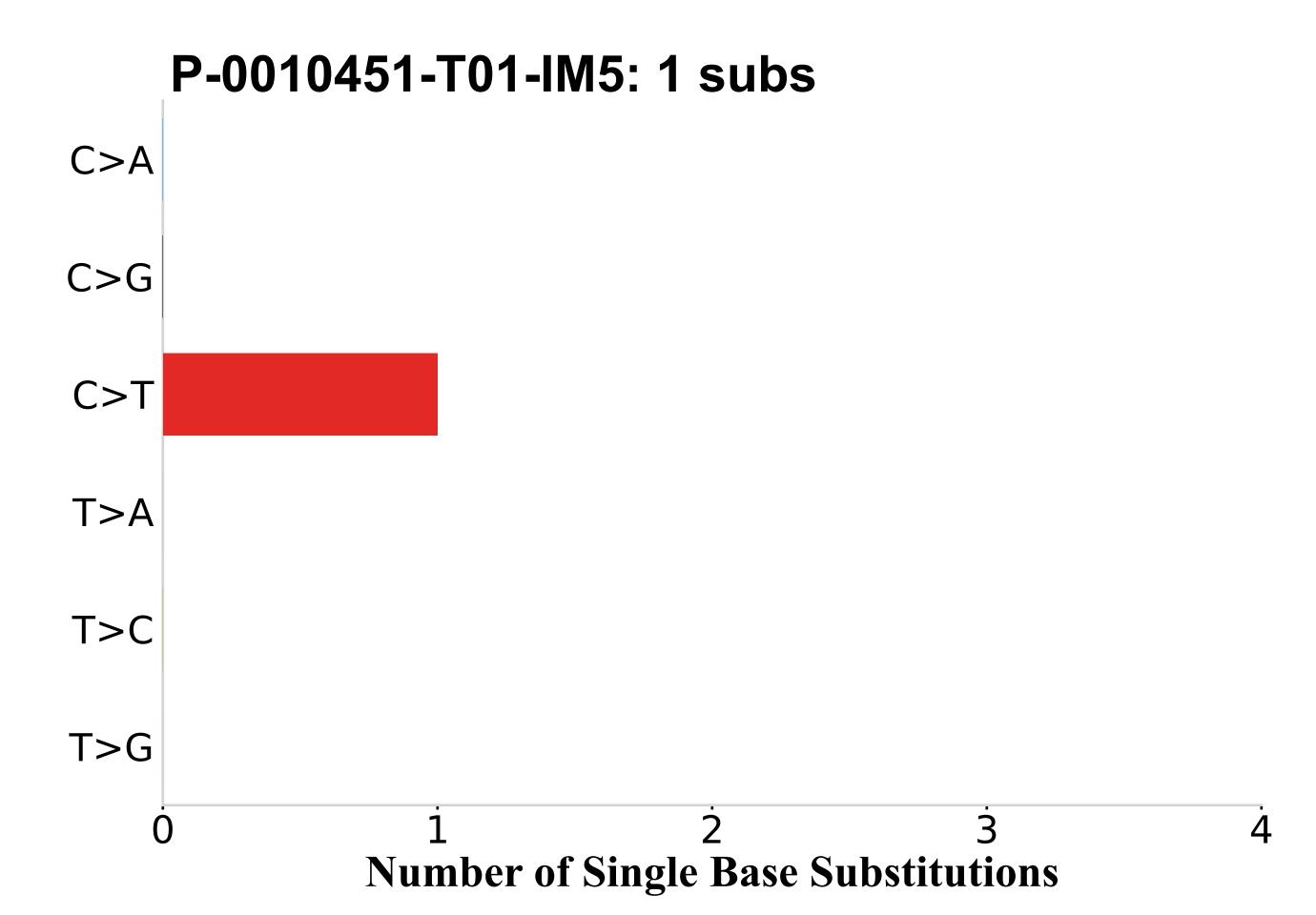


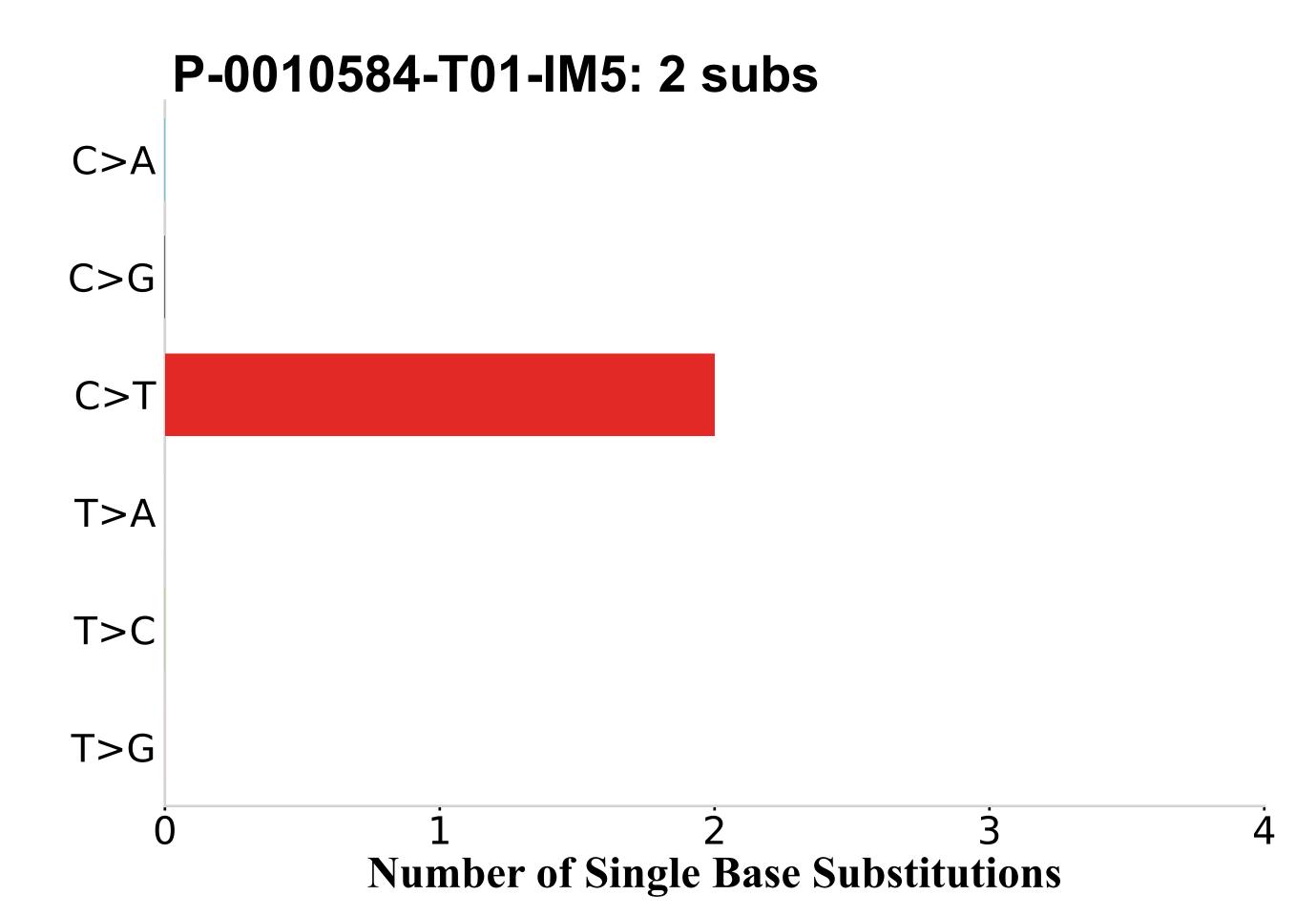
P-0010226-T01-IM5: 2 subs

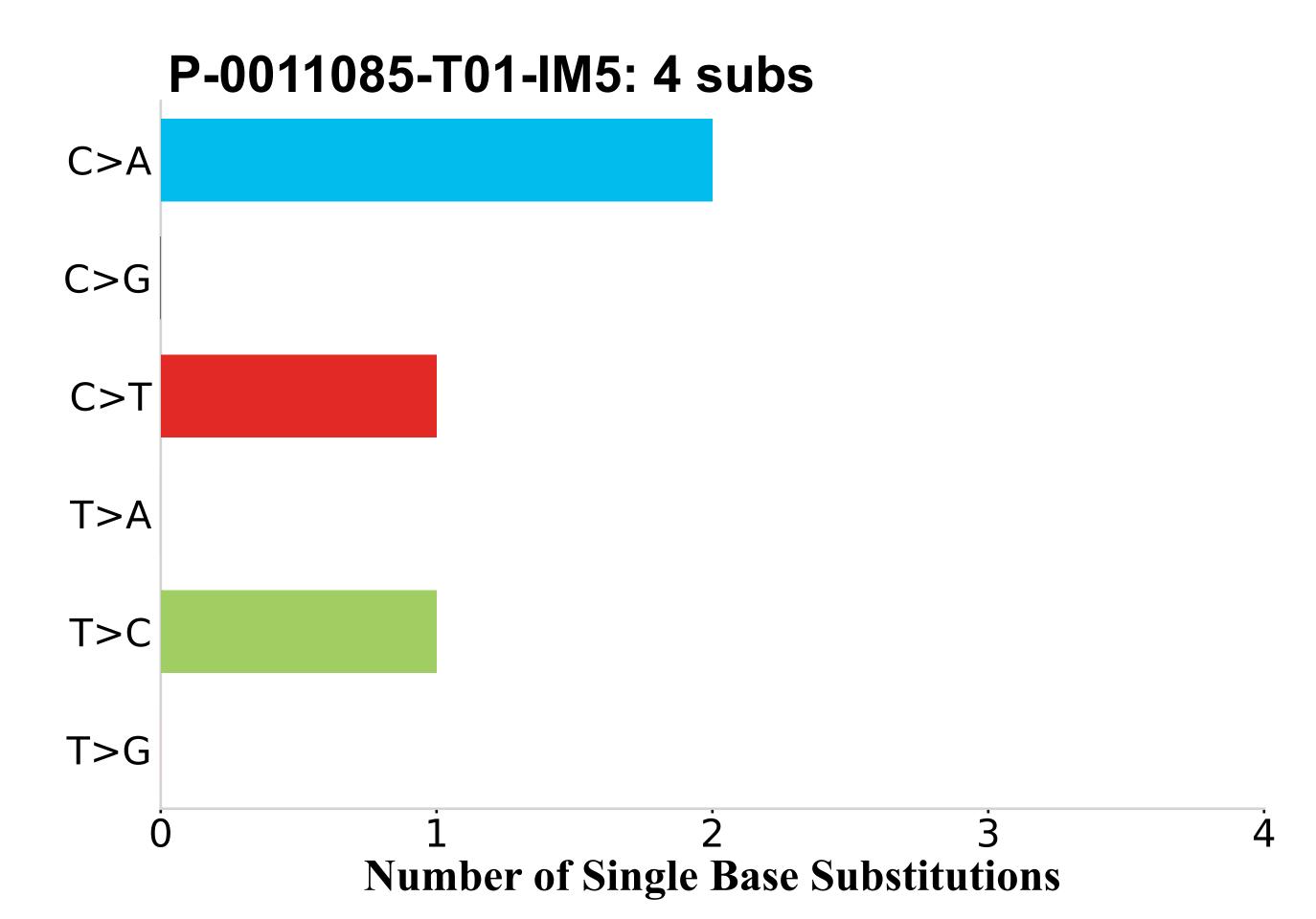


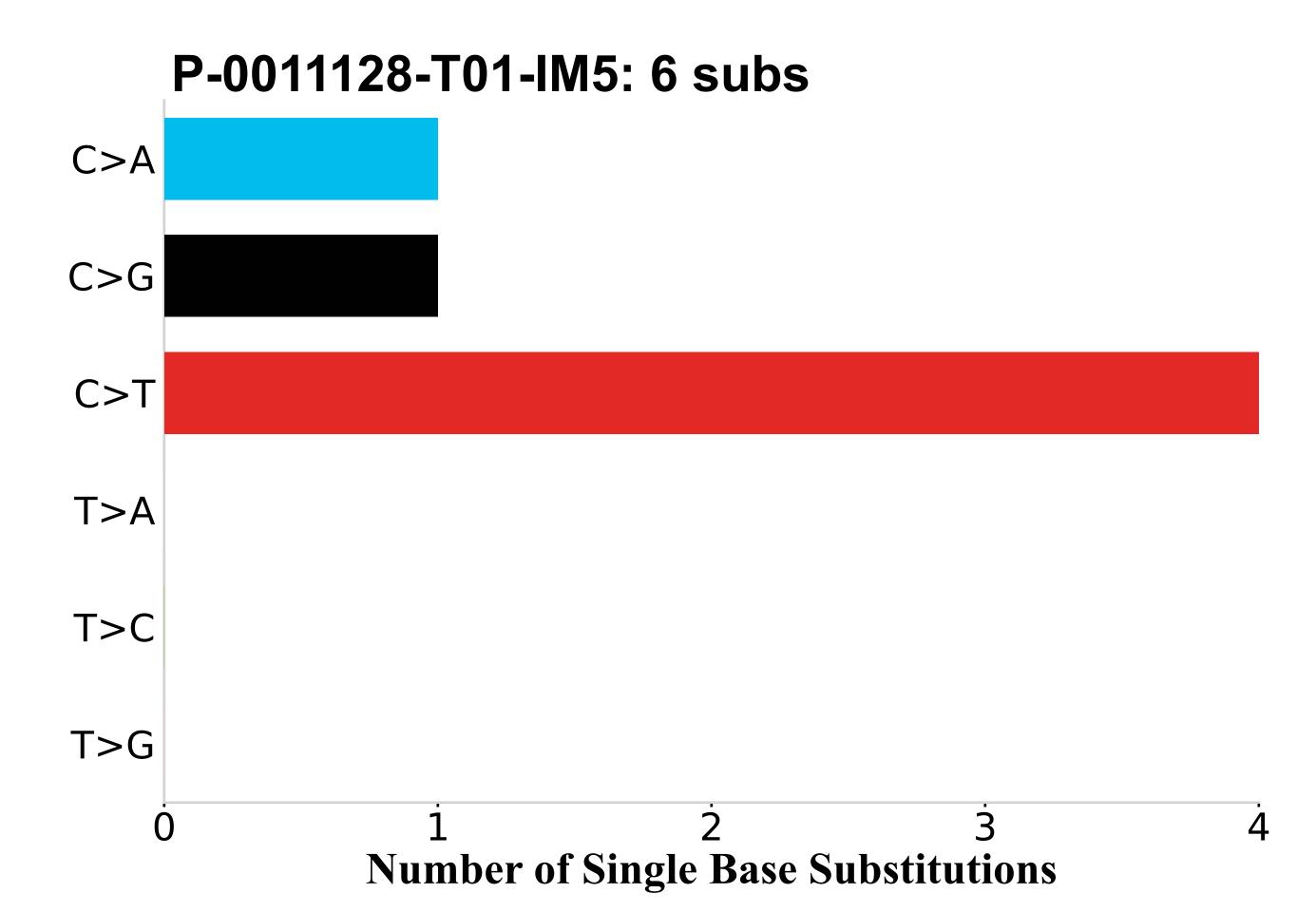


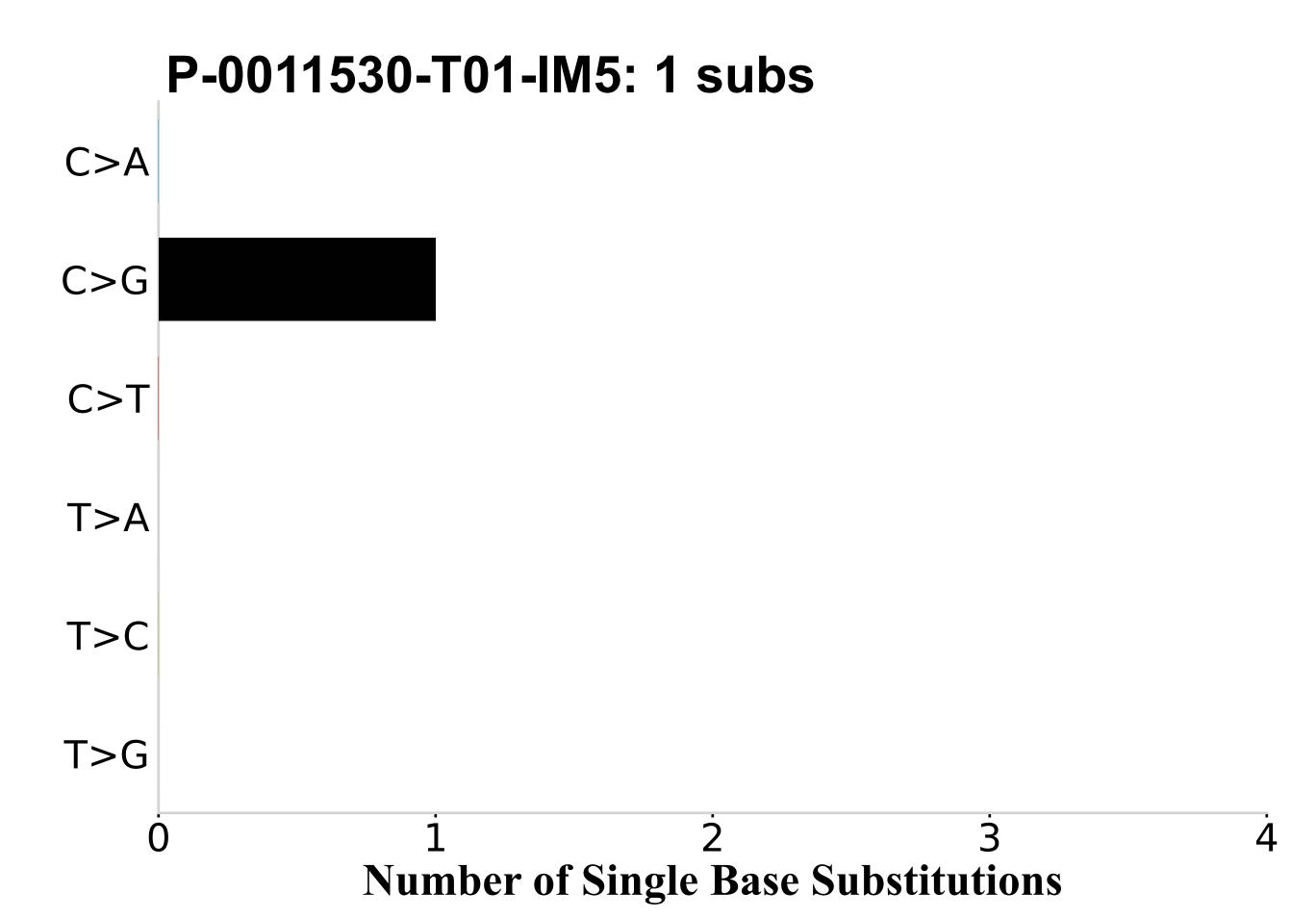






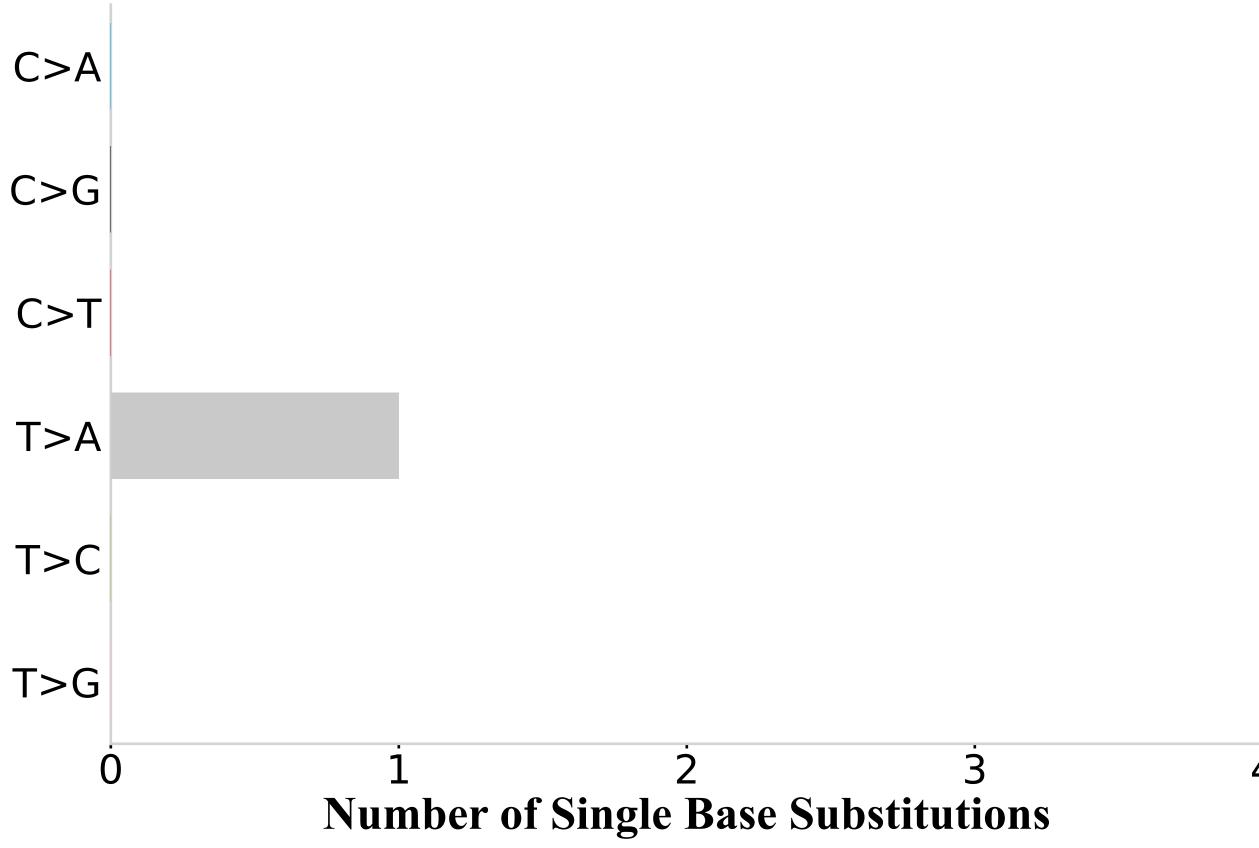


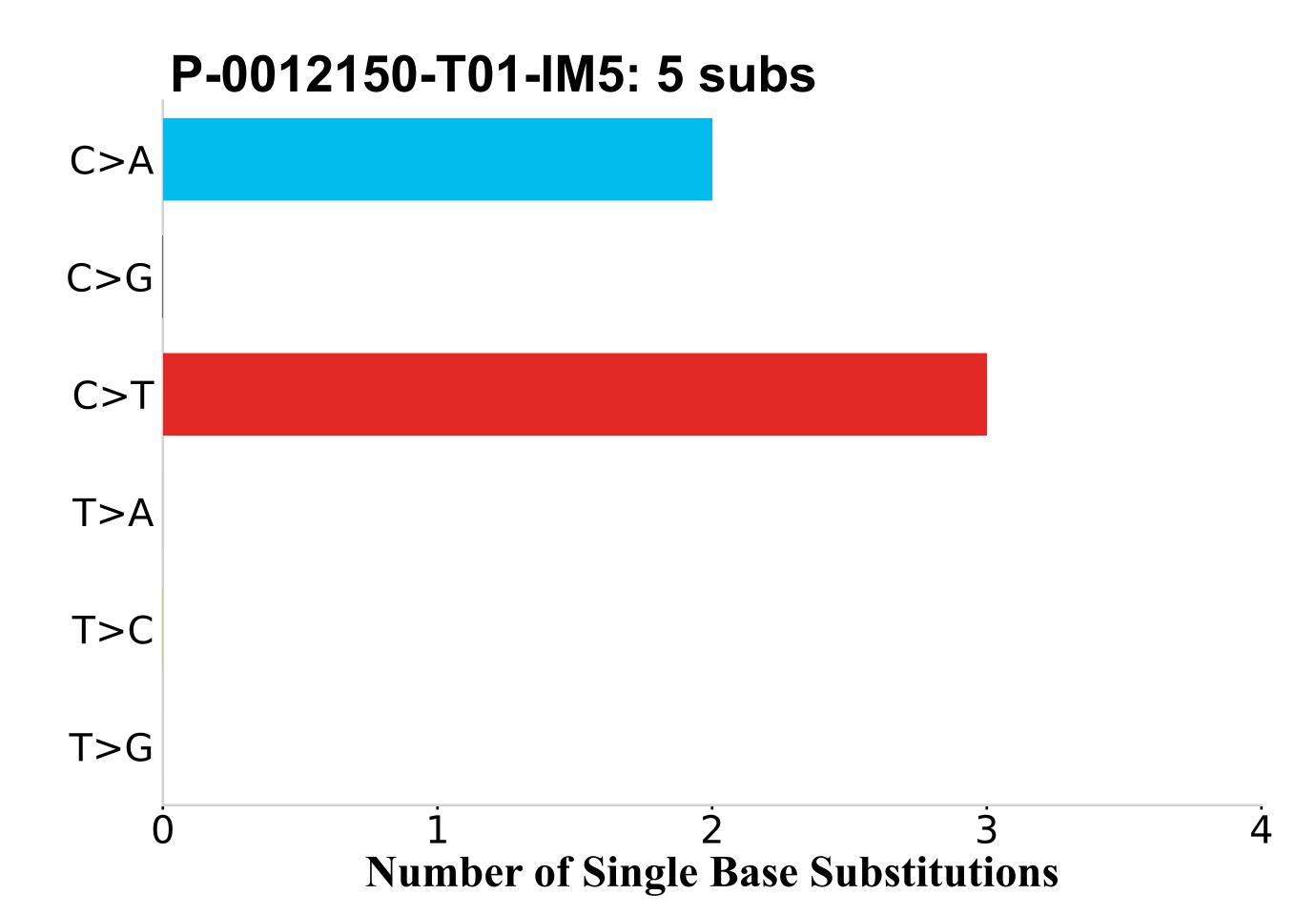




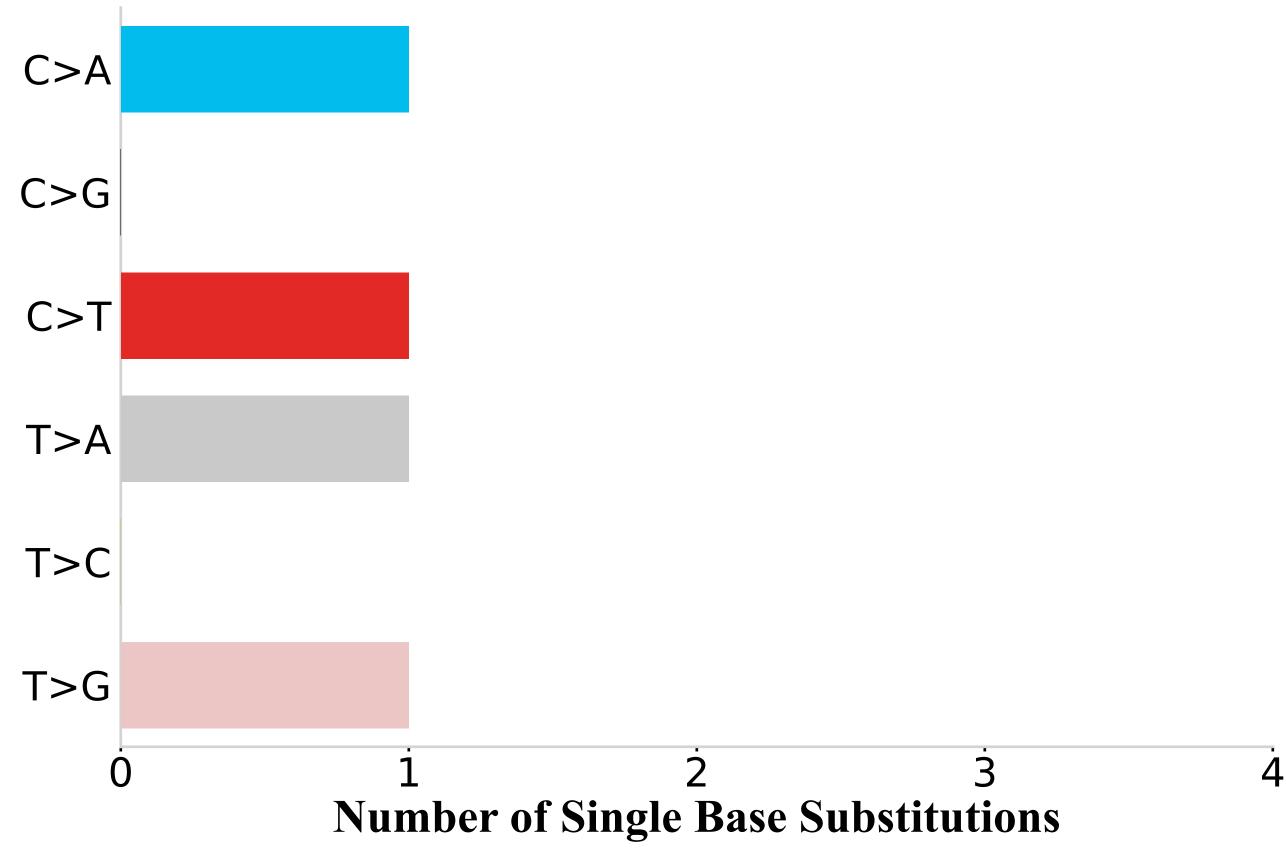
P-0012068-T01-IM5: 5 subs C>A C>G C>T T>A T>C T>G **Number of Single Base Substitutions**

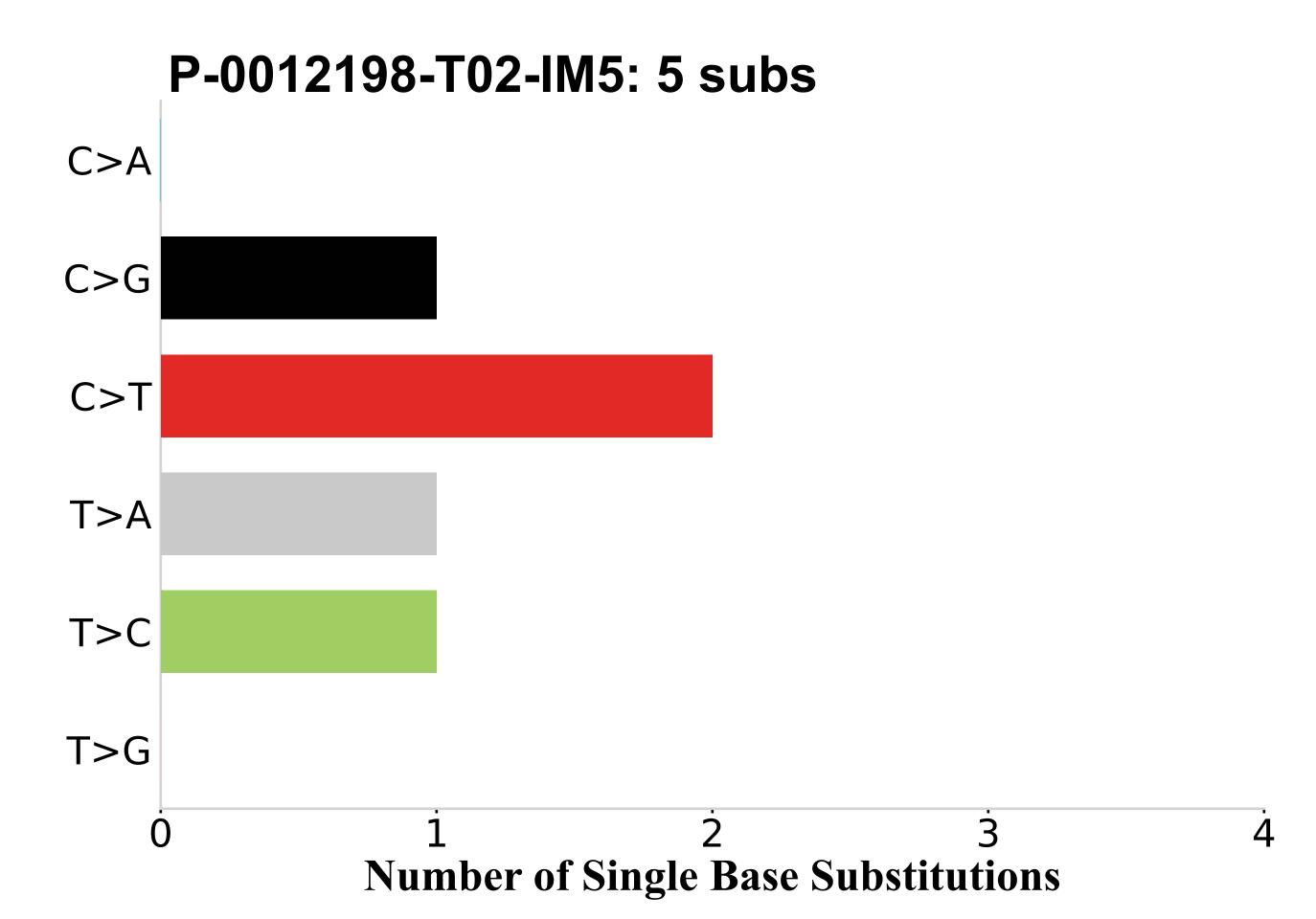




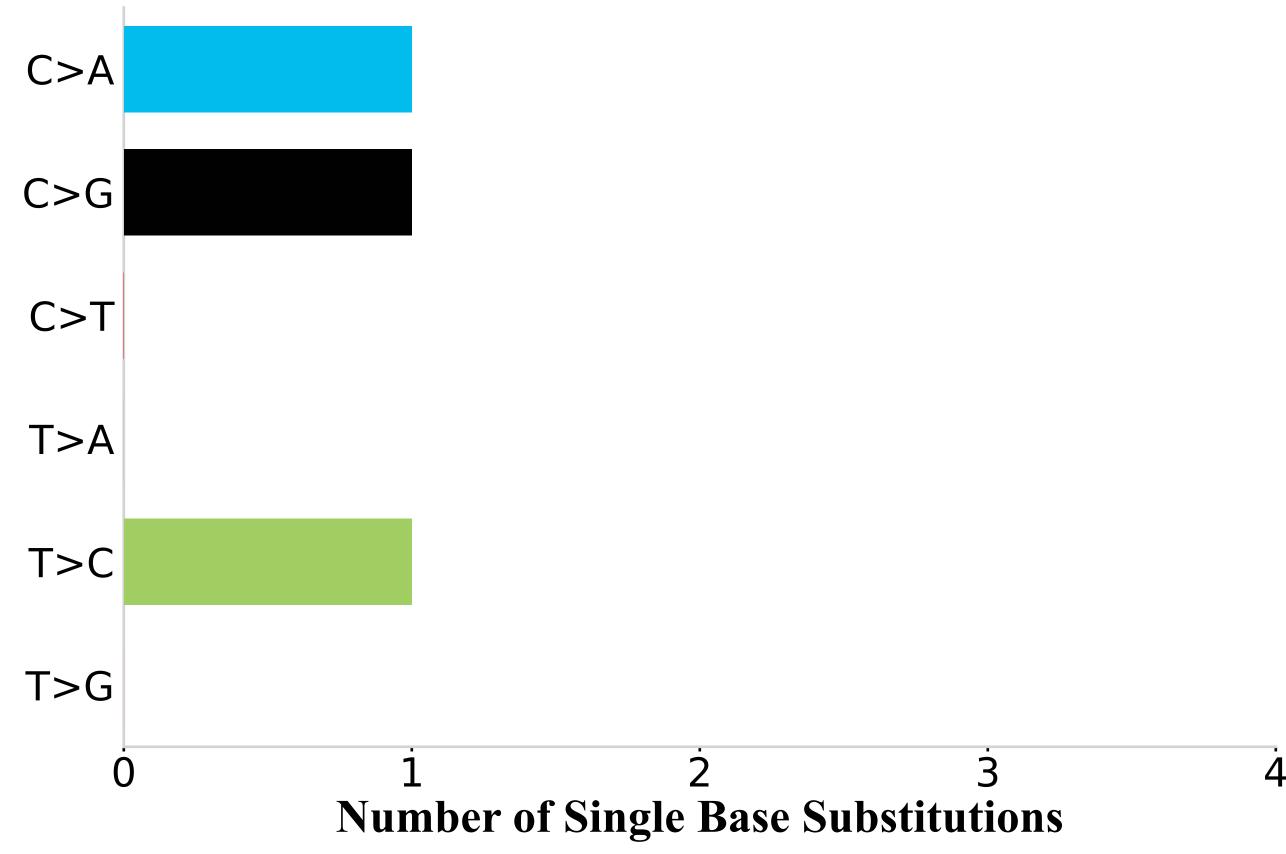


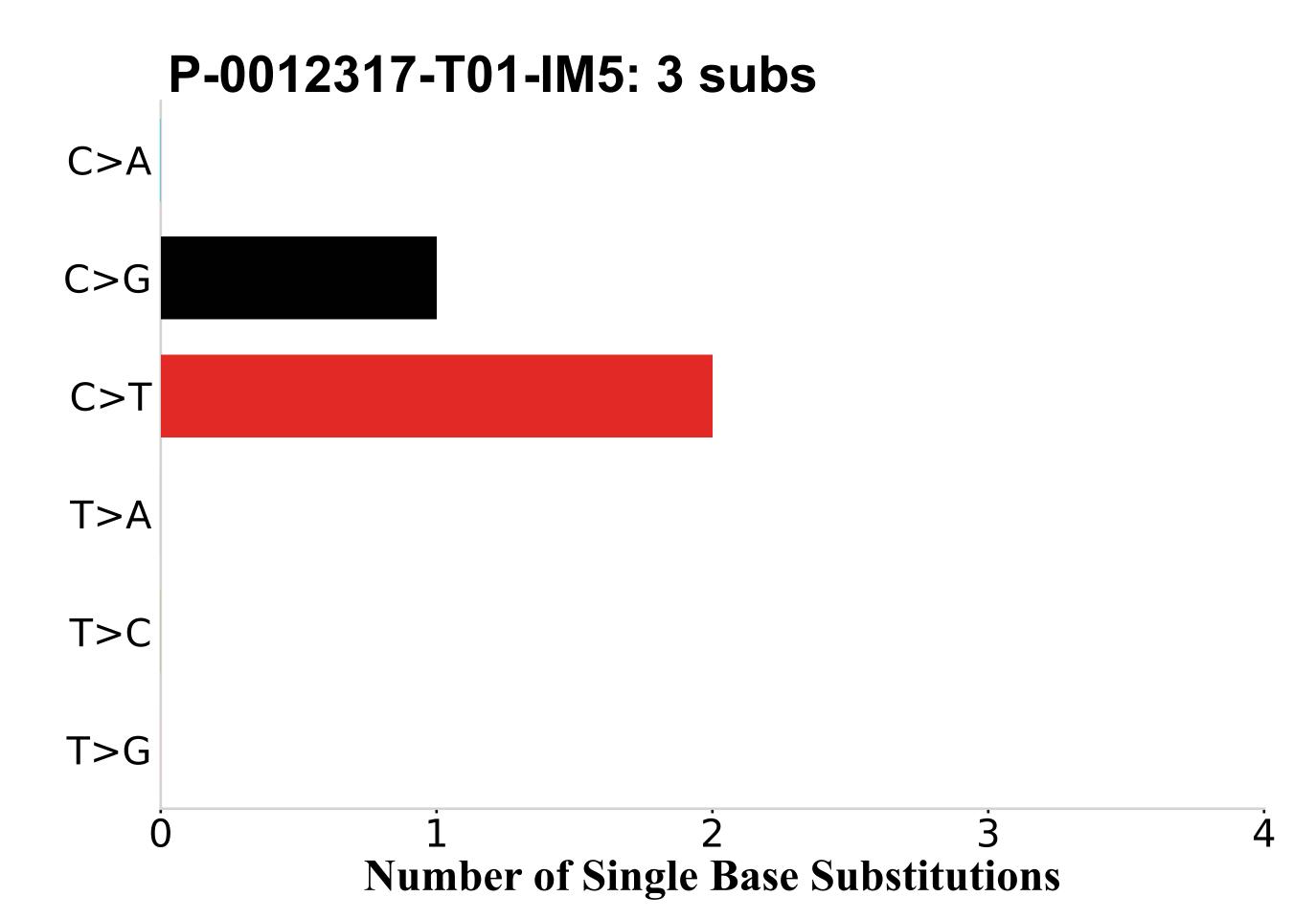
P-0012182-T01-IM5: 4 subs

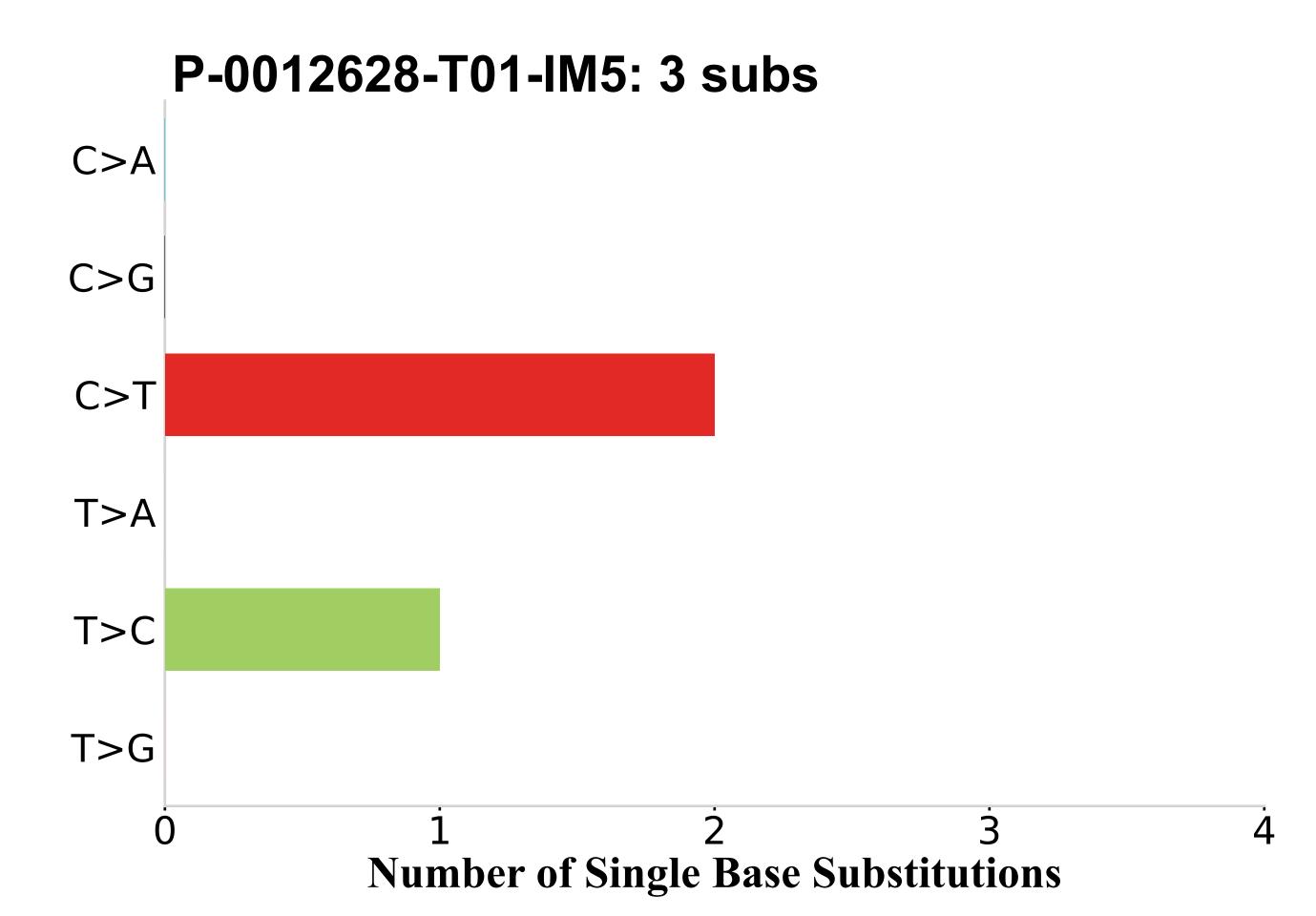


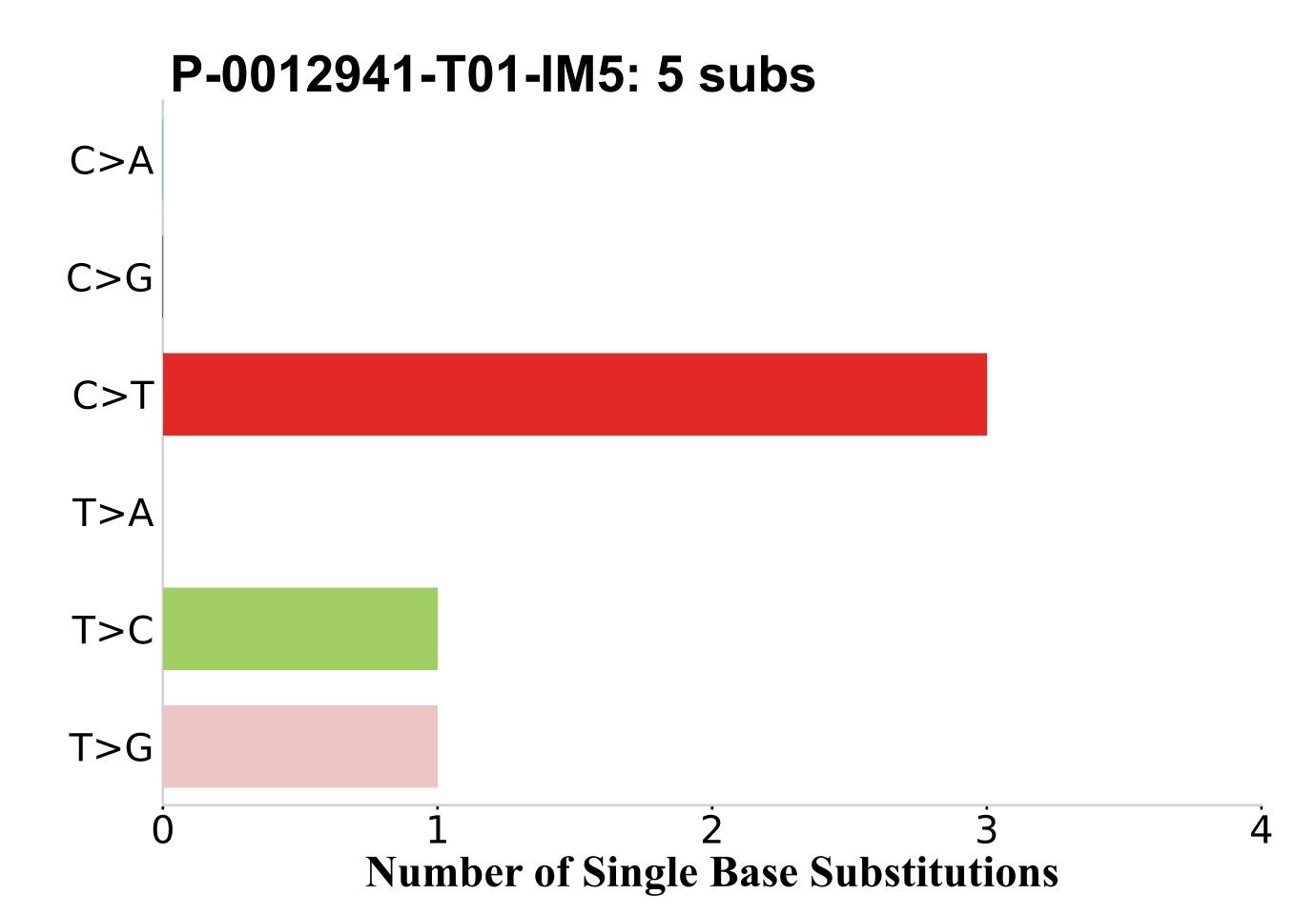


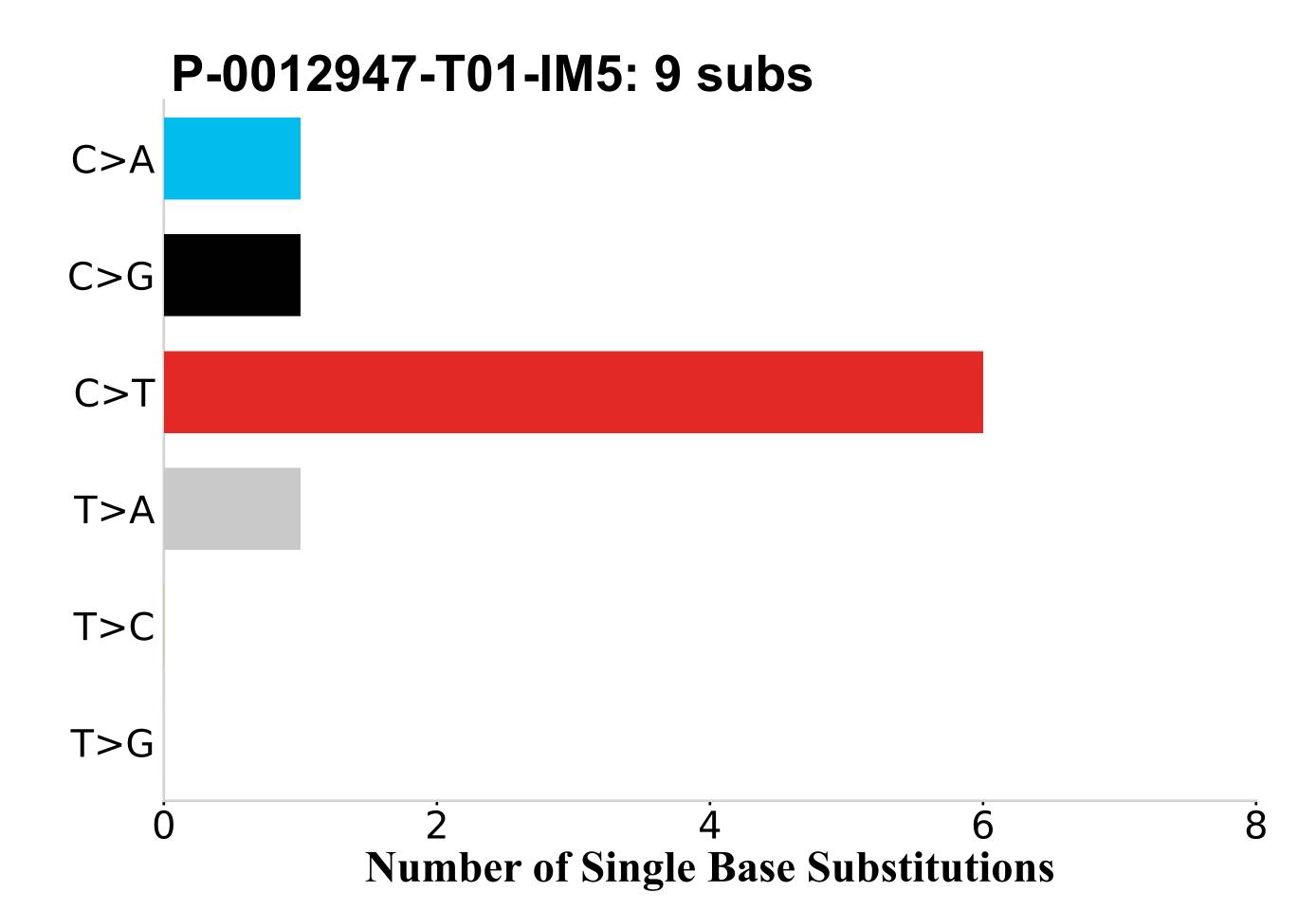
P-0012276-T01-IM5: 3 subs

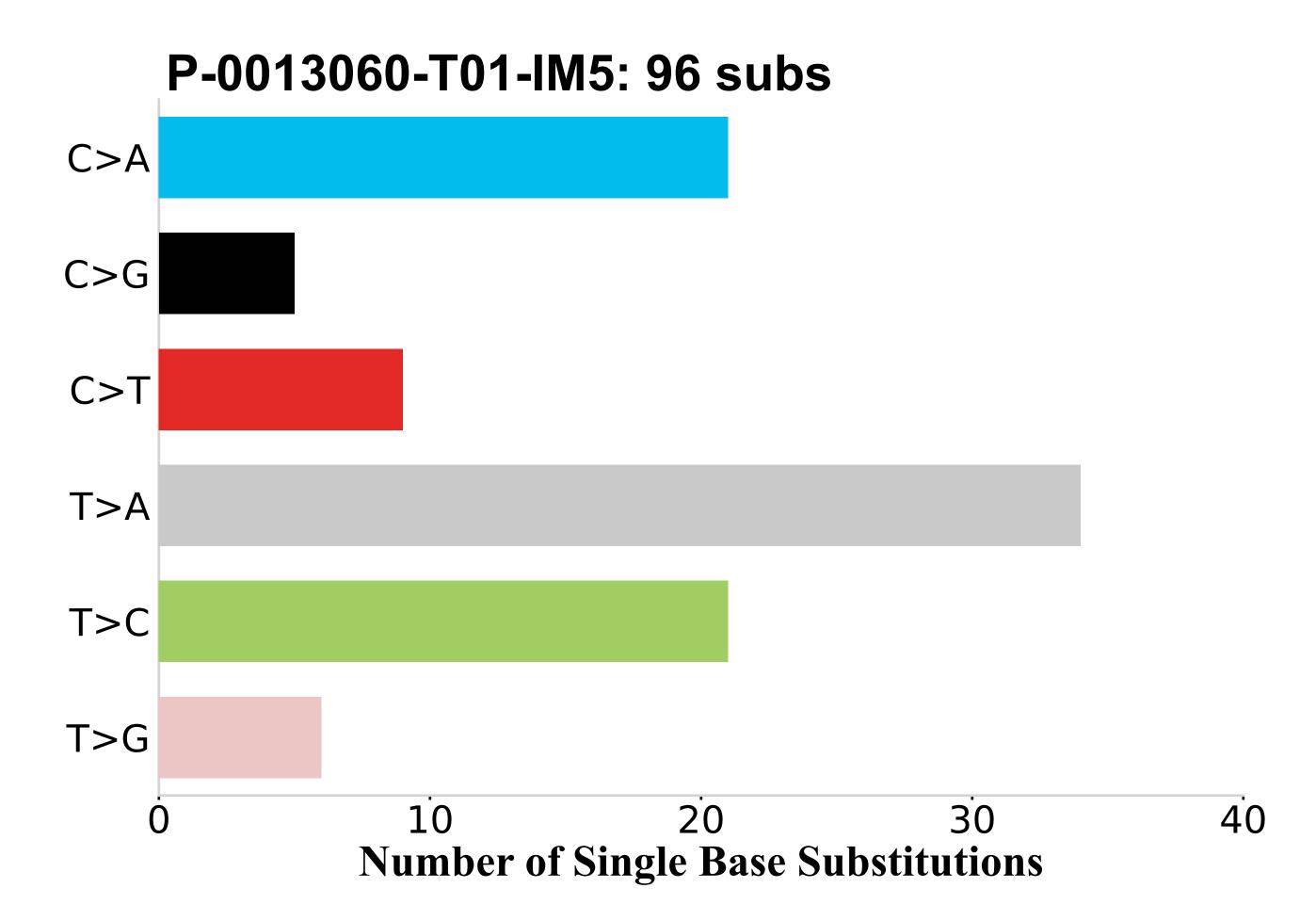


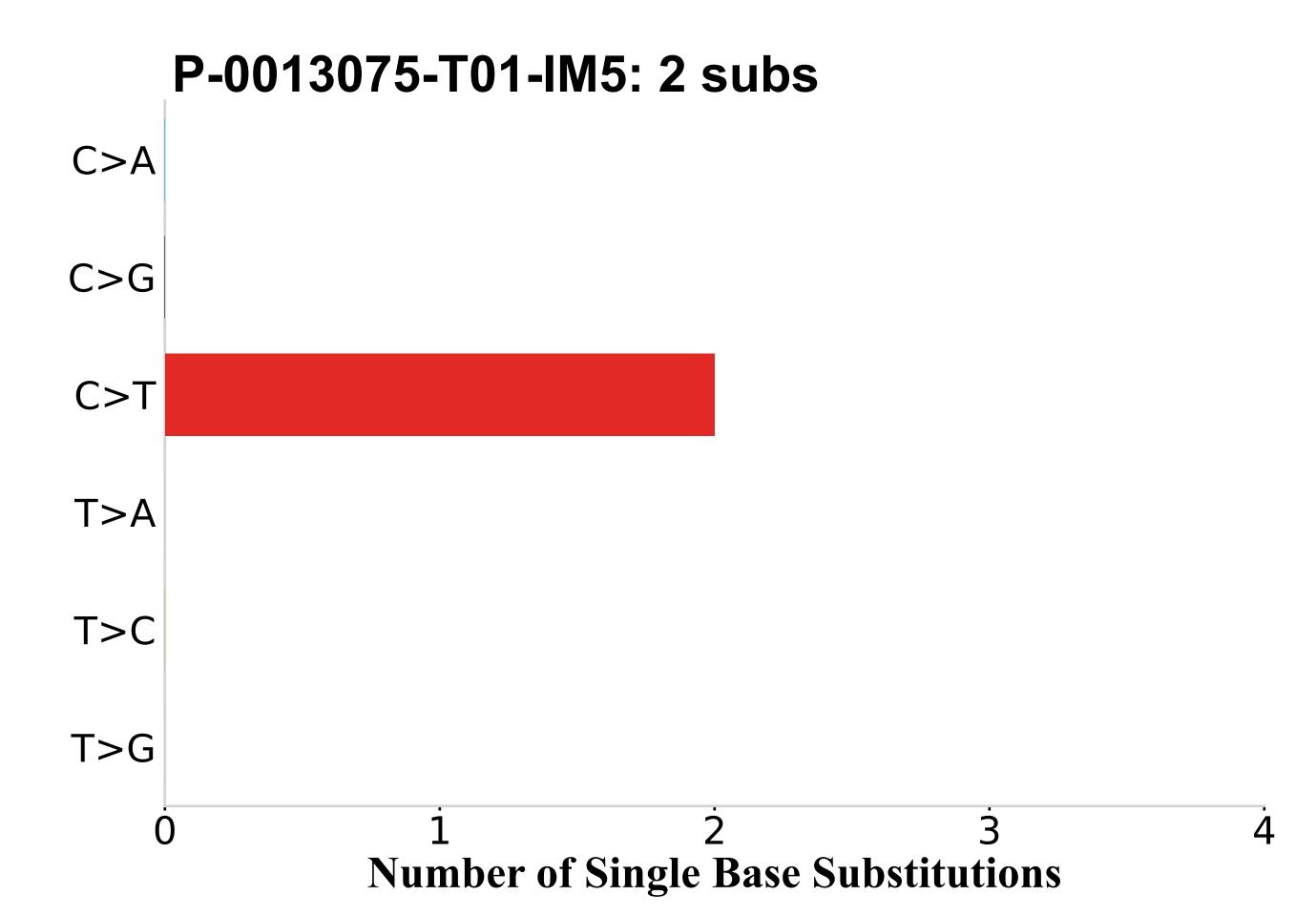


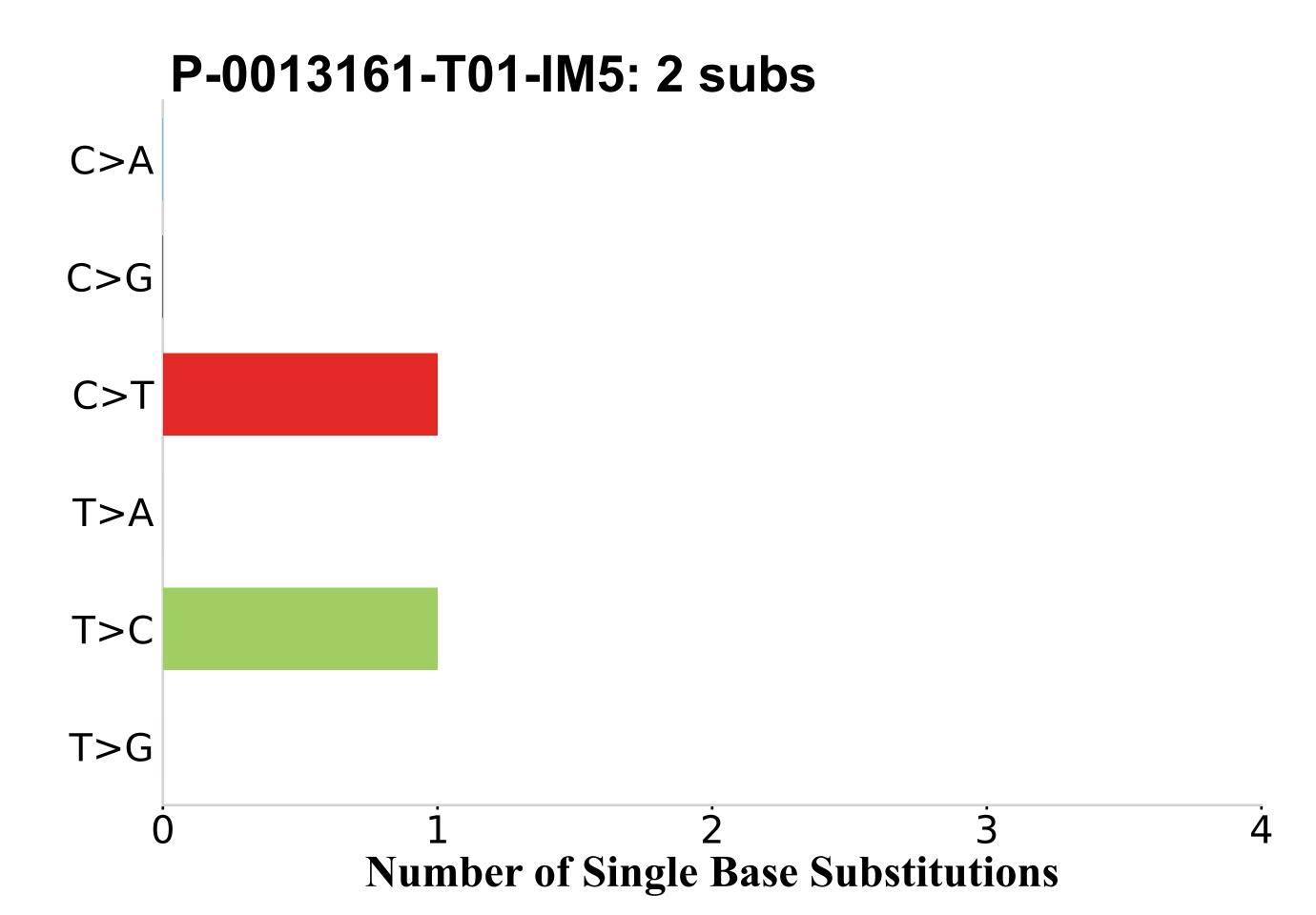


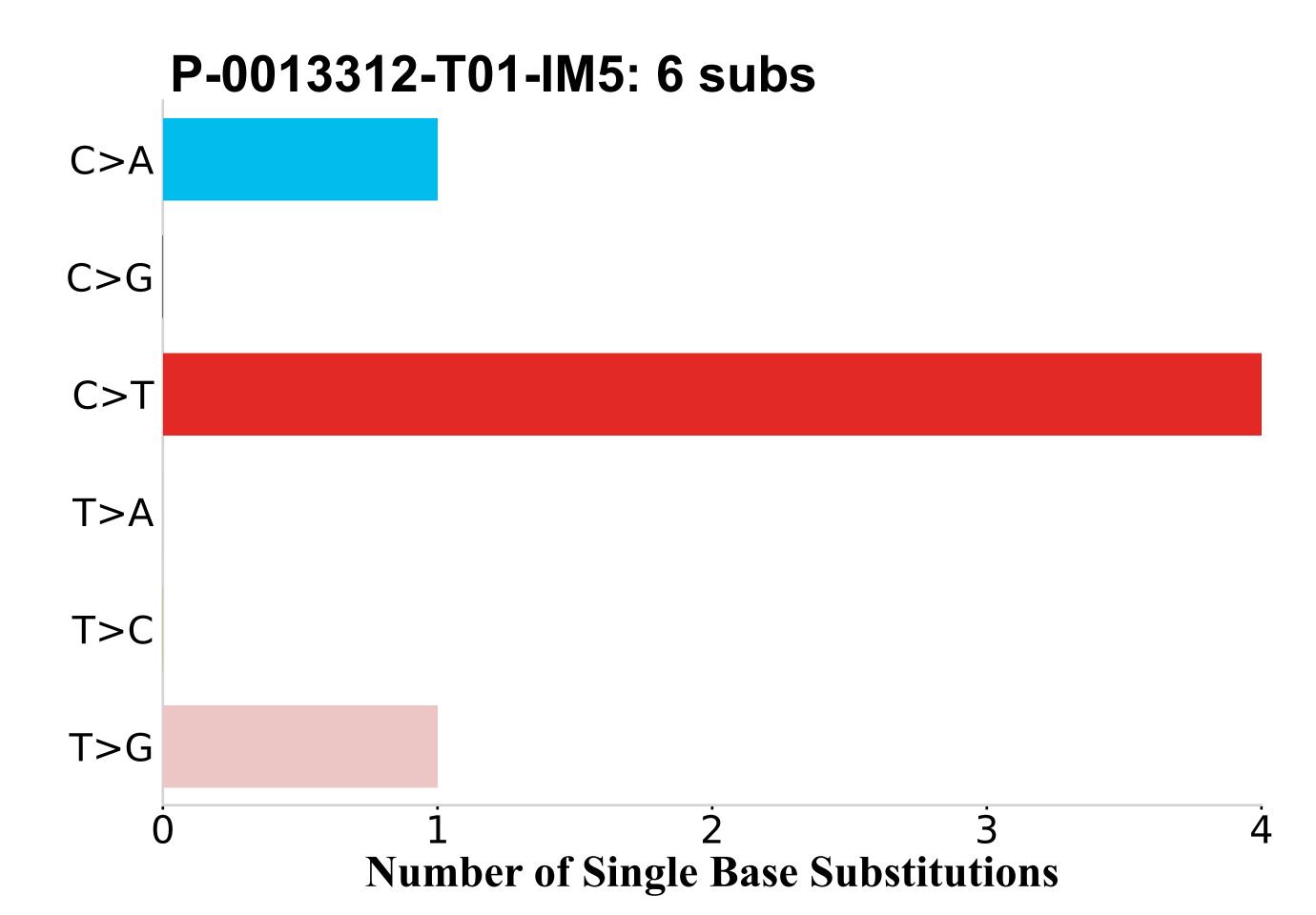


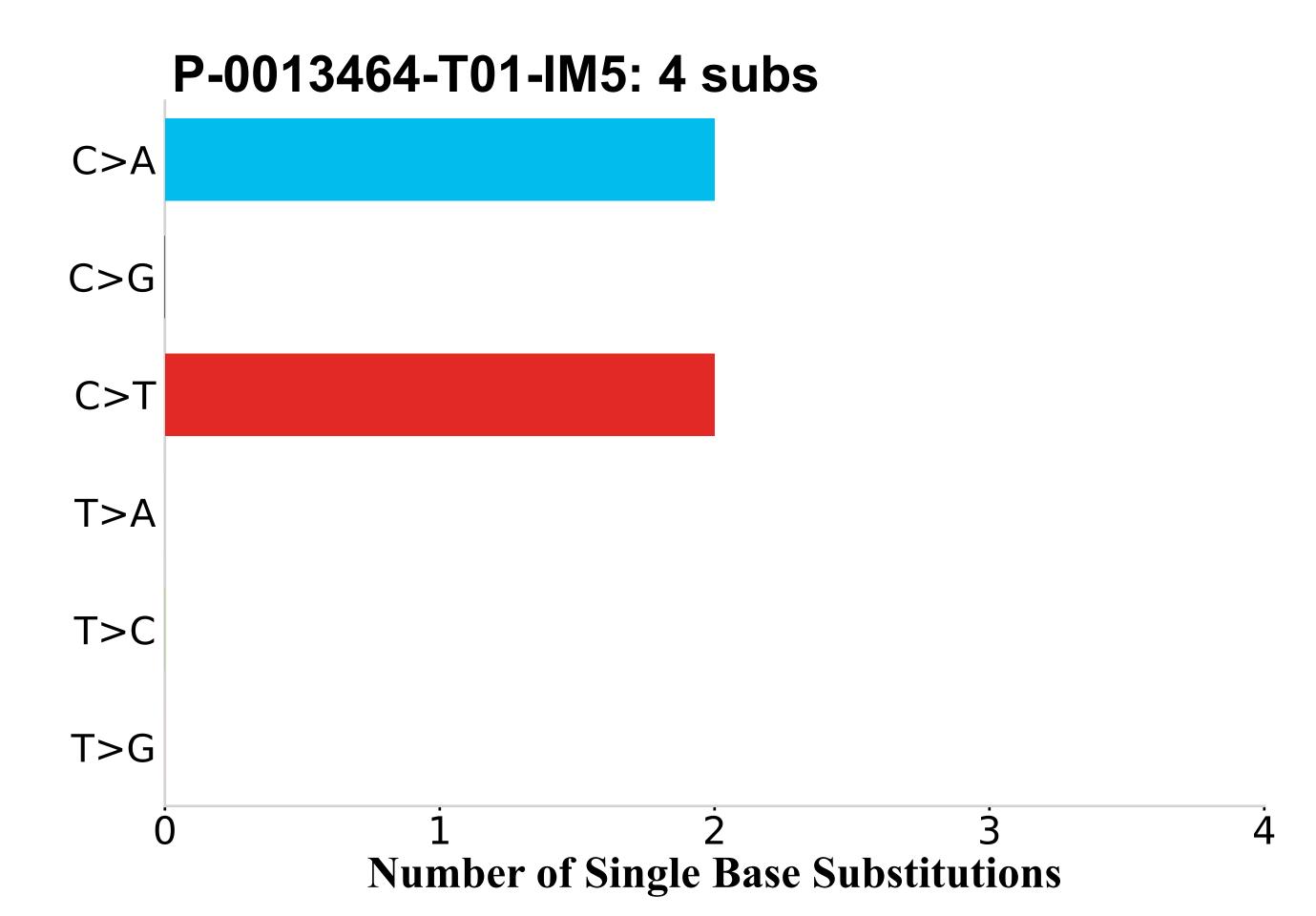


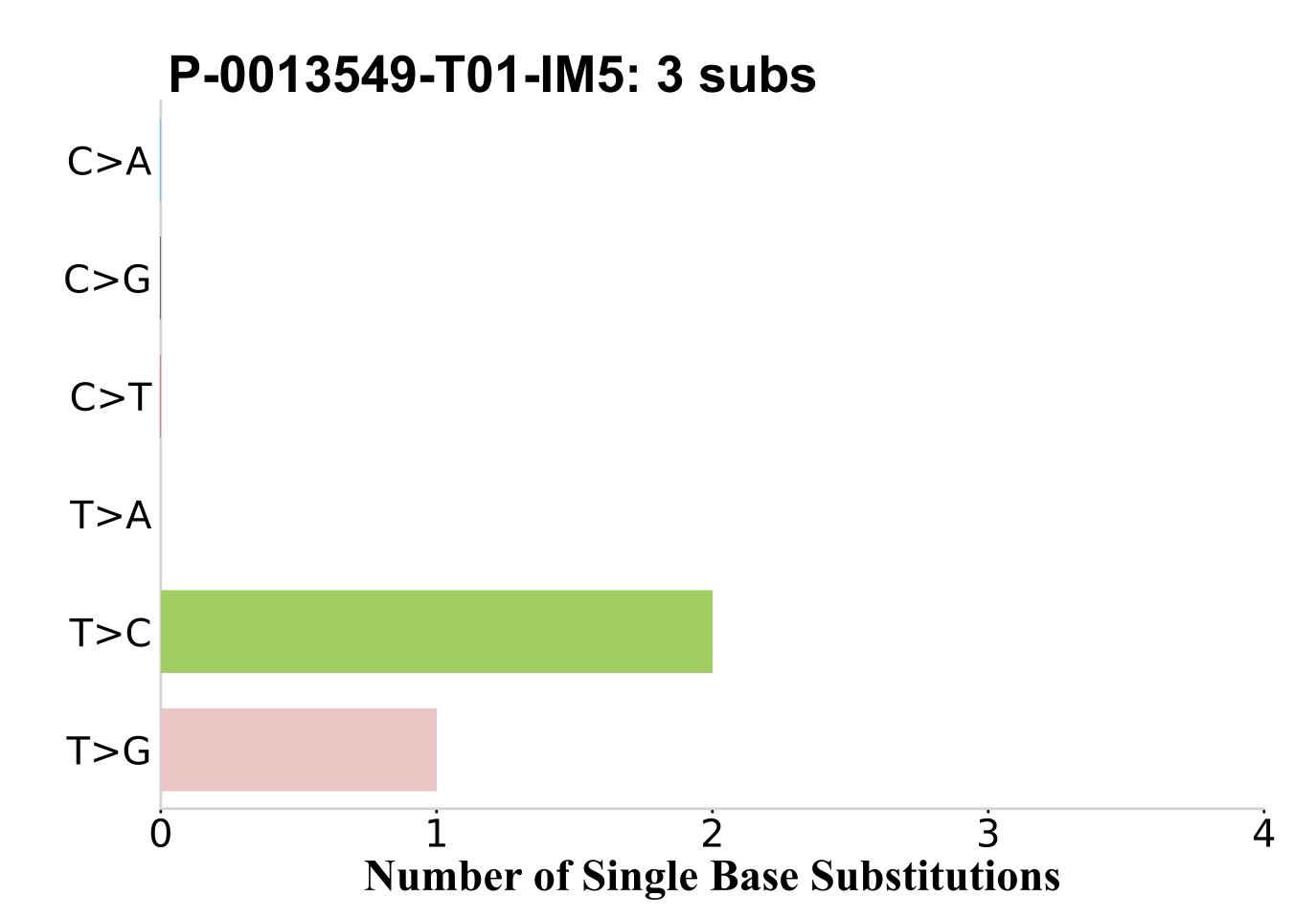




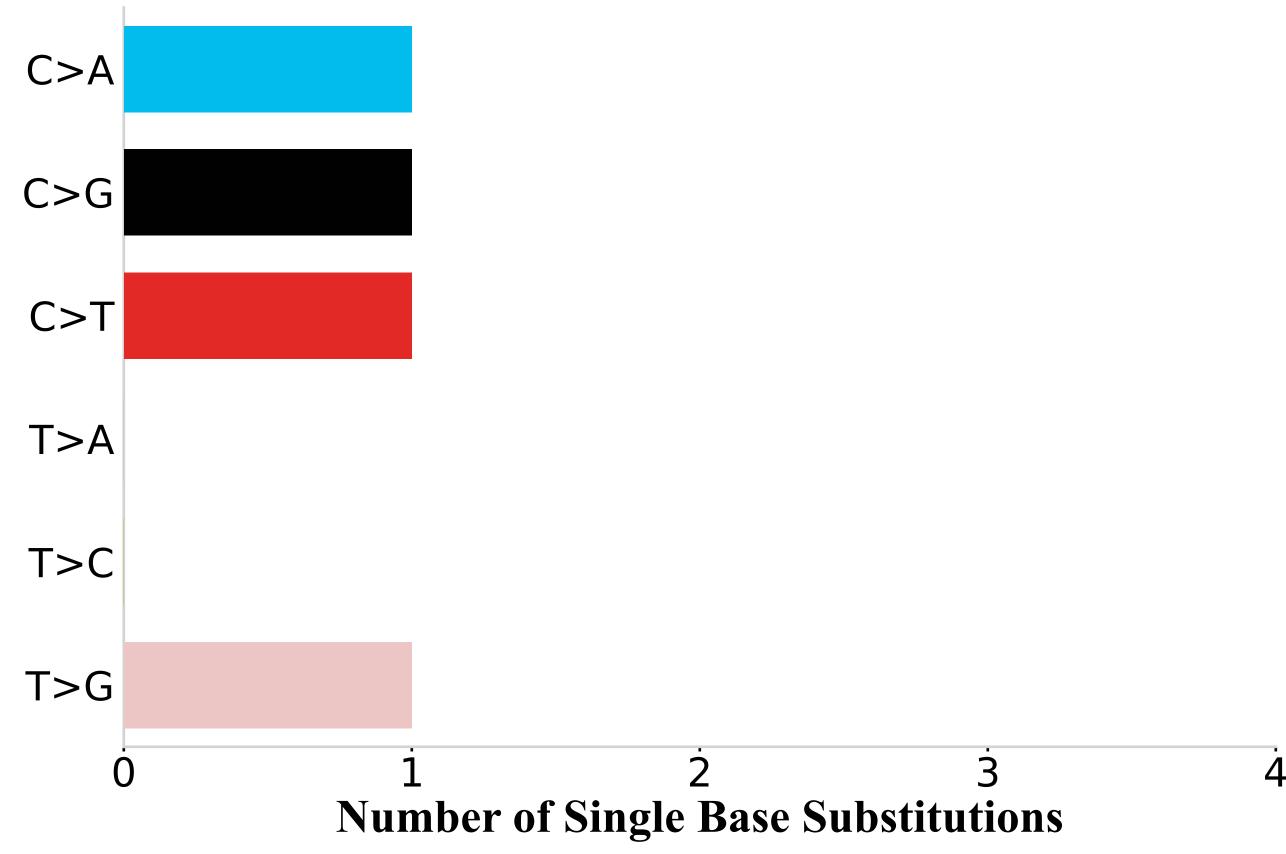




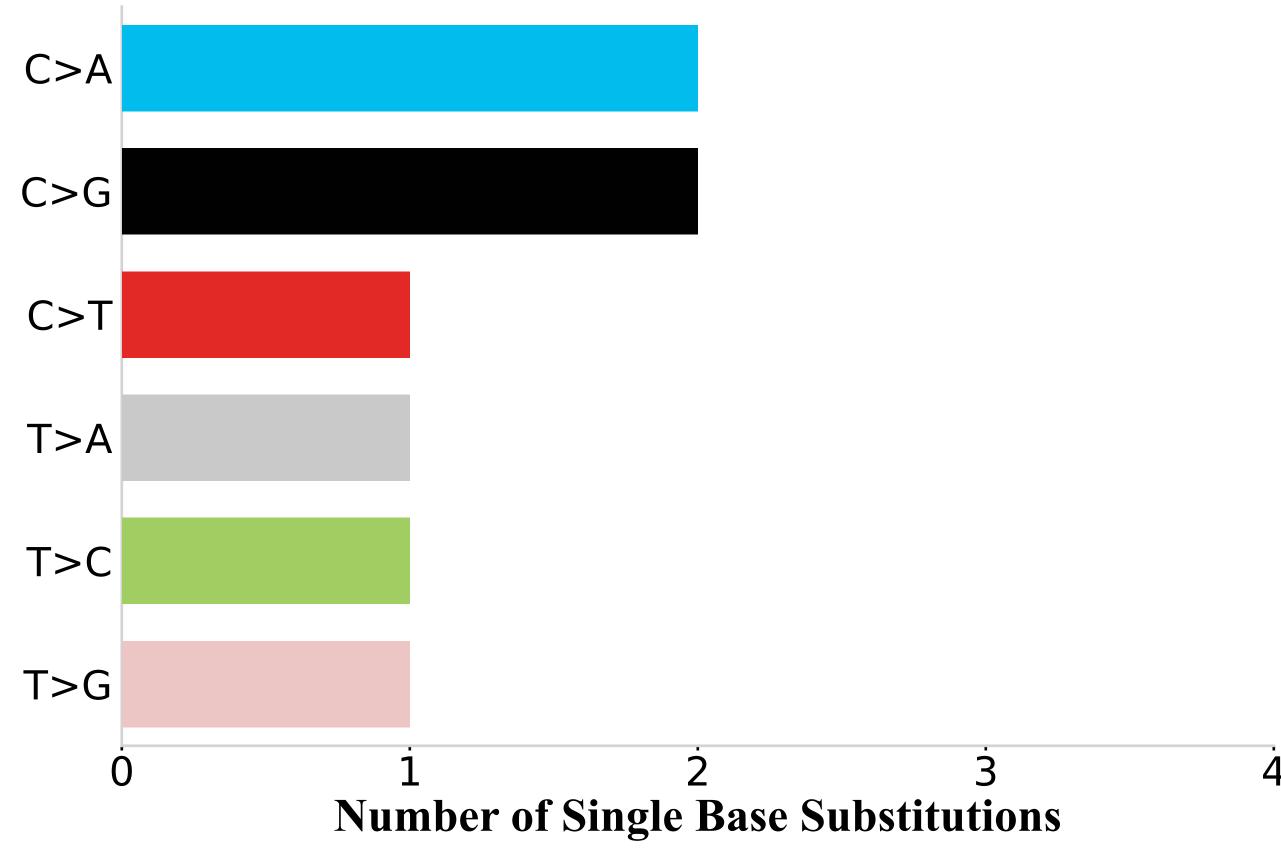


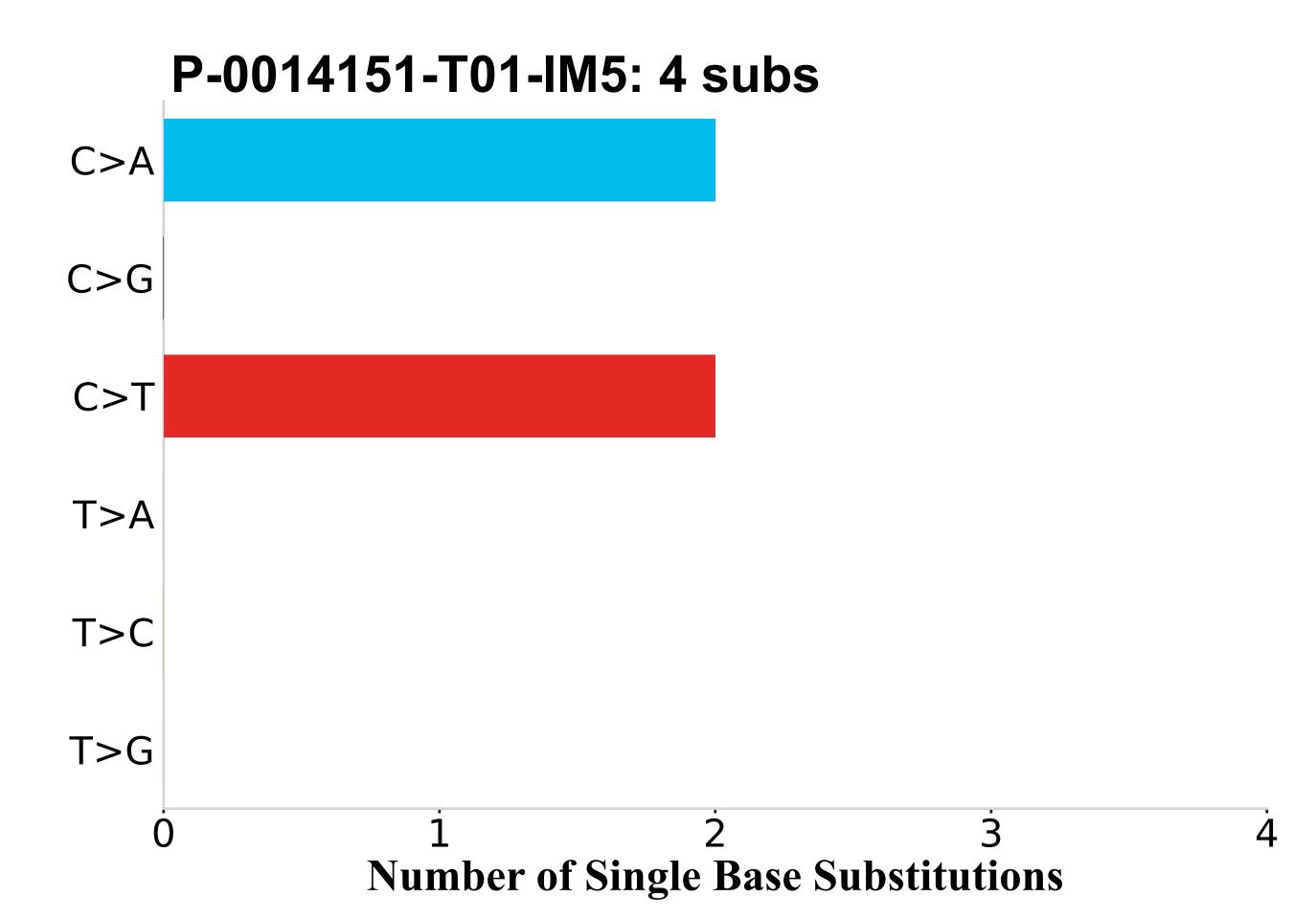


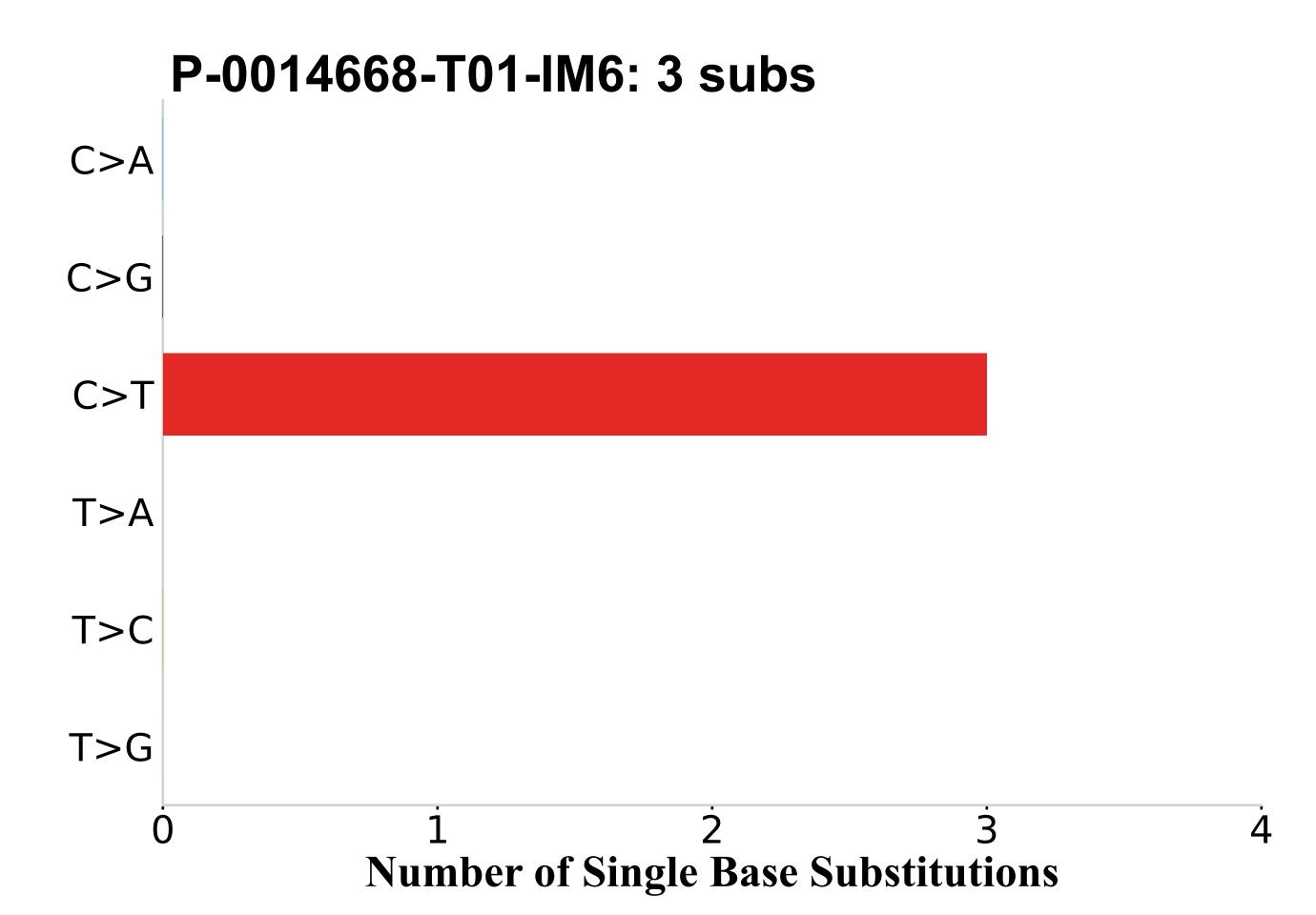
P-0013553-T01-IM5: 4 subs



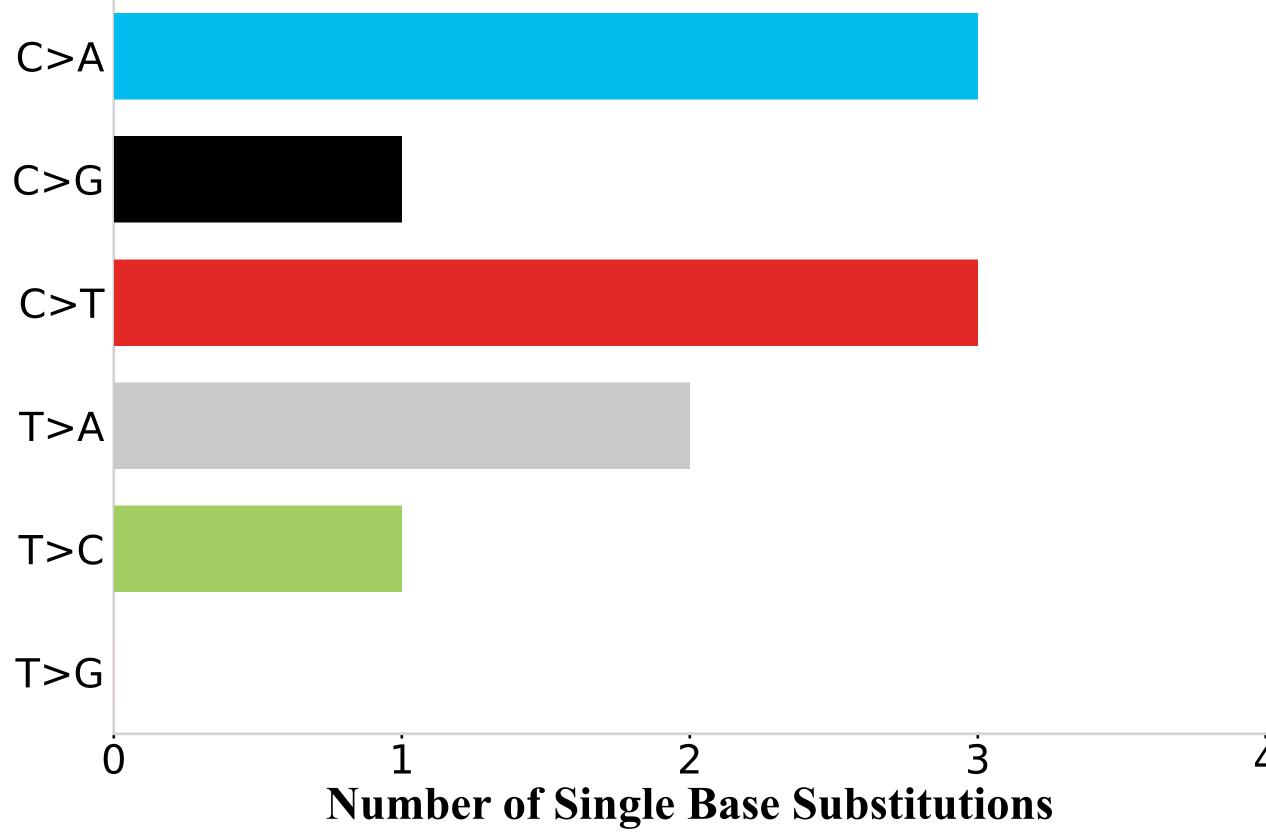




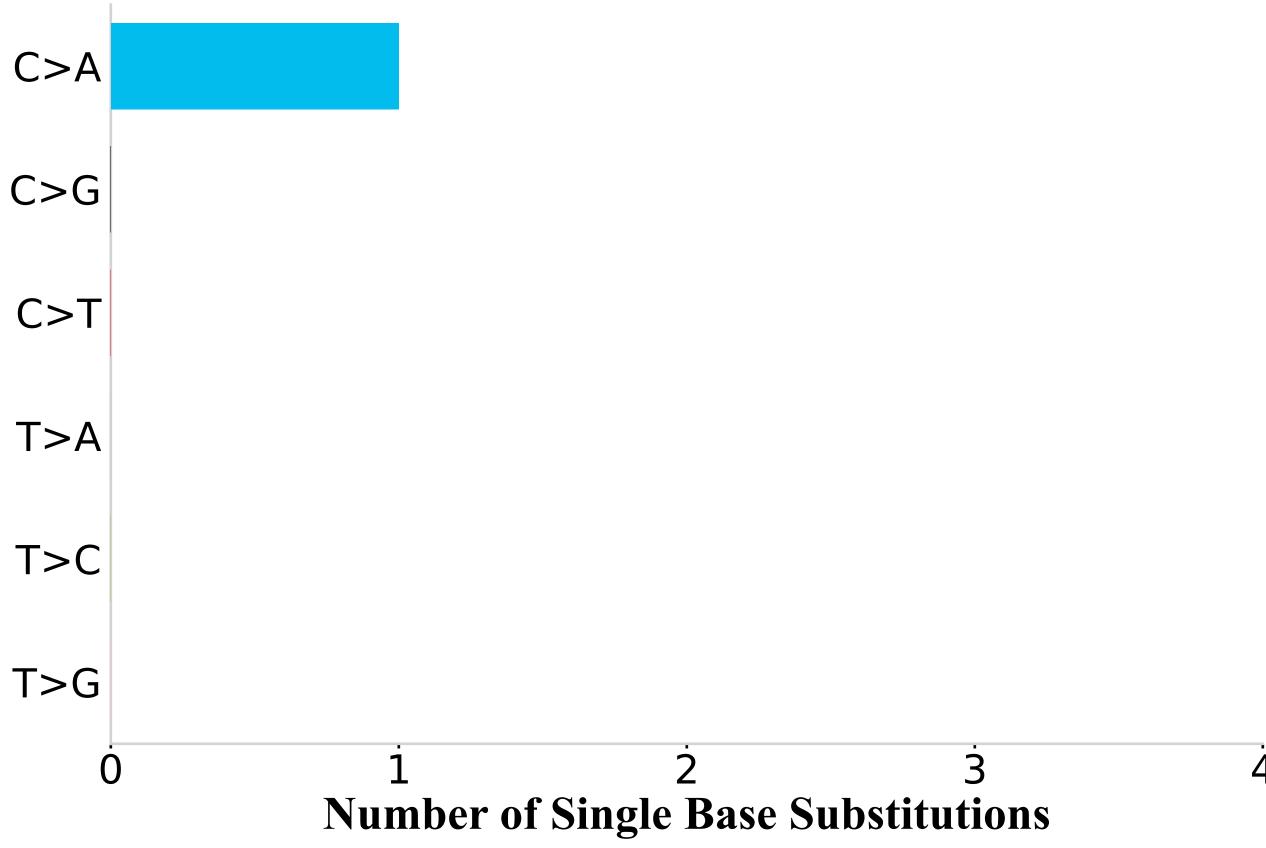


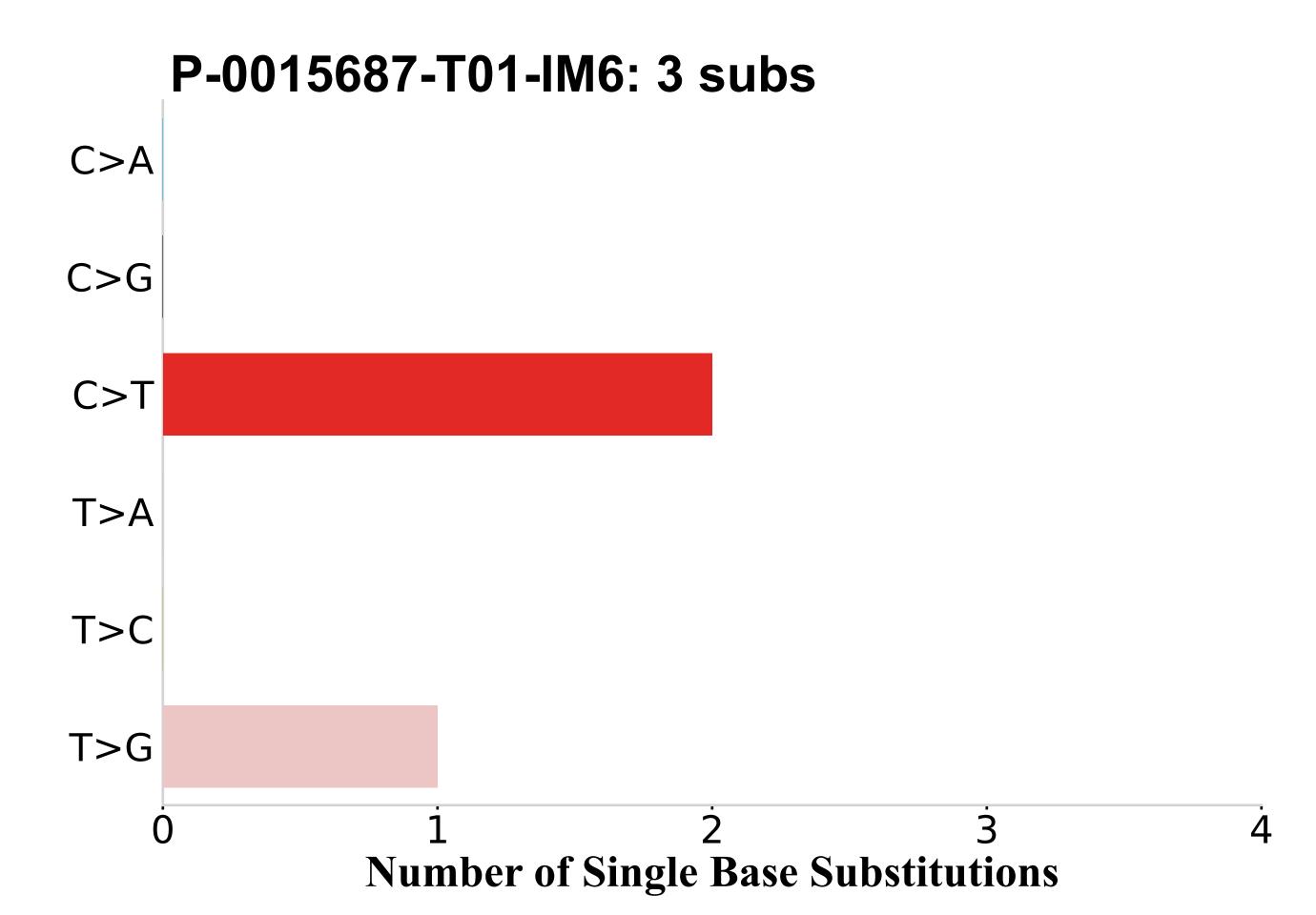


P-0015203-T01-IM6: 10 subs

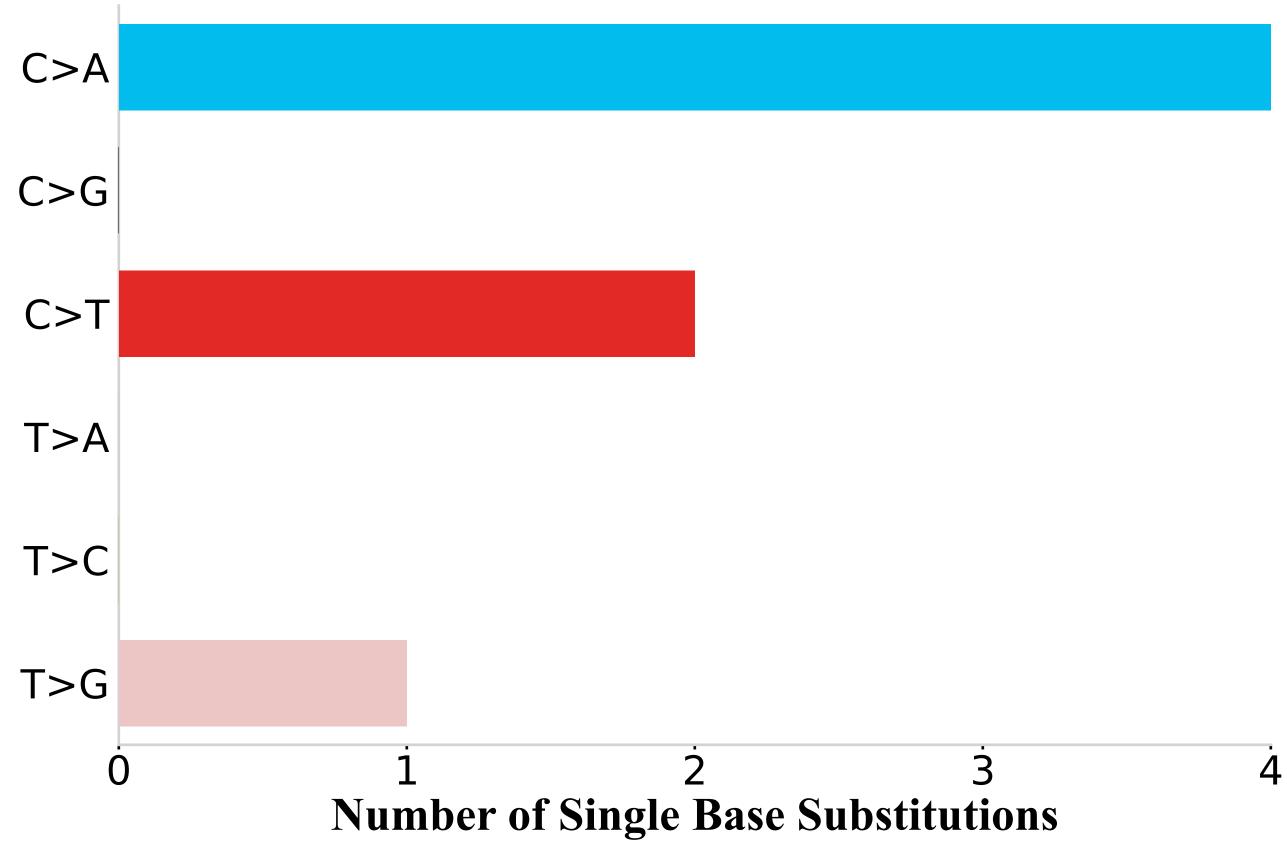


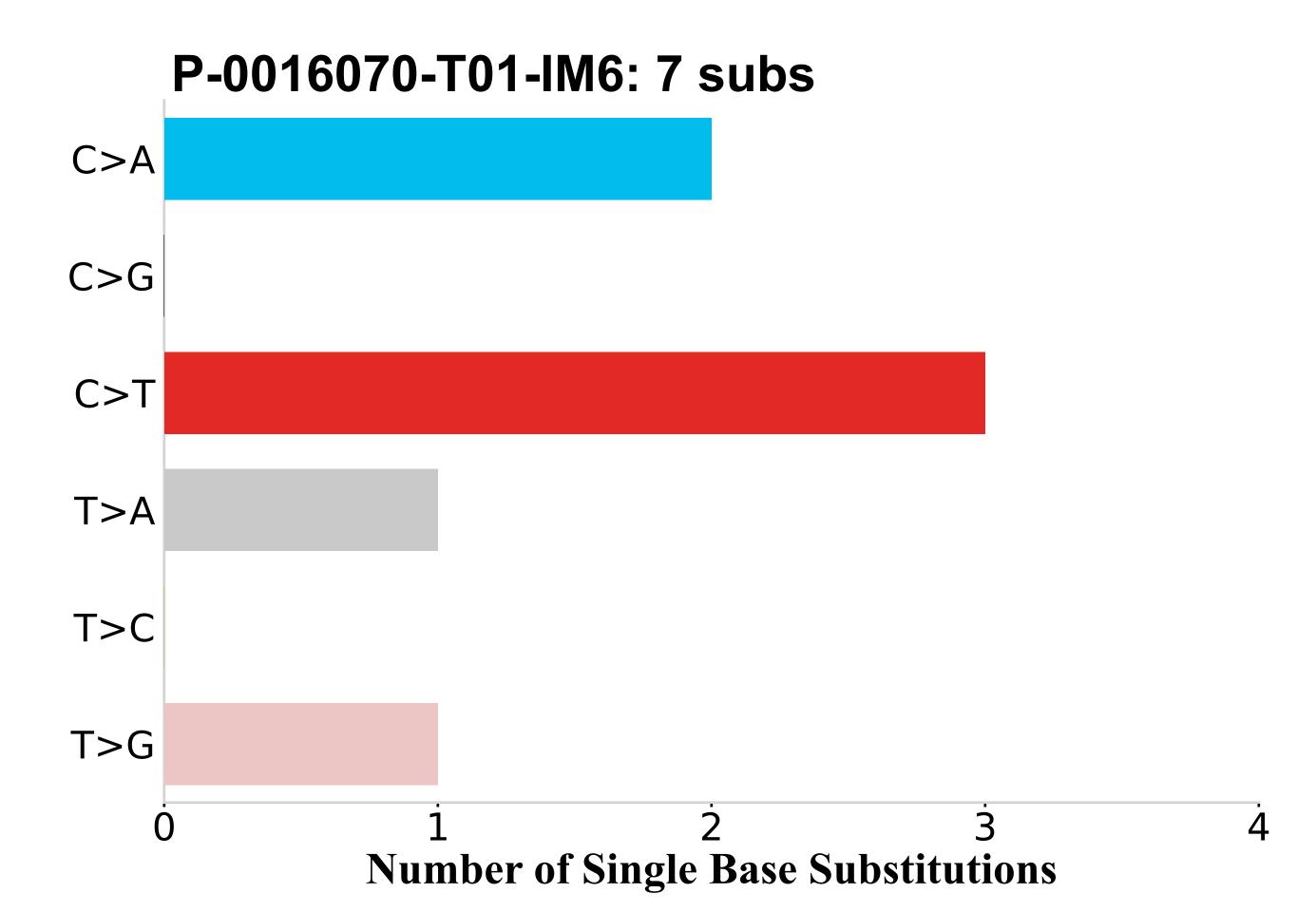


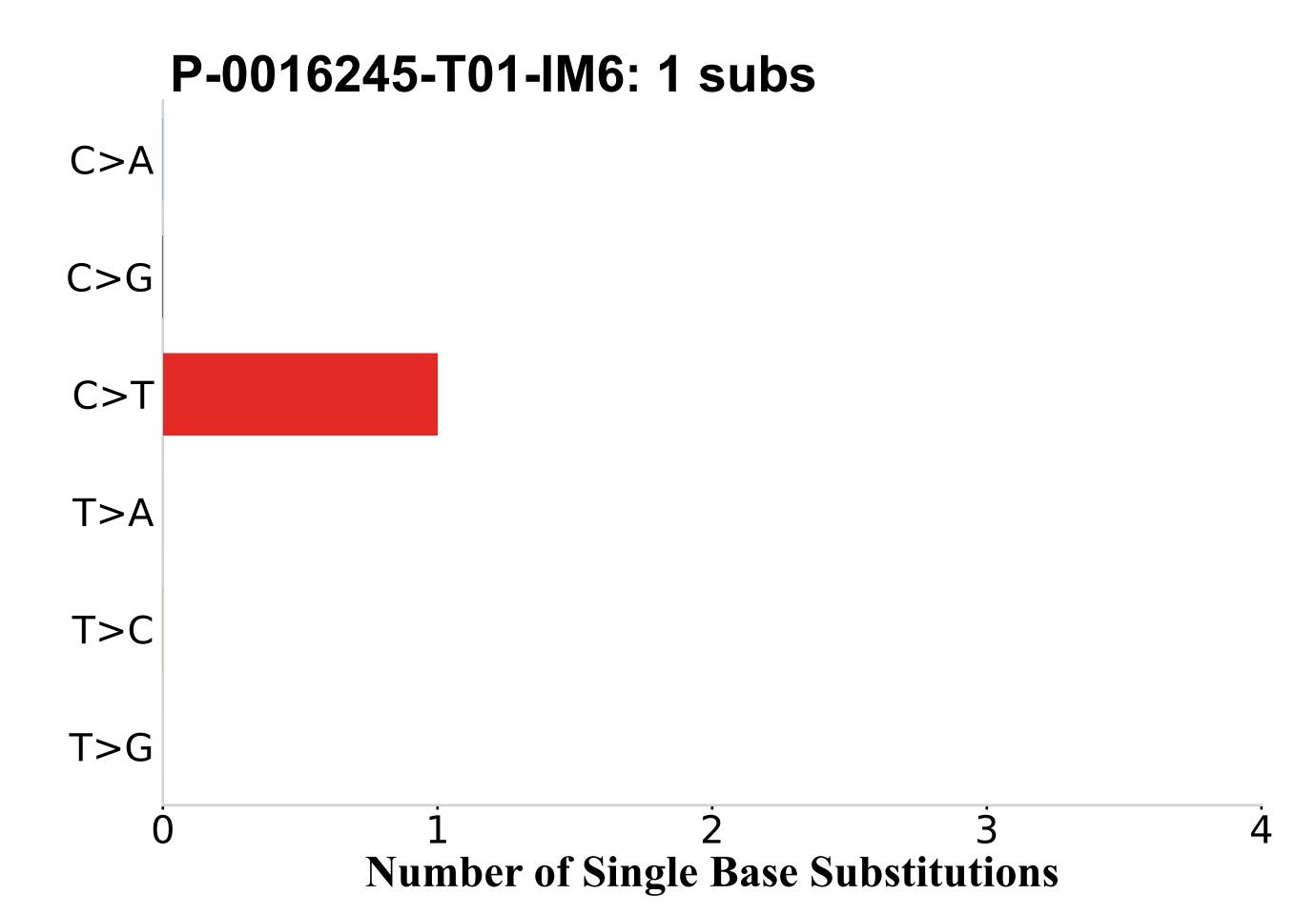


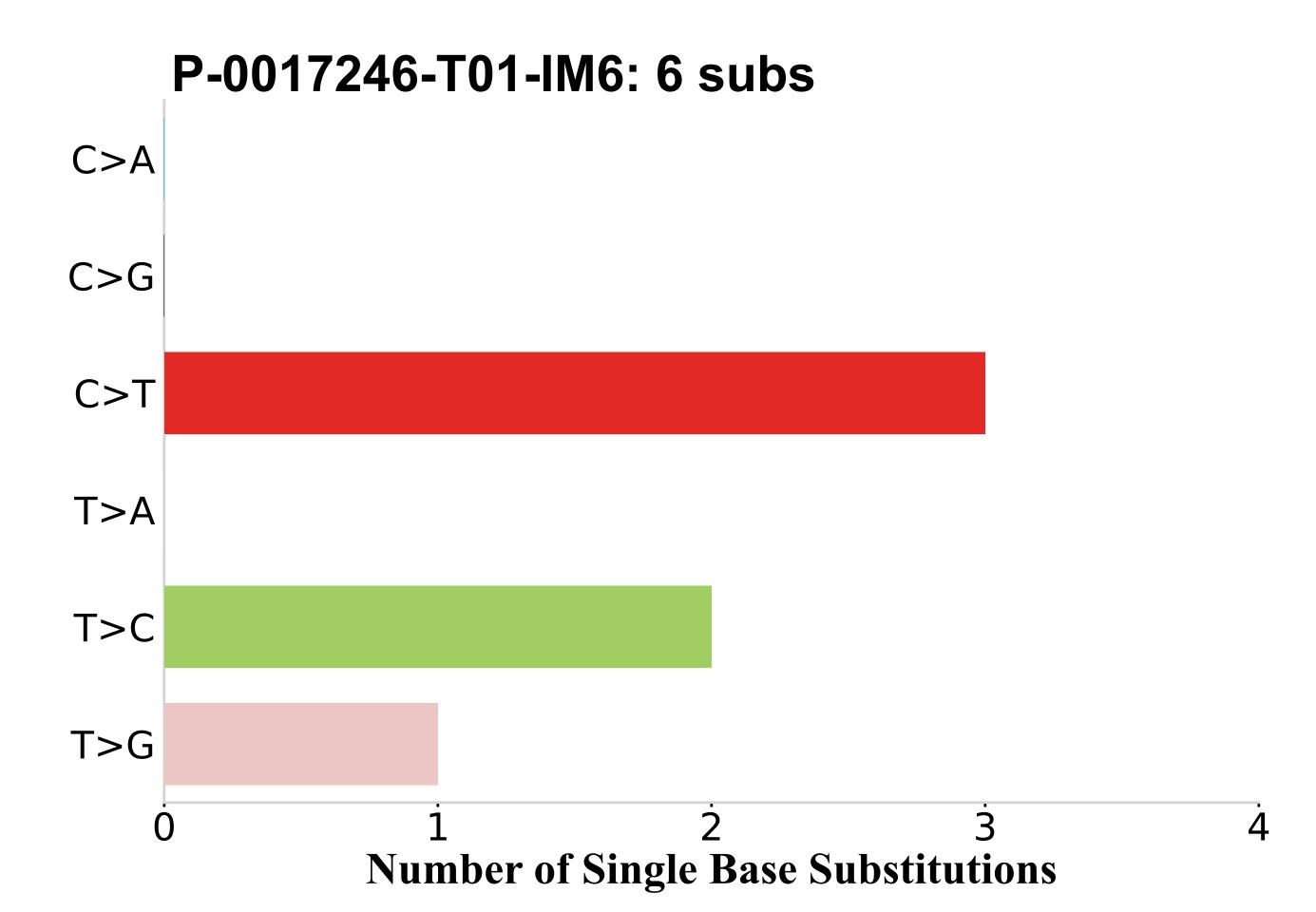


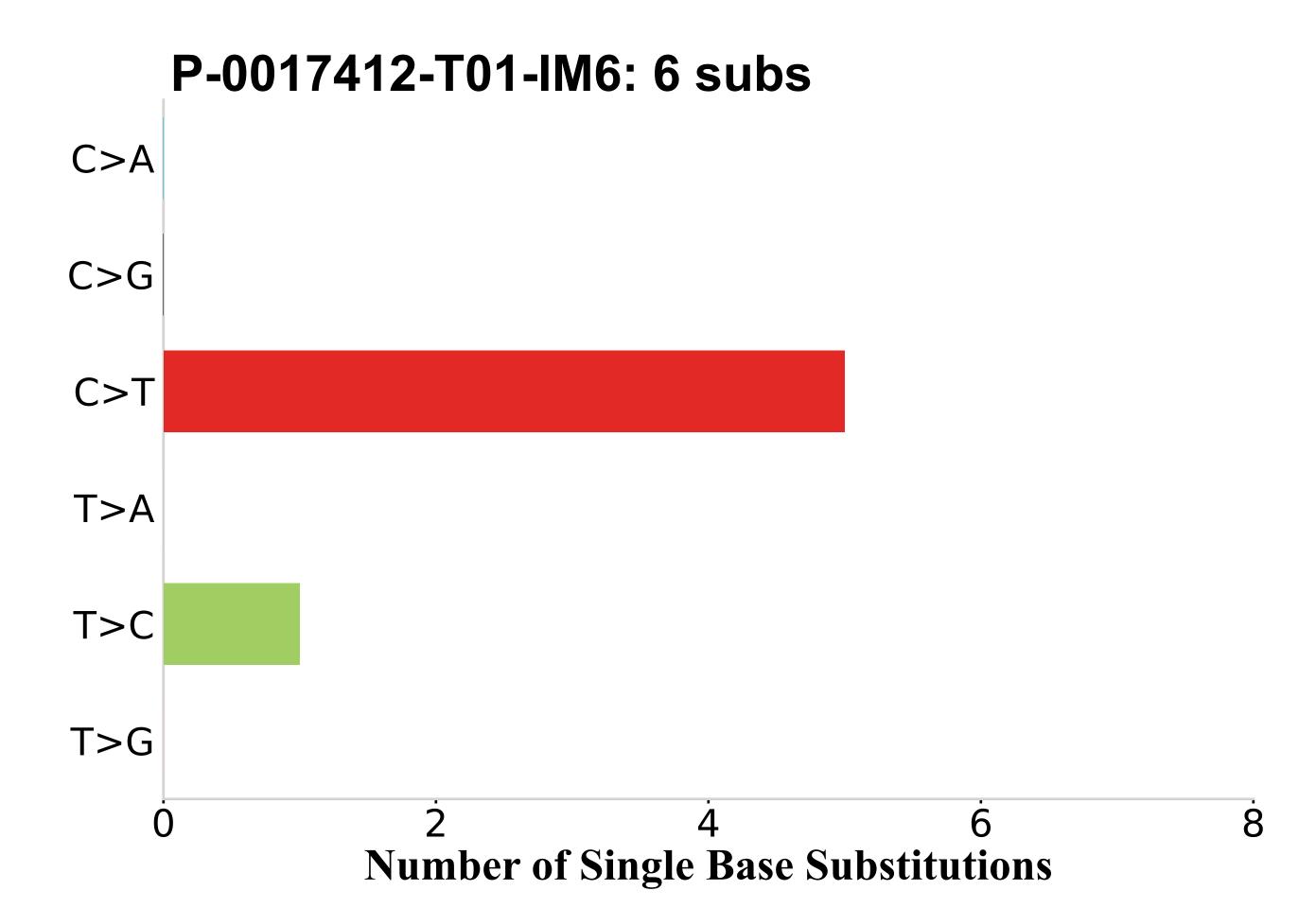






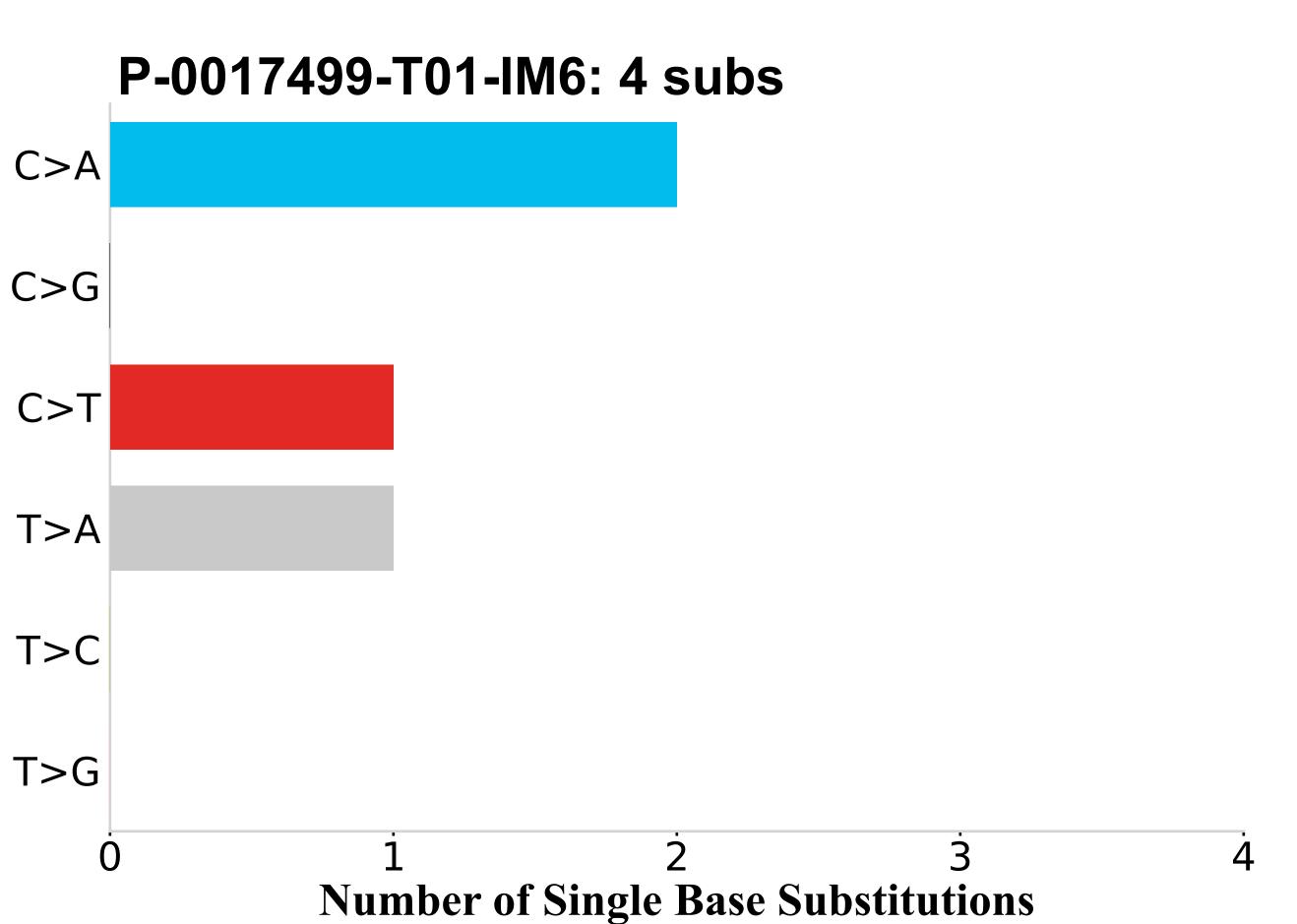


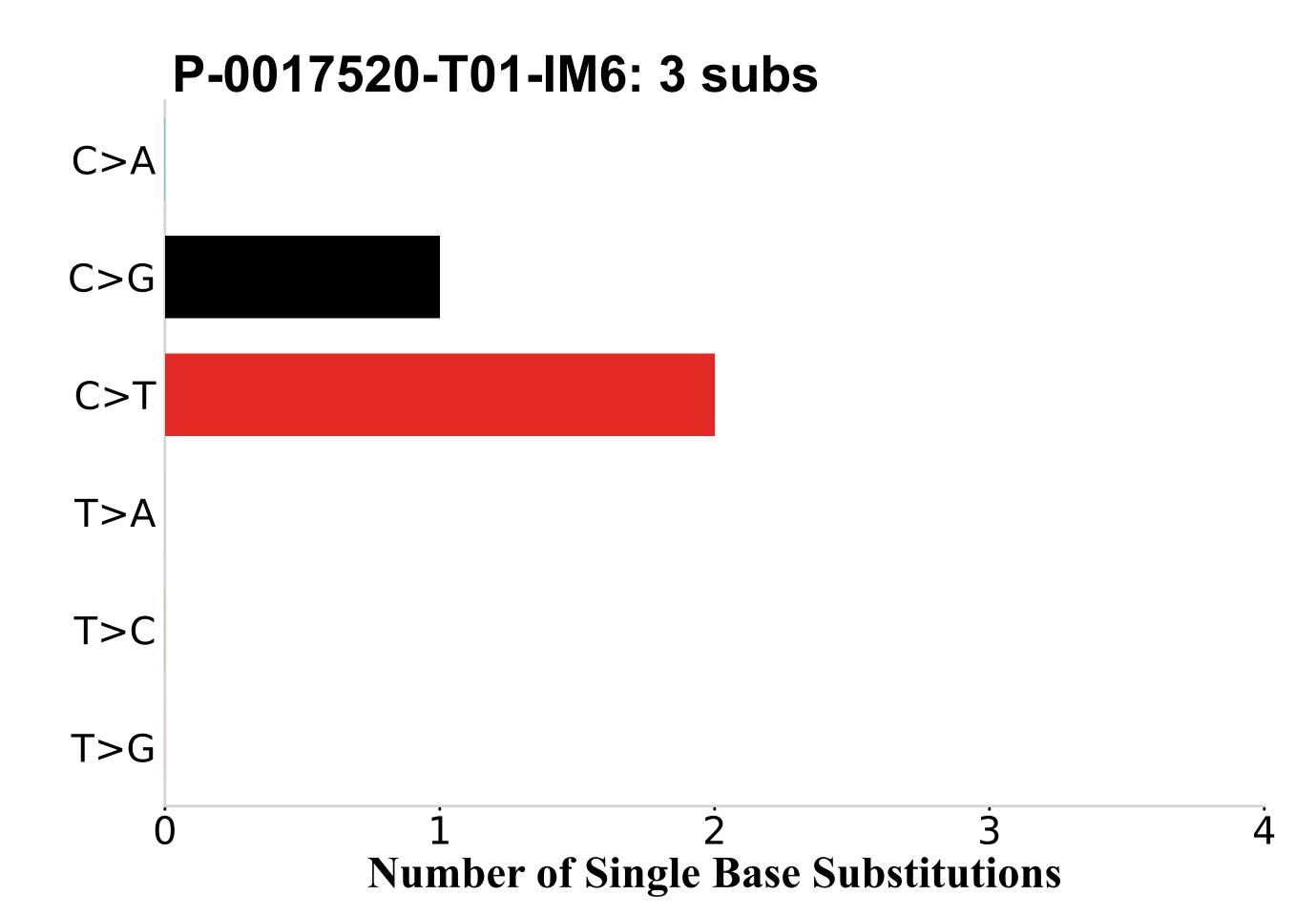




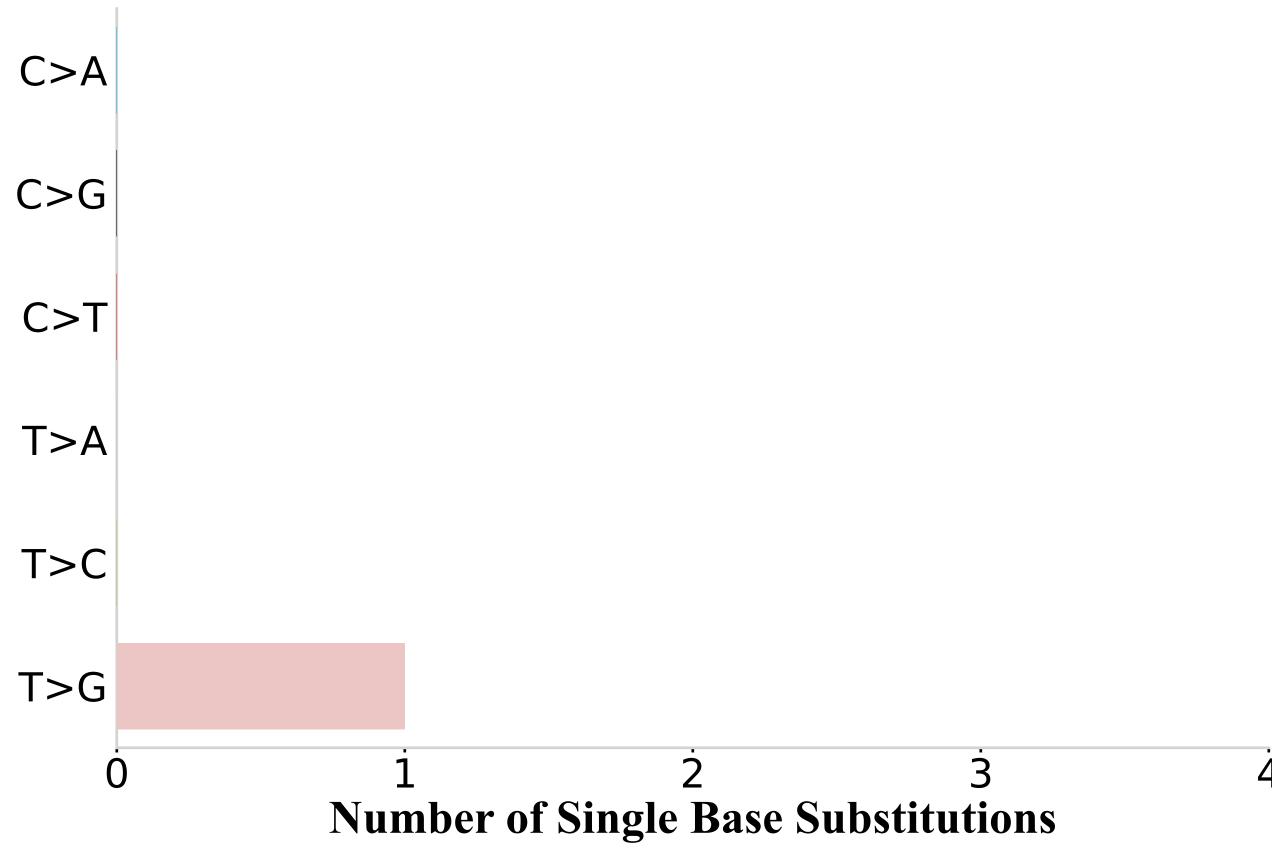
P-0017415-T01-IM6: 9 subs C>A C>G C>T T>AT>C T>G

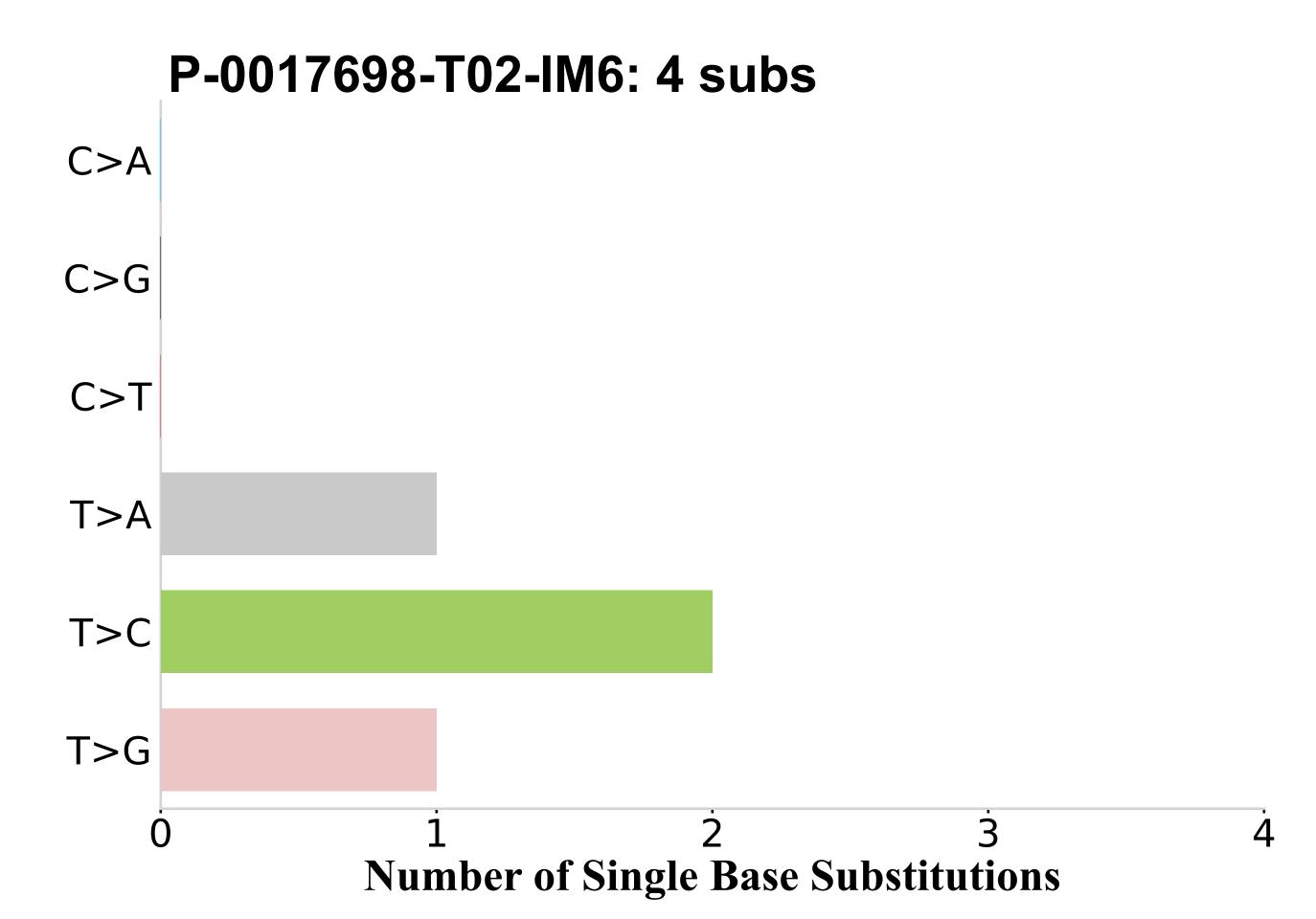
Number of Single Base Substitutions



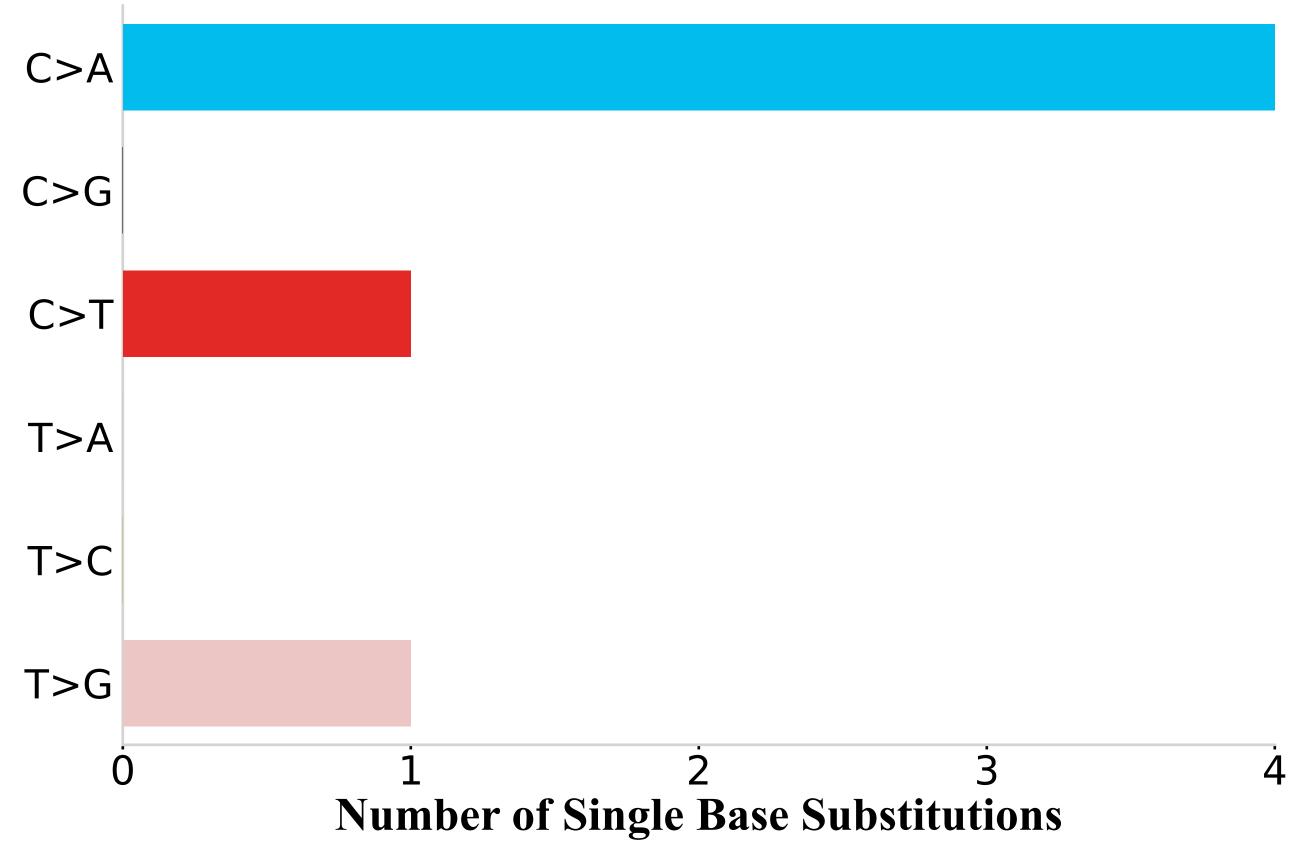


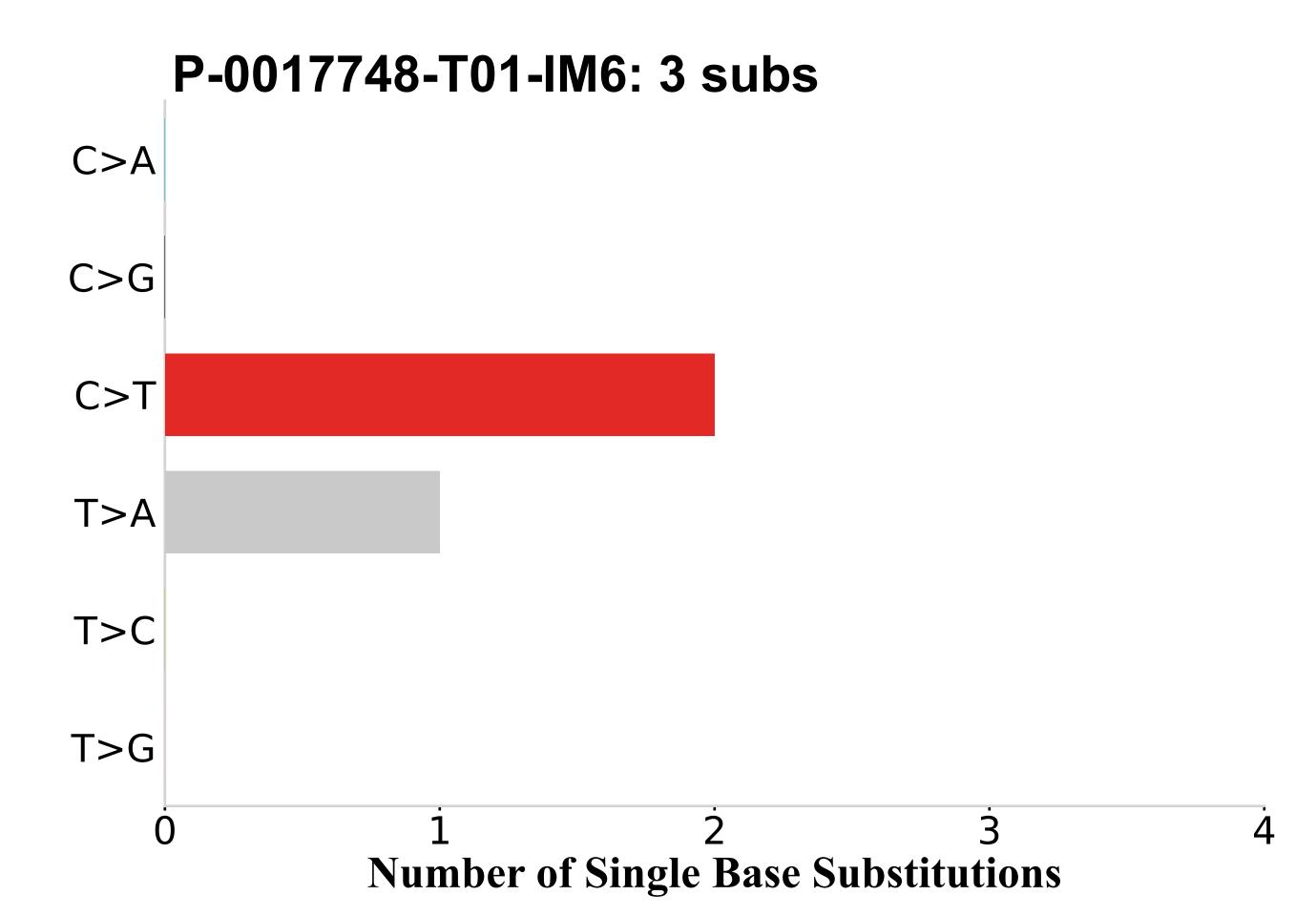
P-0017630-T02-IM6: 1 subs

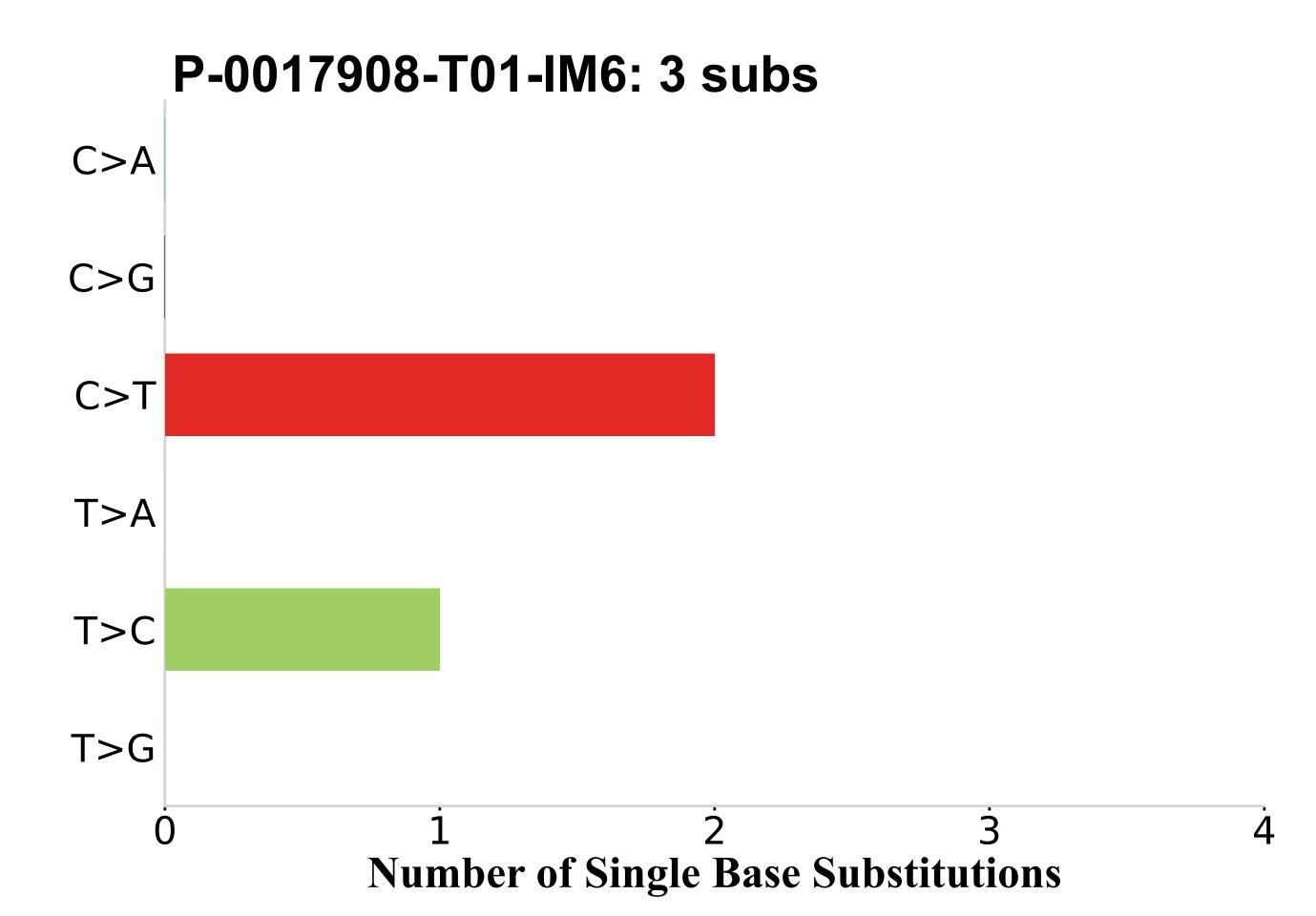


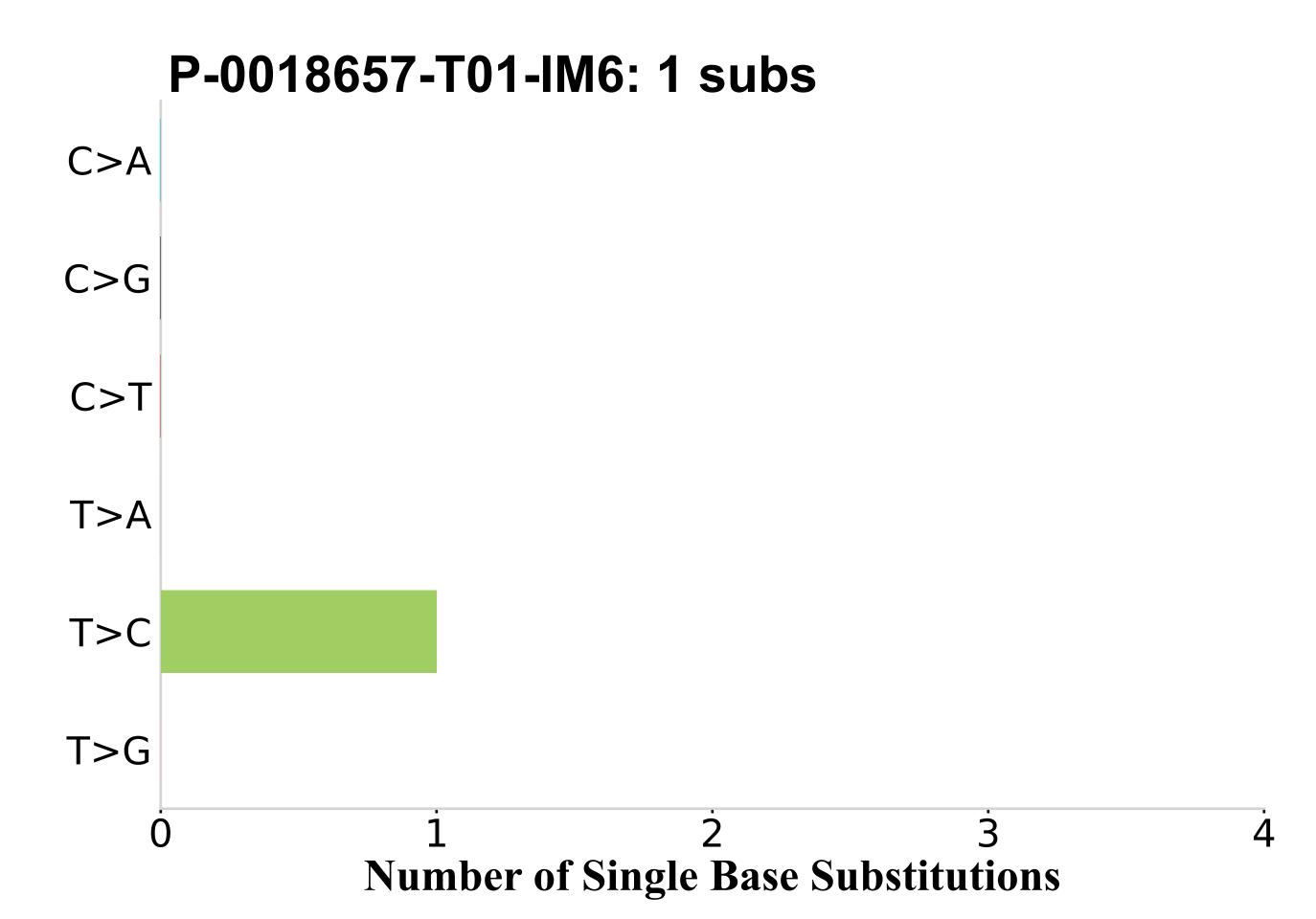


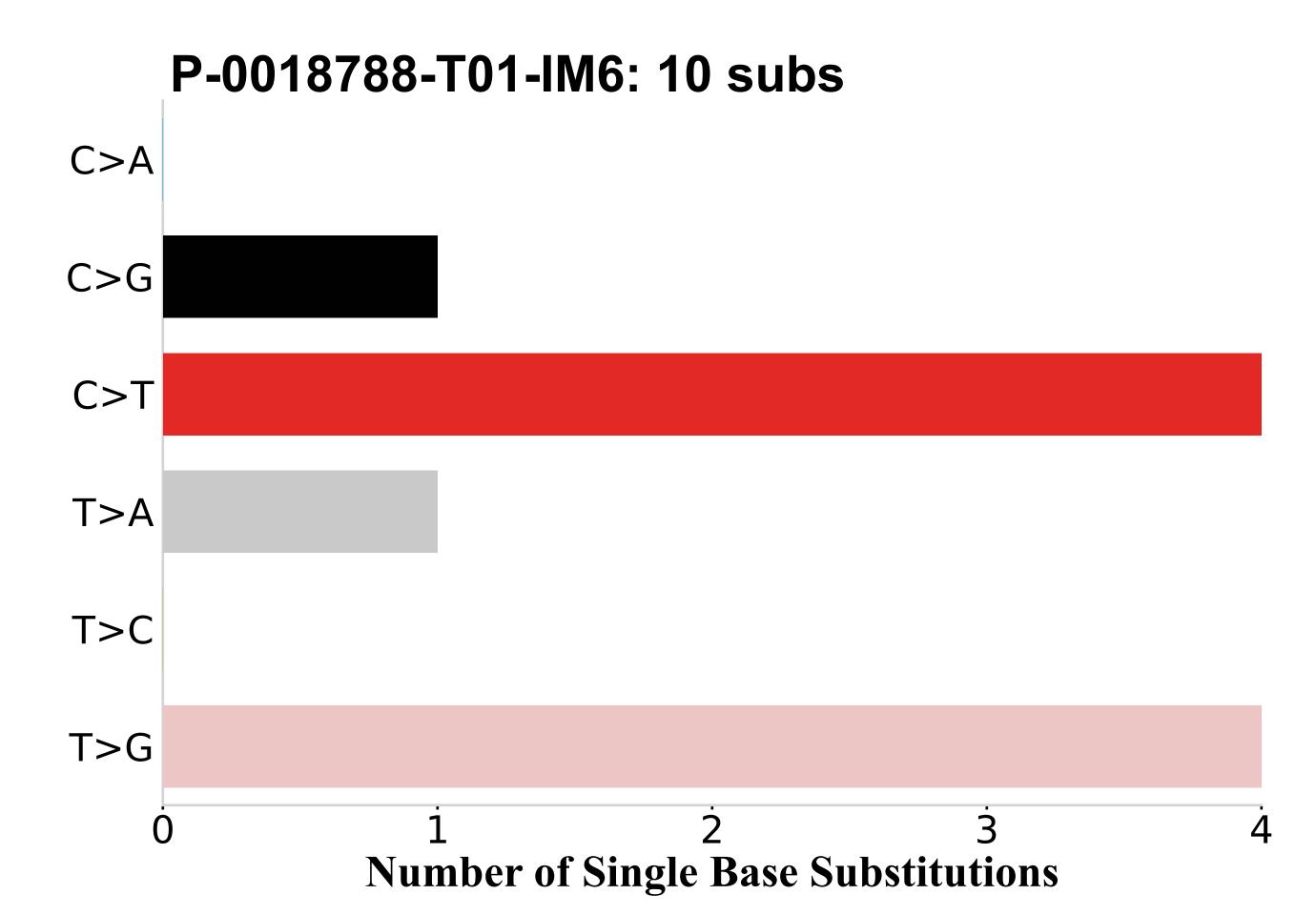


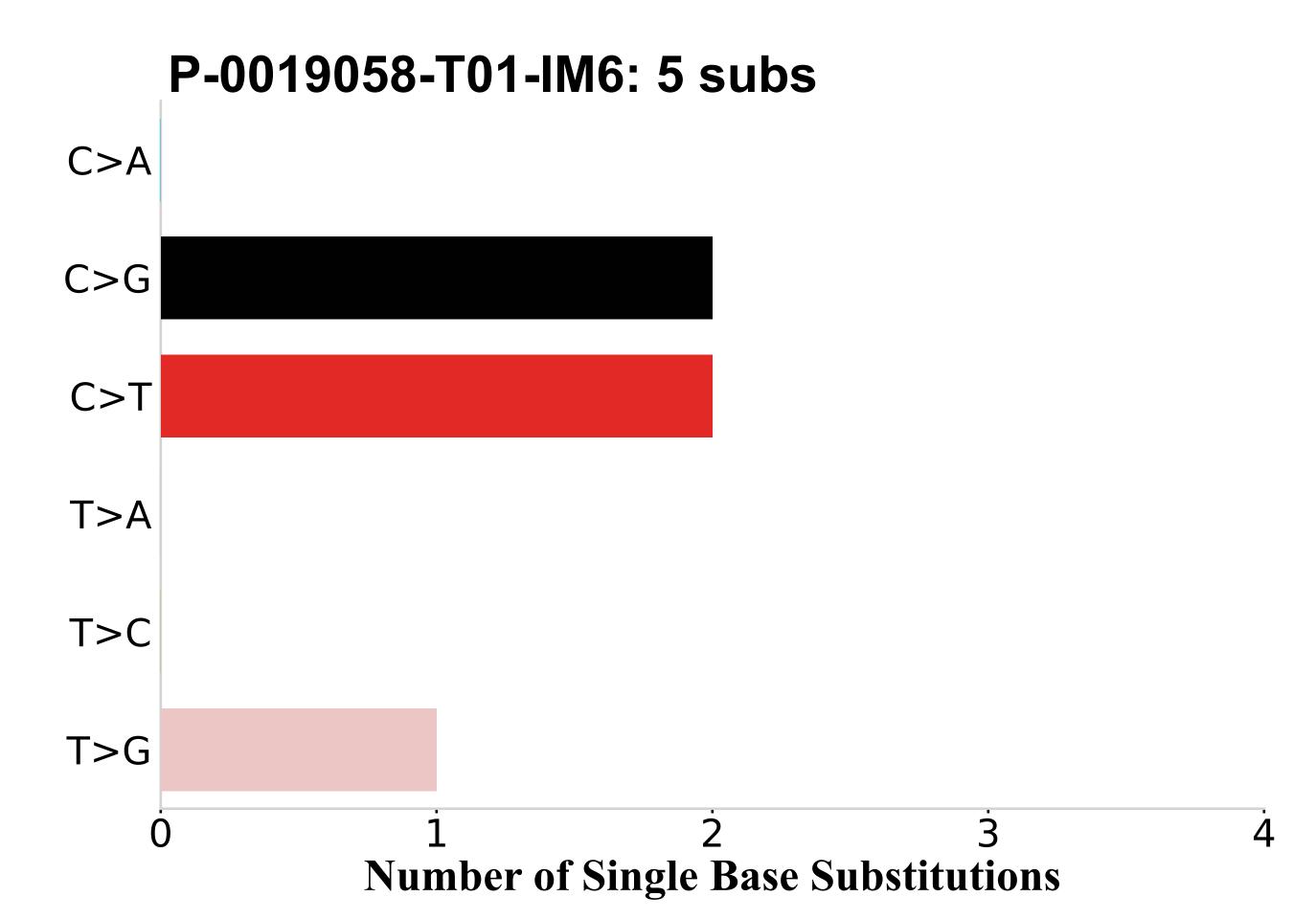


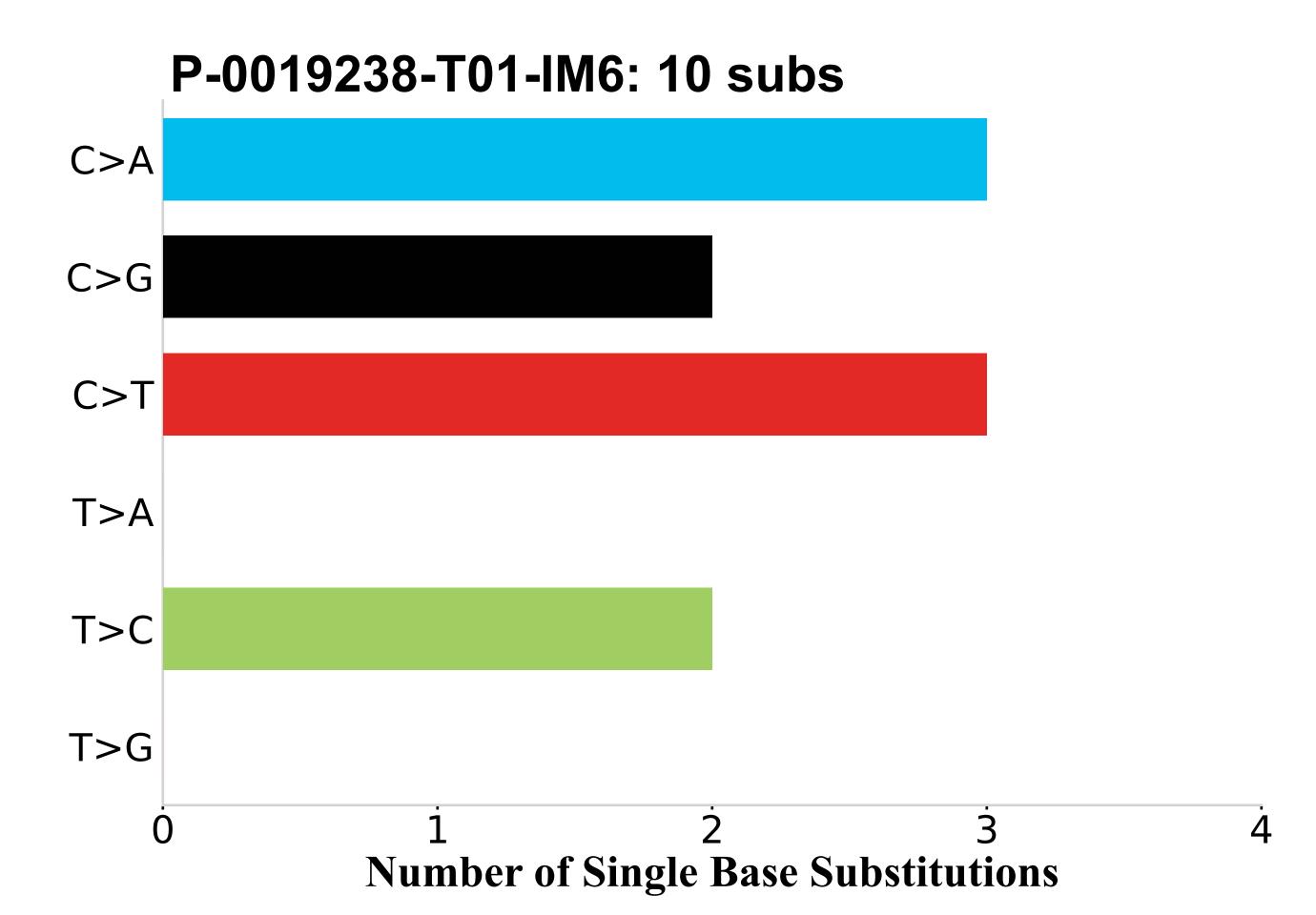


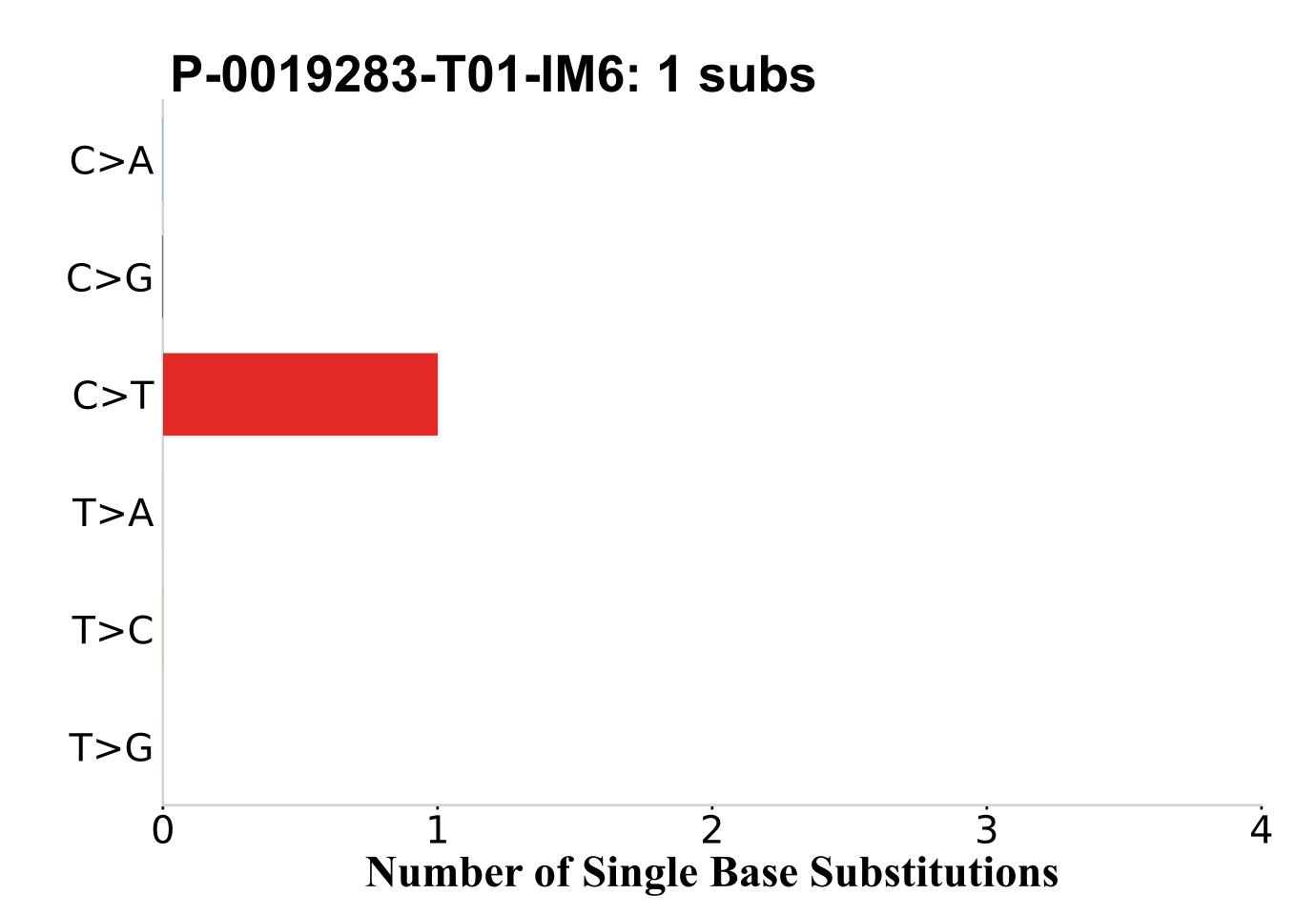


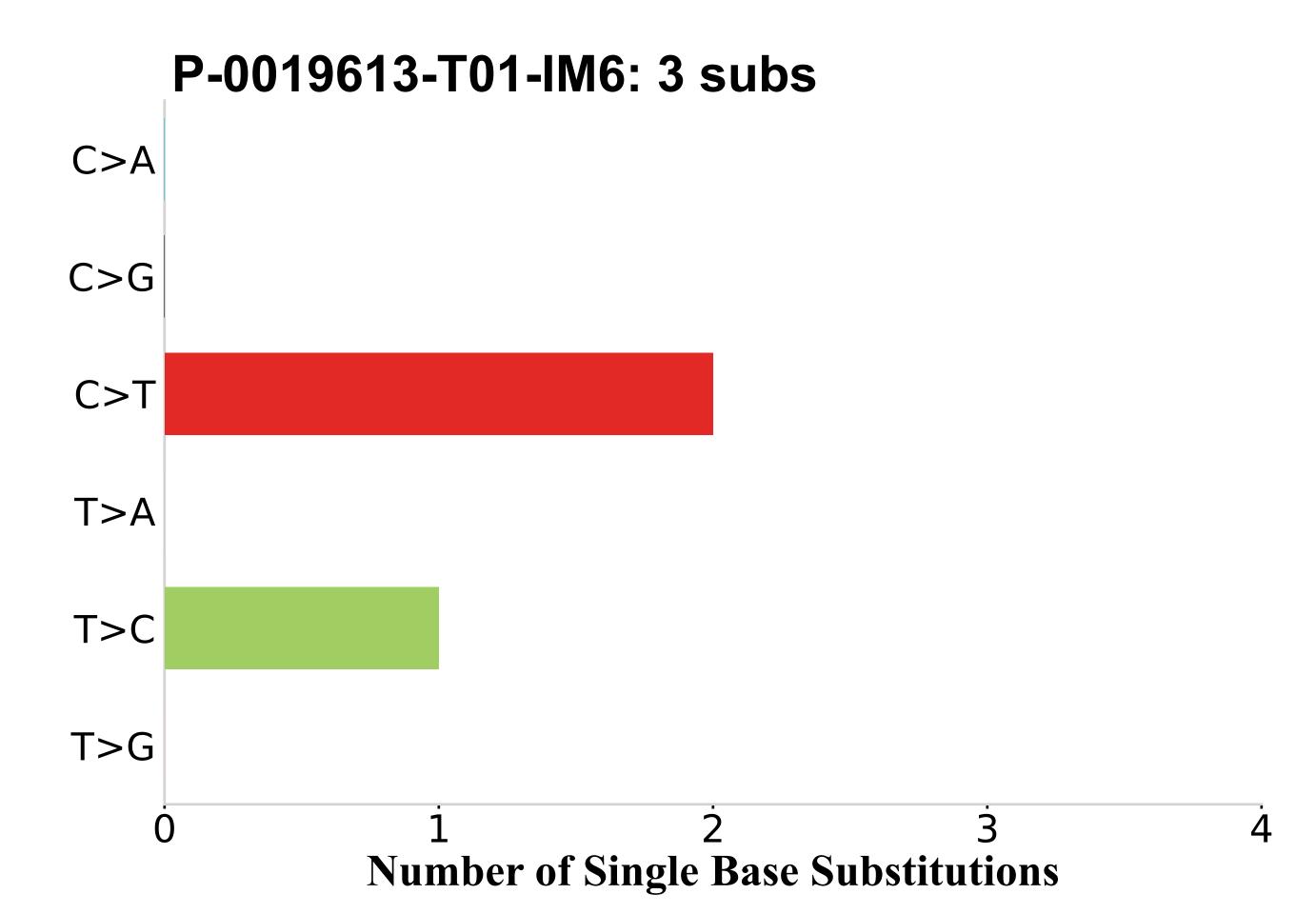


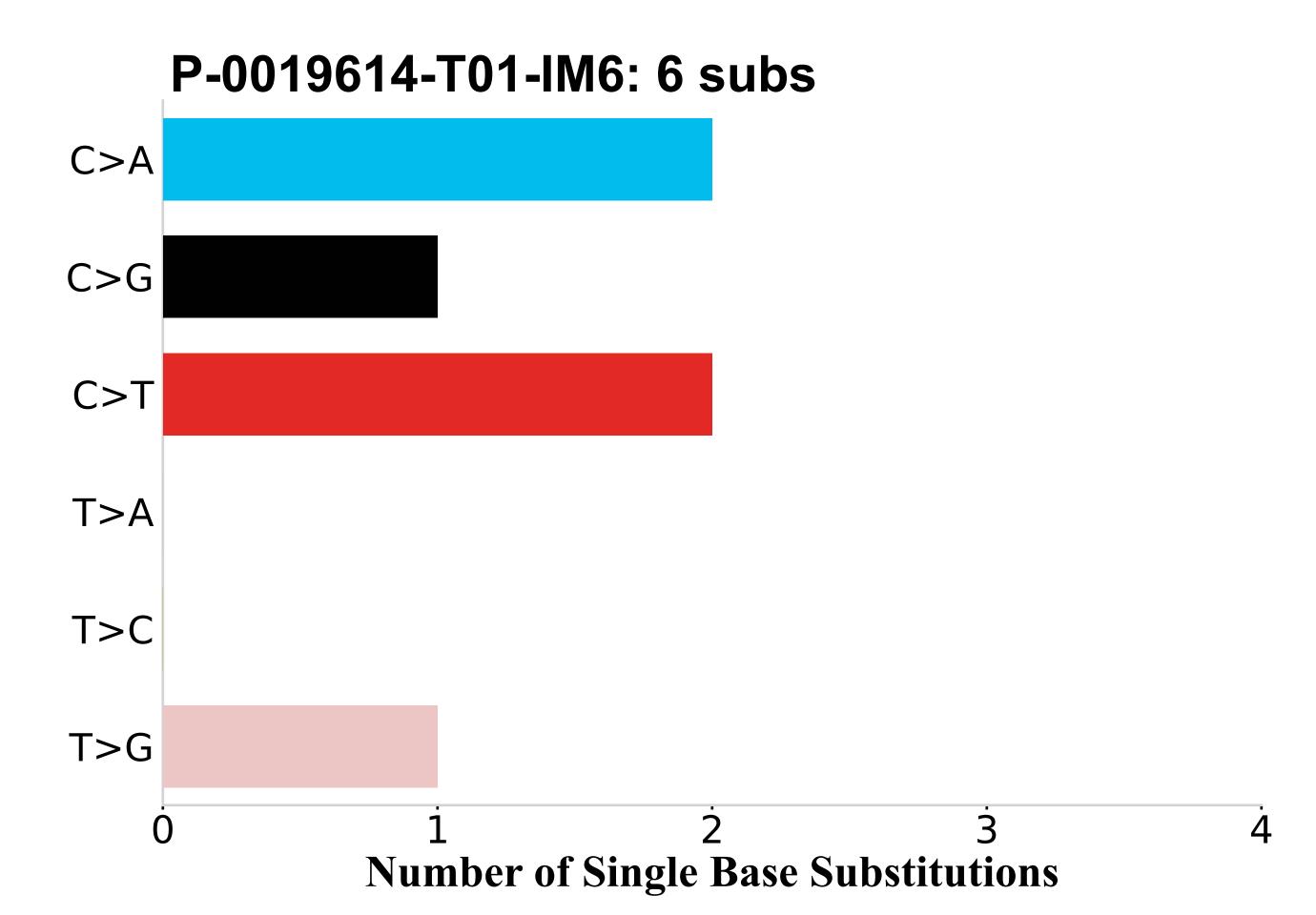


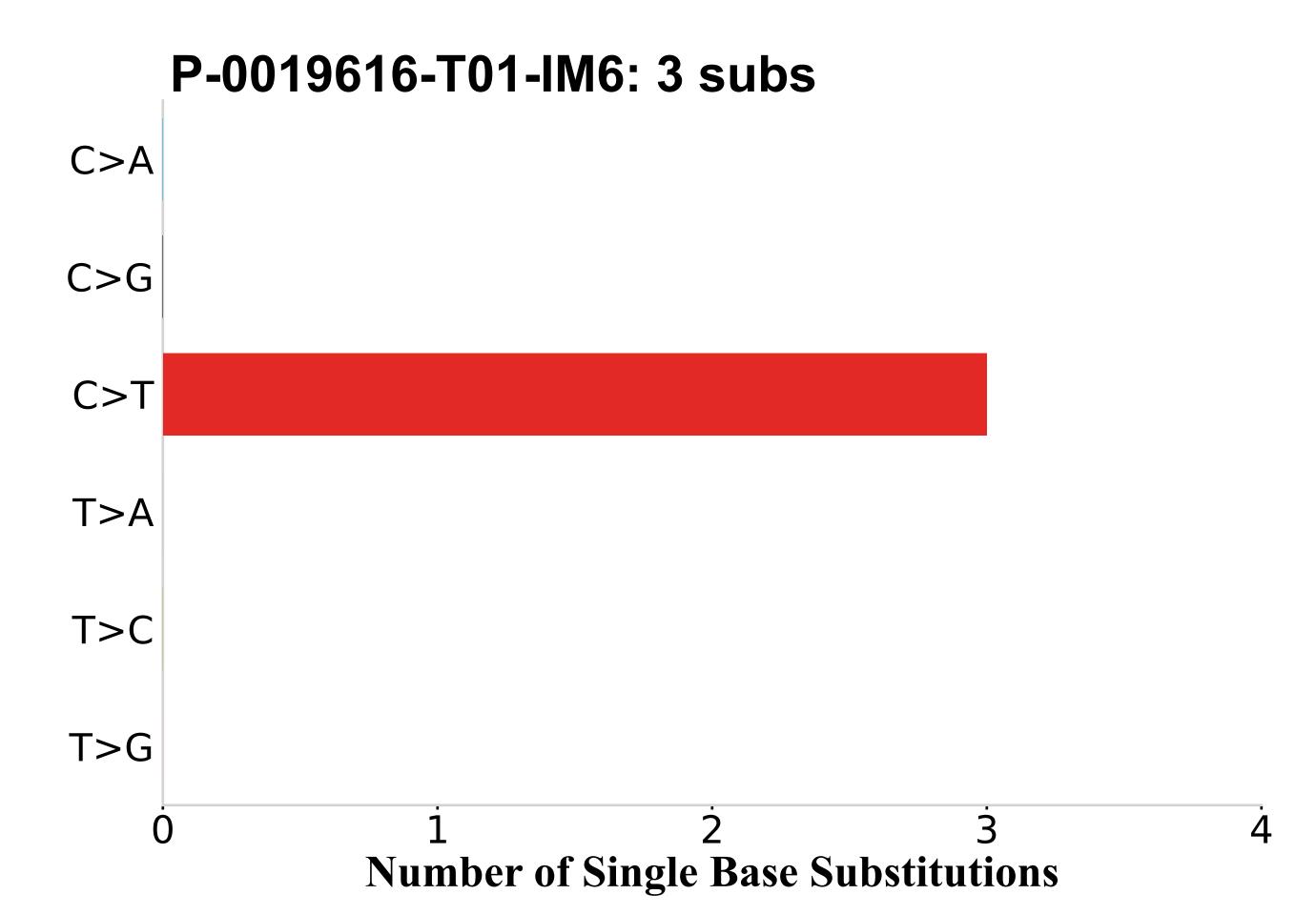




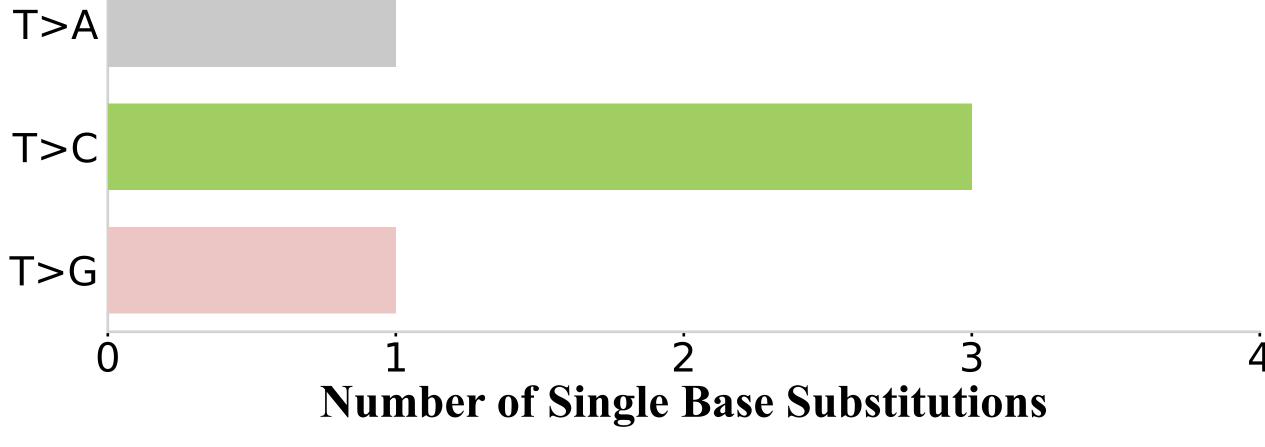


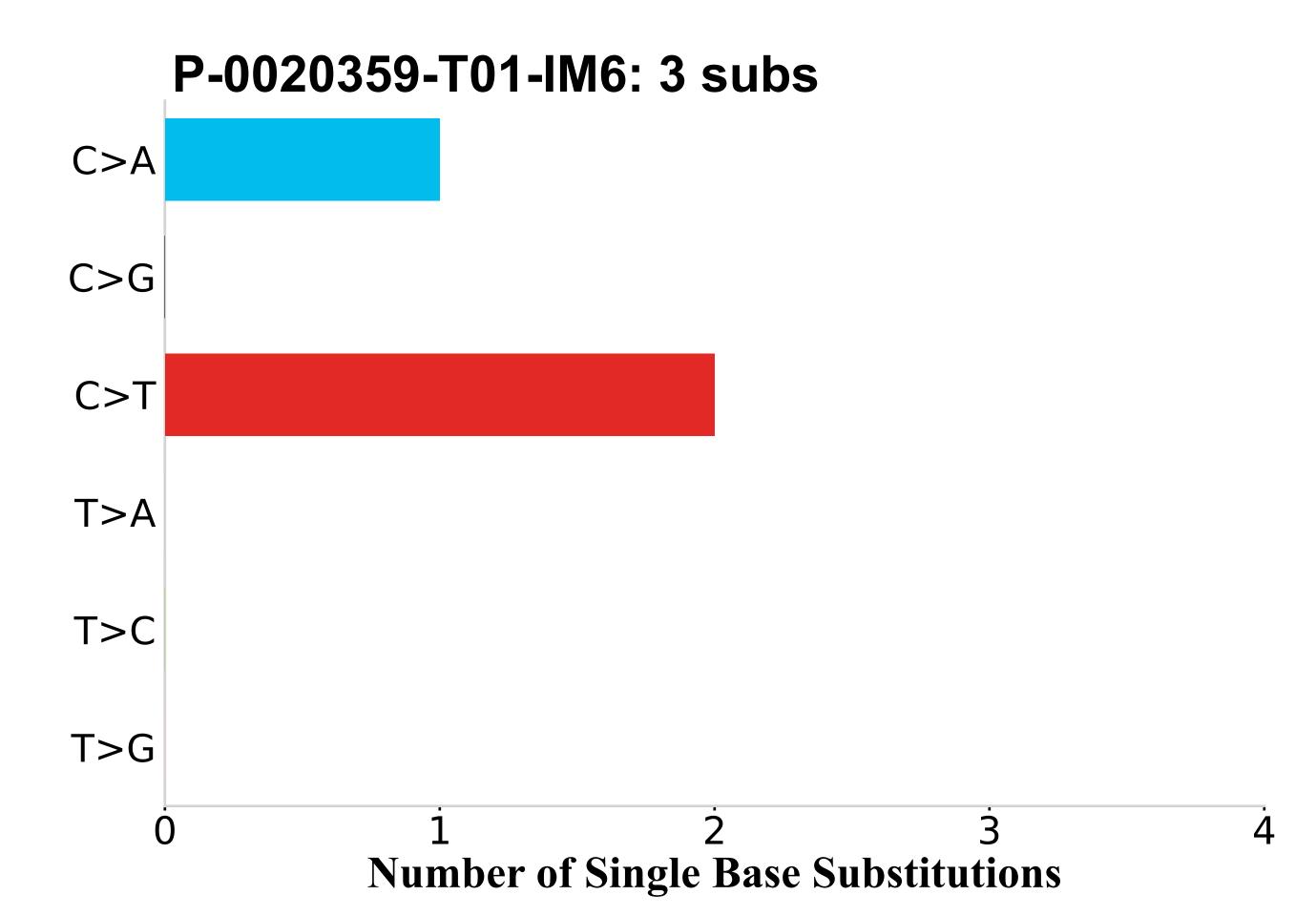




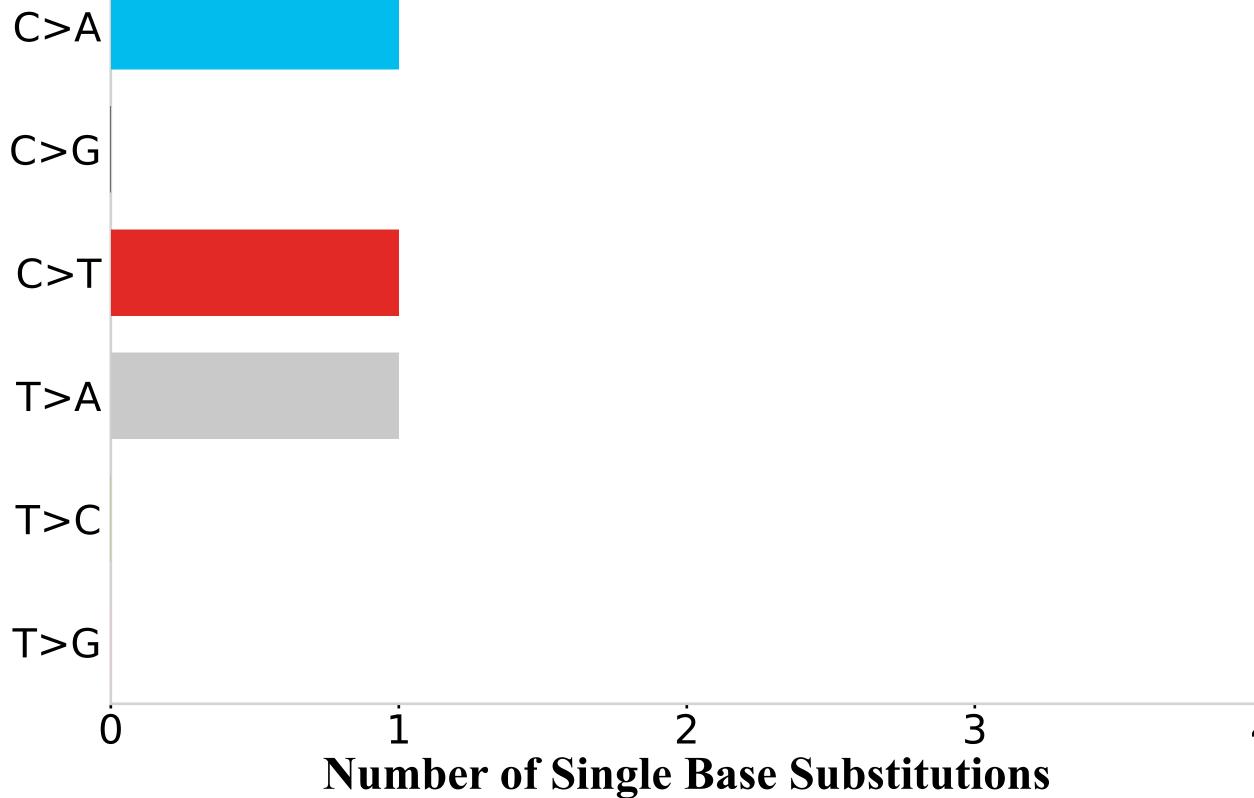


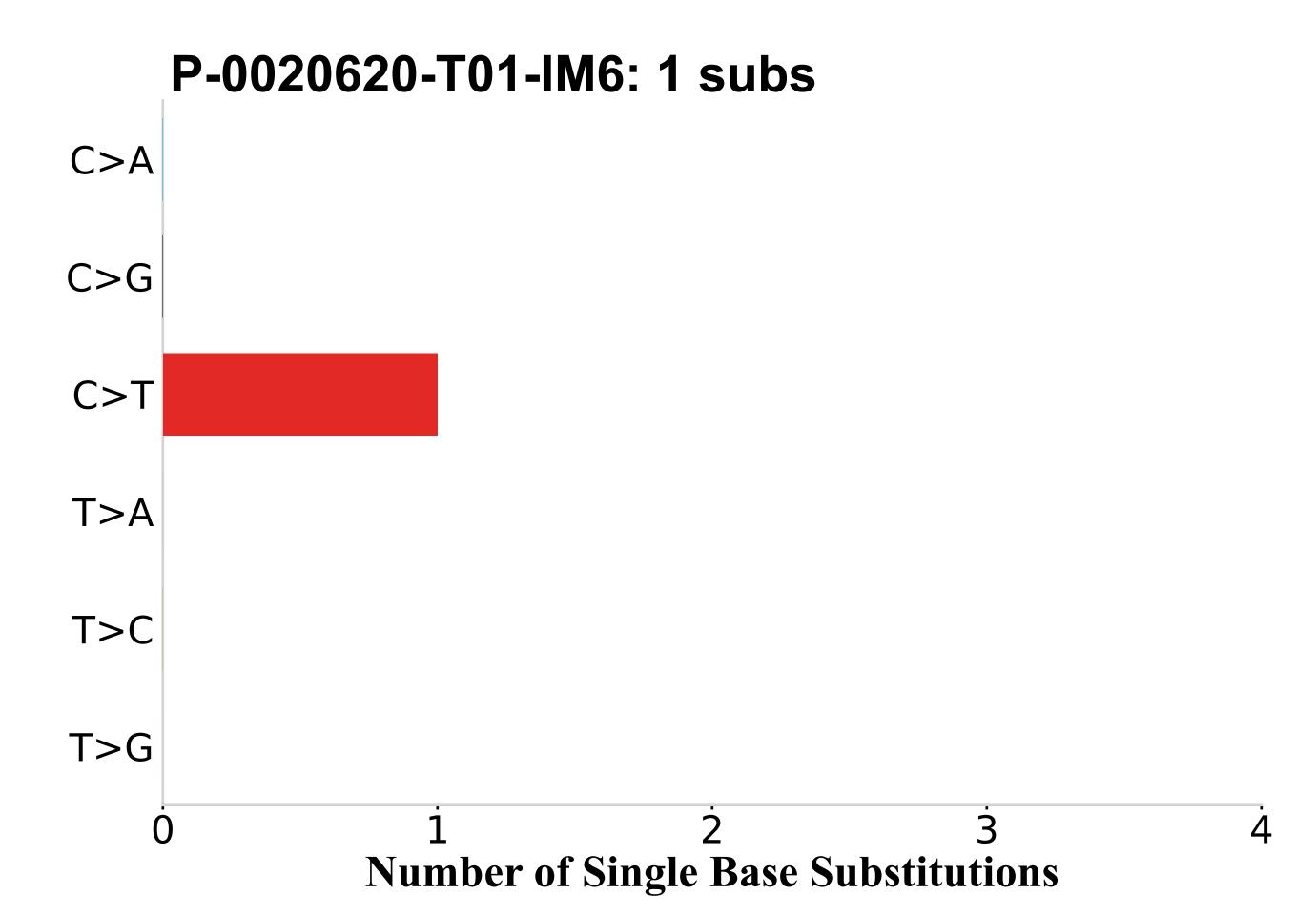
P-0020304-T01-IM6: 10 subs C>A C>G C>T

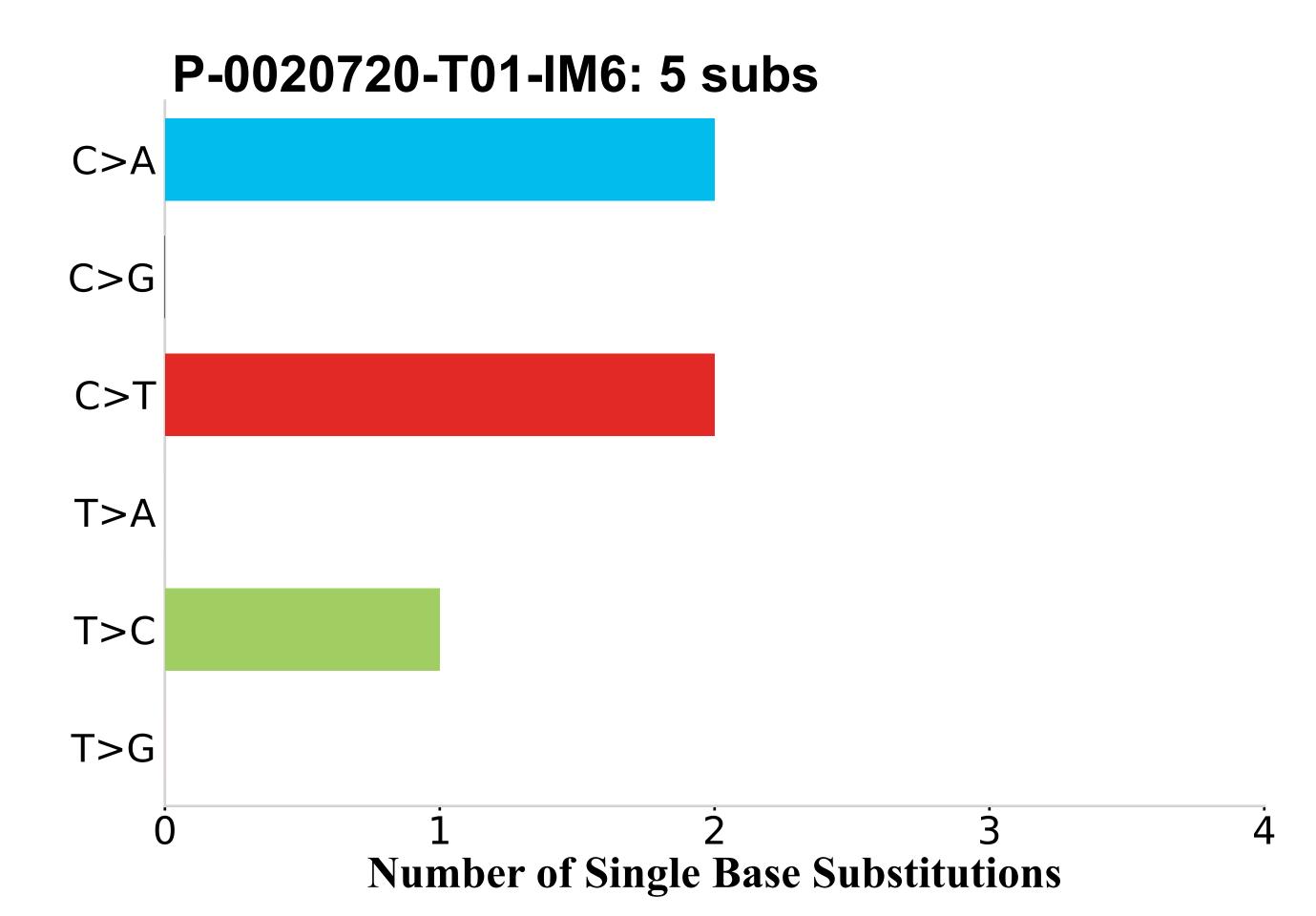


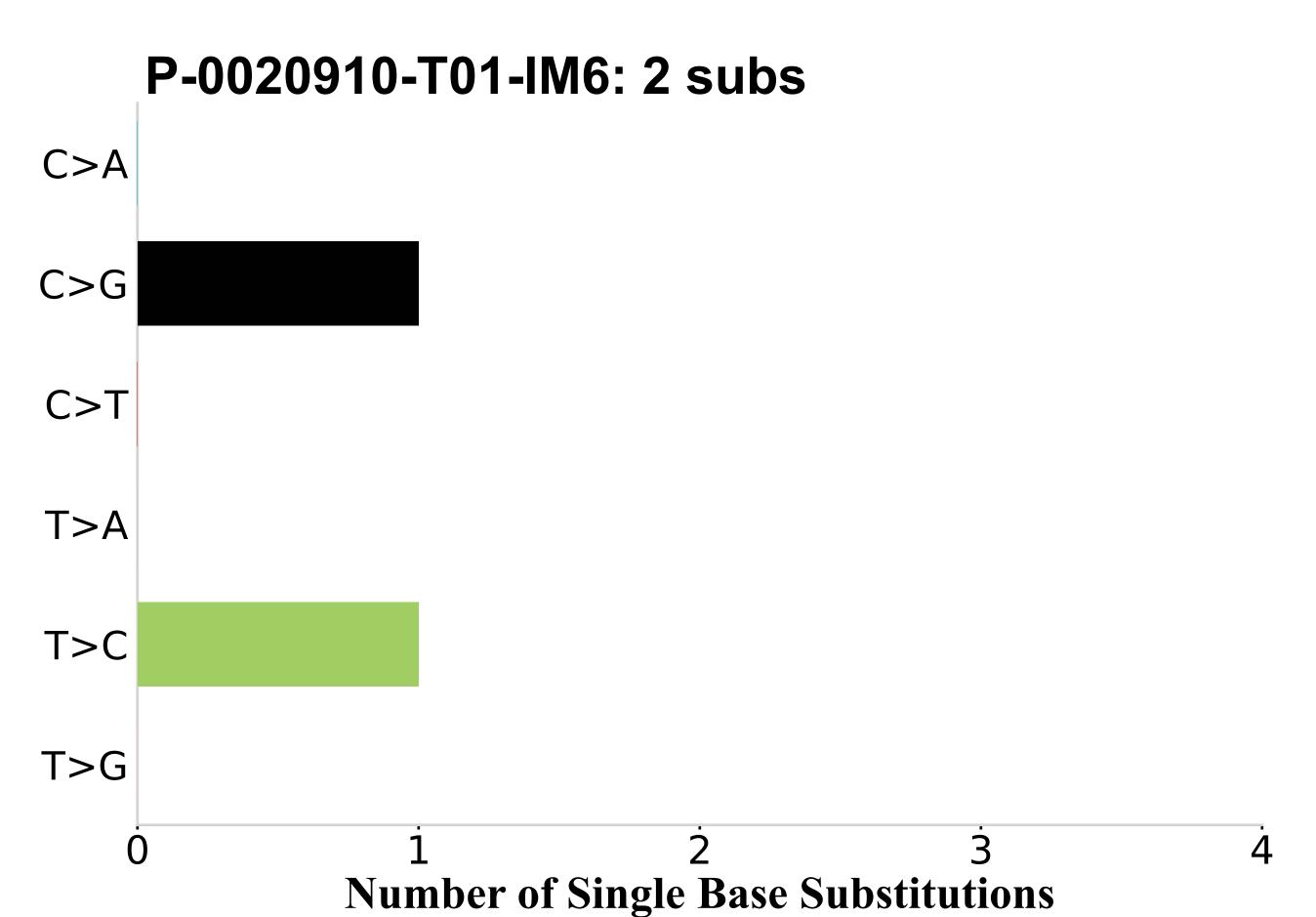


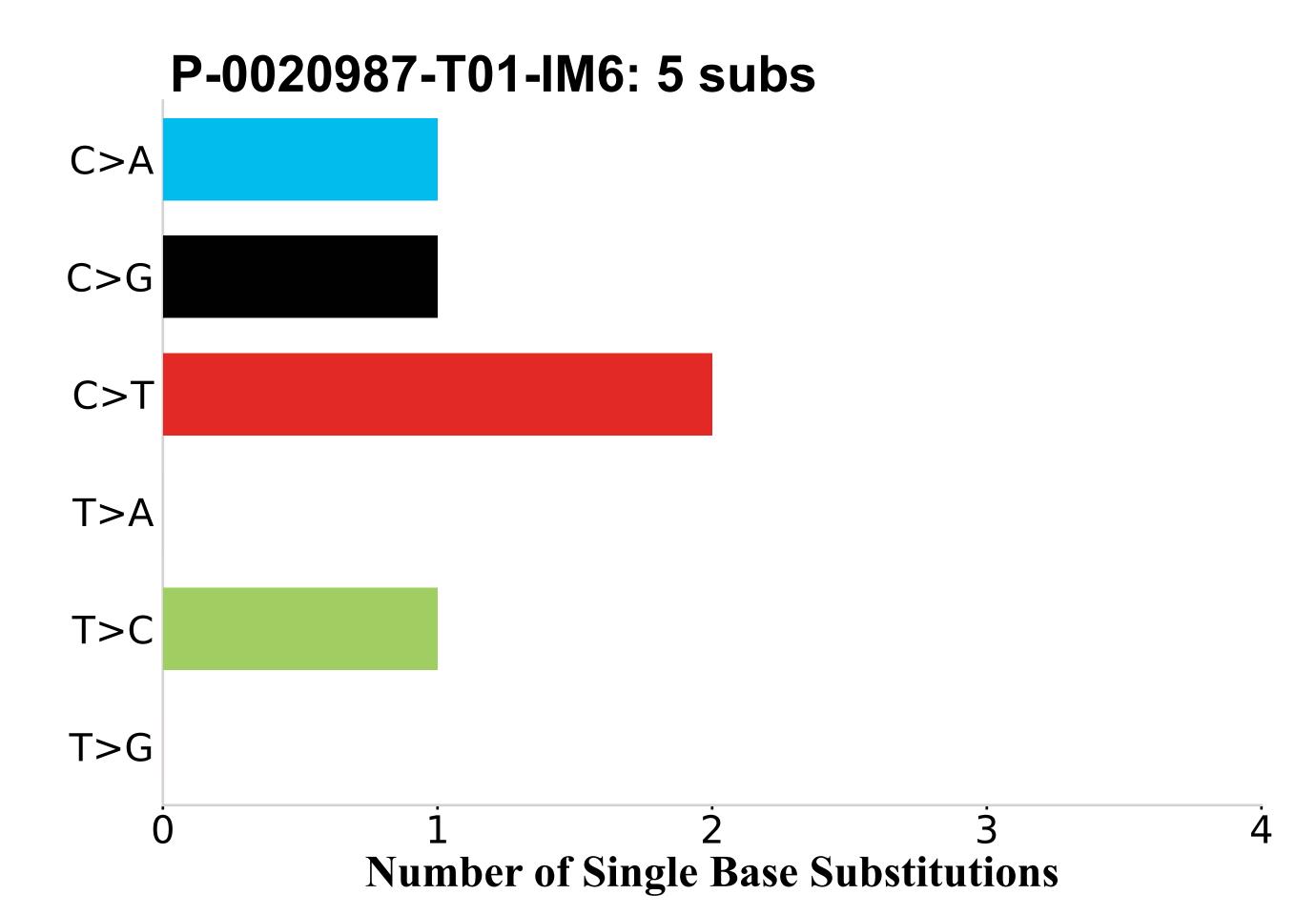
P-0020428-T01-IM6: 3 subs

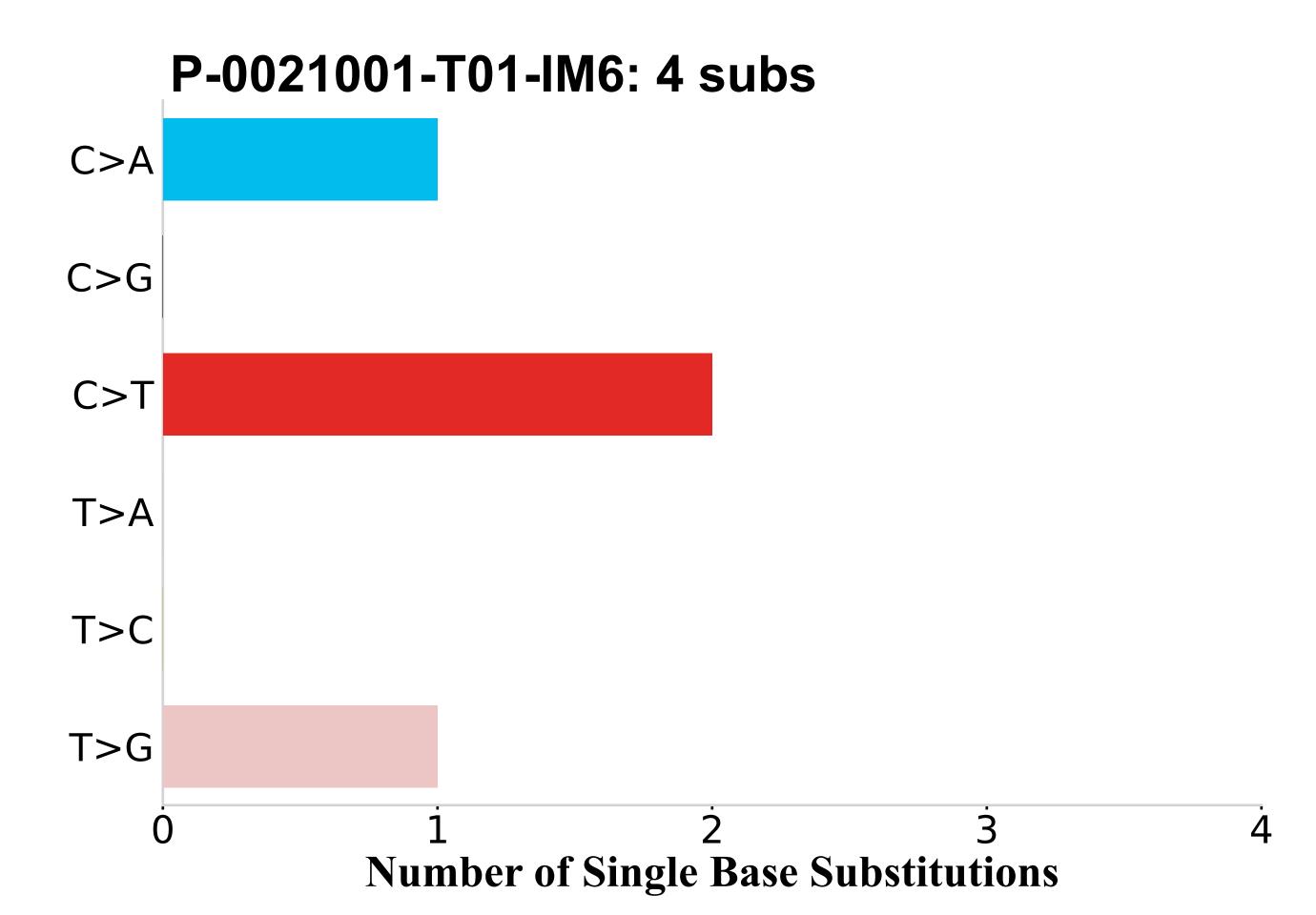




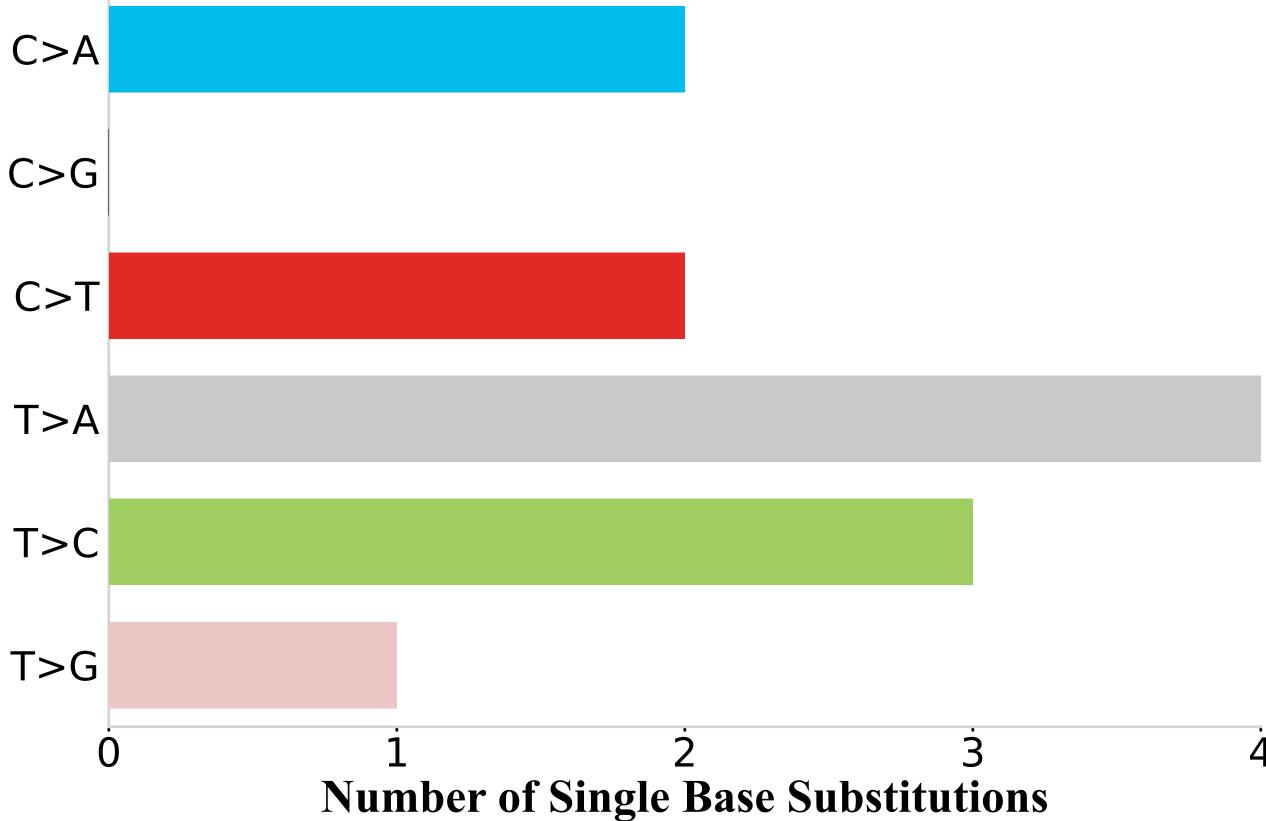


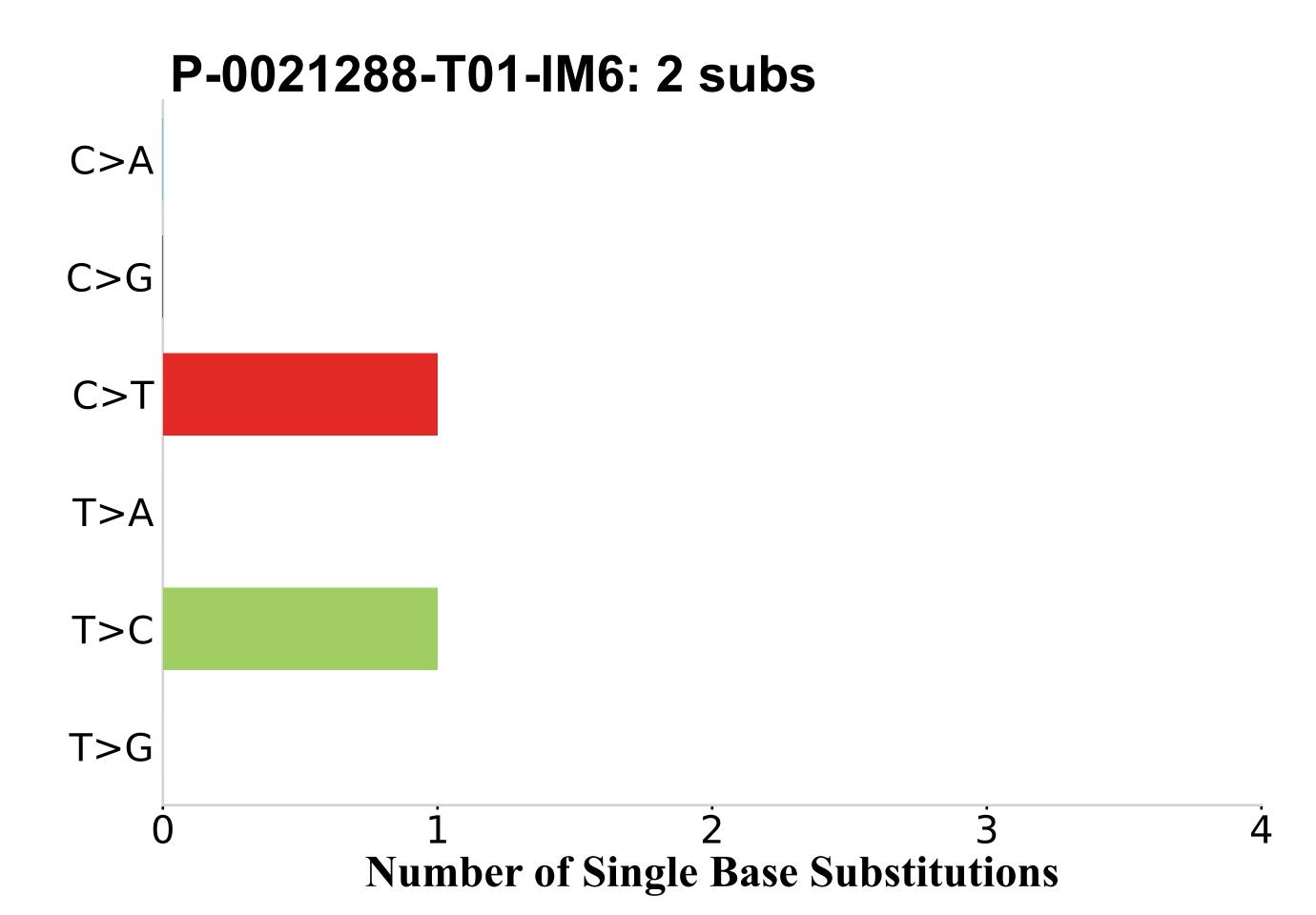


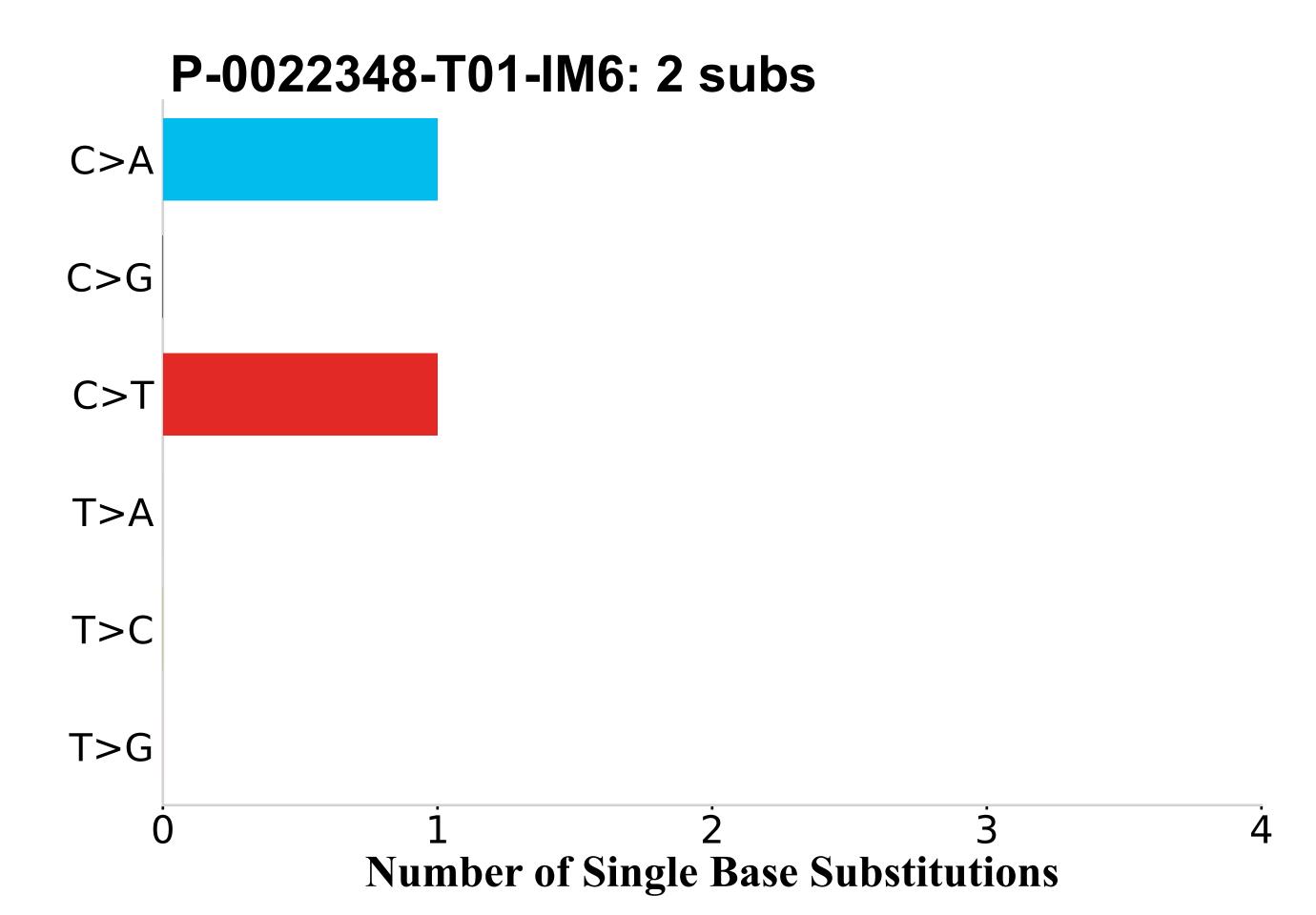


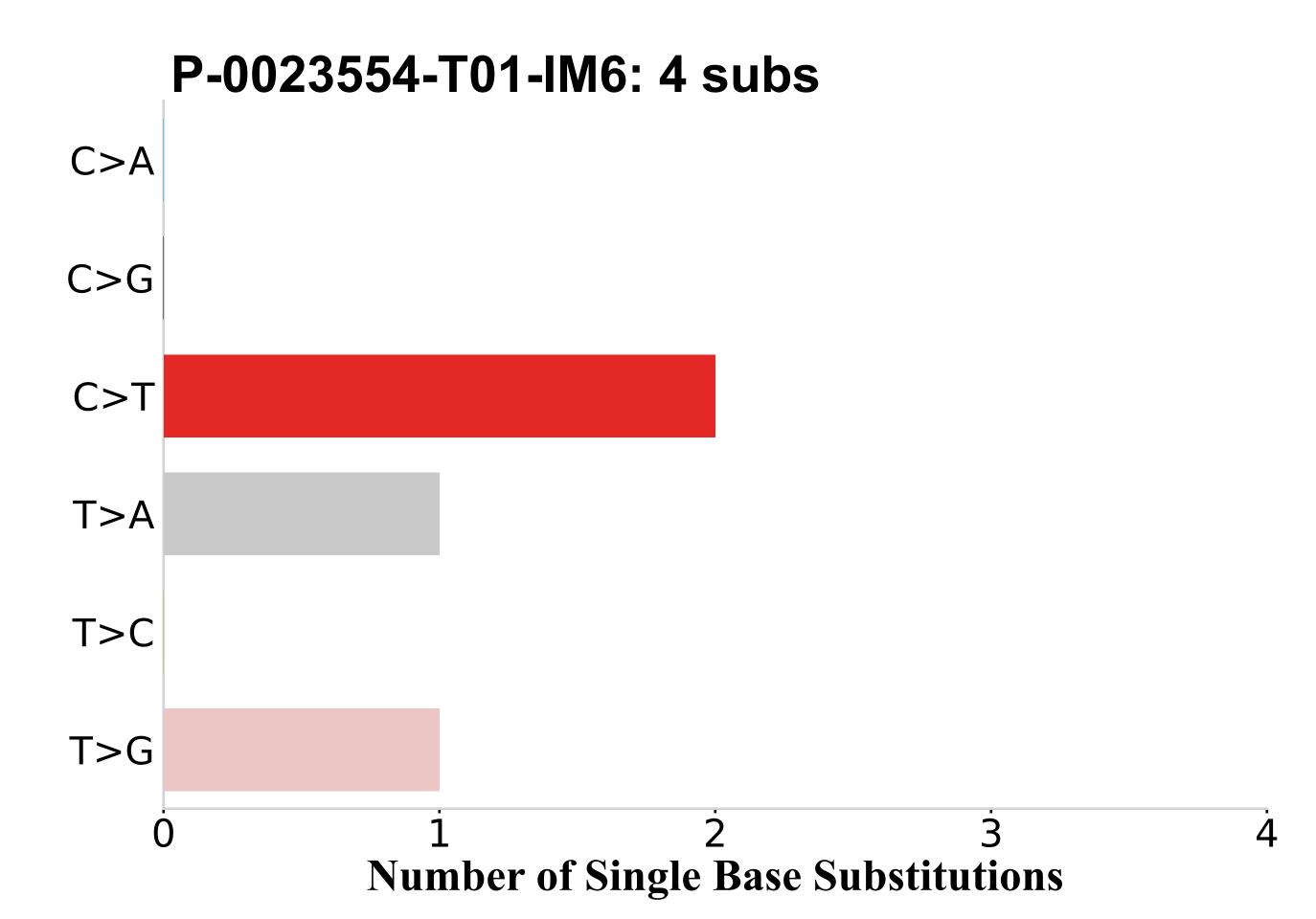


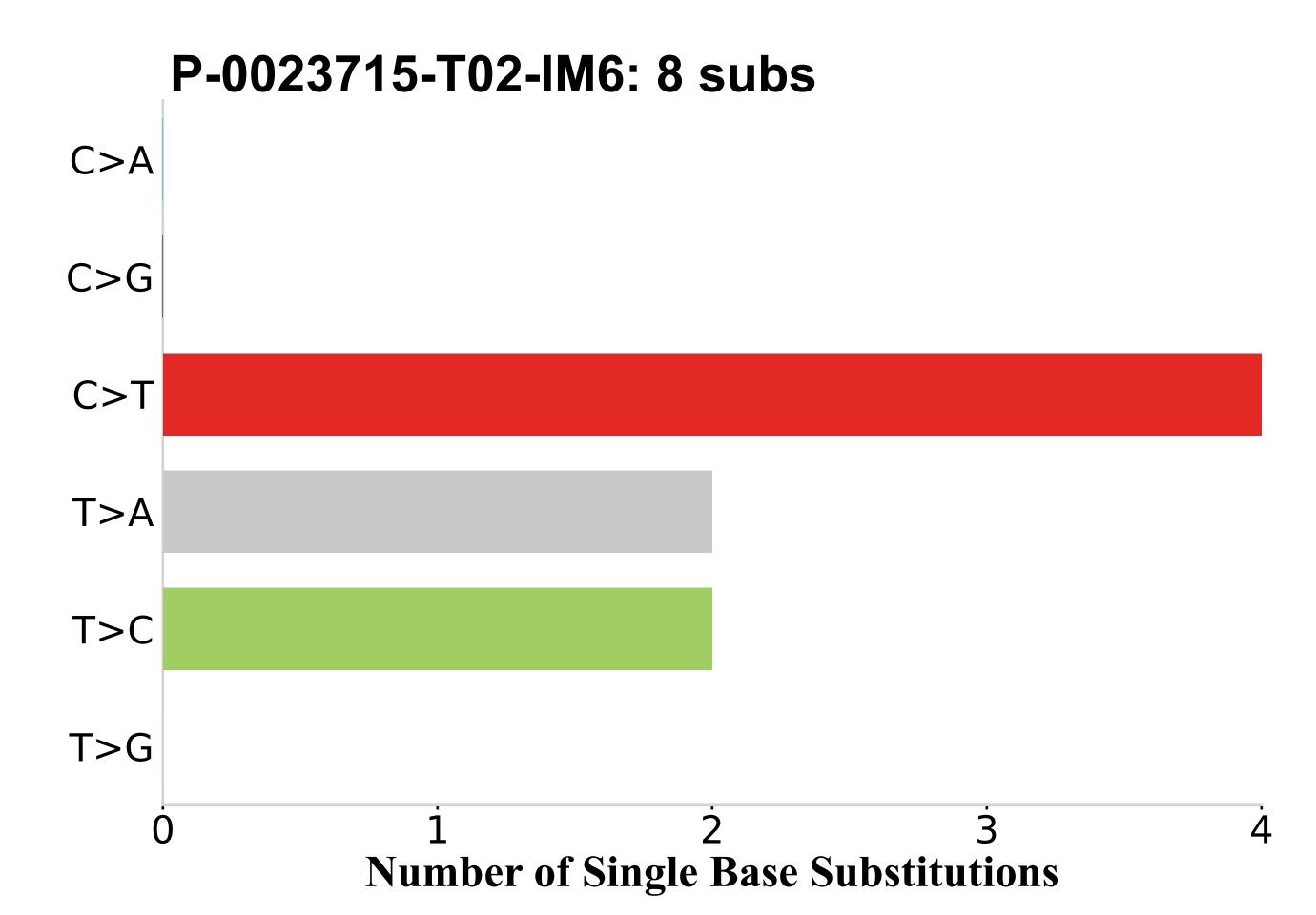
P-0021059-T01-IM6: 12 subs

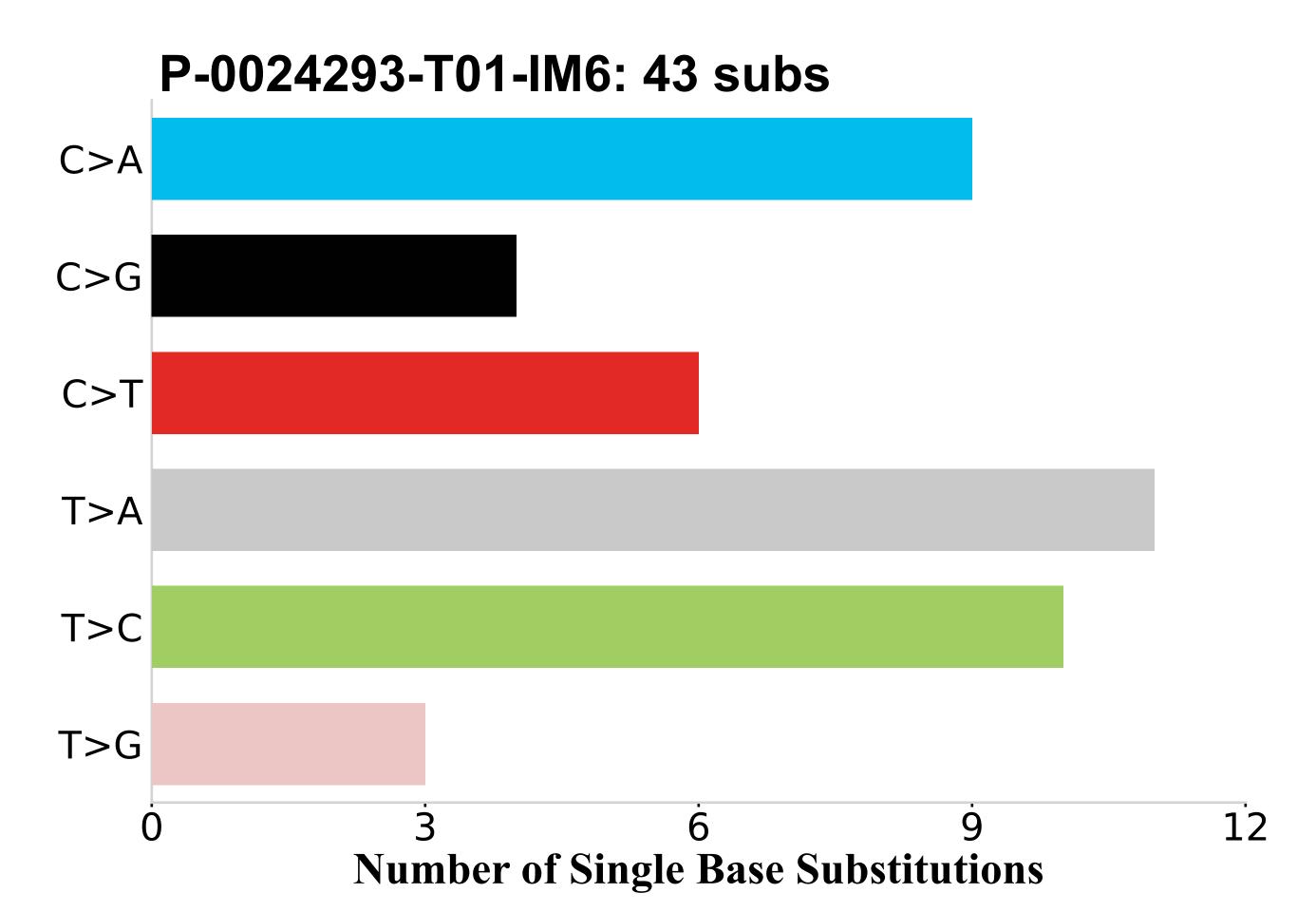


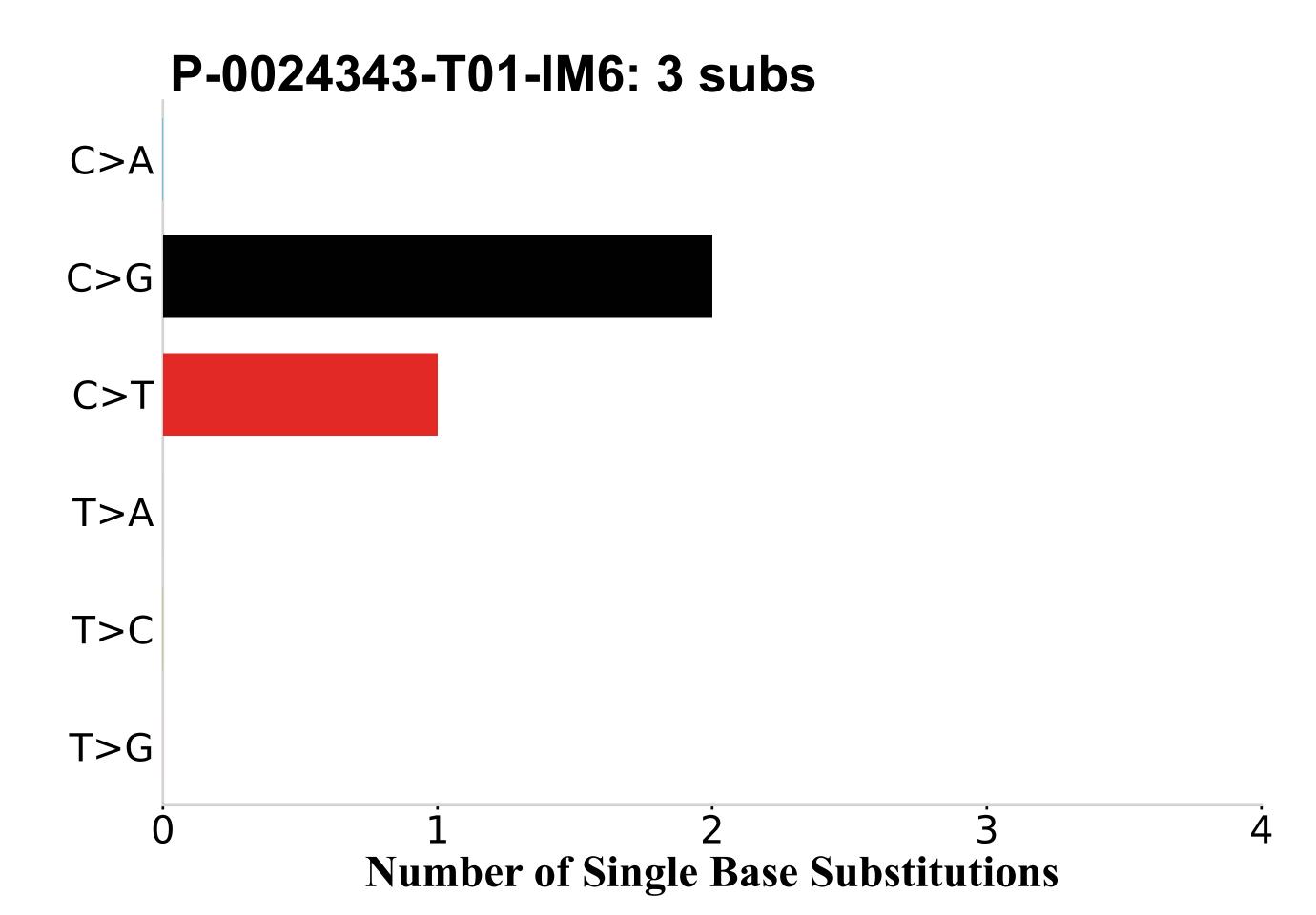


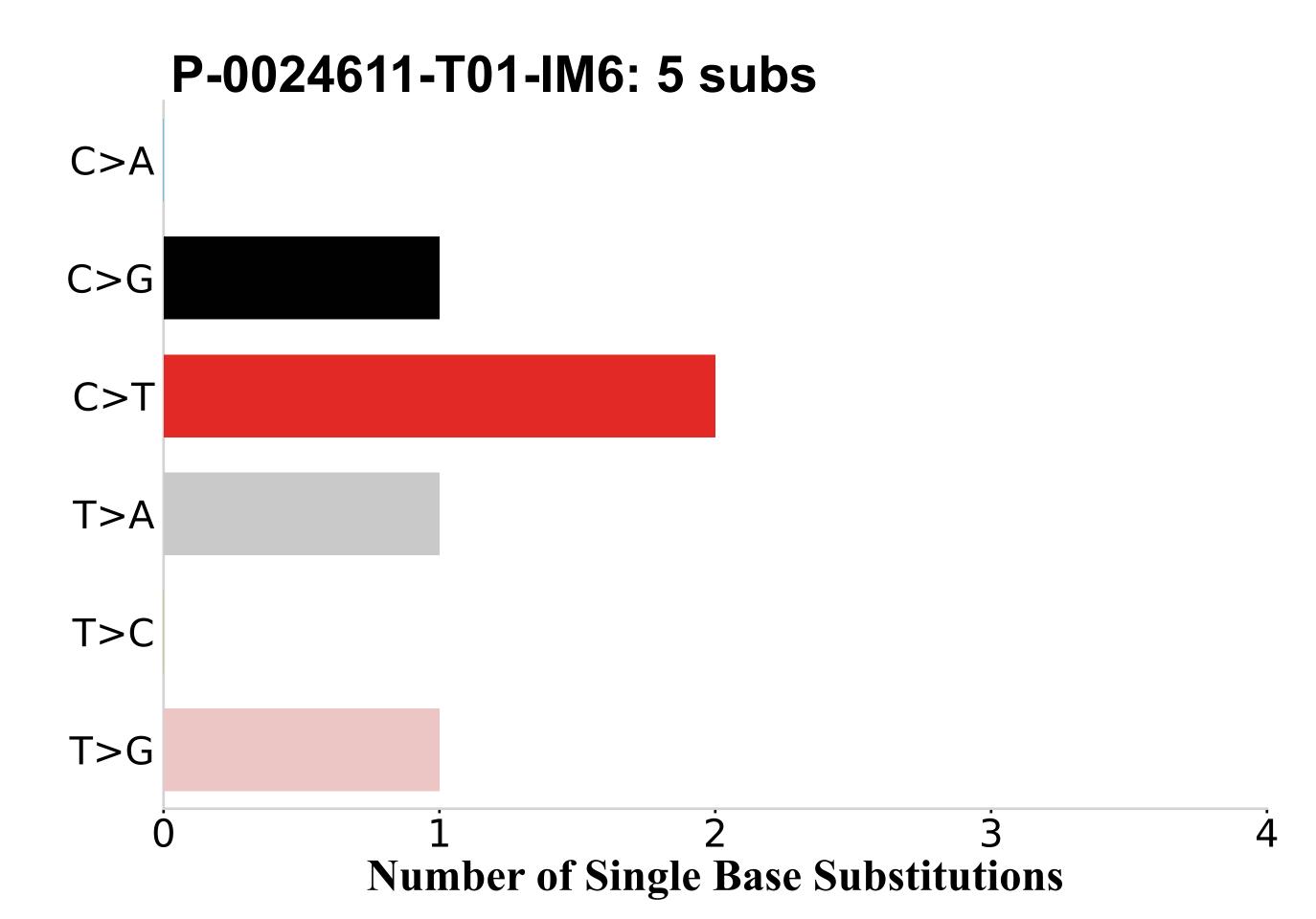


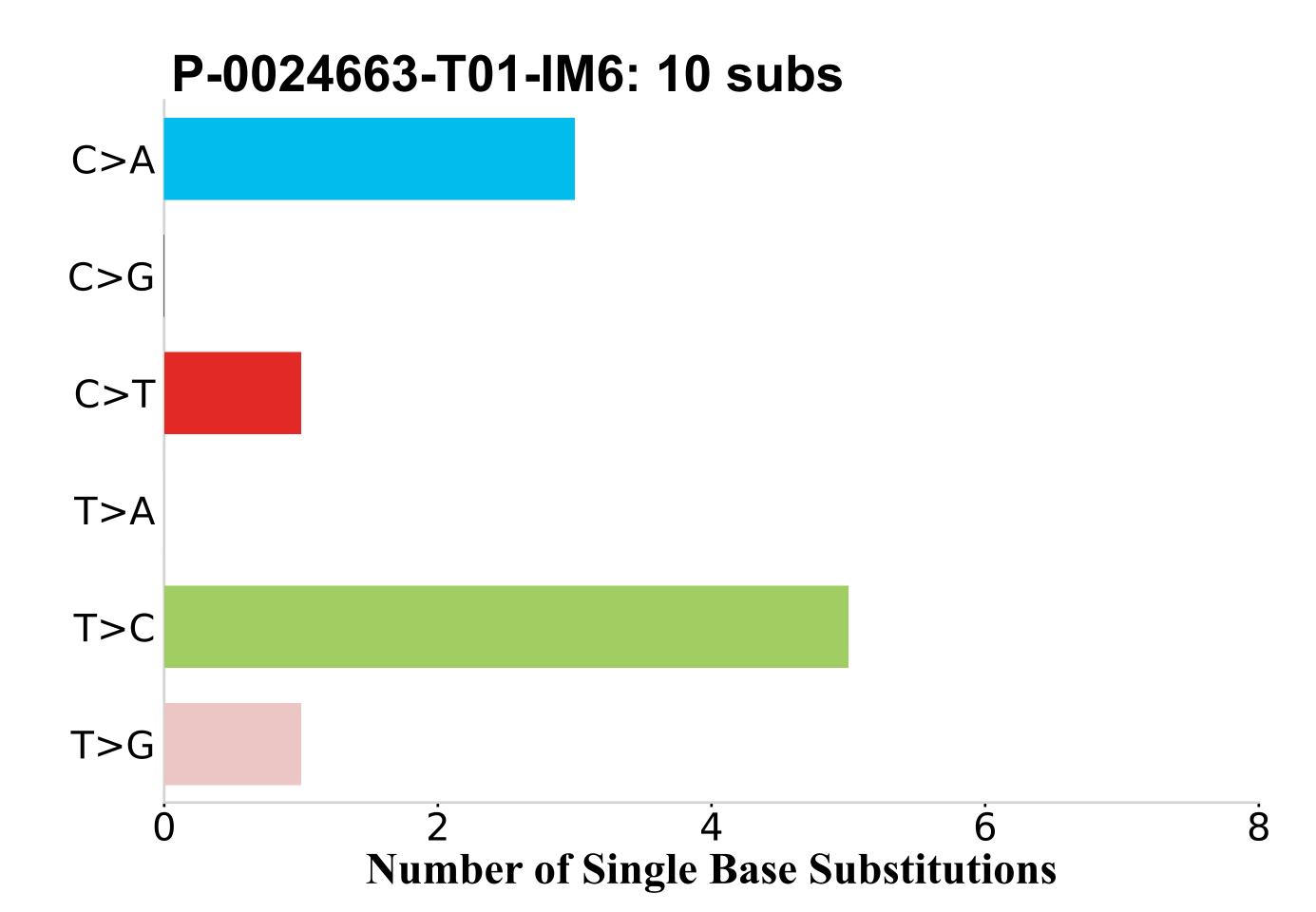


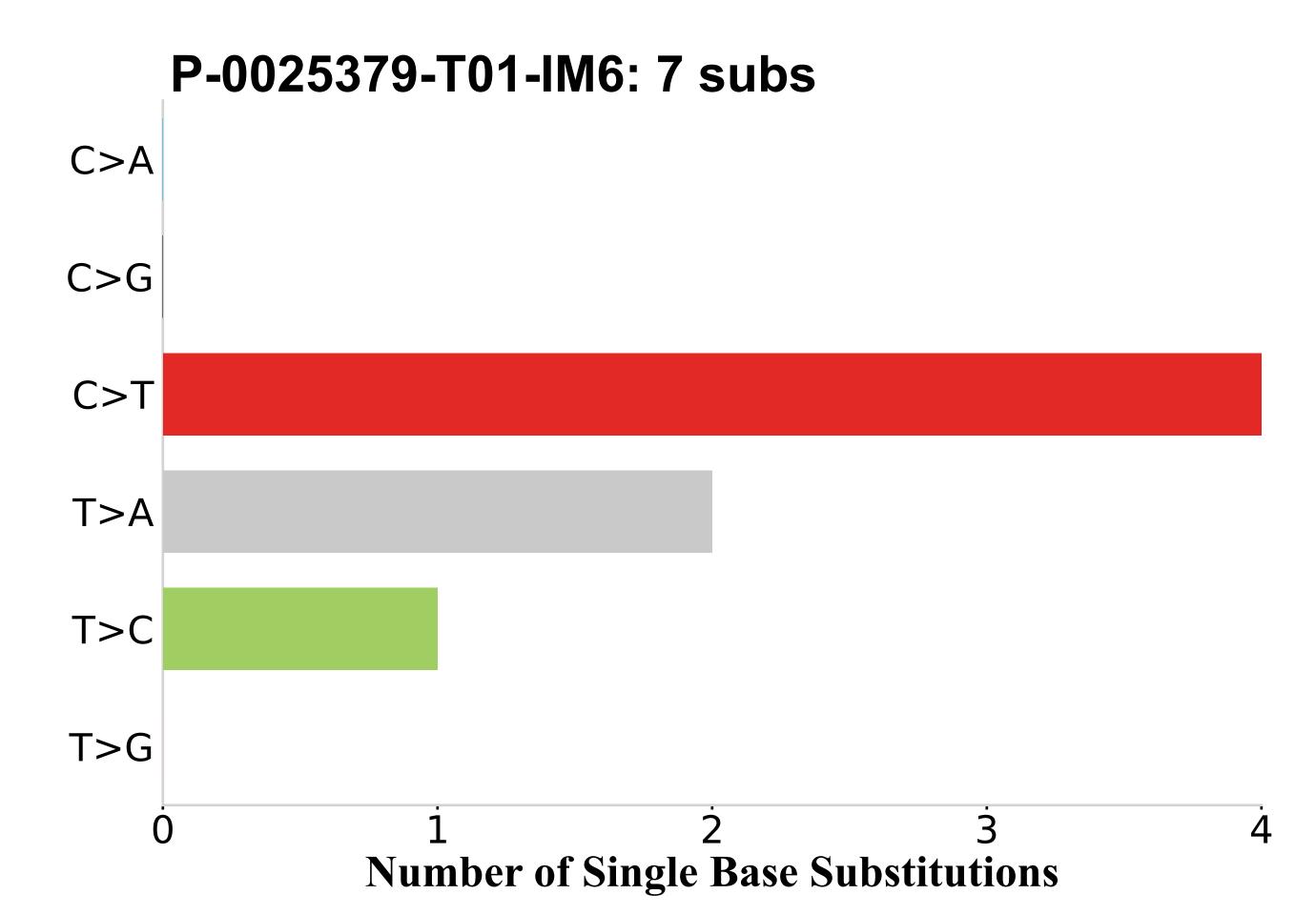


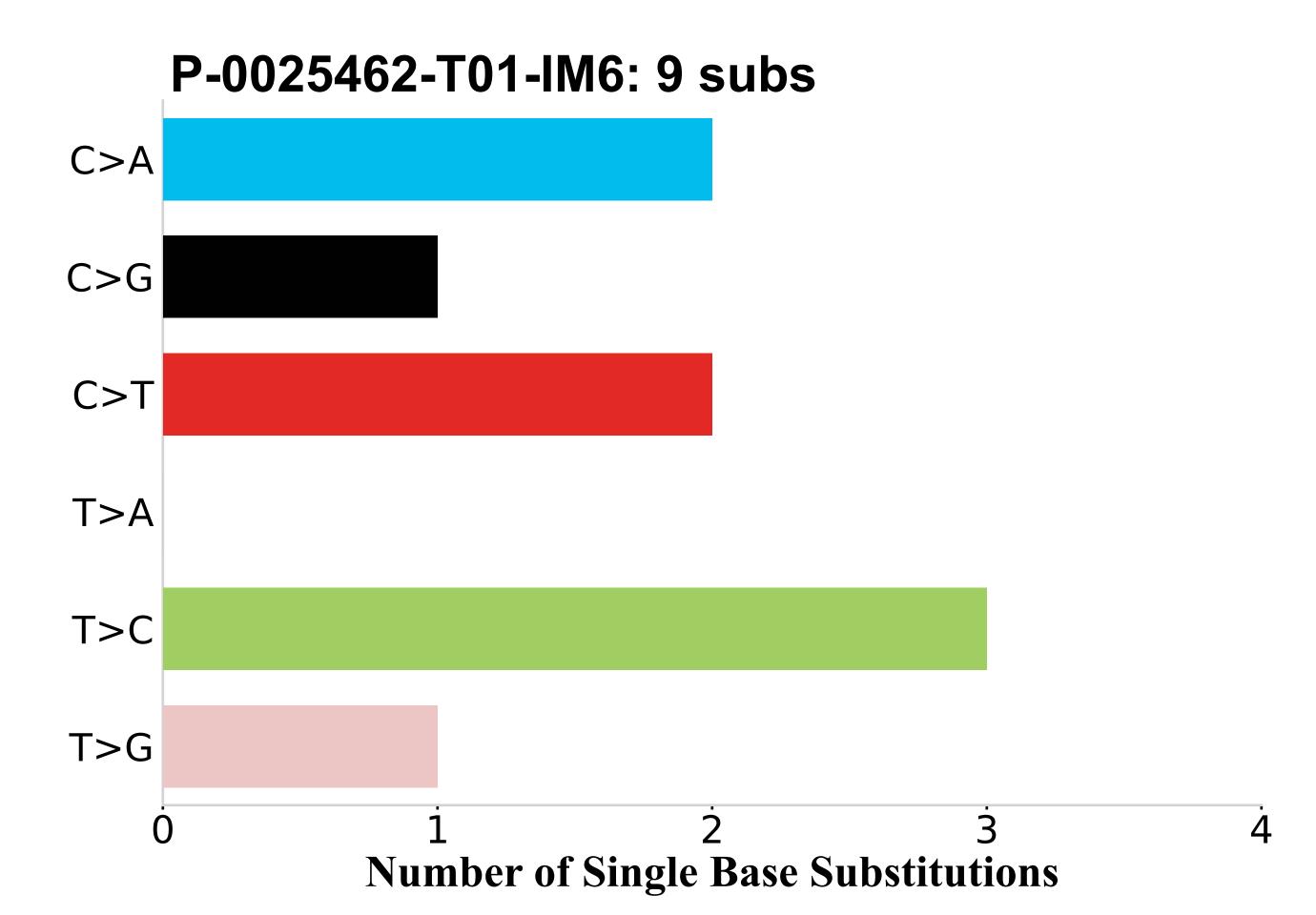


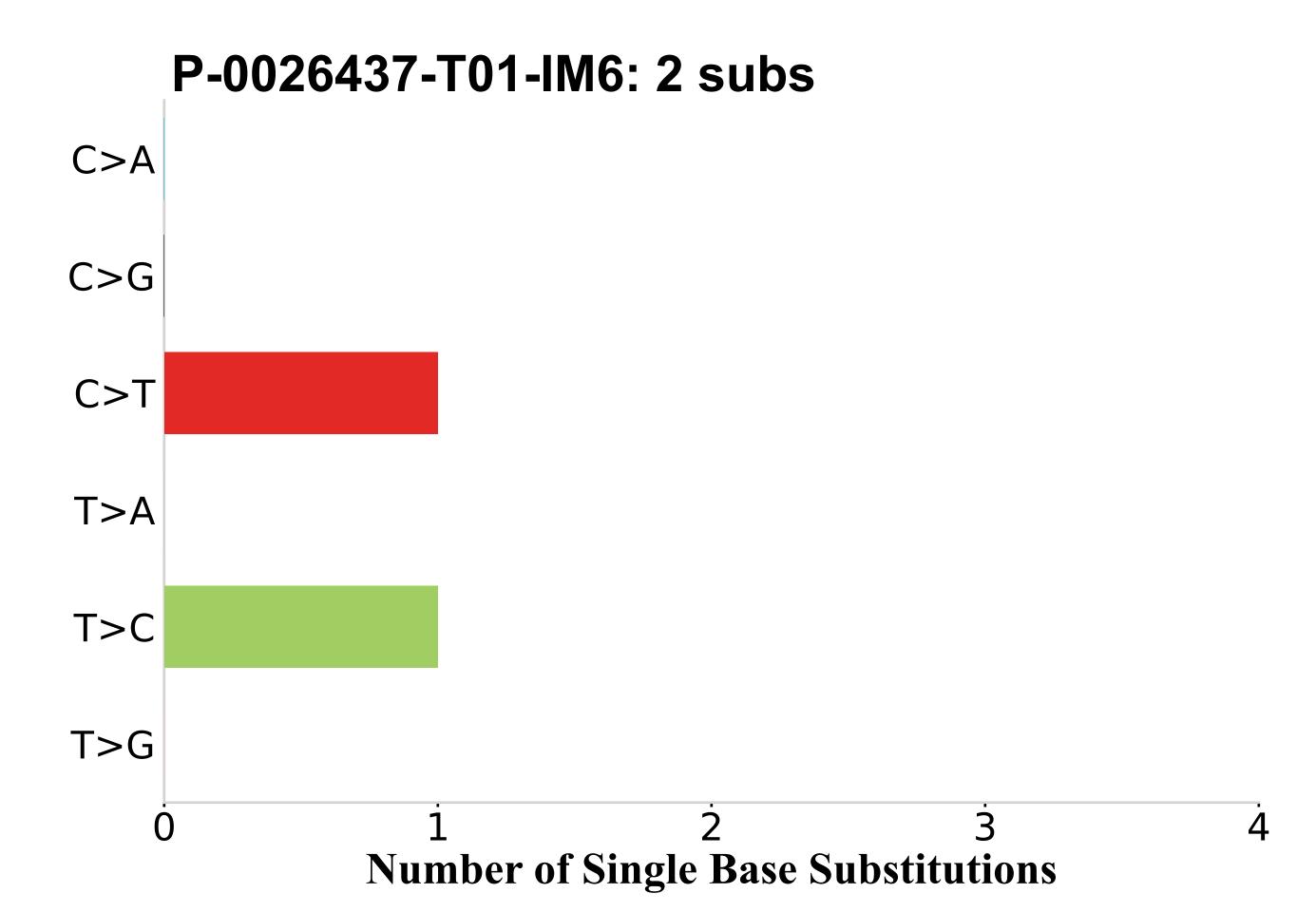




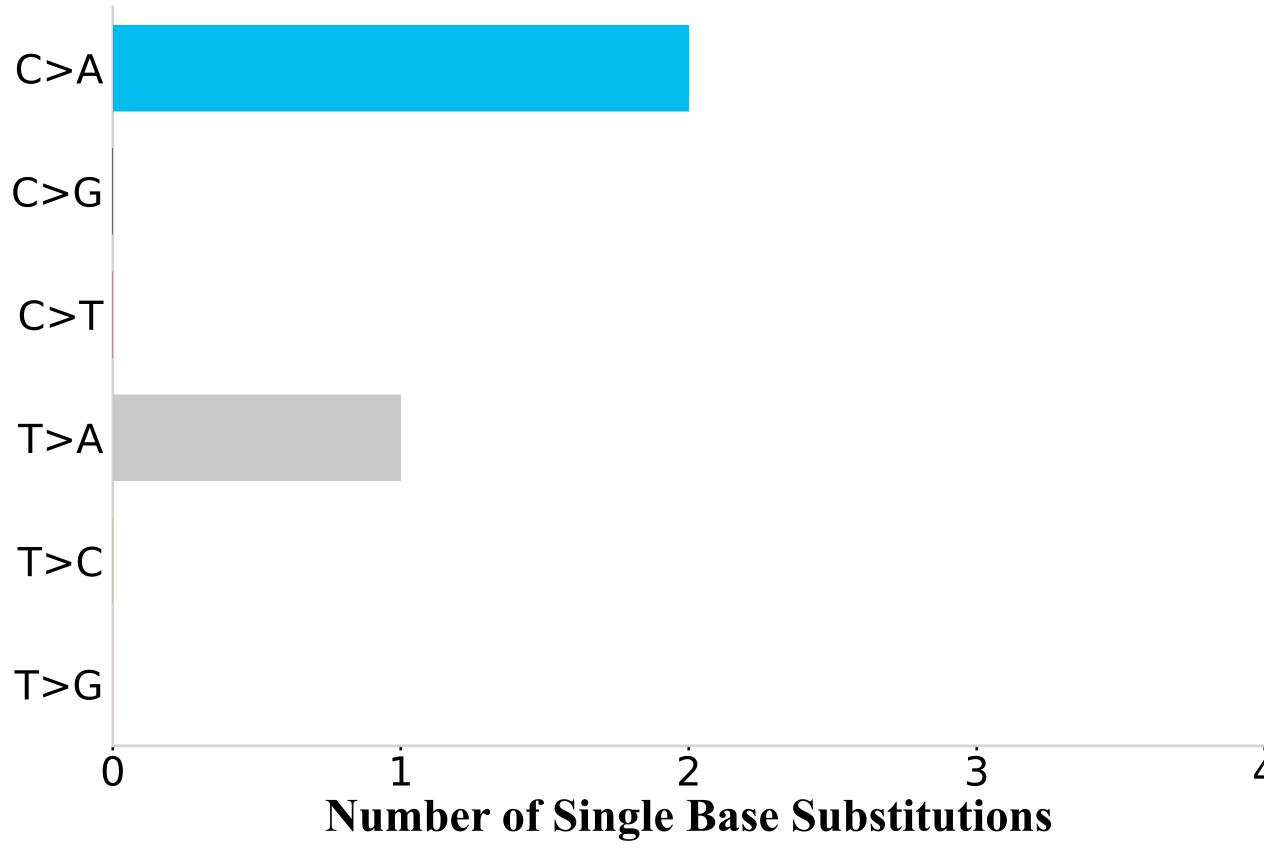


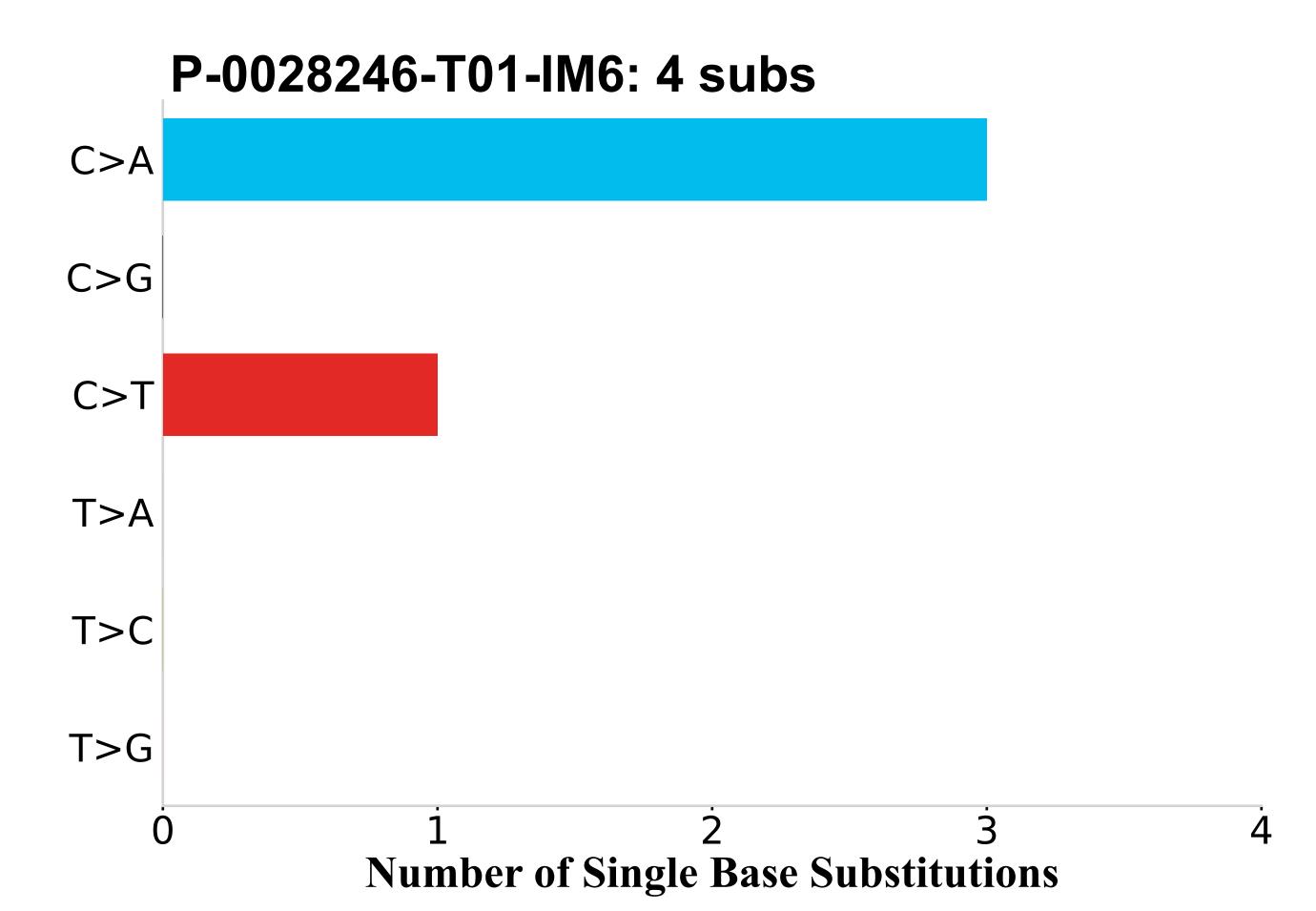


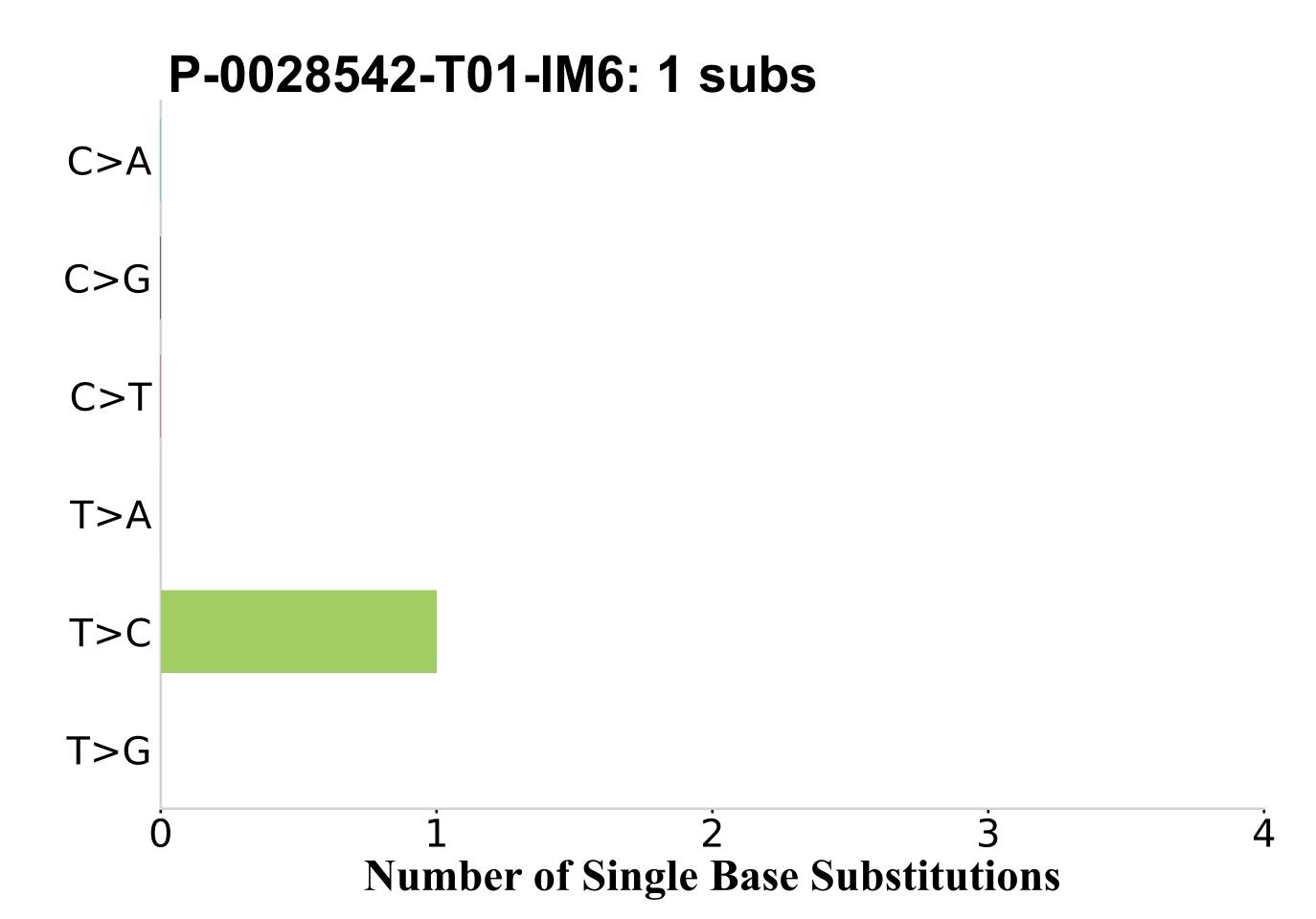


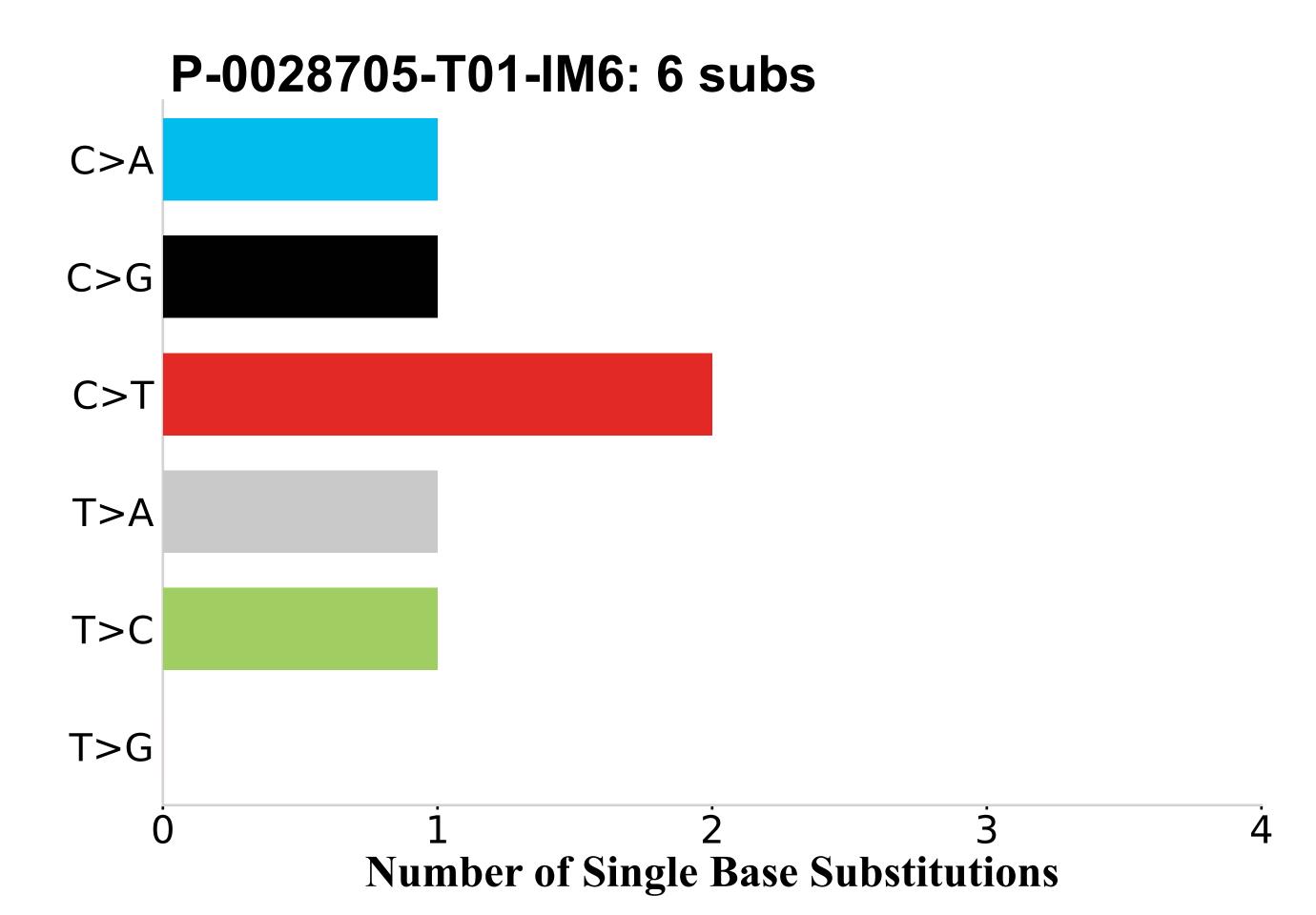


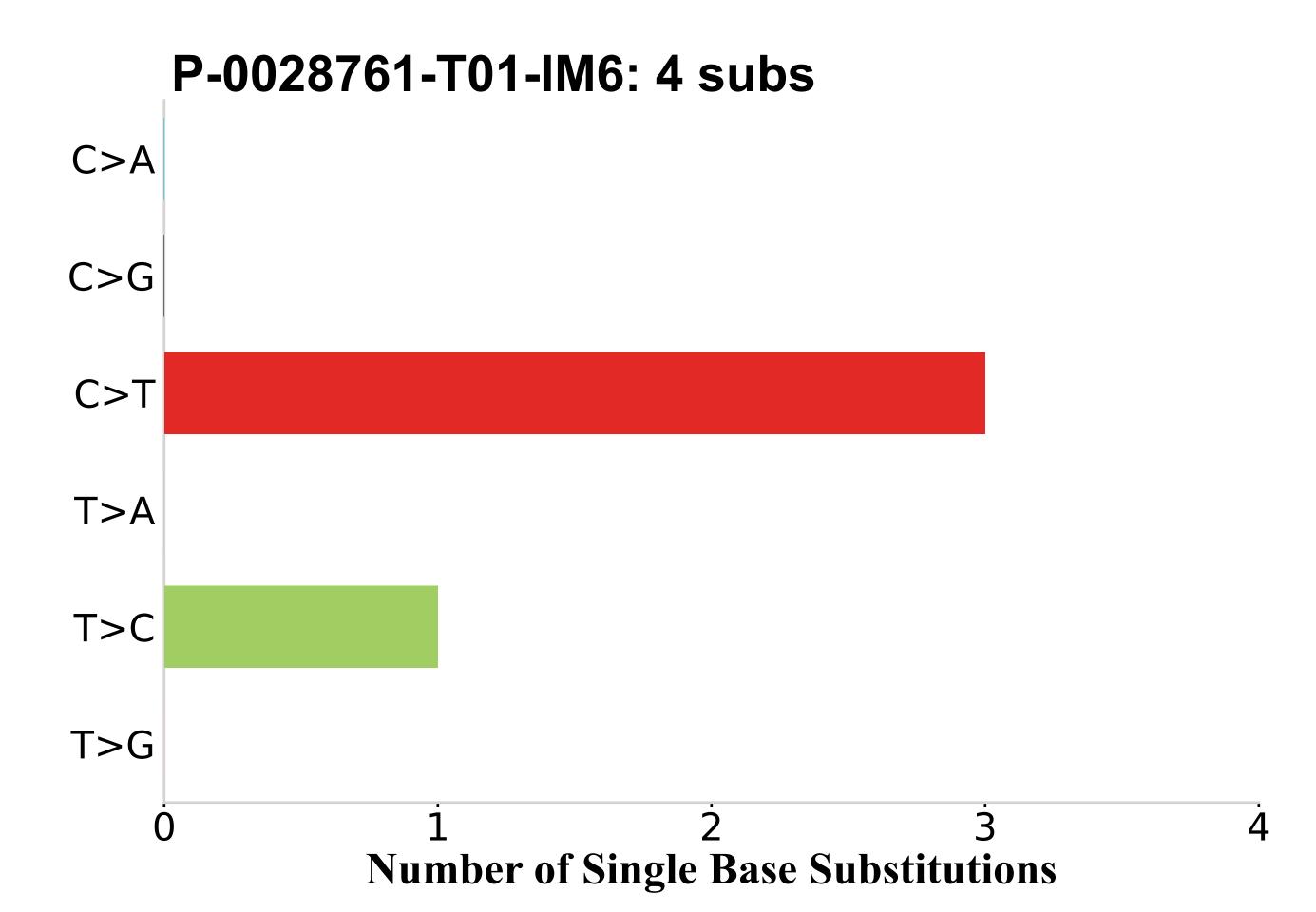




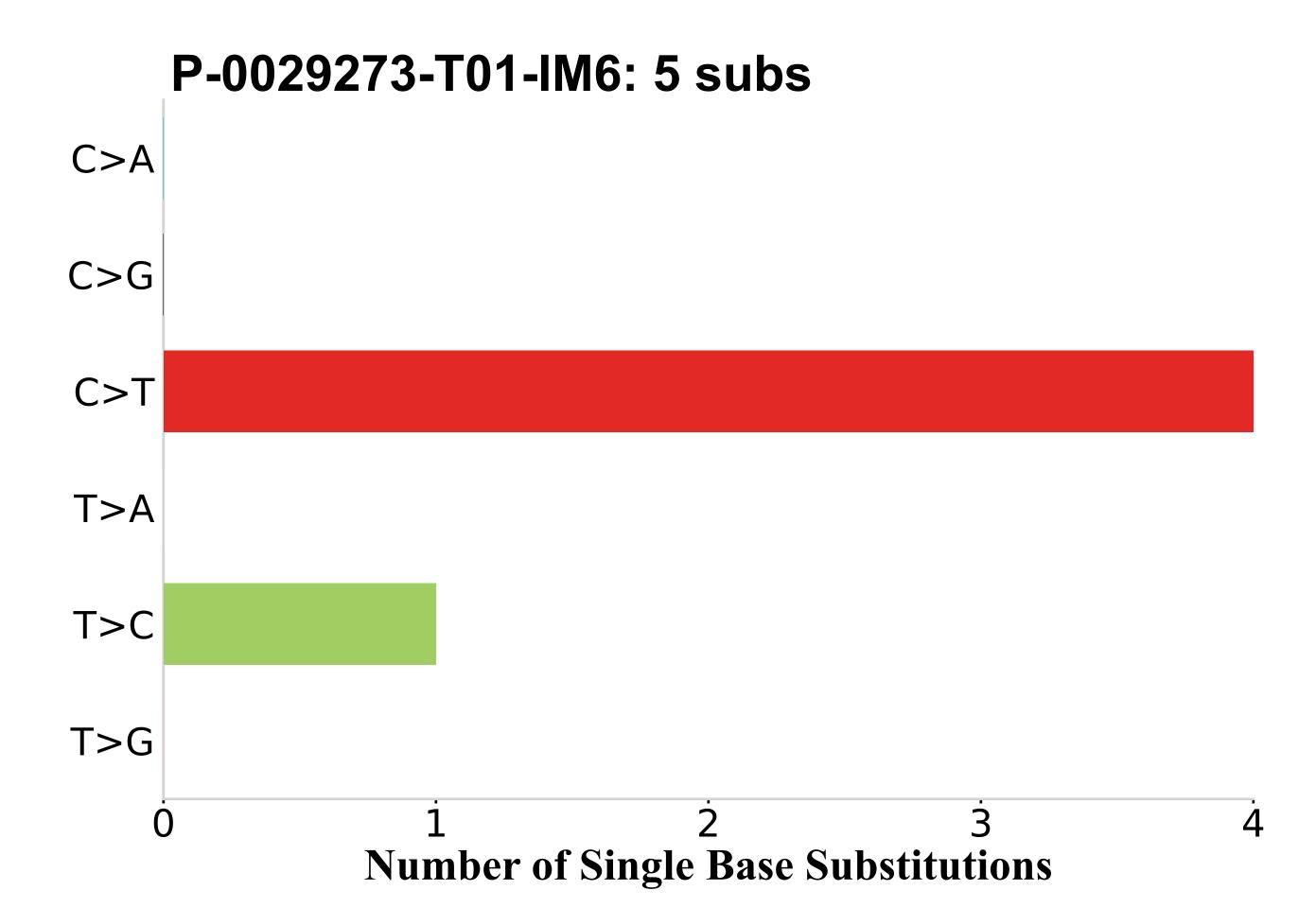


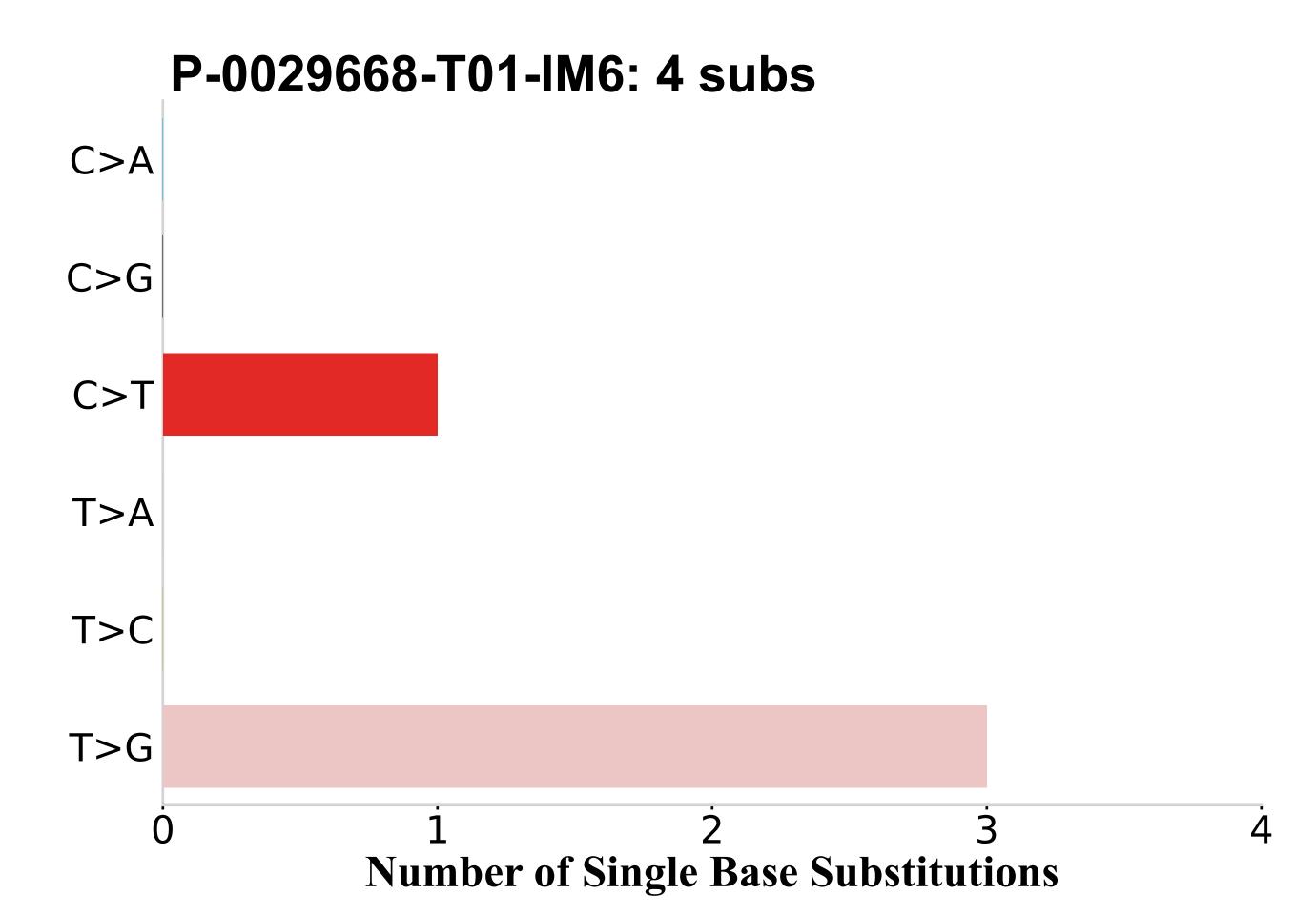


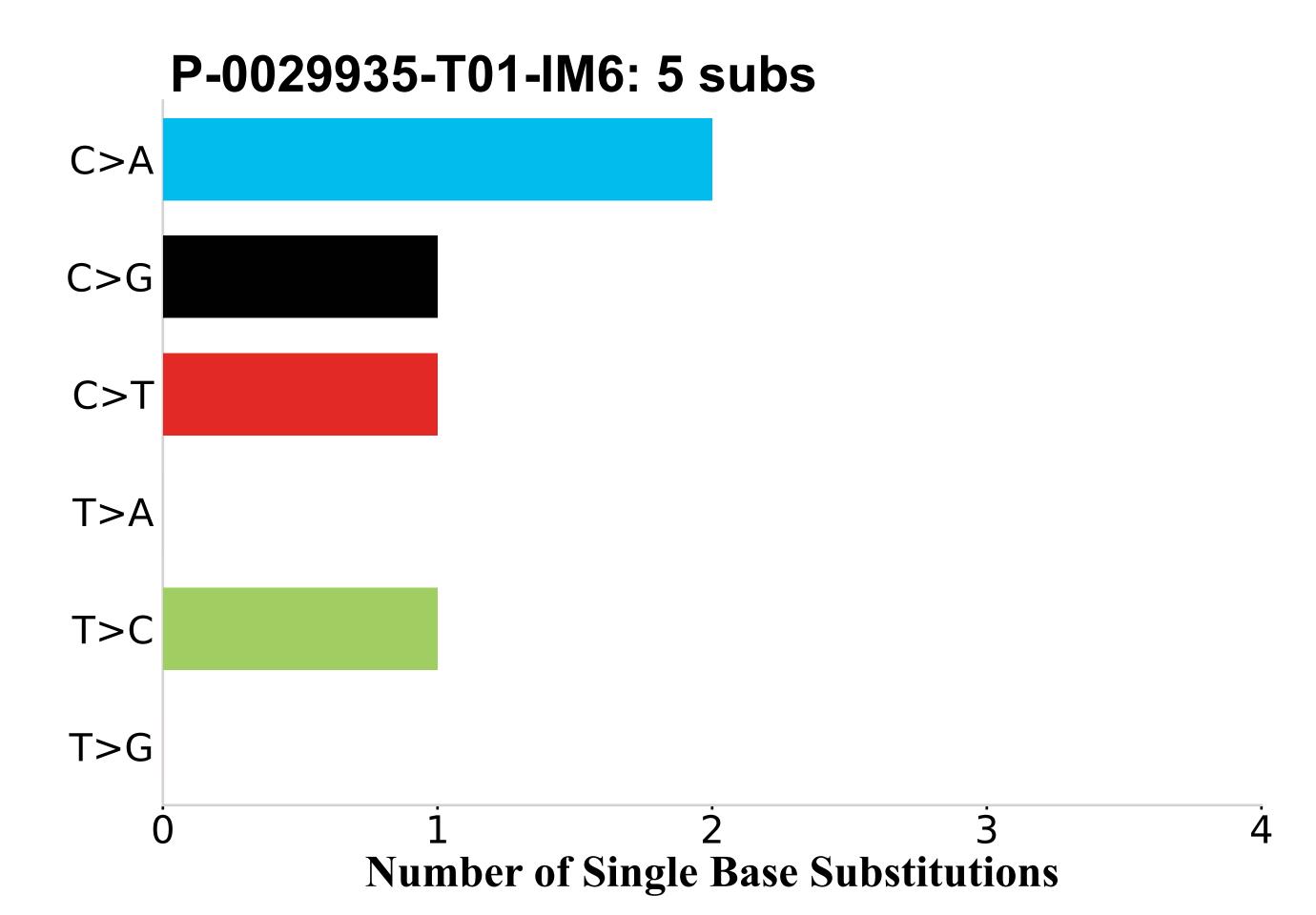


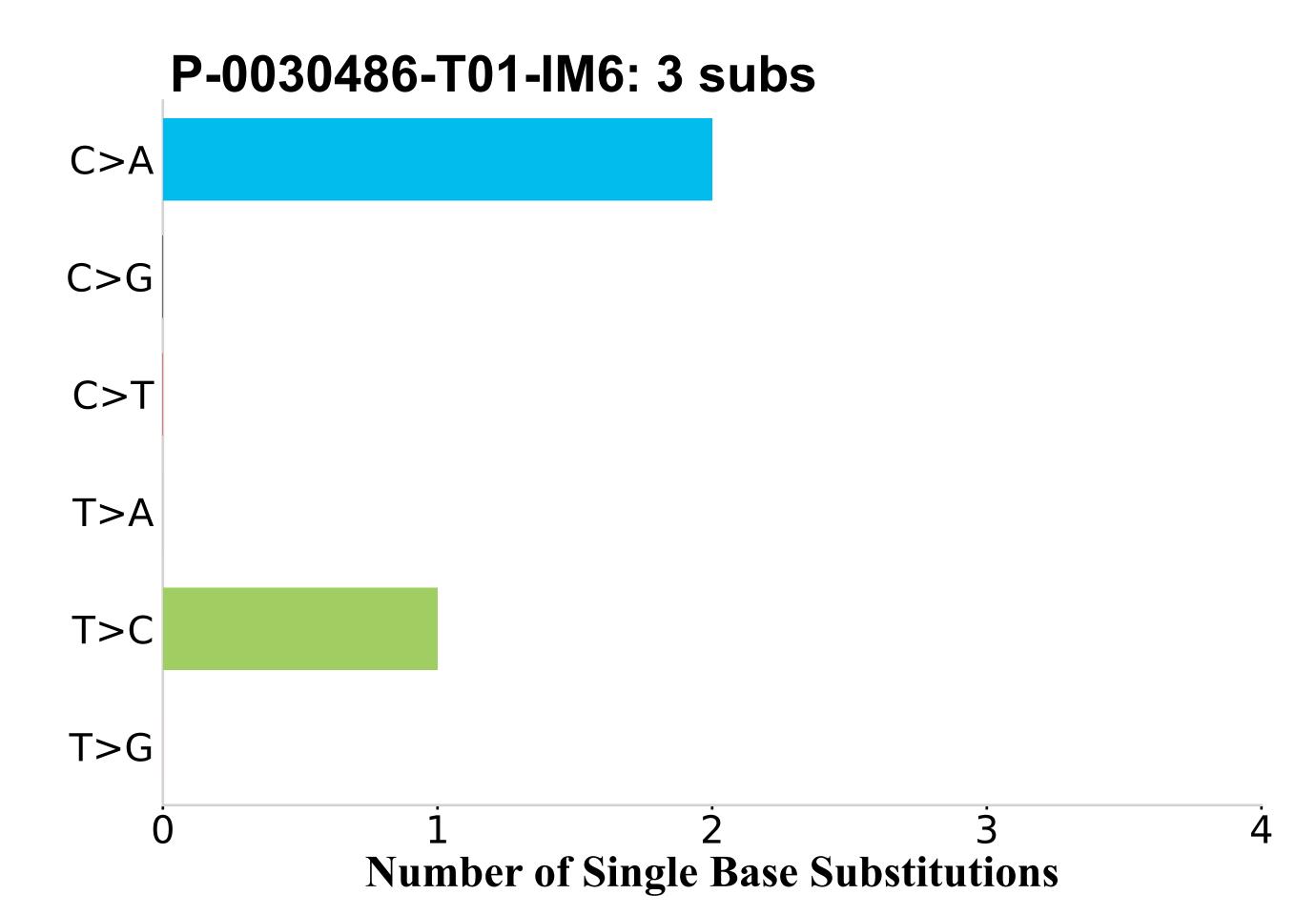


P-0029068-T01-IM6: 8 subs C>A C>G C>T T>A T>C T>G **Number of Single Base Substitutions**

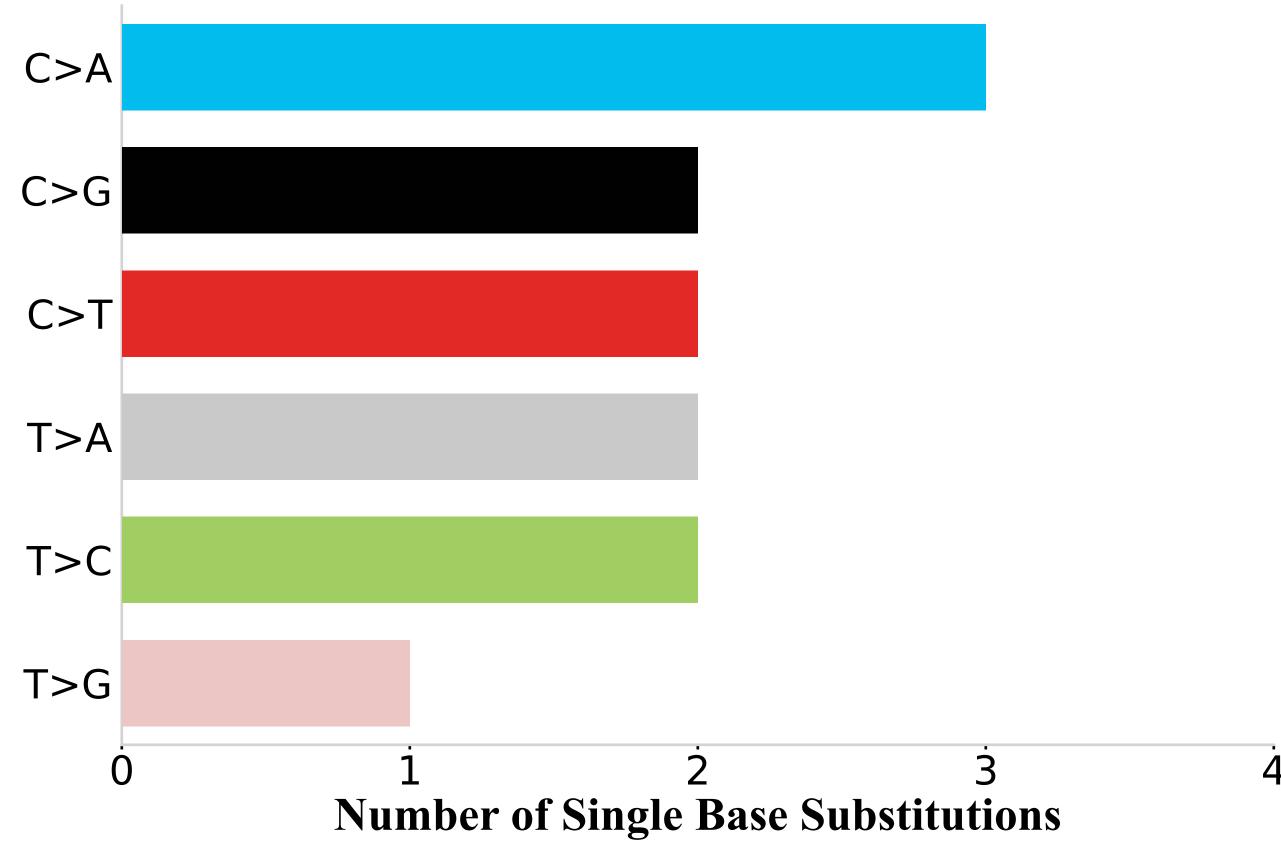




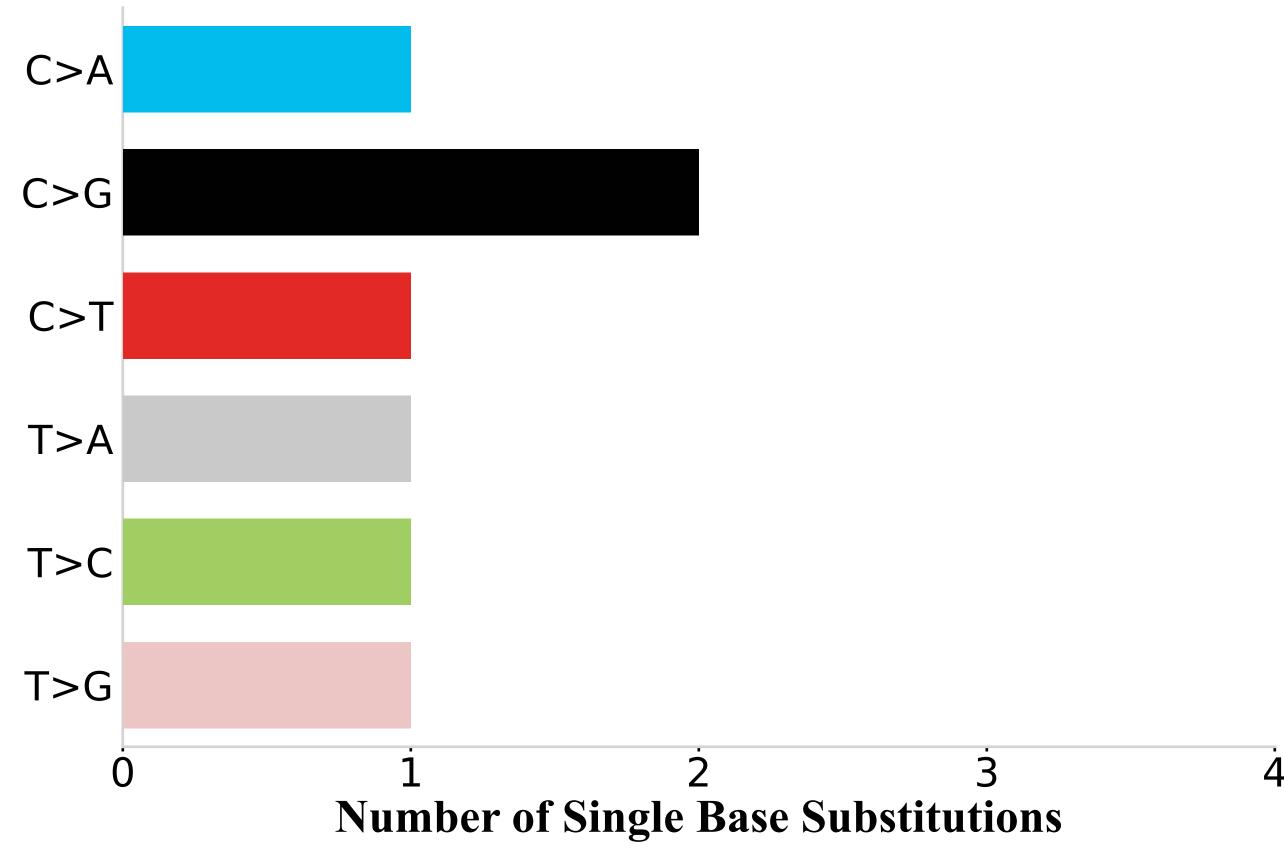


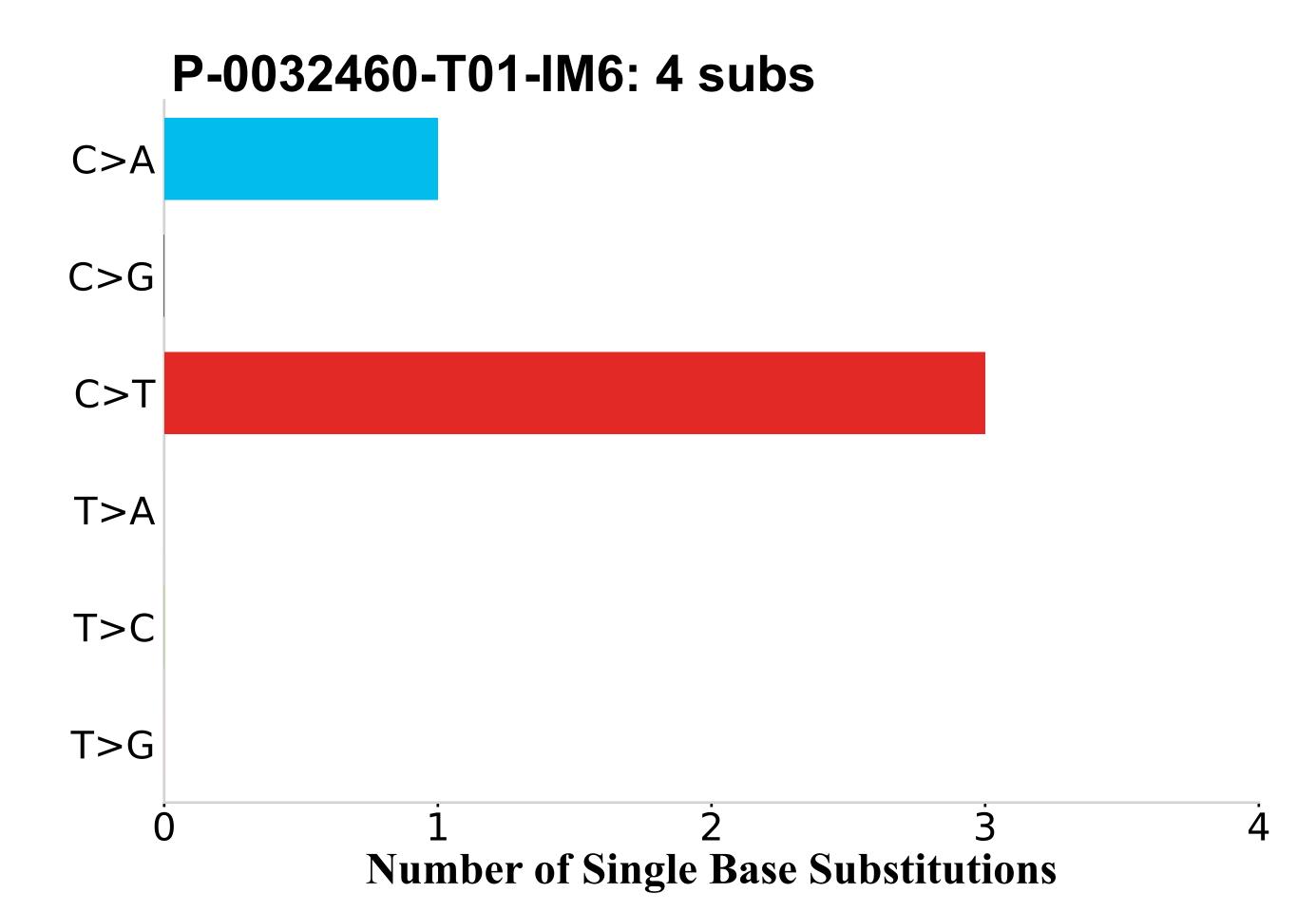


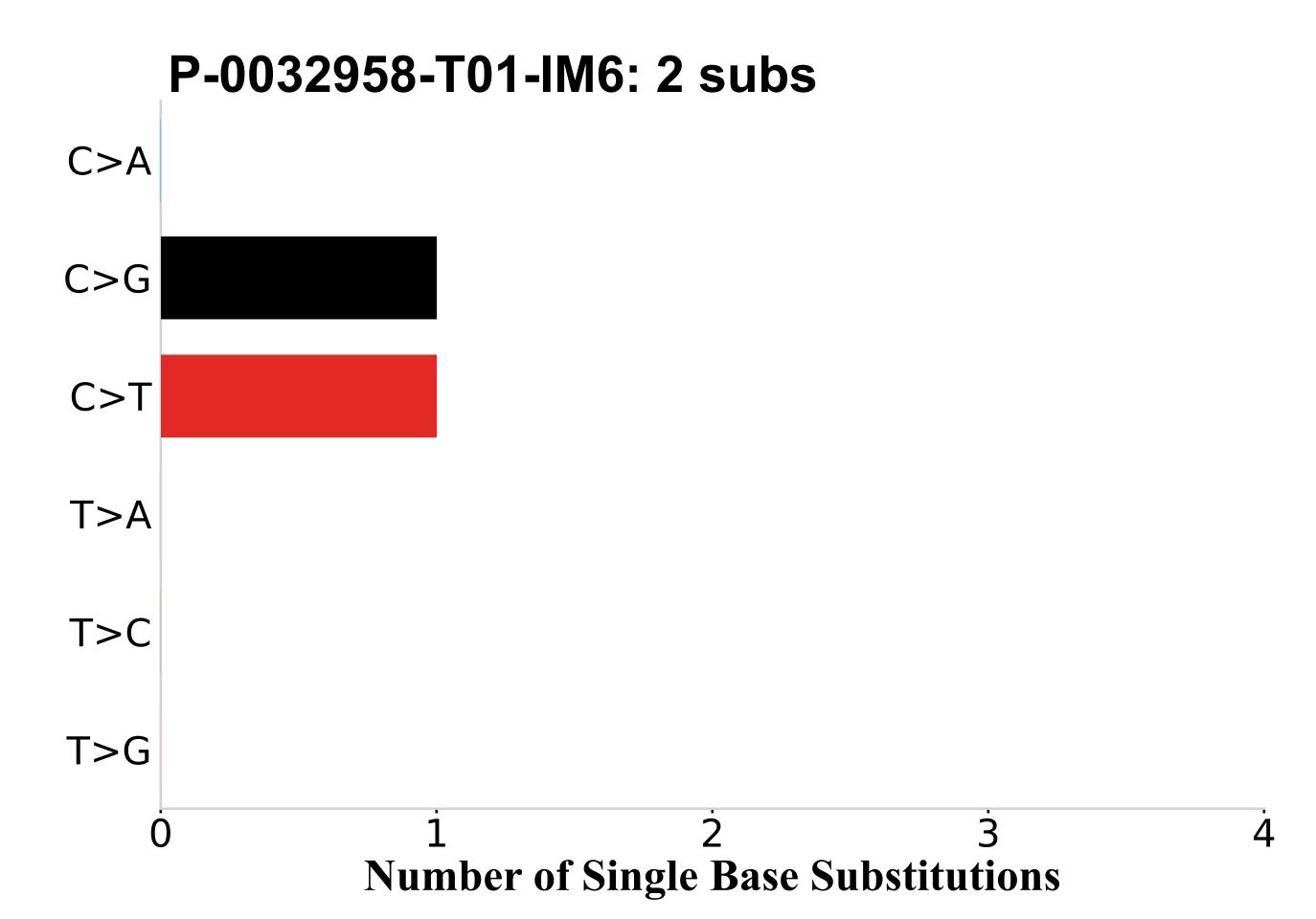


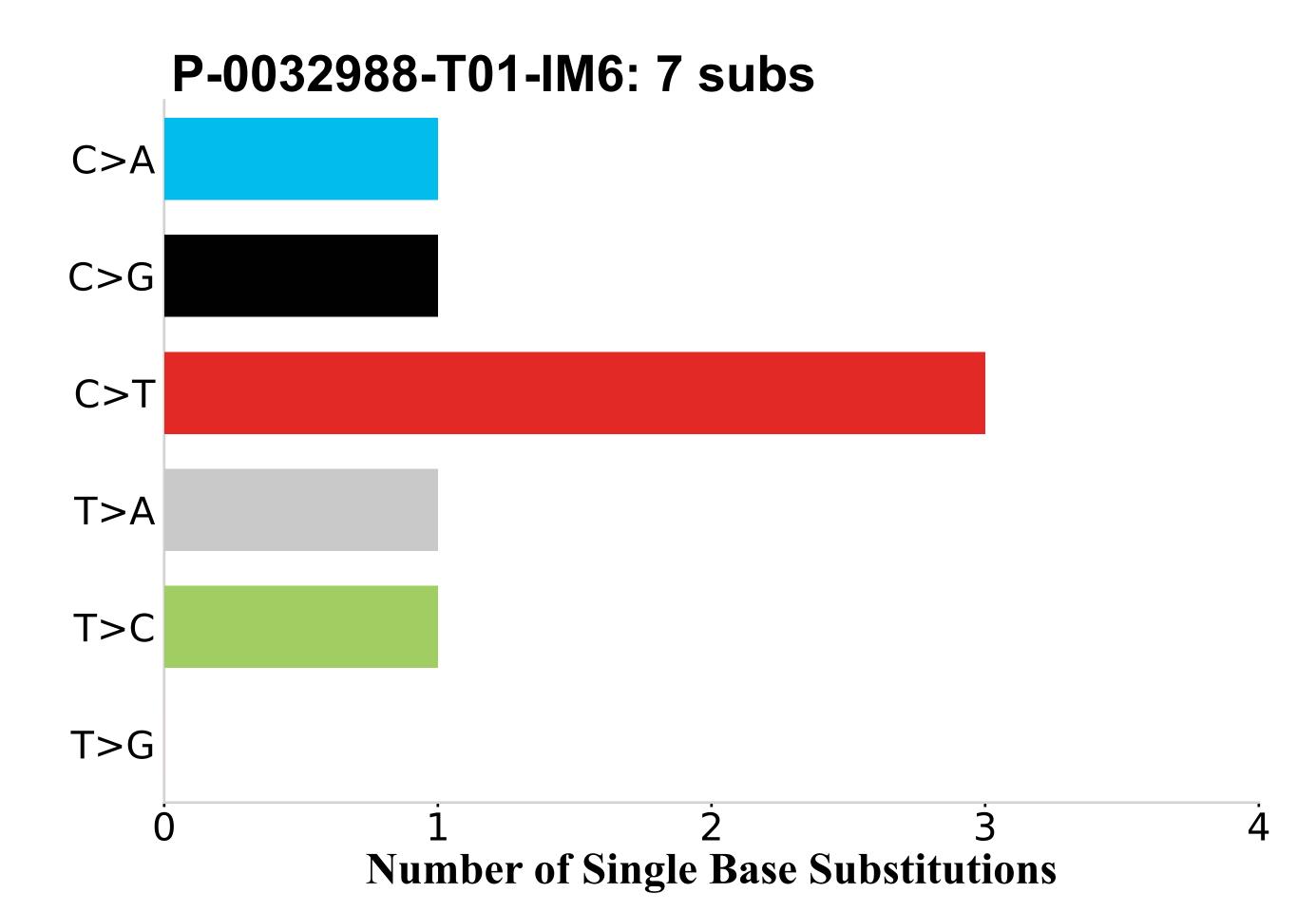


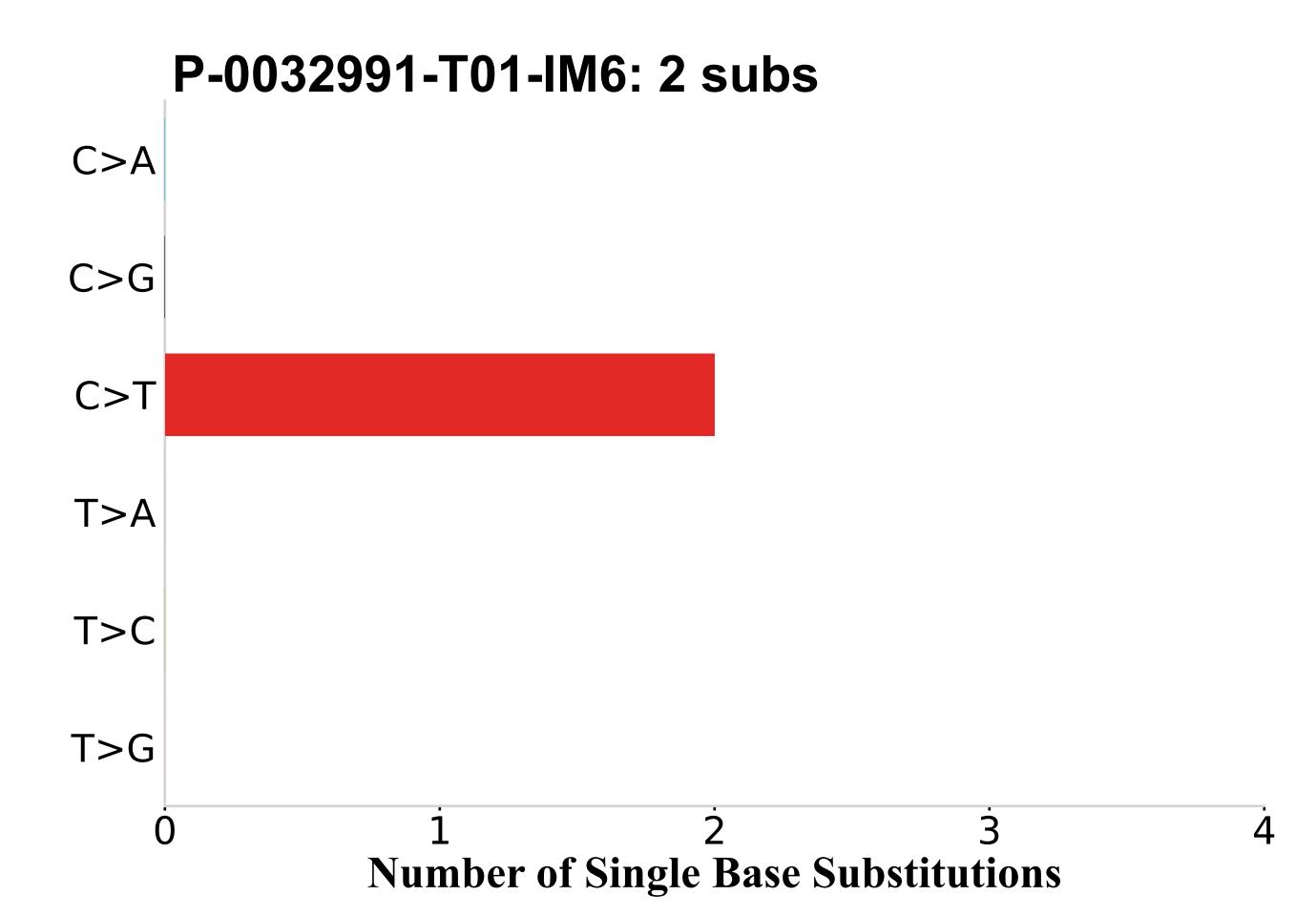
P-0031657-T01-IM6: 7 subs



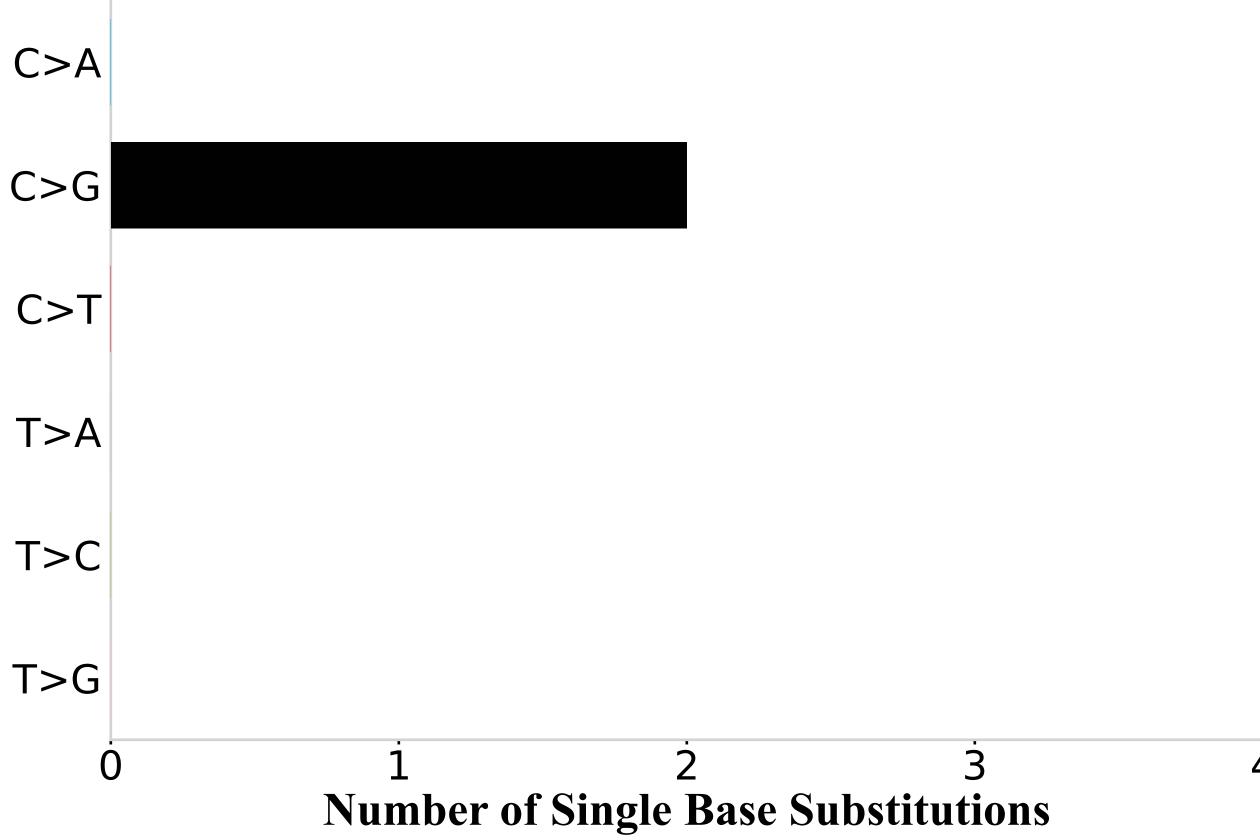




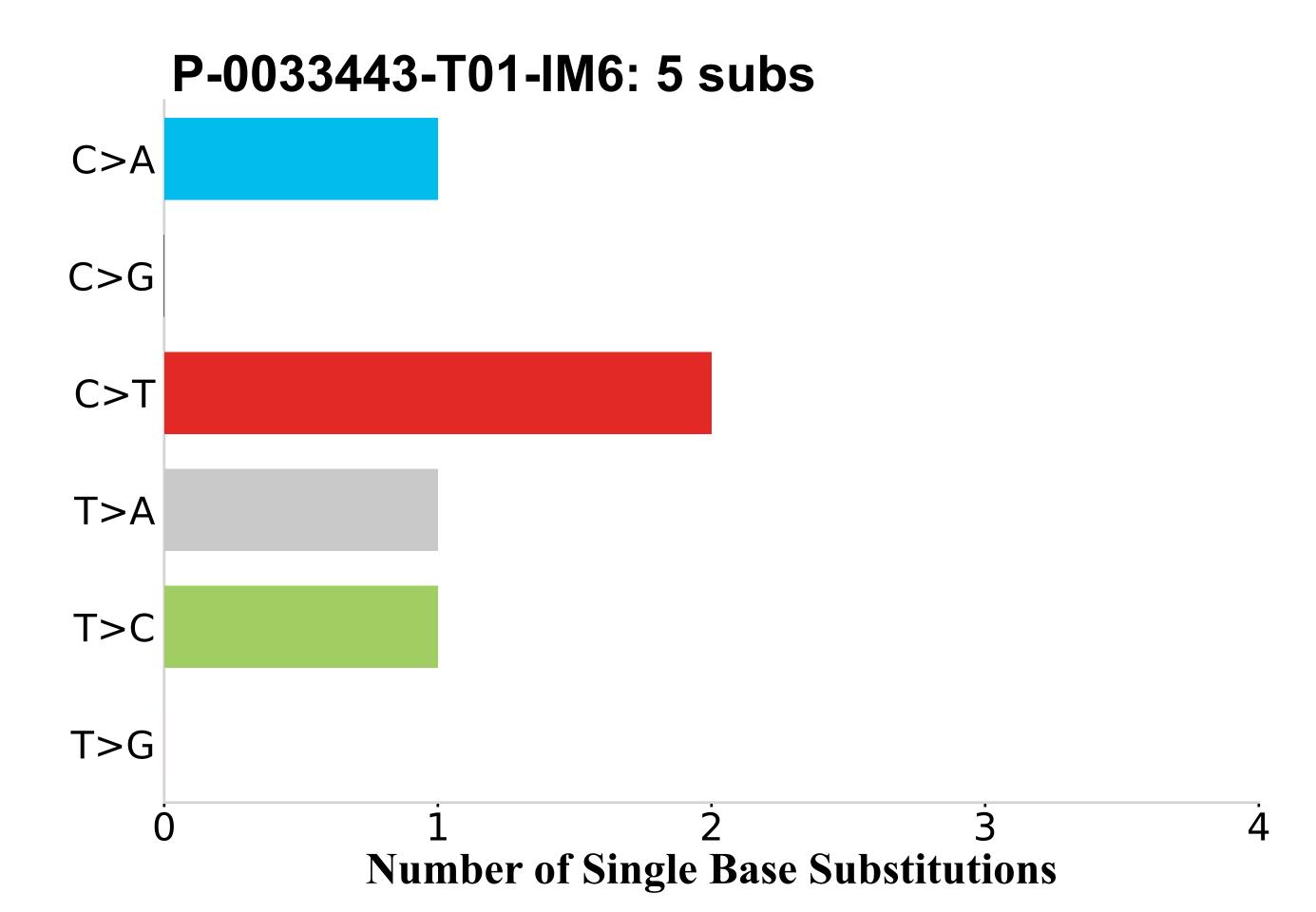




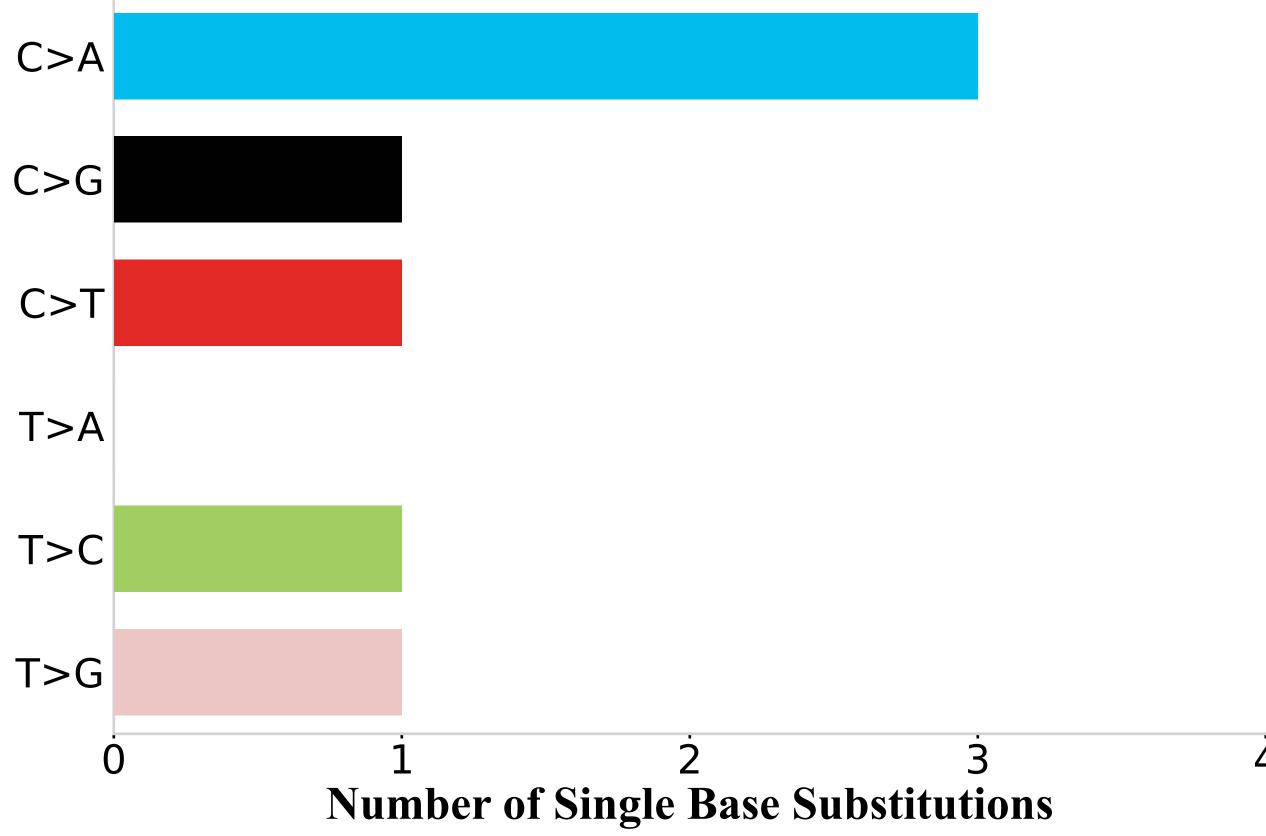


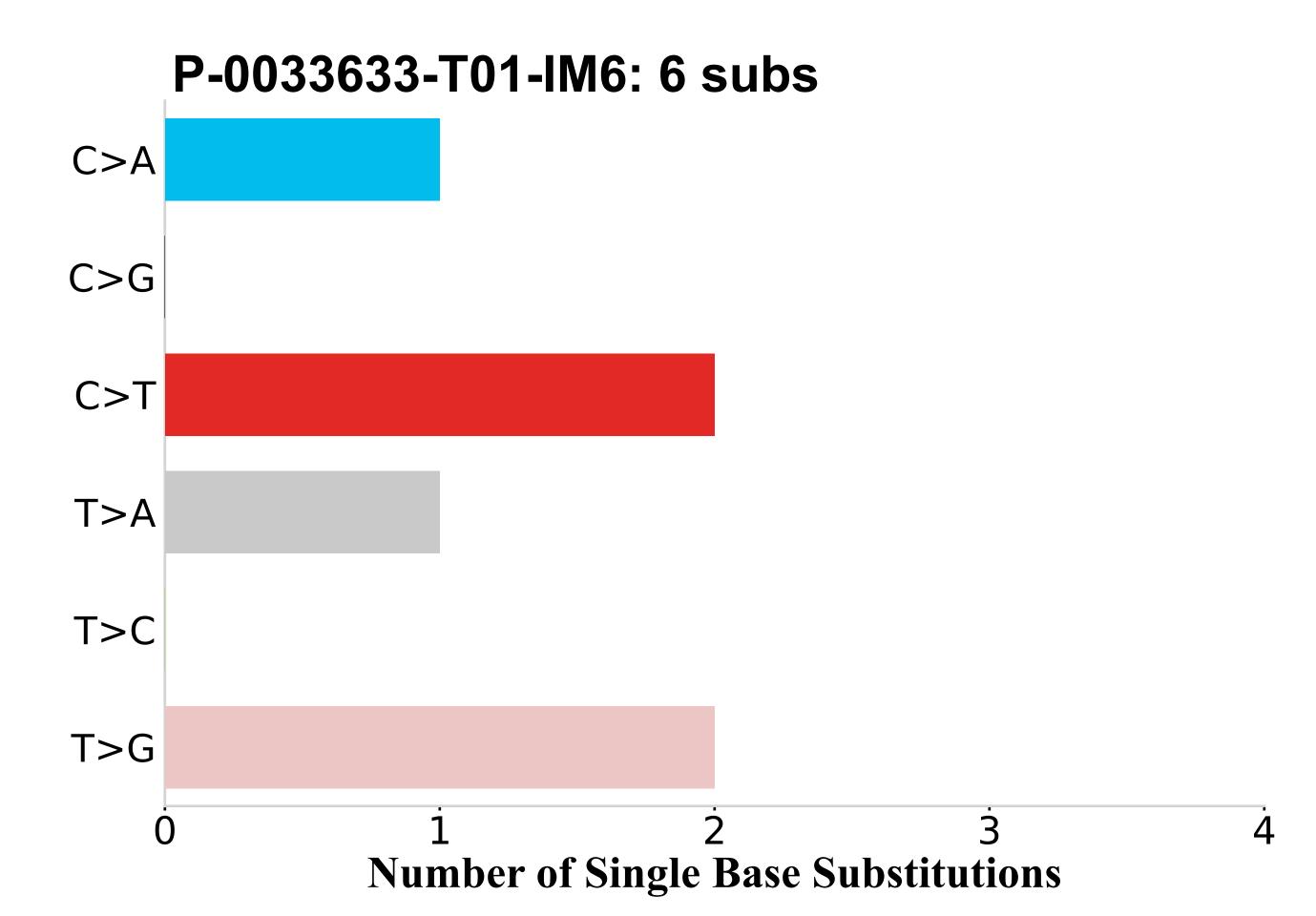


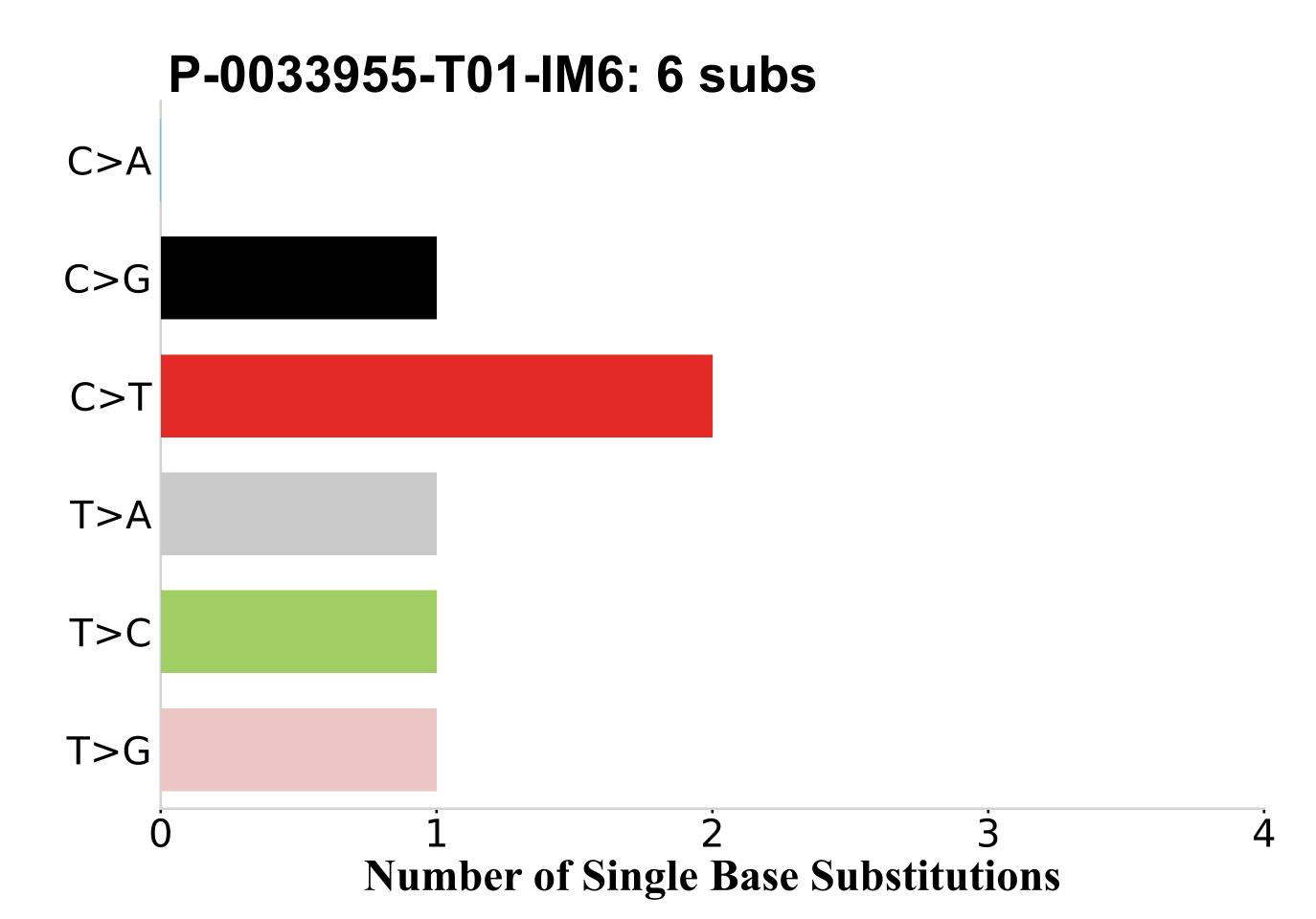


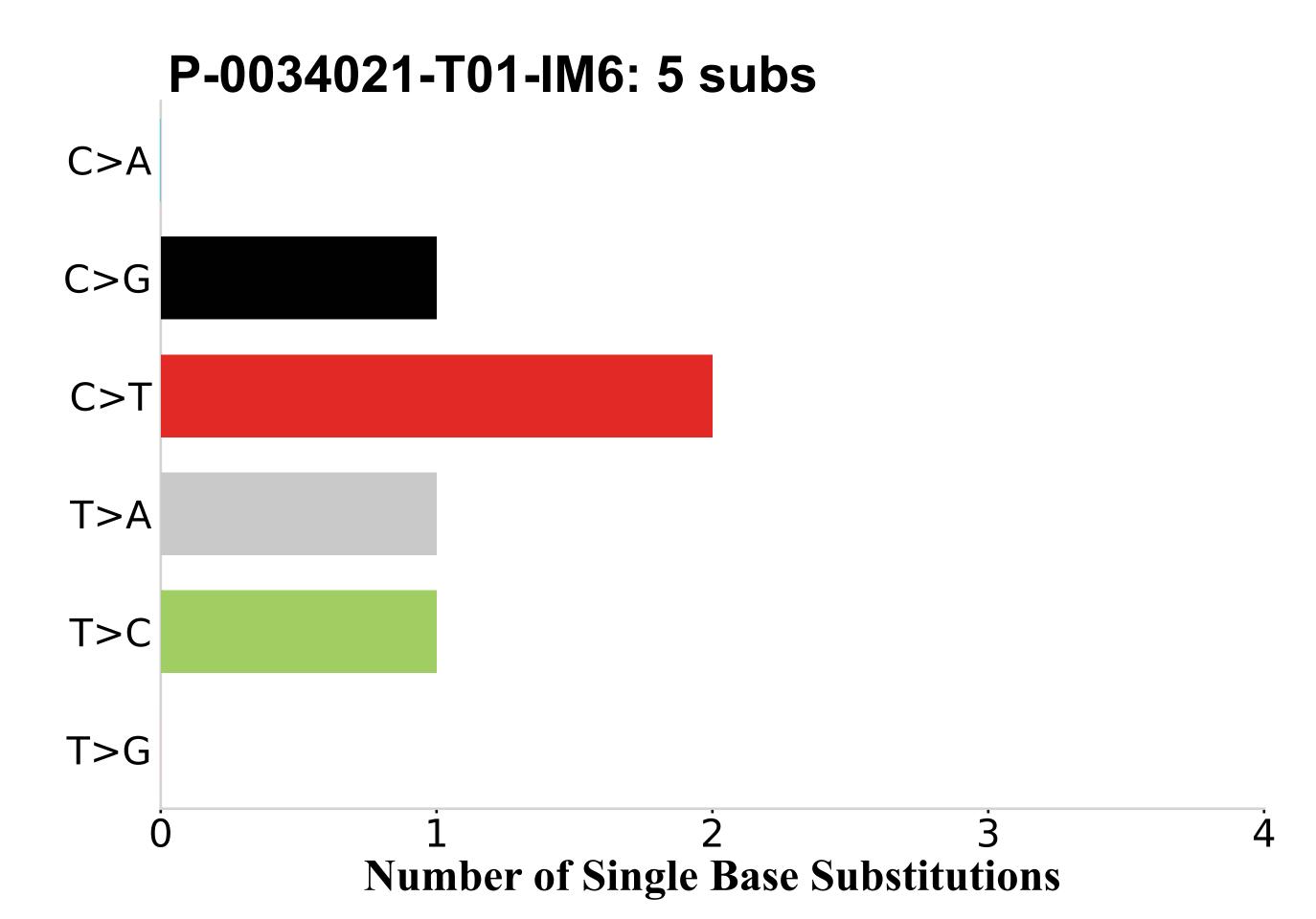




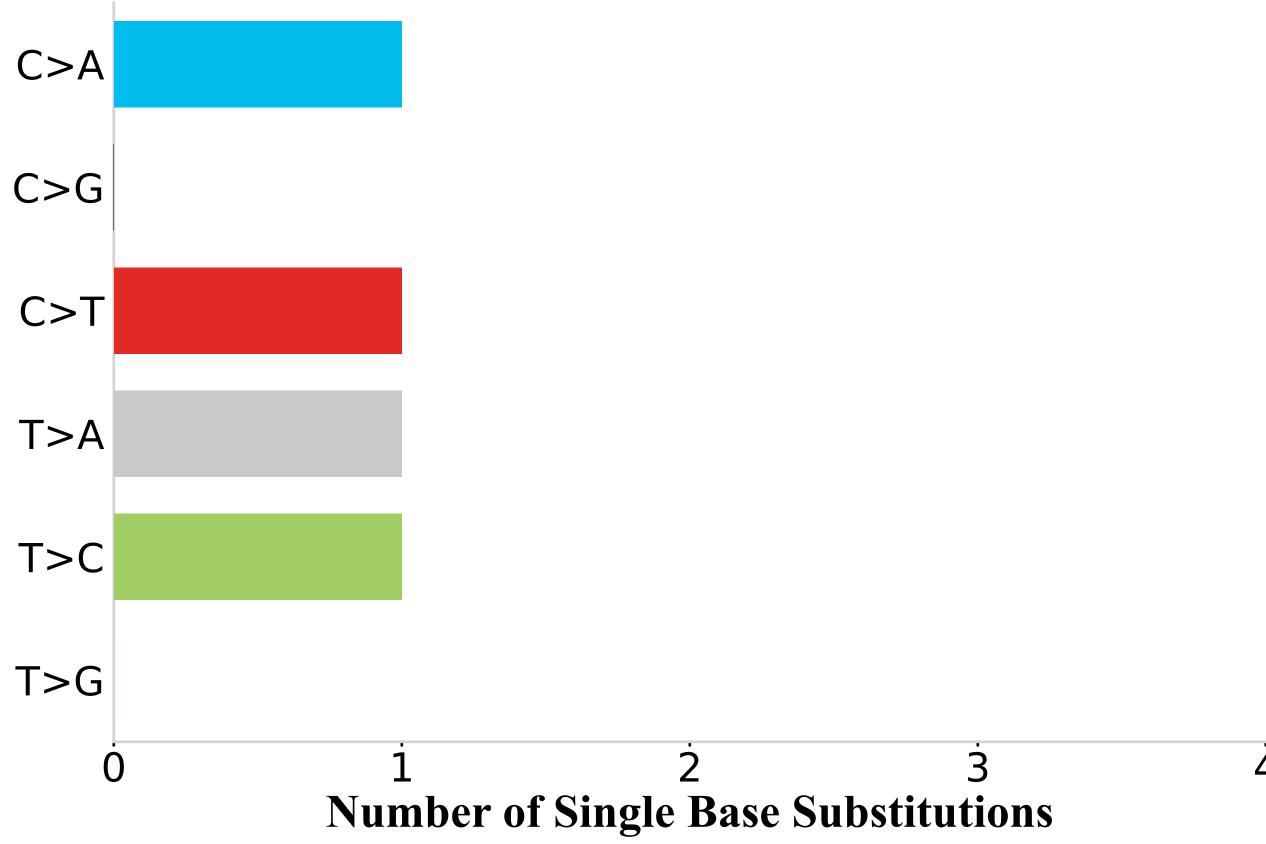


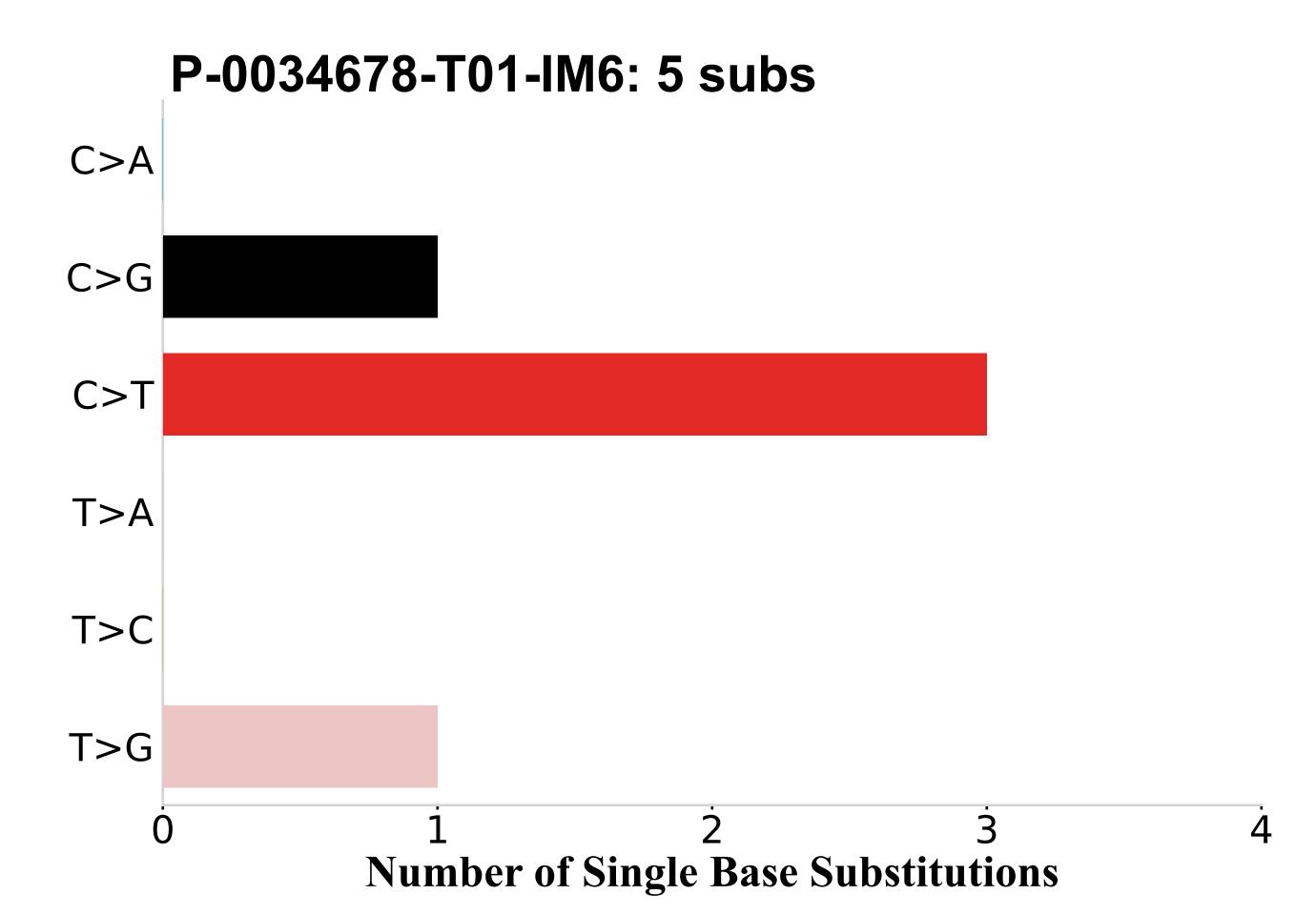




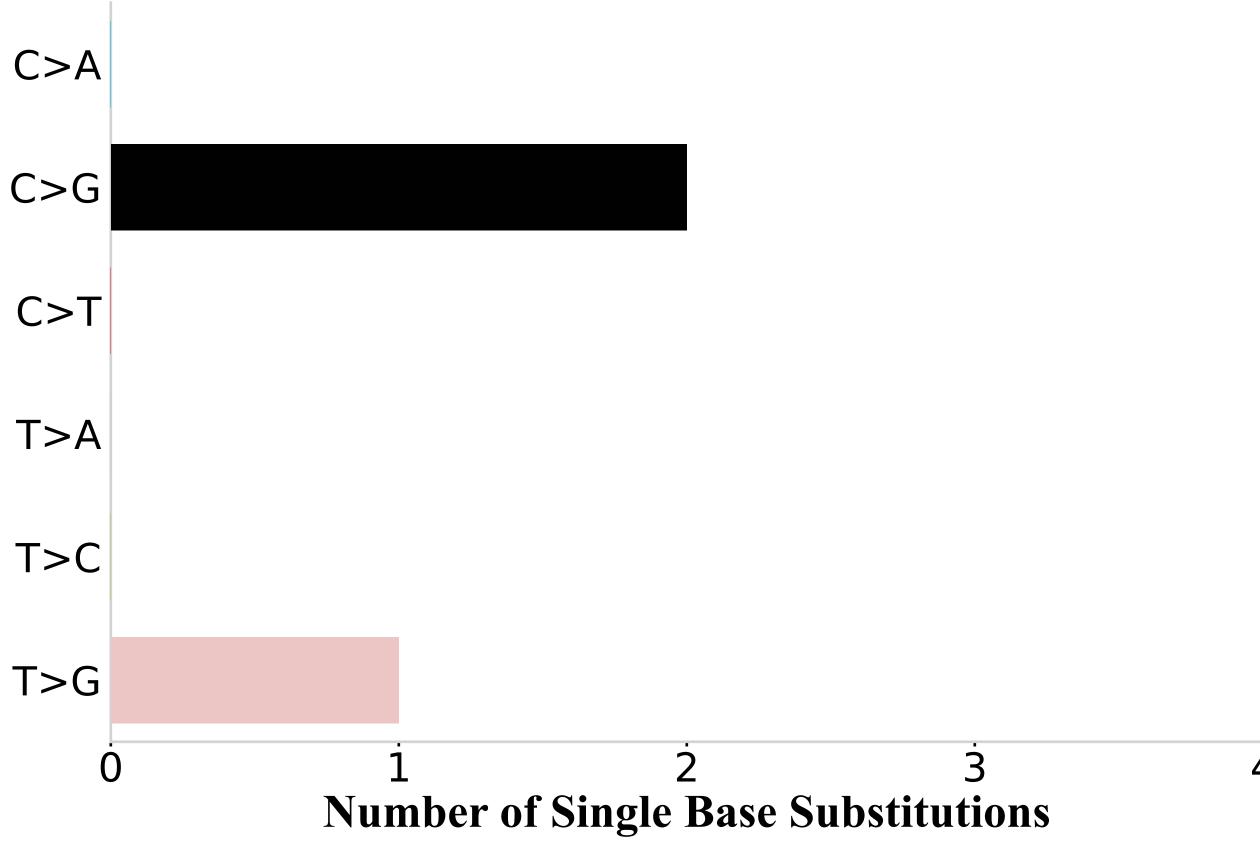


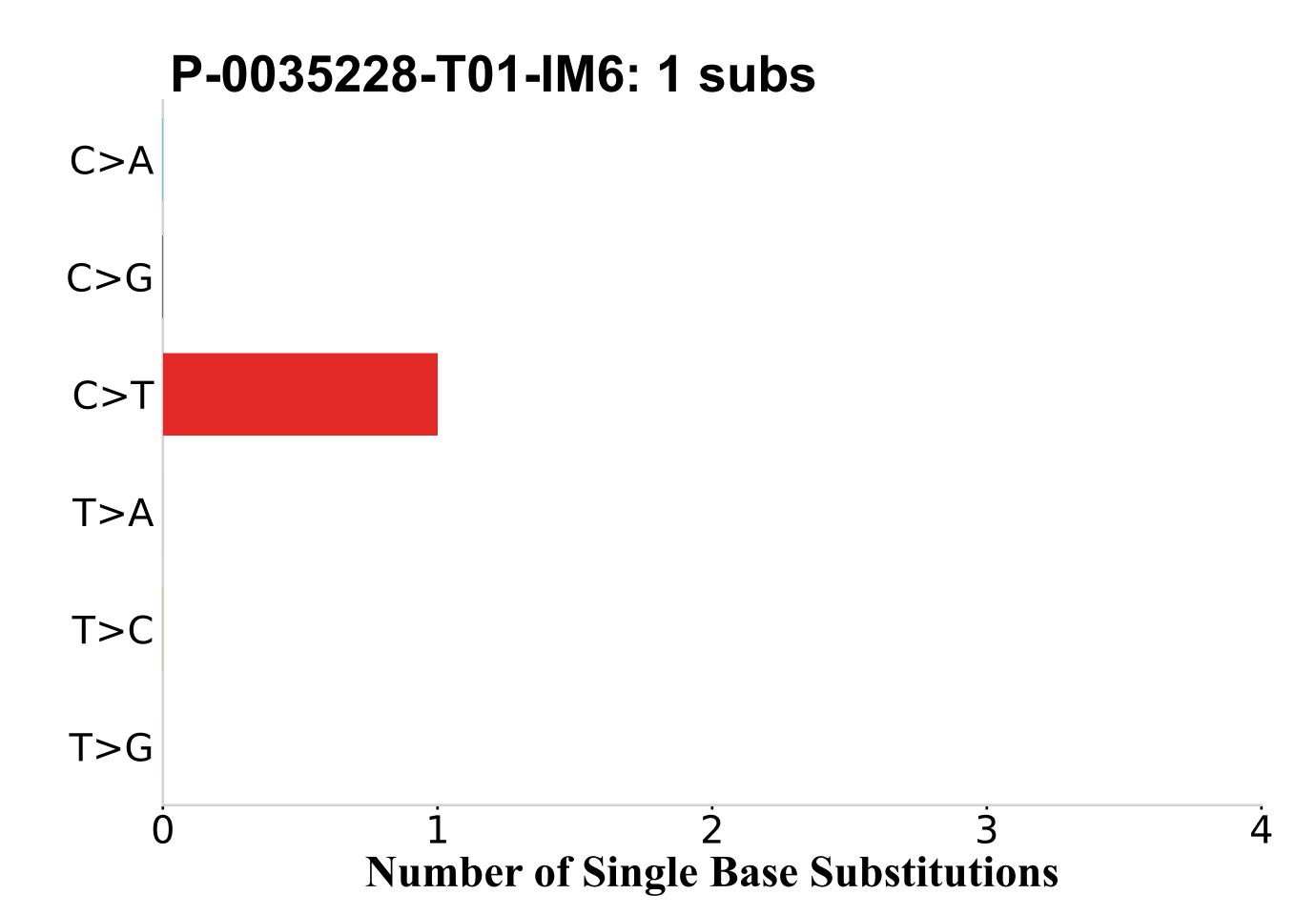


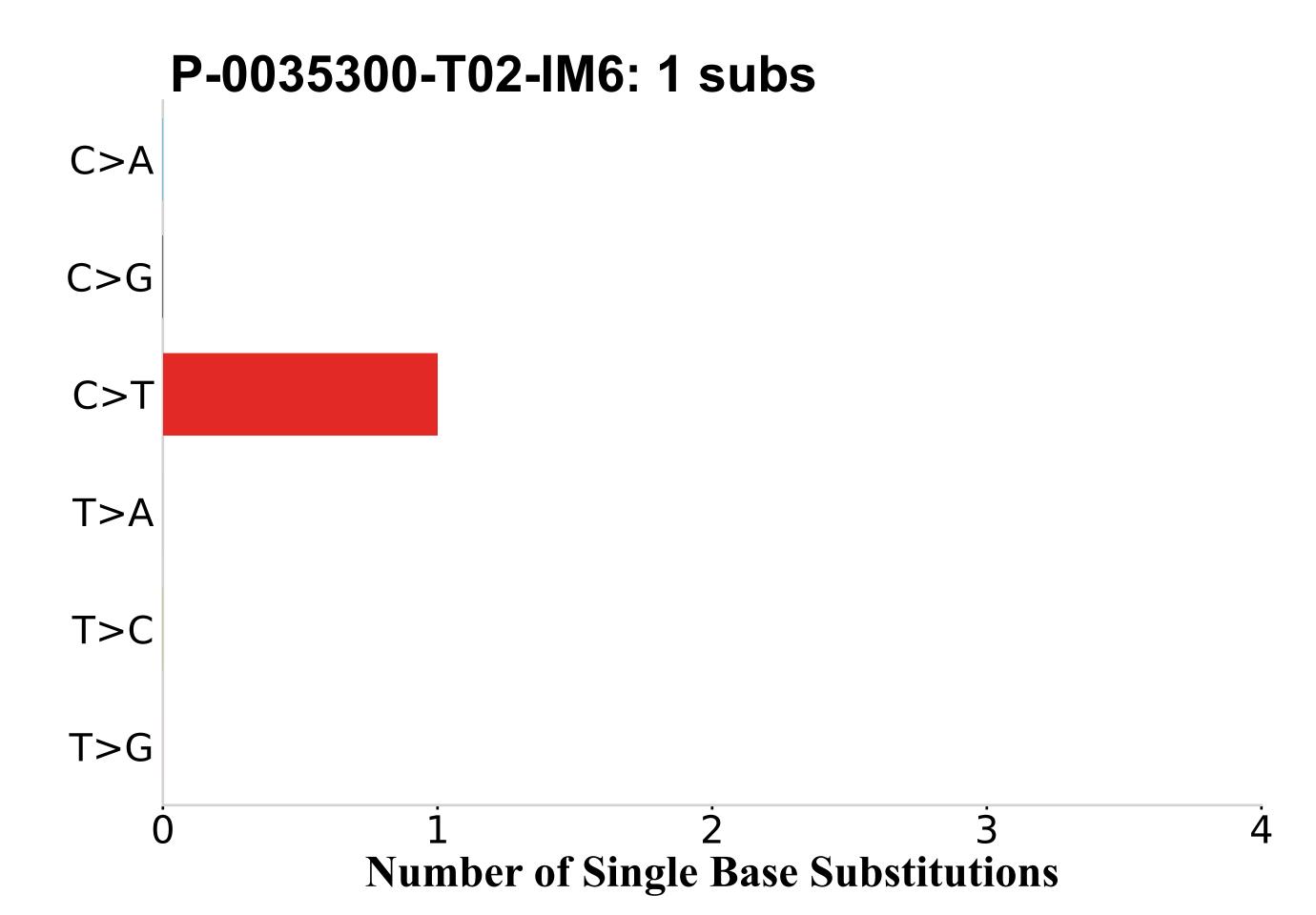


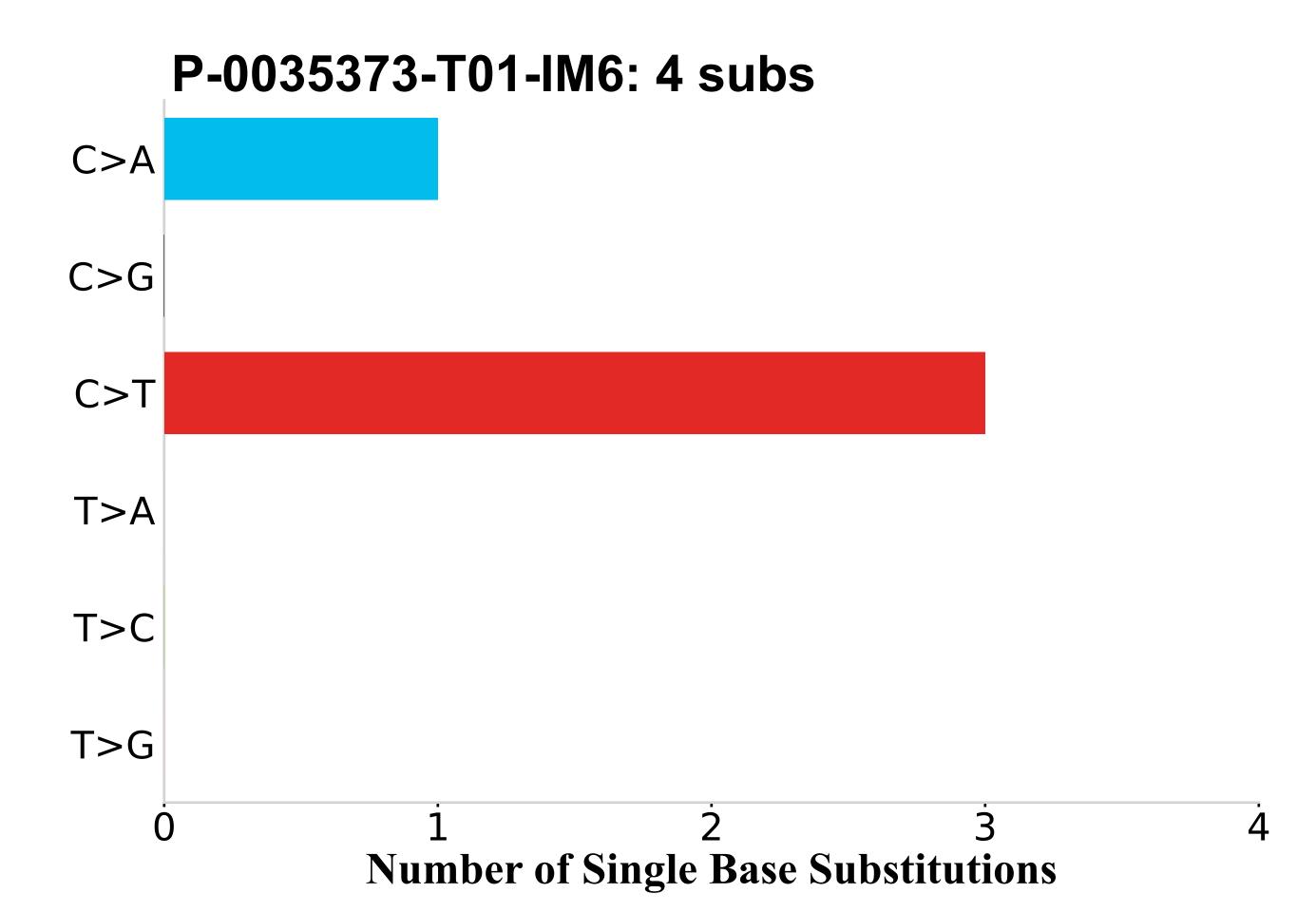


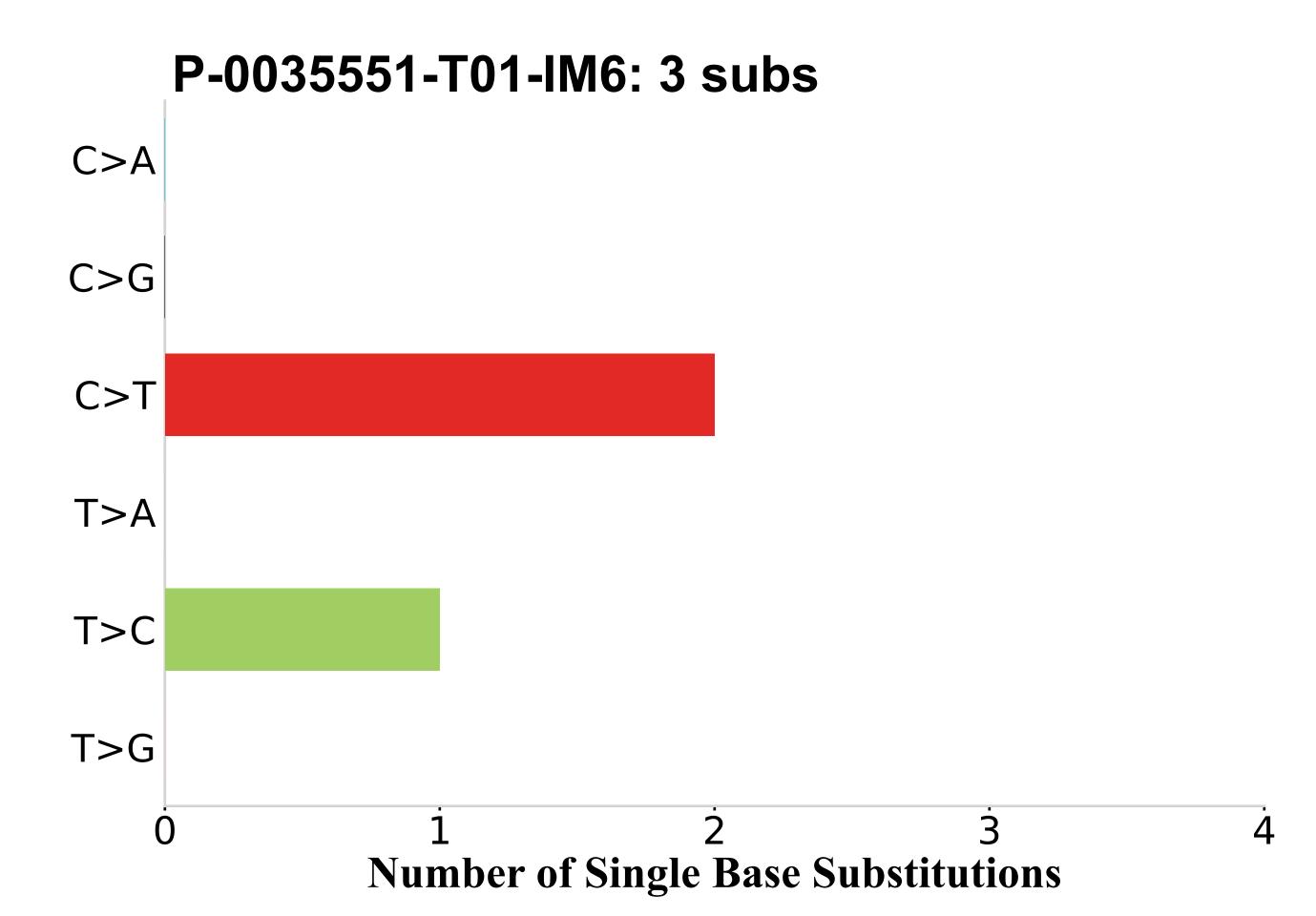


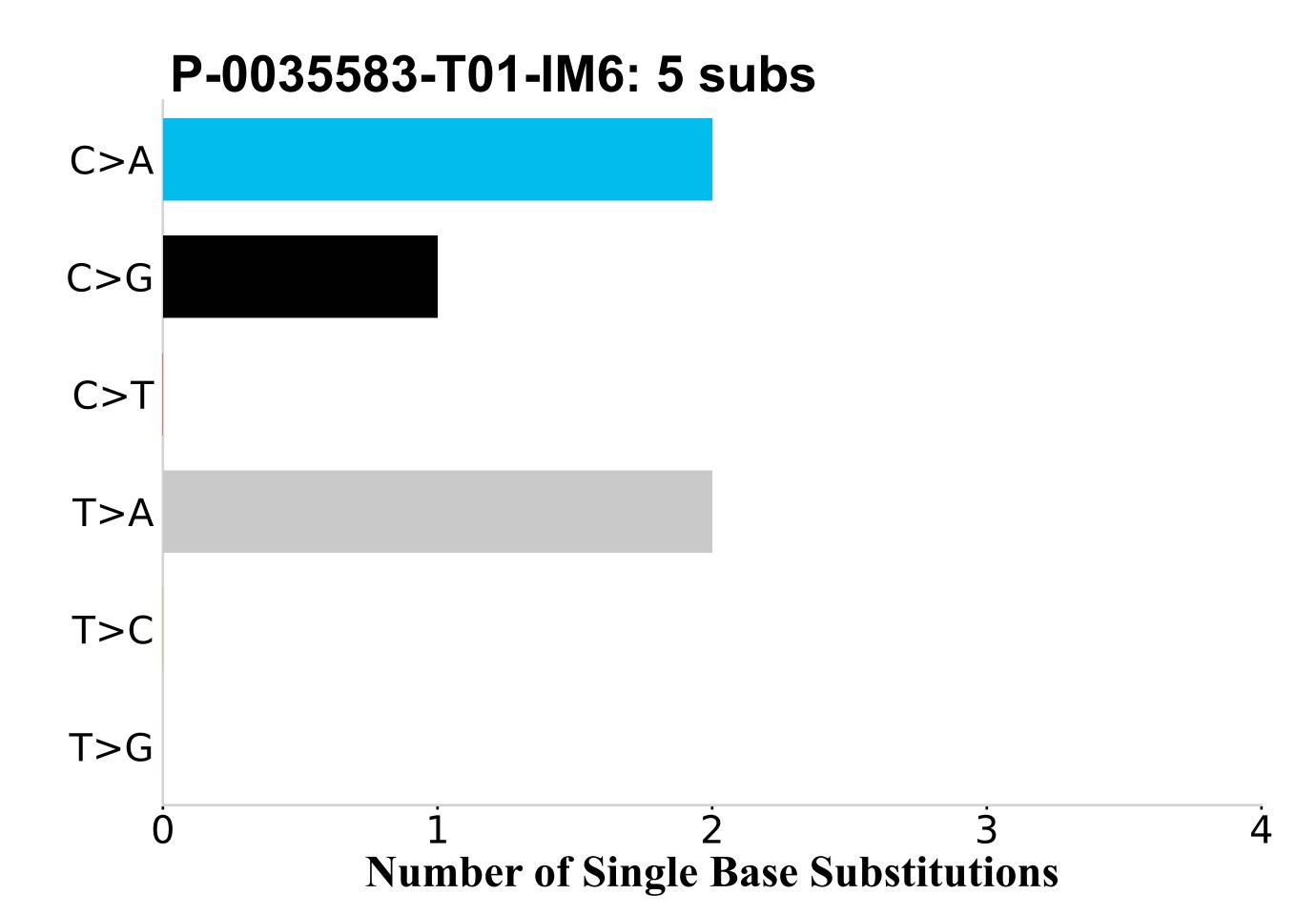


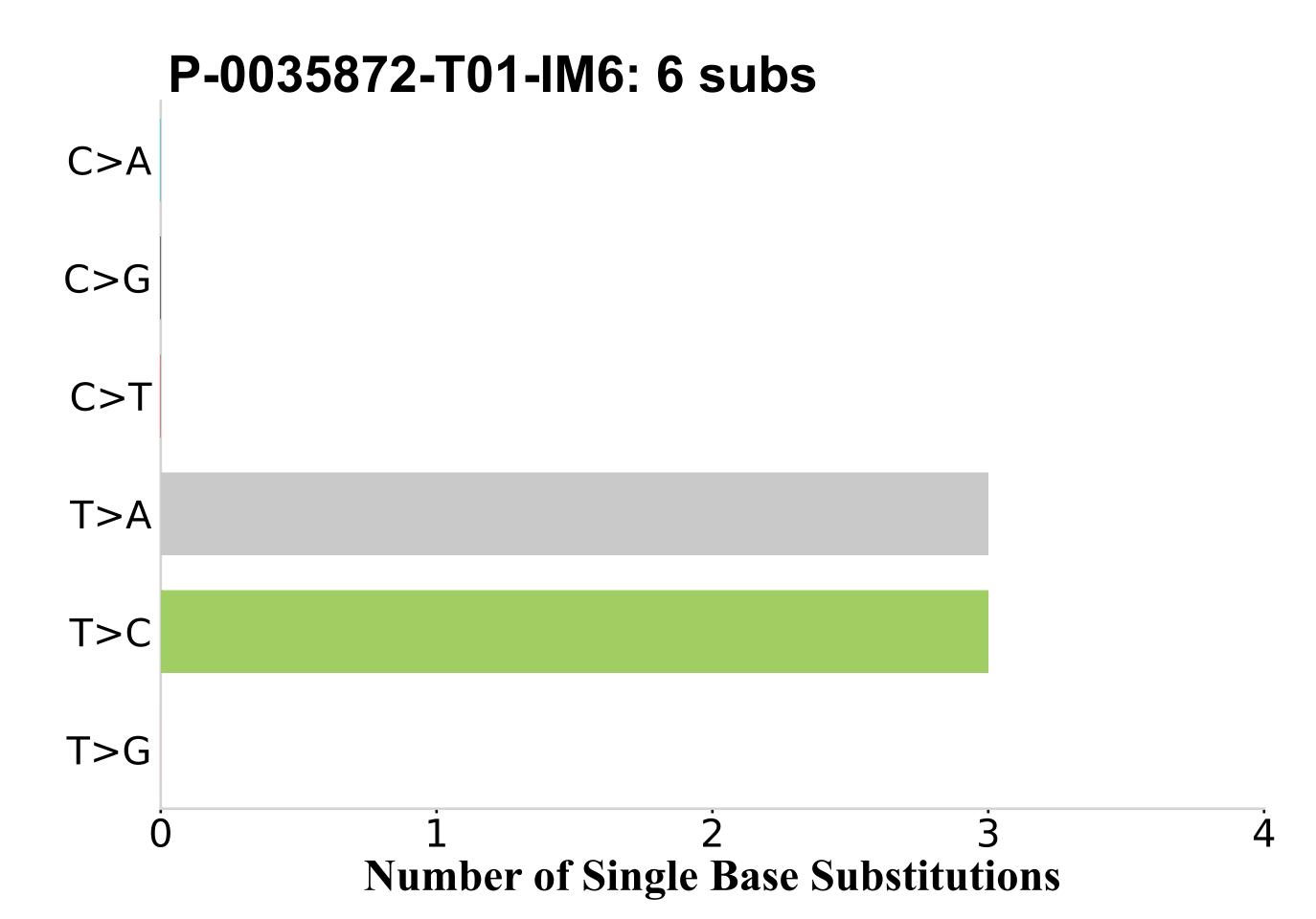




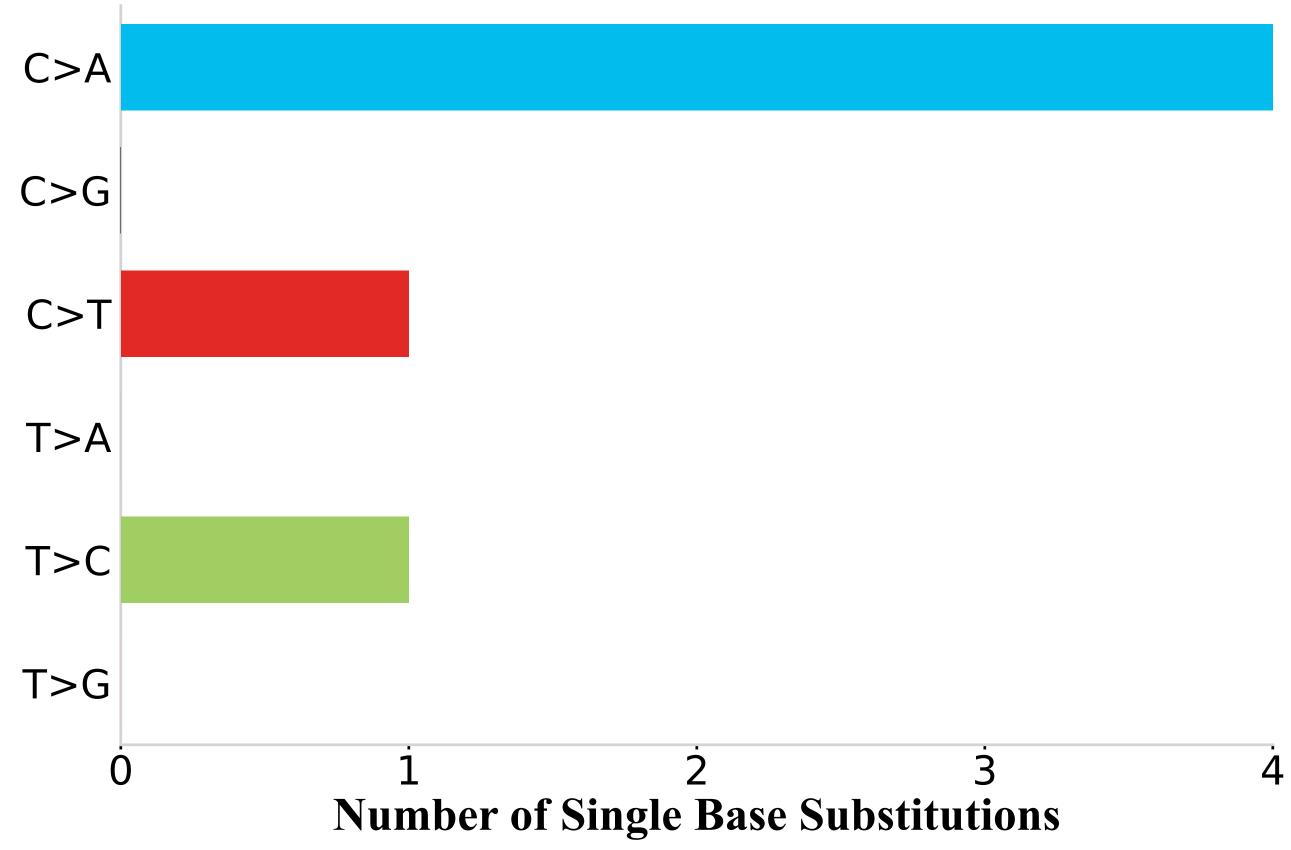


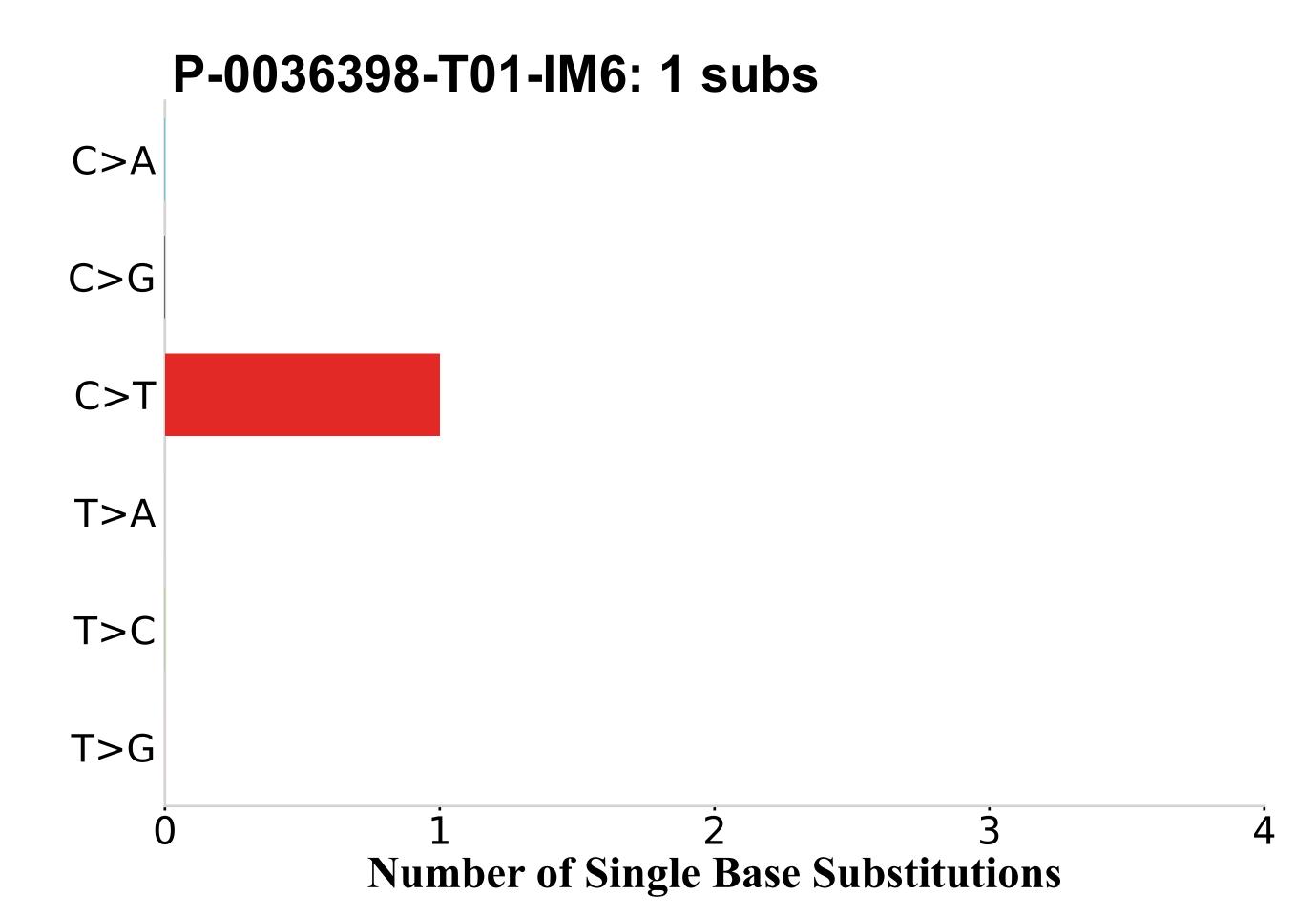


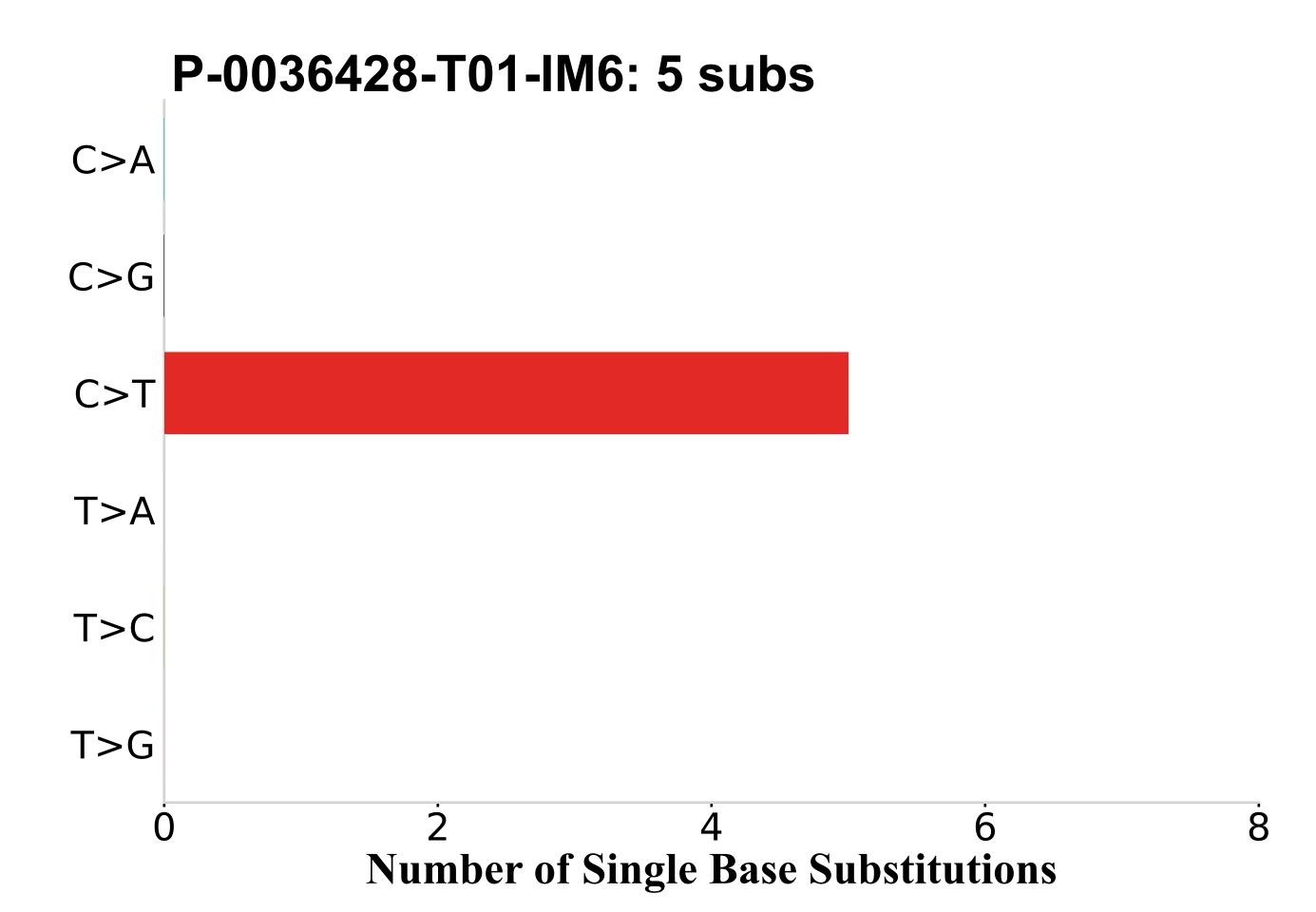




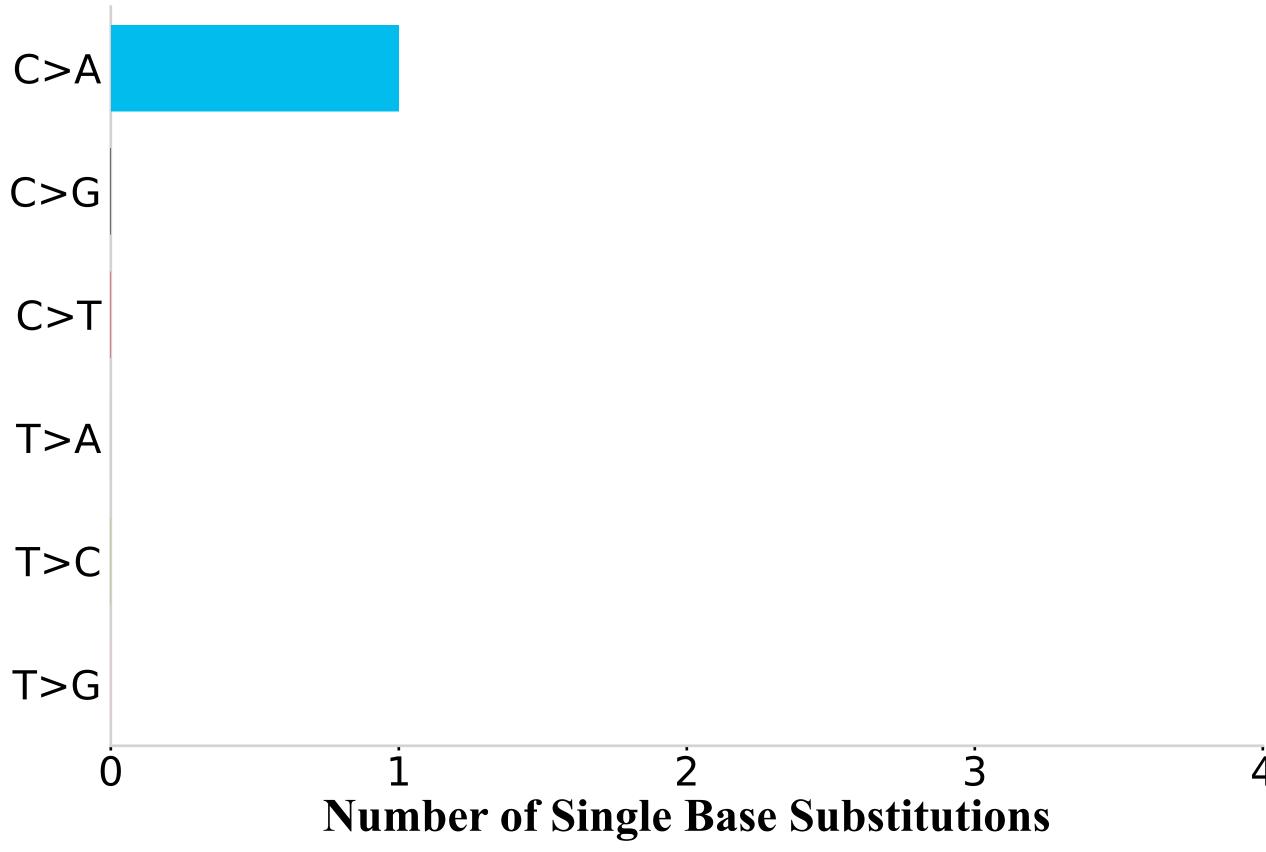


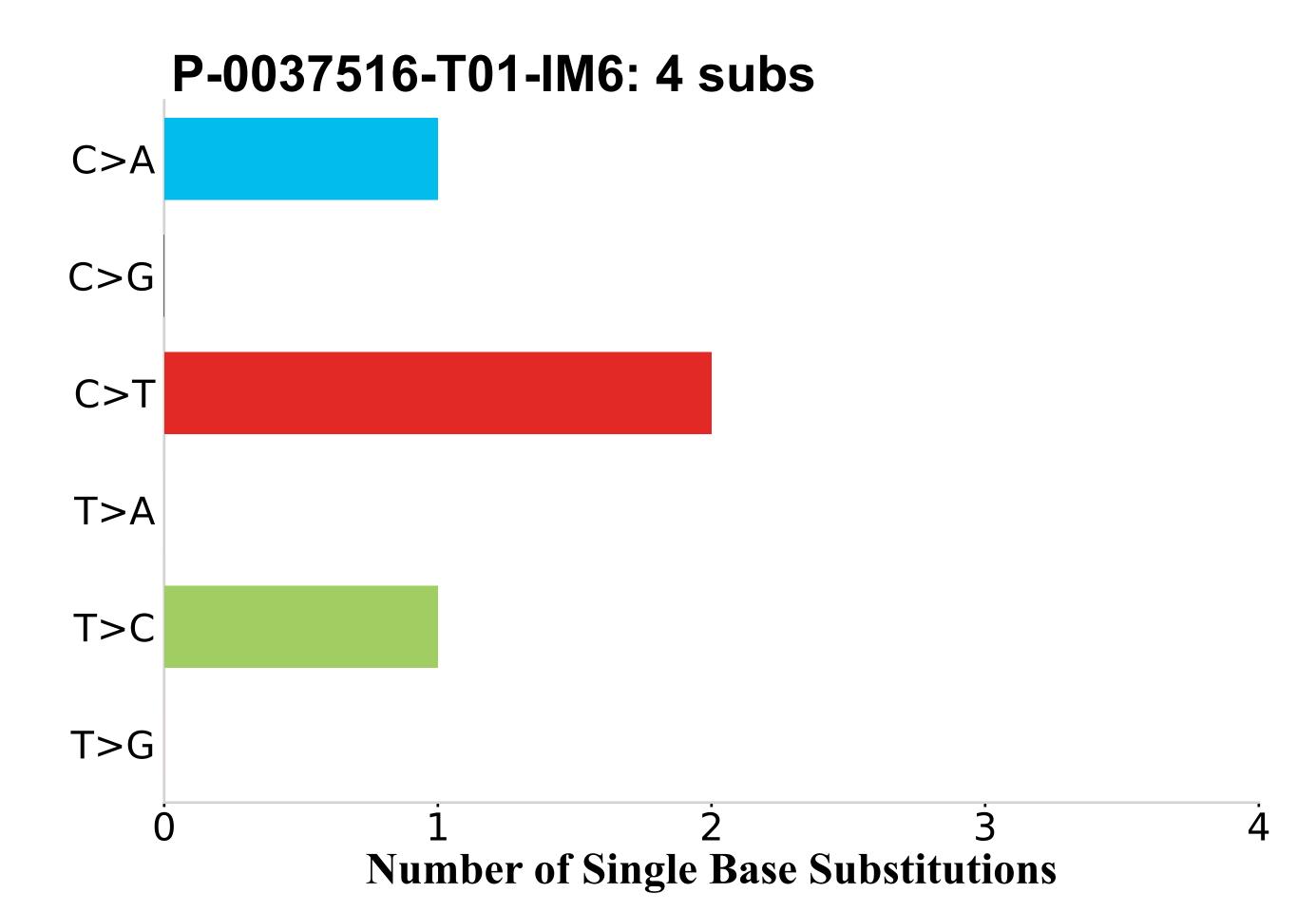


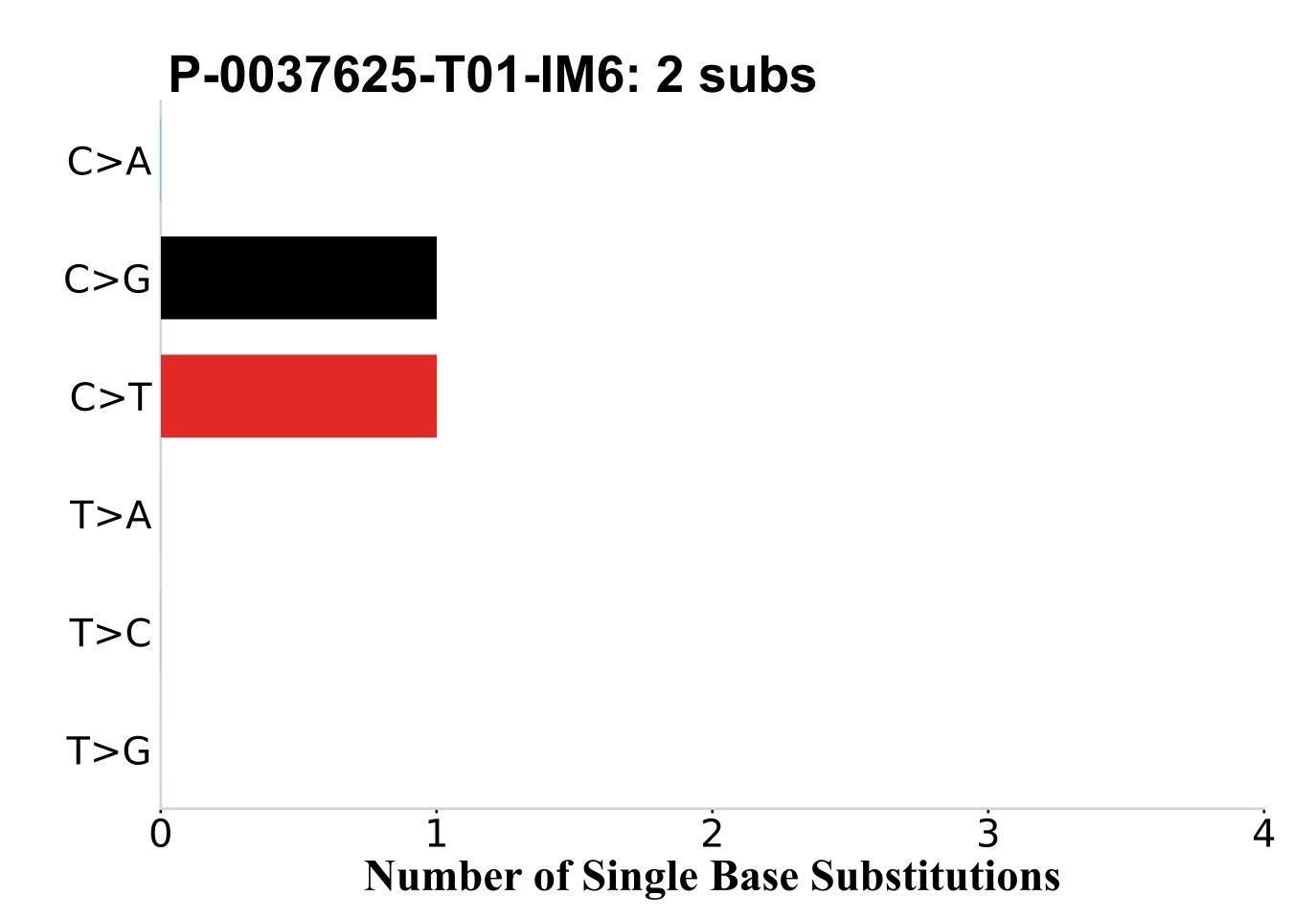


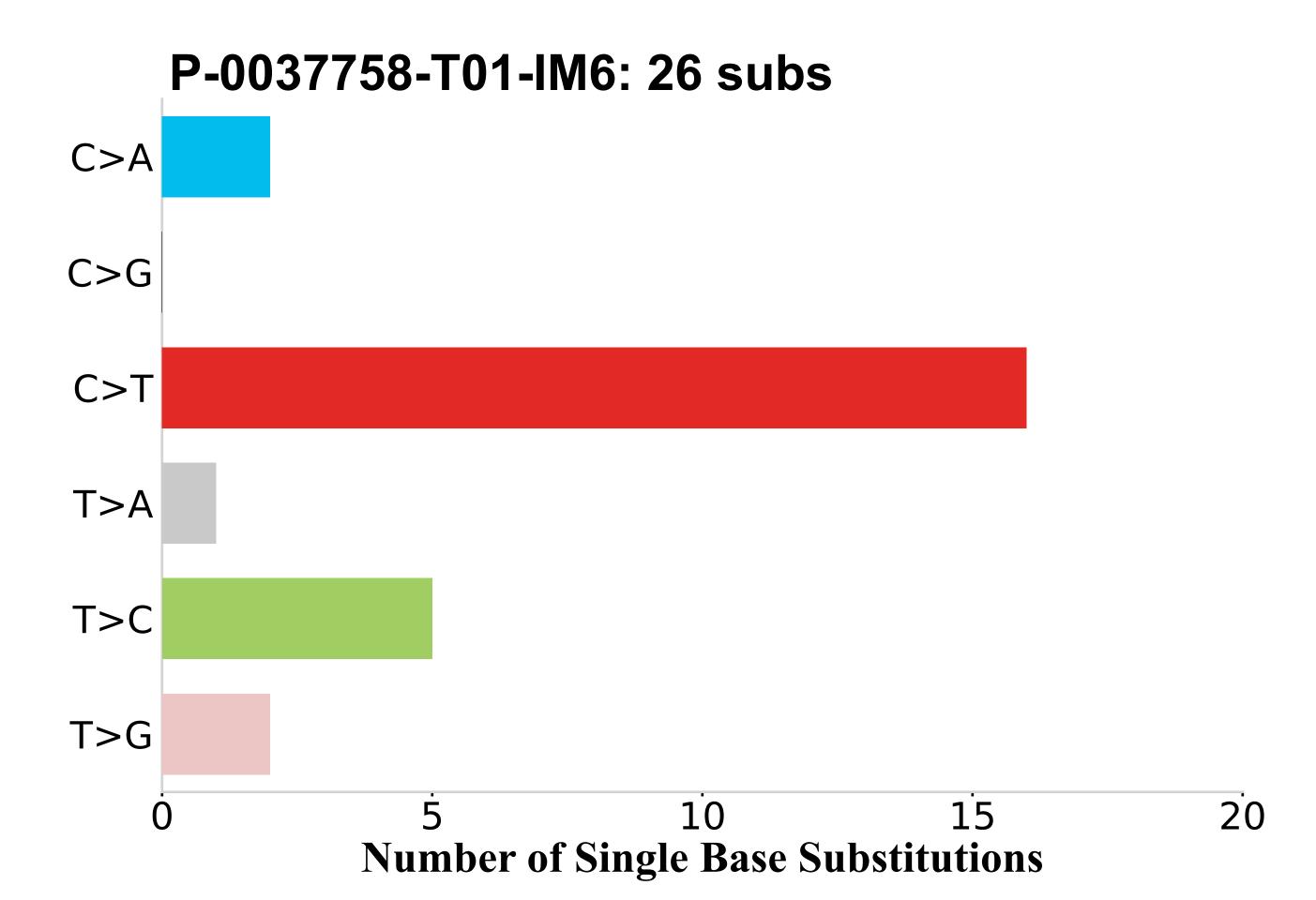


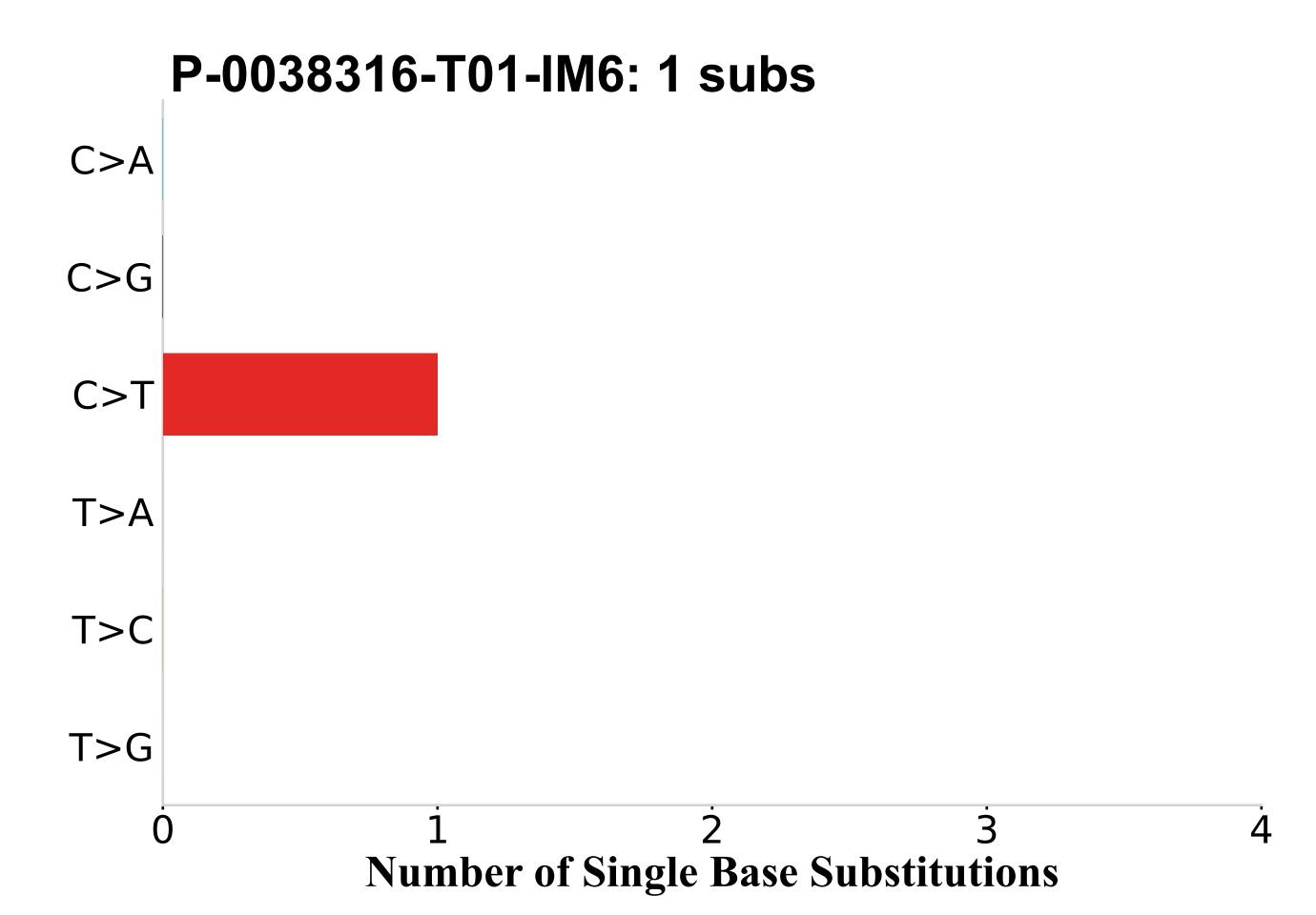


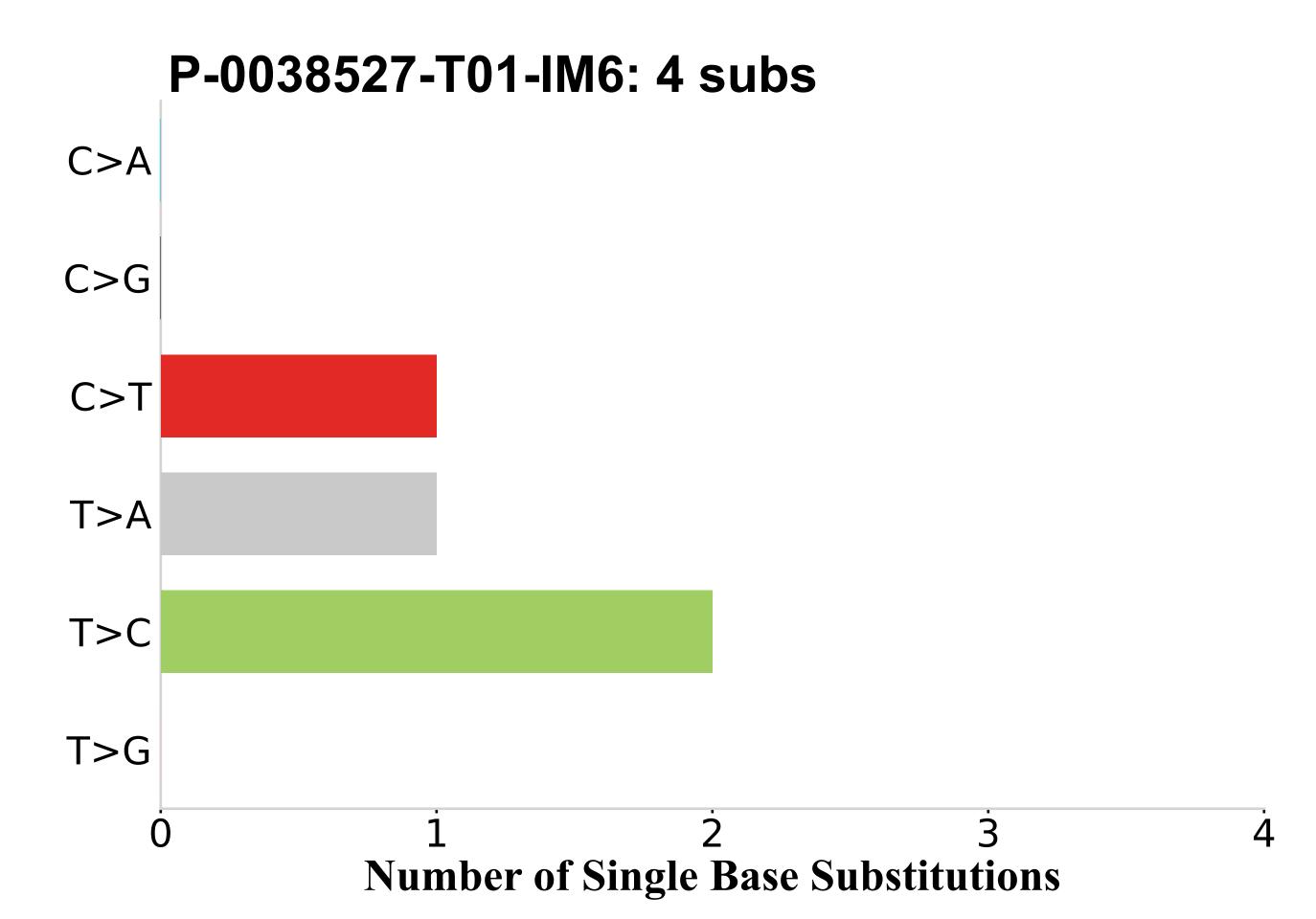


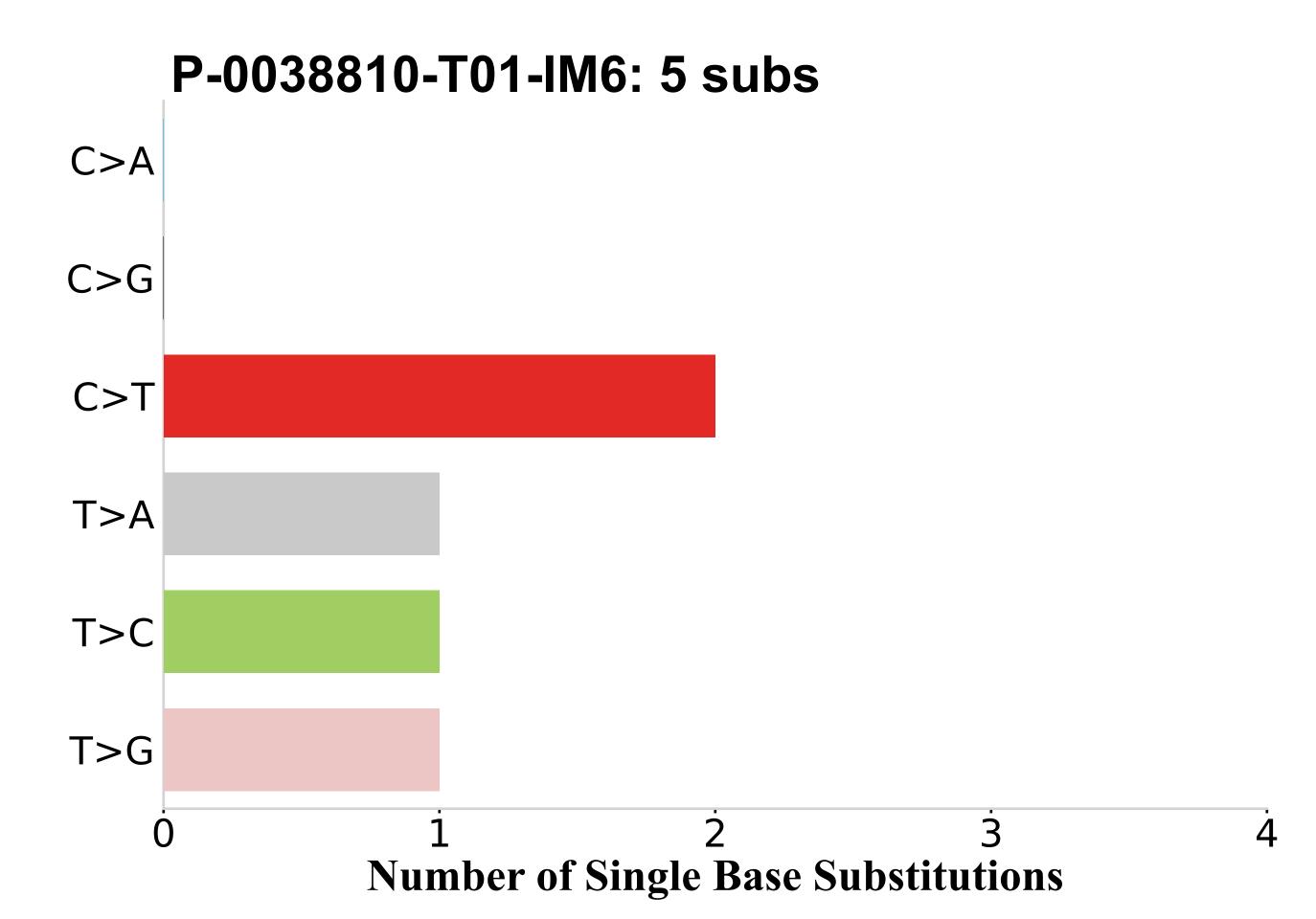


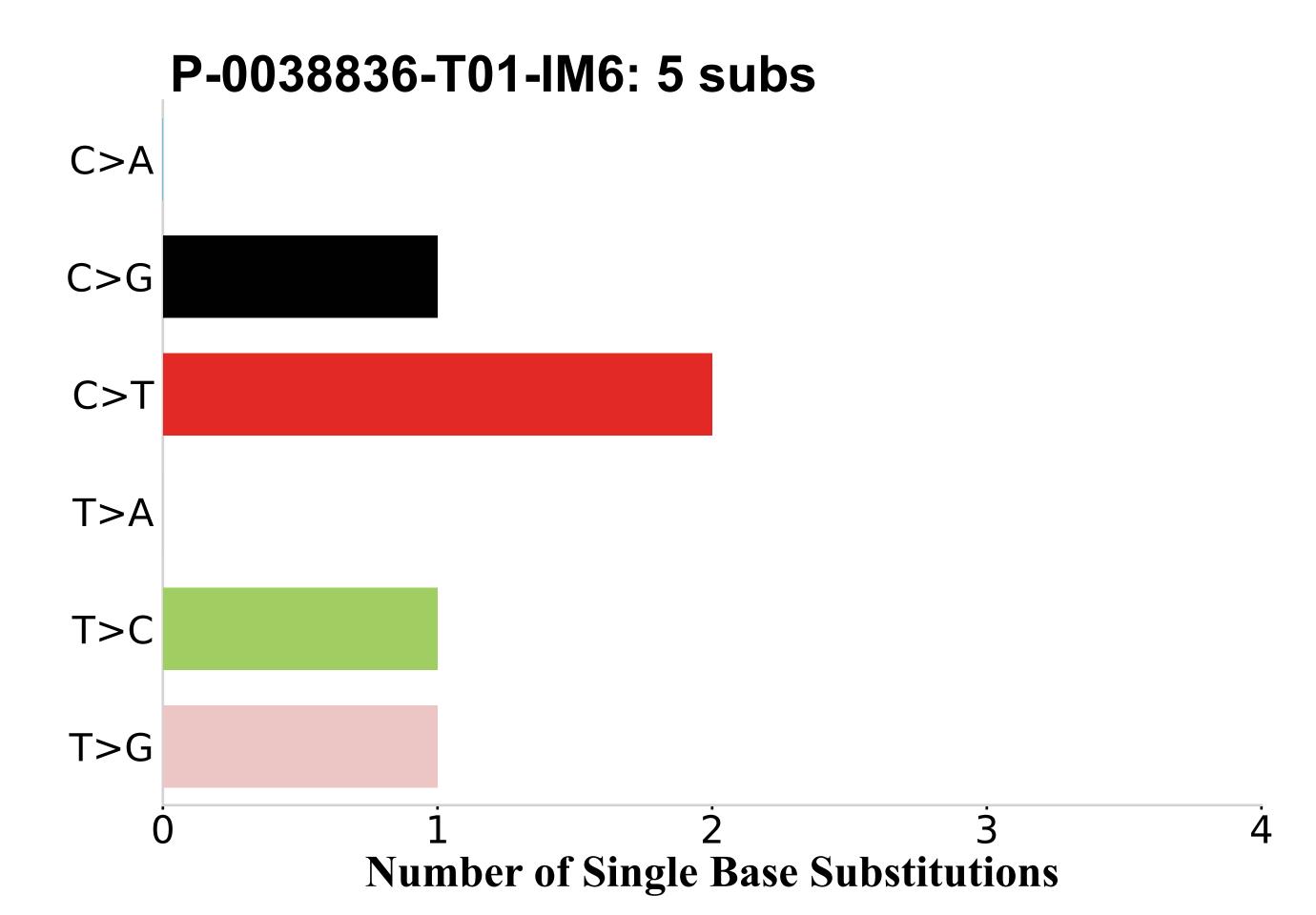


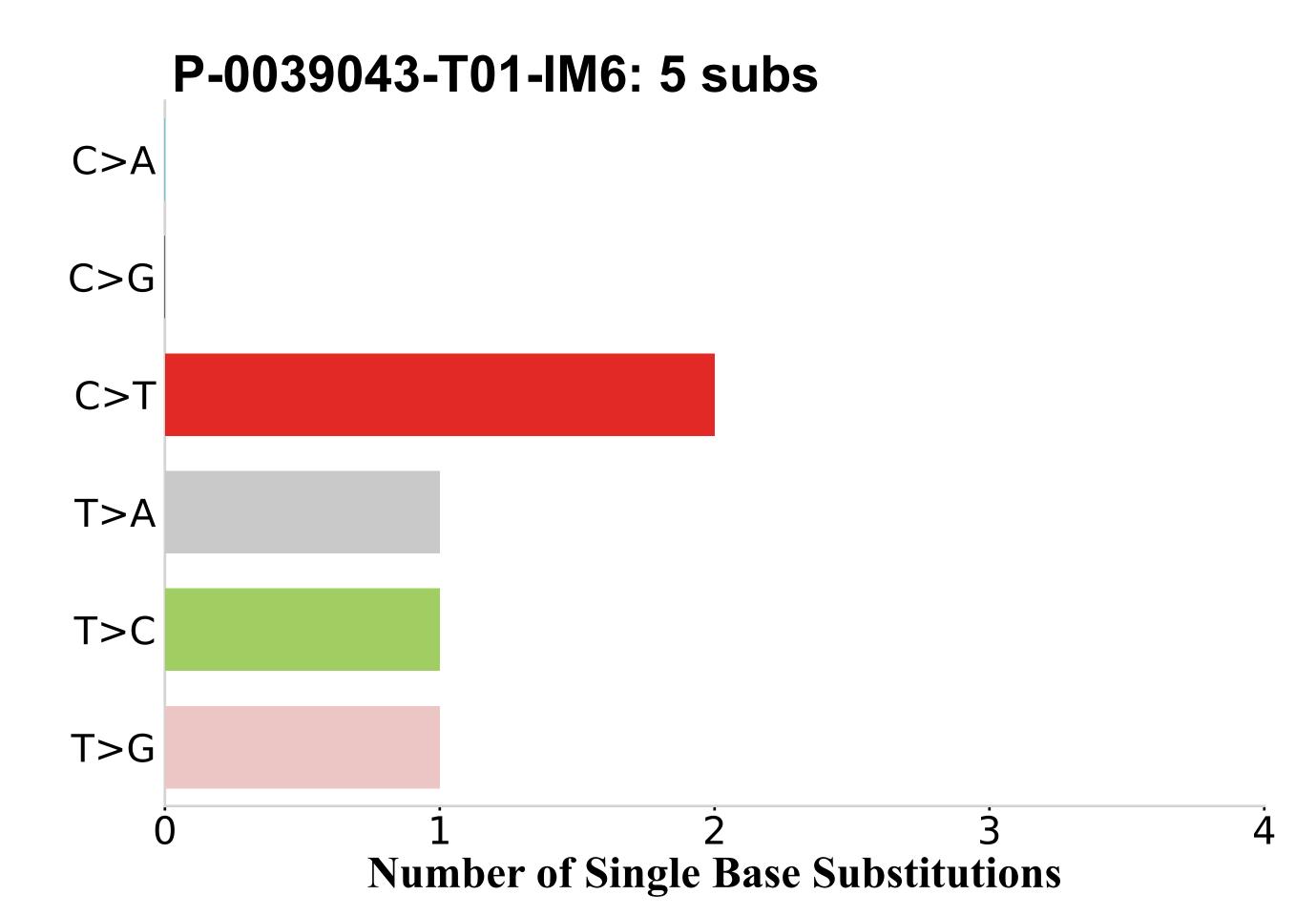








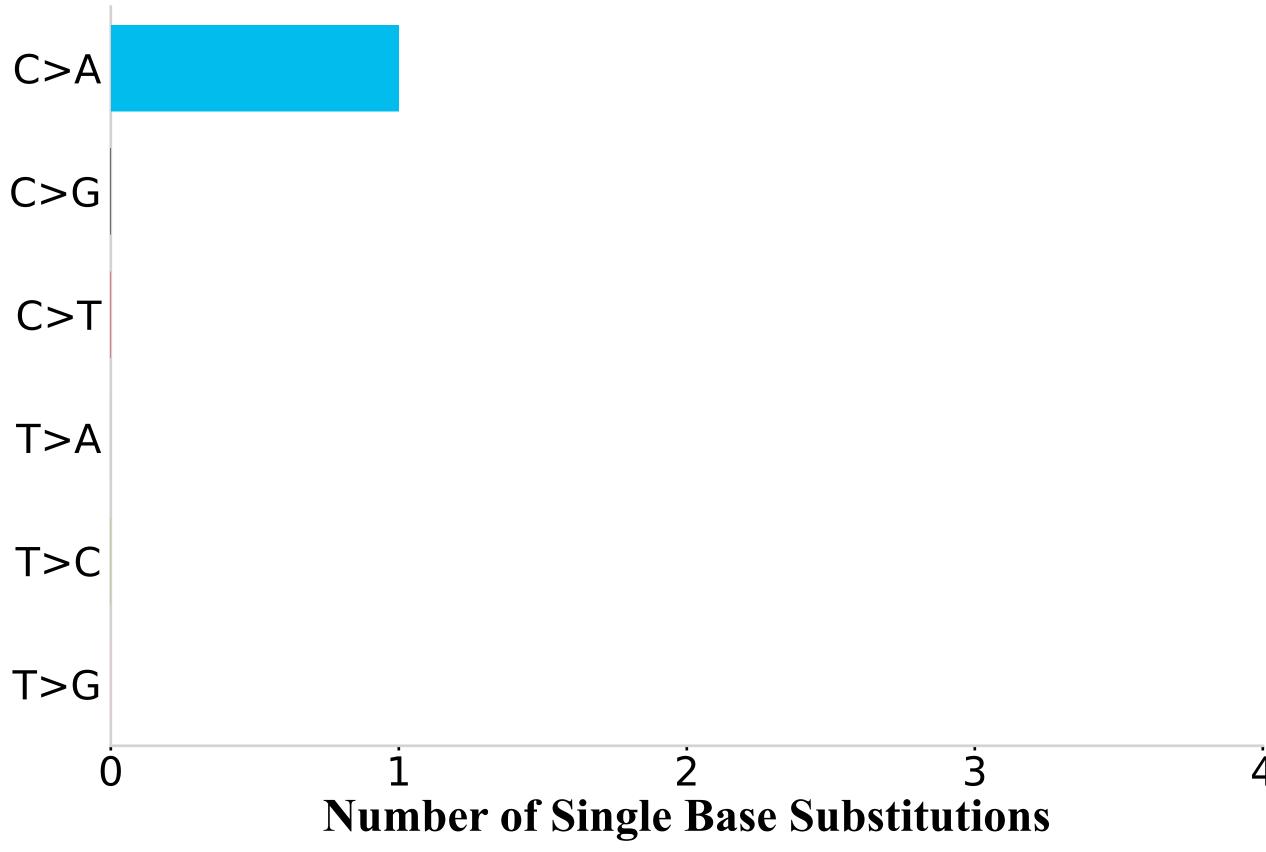


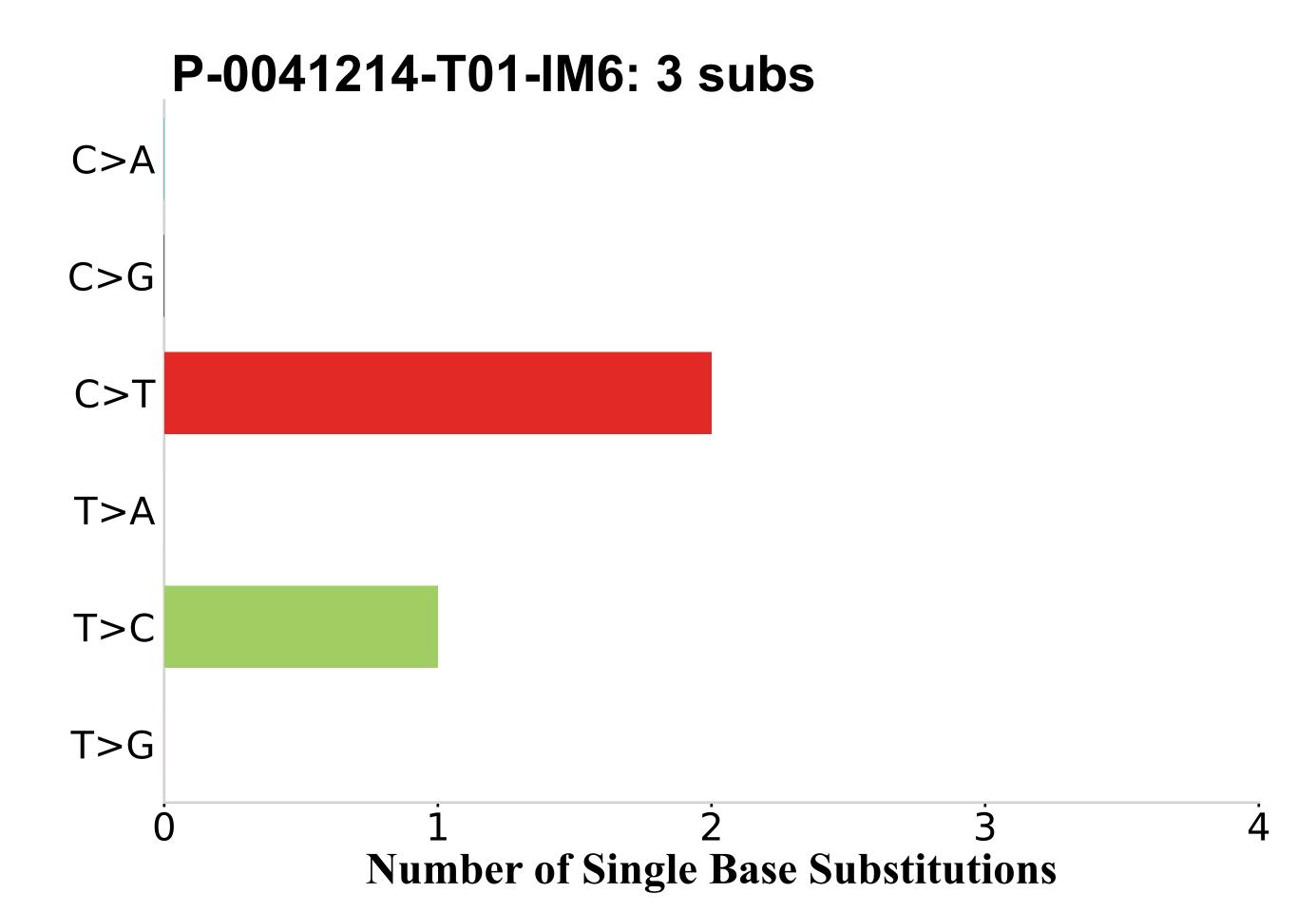


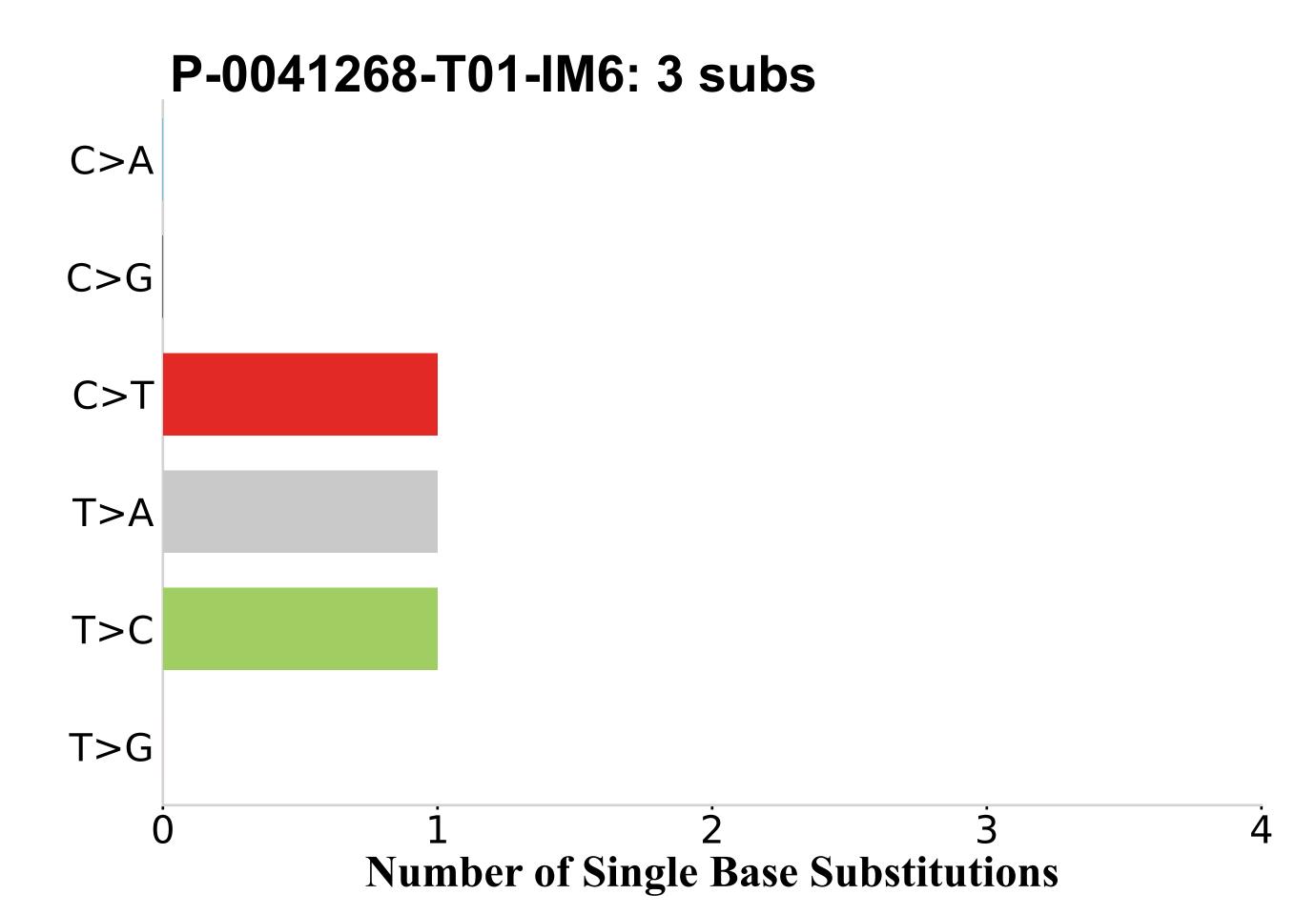
P-0039634-T01-IM6: 4 subs C>A C>G C>T T>A T>C T>G

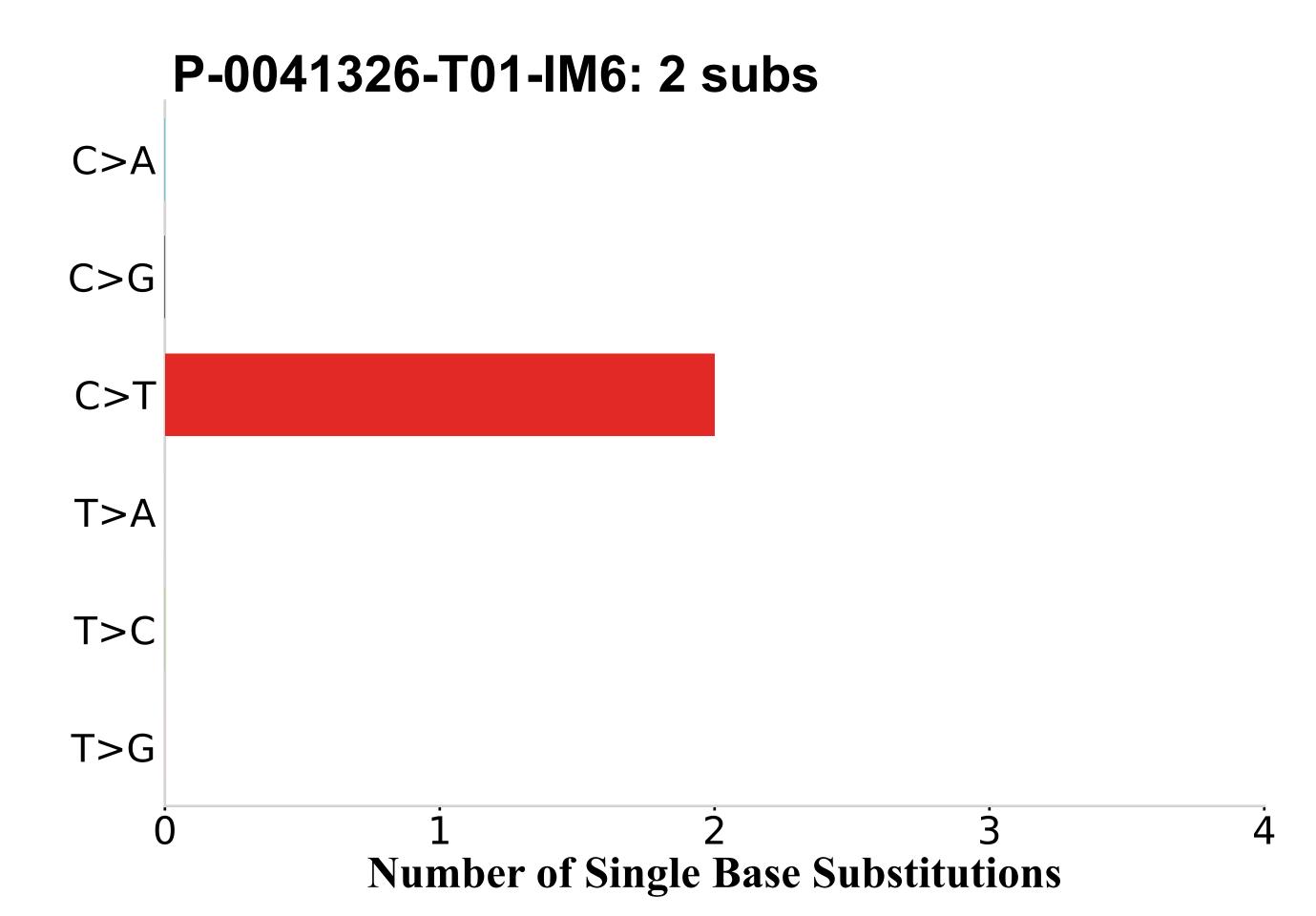
Number of Single Base Substitutions

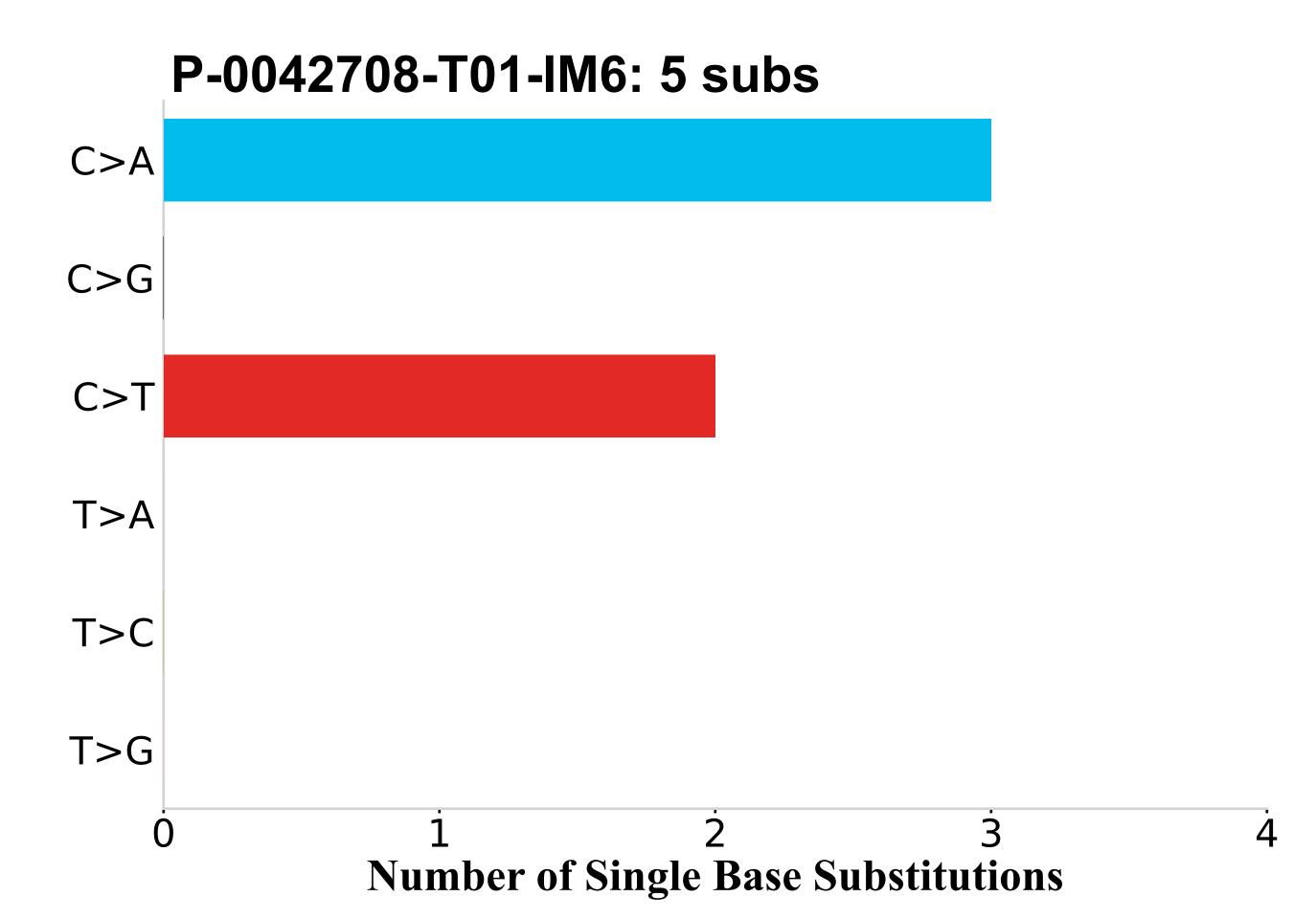




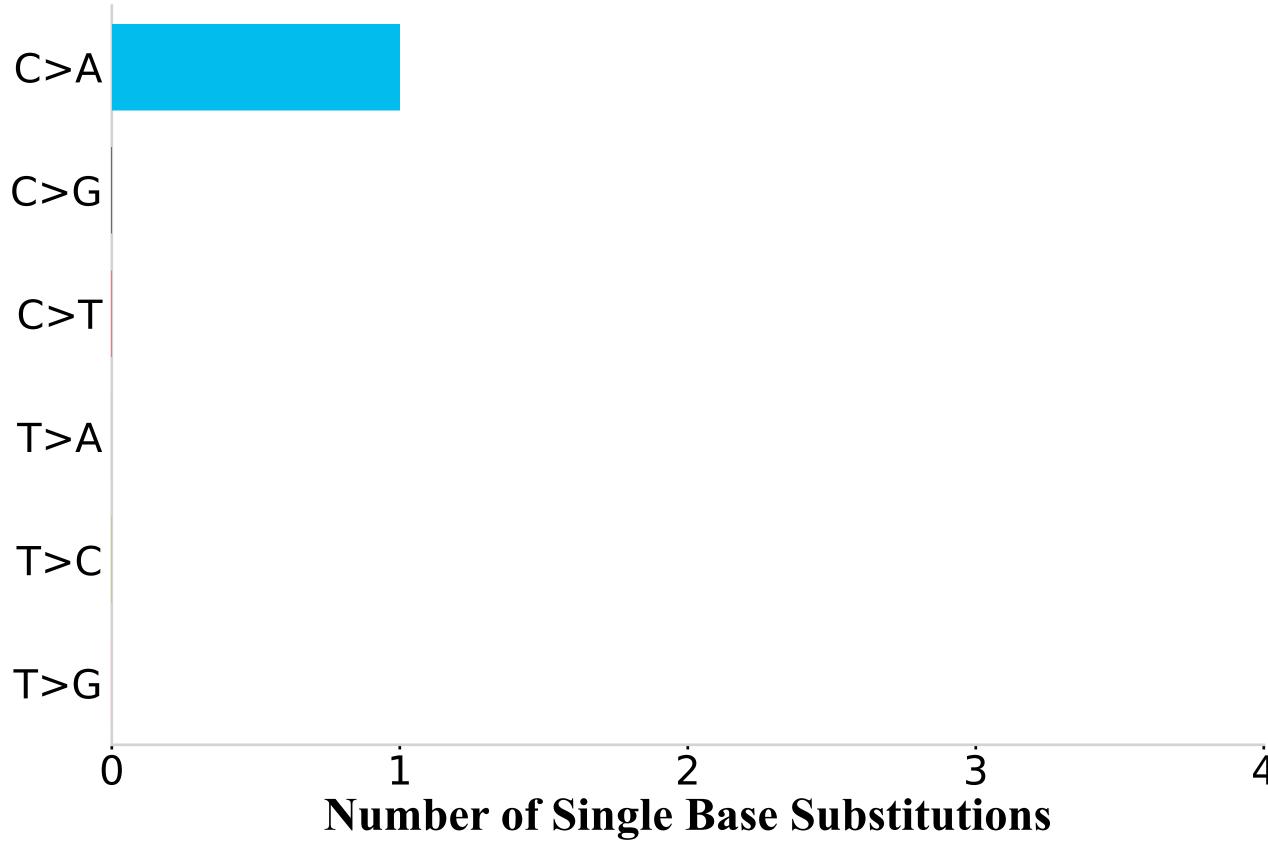




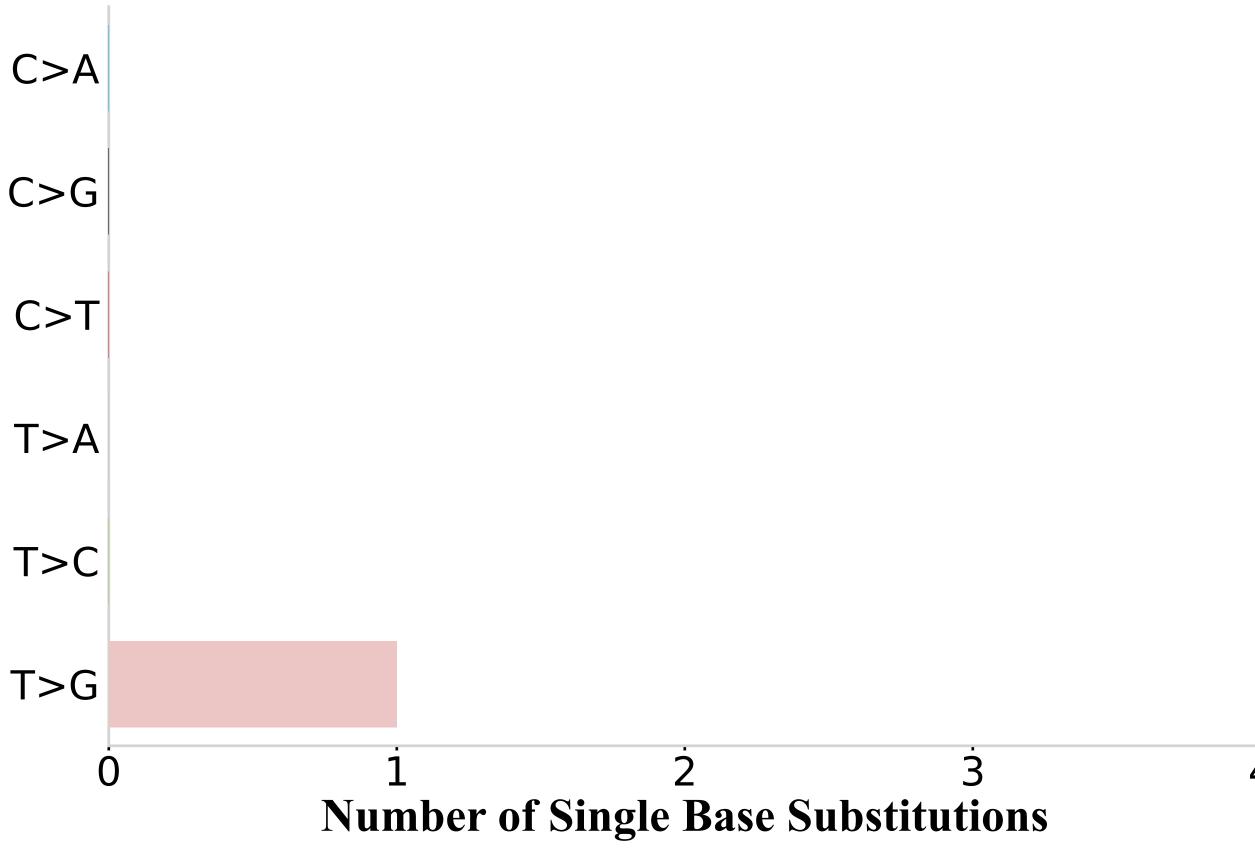


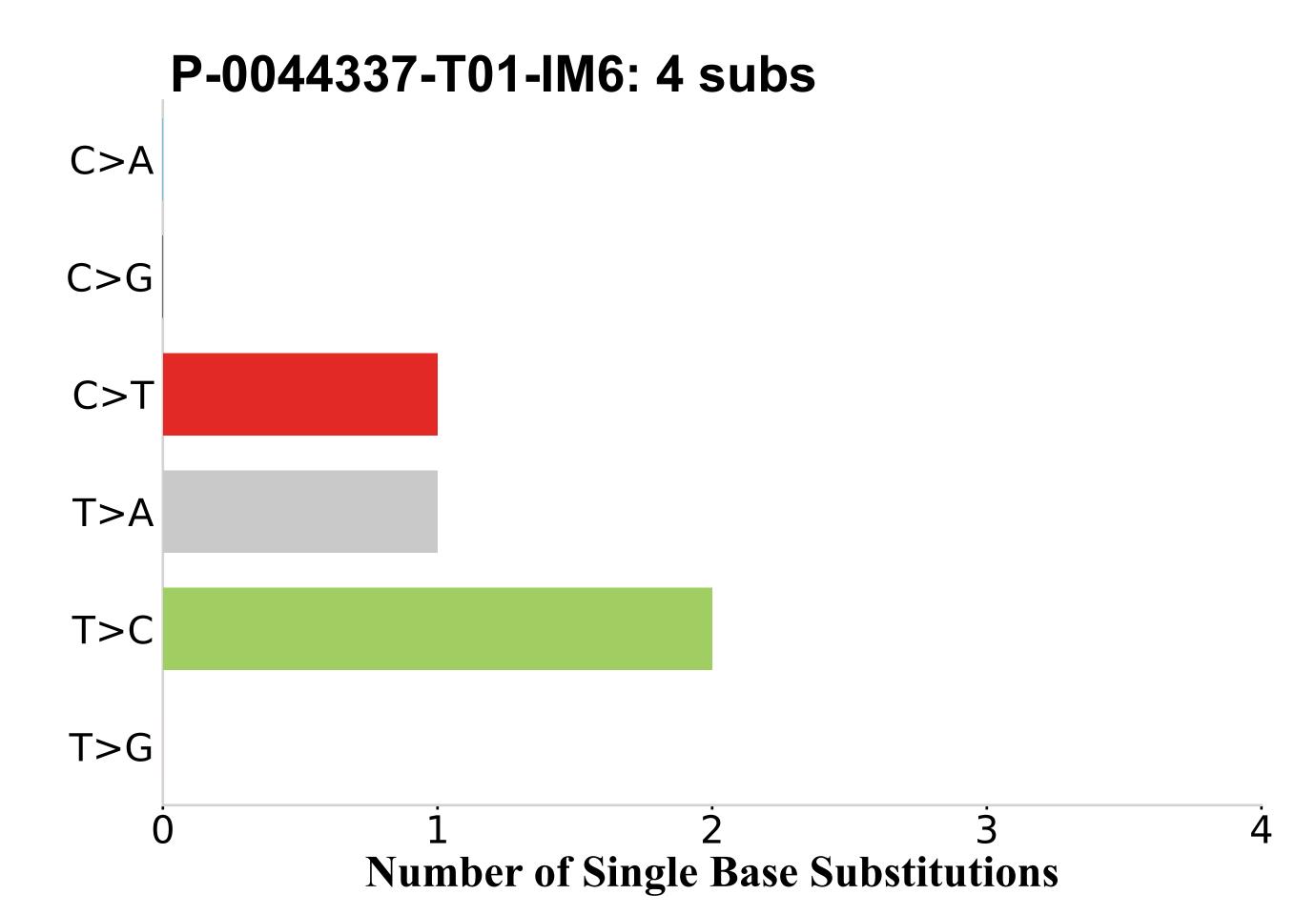


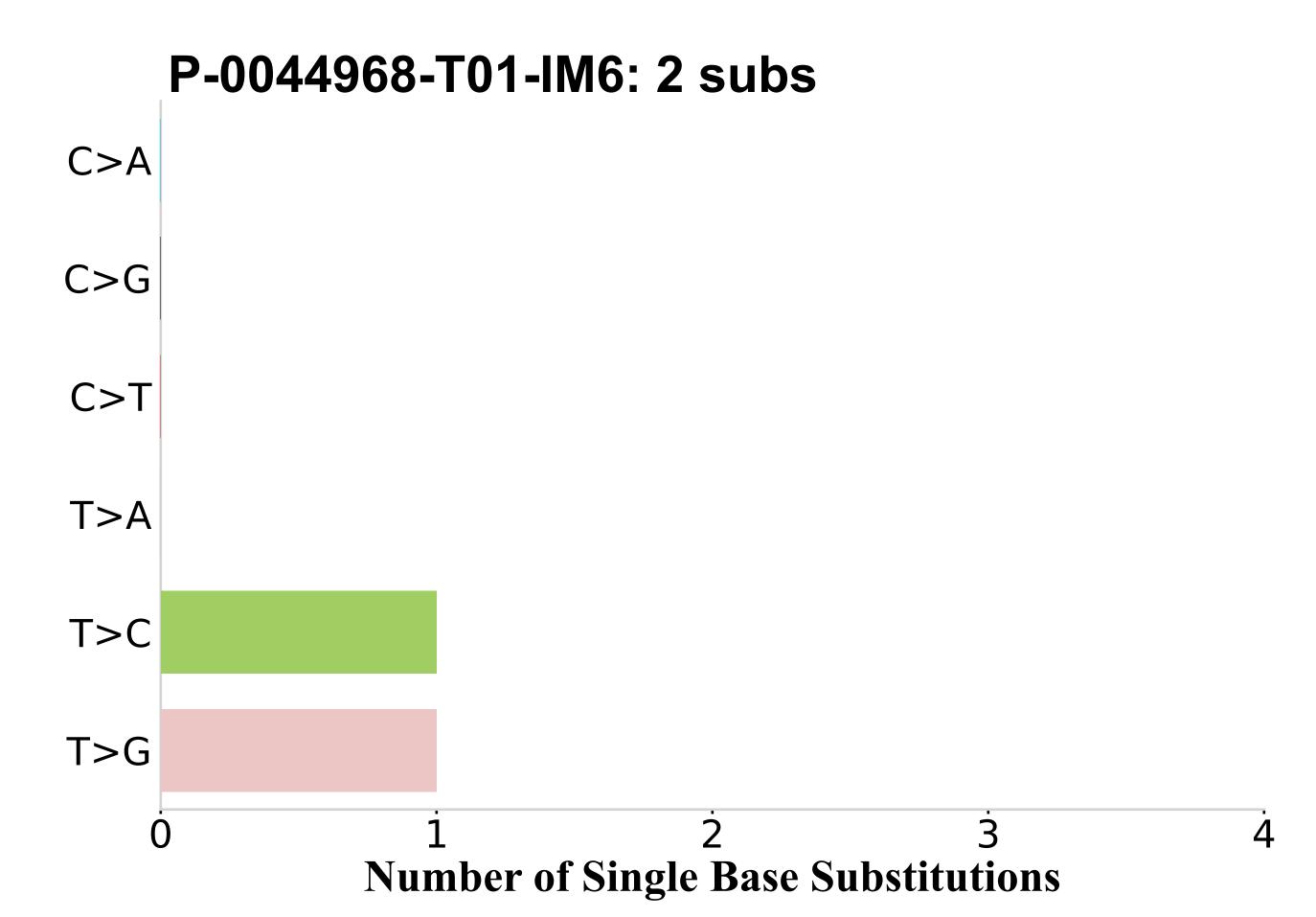


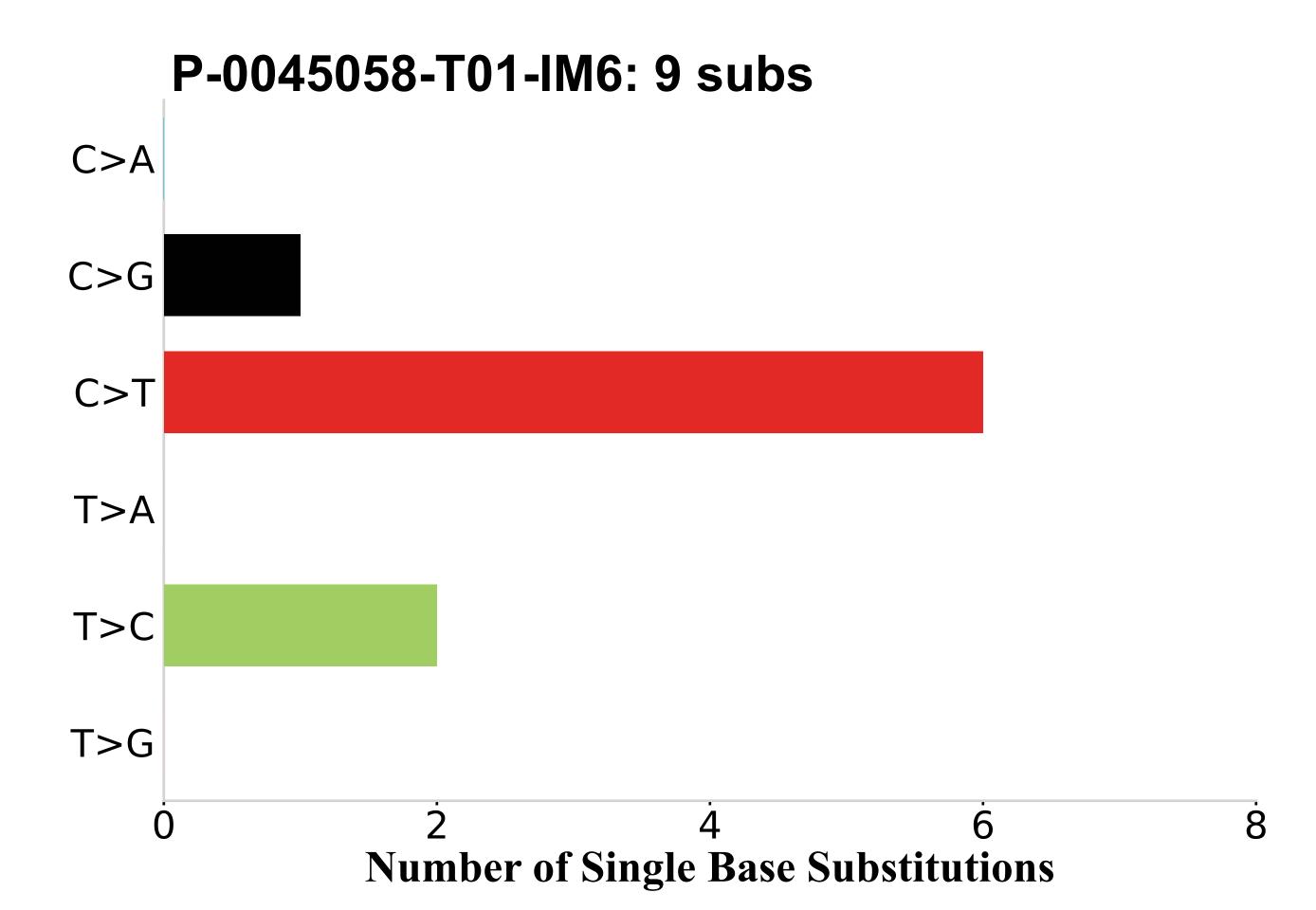


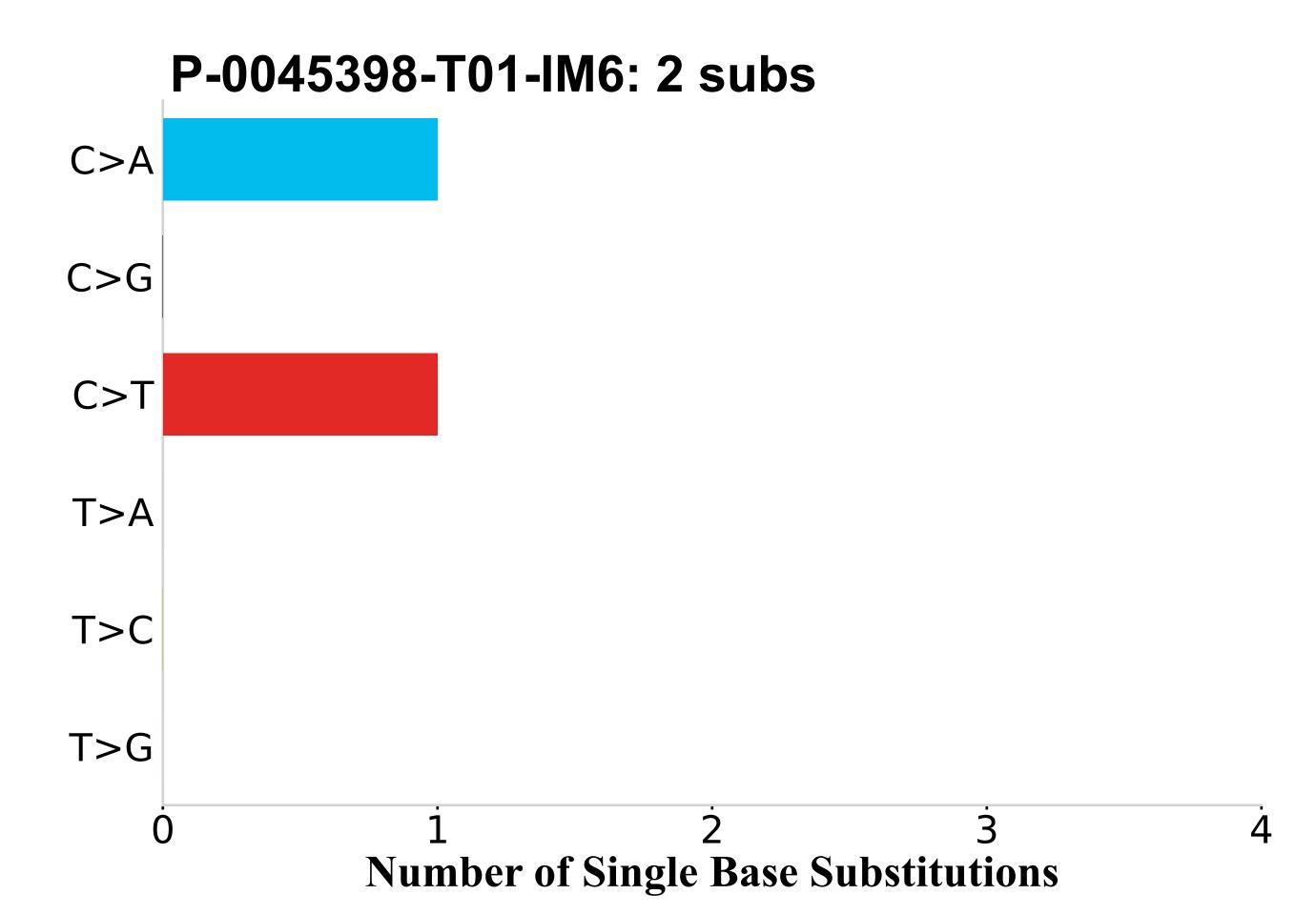


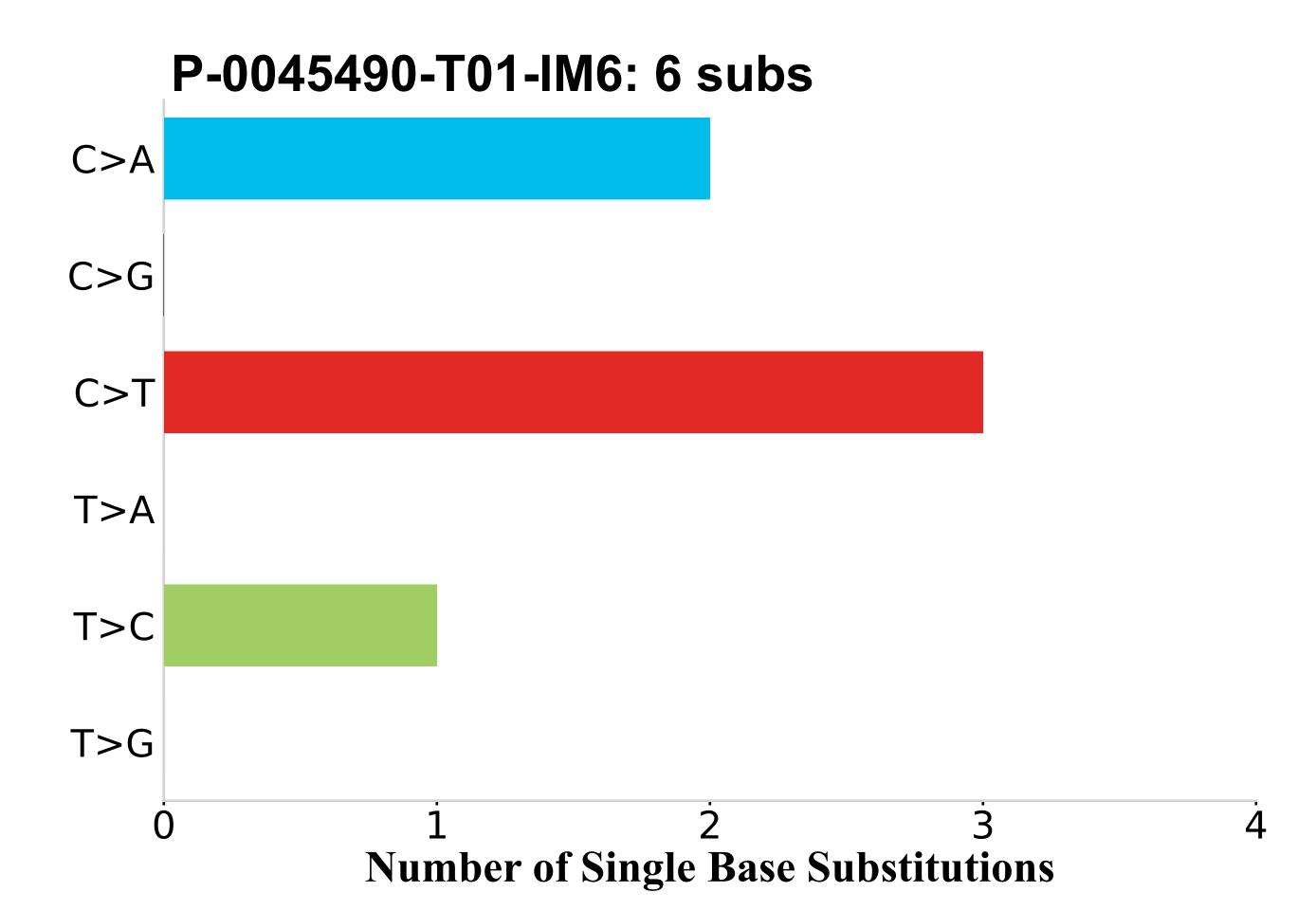


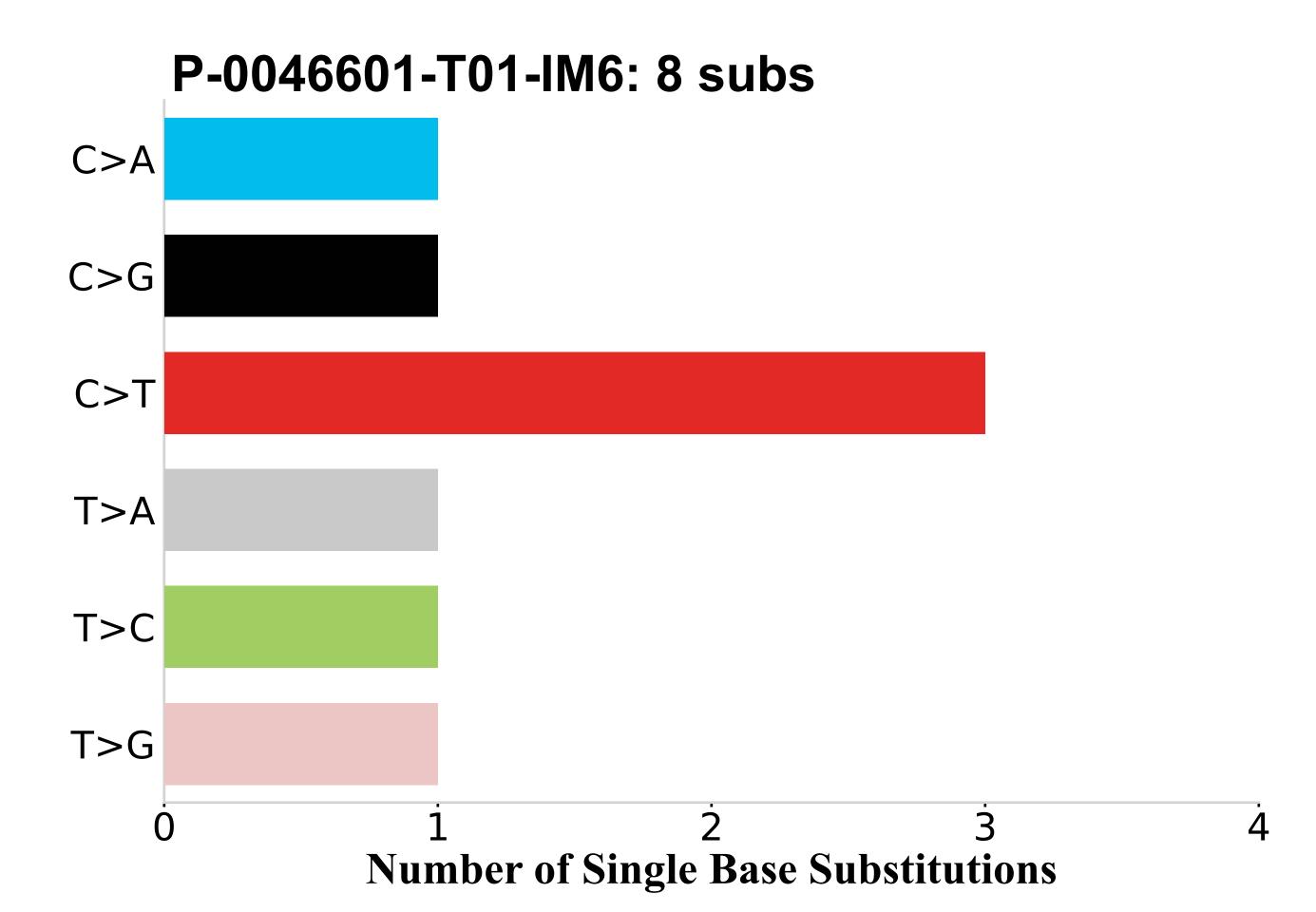




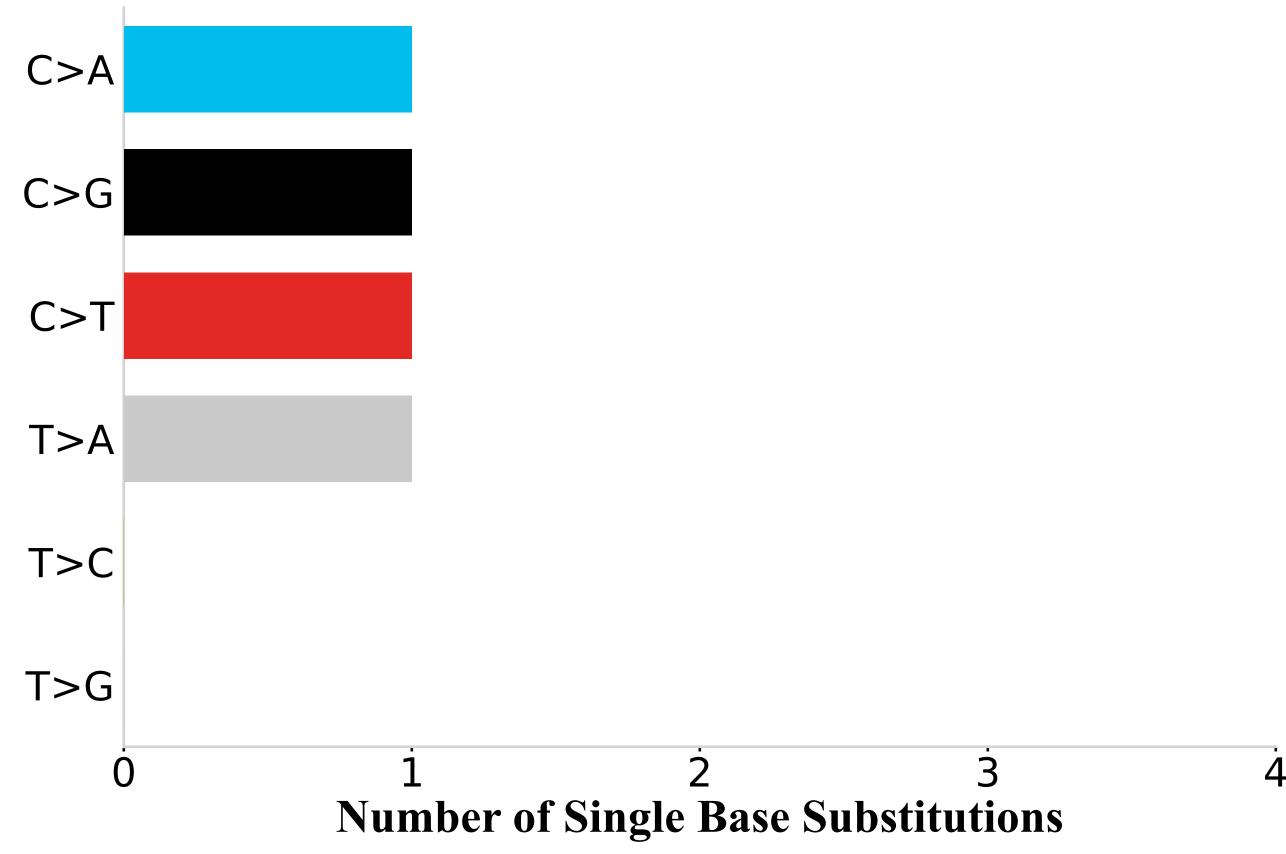


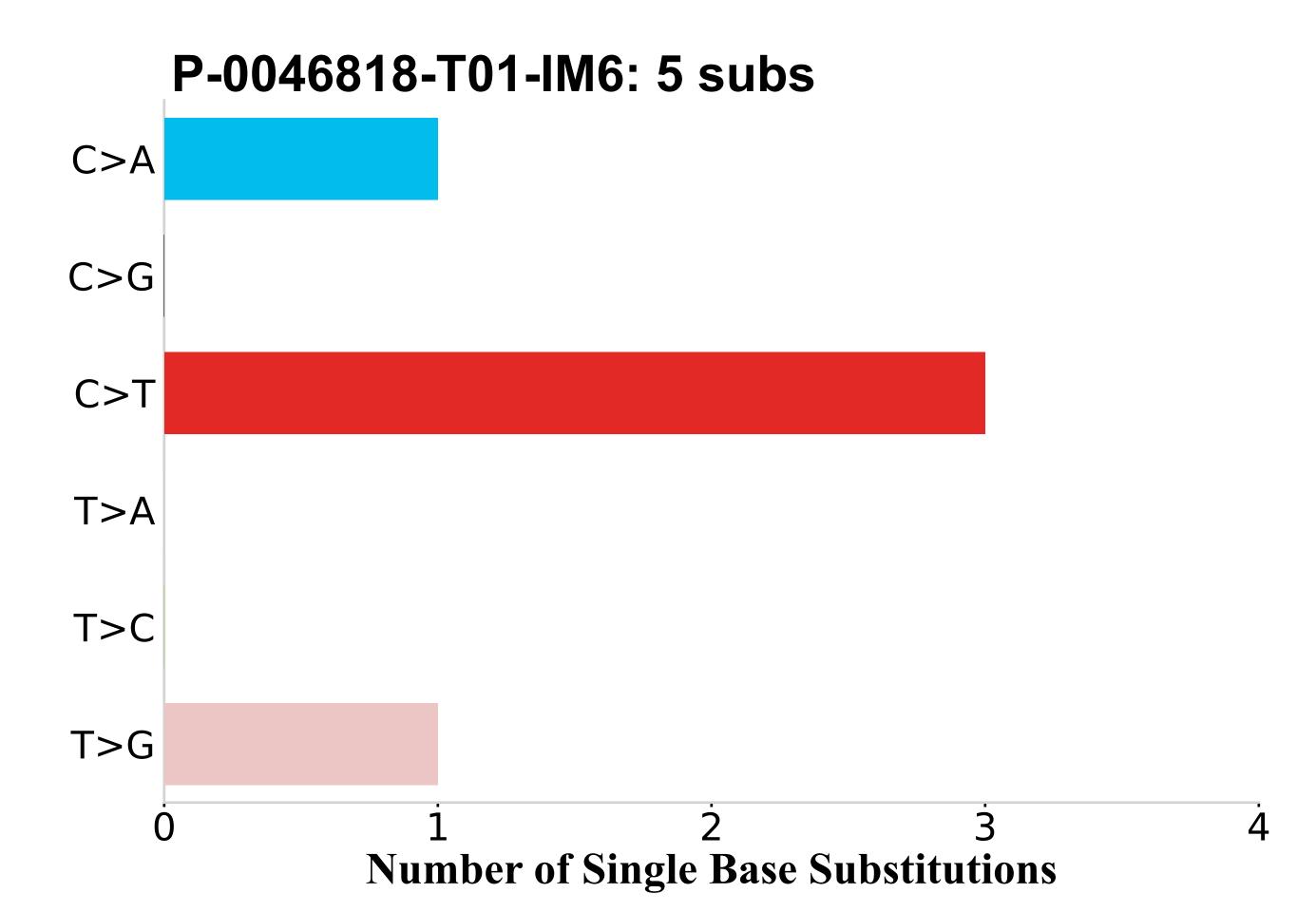


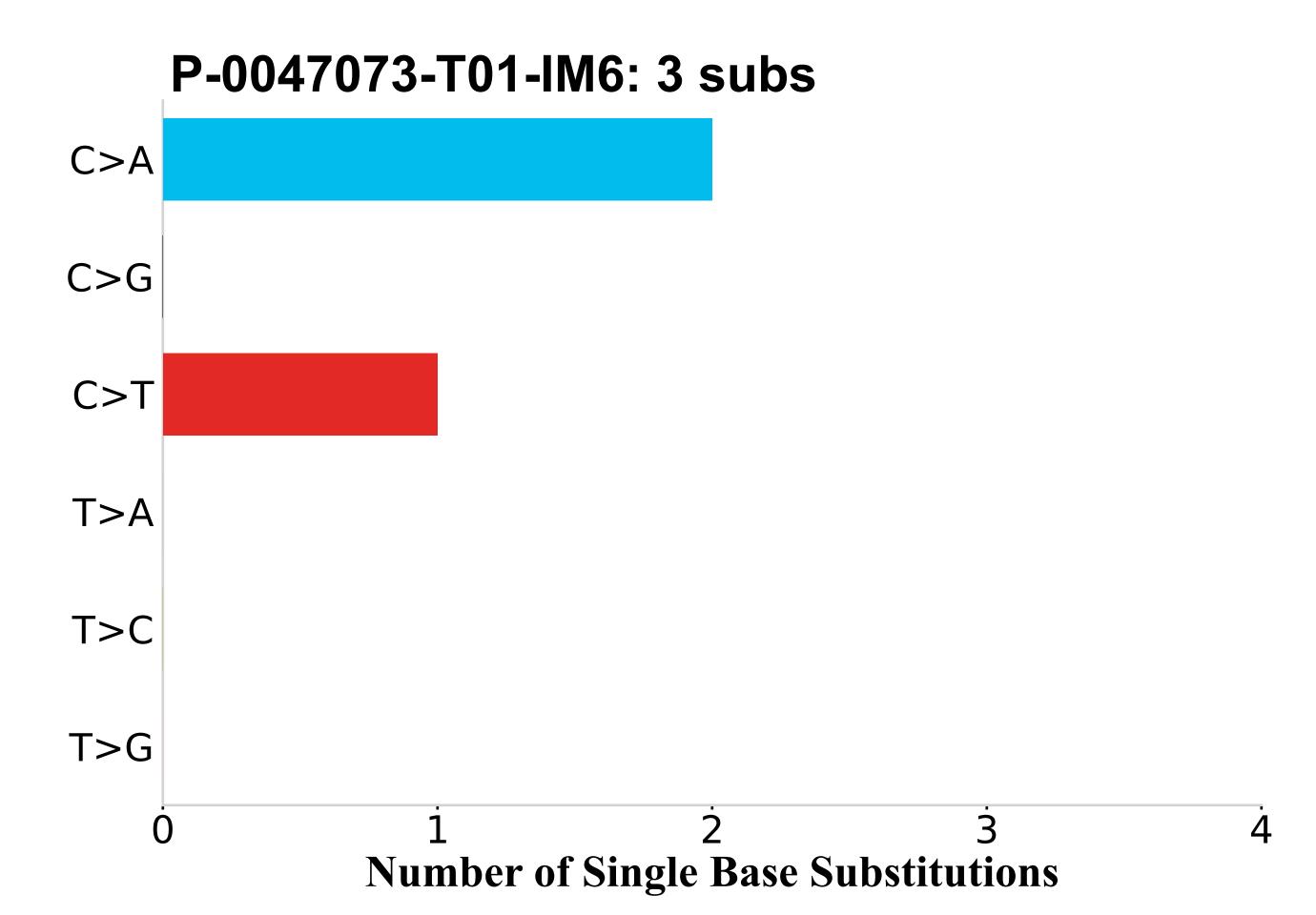


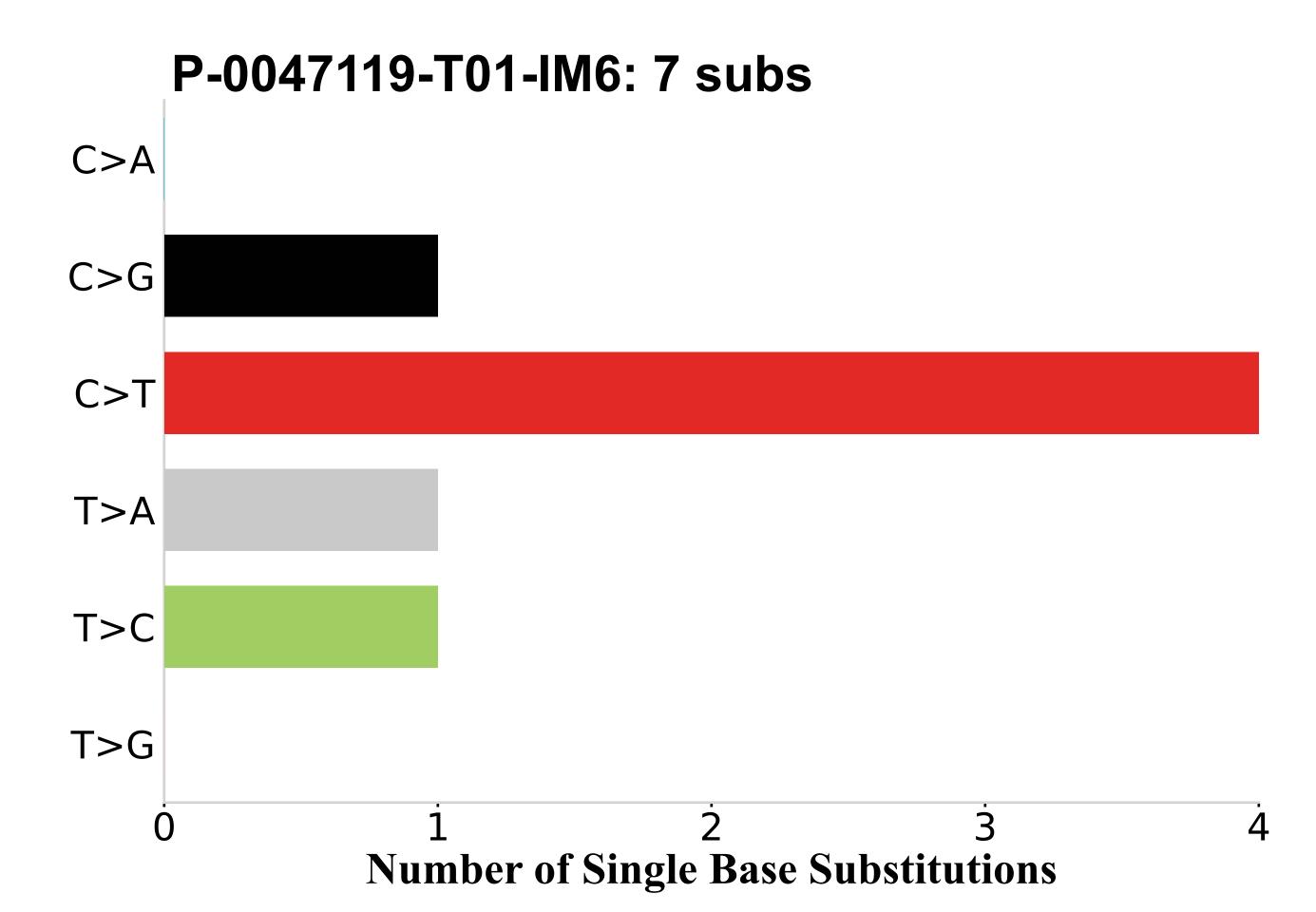


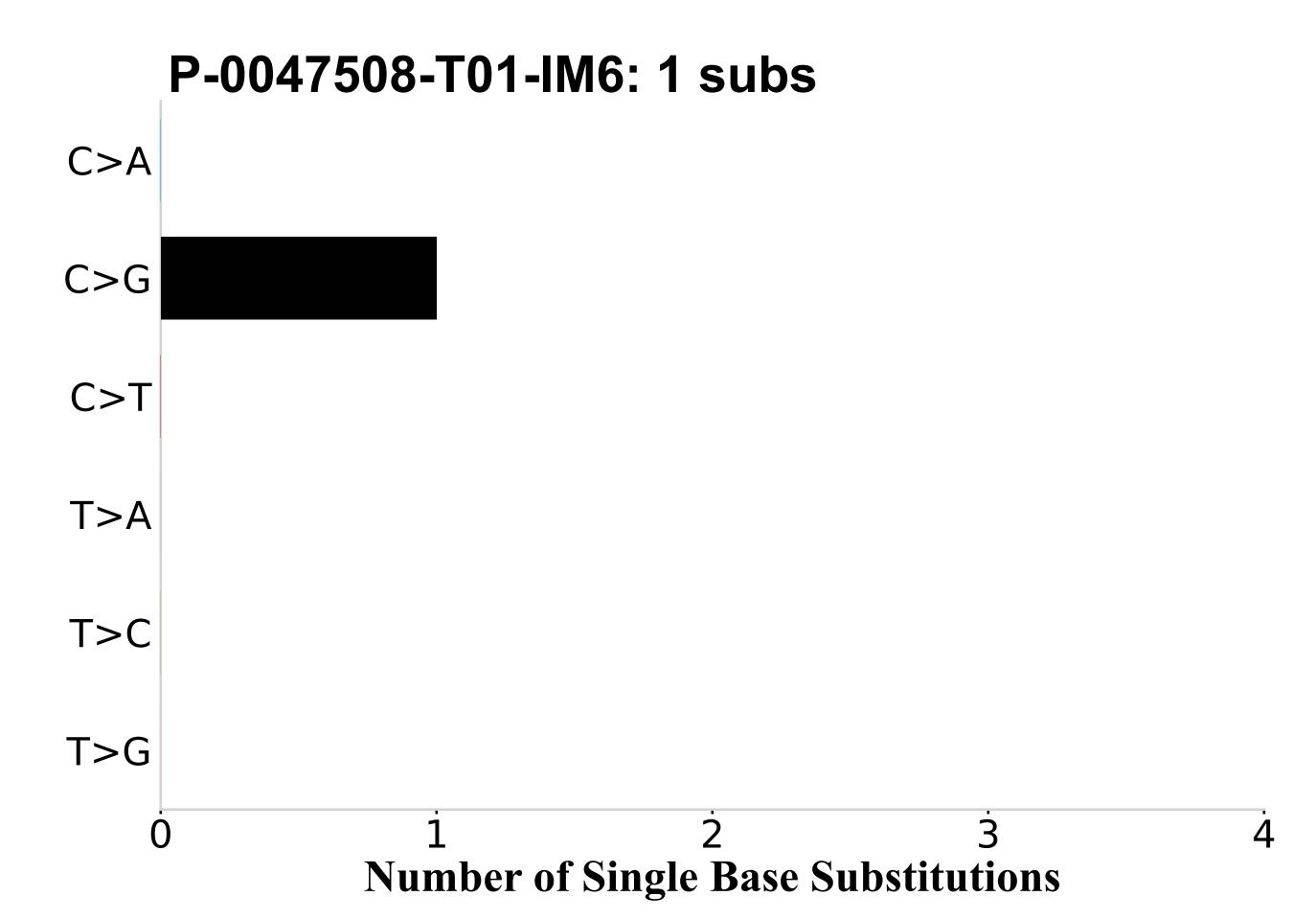
P-0046650-T01-IM6: 4 subs











P-0048569-T02-IM6: 5 subs C>A C>G C>T T>A T>C T>G **Number of Single Base Substitutions**

P-0048802-T01-IM6: 3 subs C>A C>G C>T T>A T>C T>G **Number of Single Base Substitutions**



