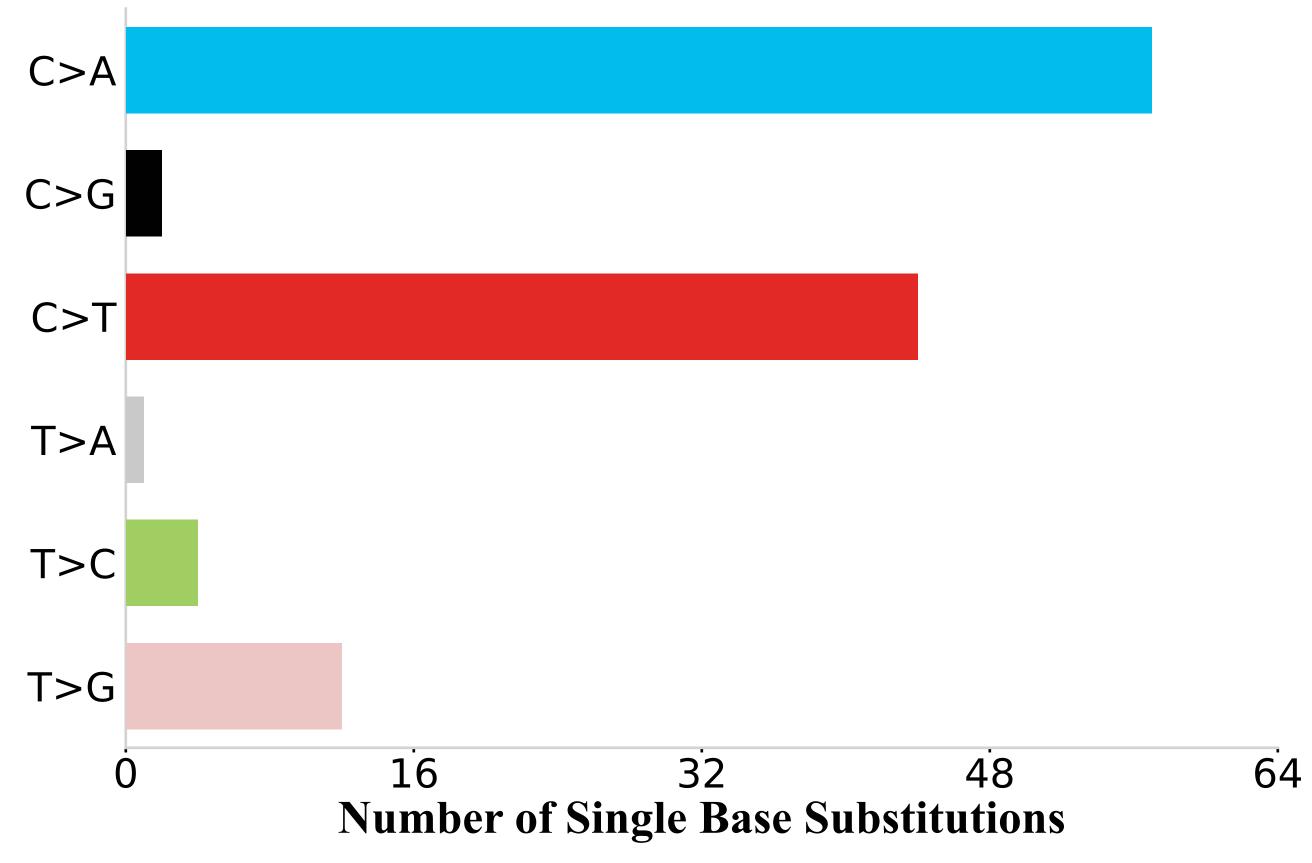
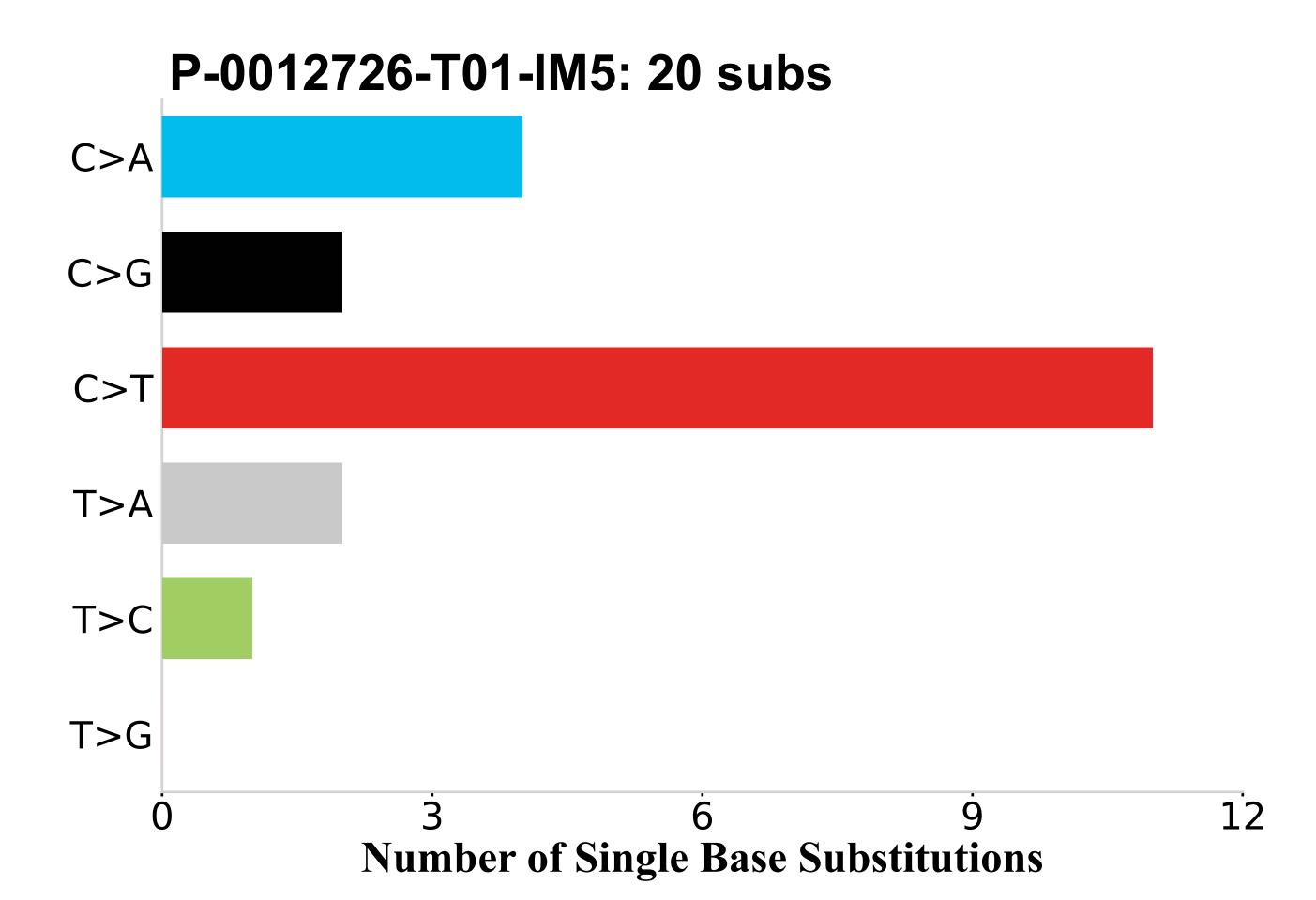
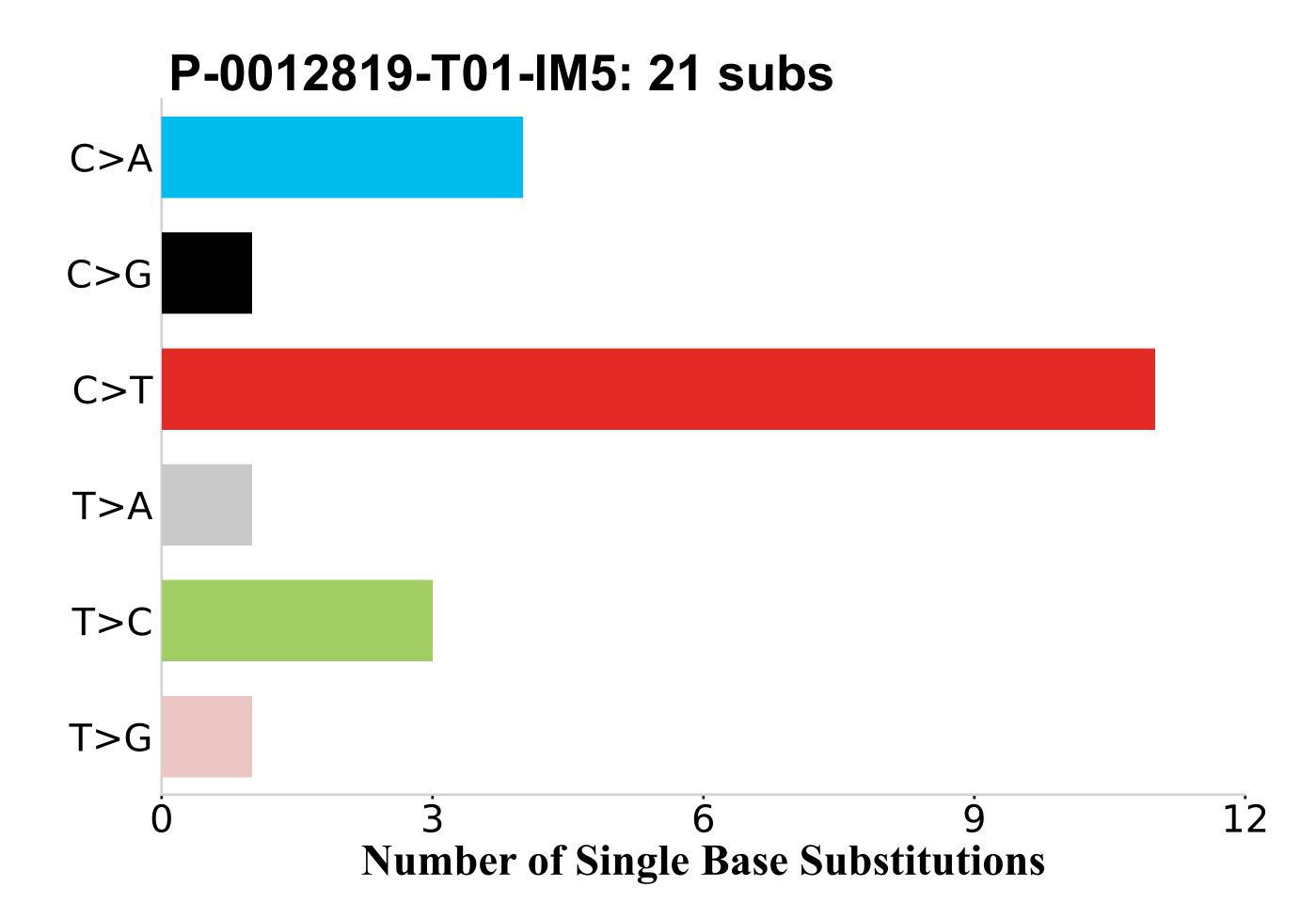
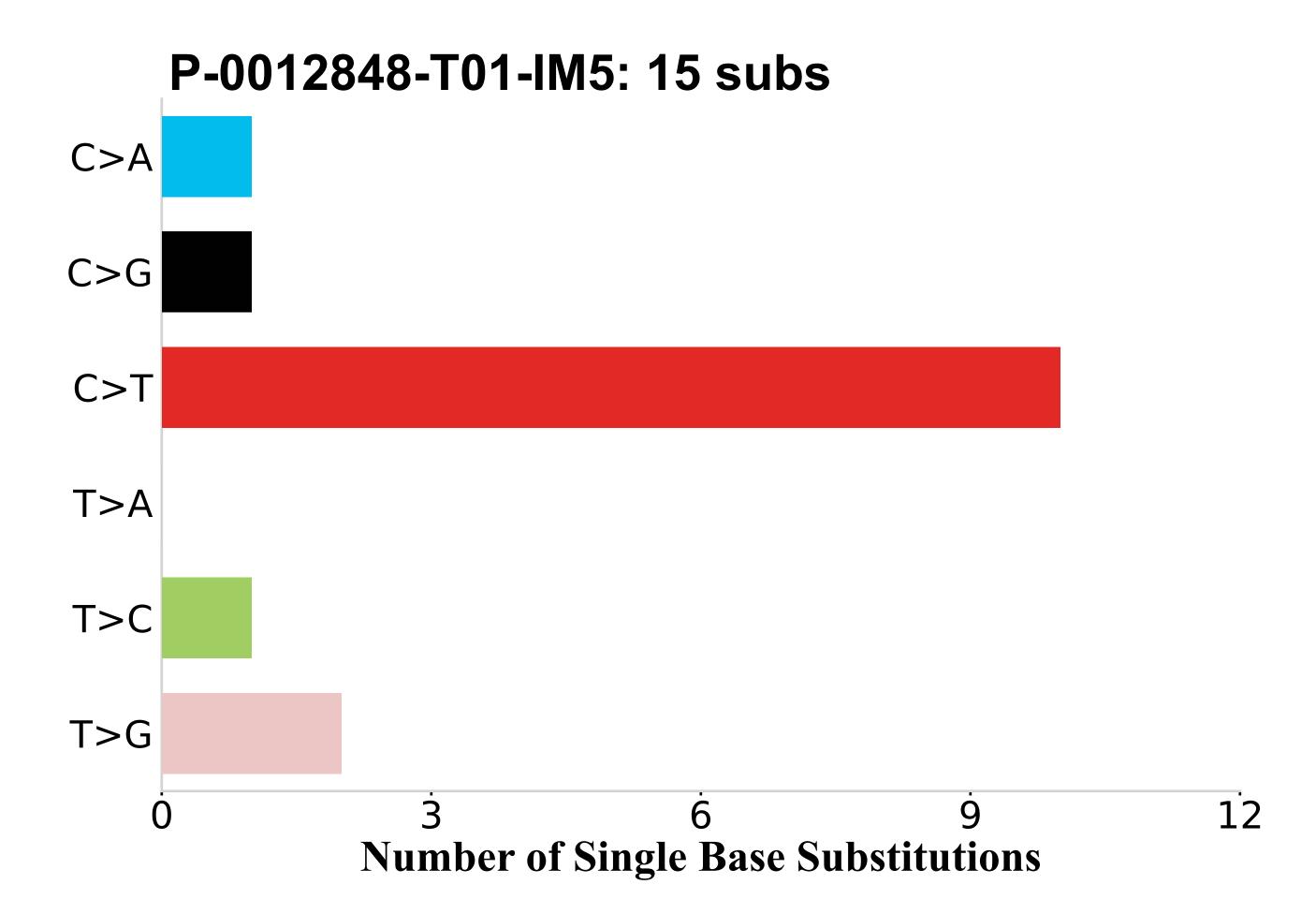


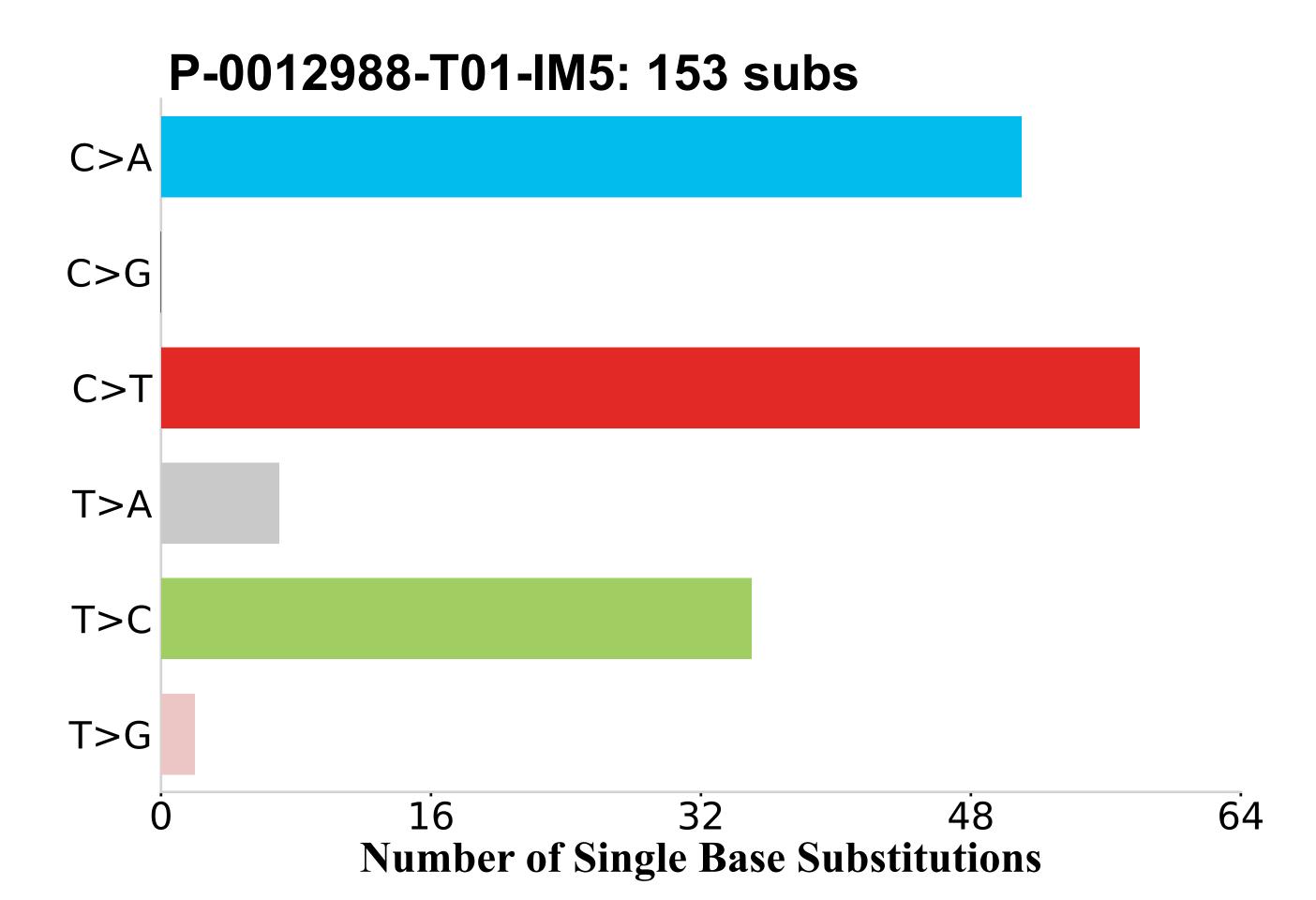
P-0012670-T01-IM5: 120 subs

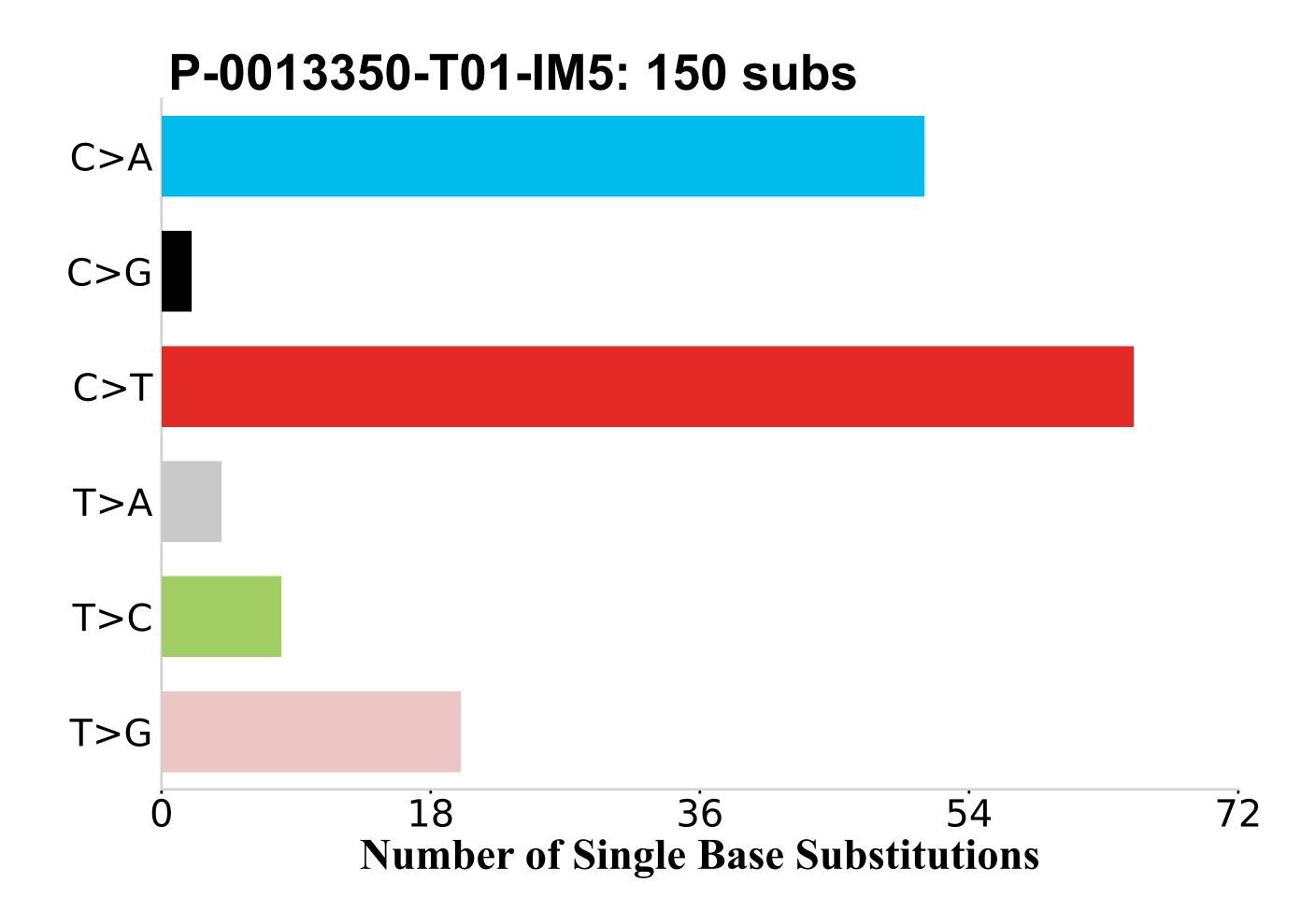


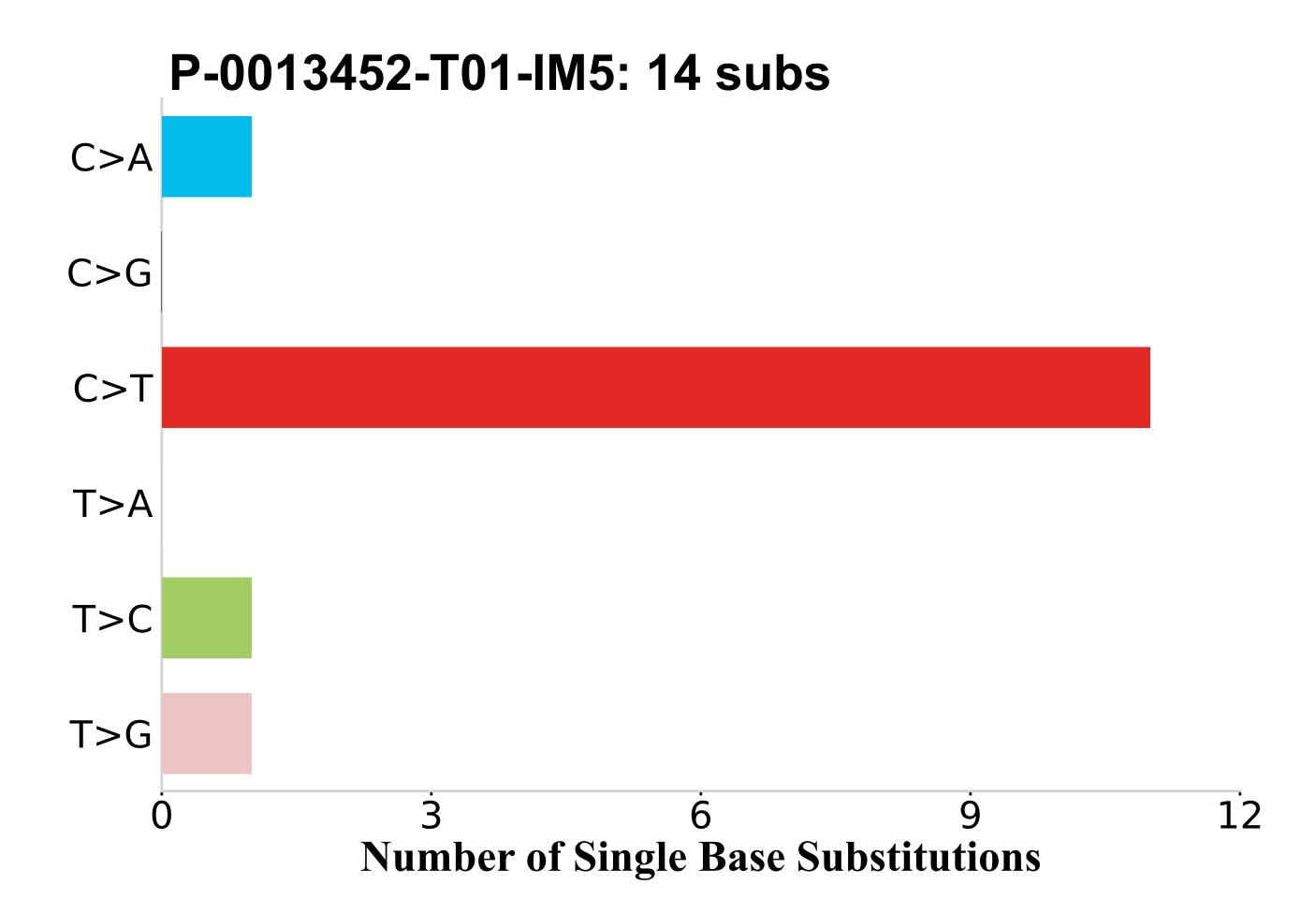


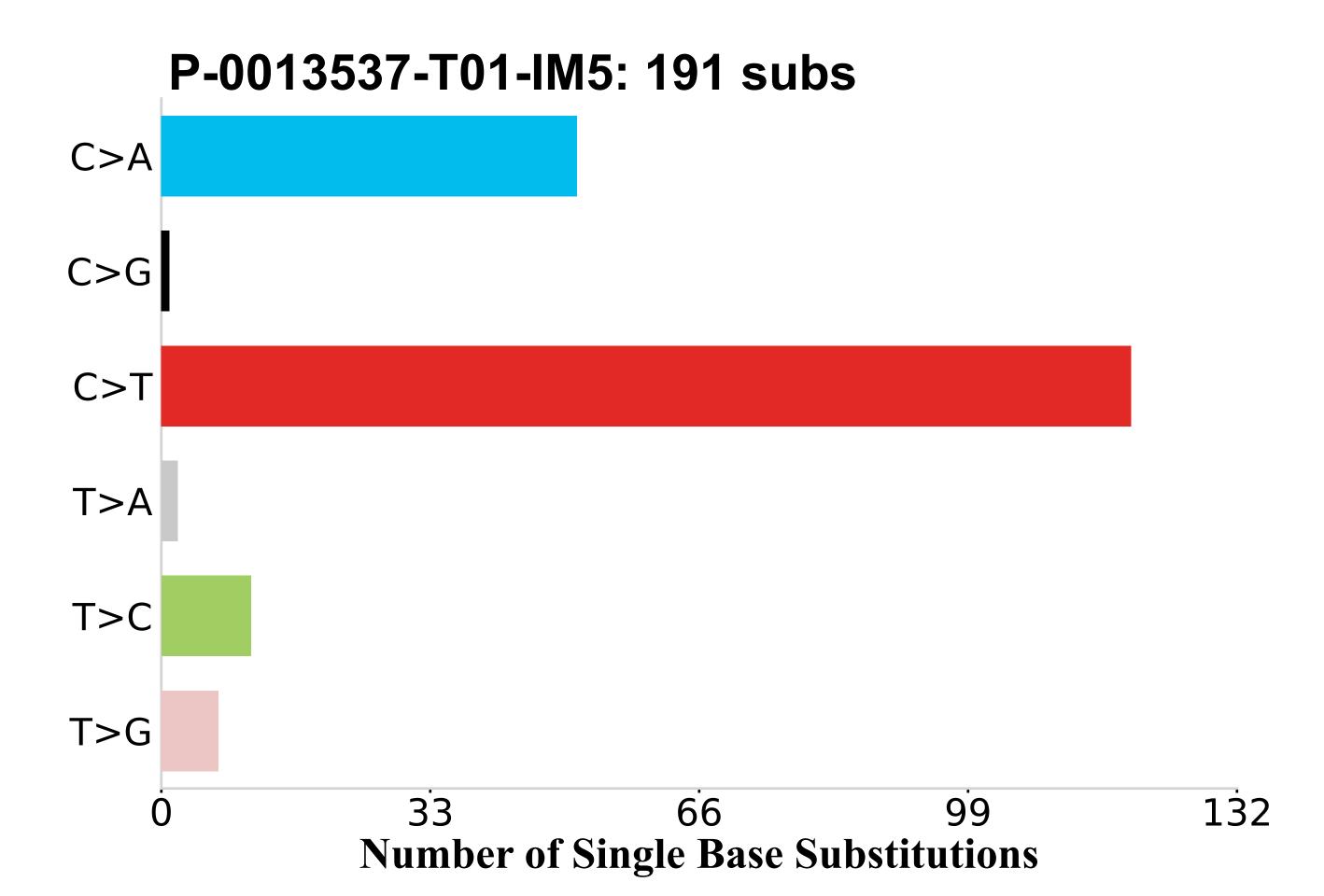


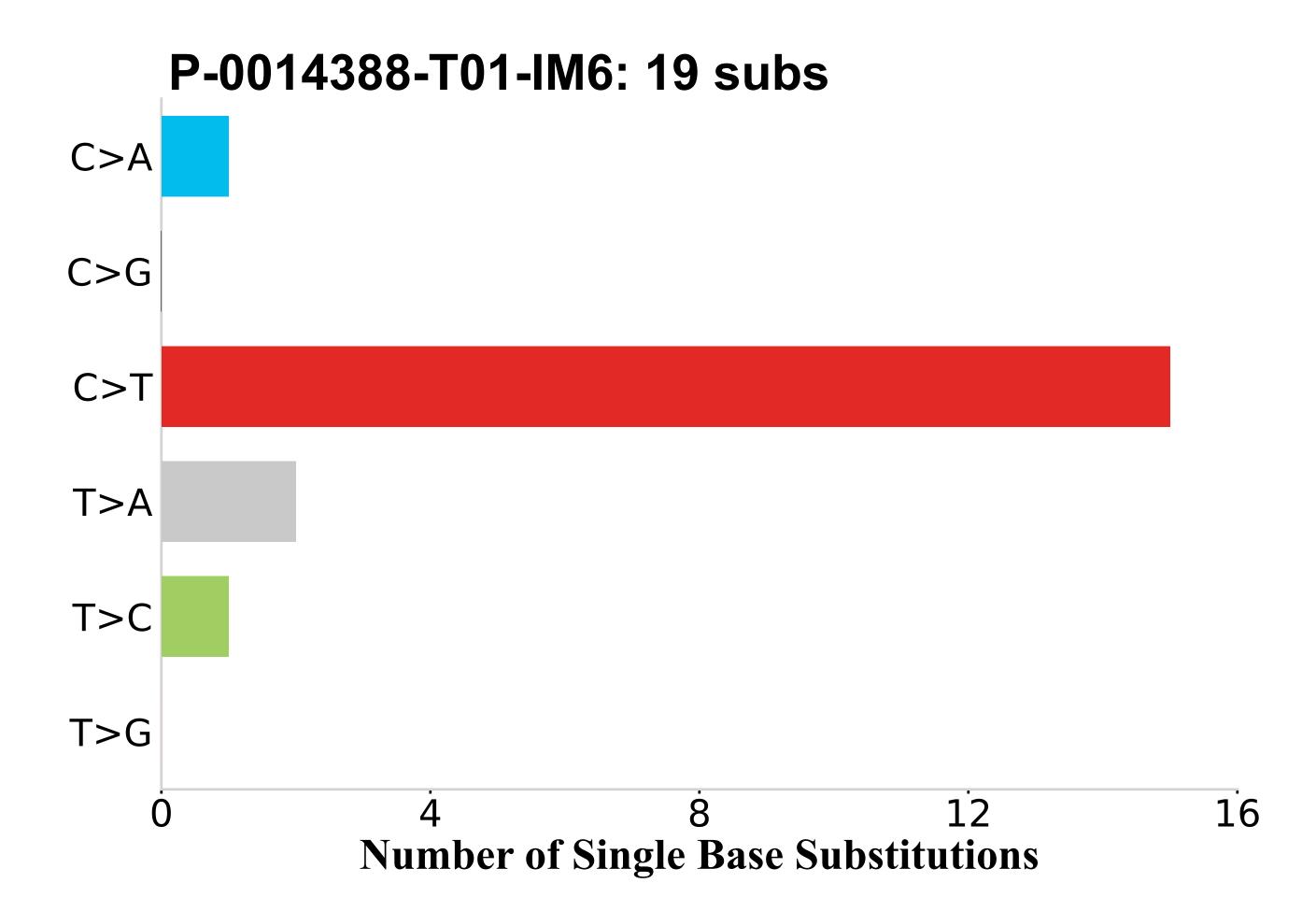


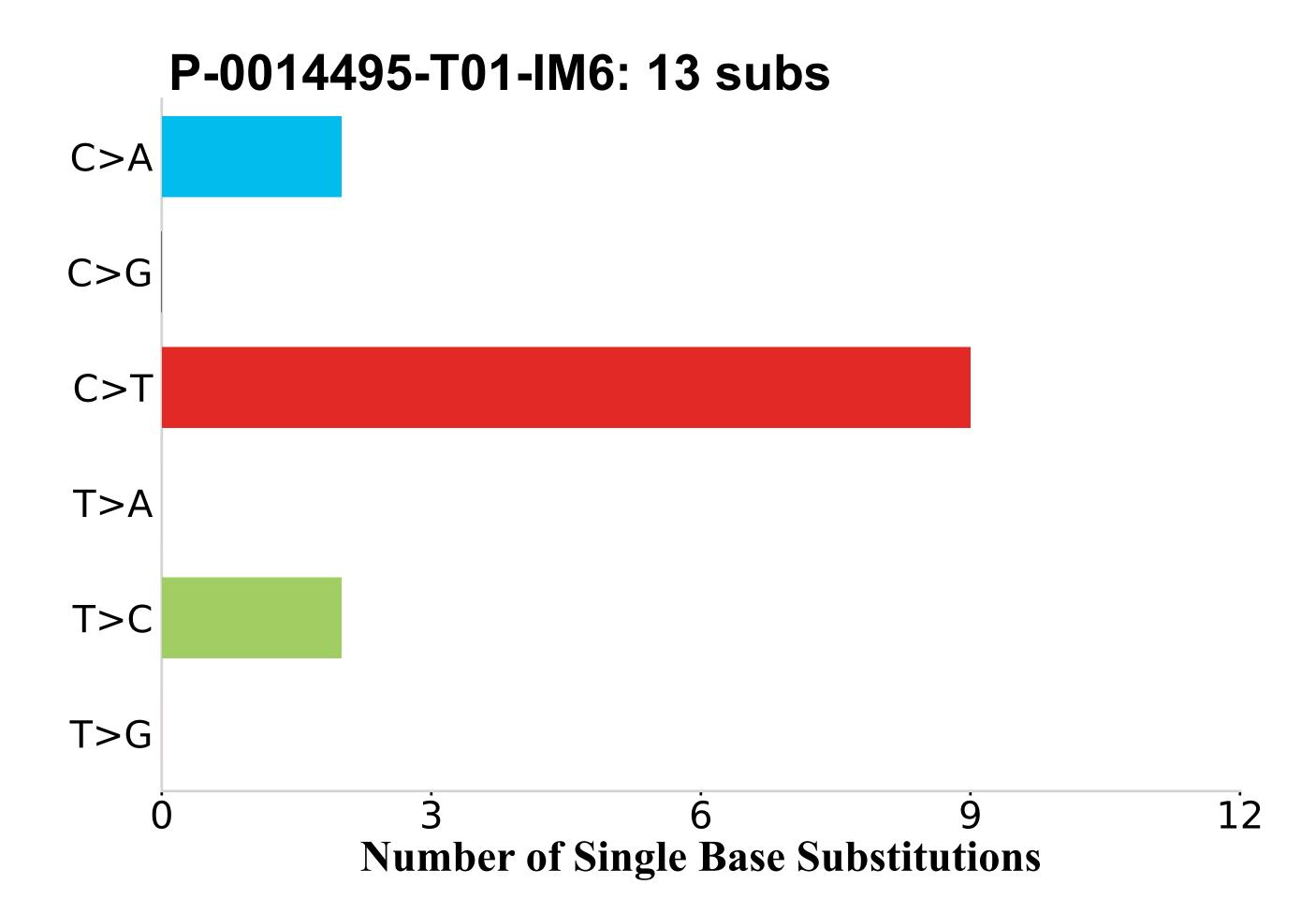


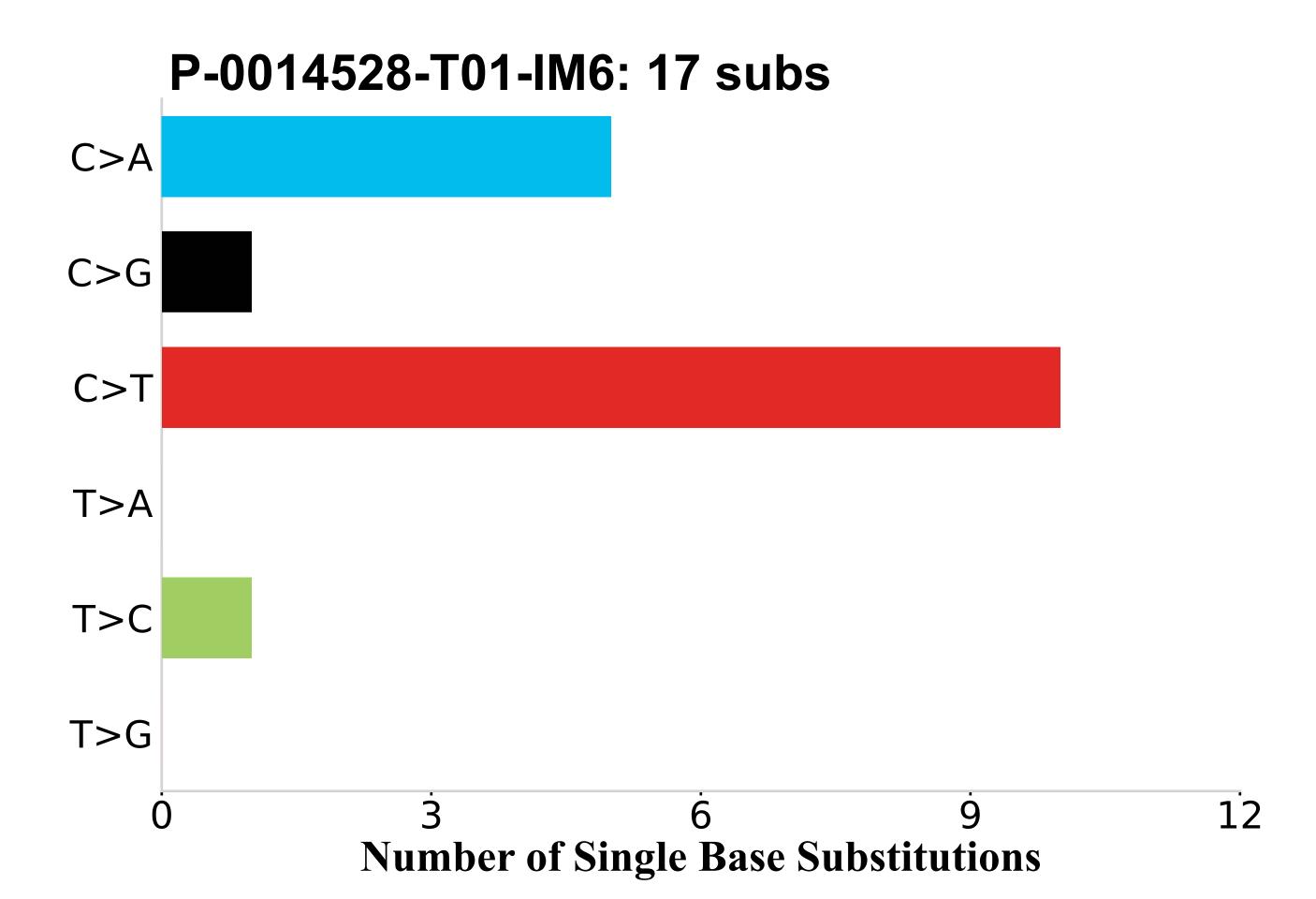


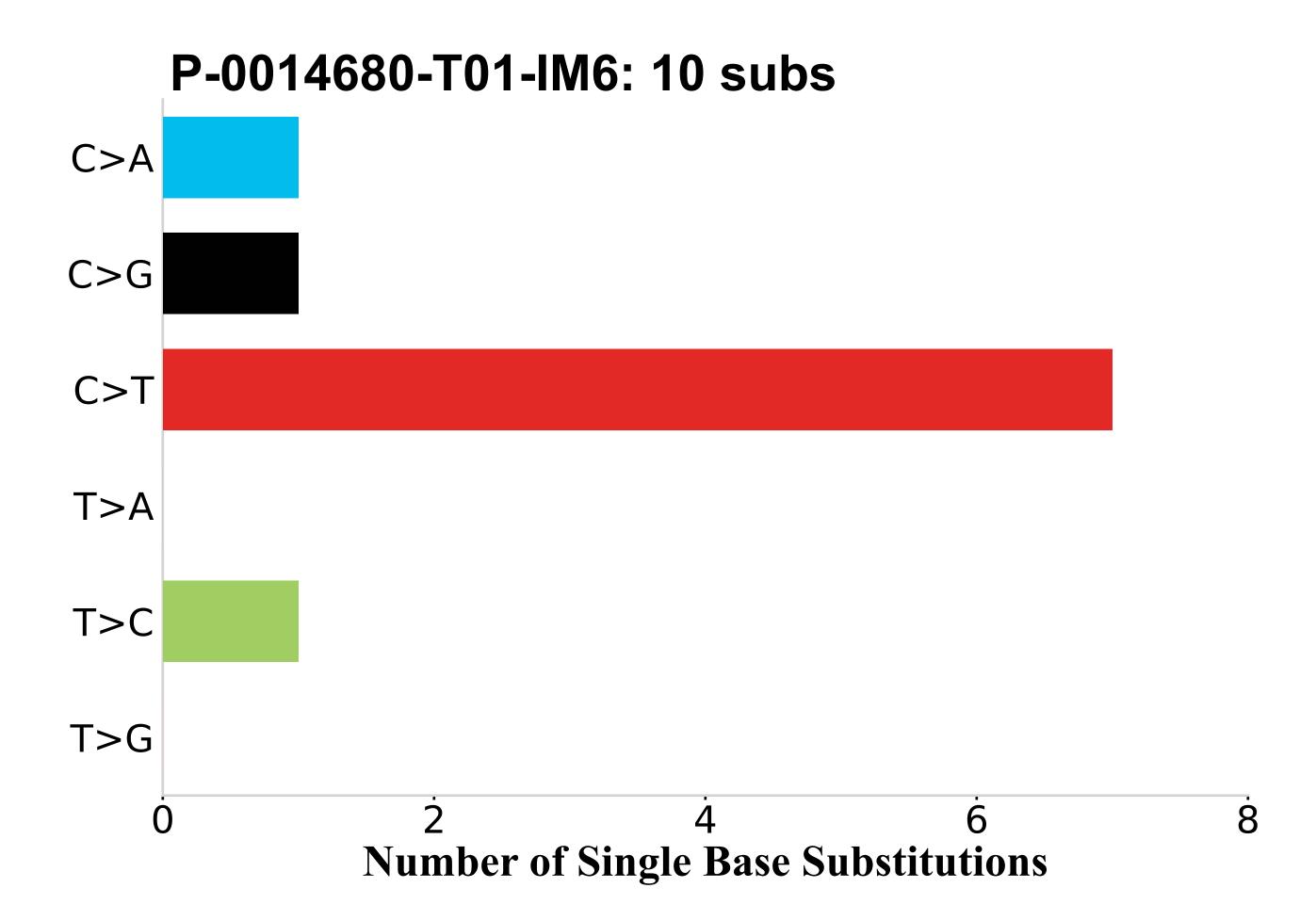


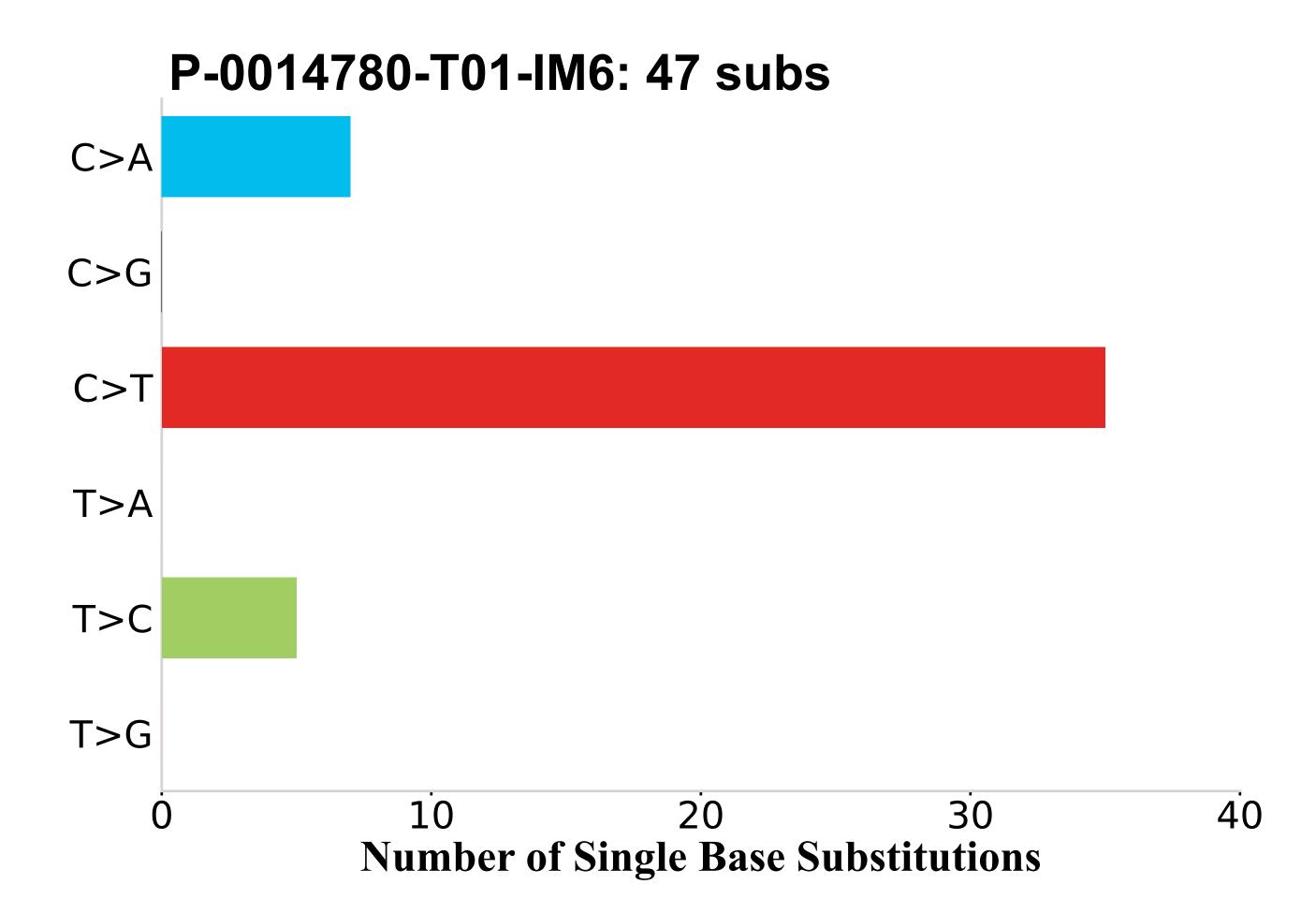


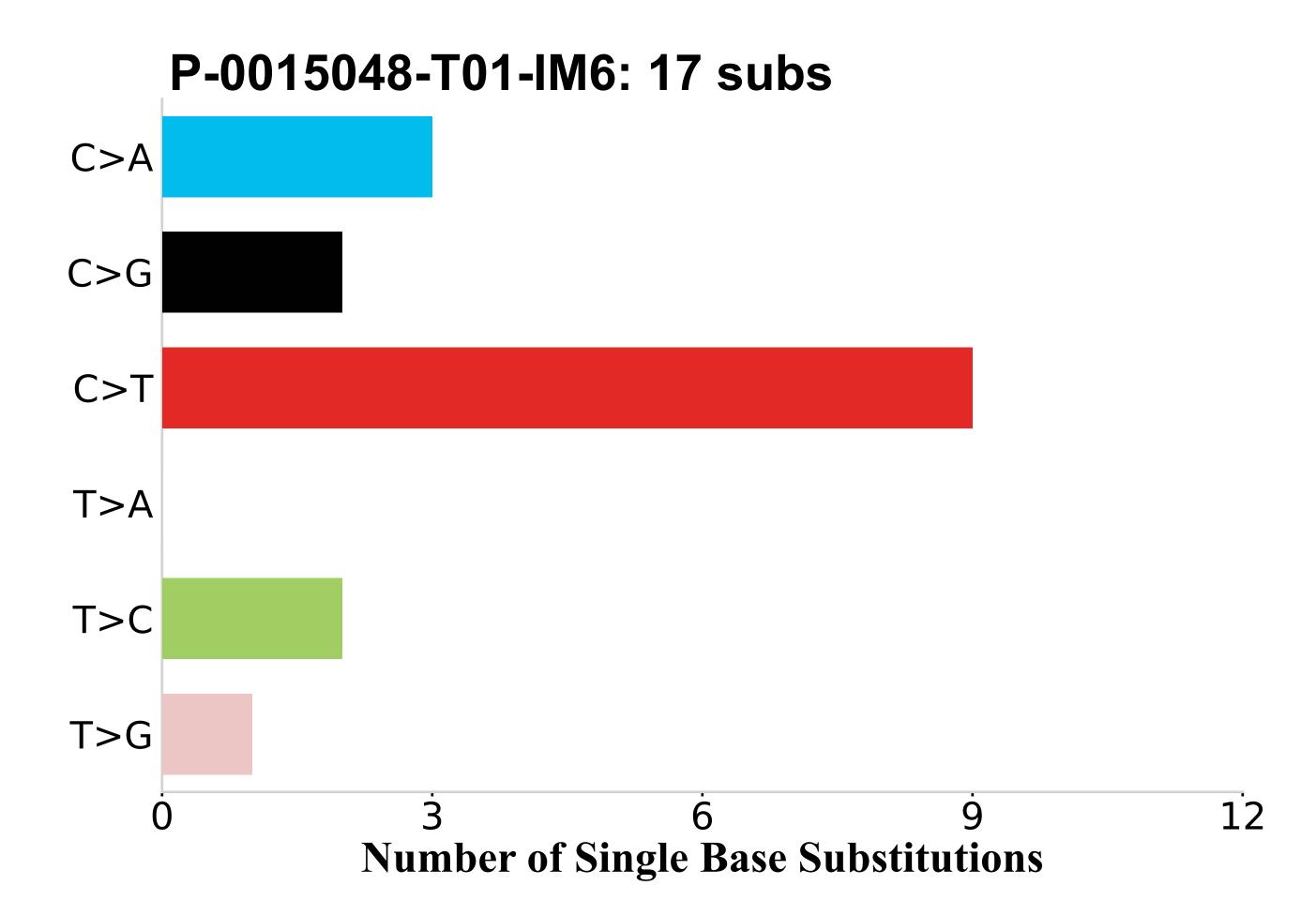


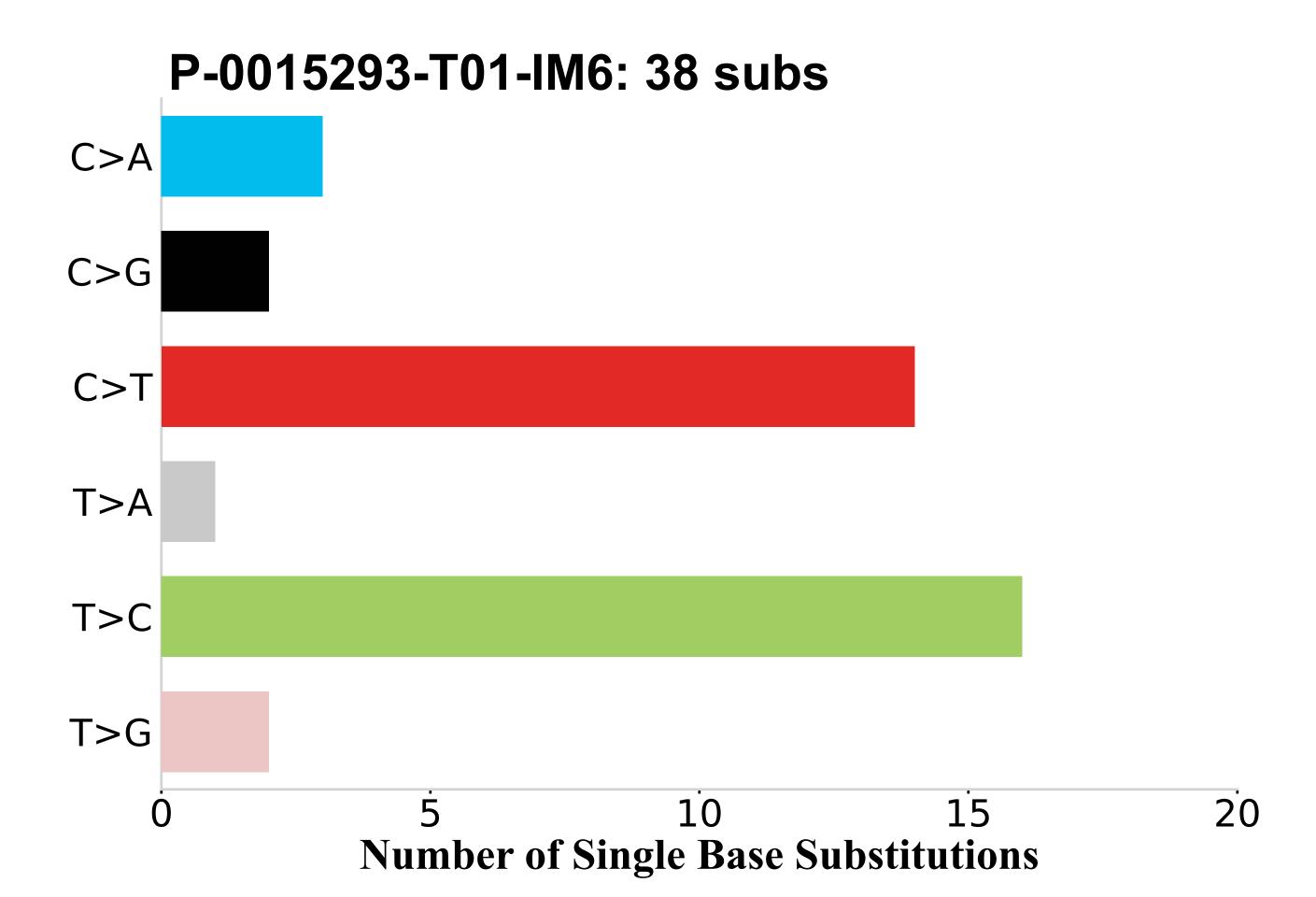


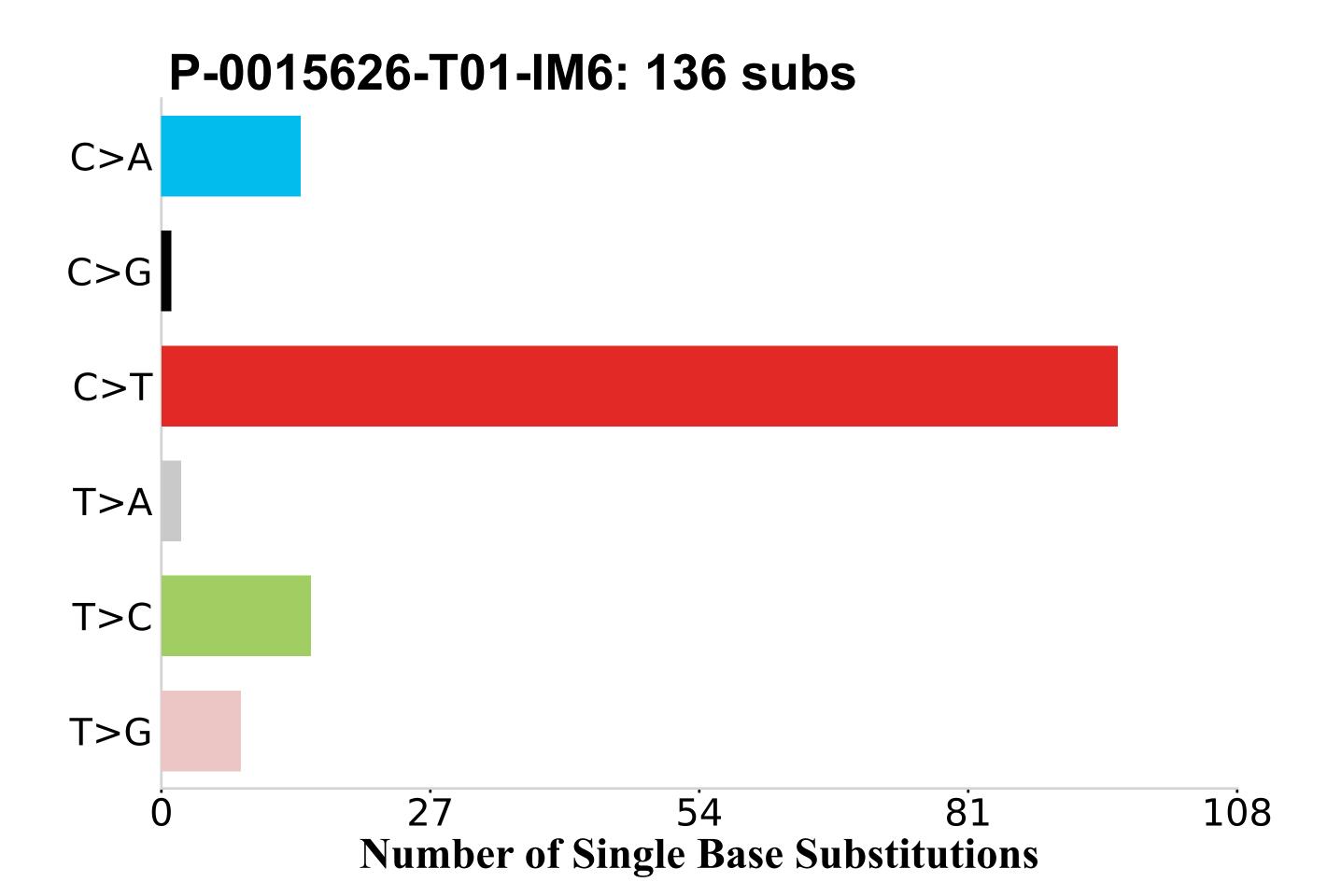


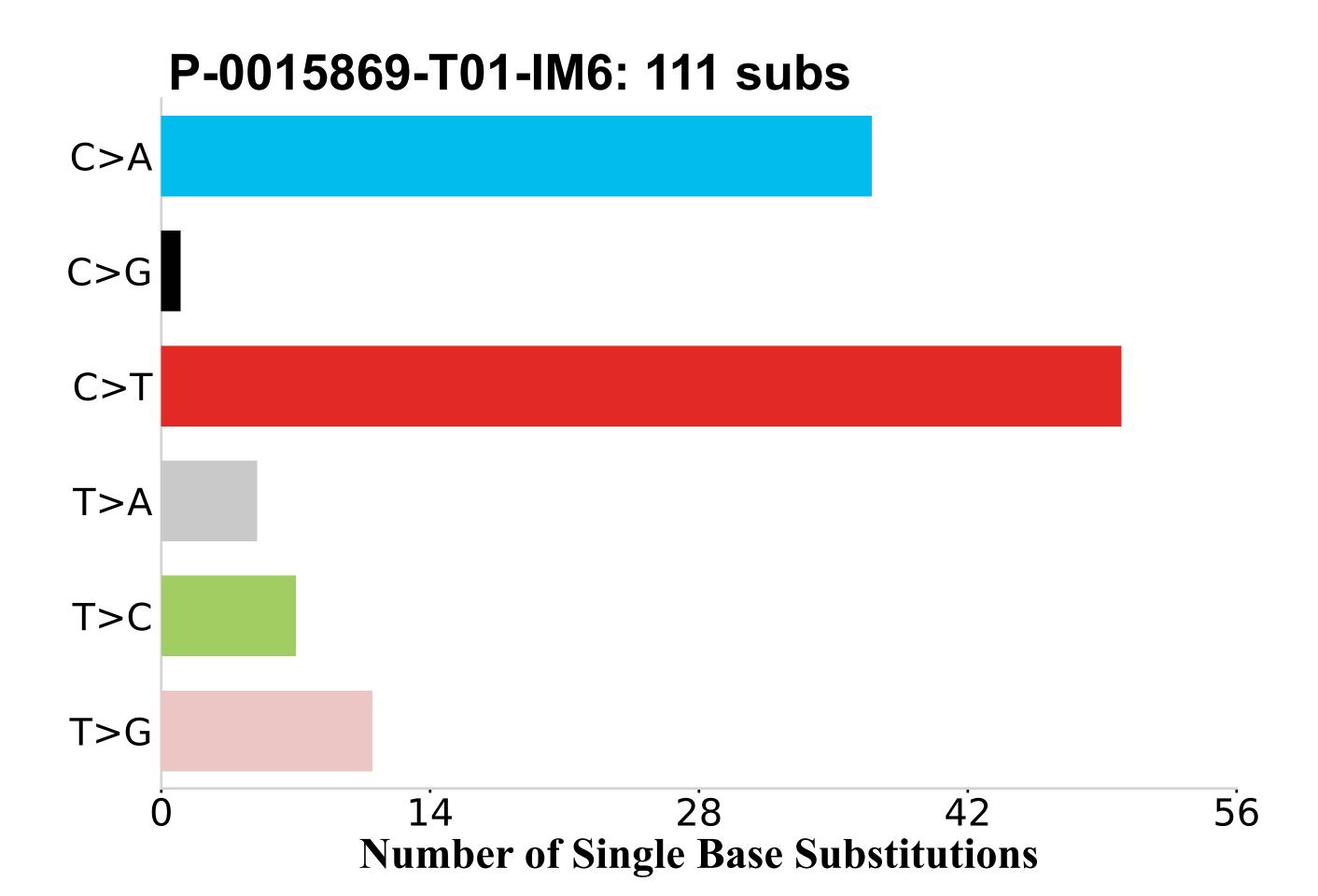


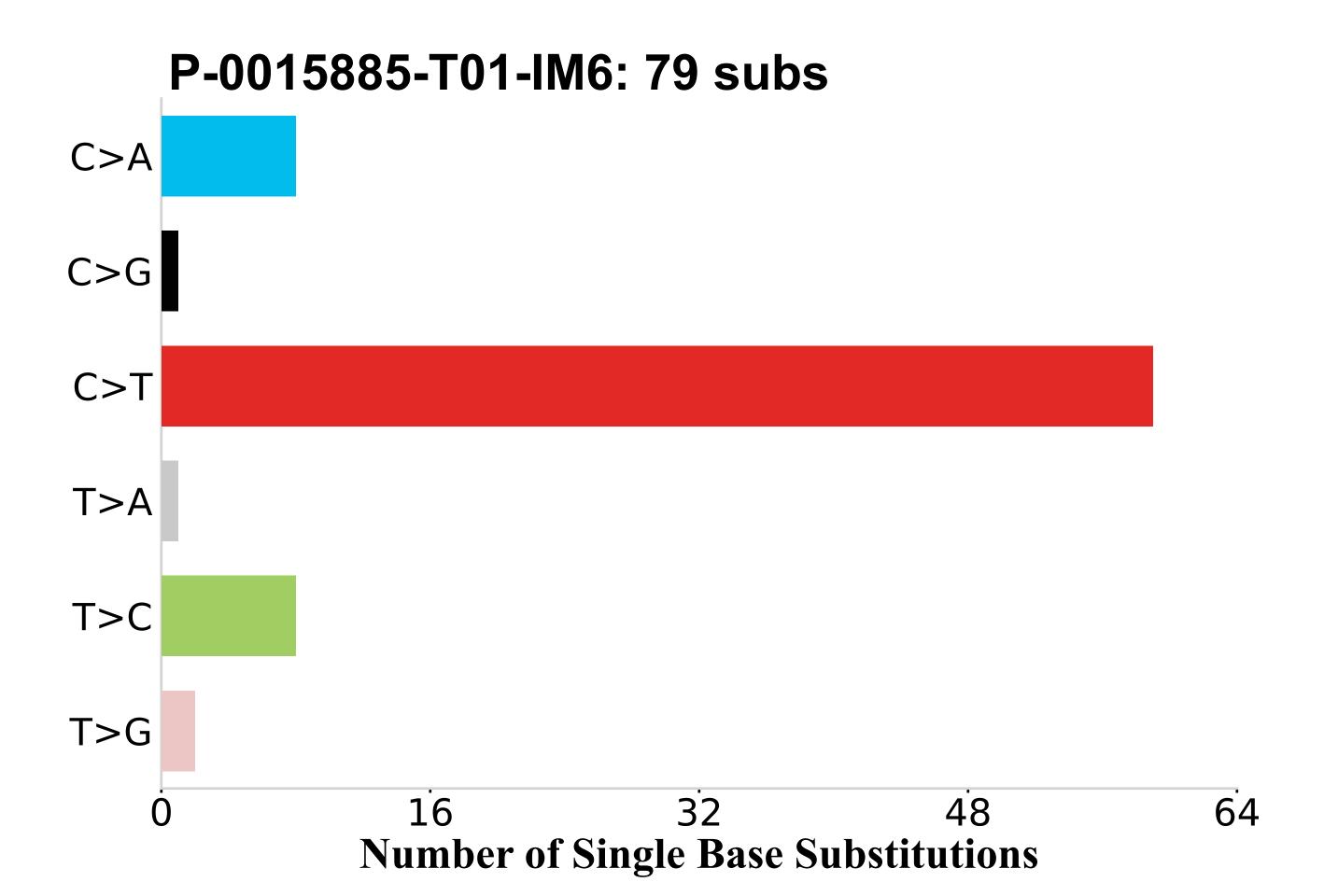


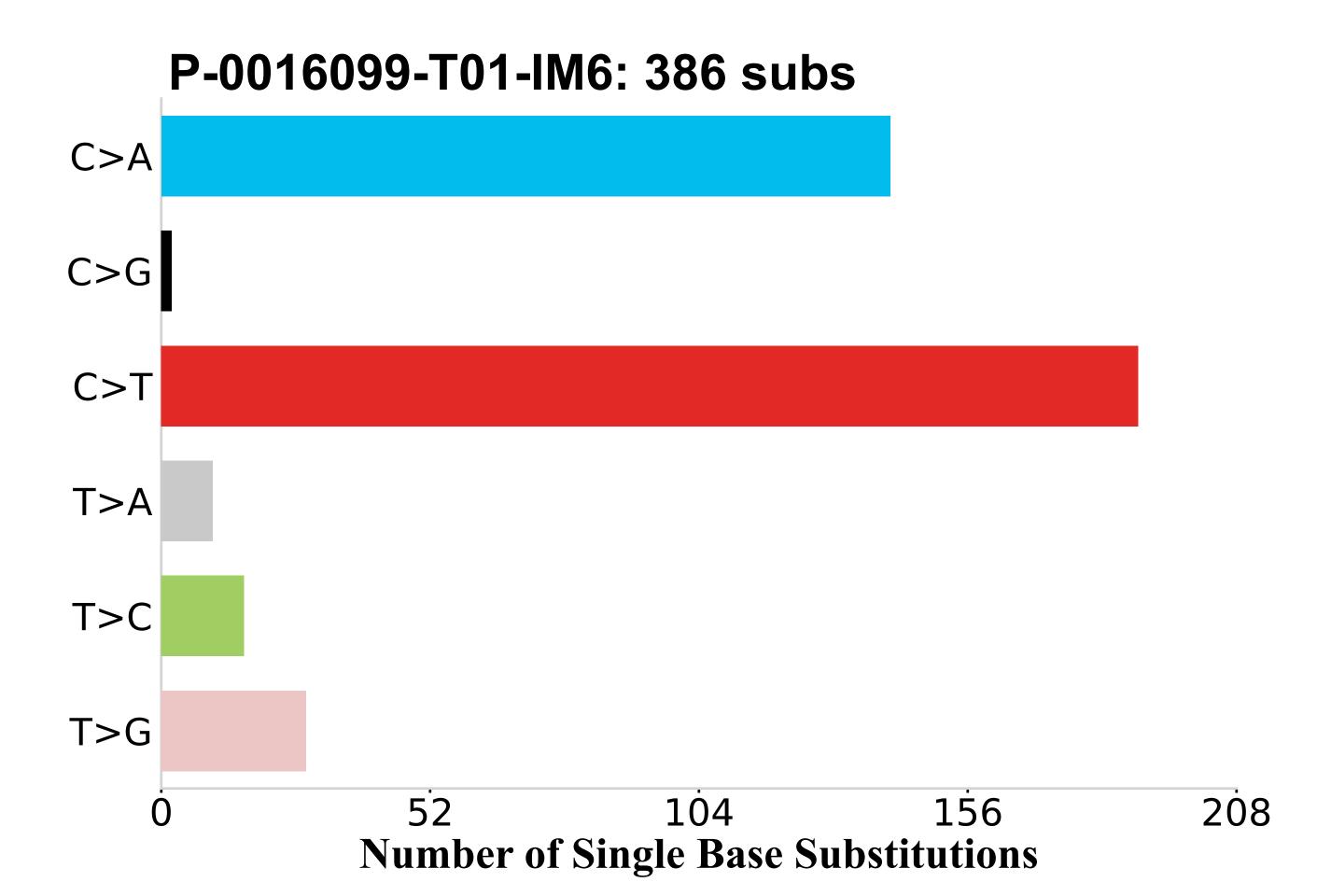


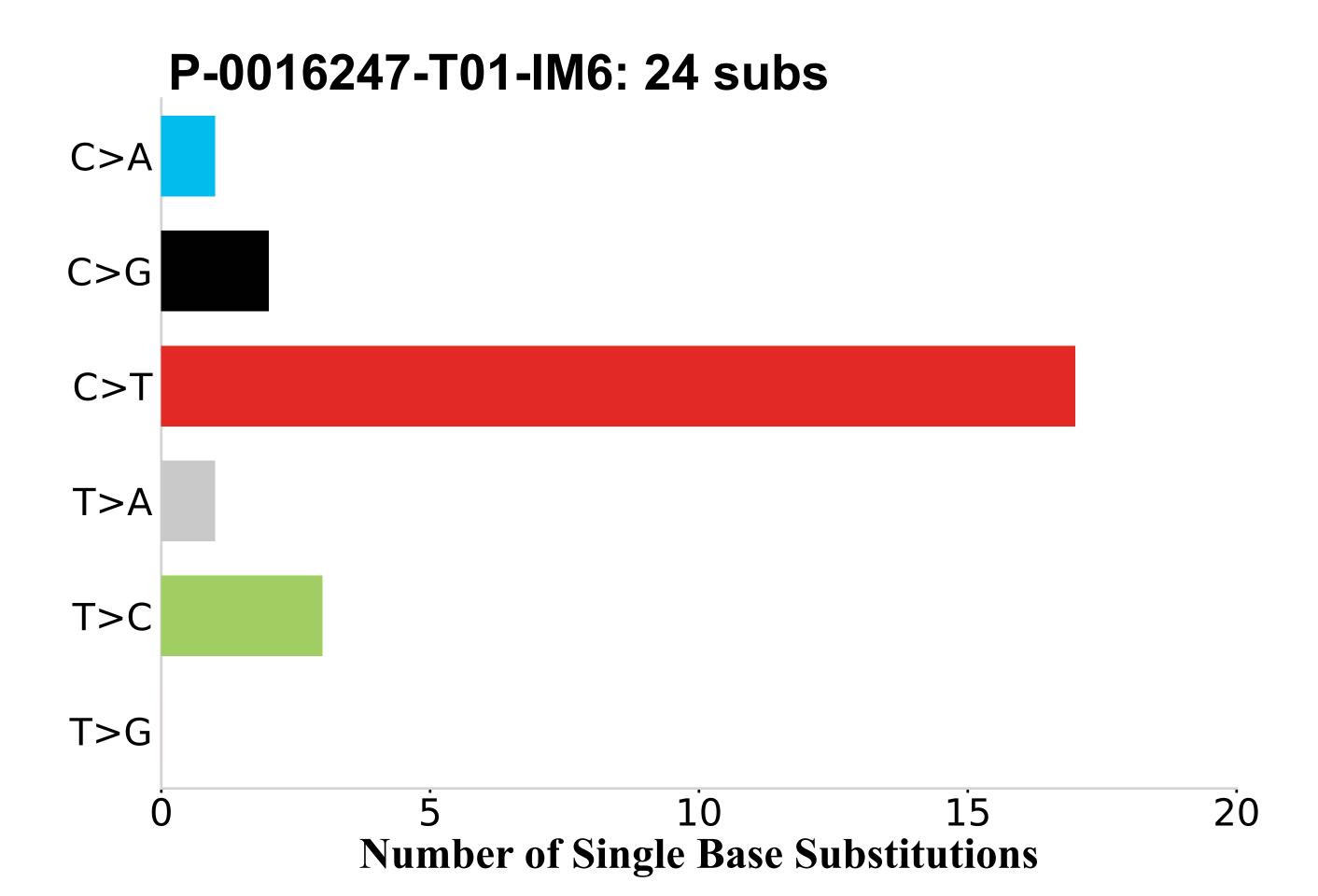


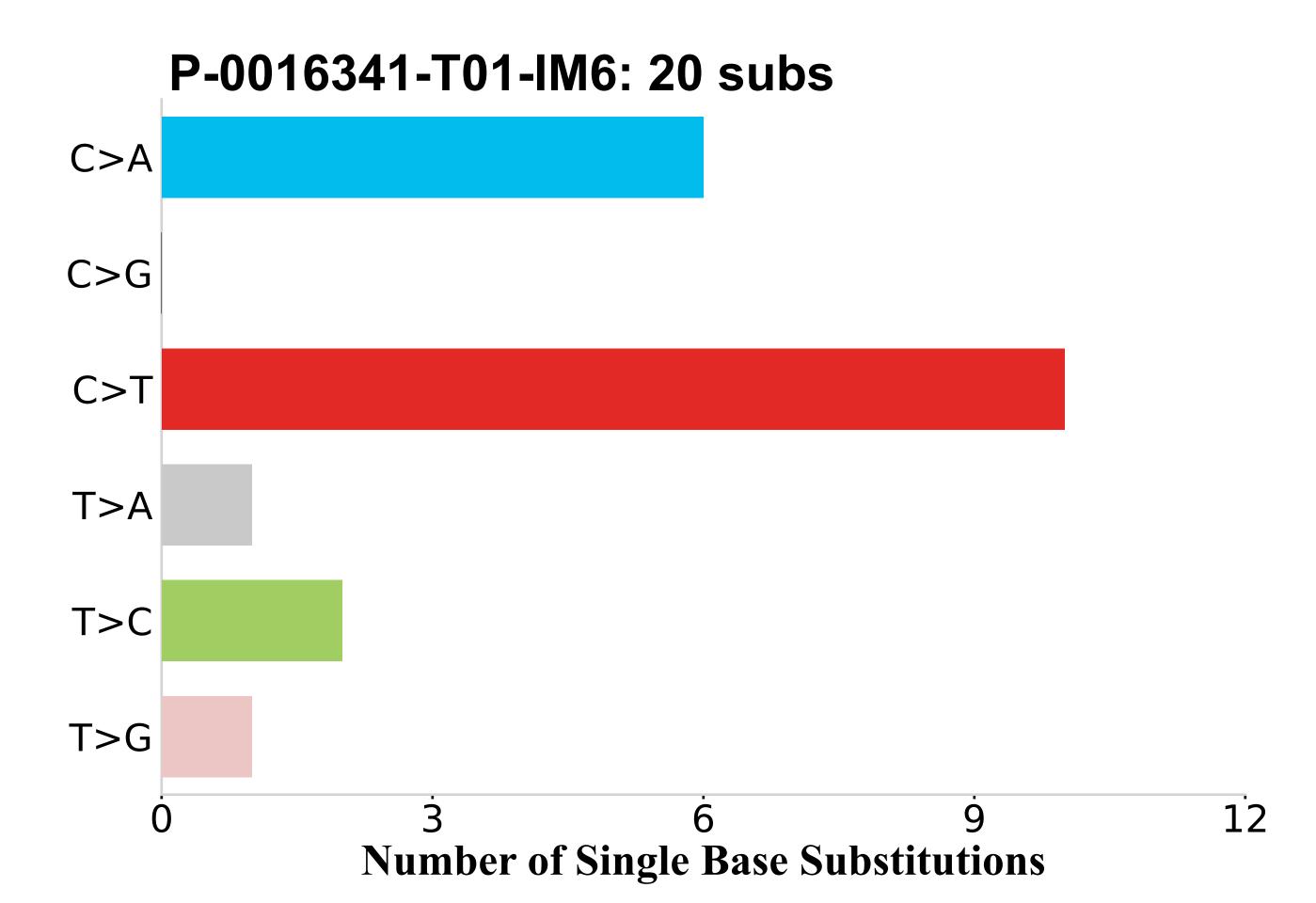


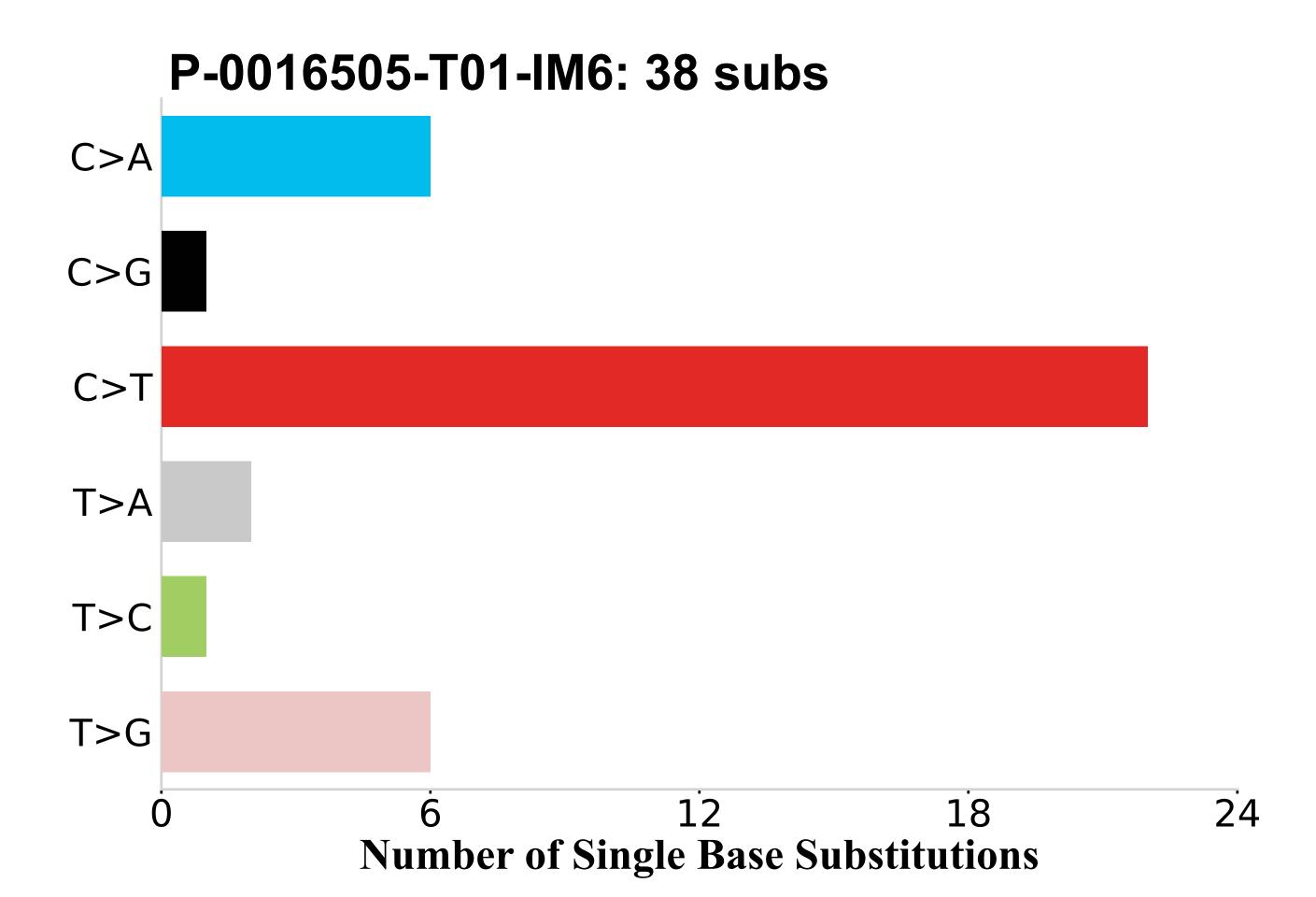


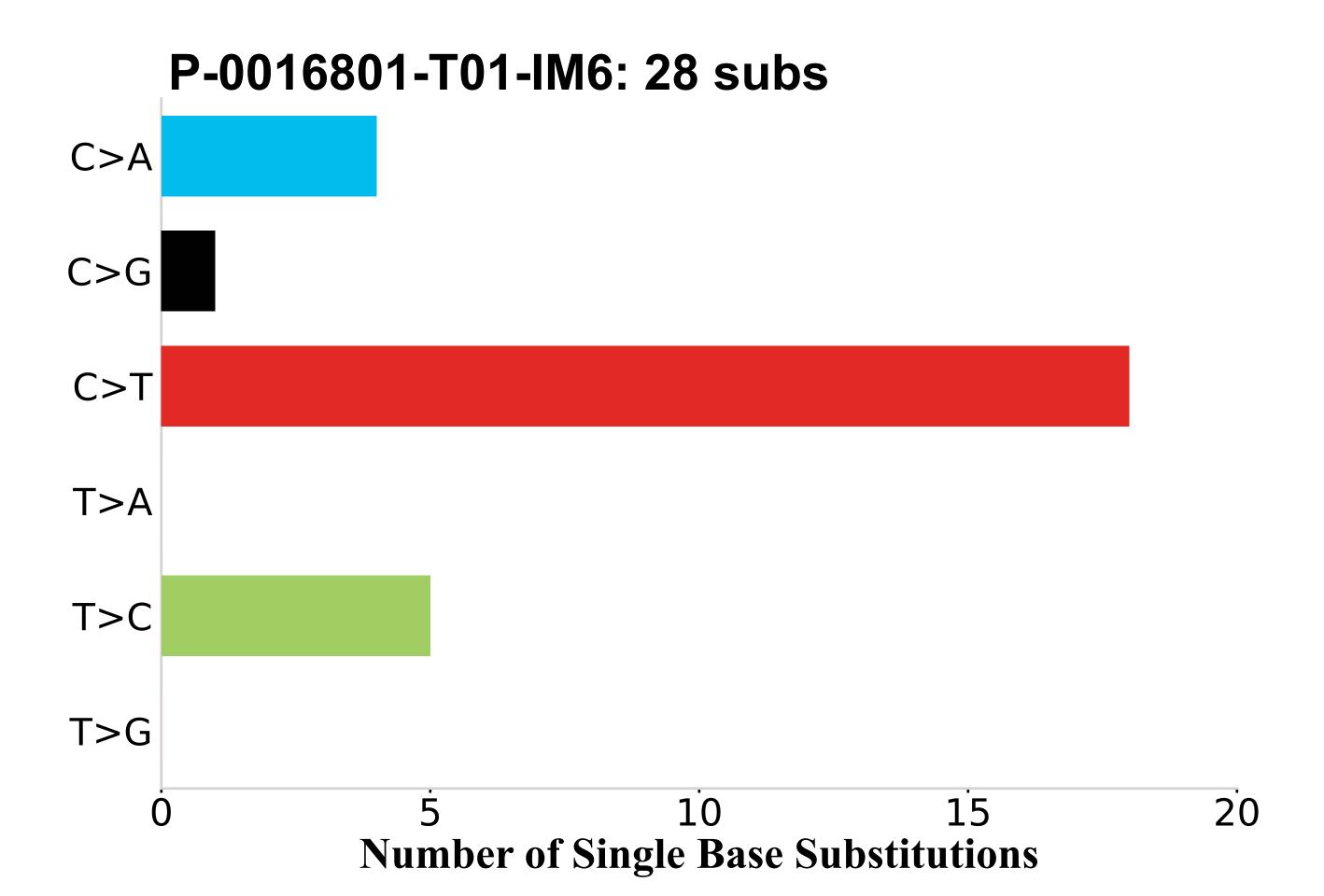






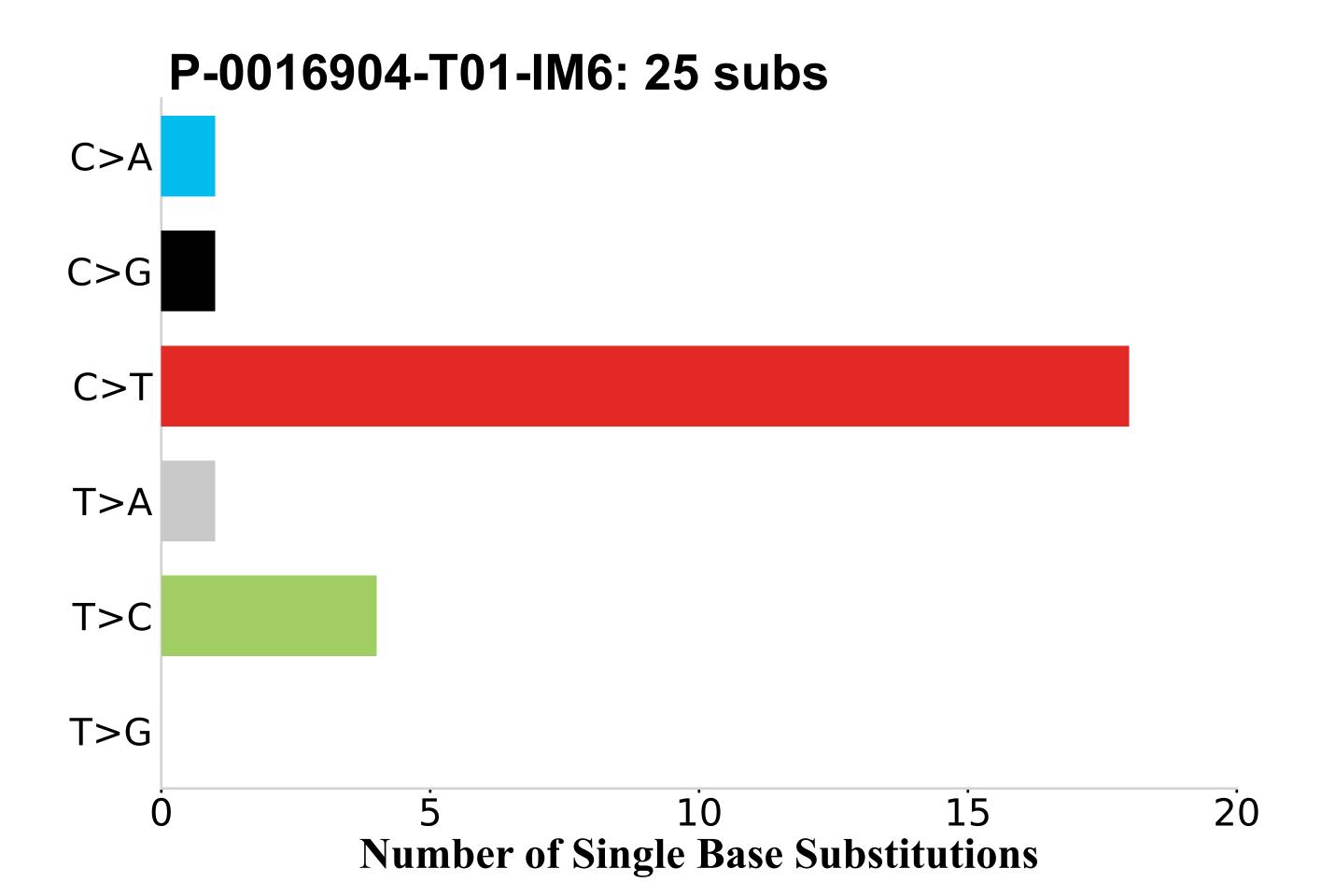


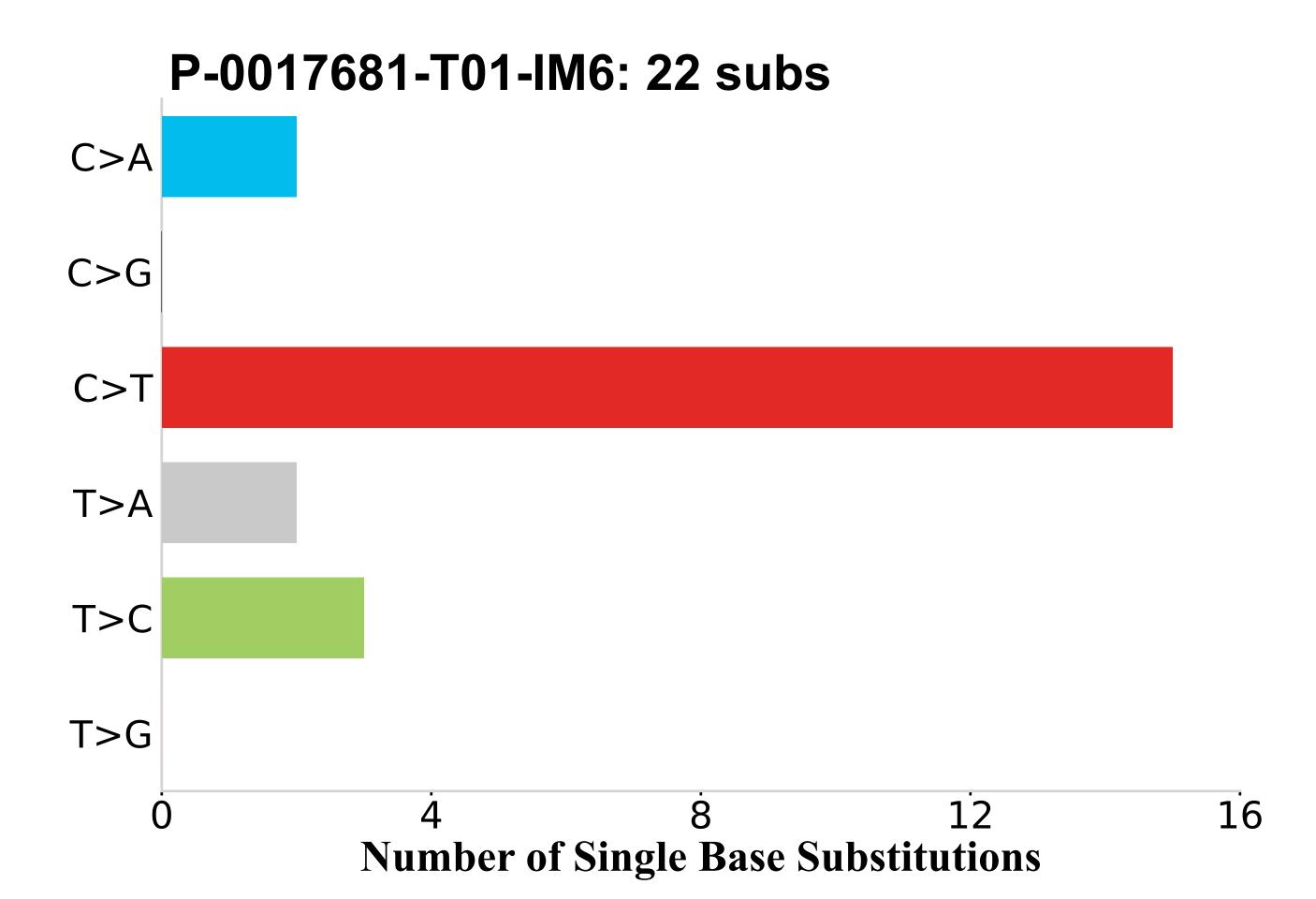


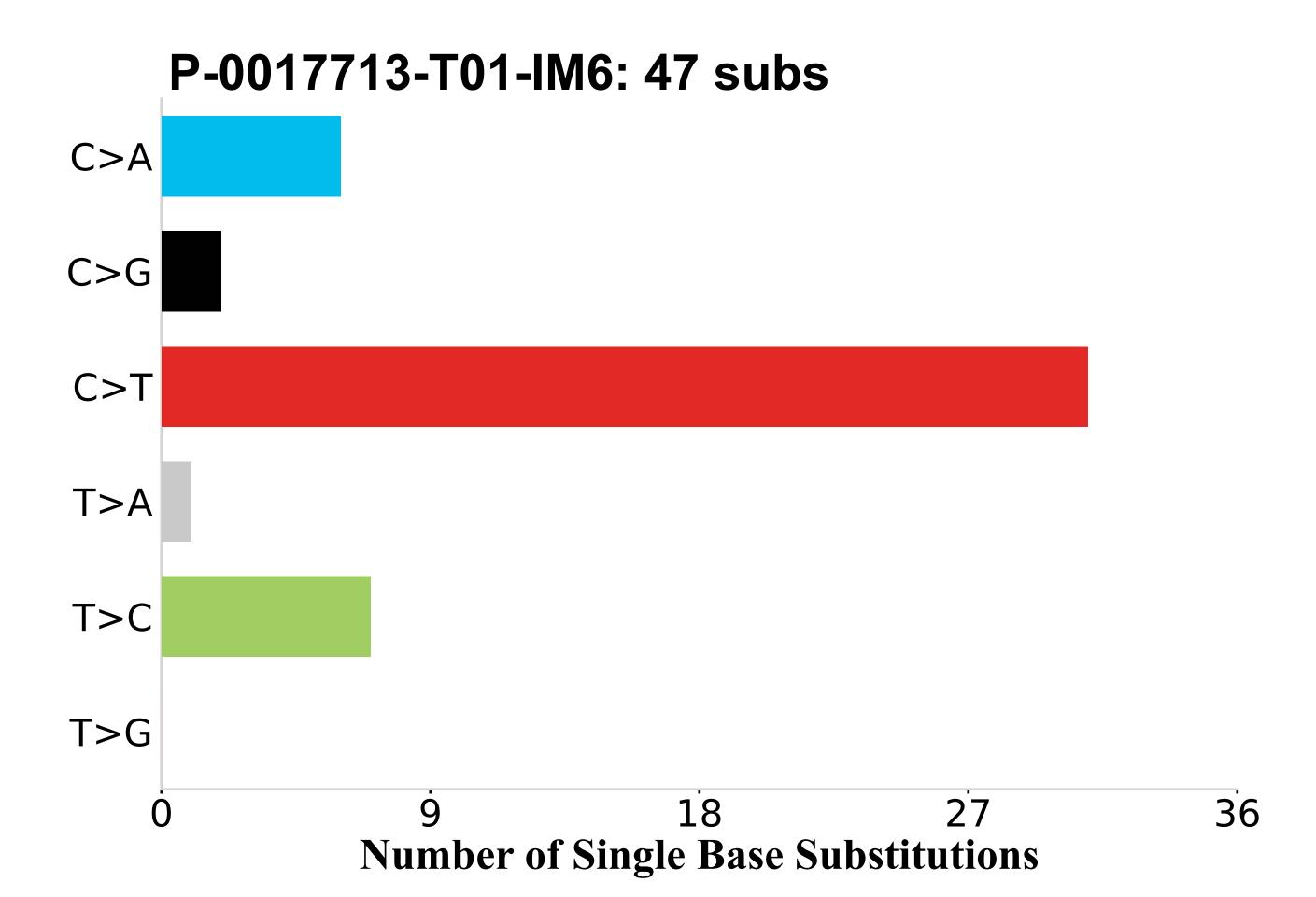


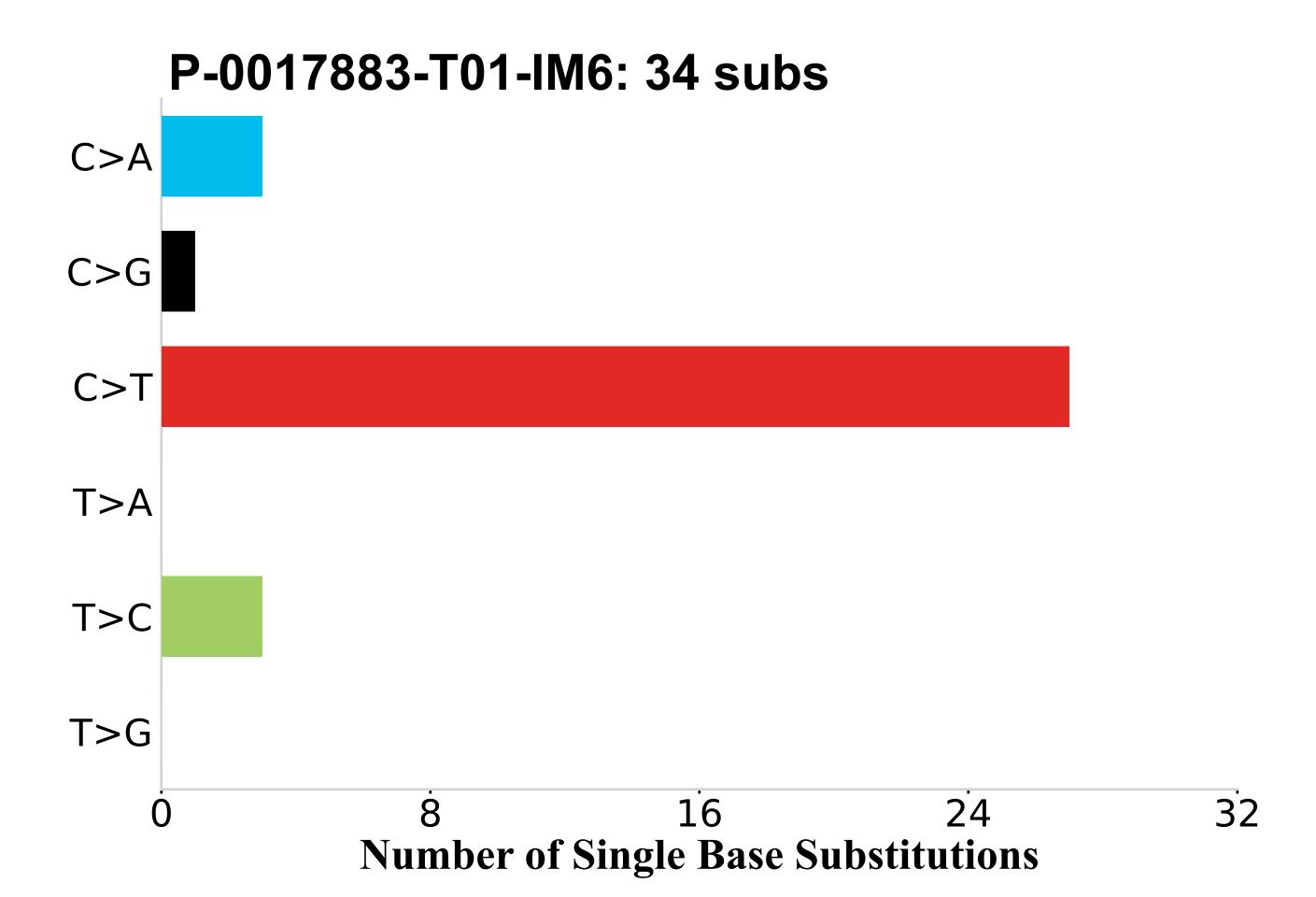
P-0016825-T01-IM6: 56 subs C>A C>G C>T T>AT>C T>G 24 32 8 16

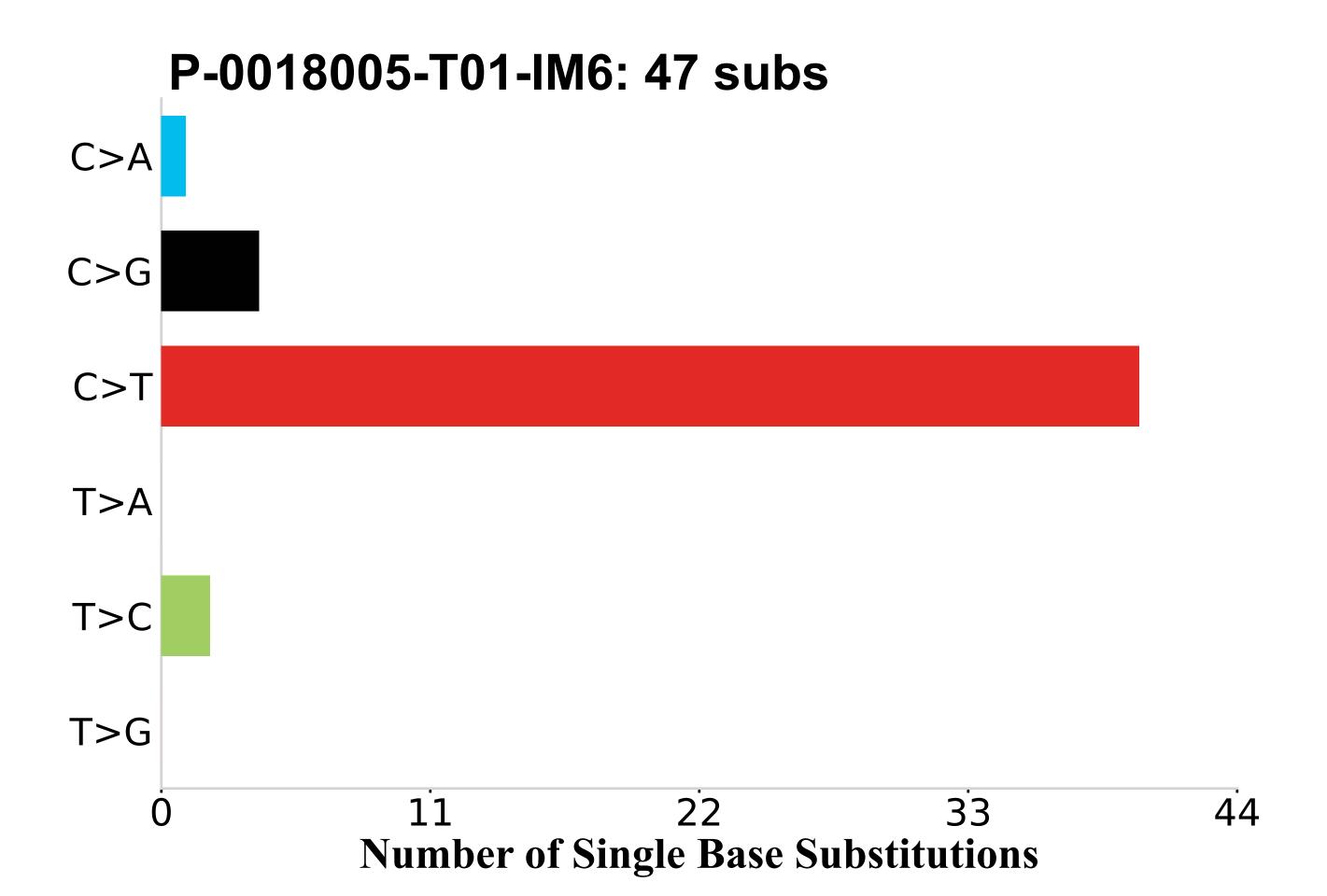
Number of Single Base Substitutions

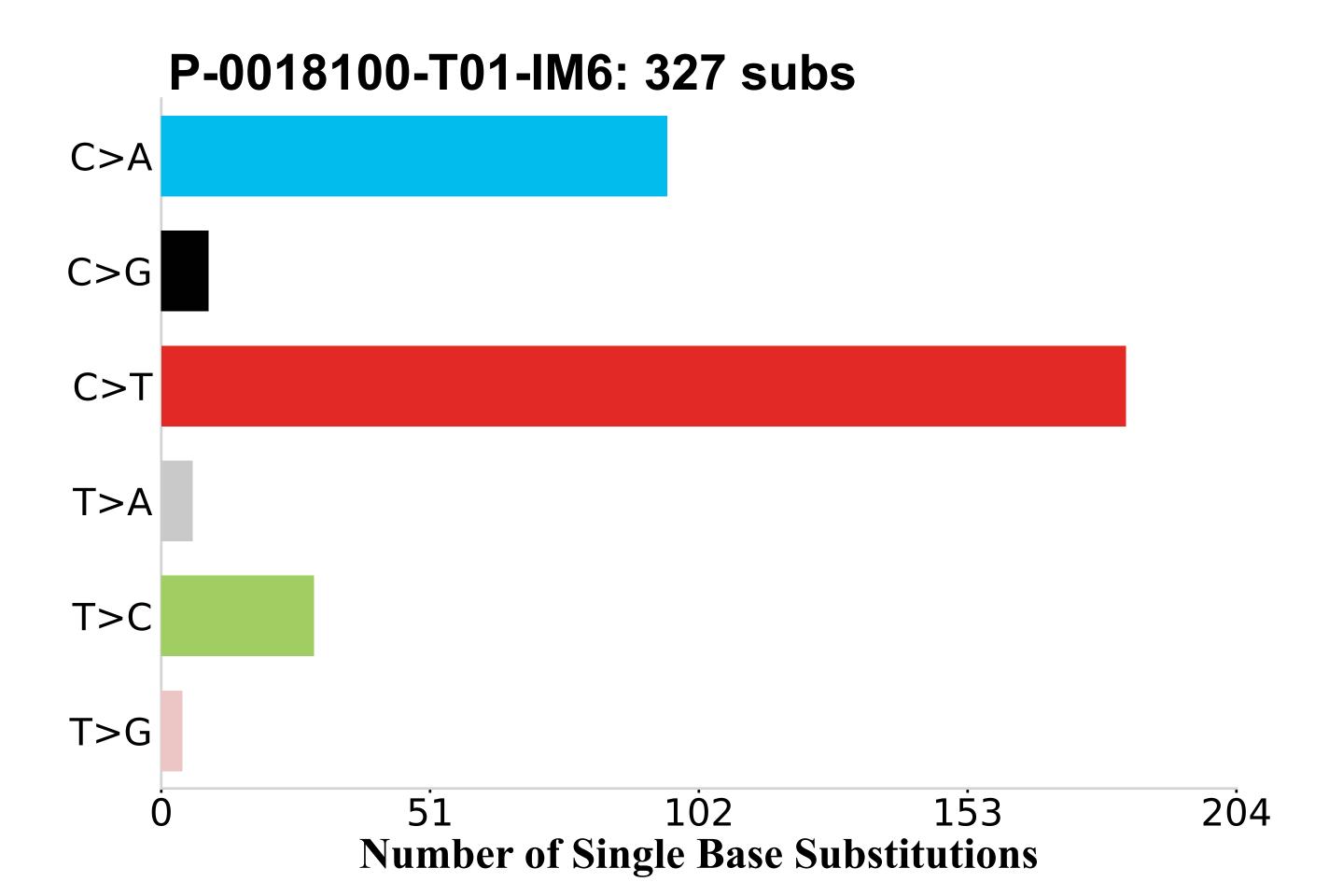


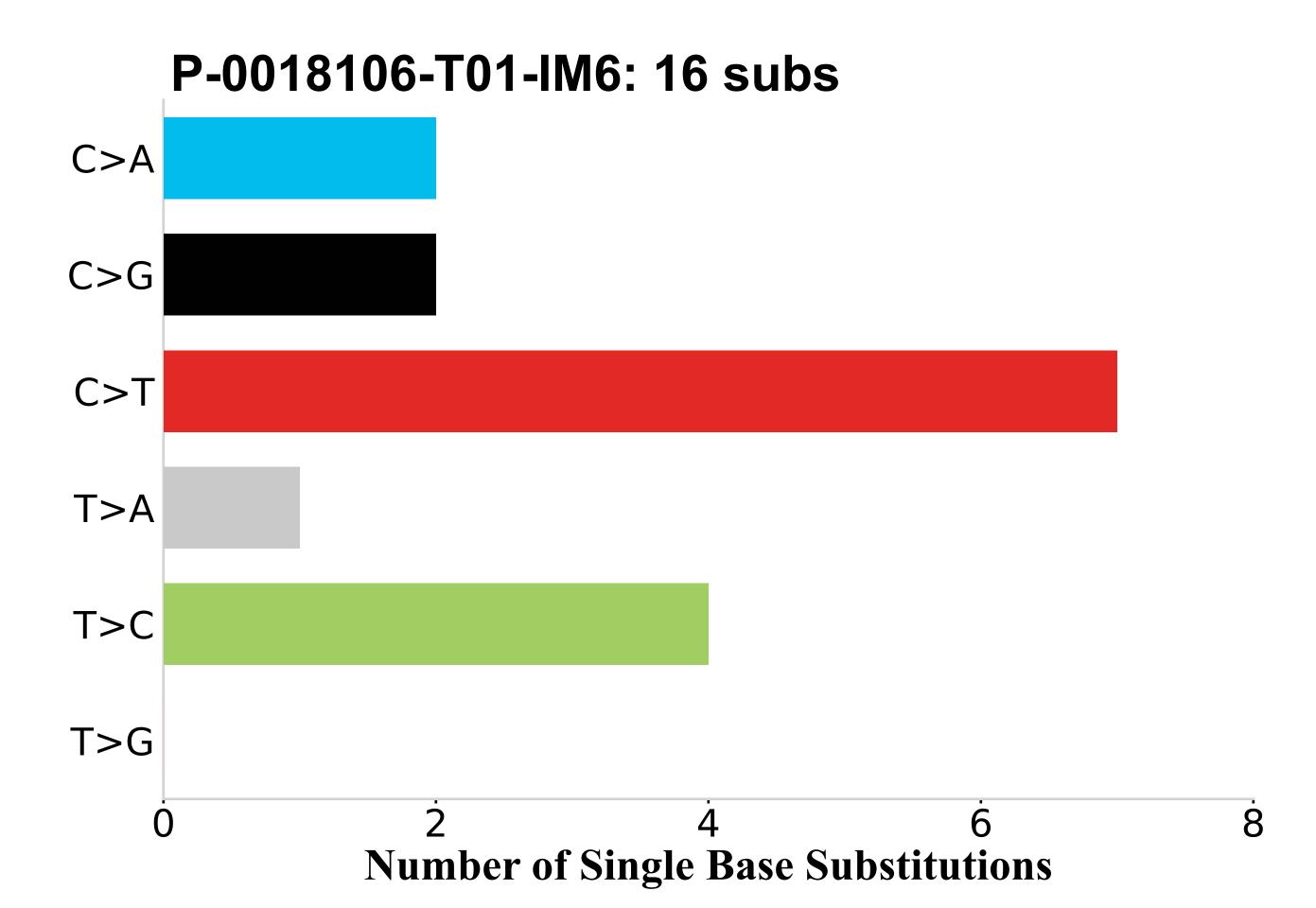


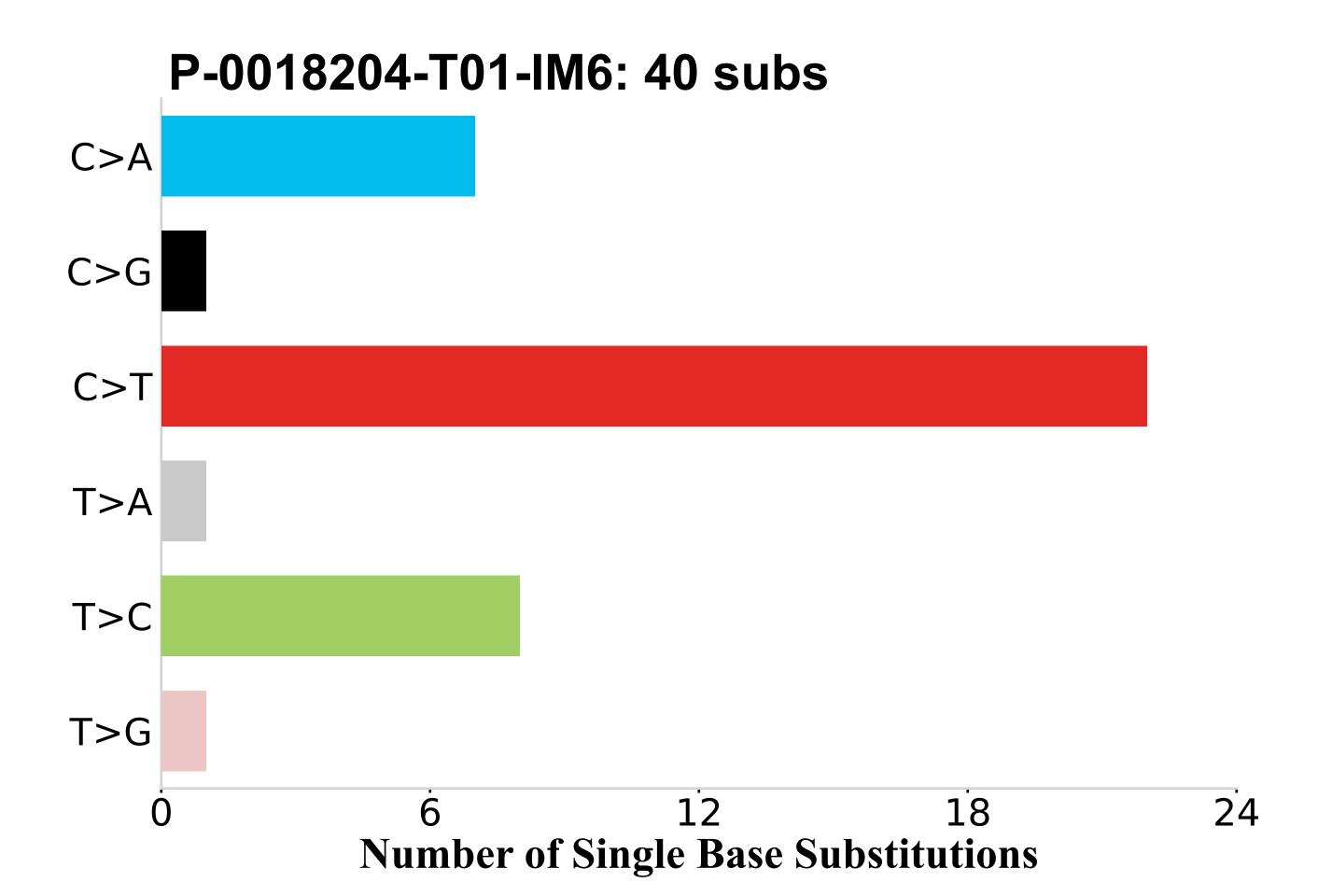


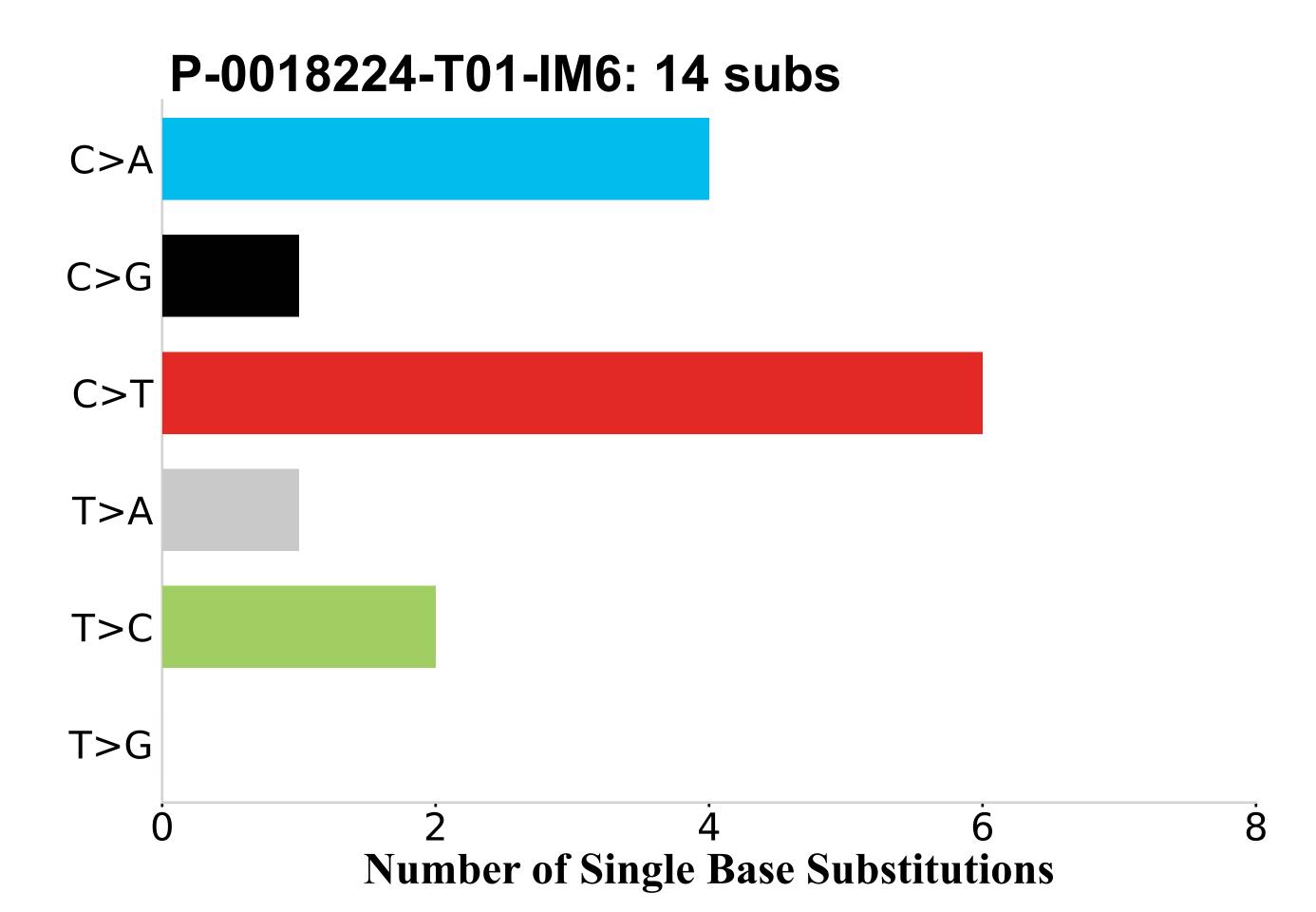


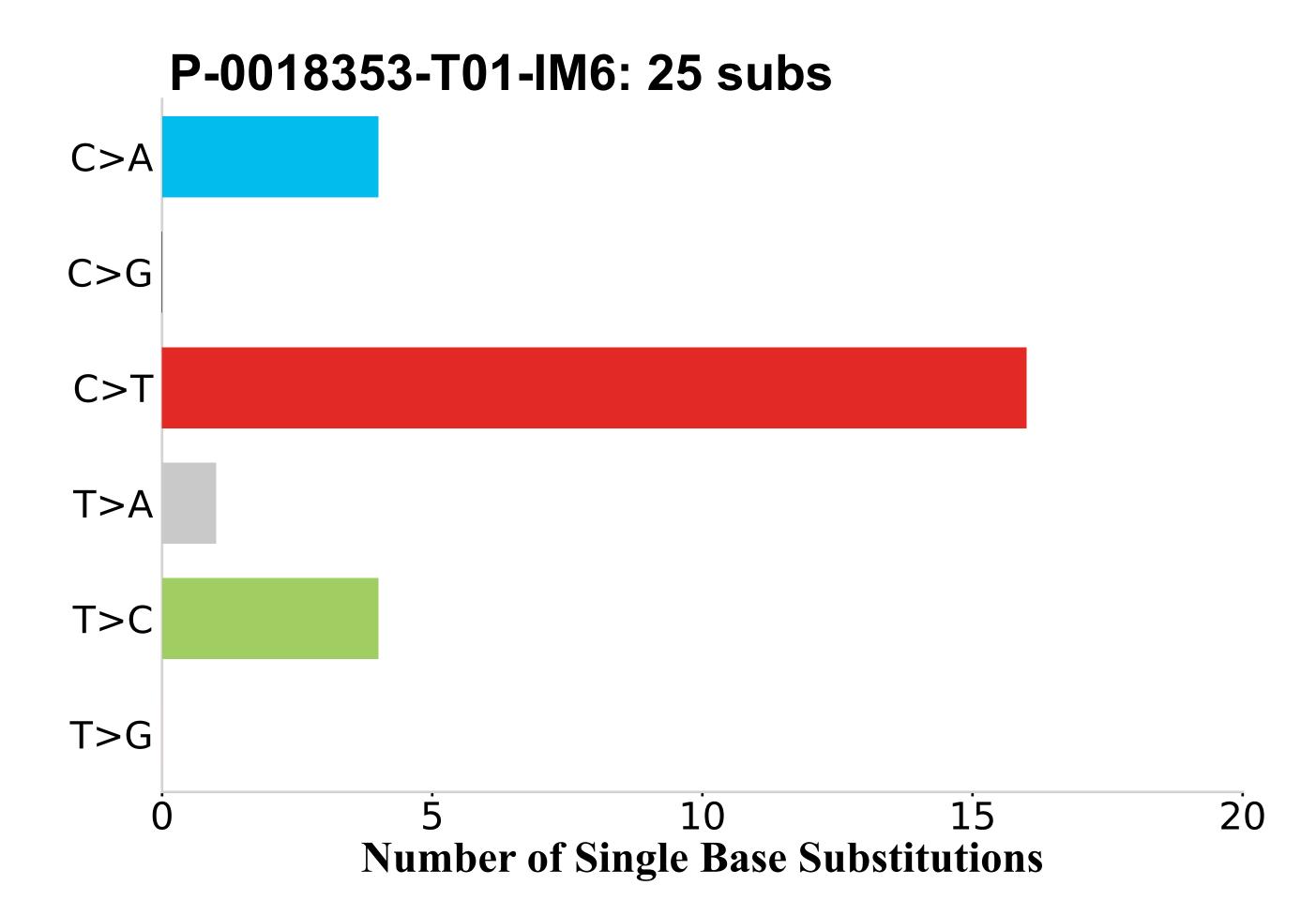




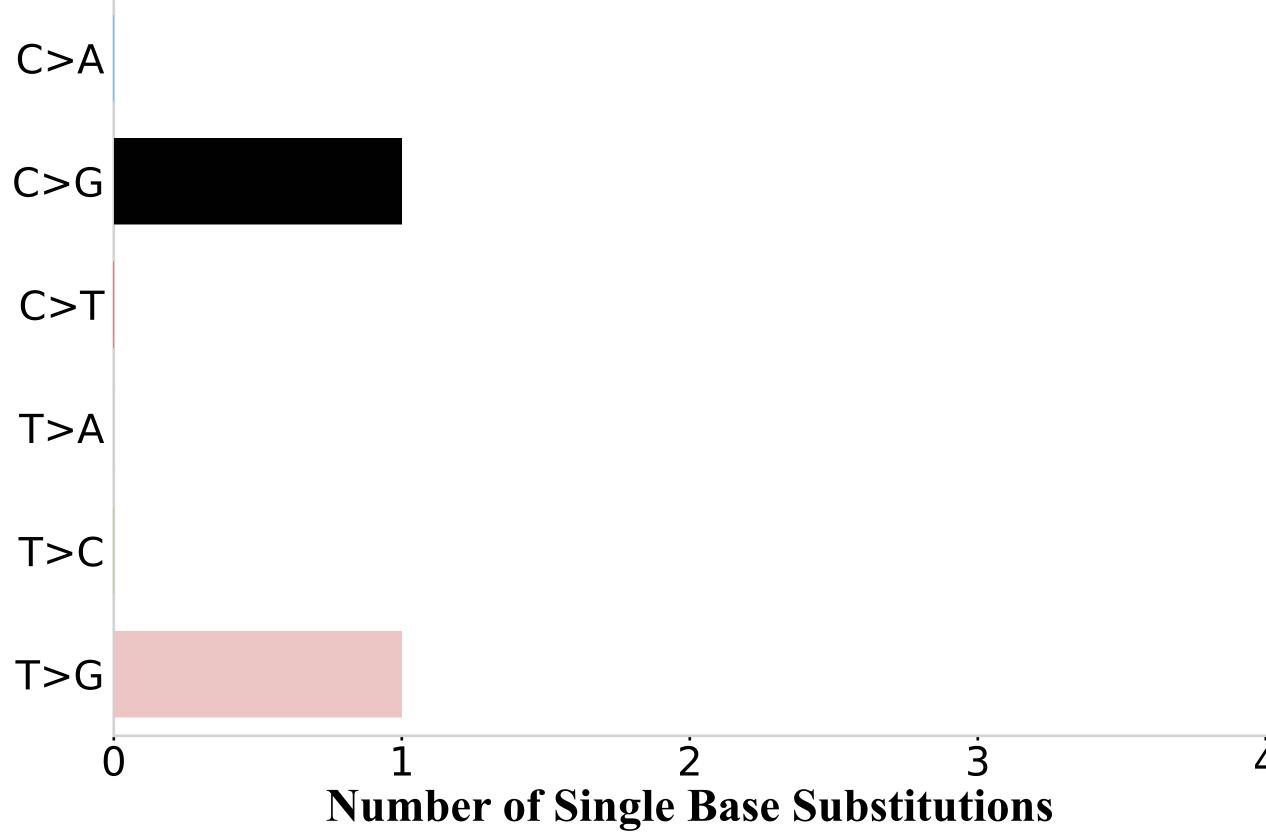


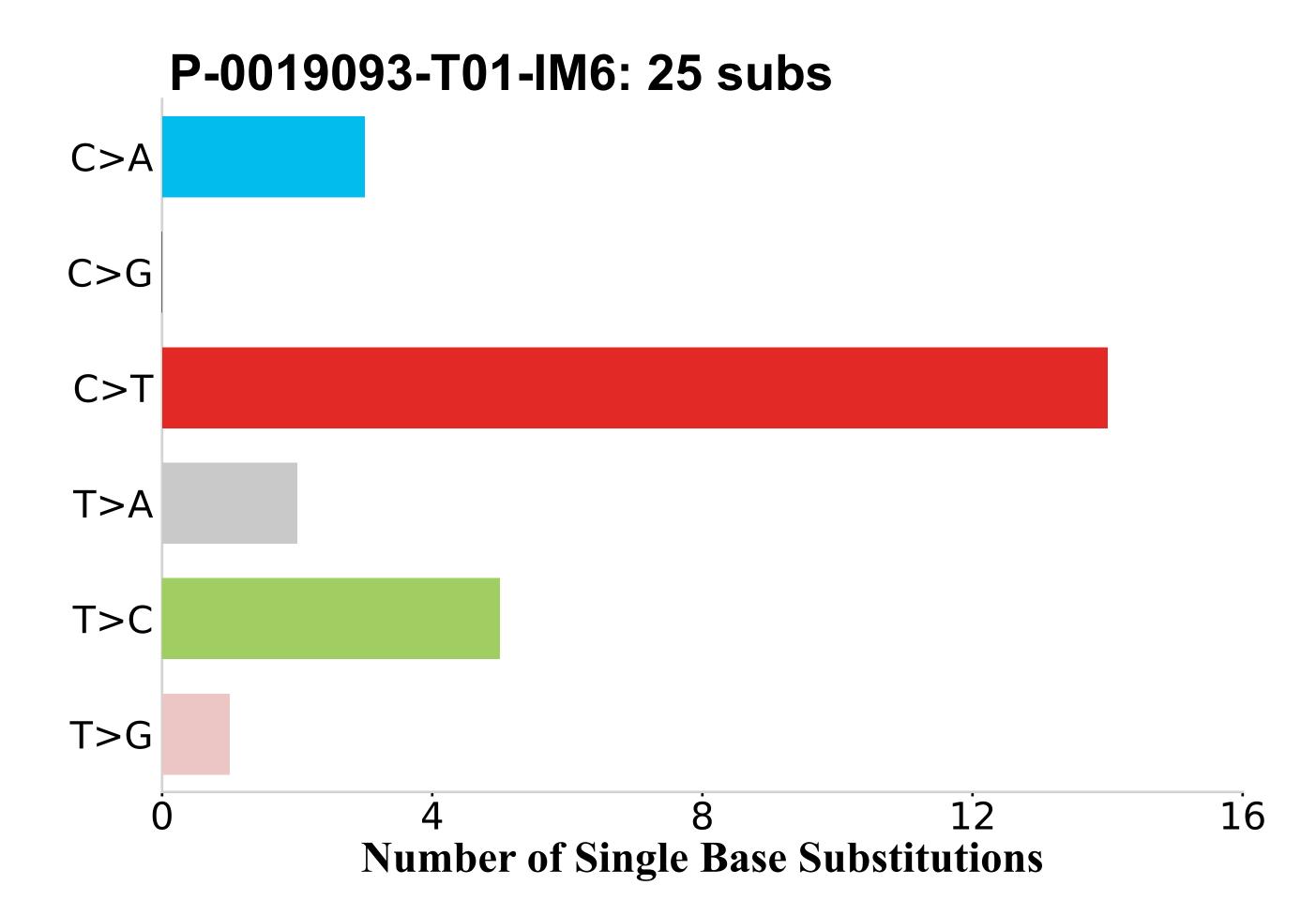


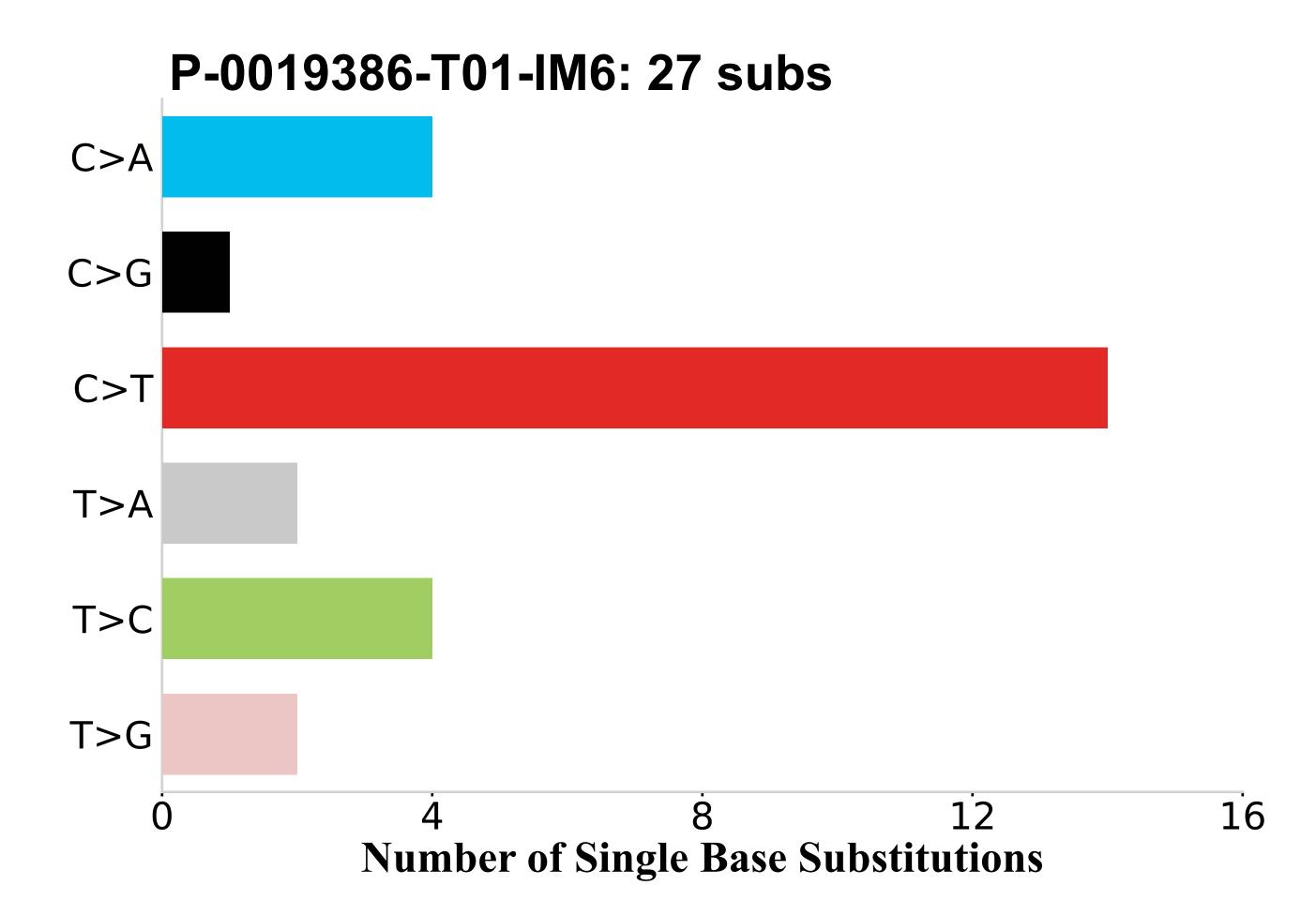


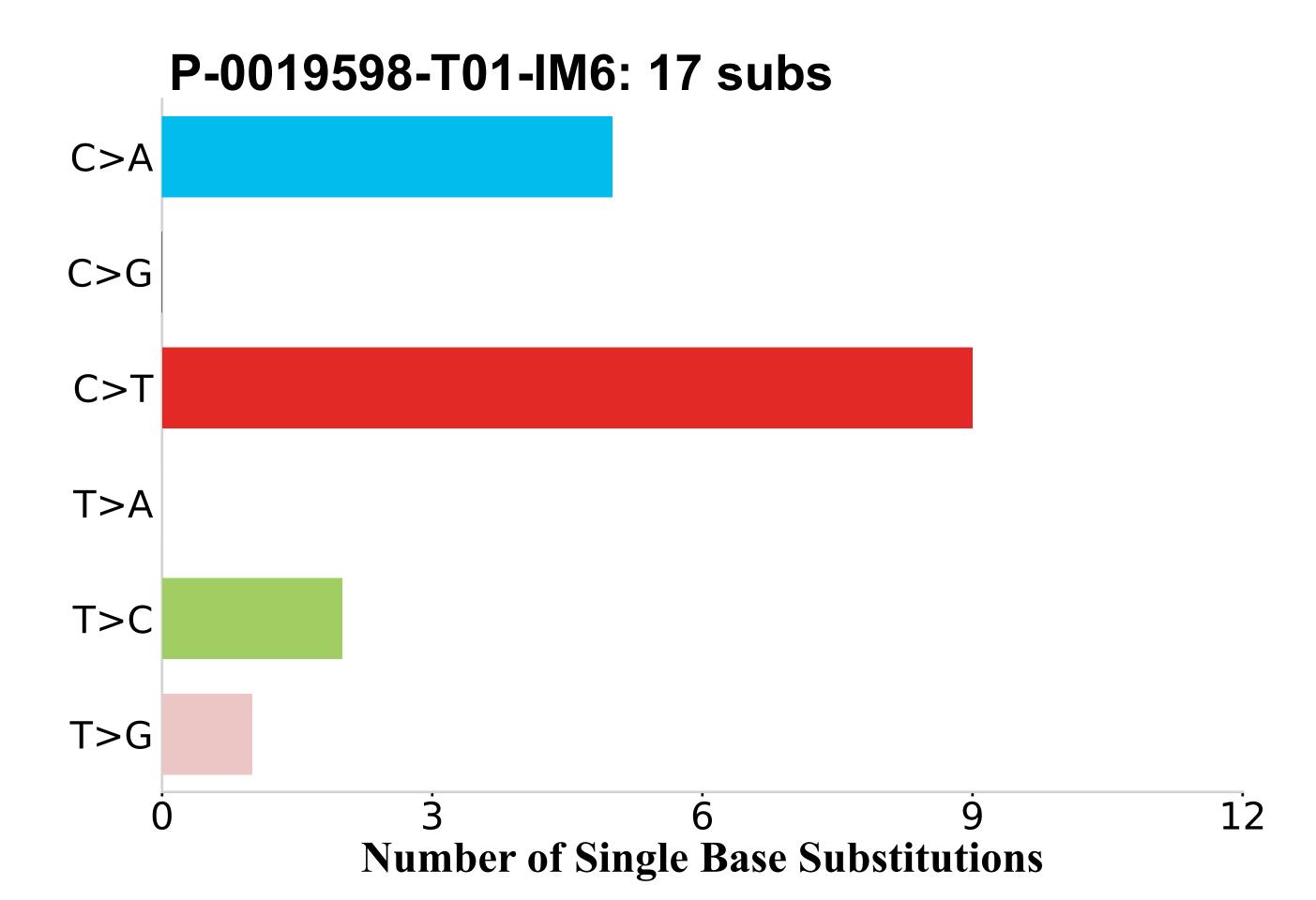


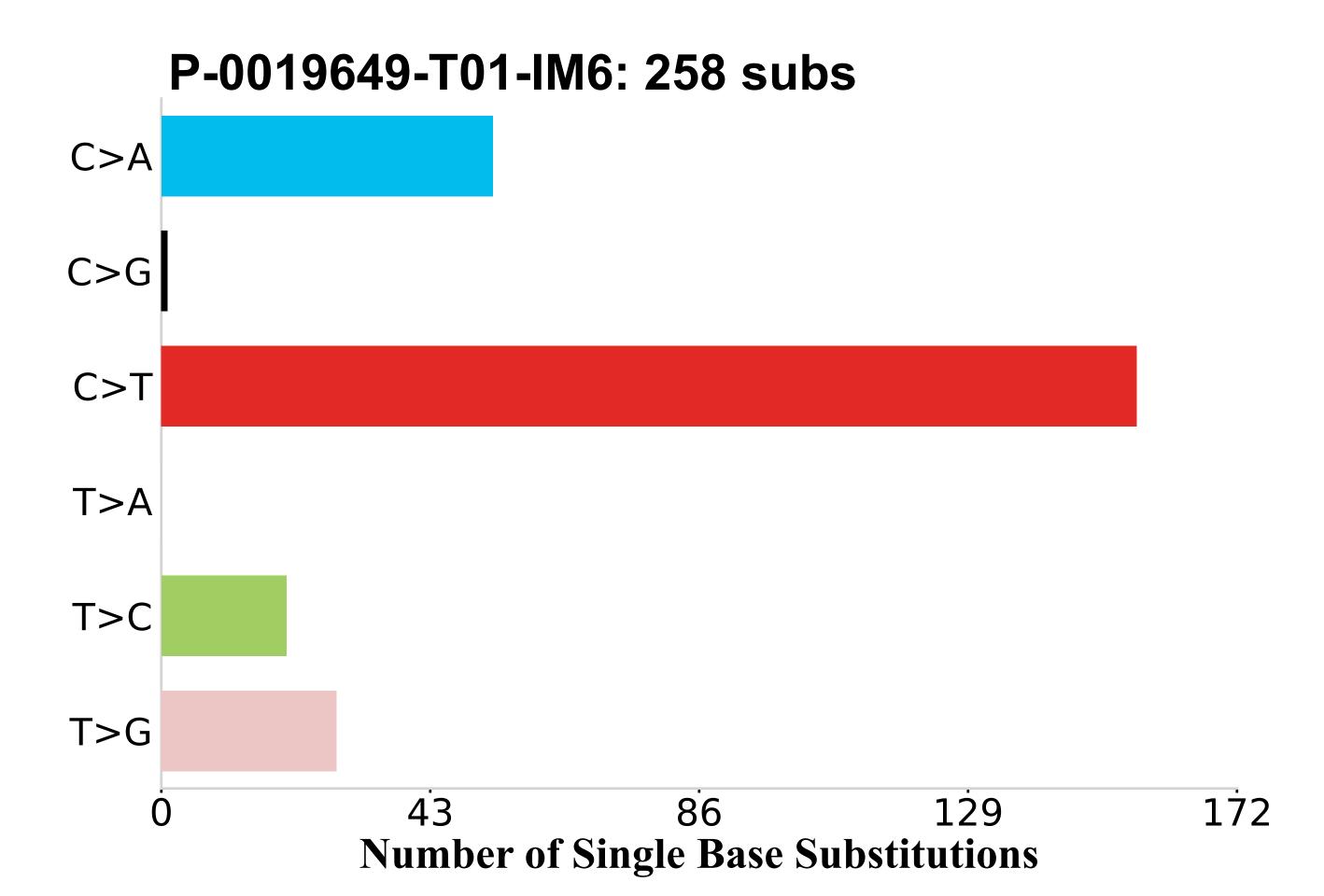
P-0018723-T01-IM6: 2 subs

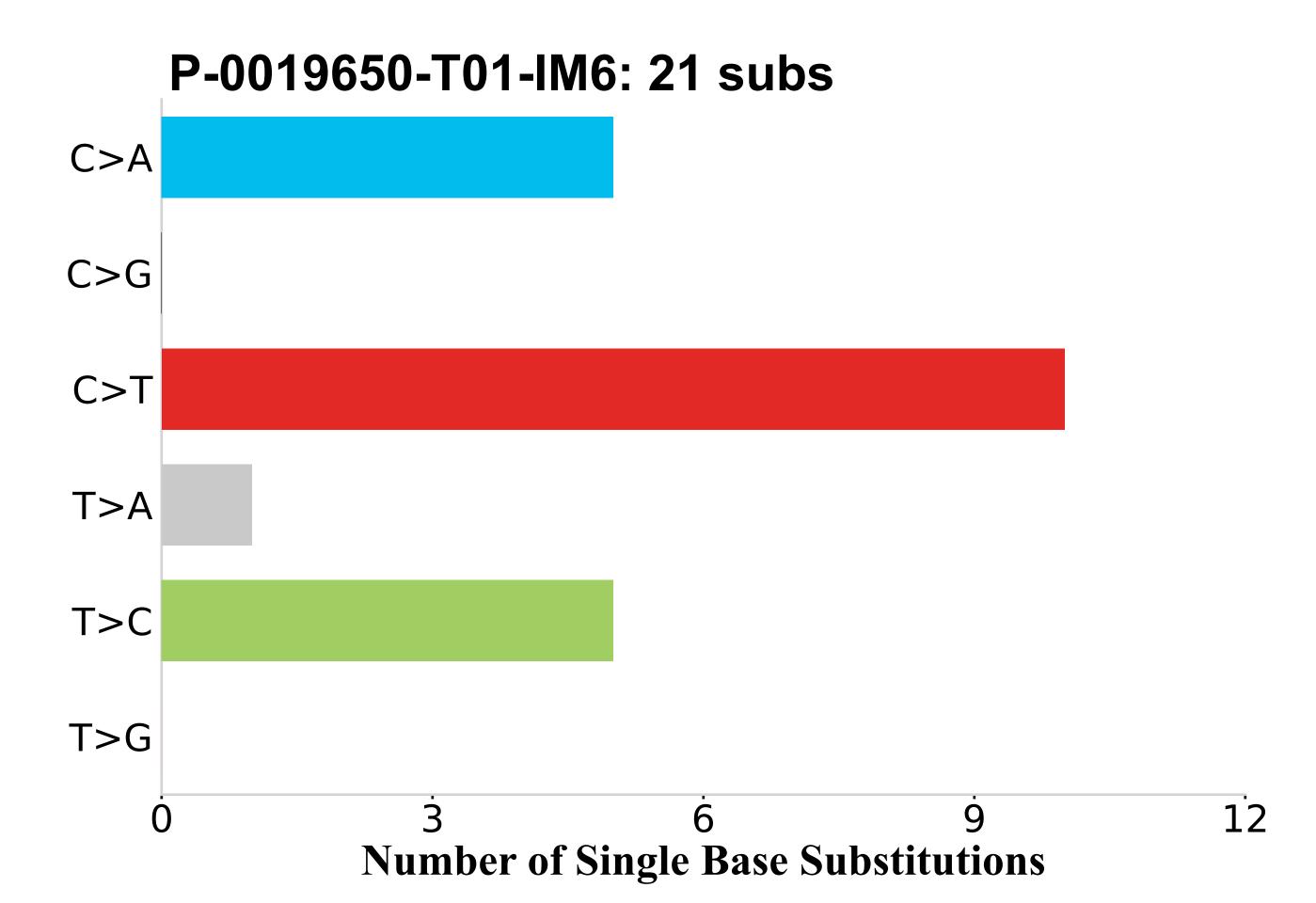


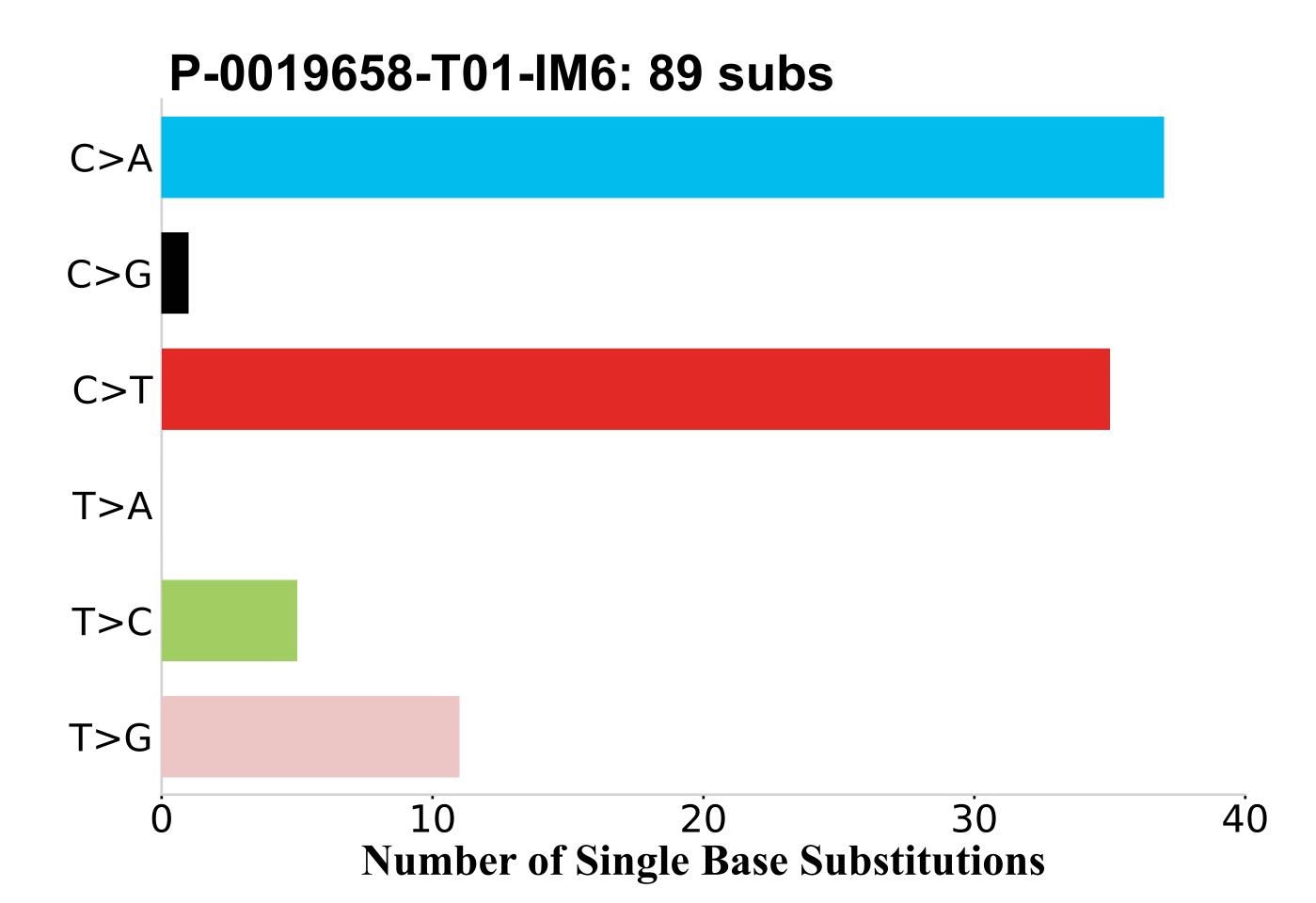


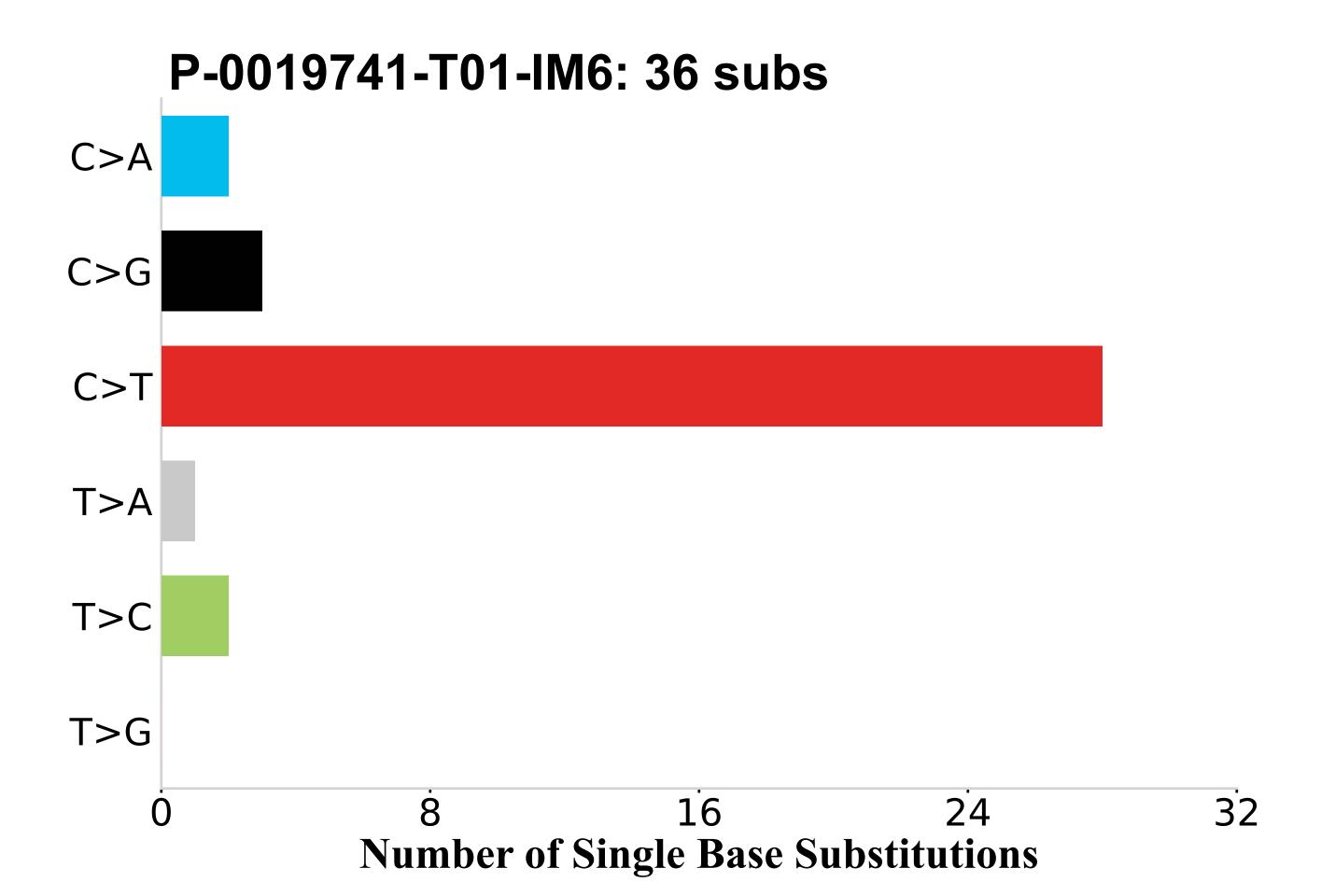


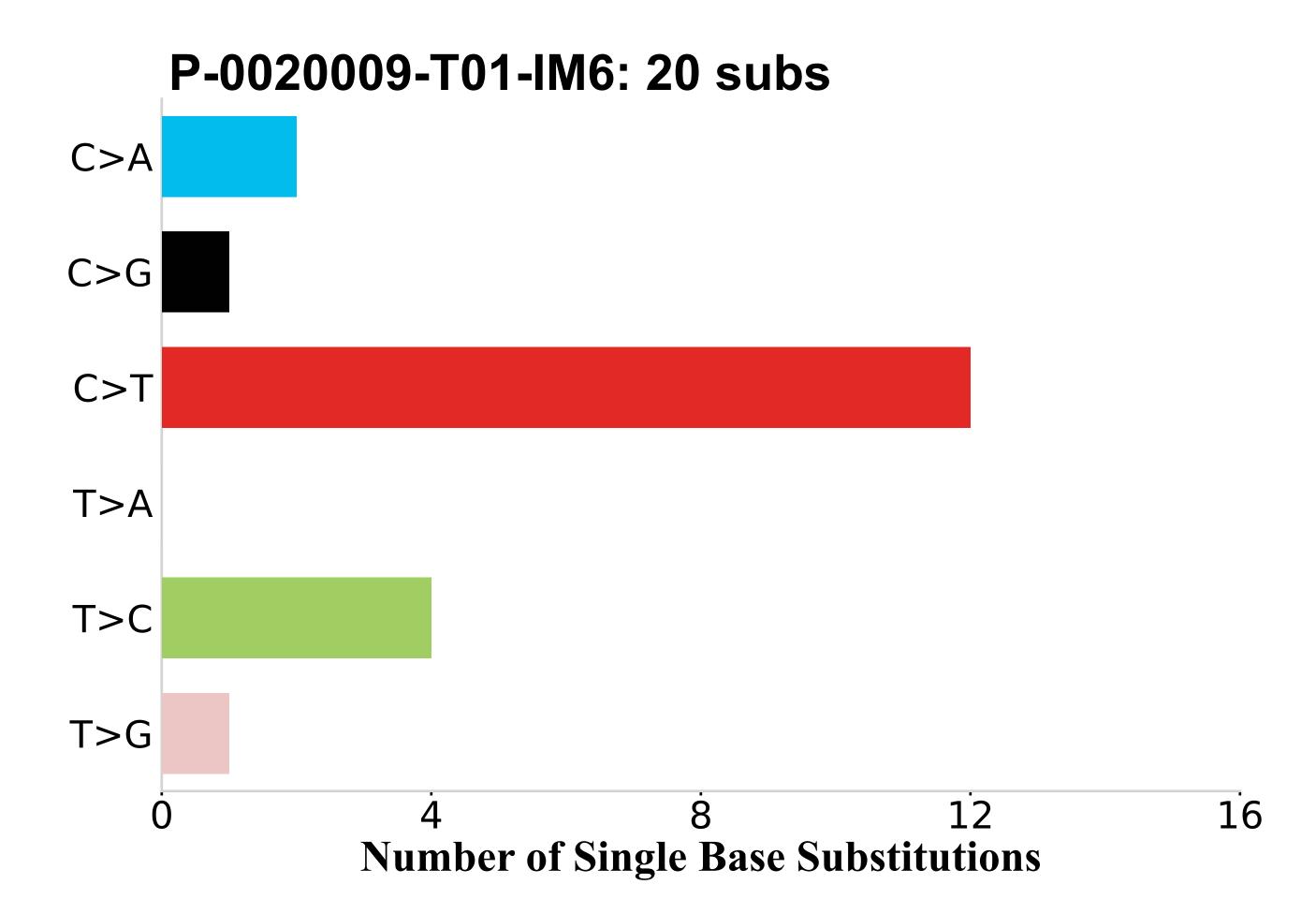


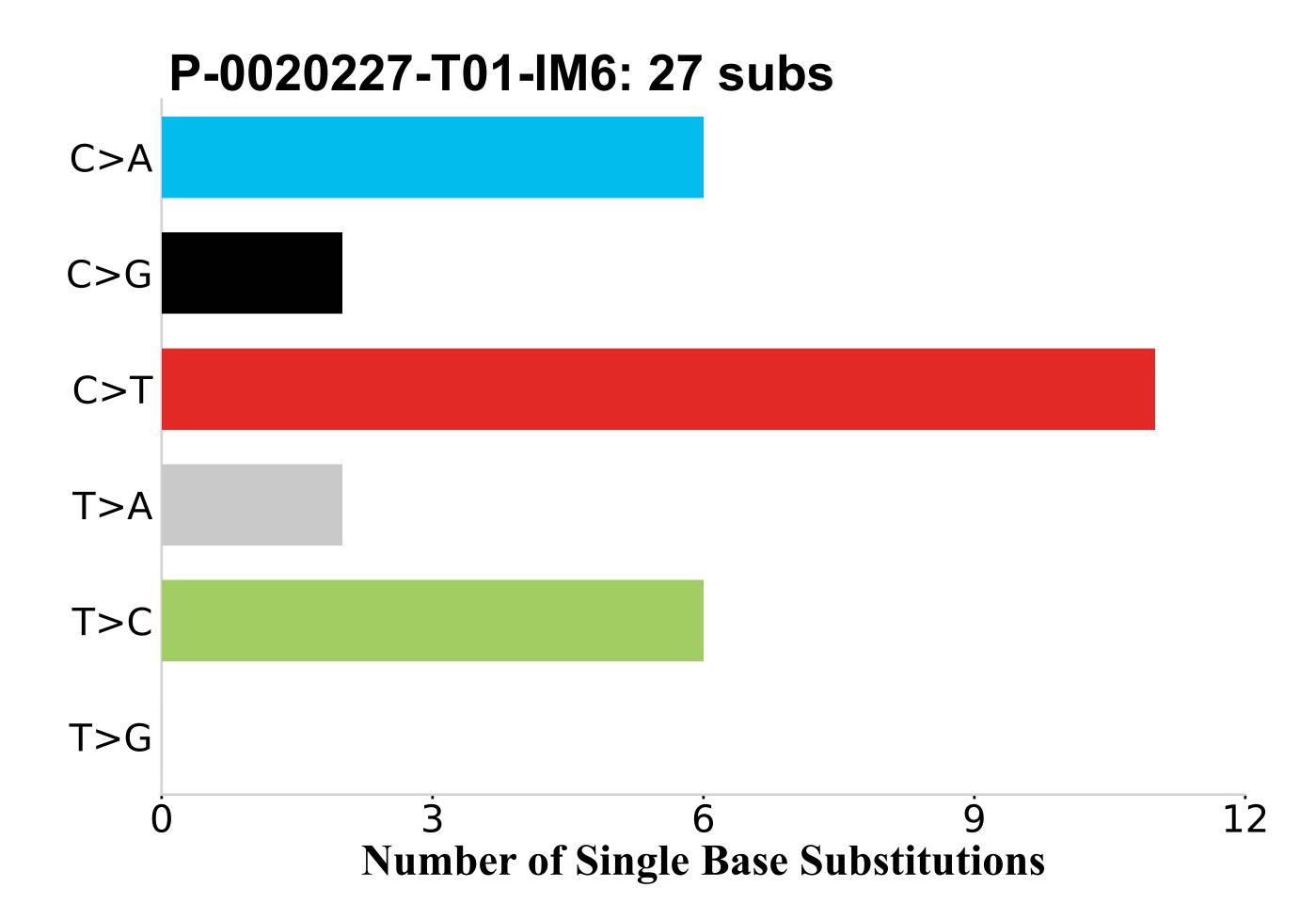


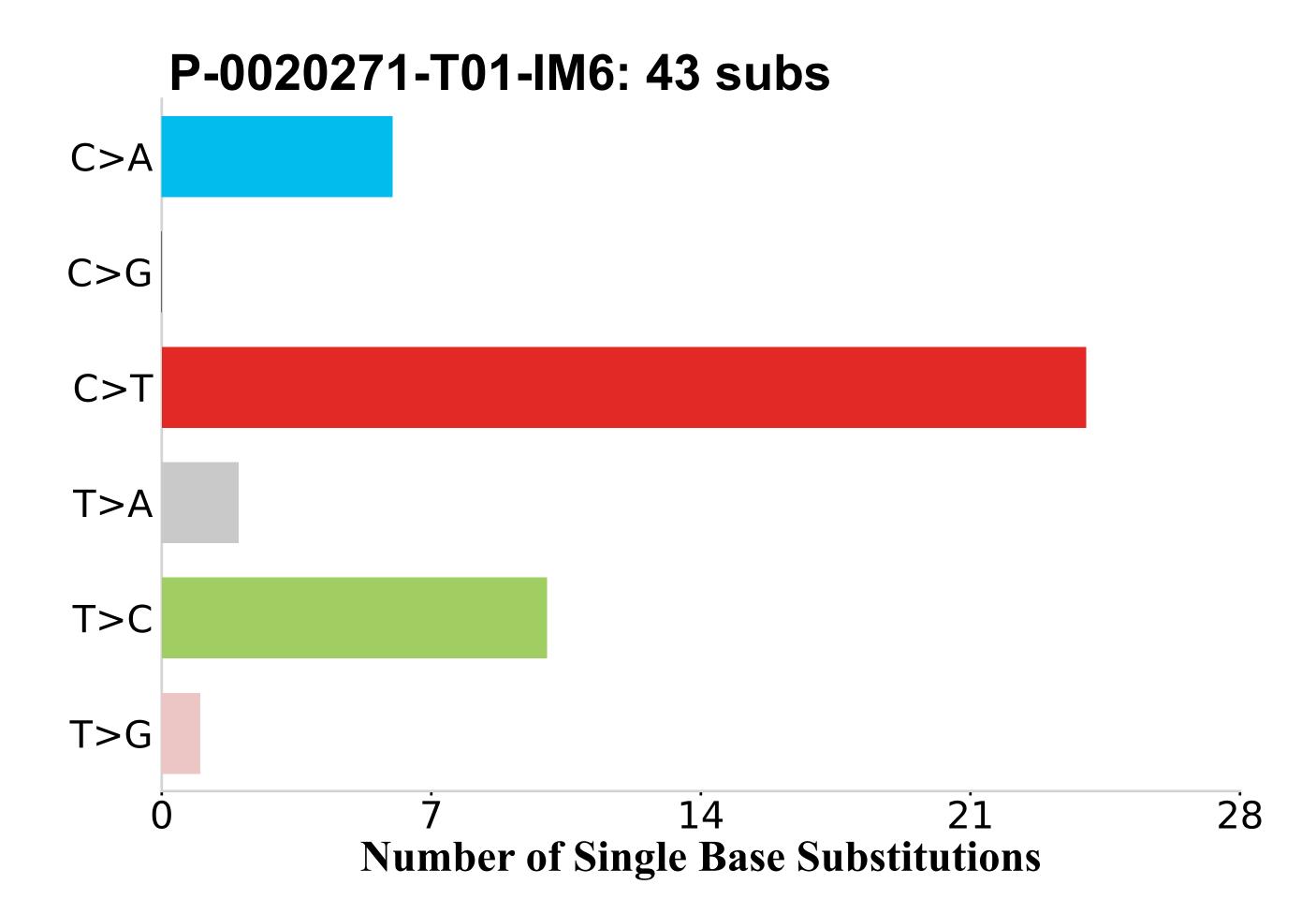


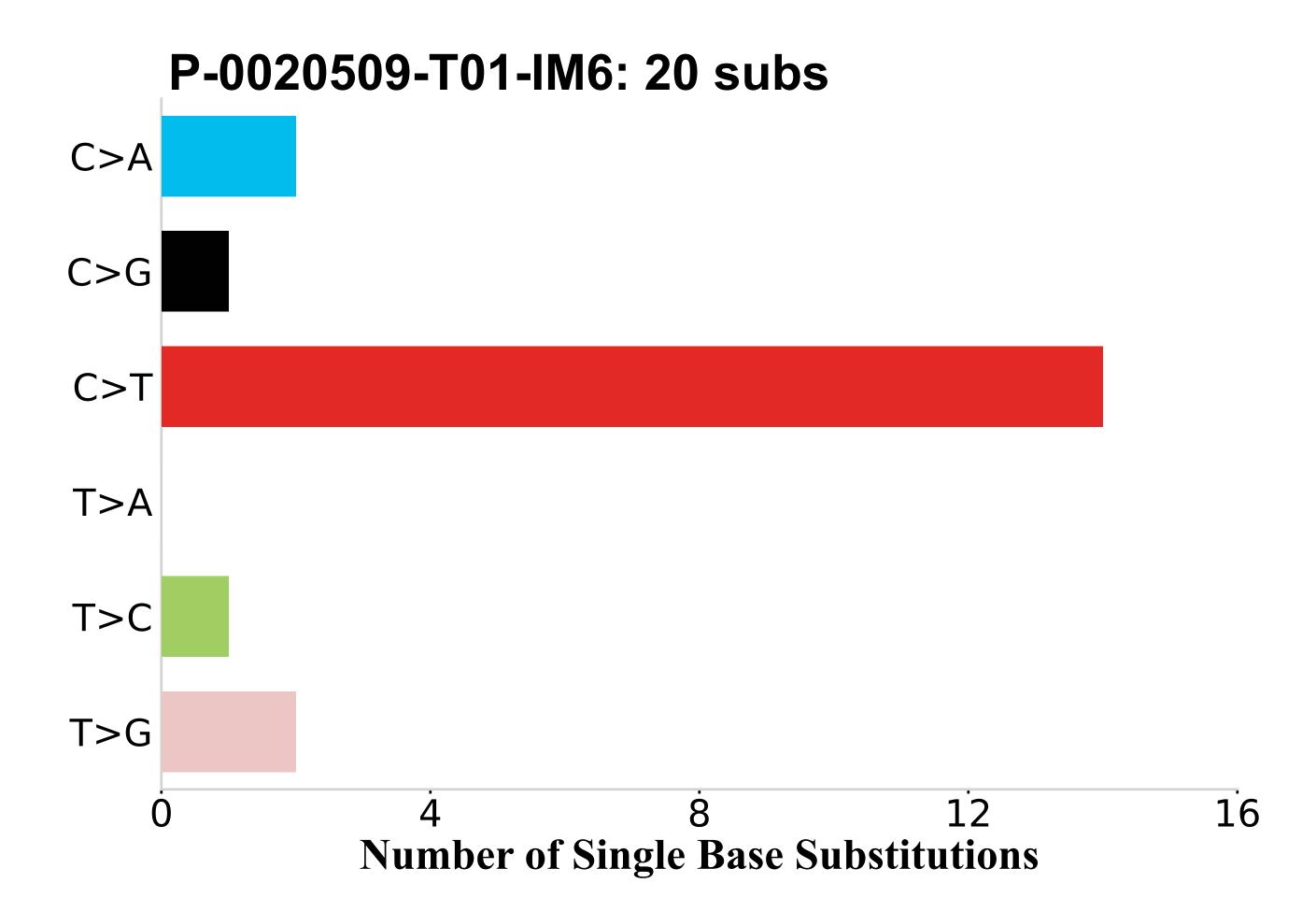


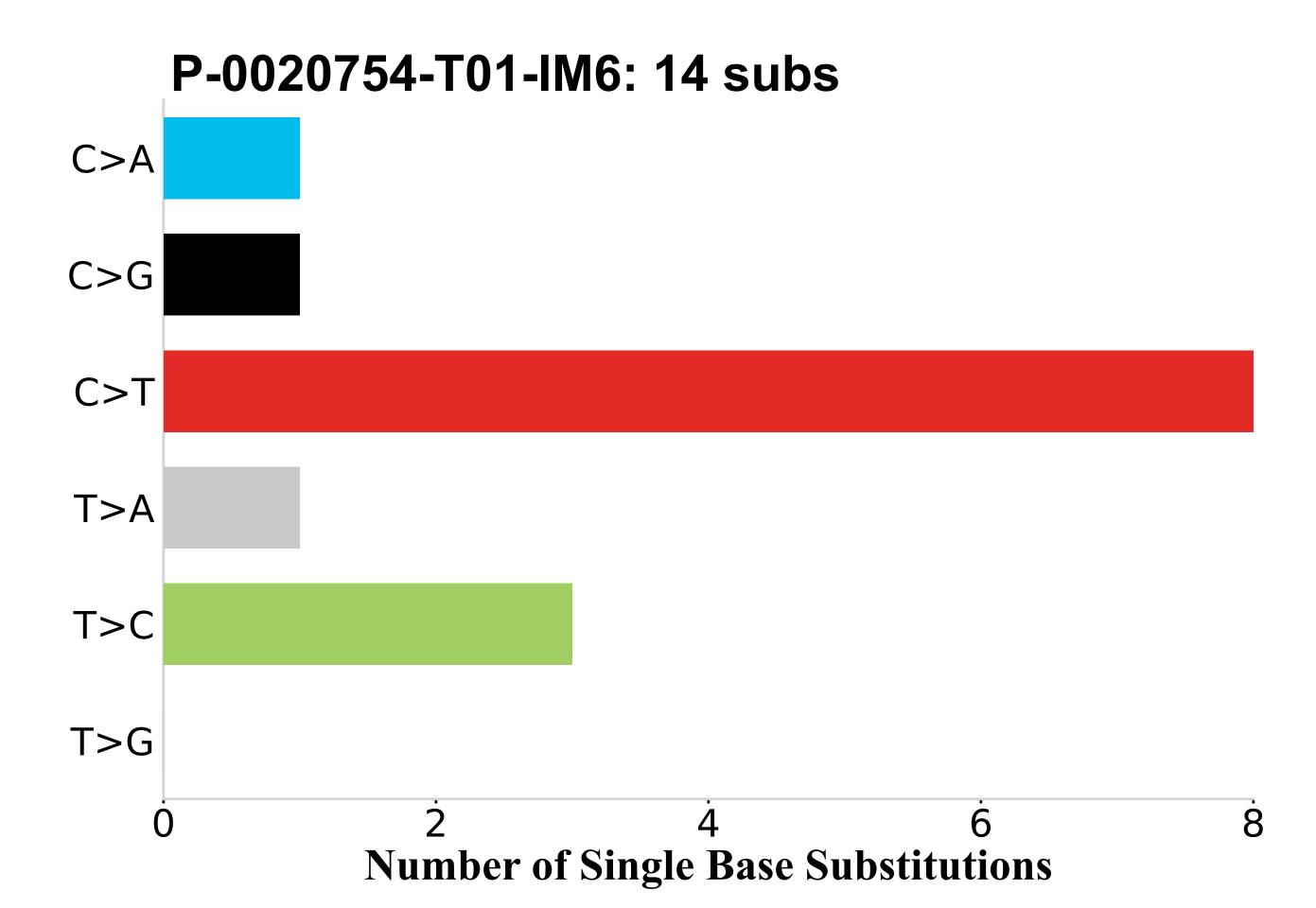


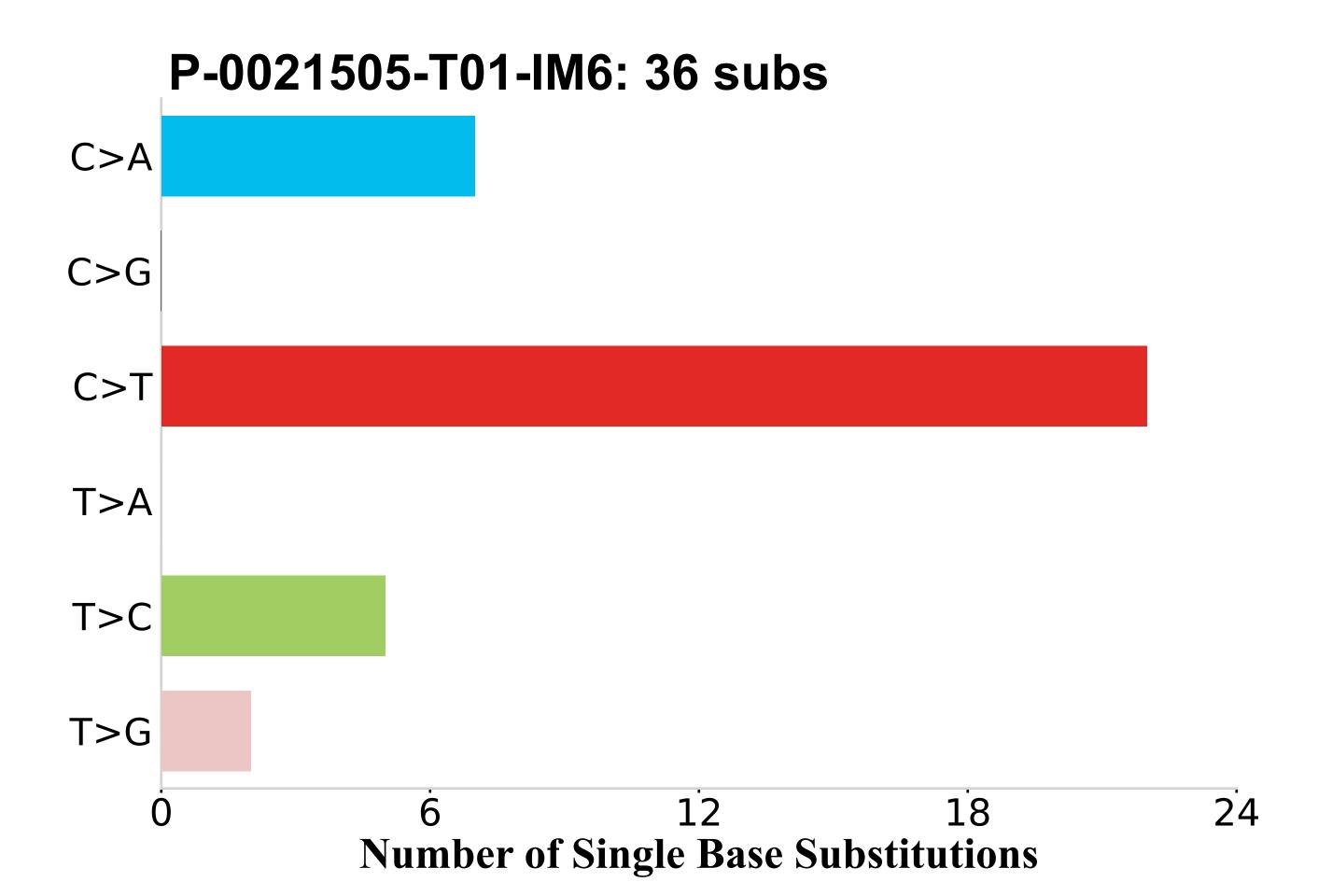


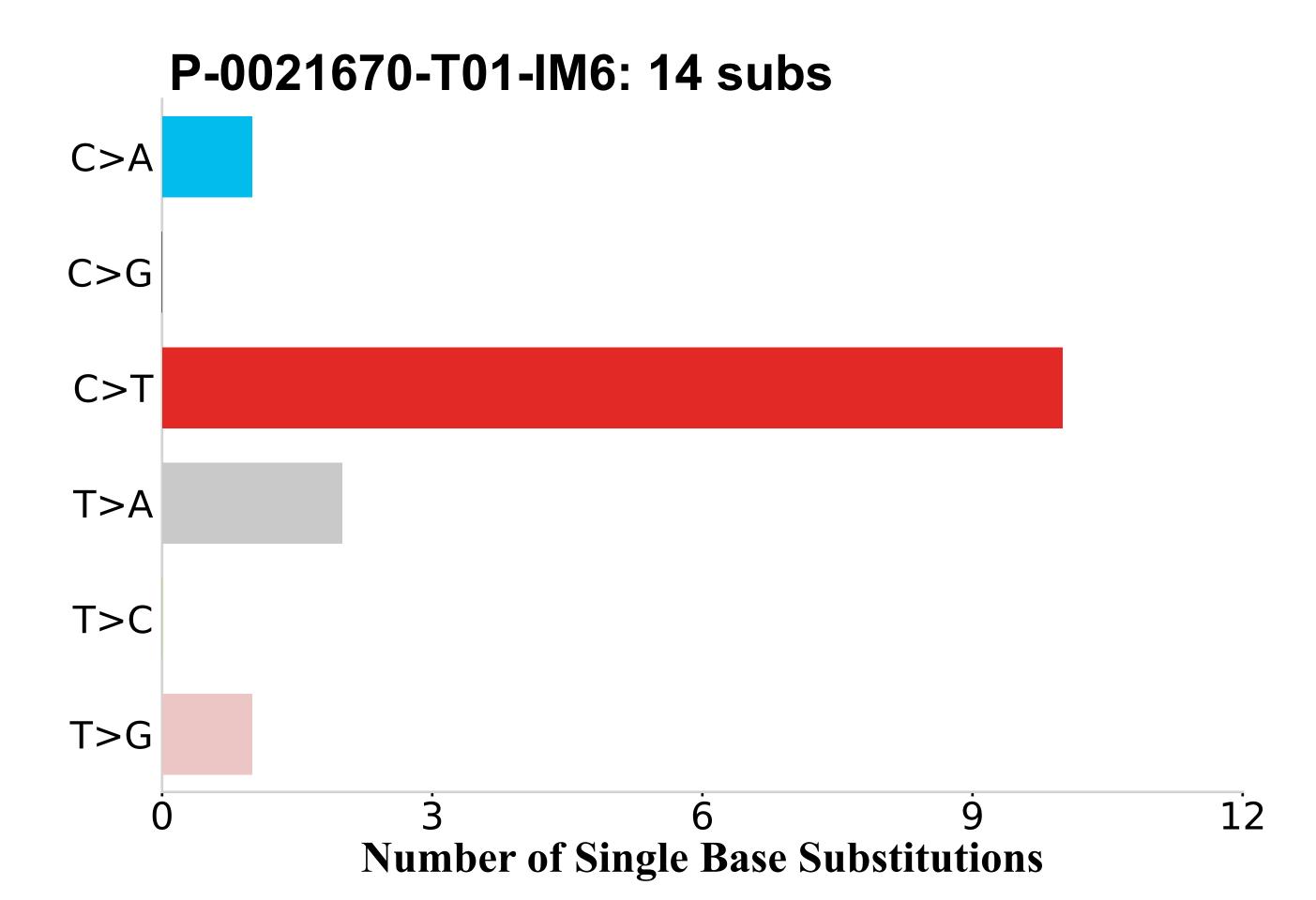


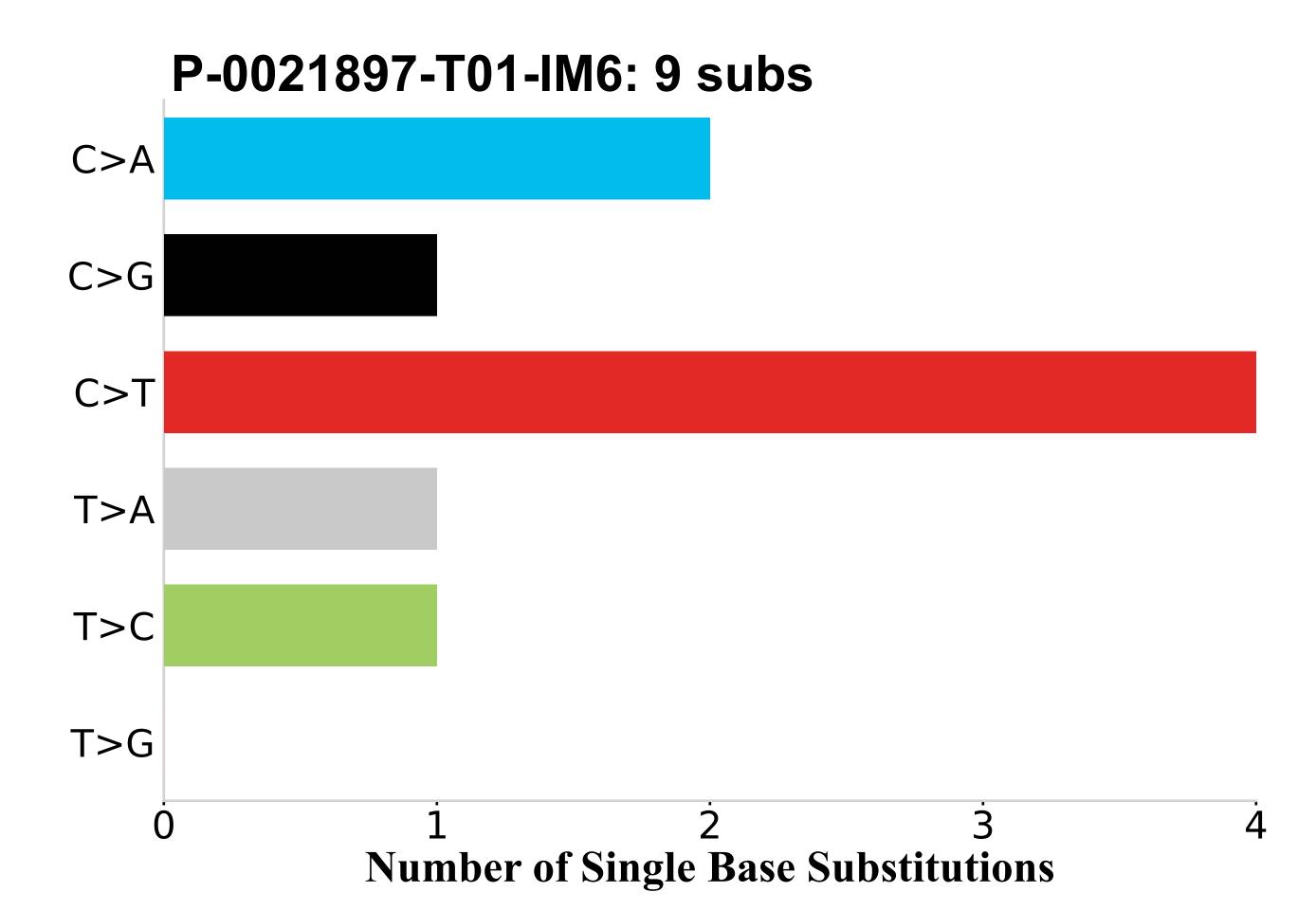


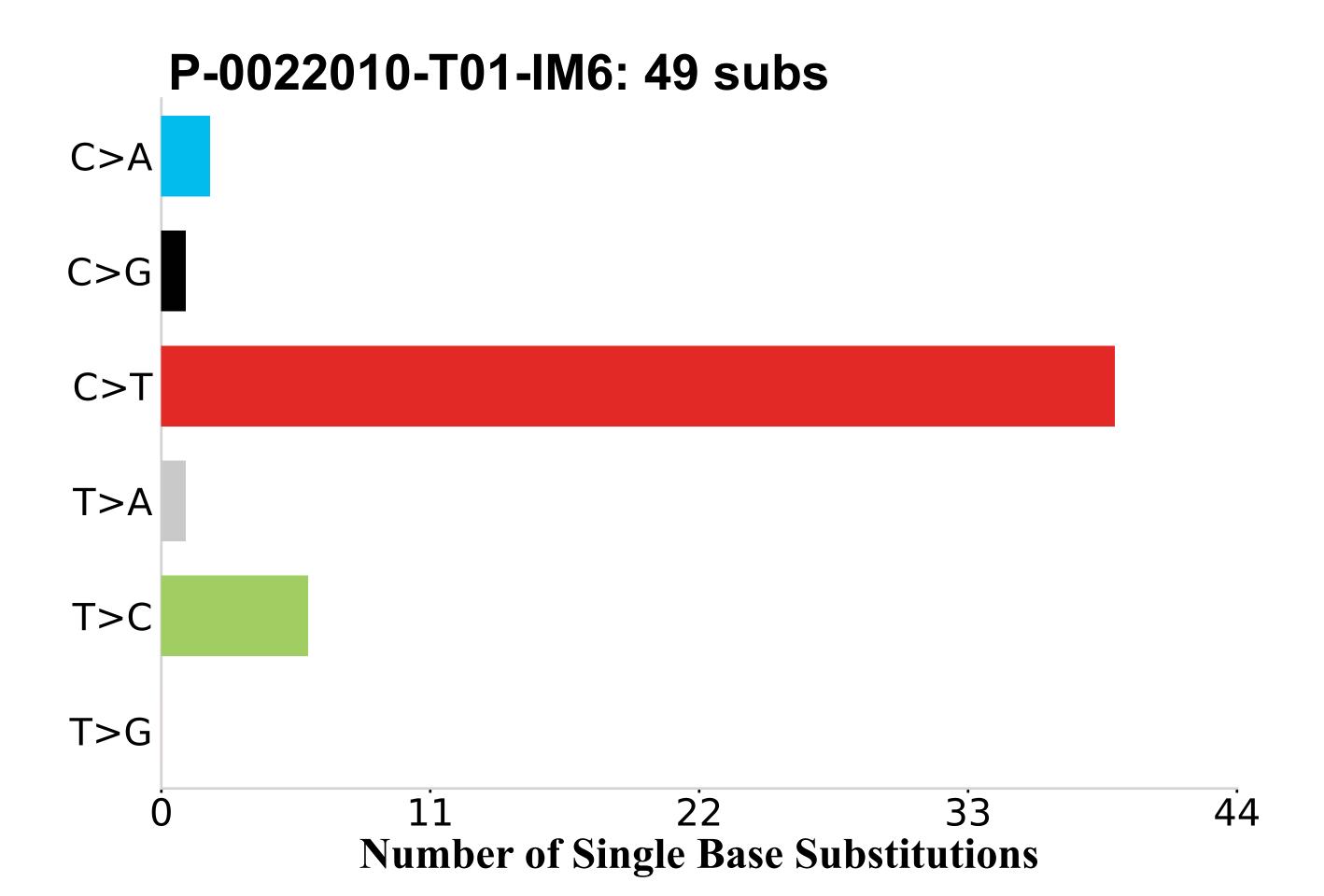




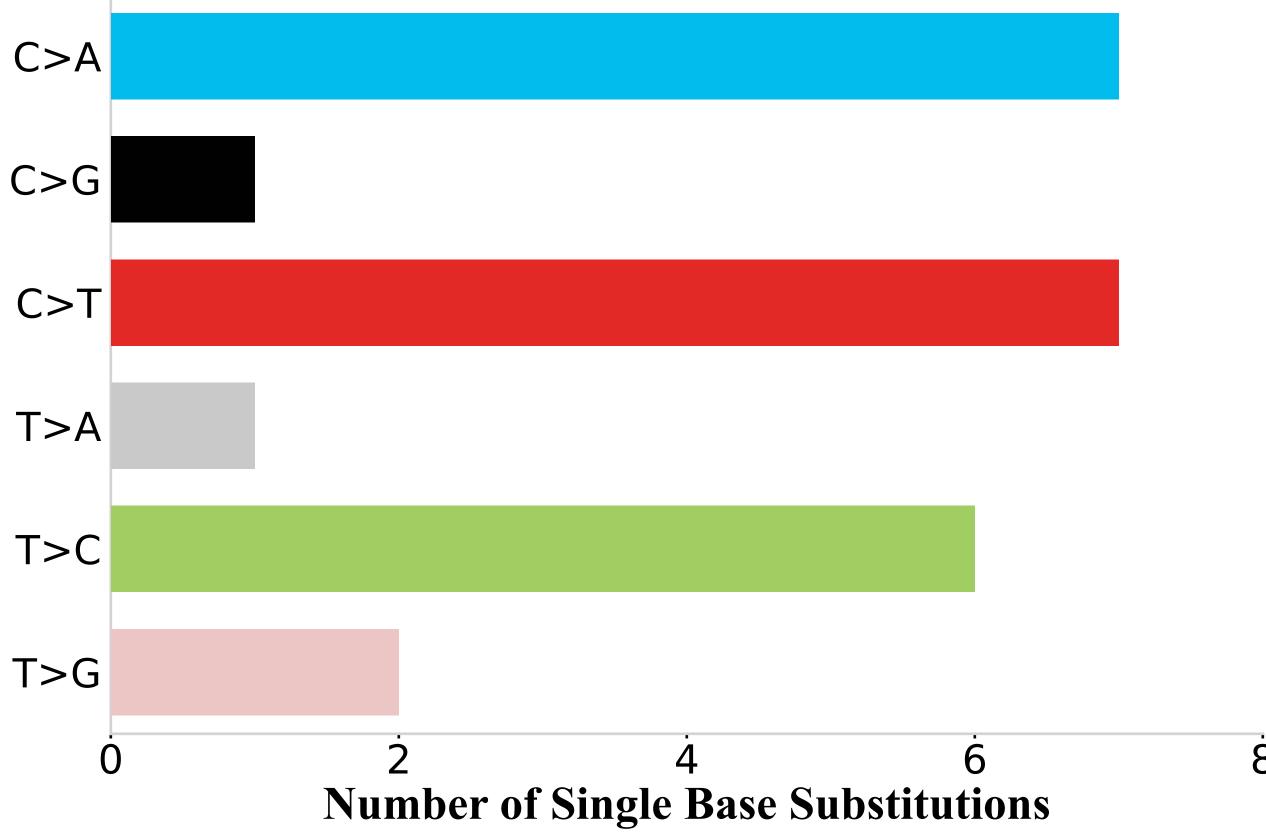


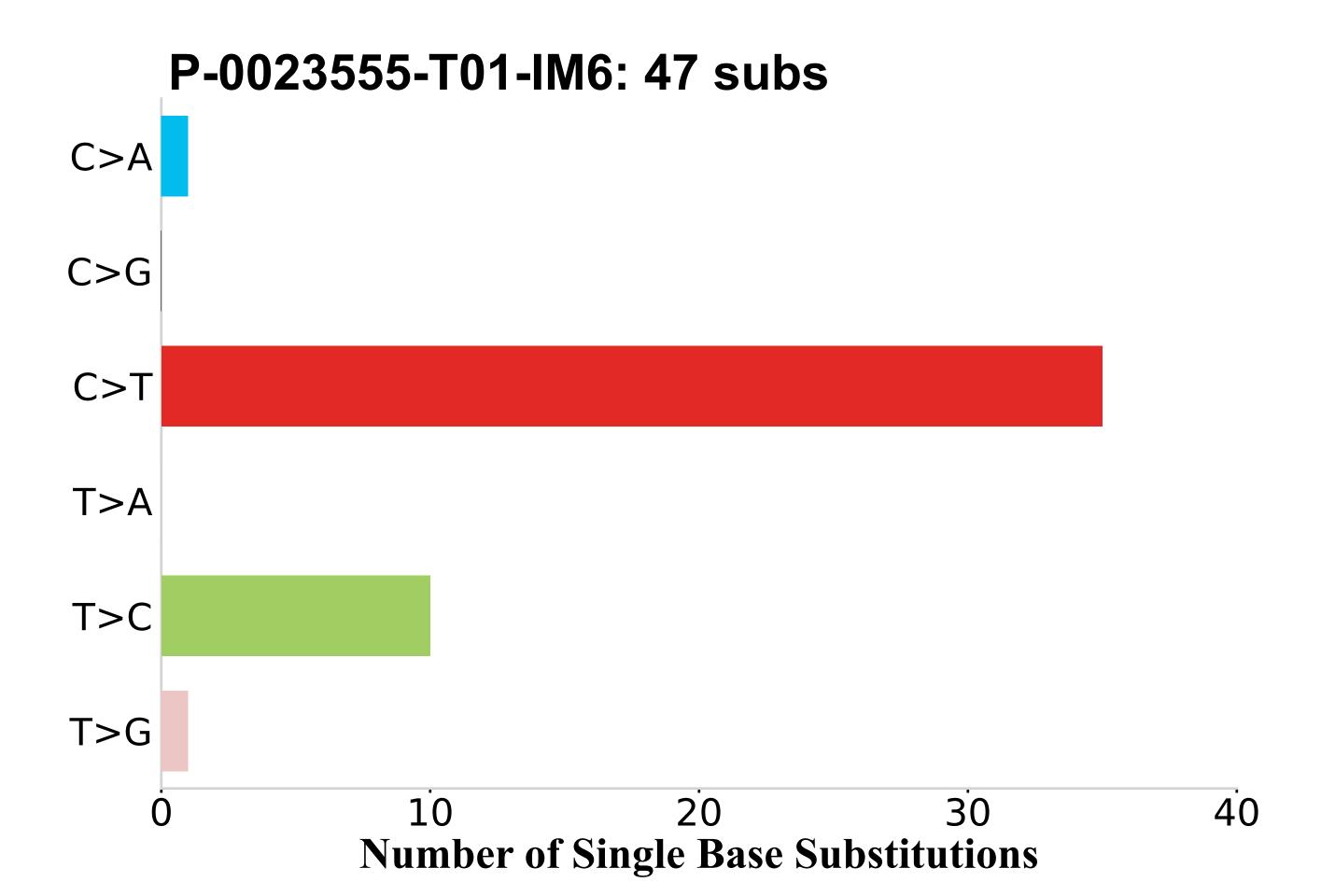


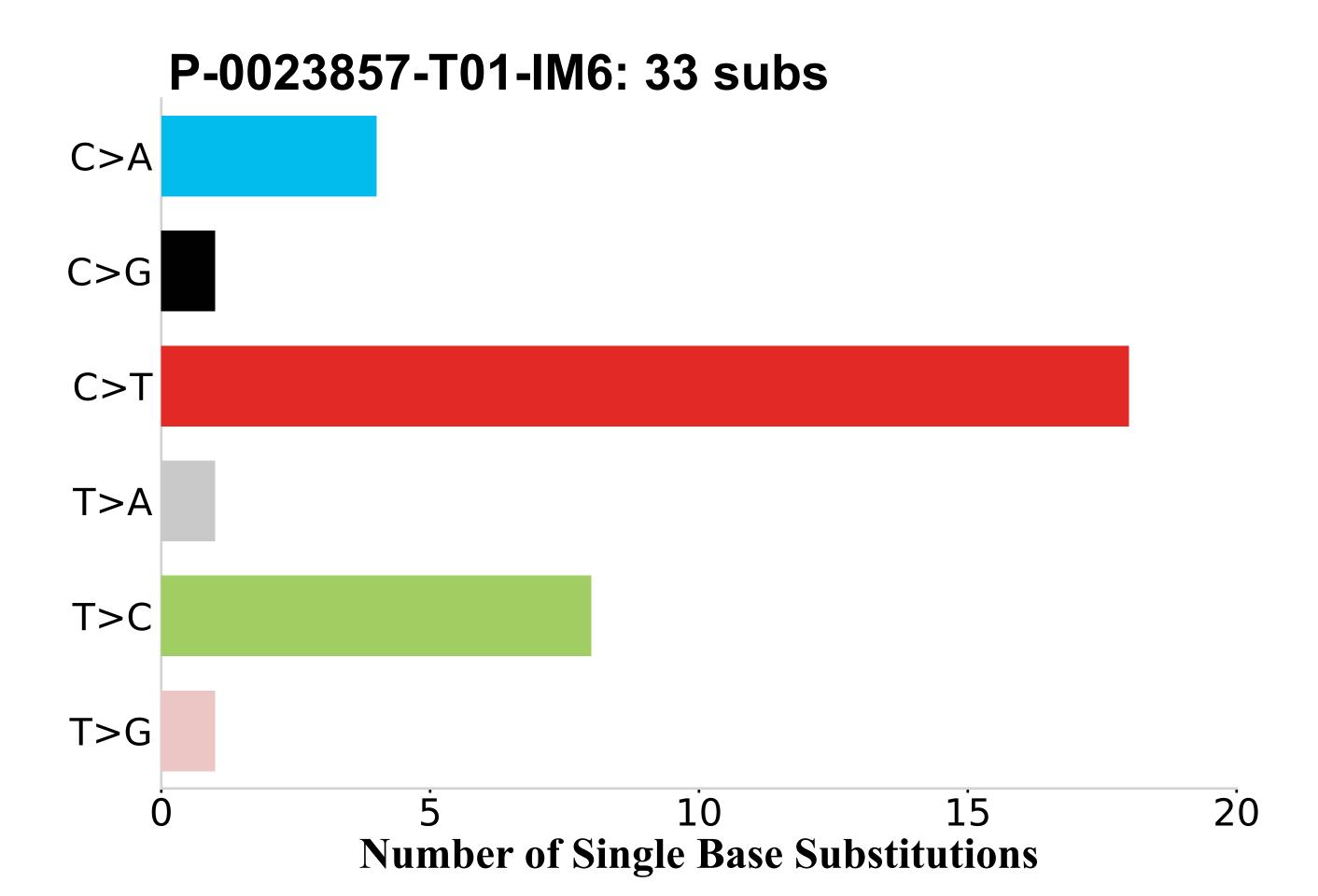


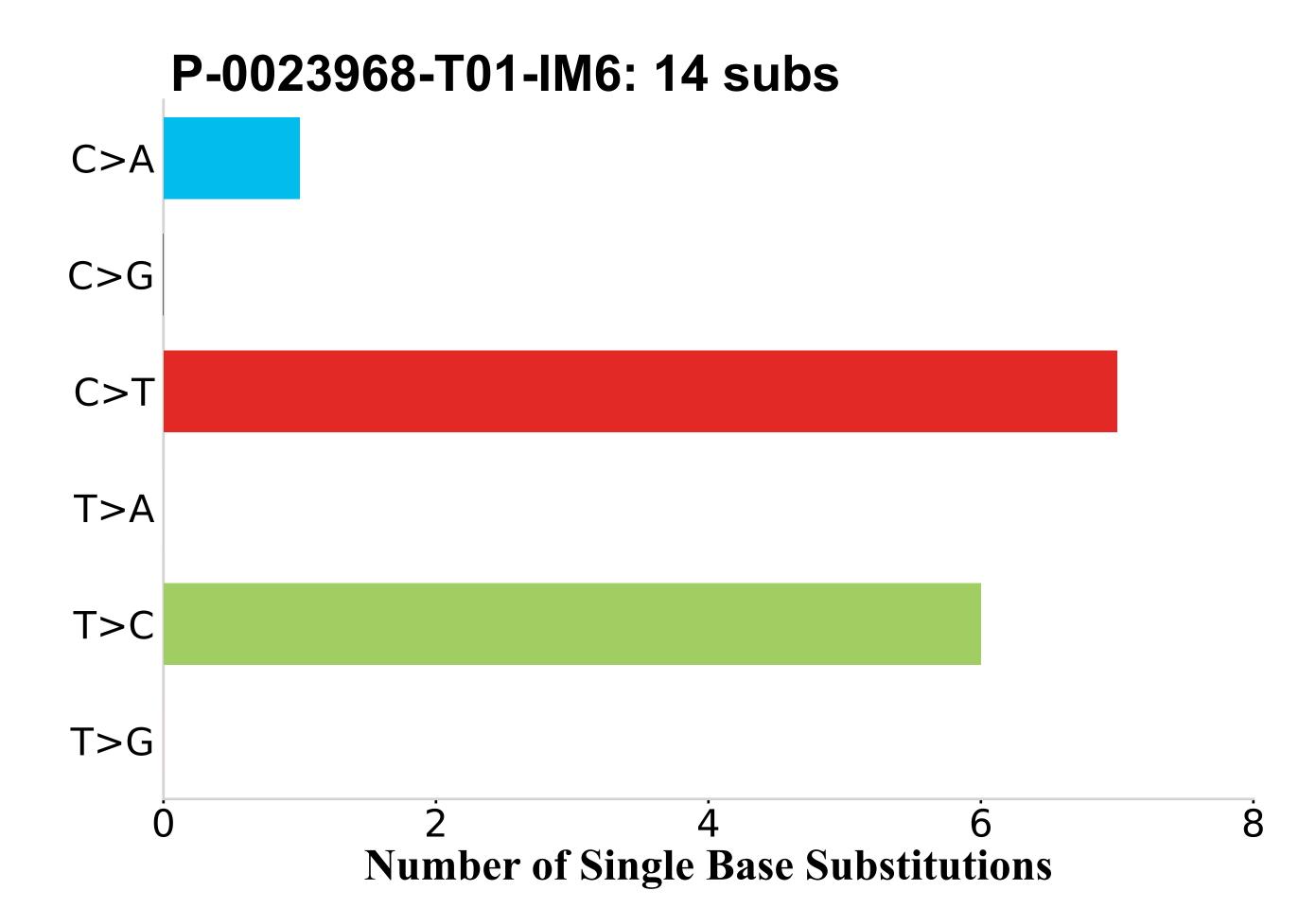


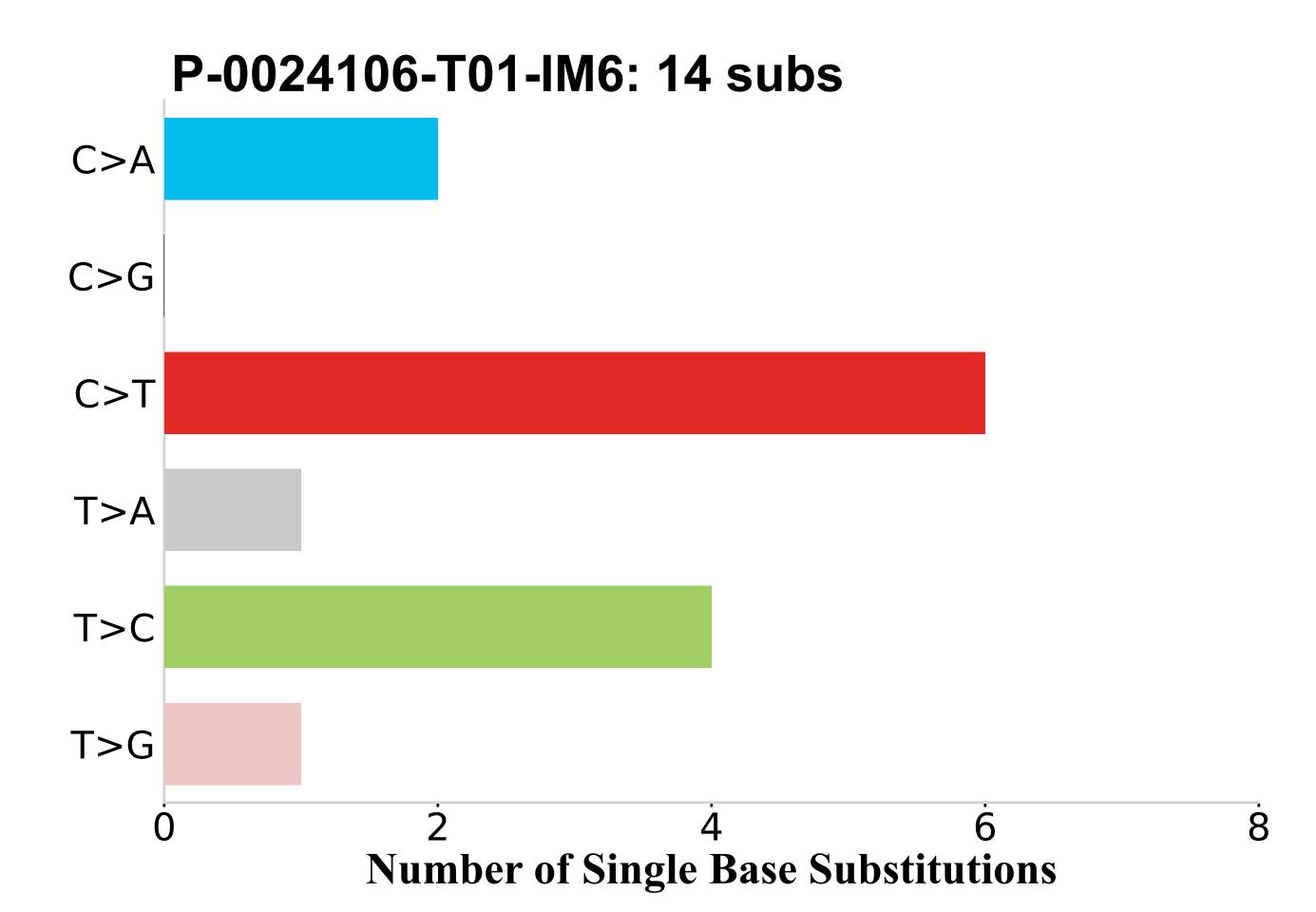
P-0023250-T01-IM6: 24 subs

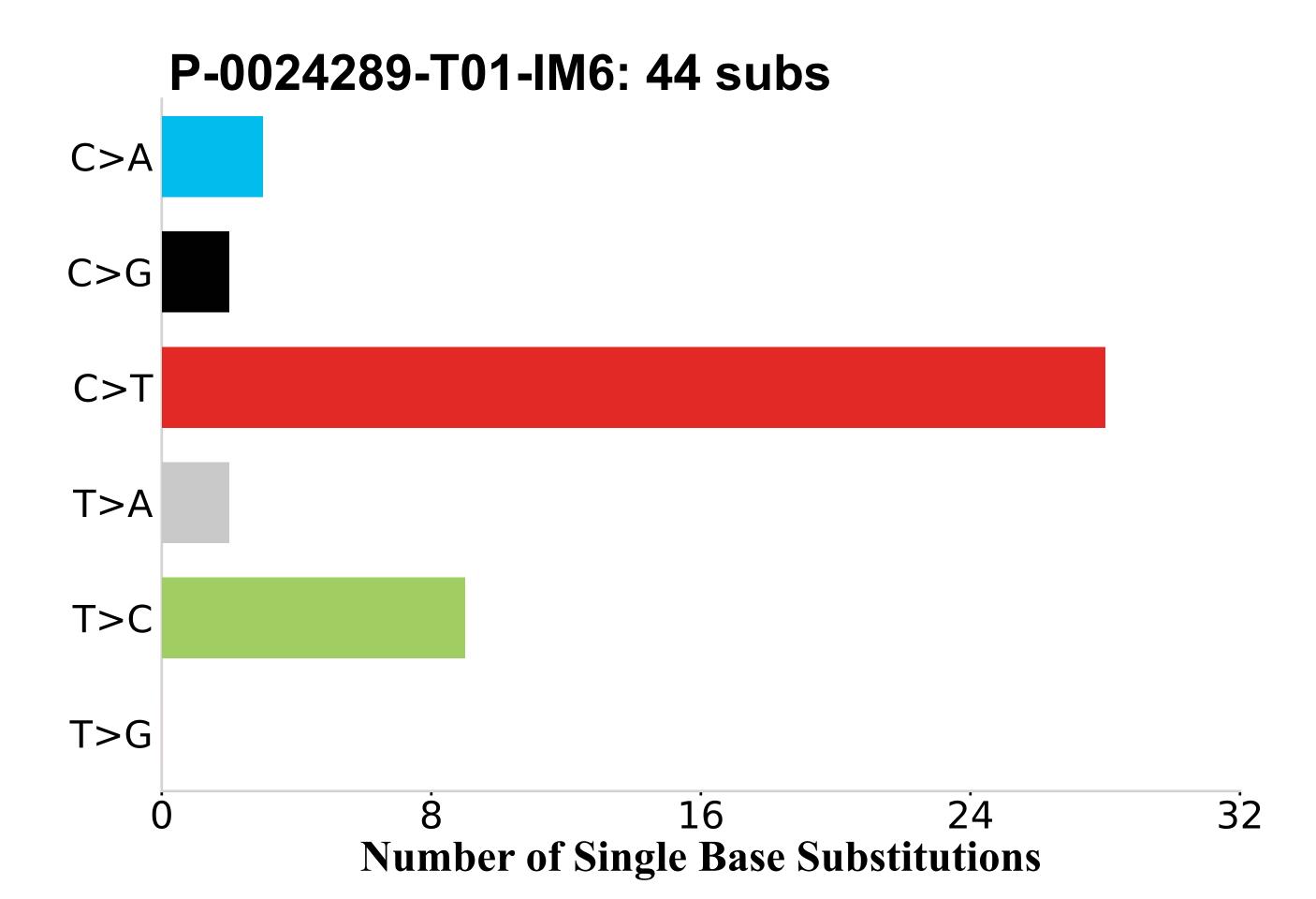




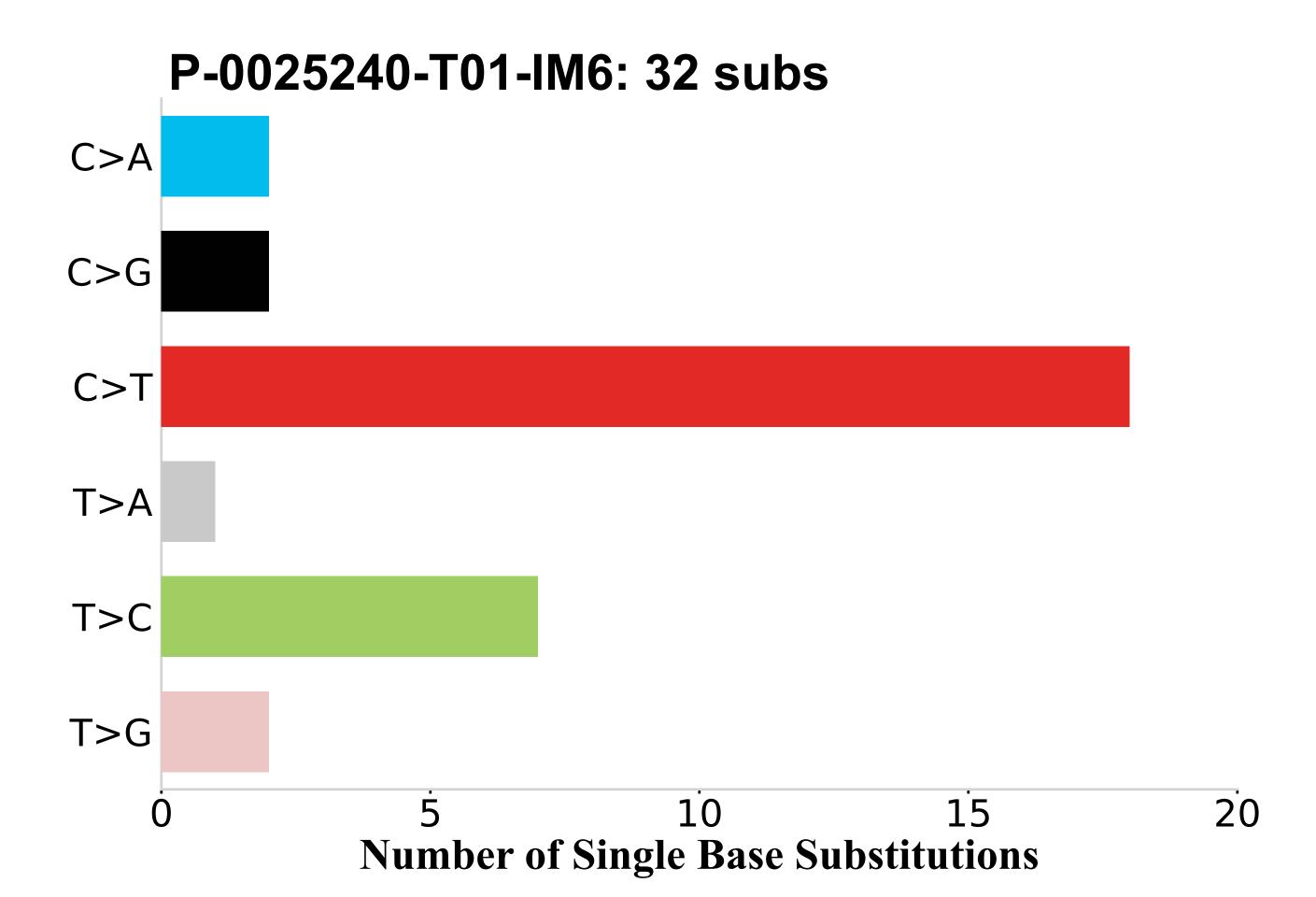


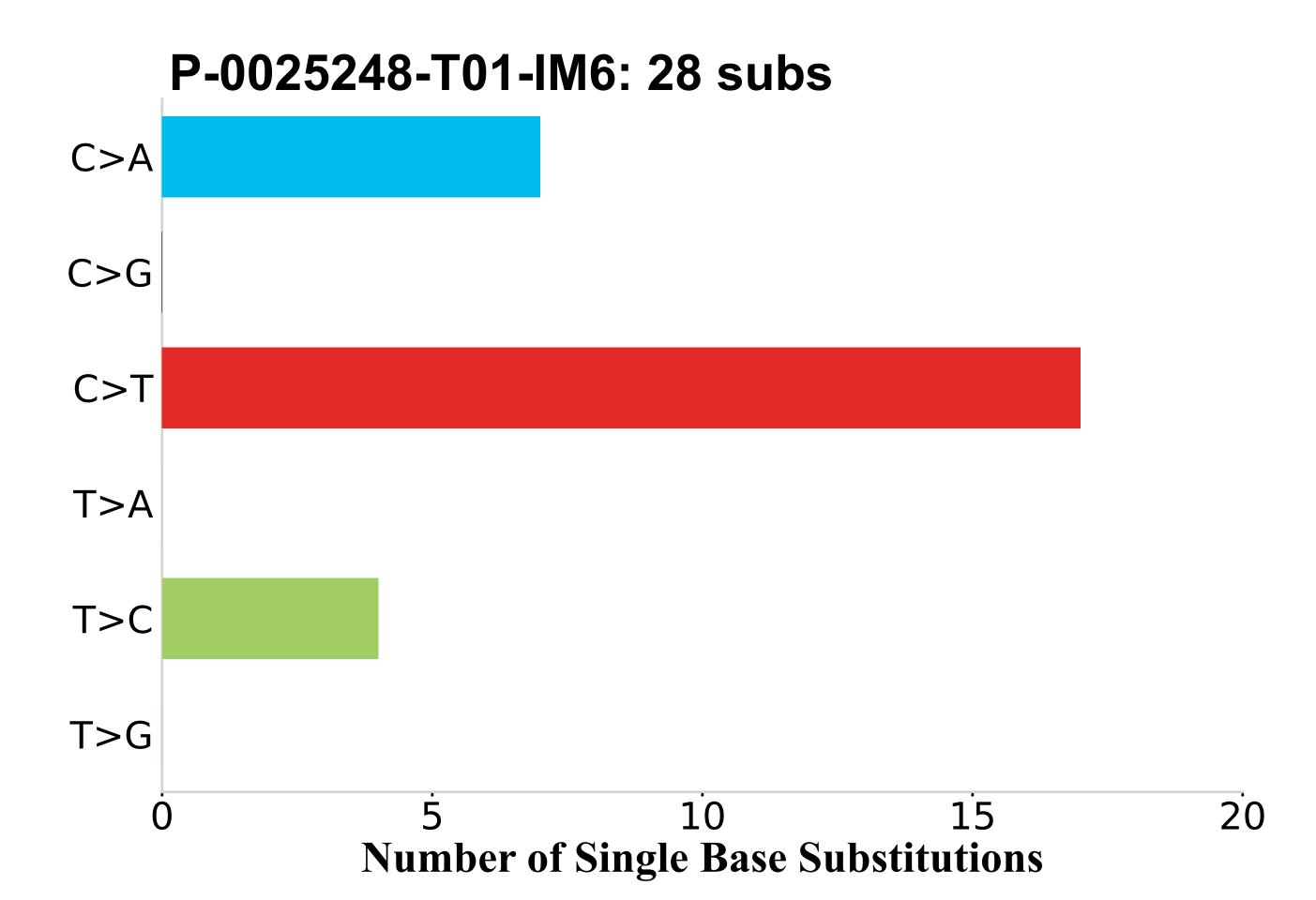


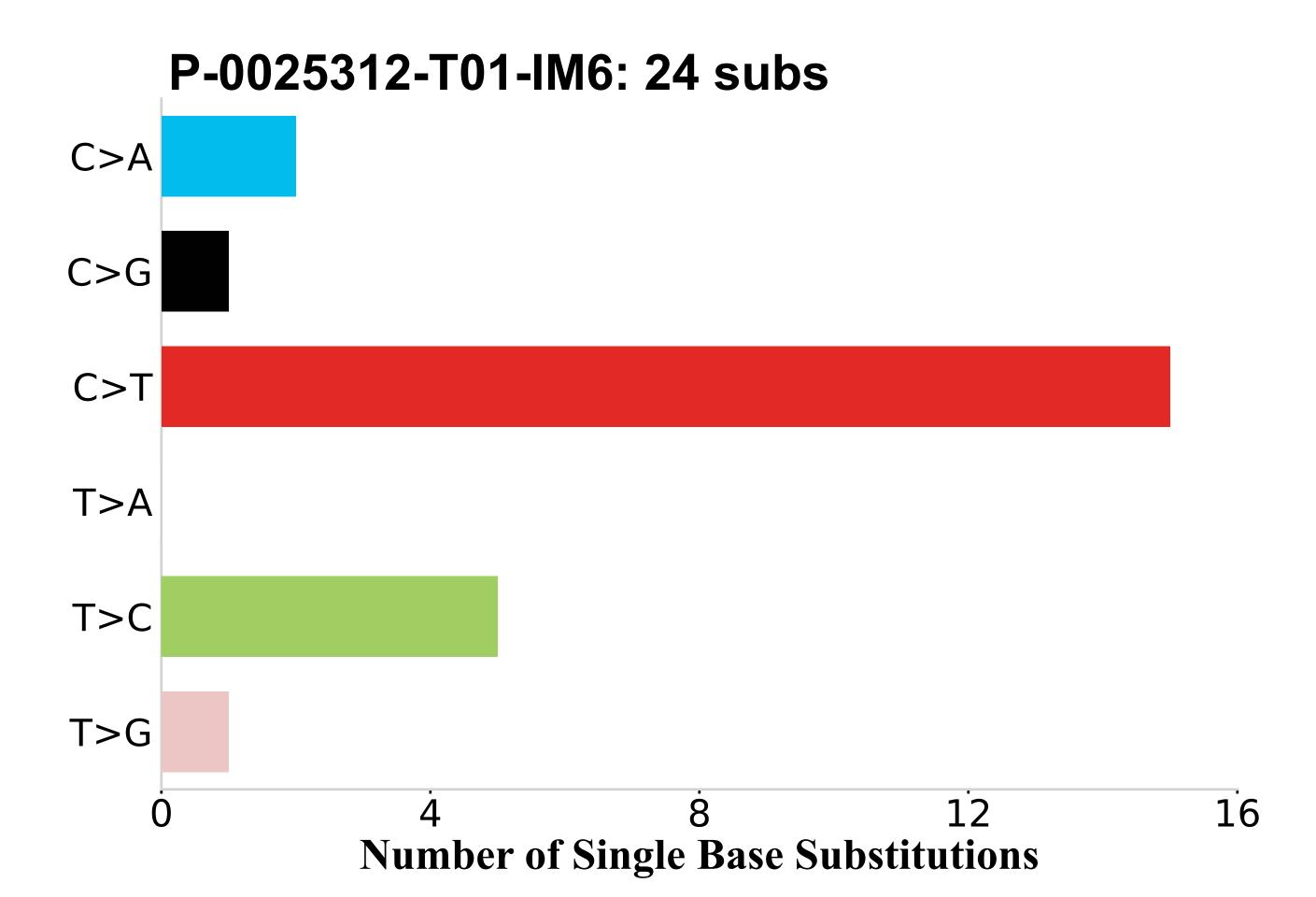


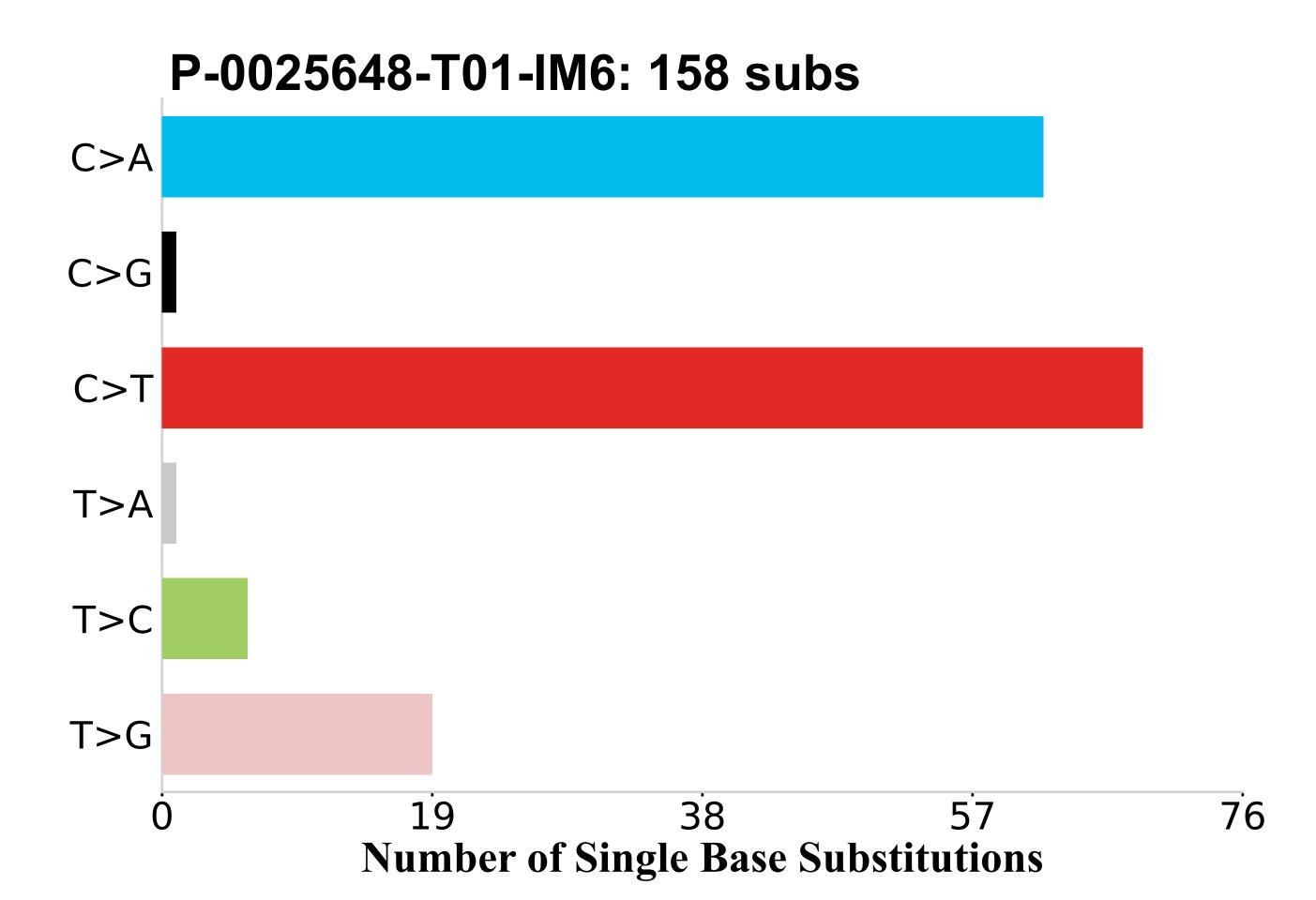


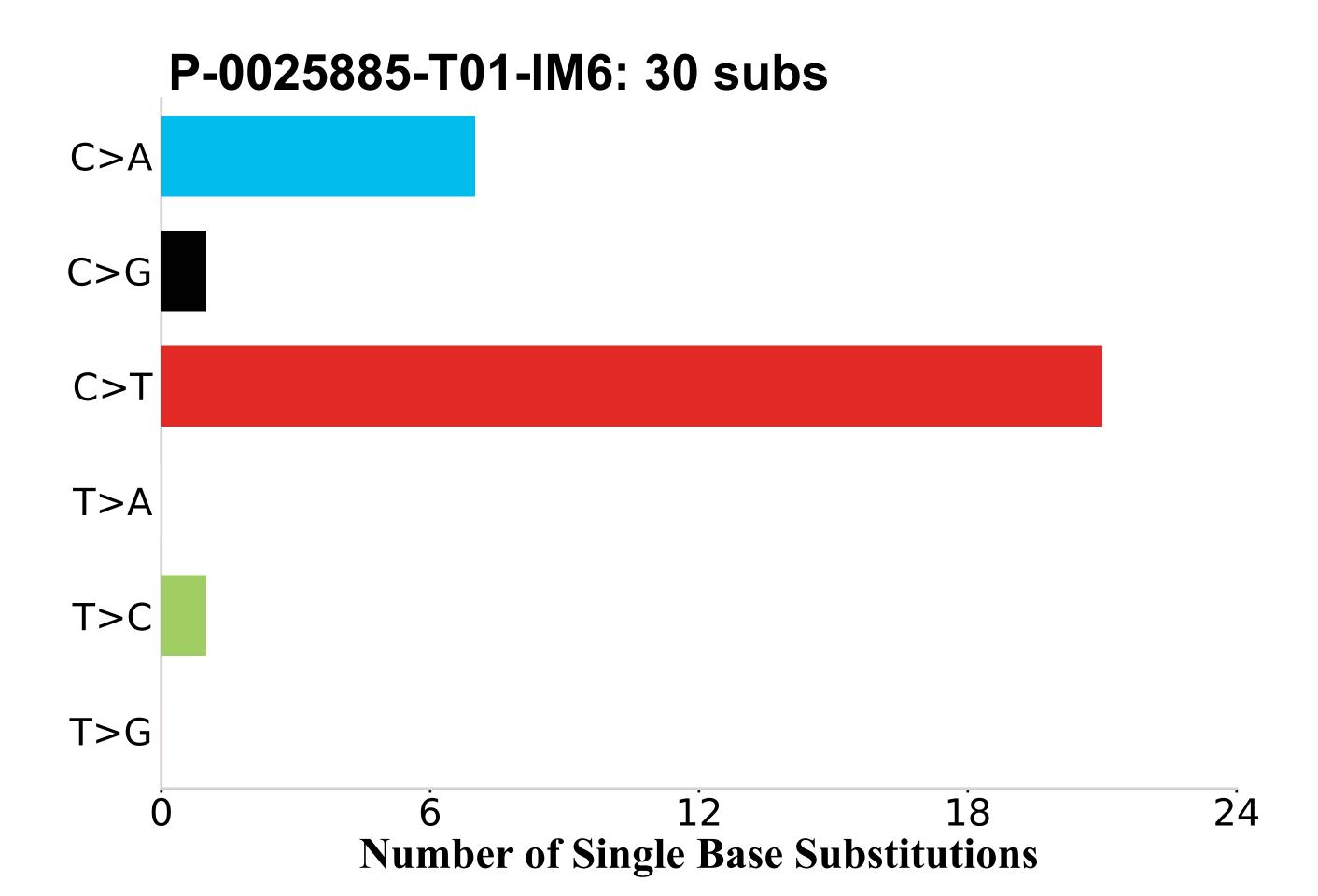
P-0025195-T01-IM6: 23 subs C>A C>G C>T T>AT>C T>G 16 12 **Number of Single Base Substitutions**

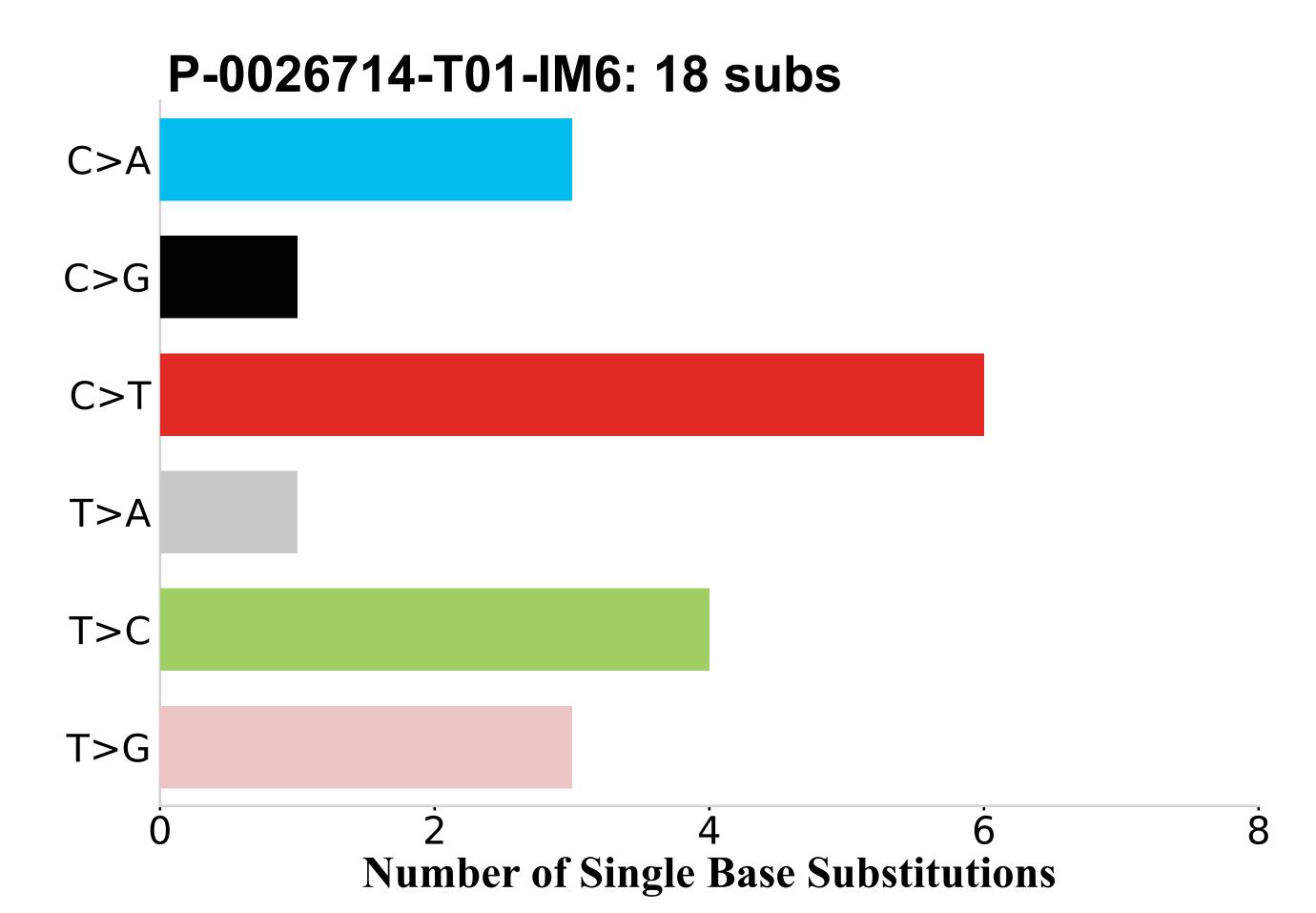


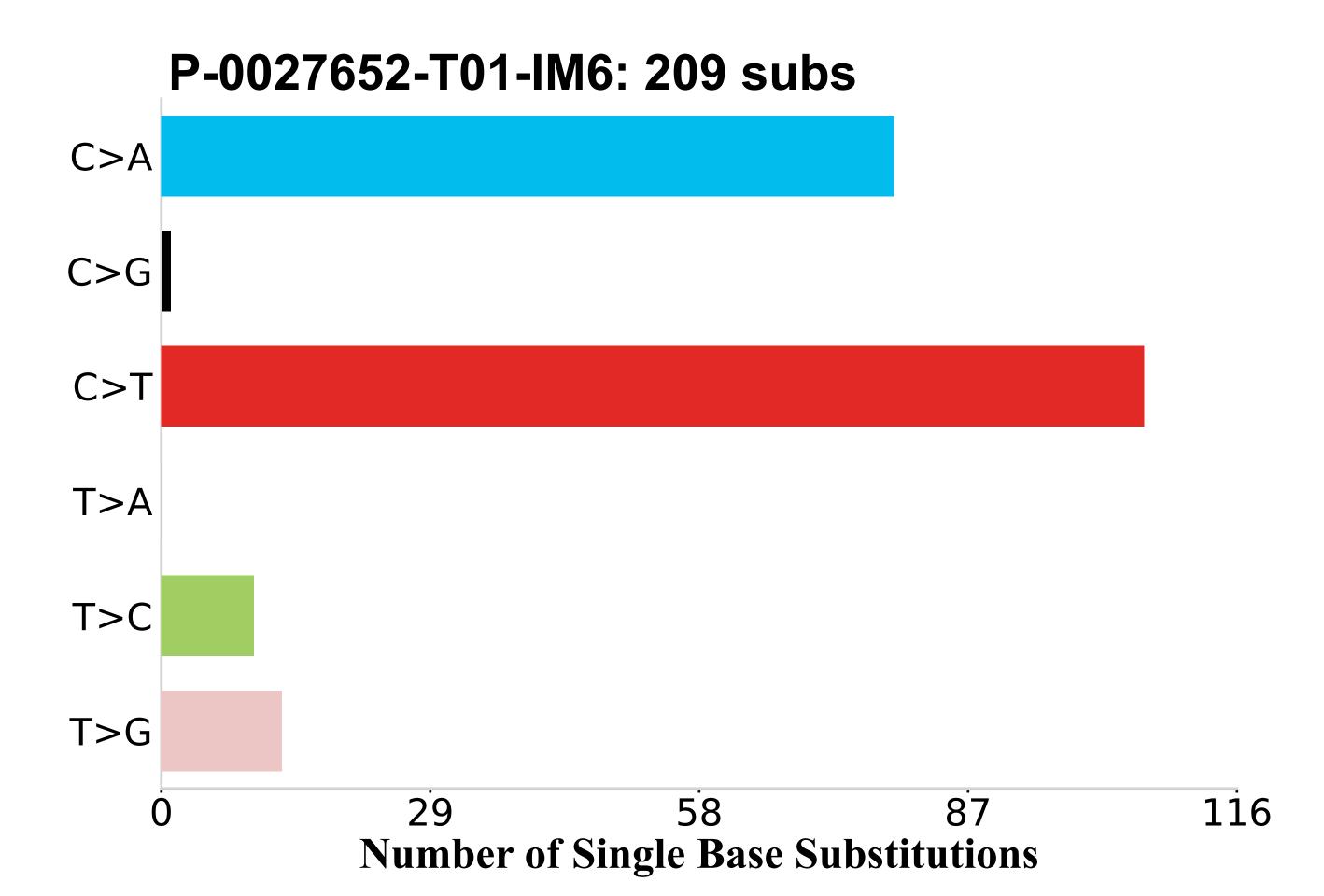


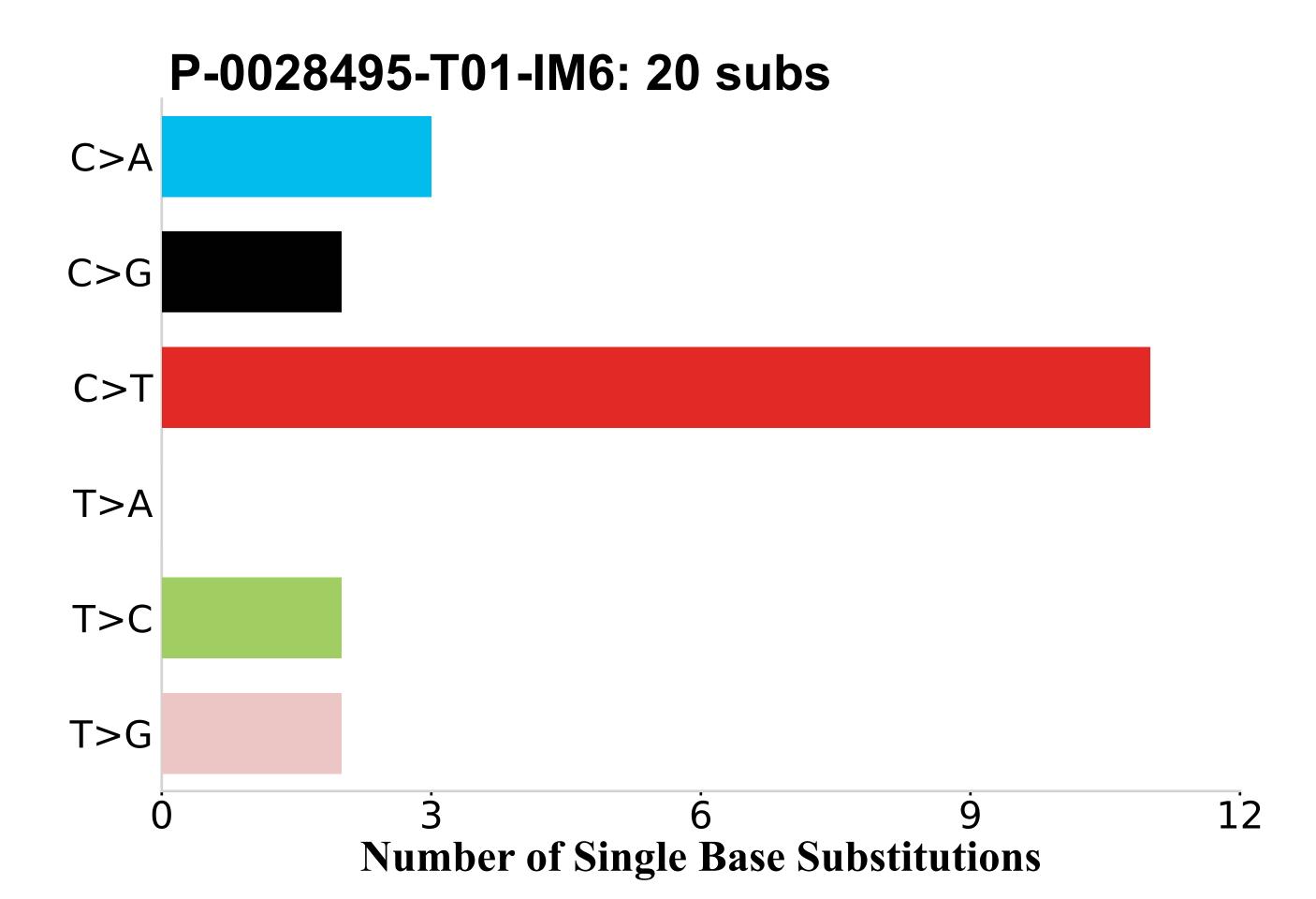


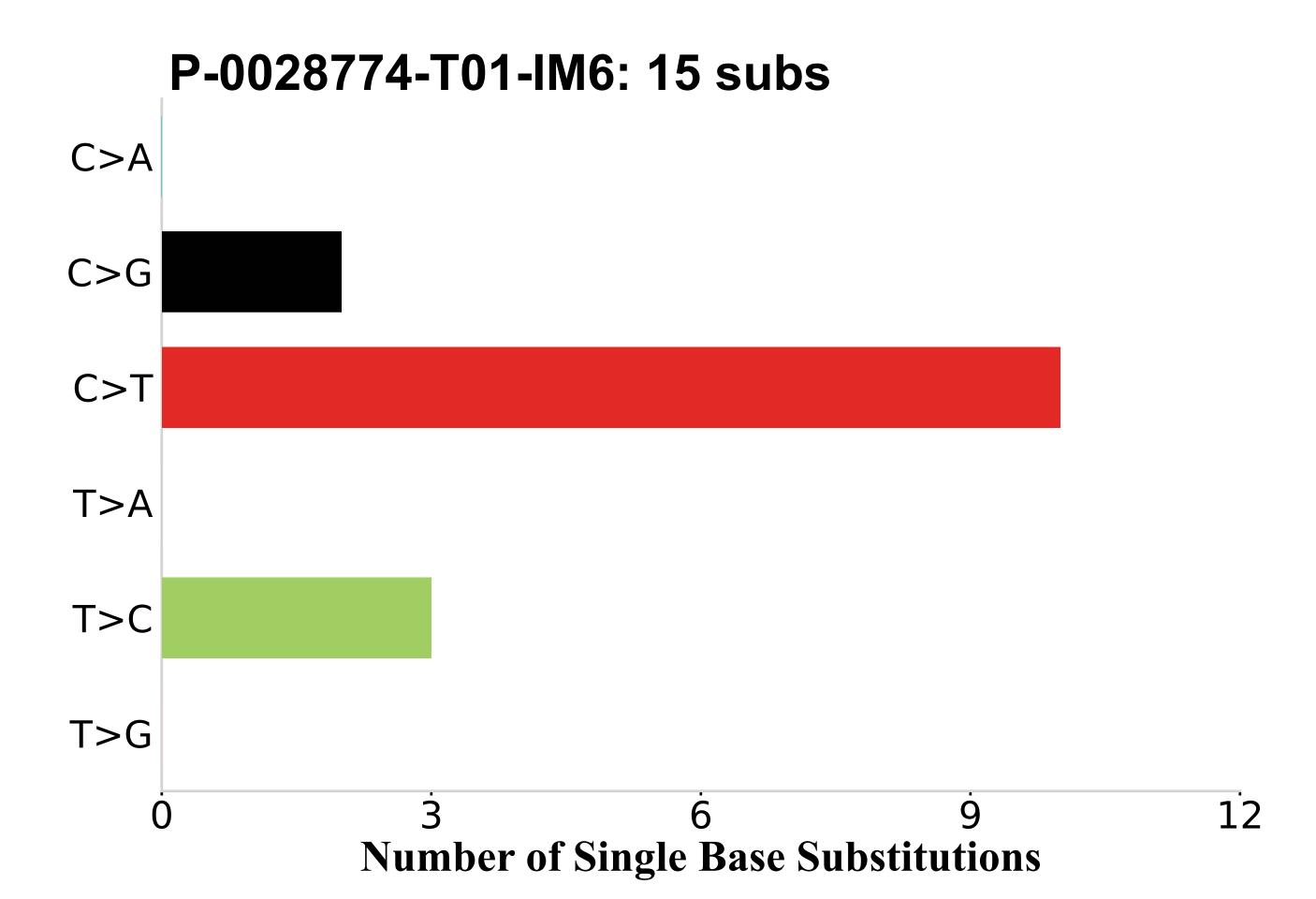


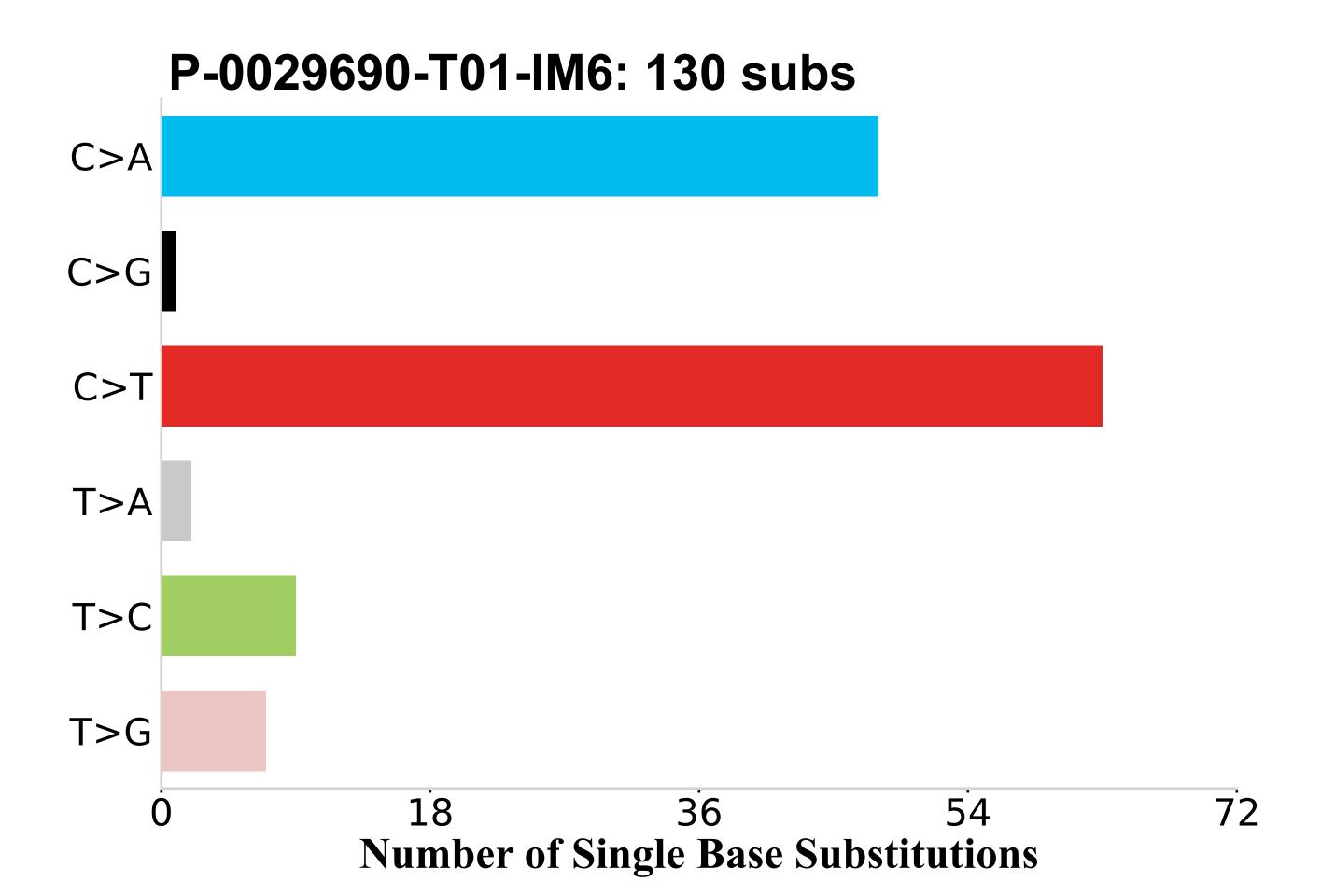


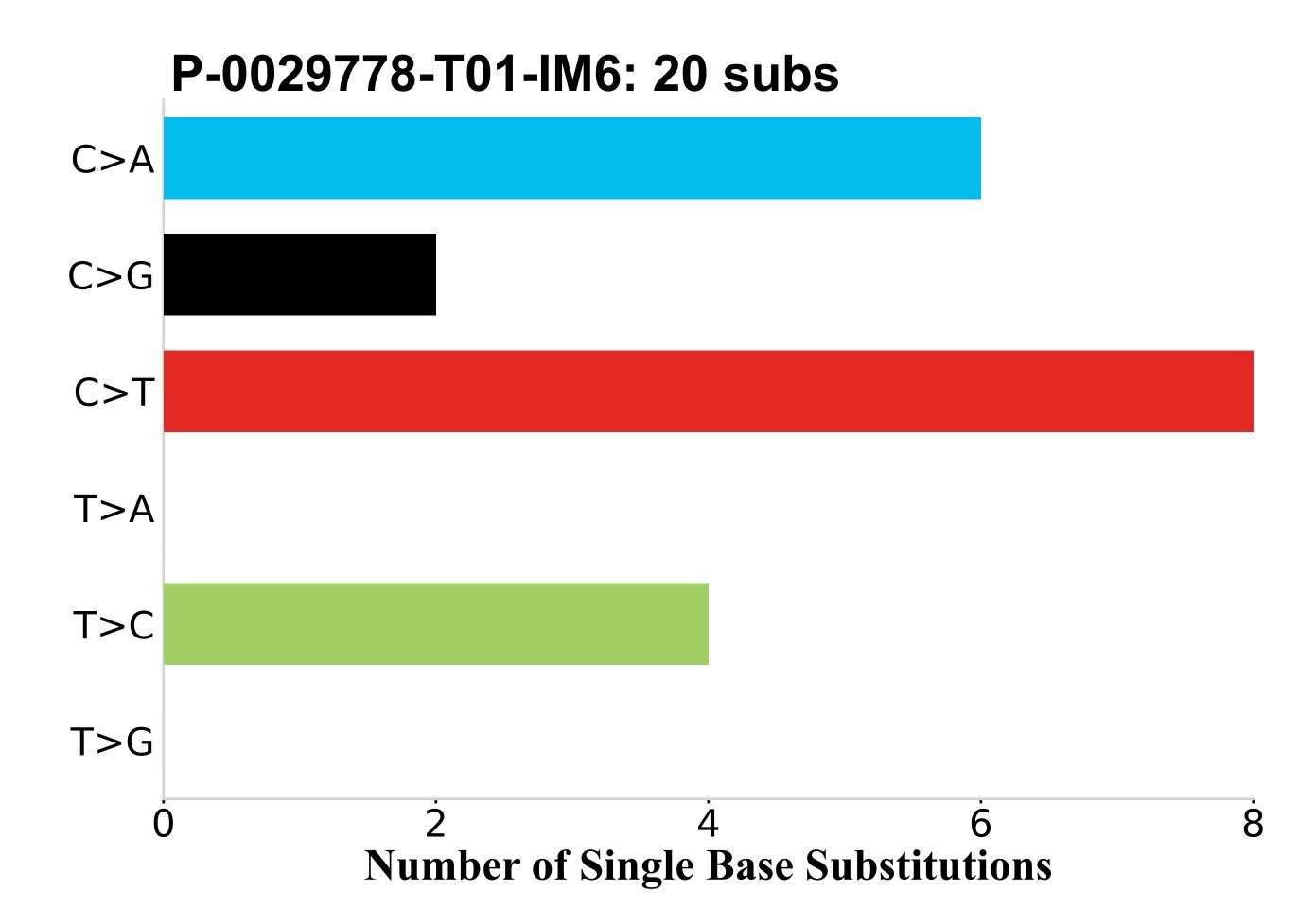








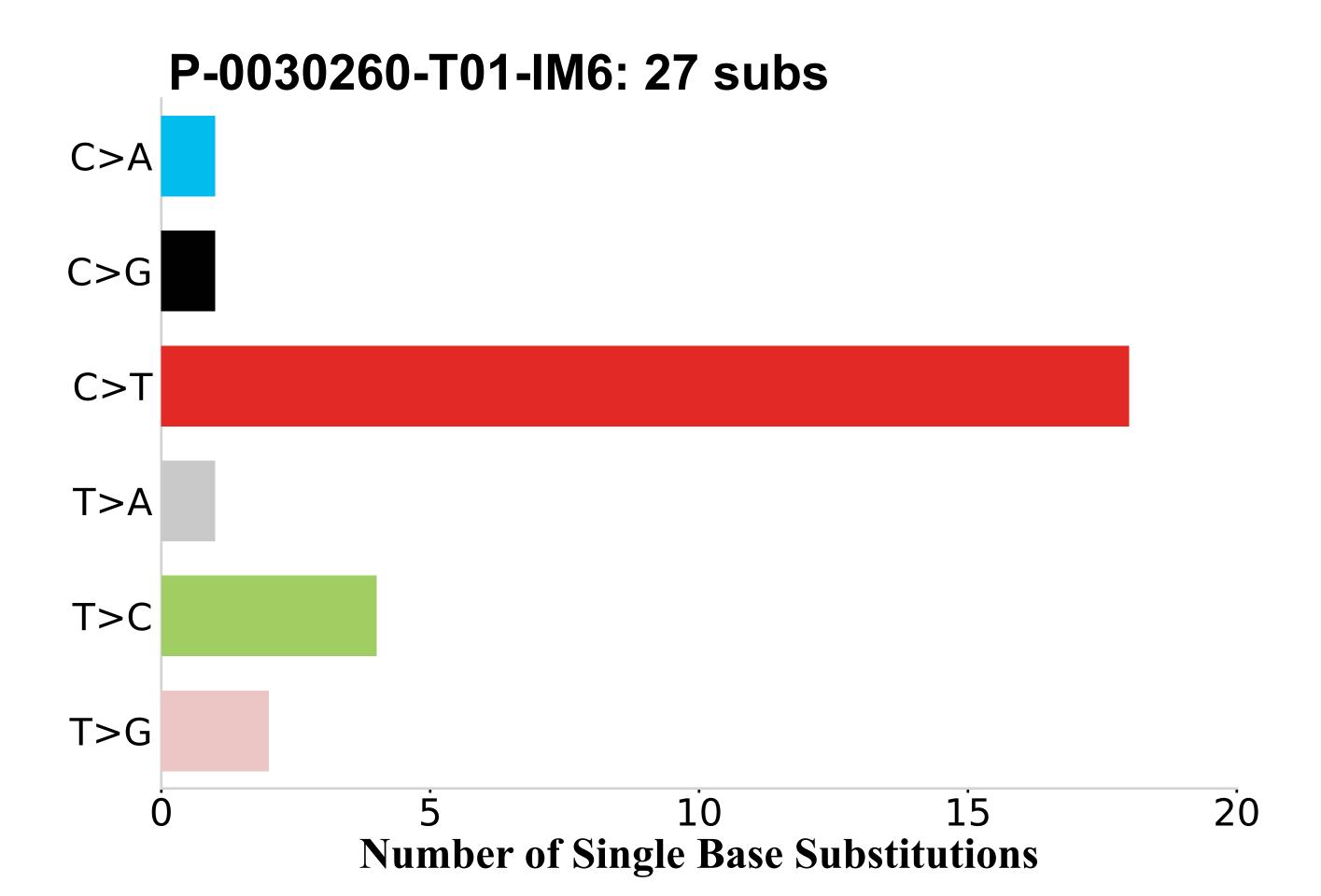




P-0029972-T01-IM6: 303 subs C>A C>G C>T T>AT>C T>G



152

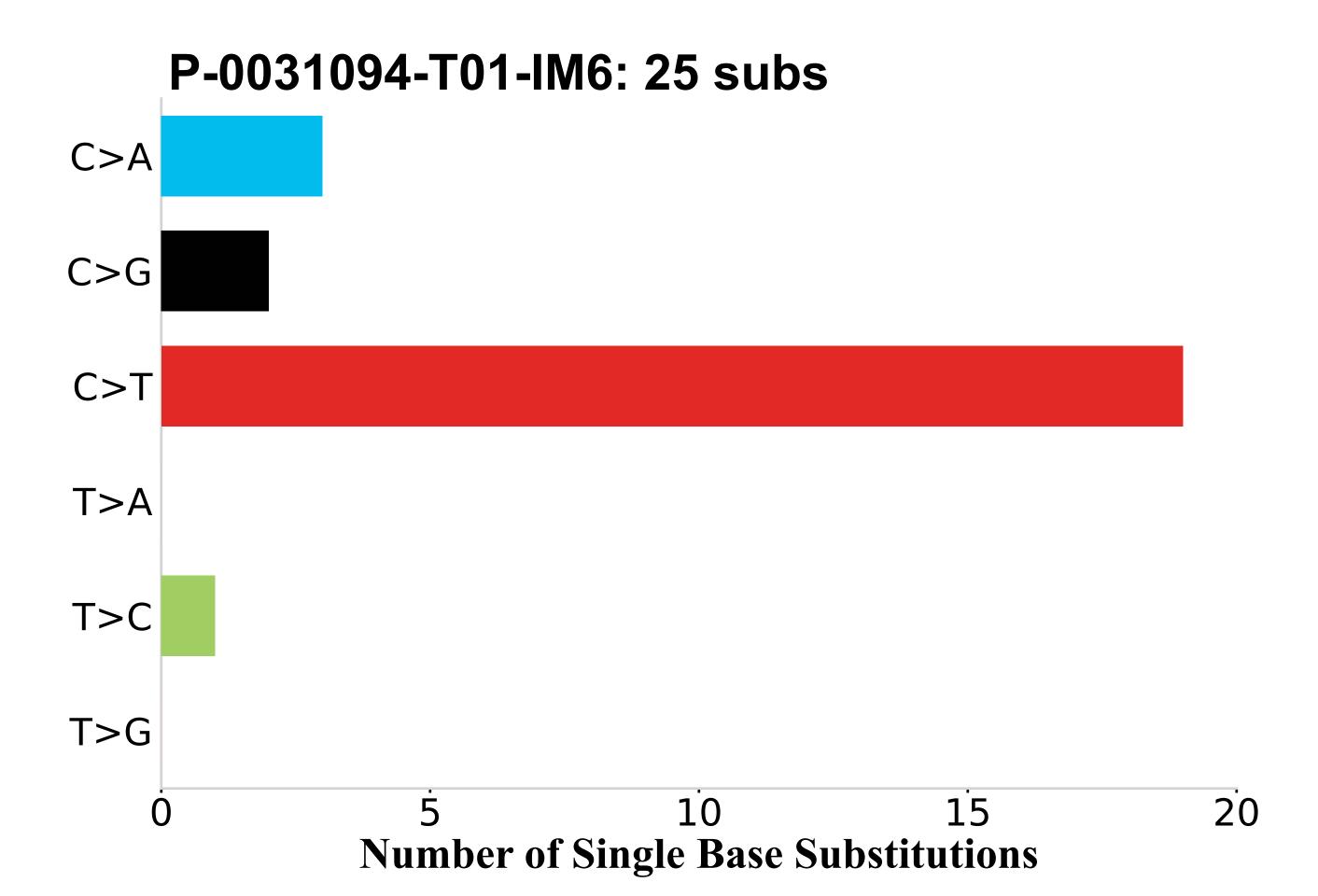


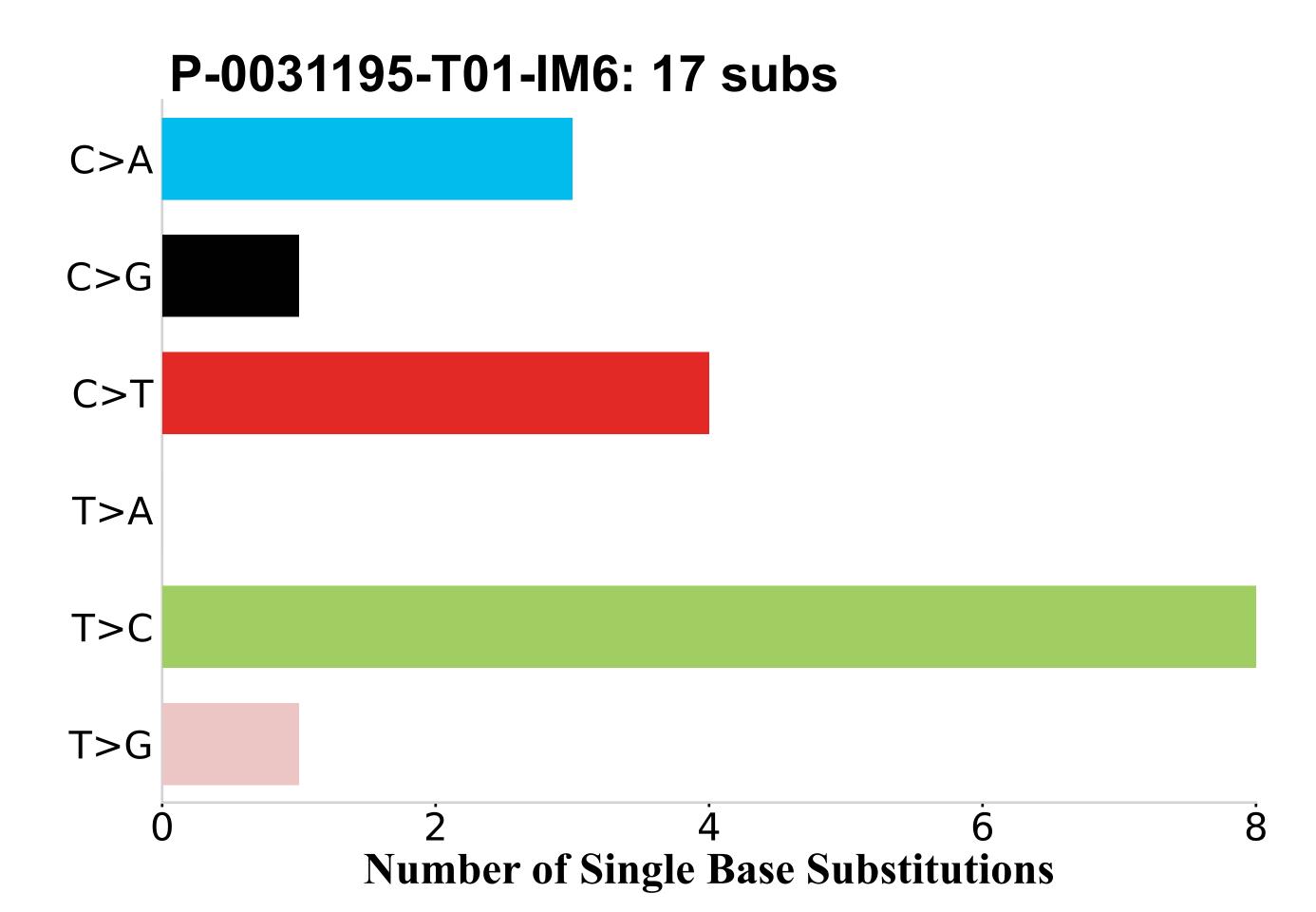
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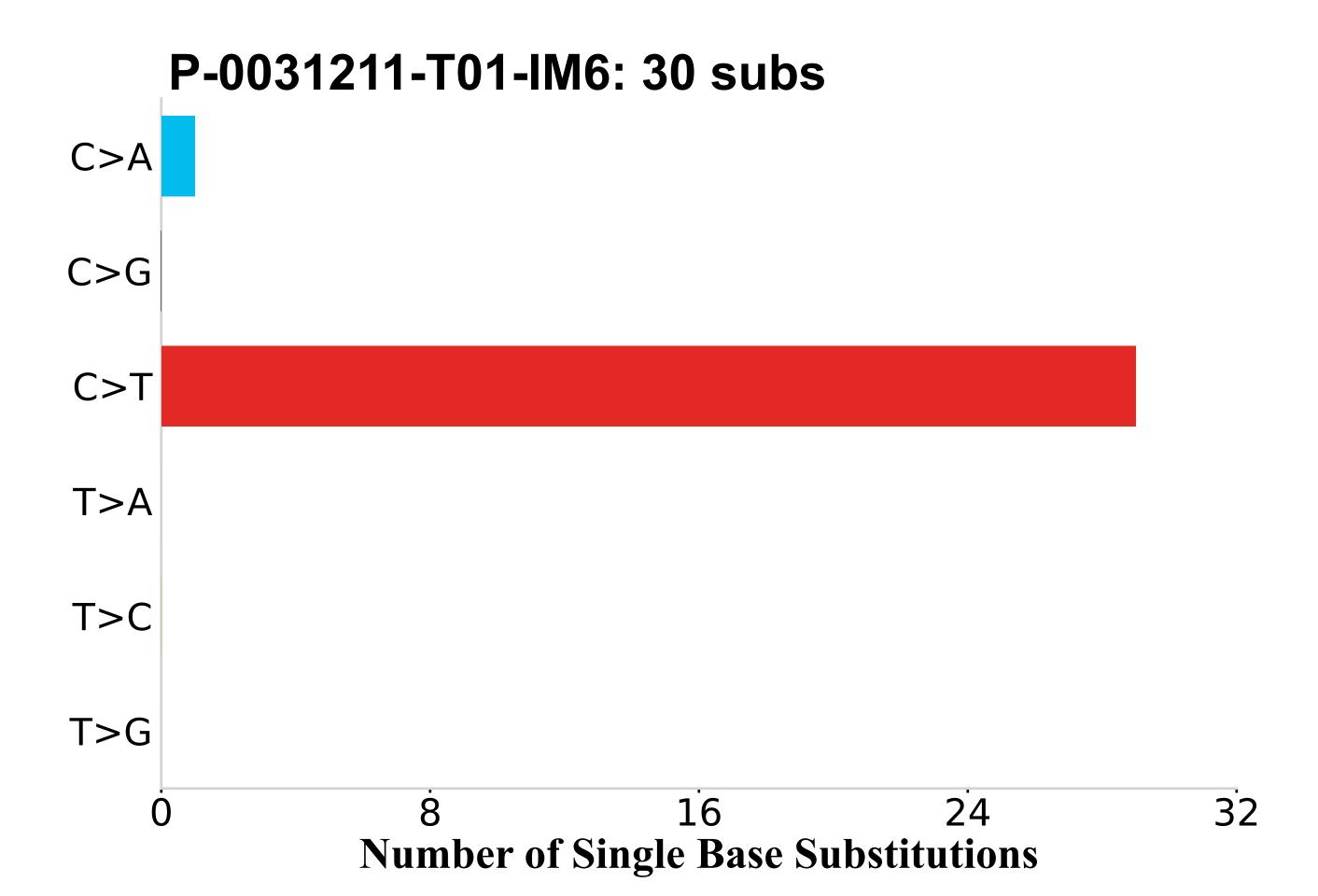
Number of Single Base Substitutions

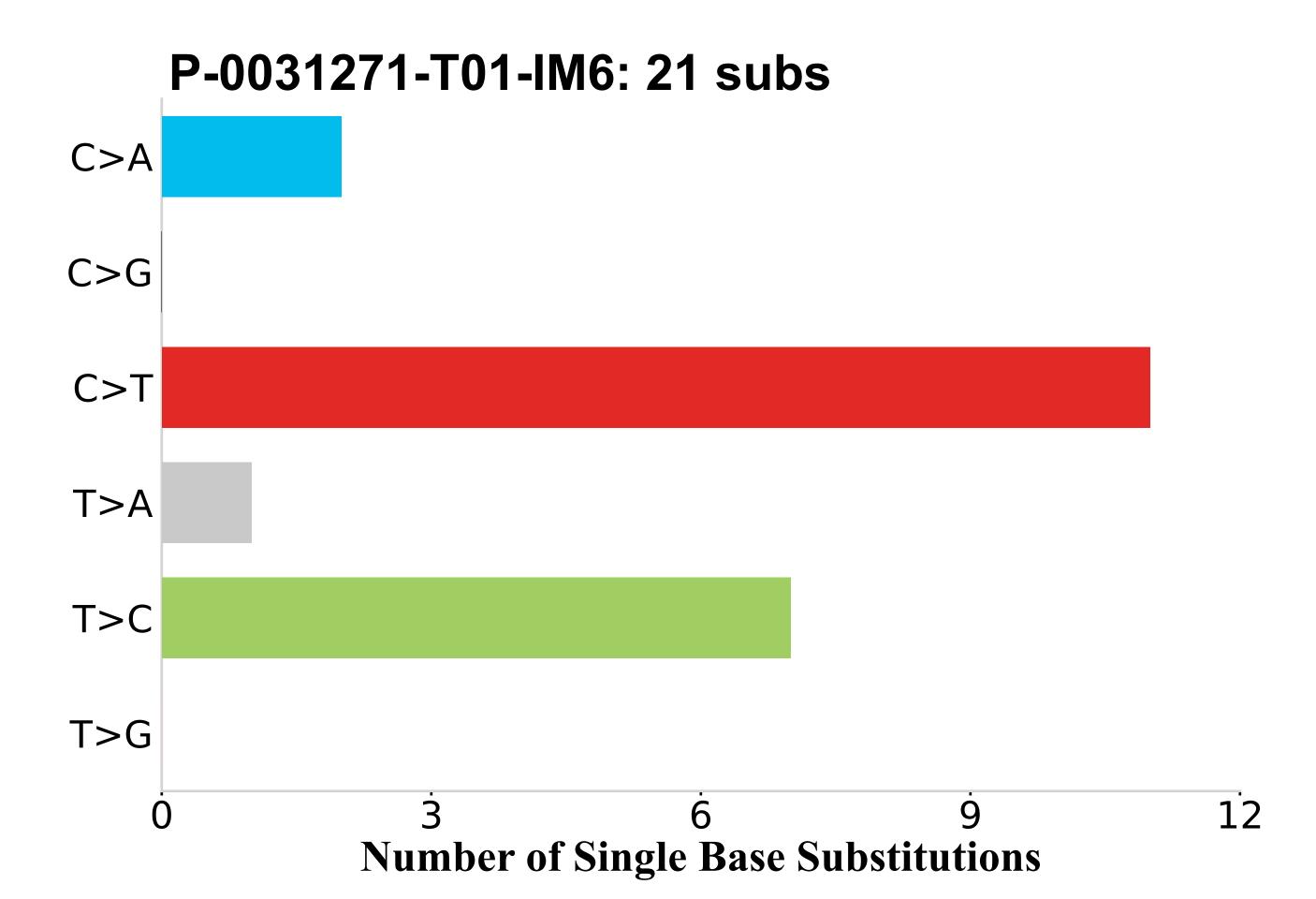
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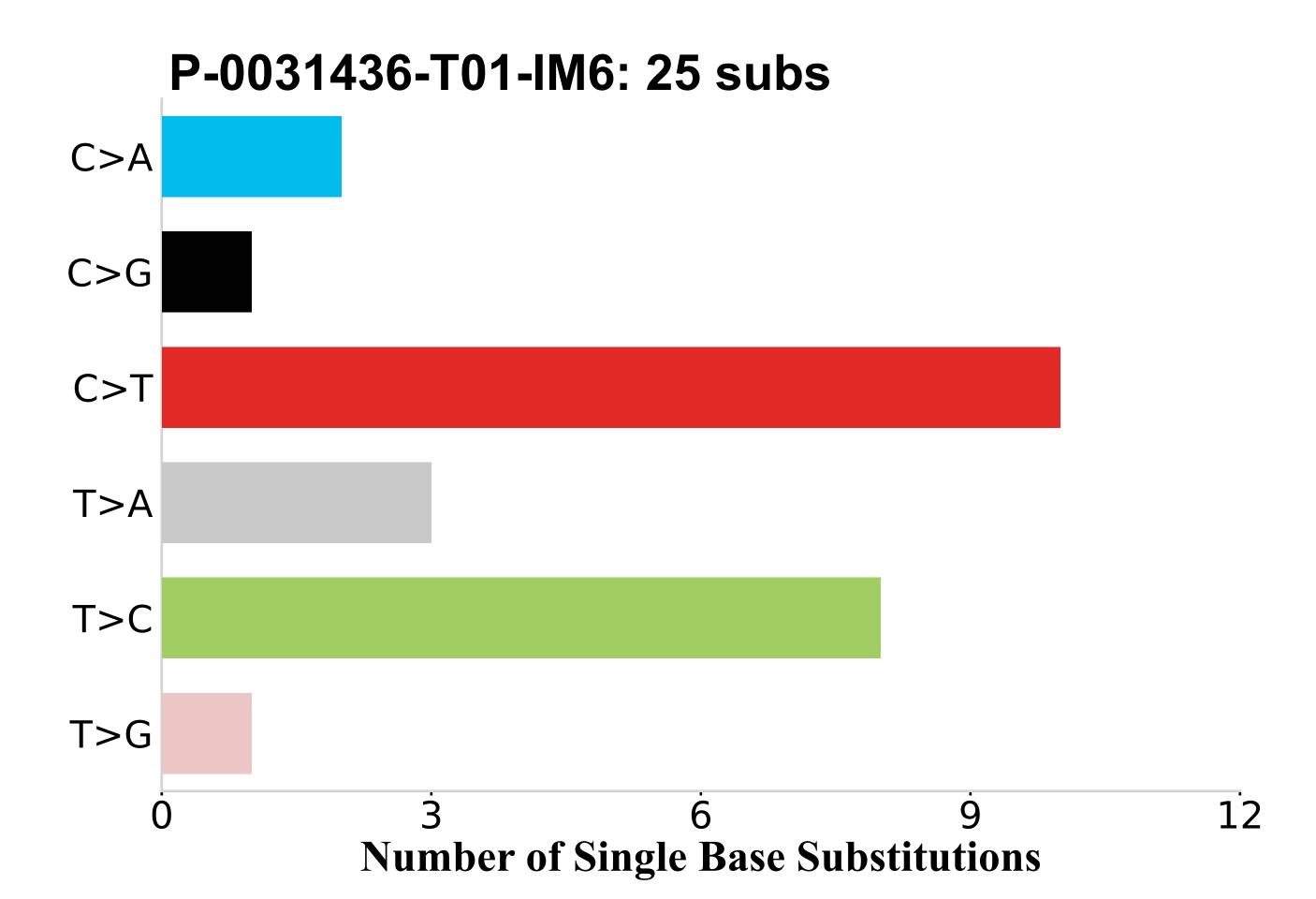
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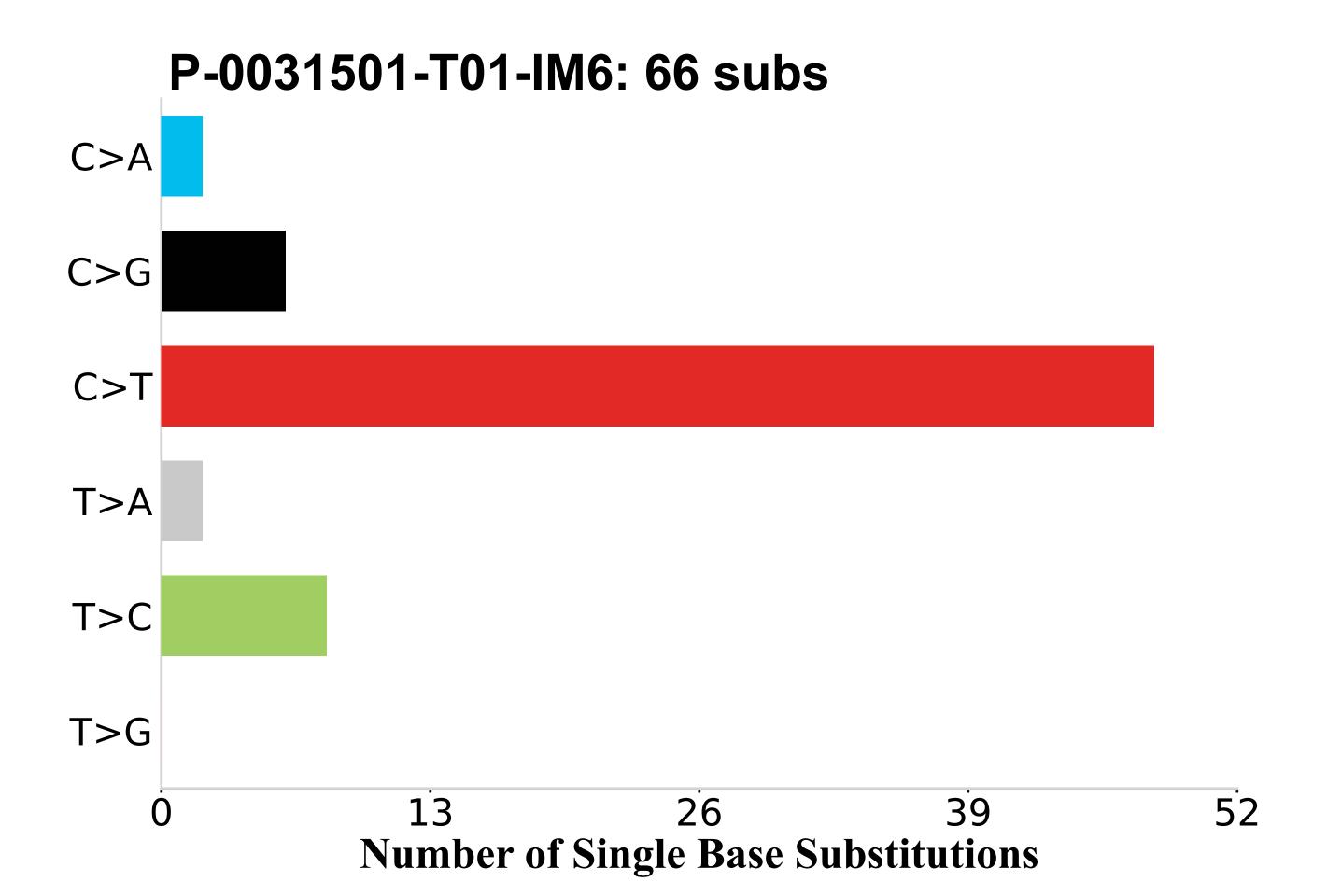


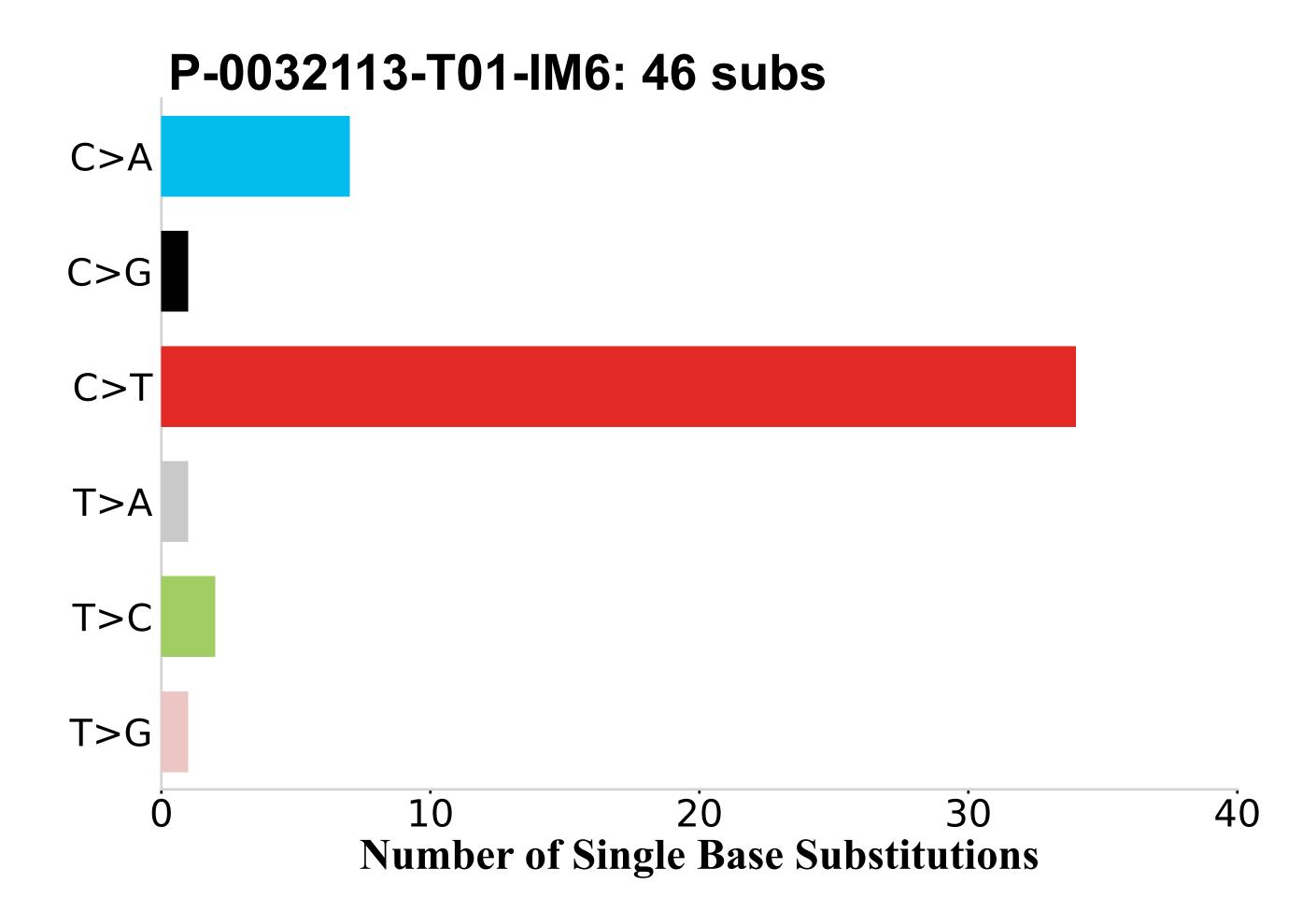


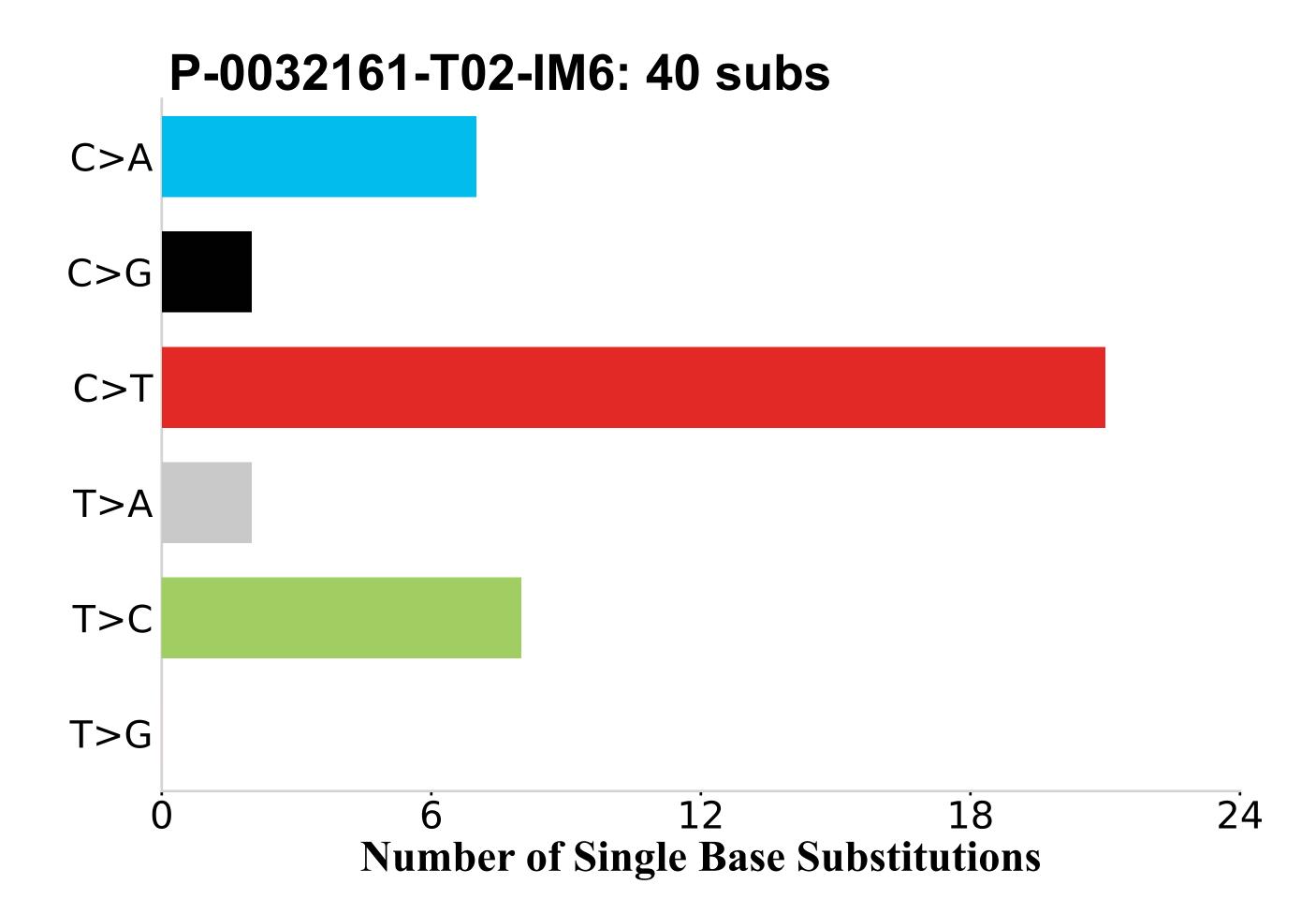


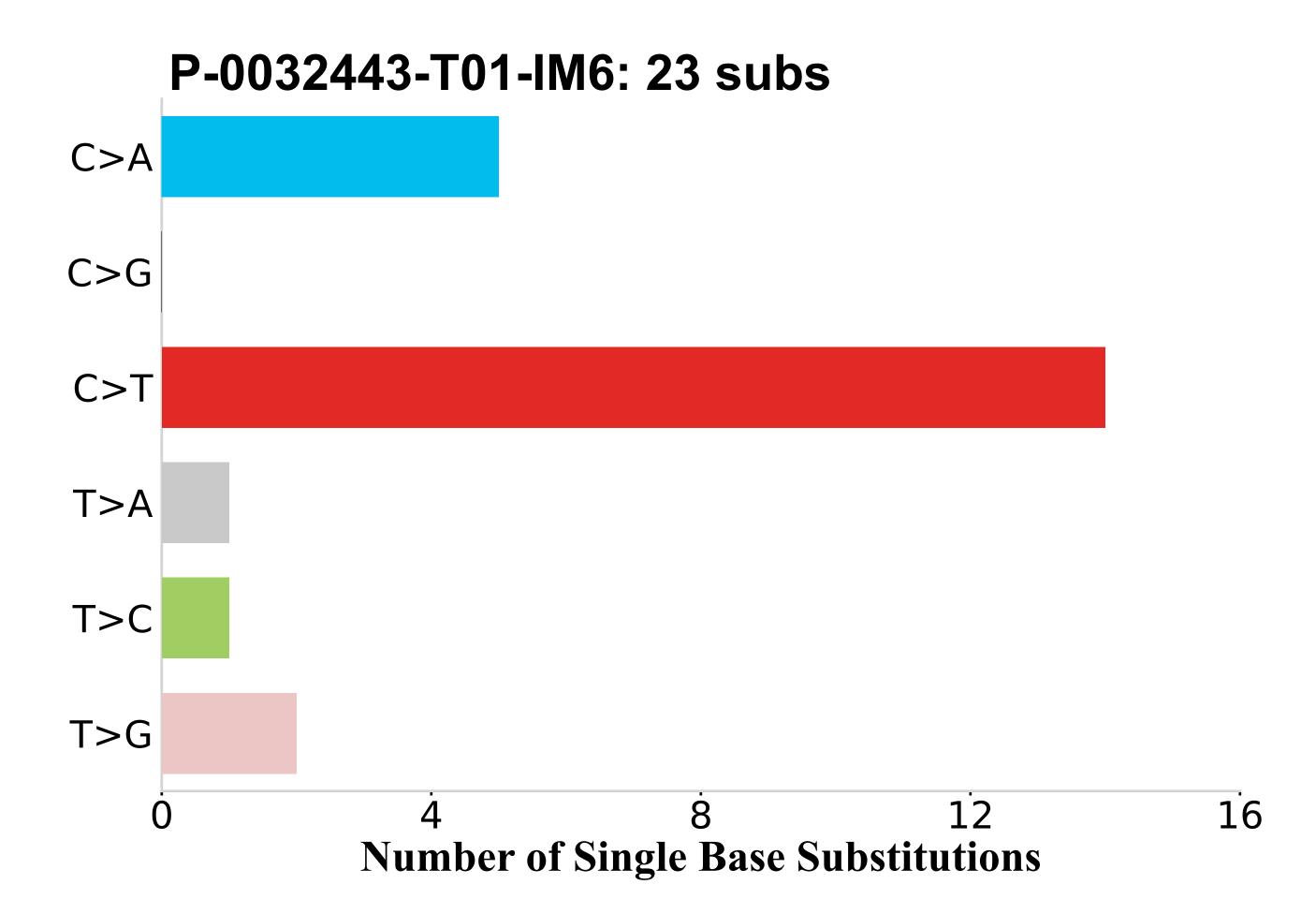


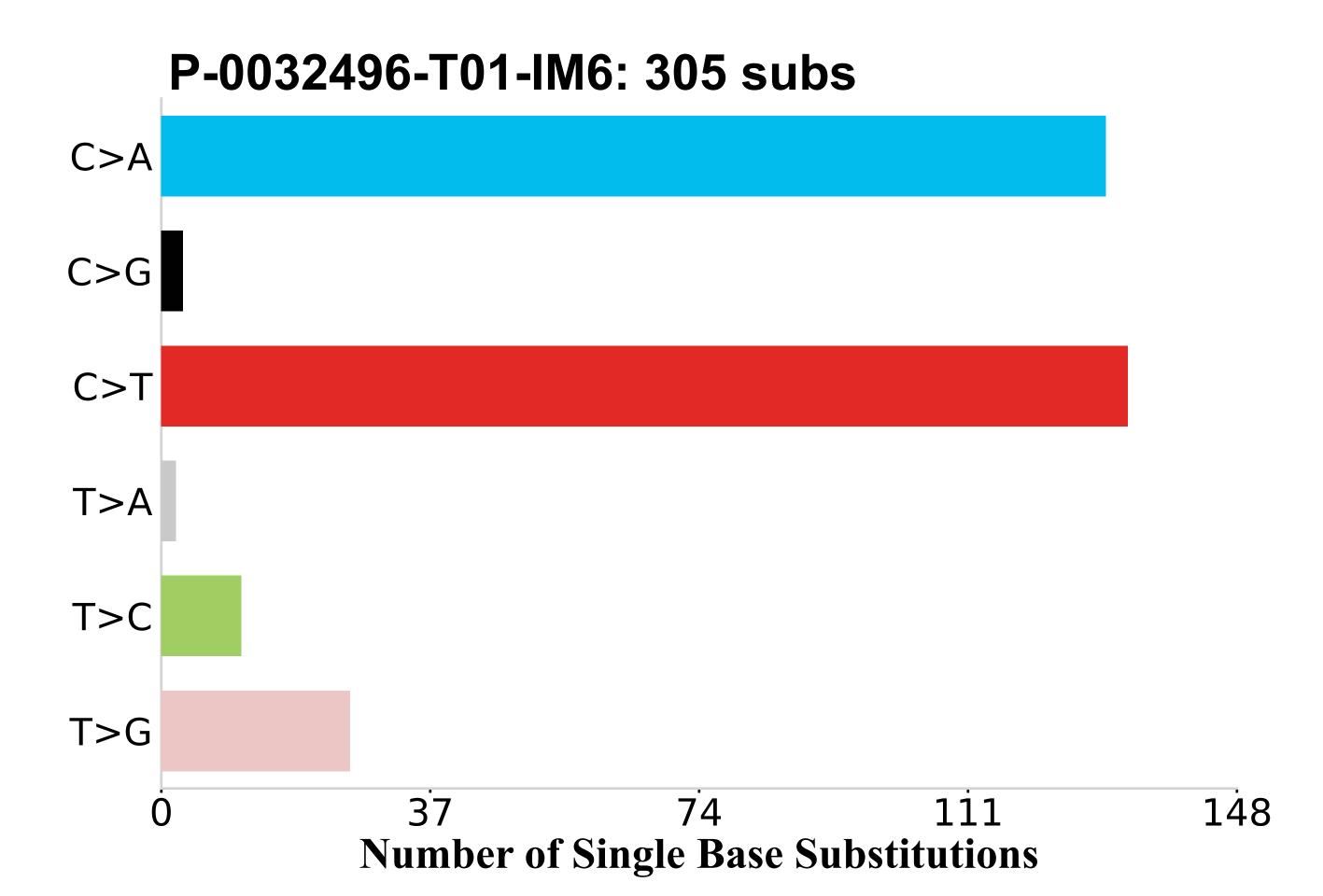


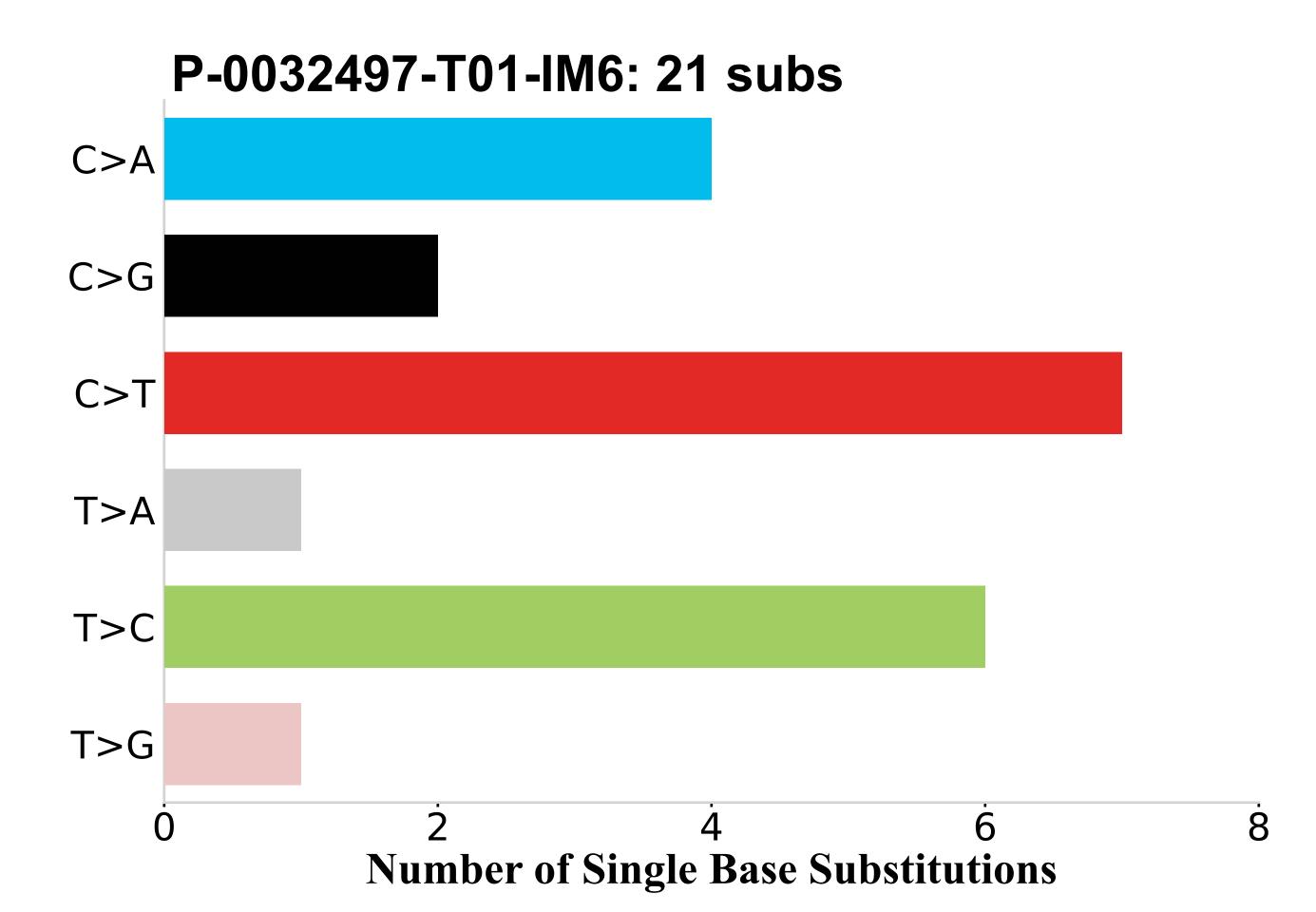


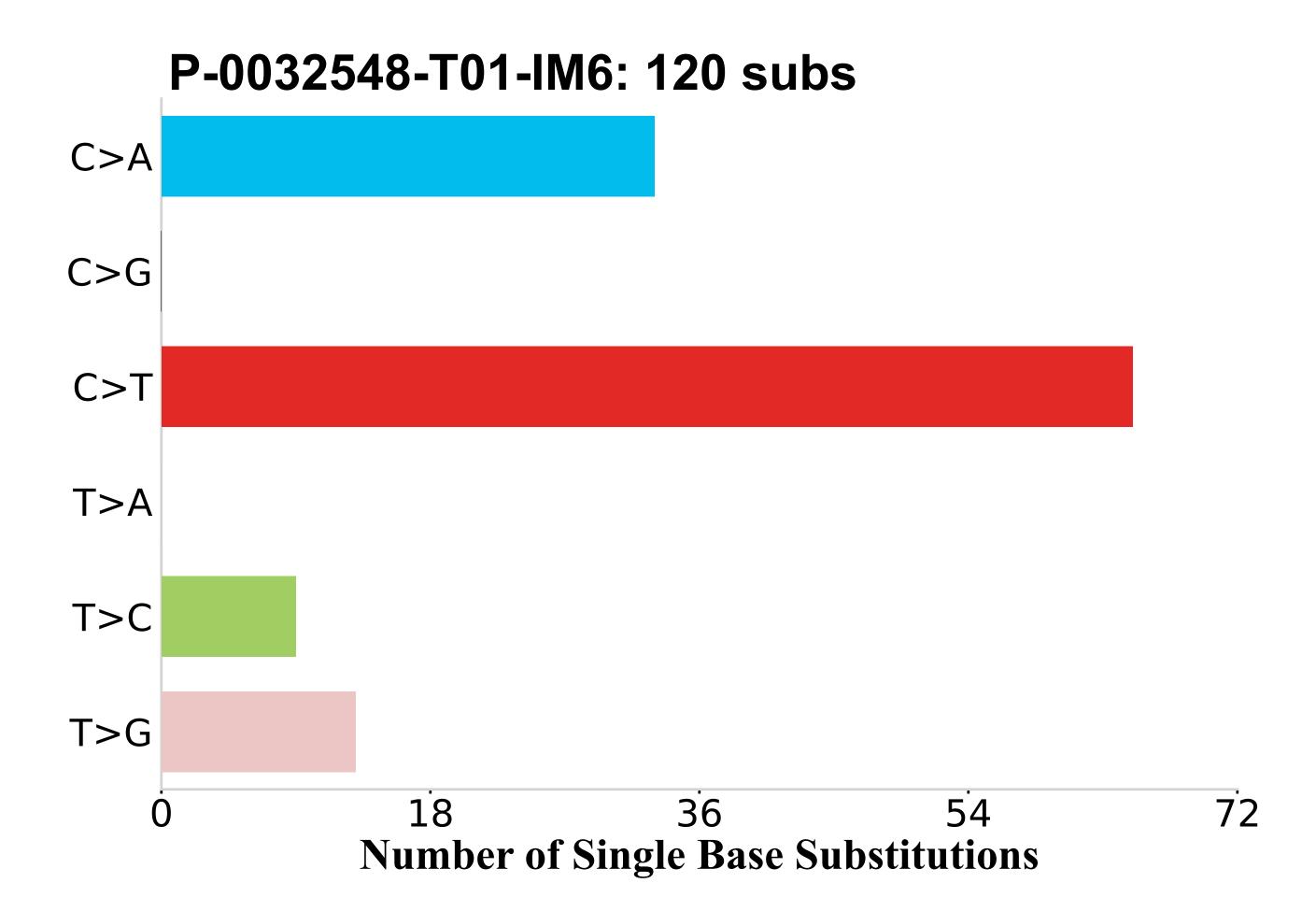


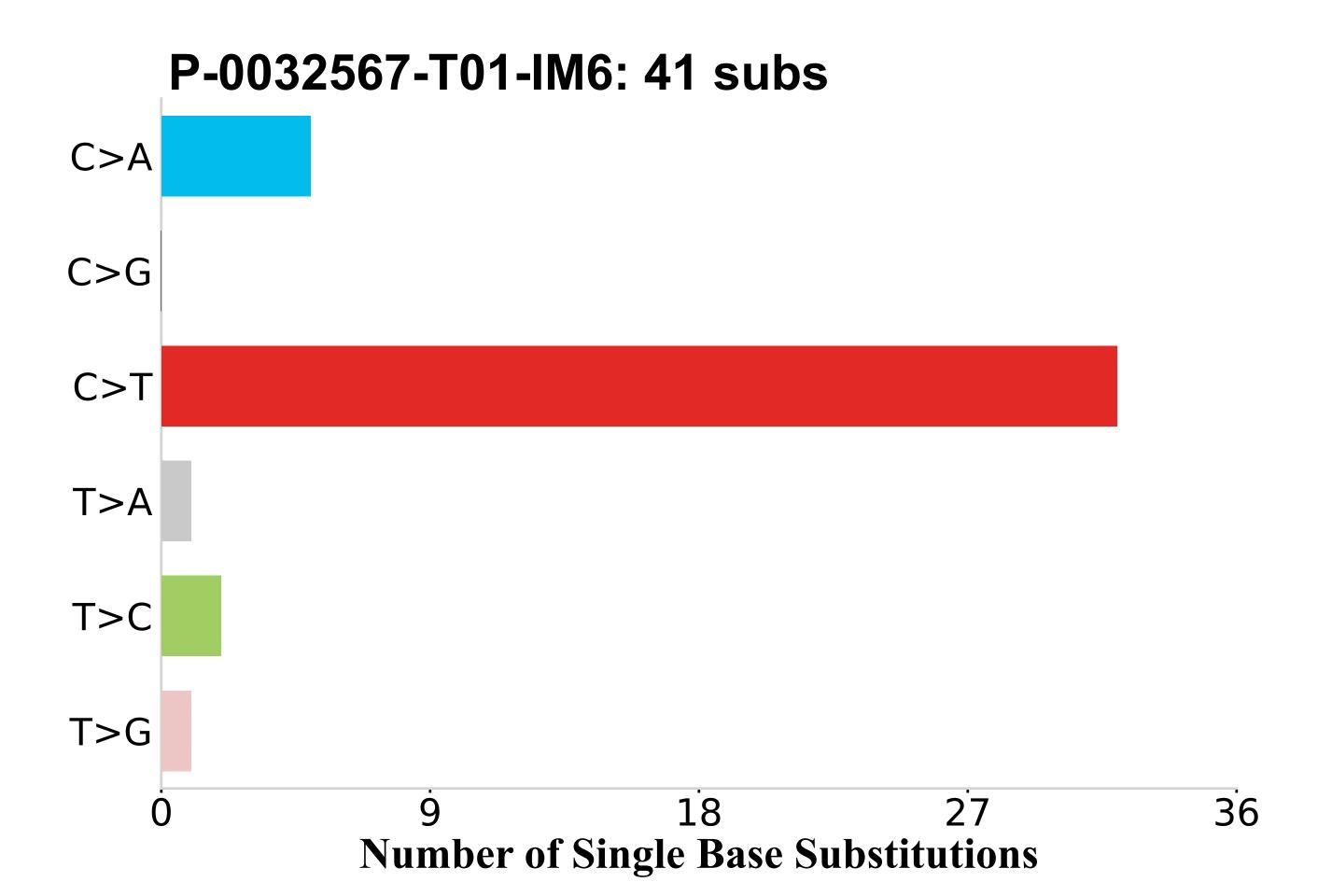


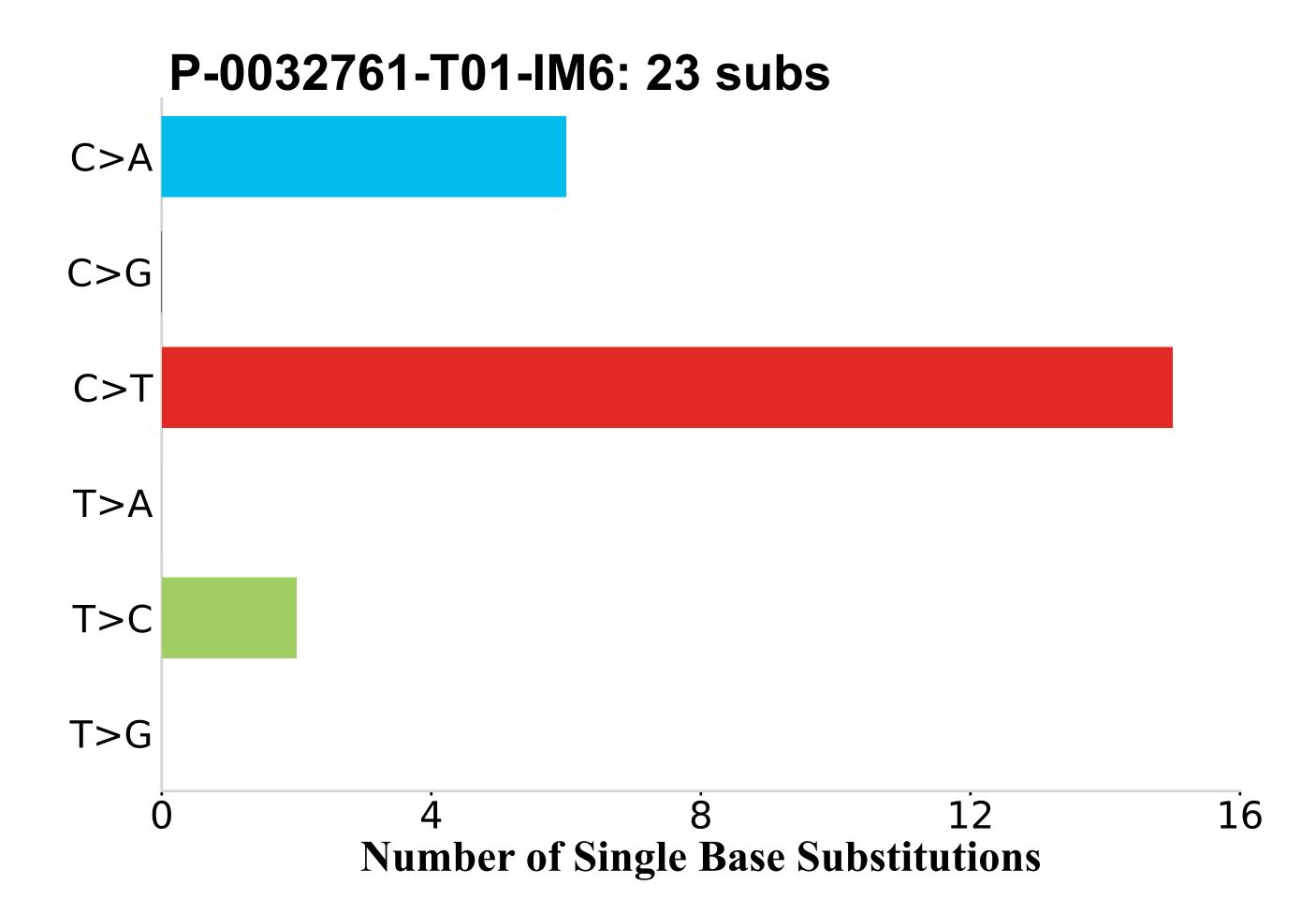


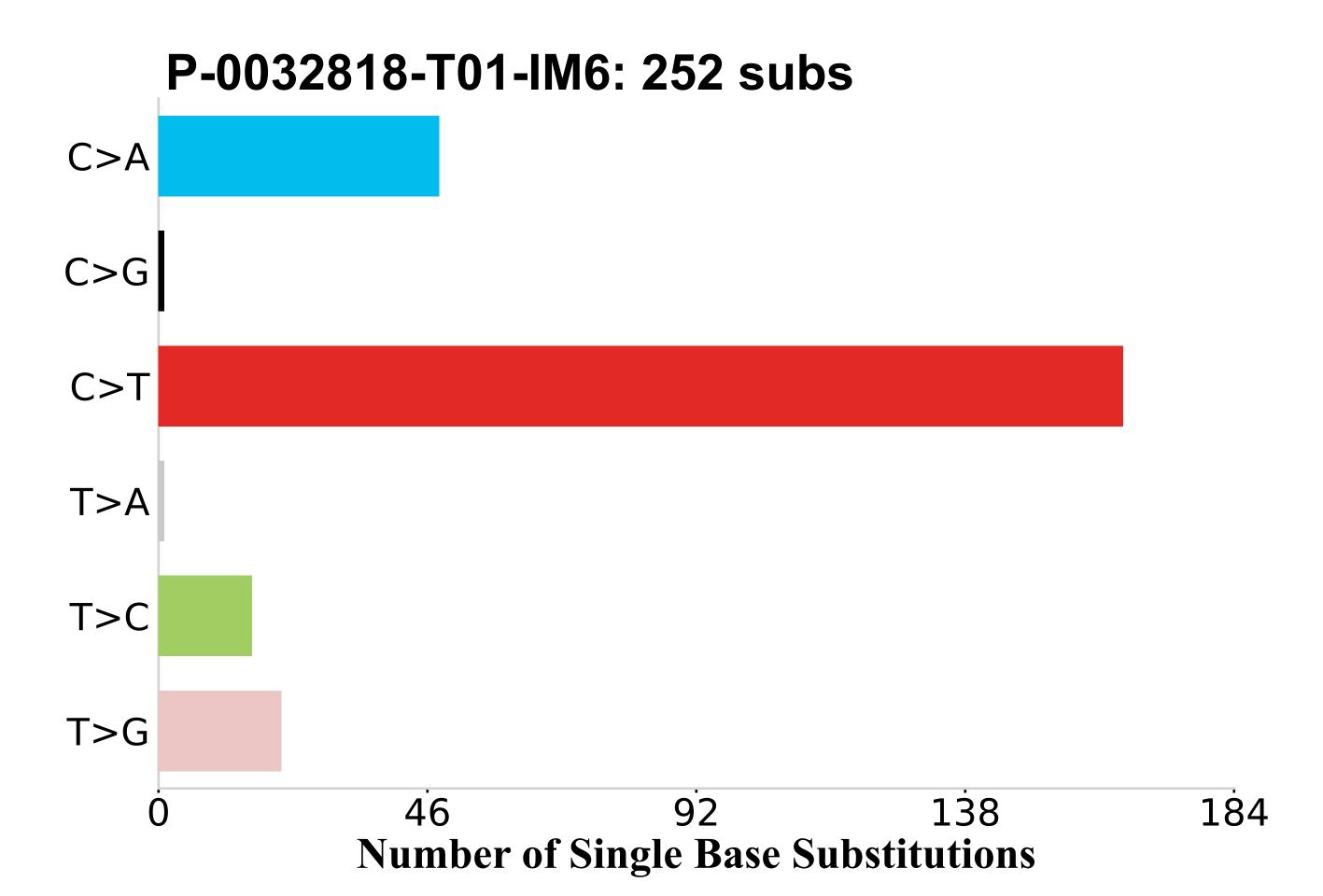


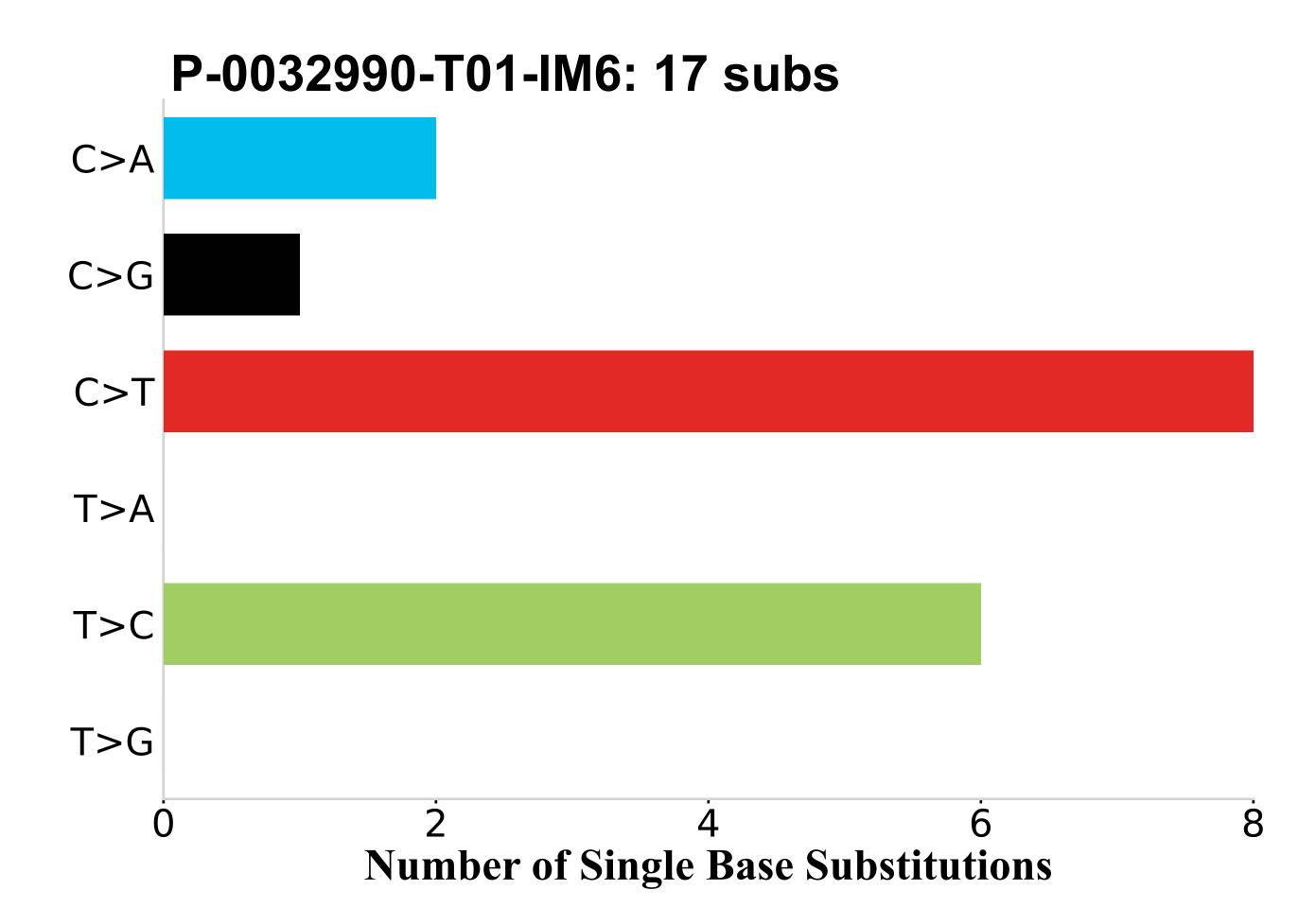


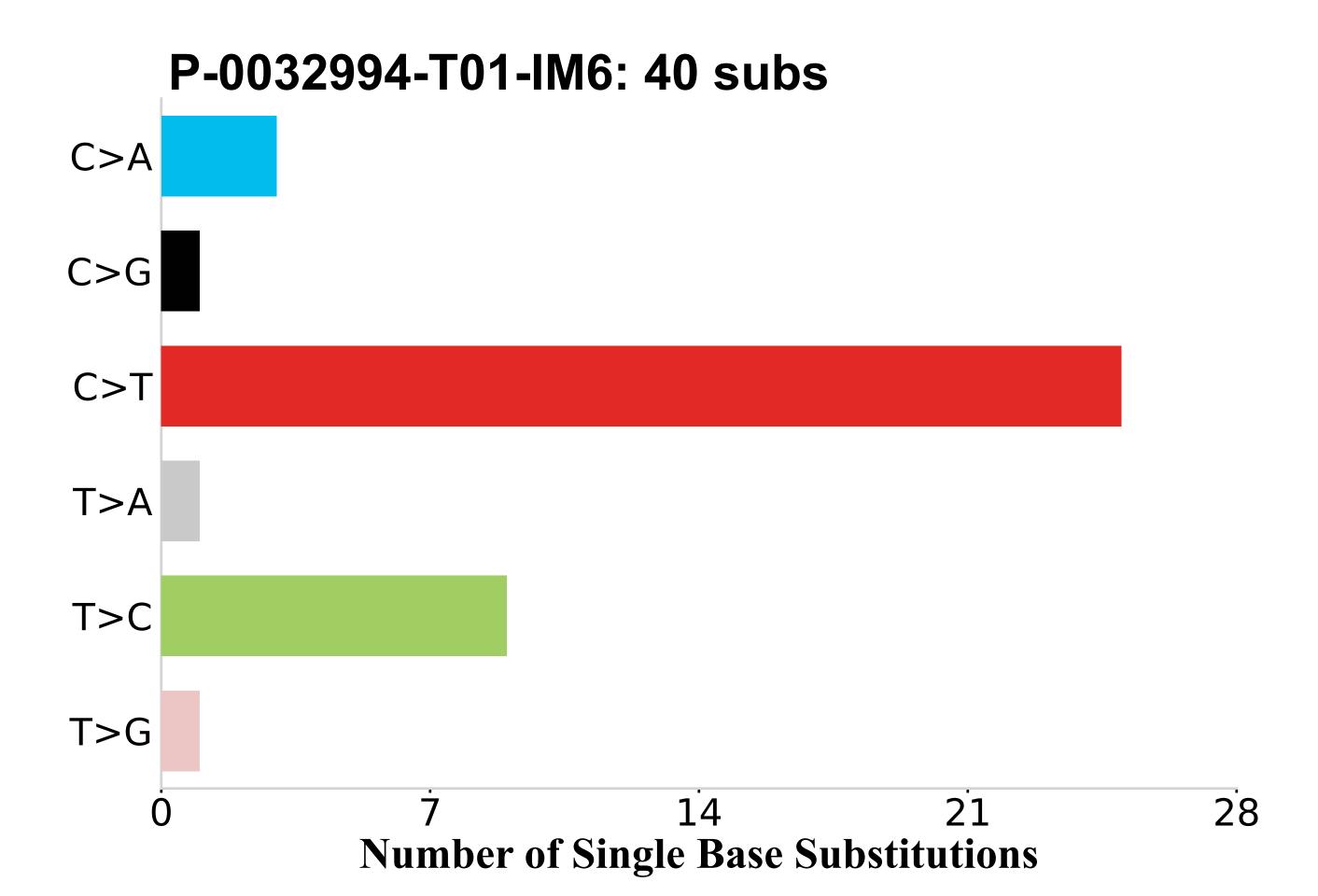


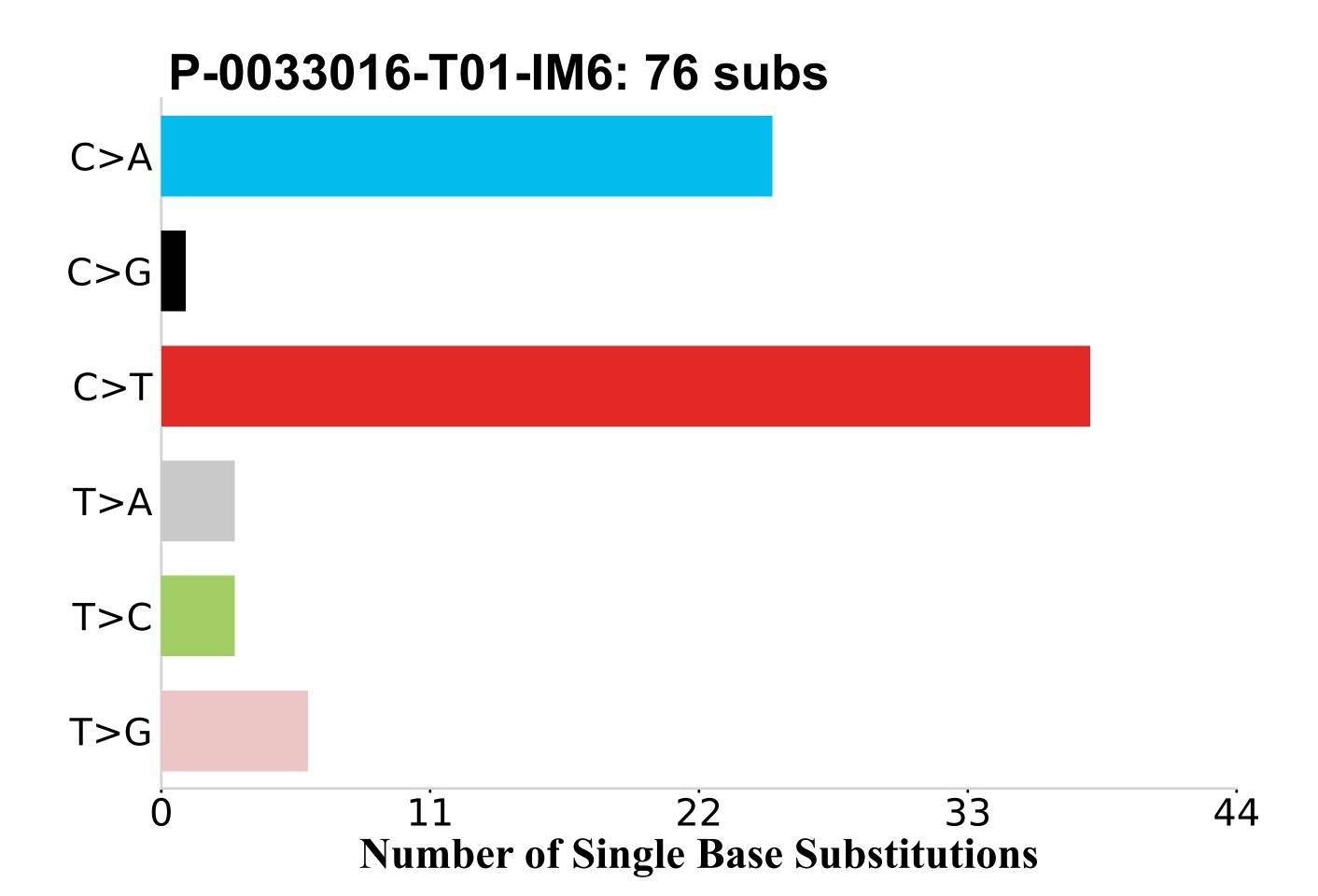


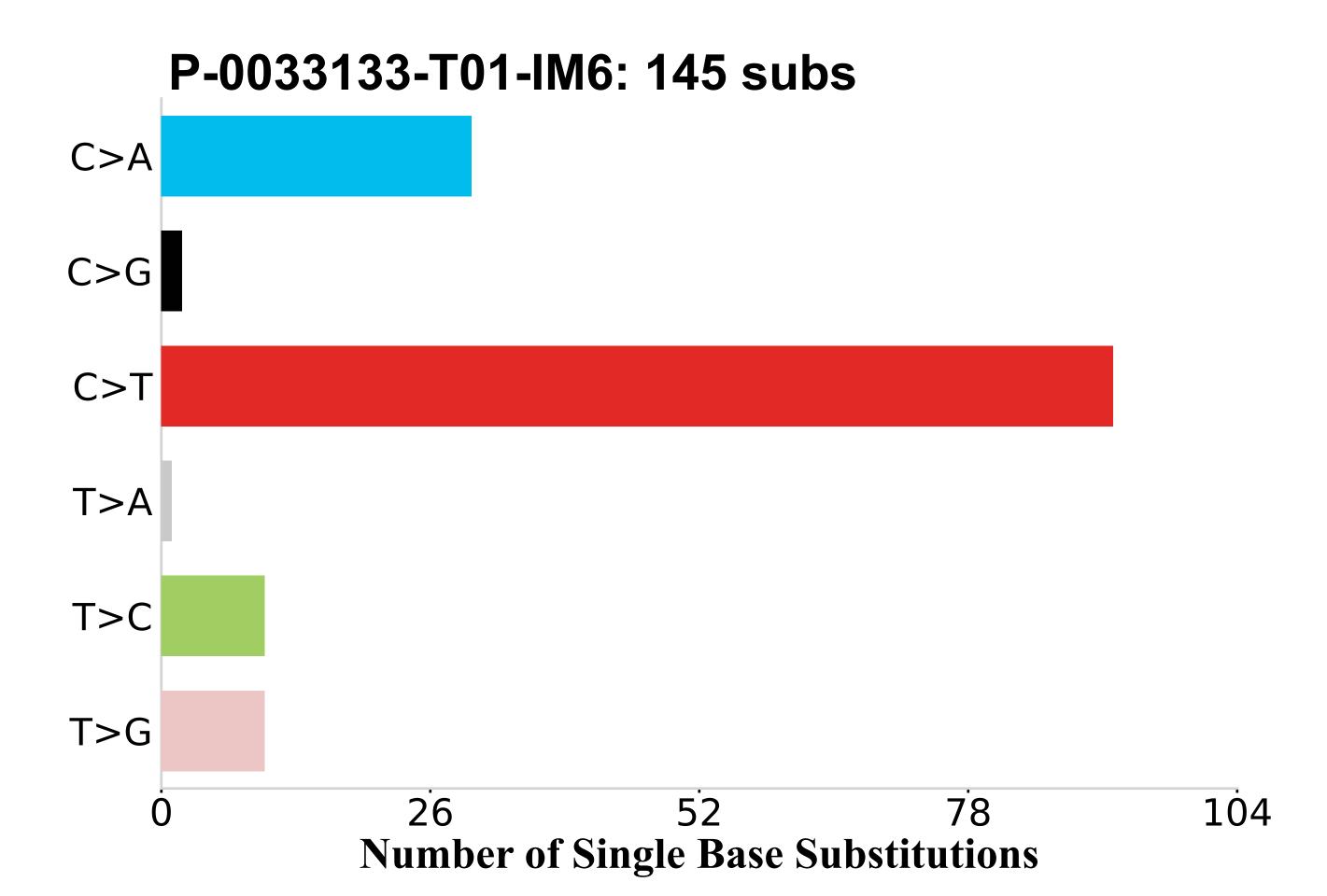


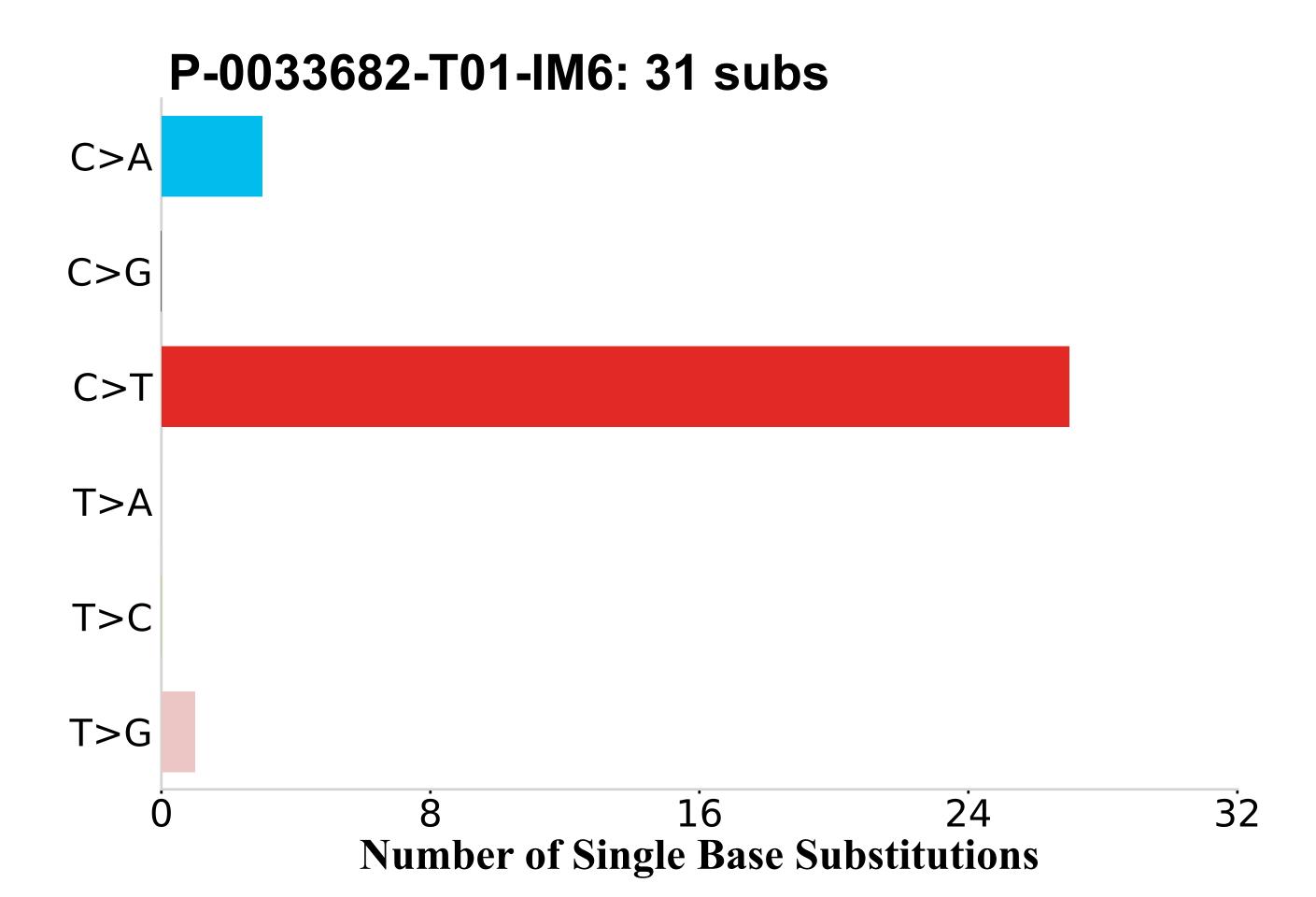


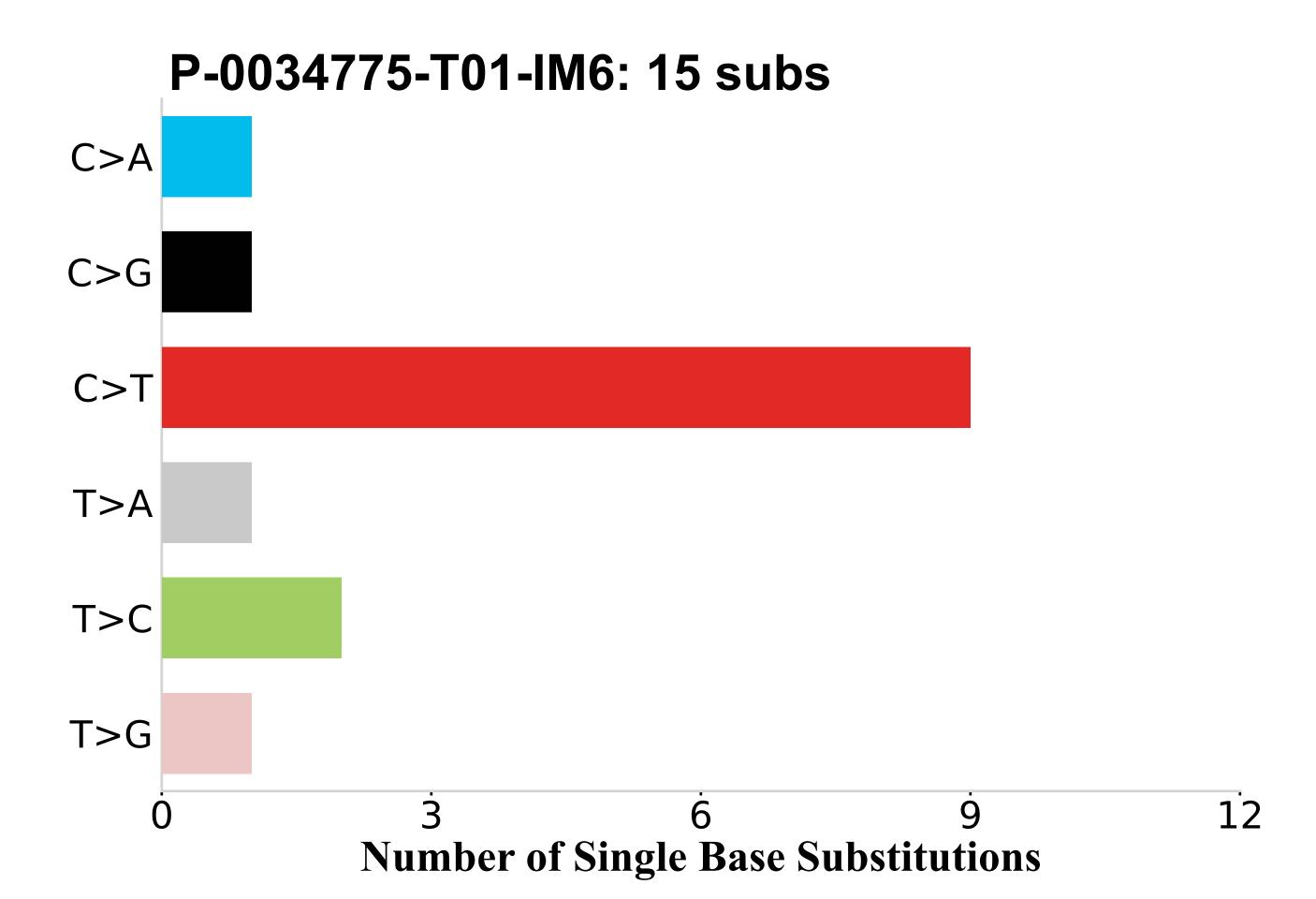


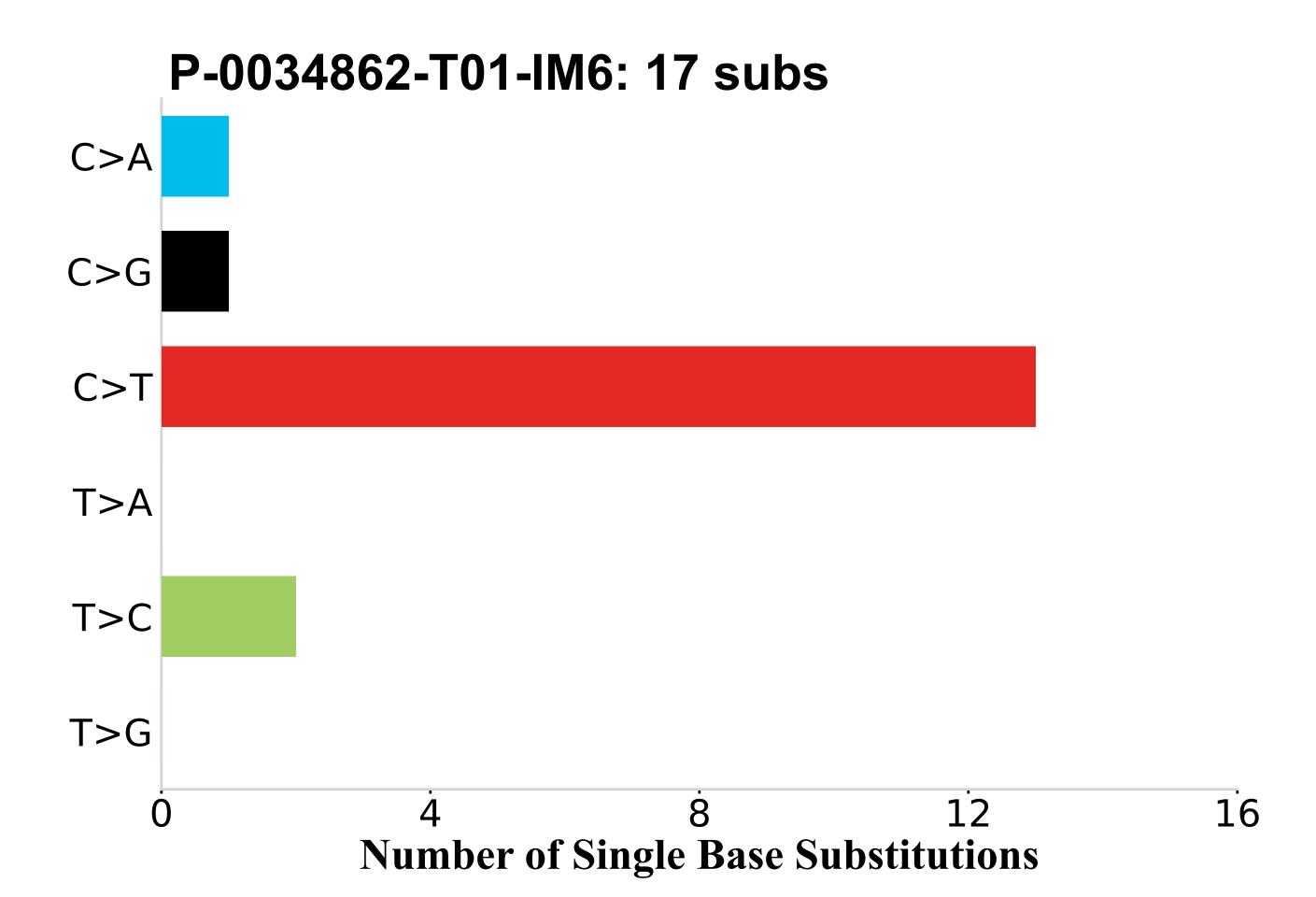


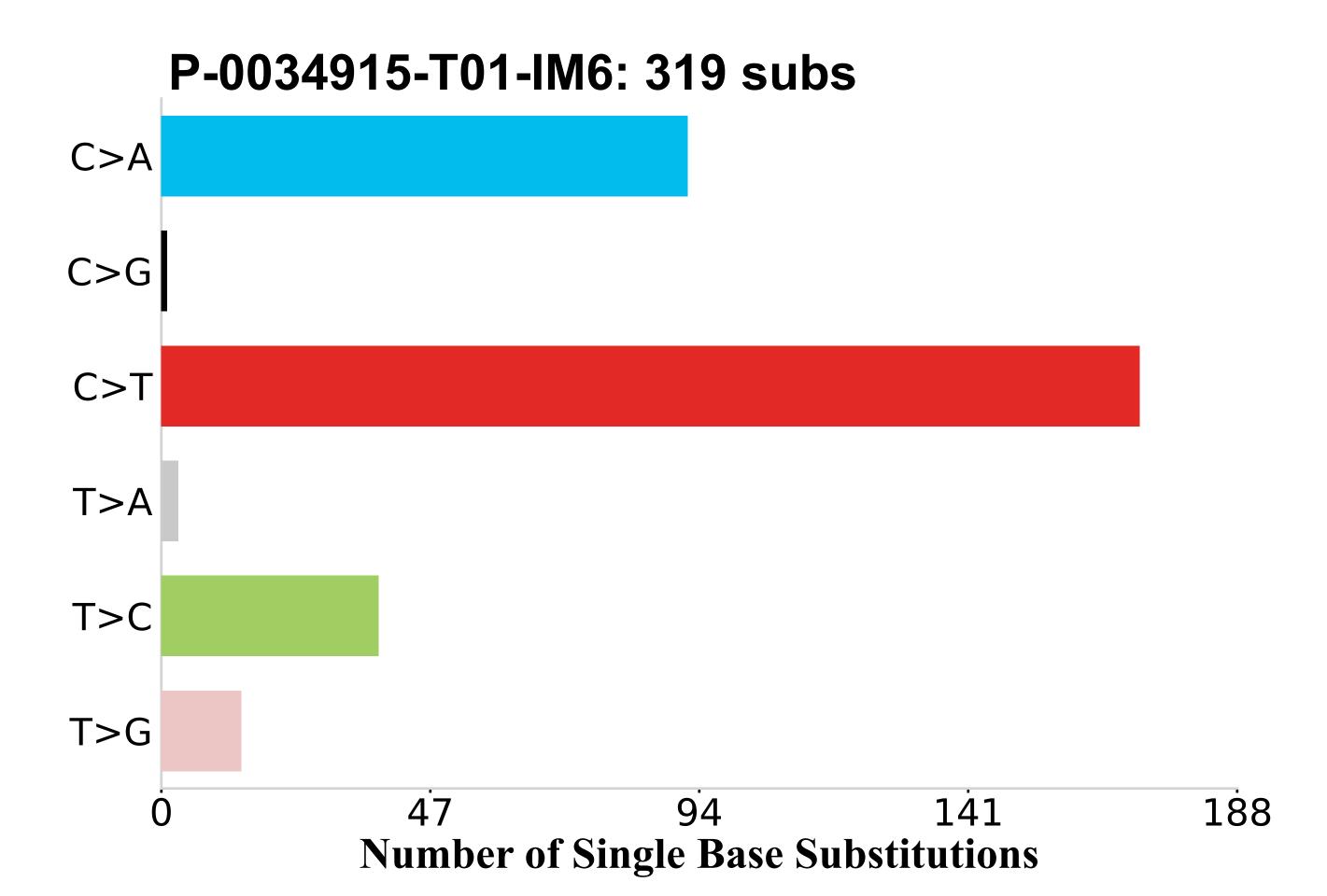


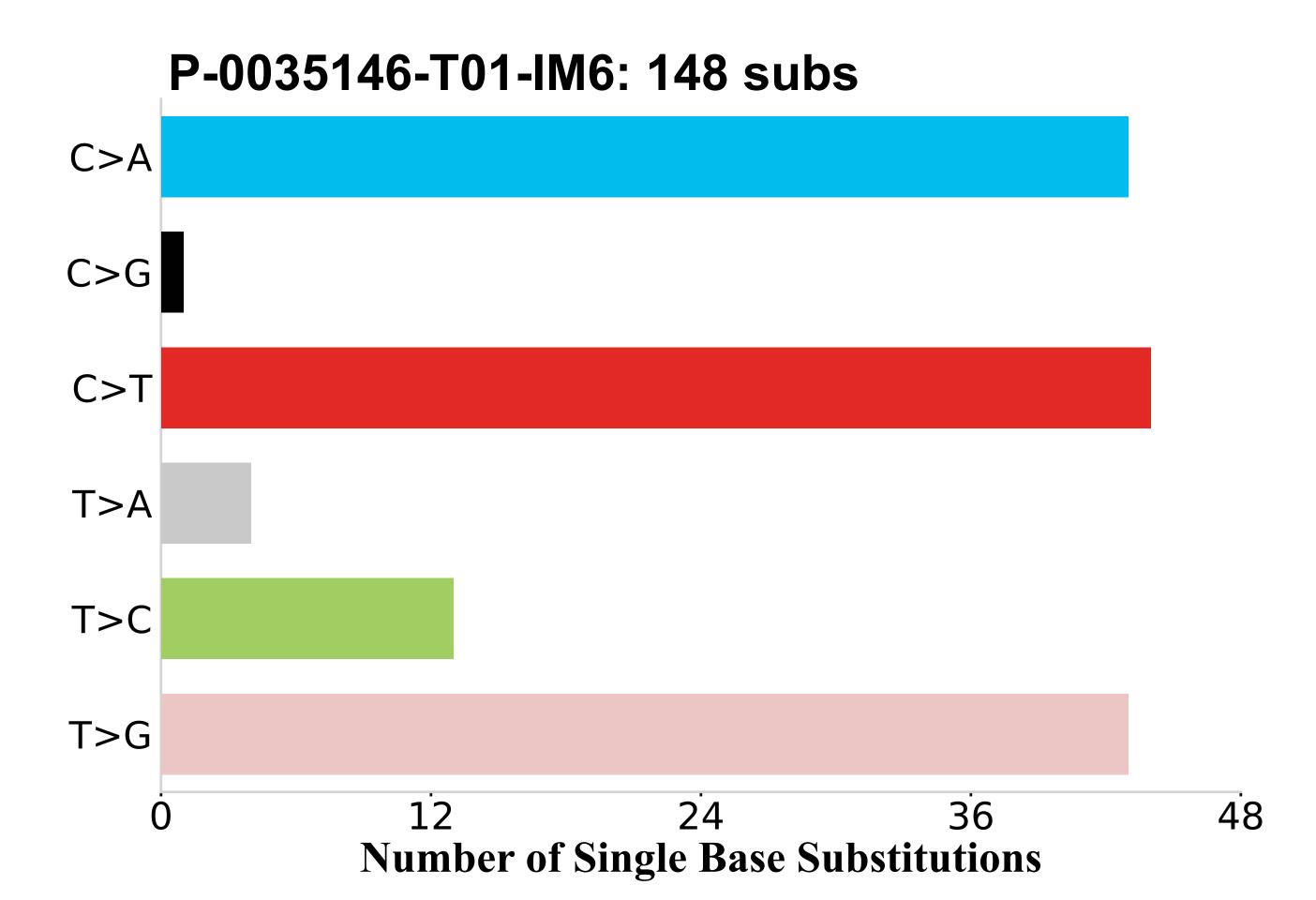


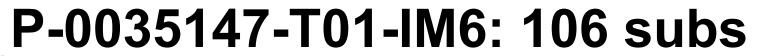


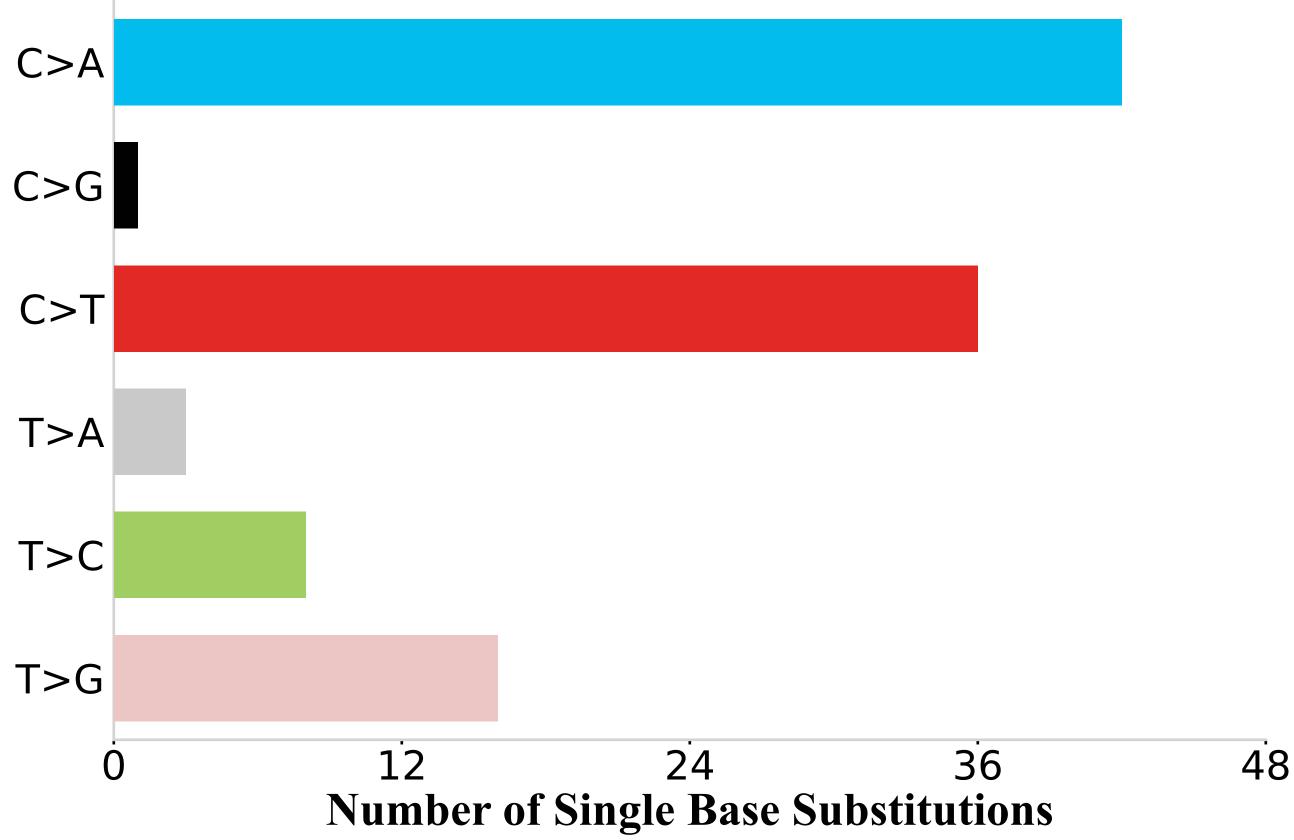


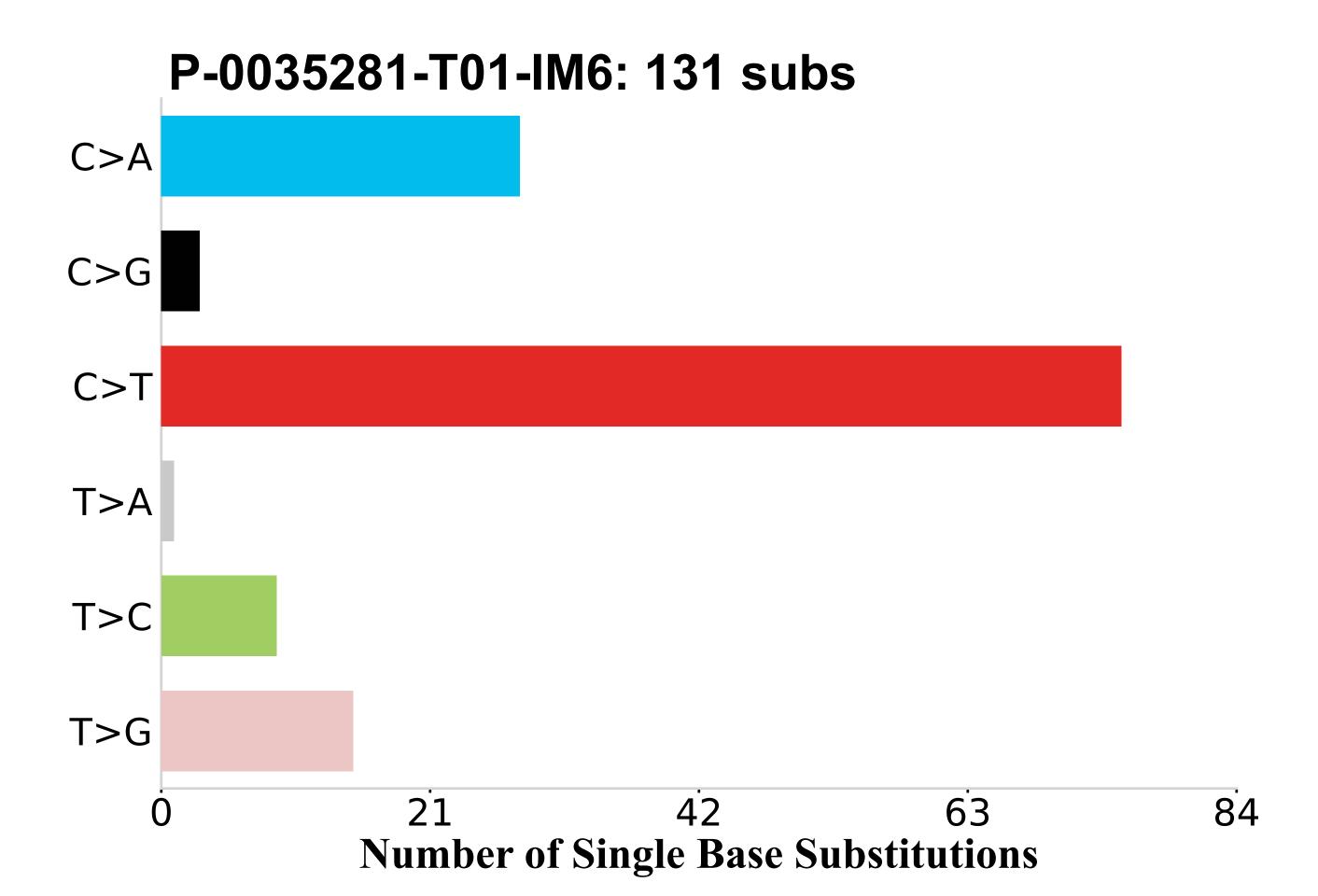


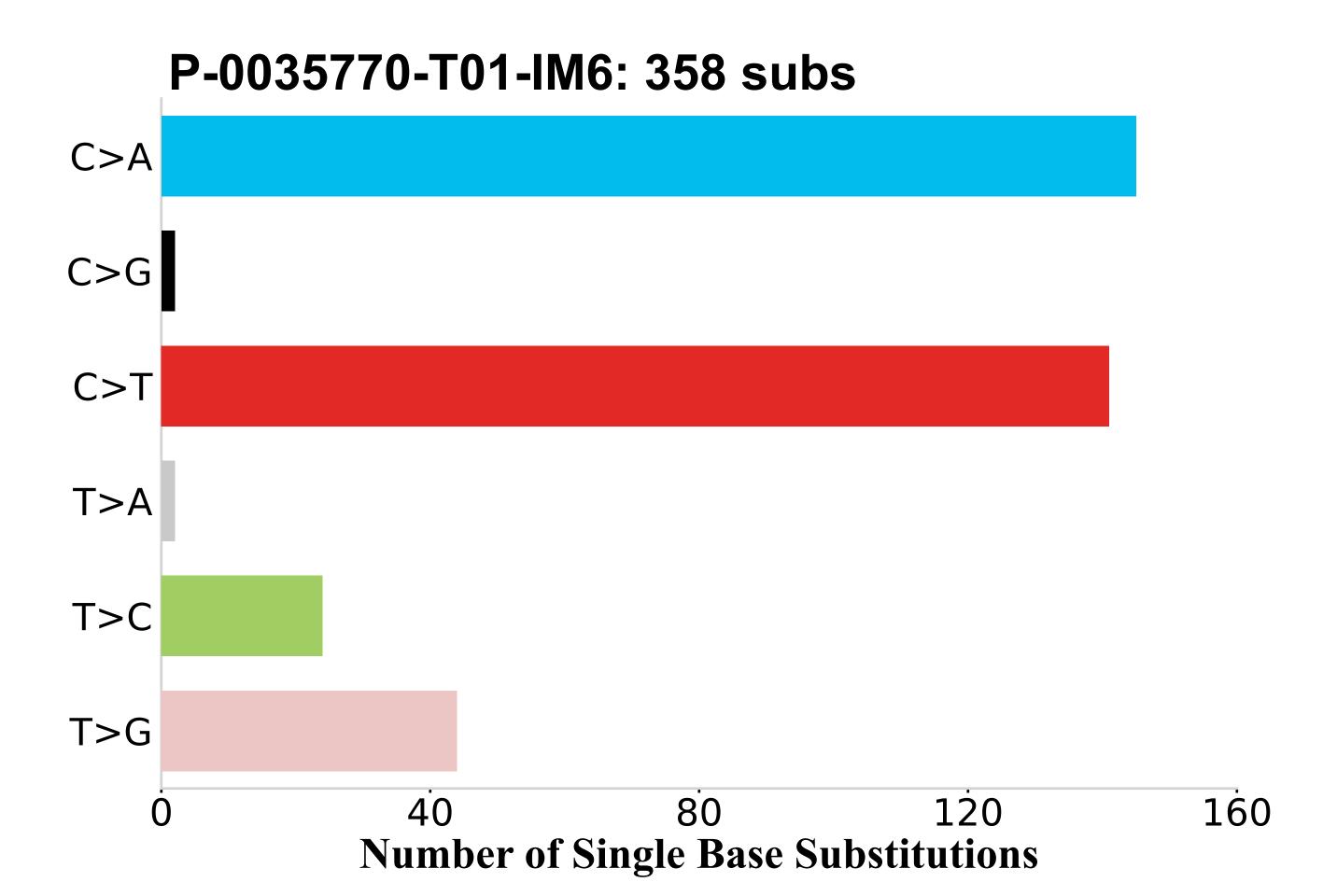


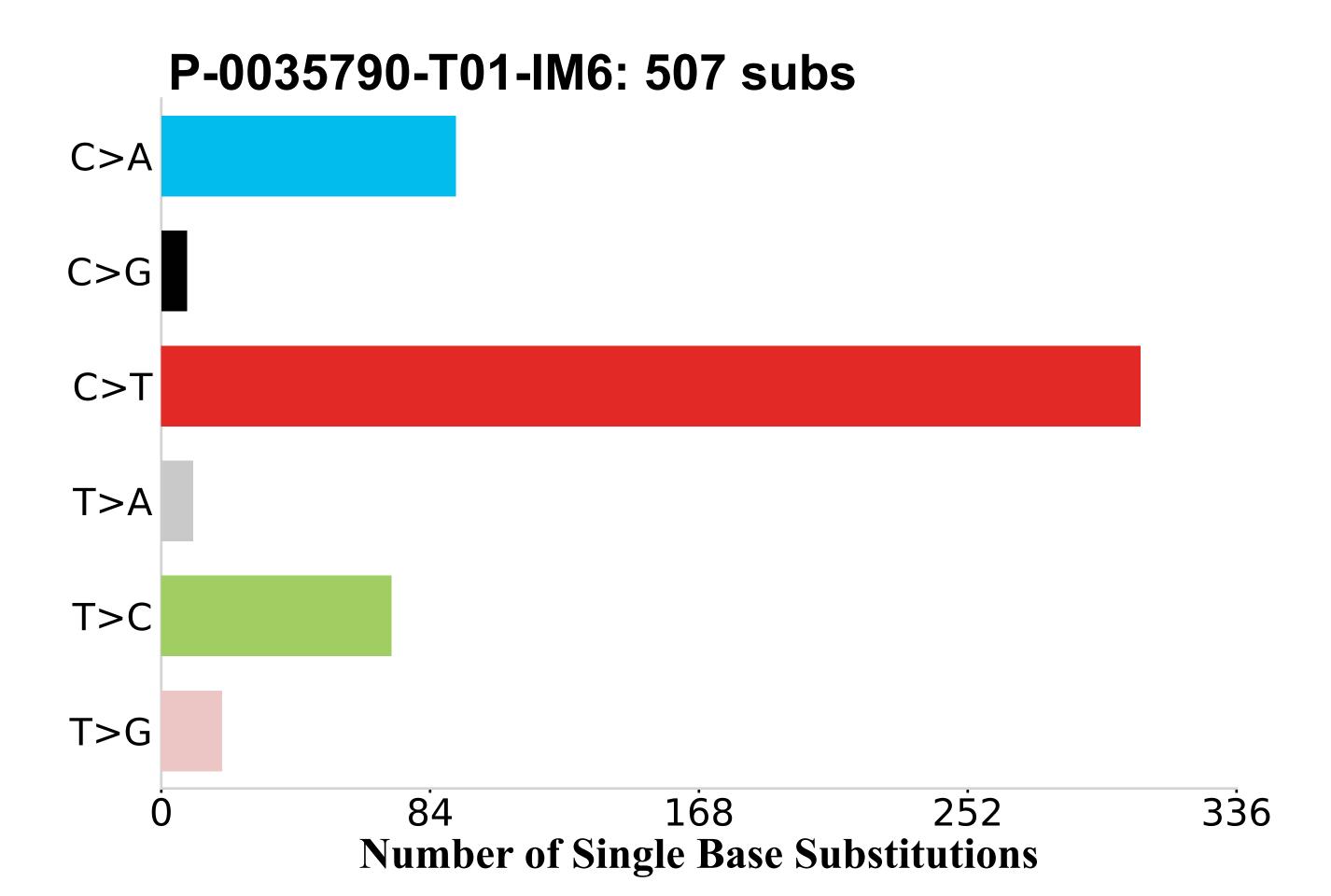


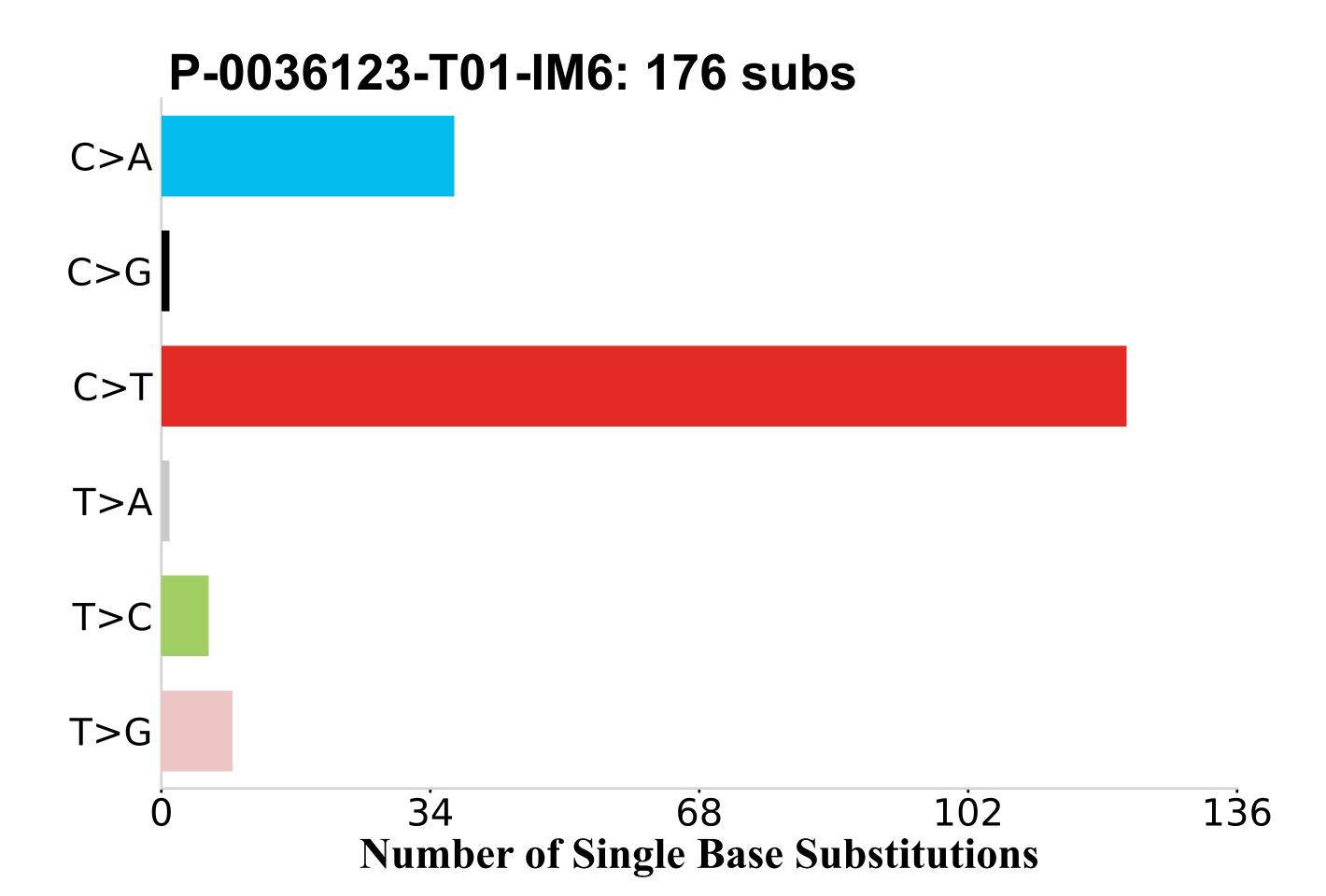


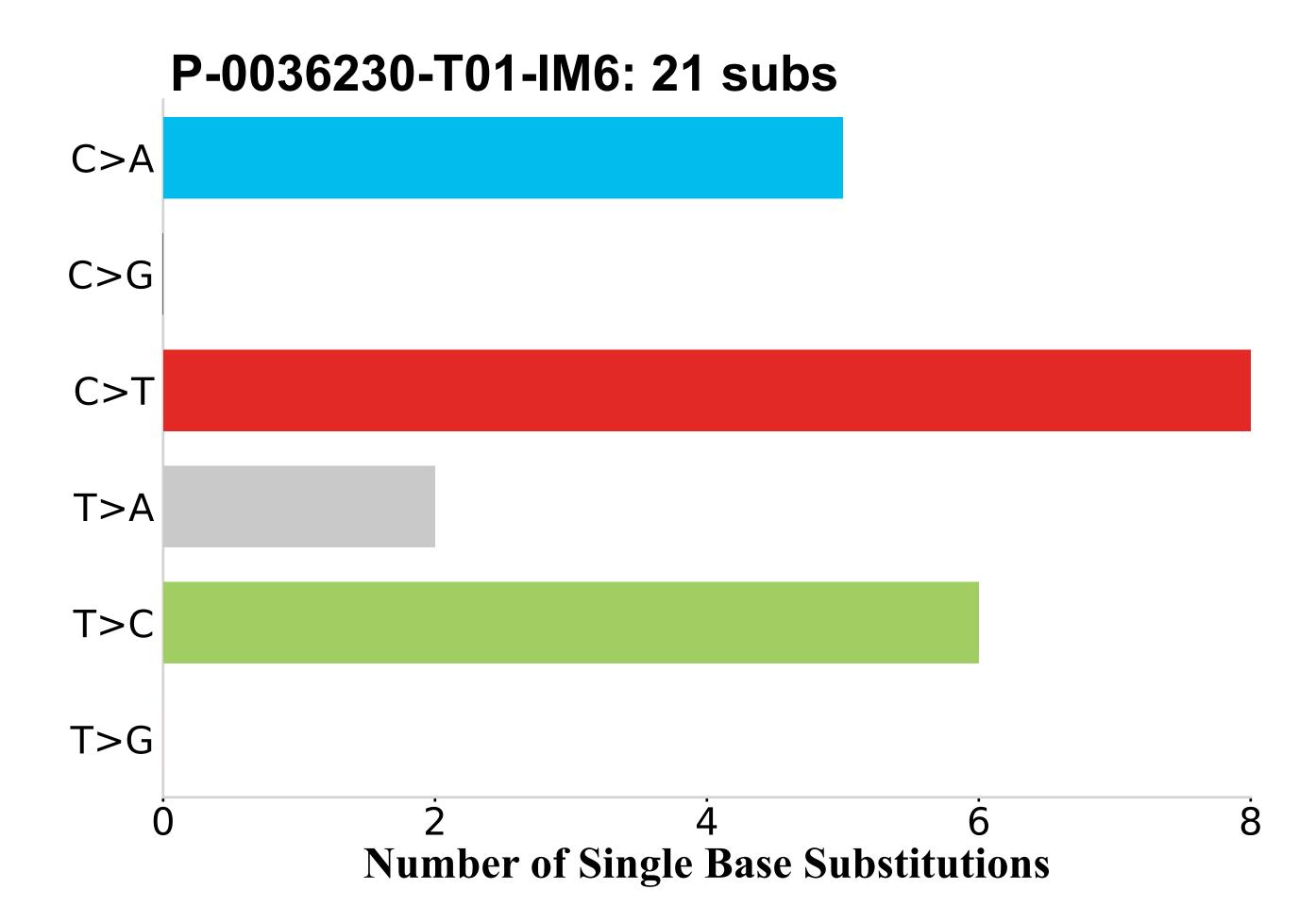


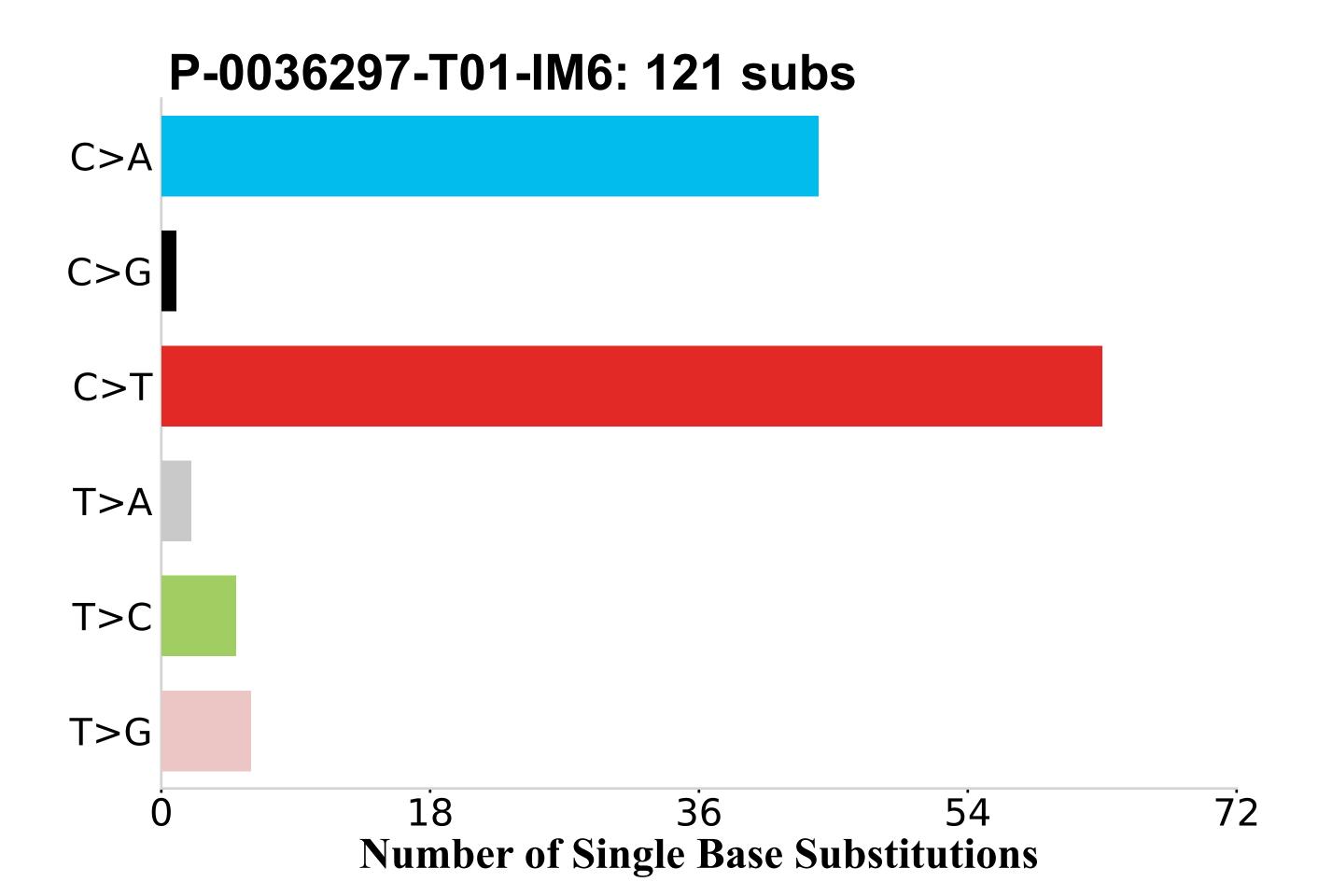


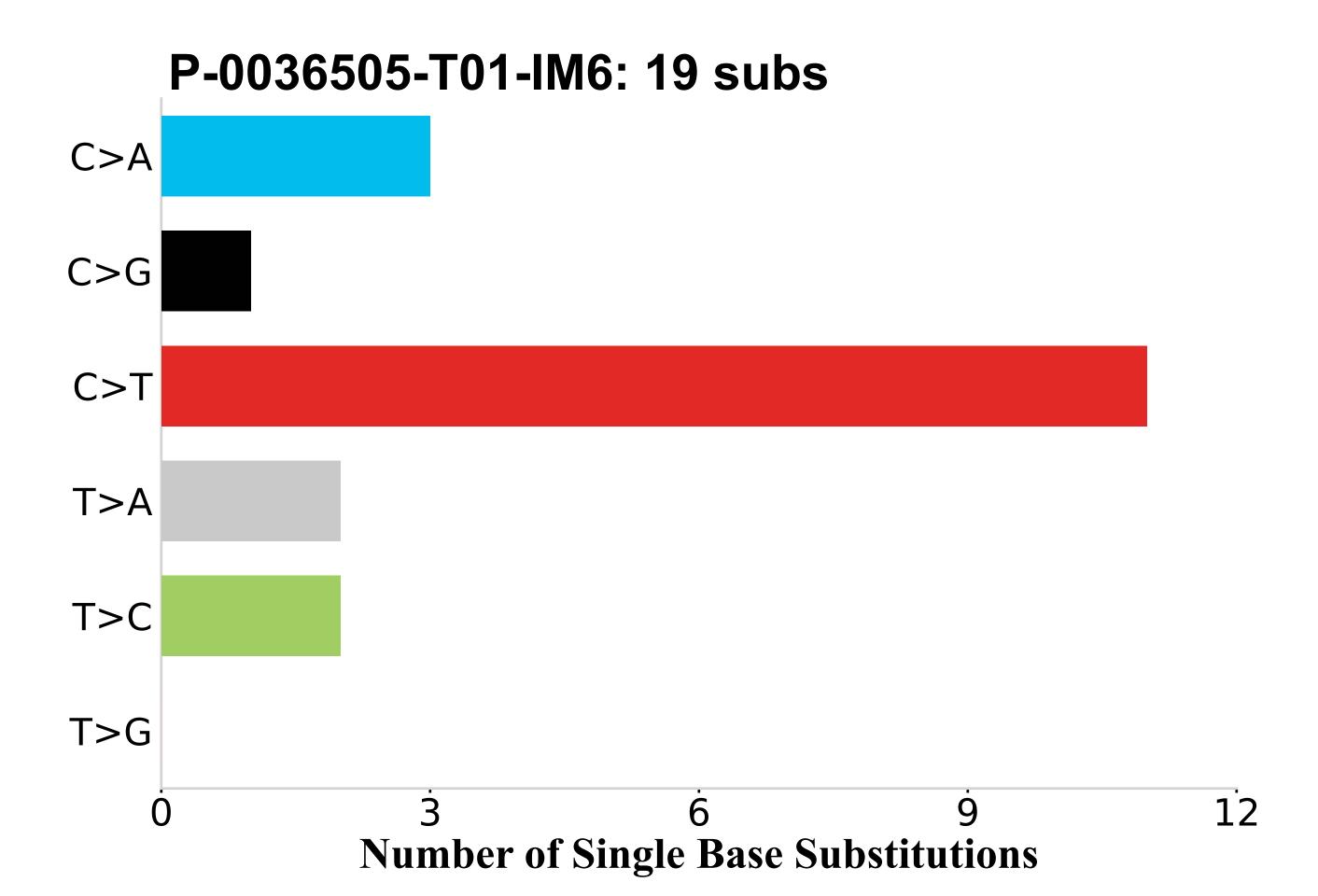


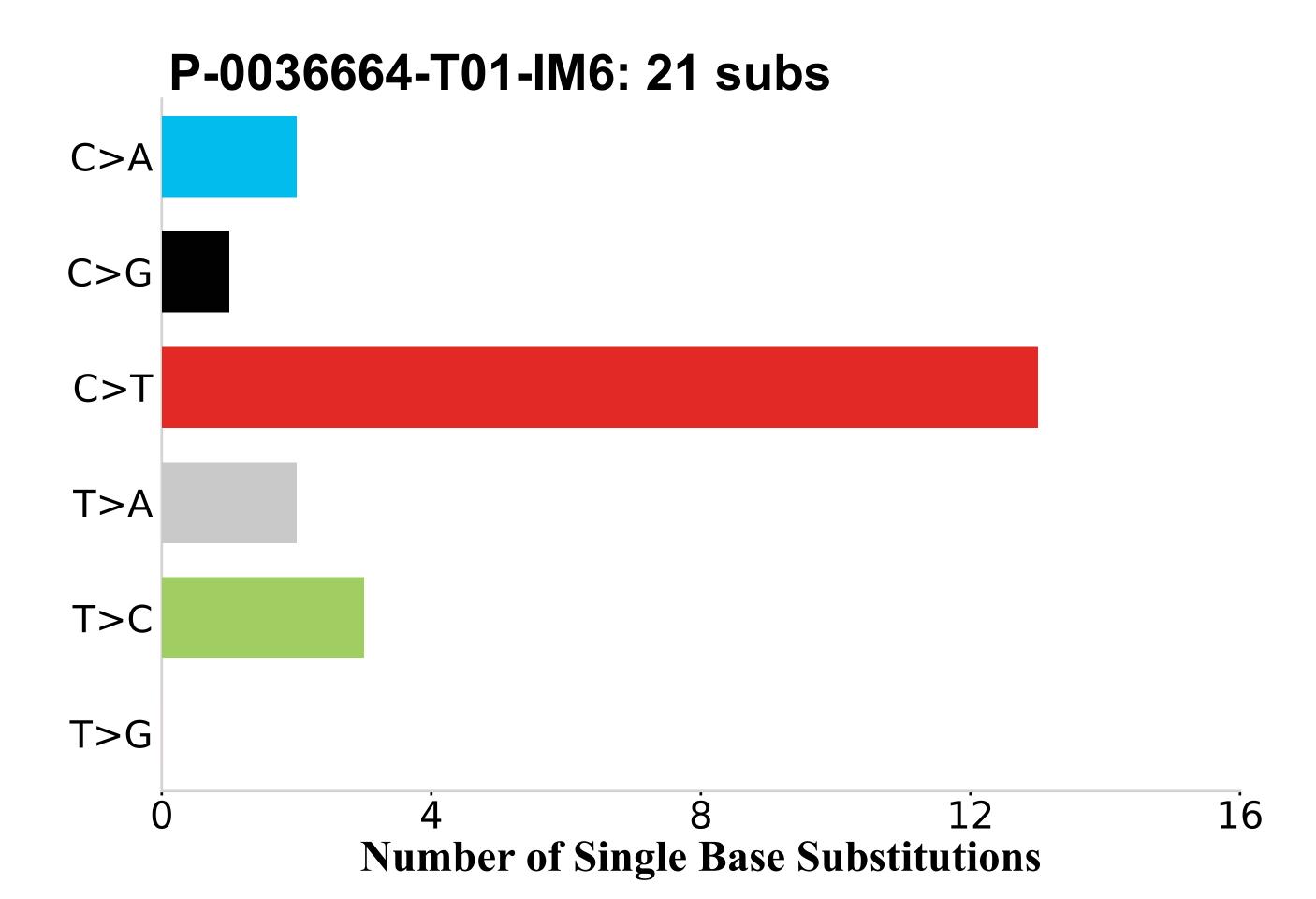


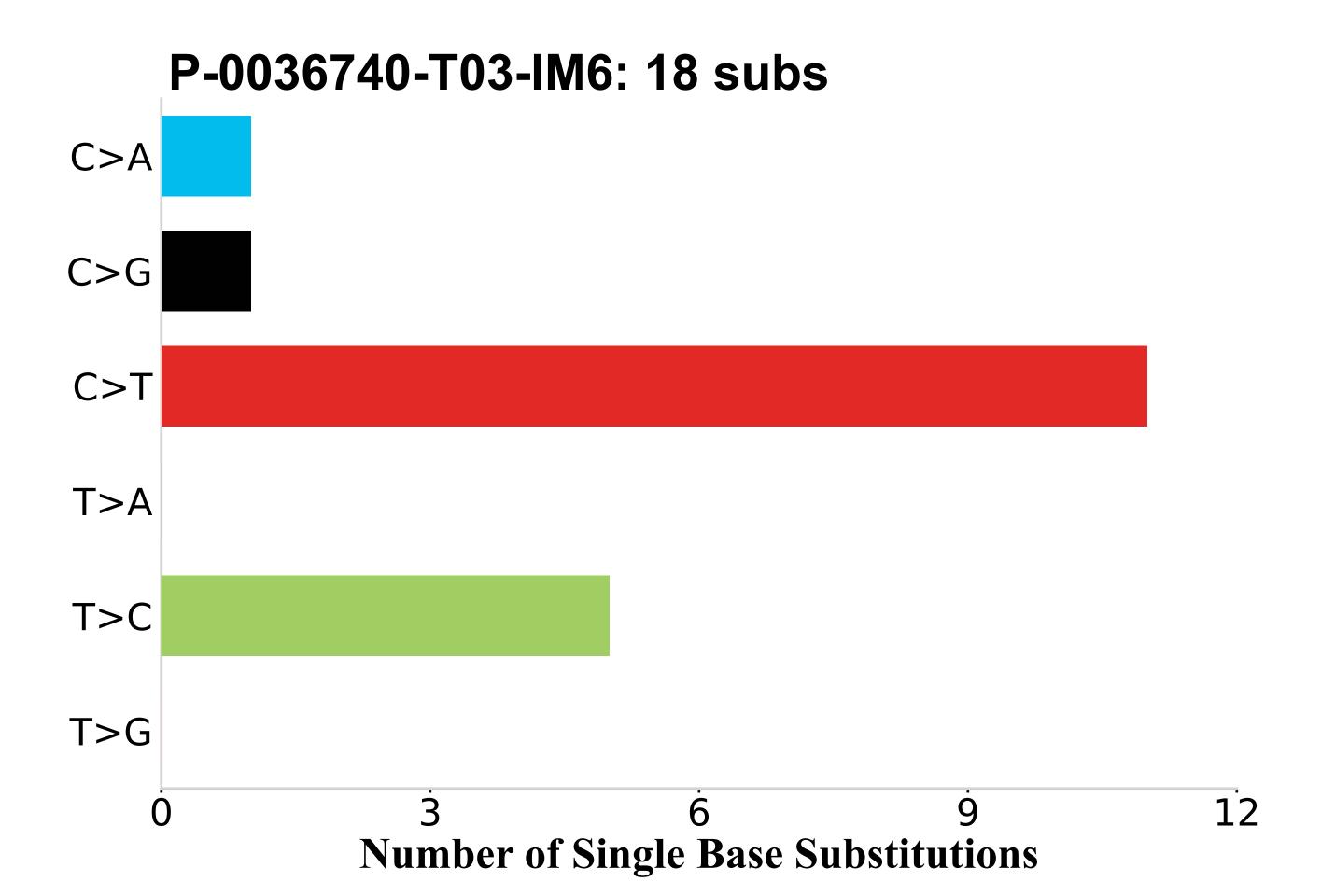


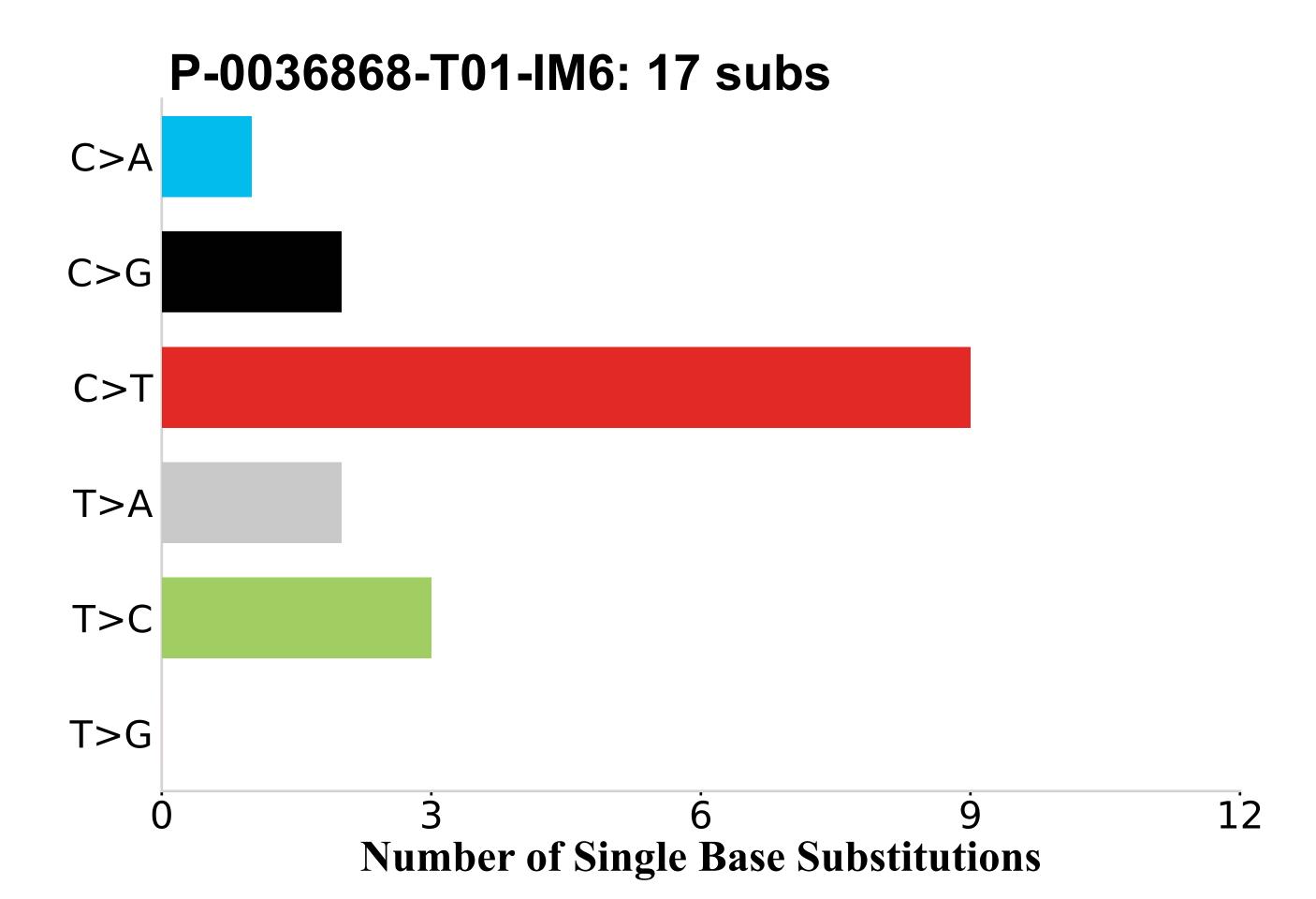


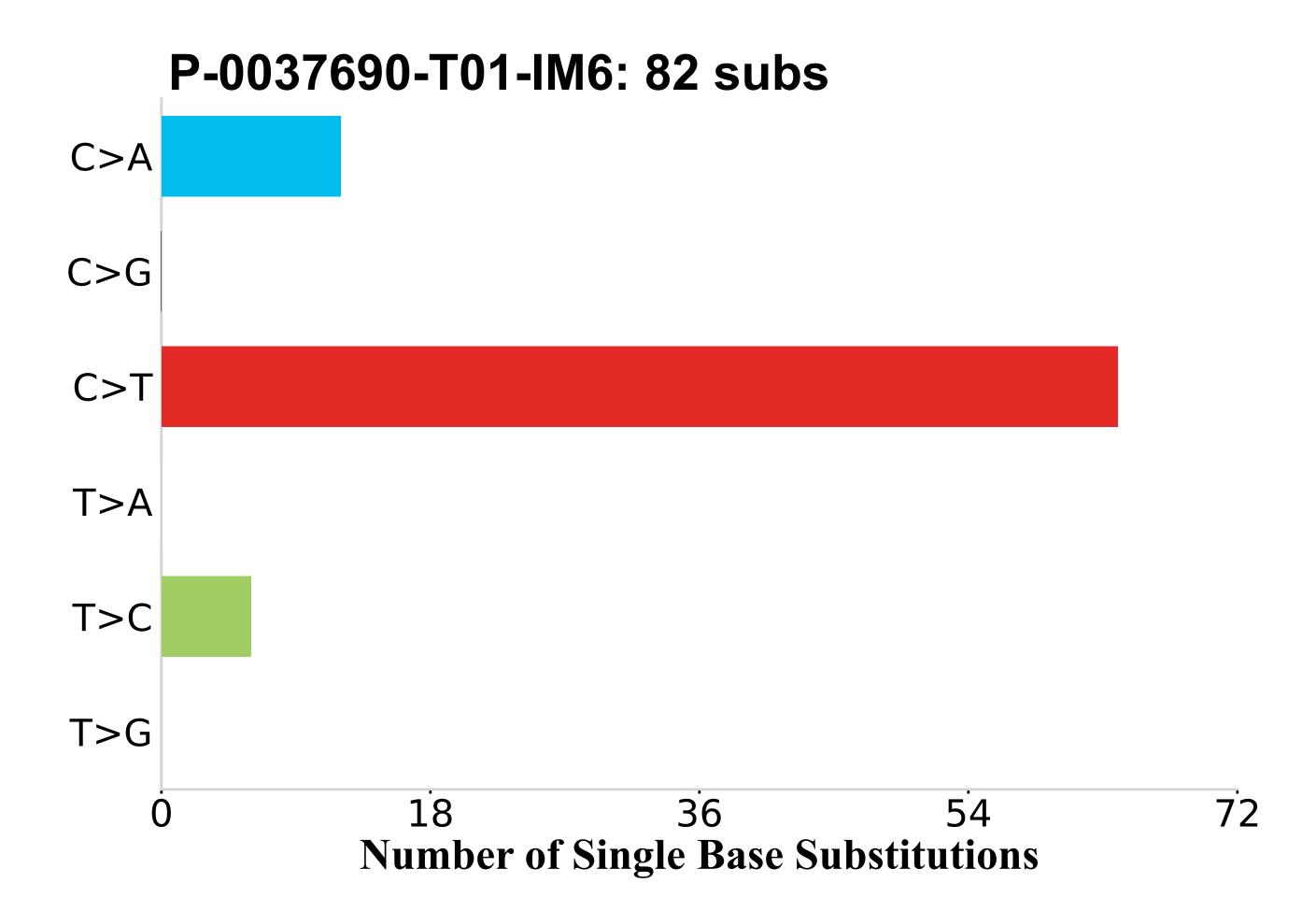


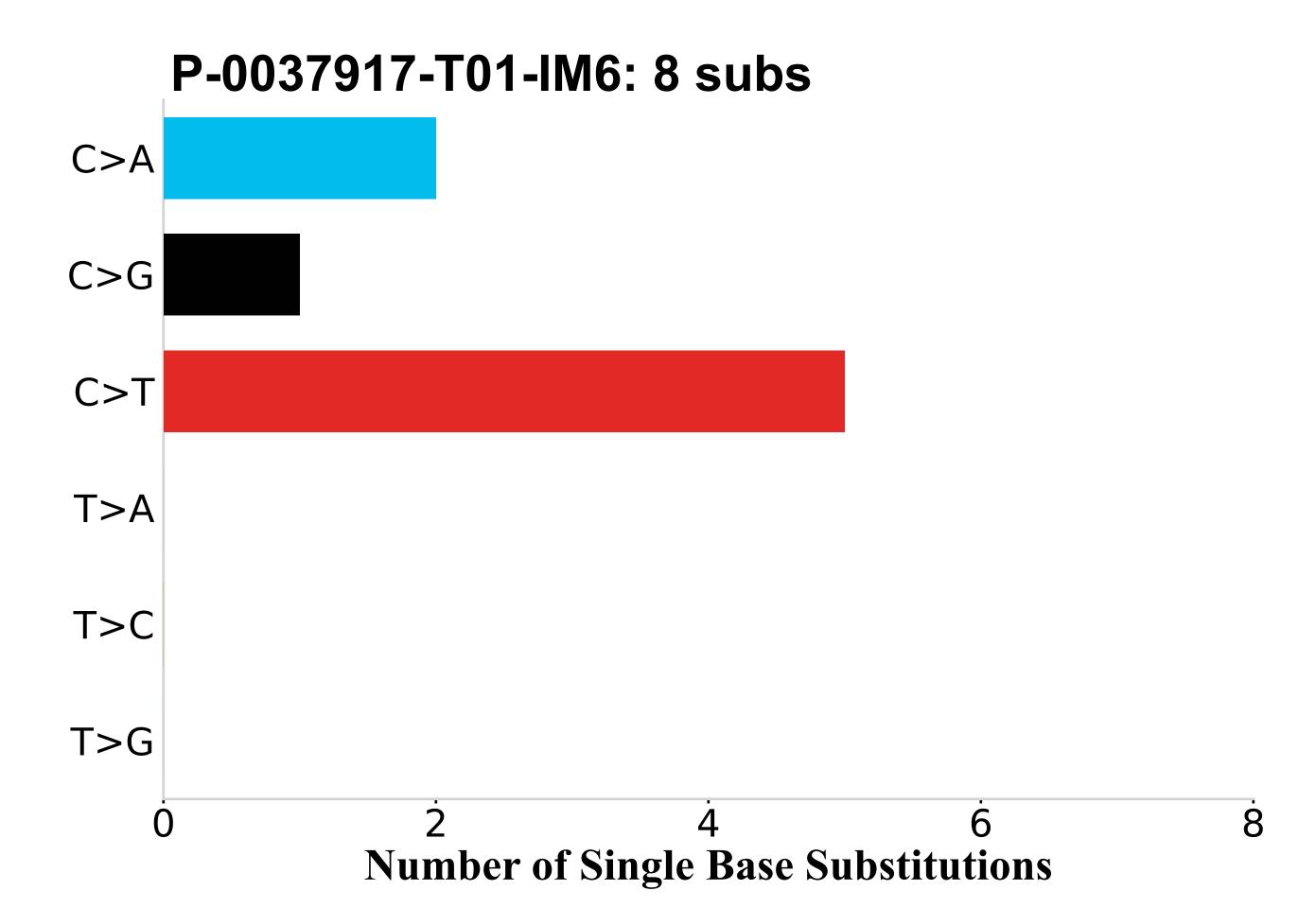


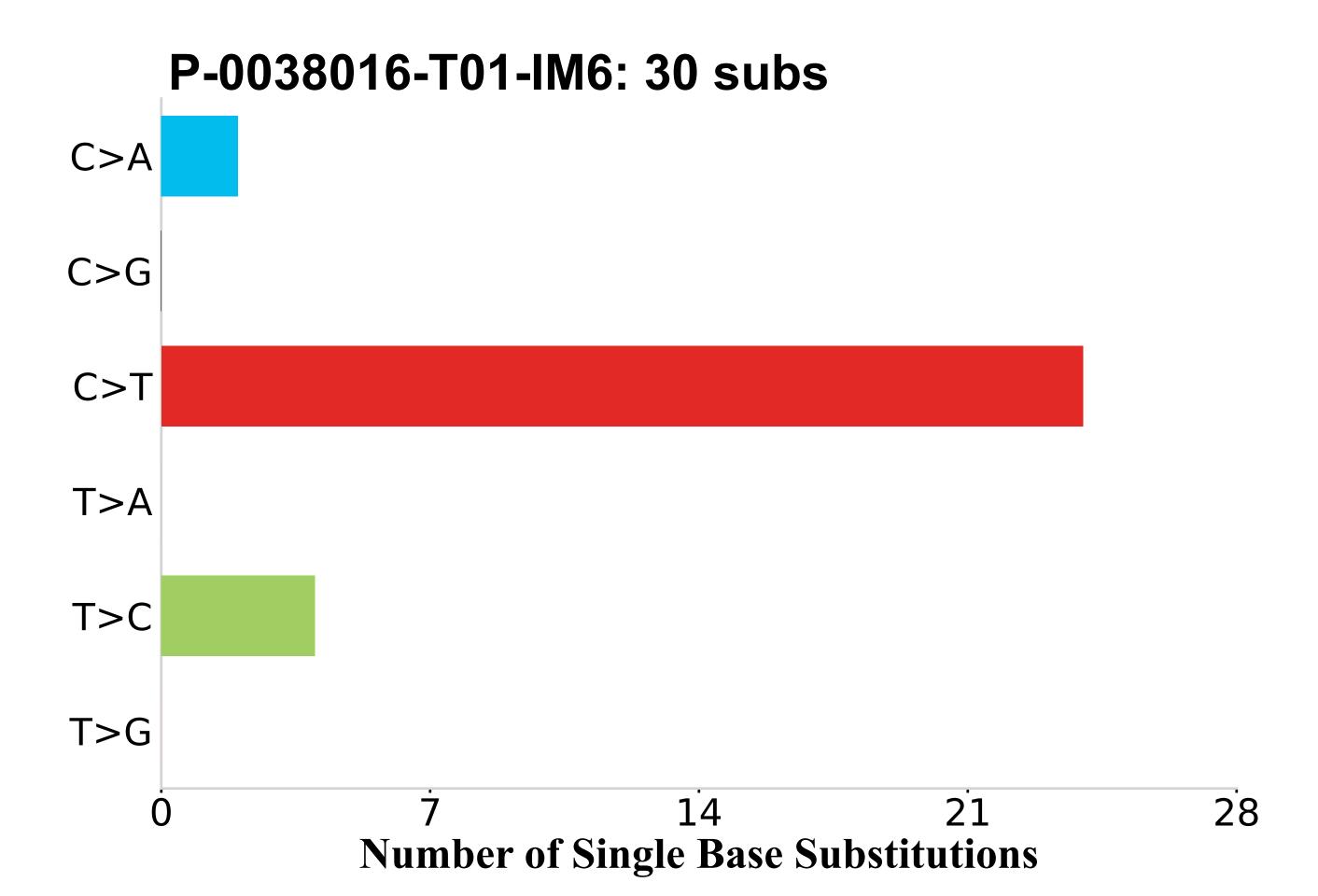


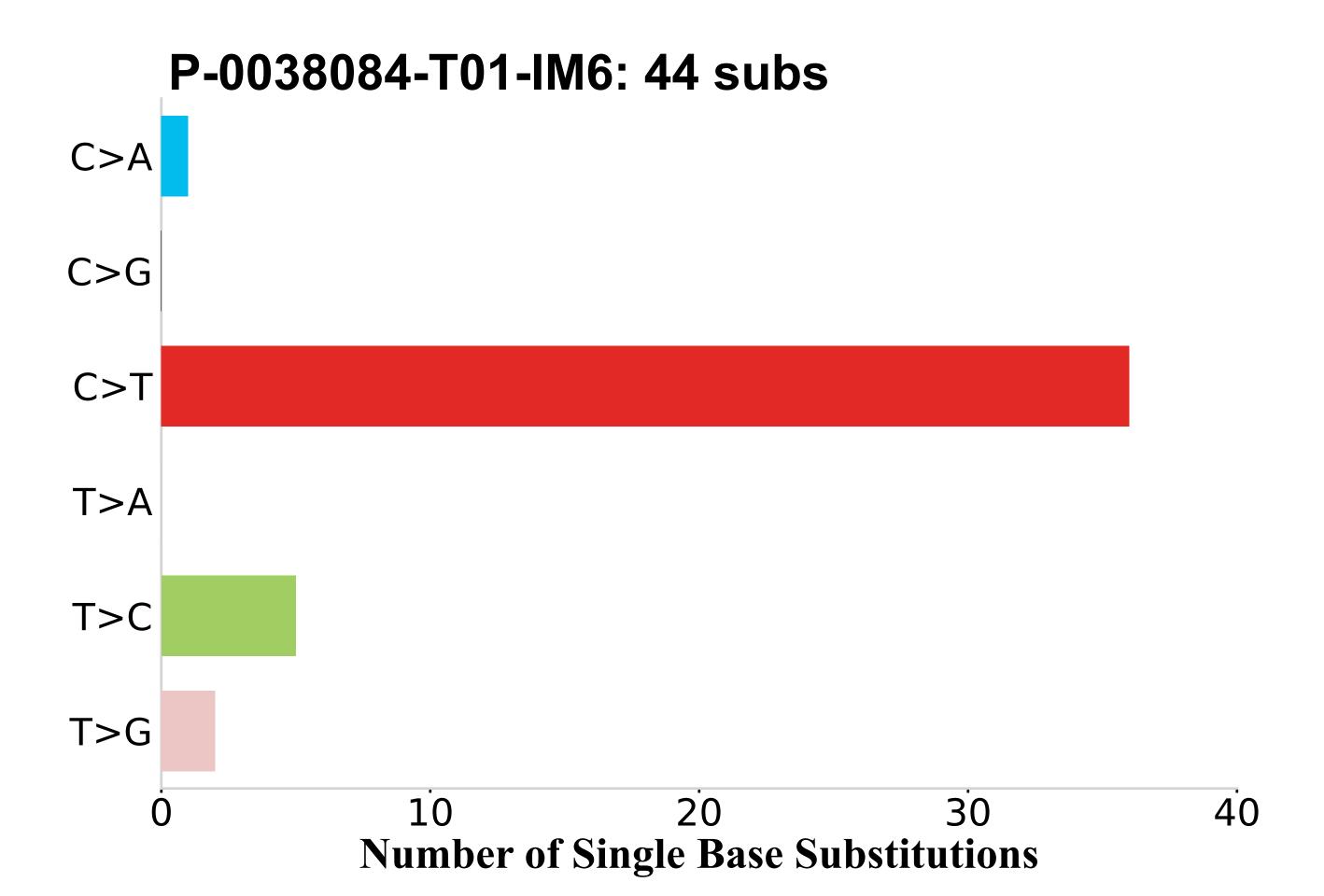


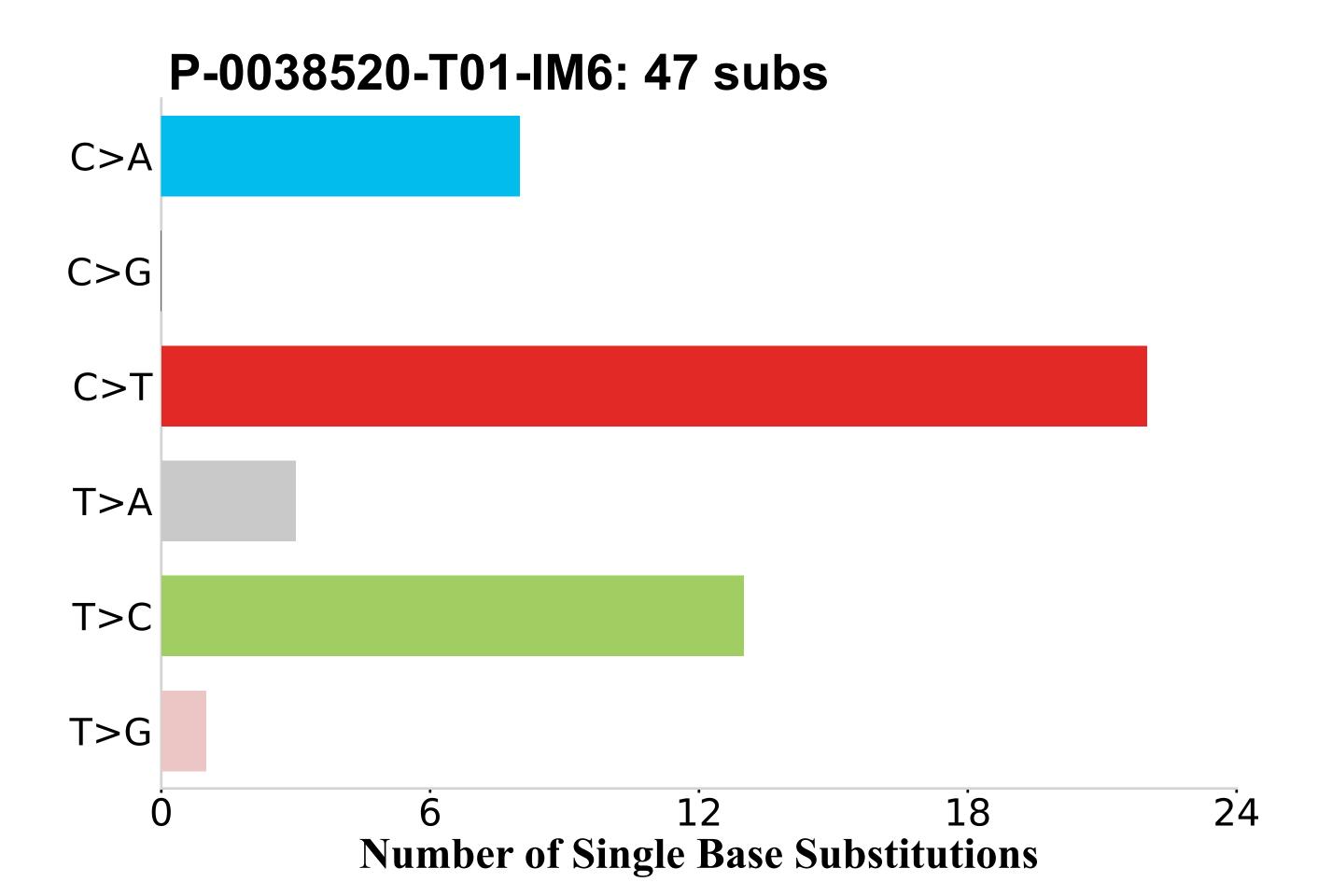


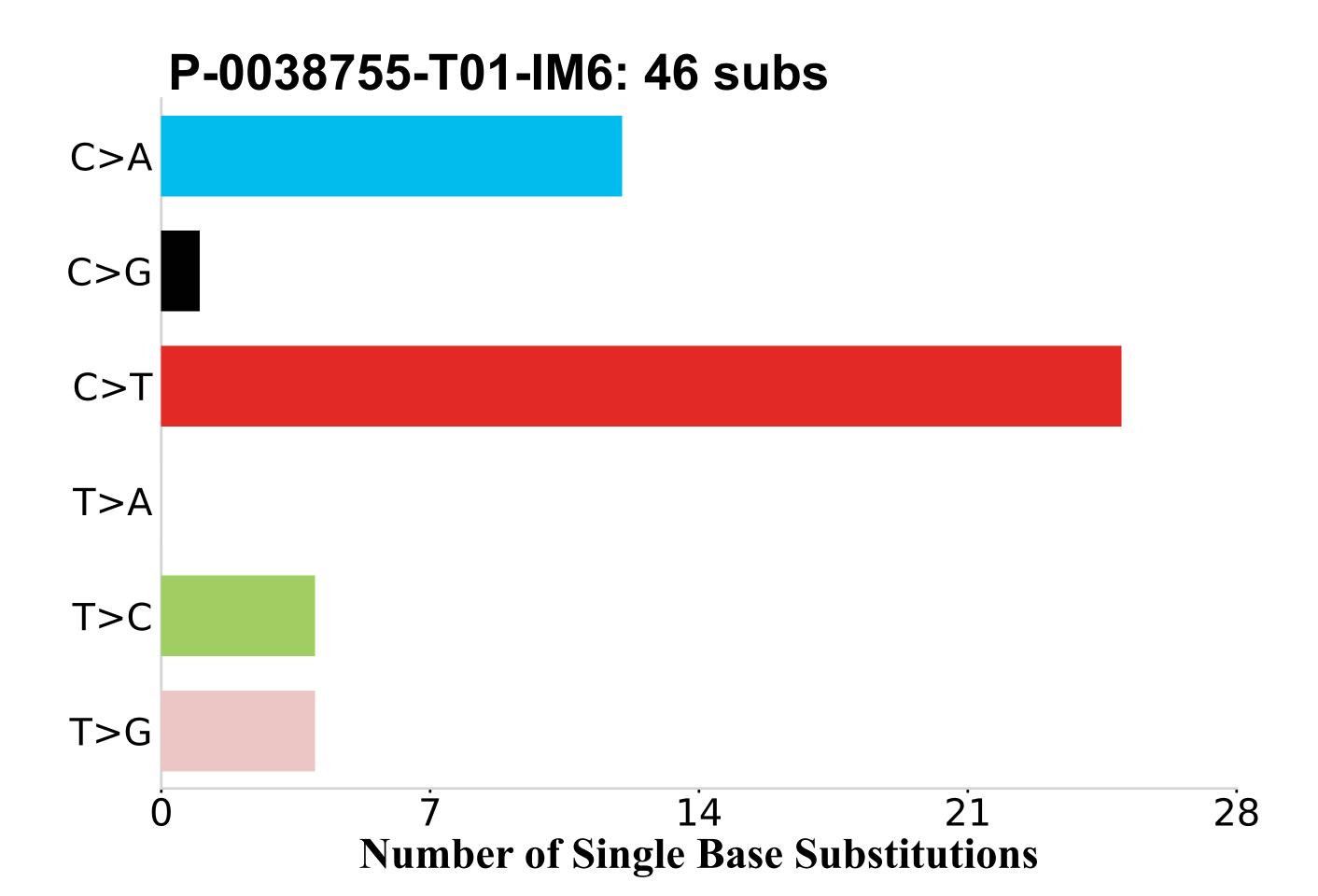


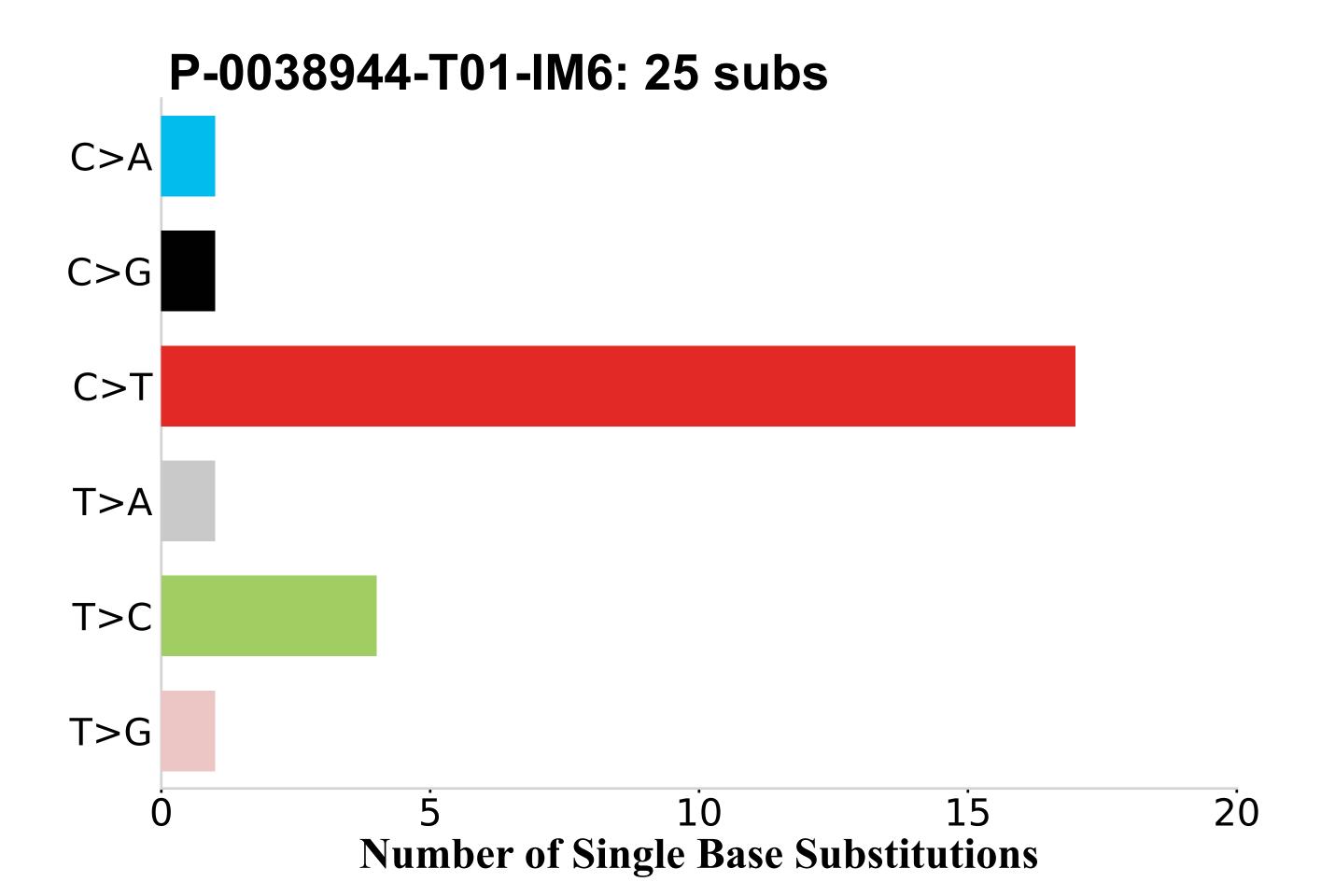


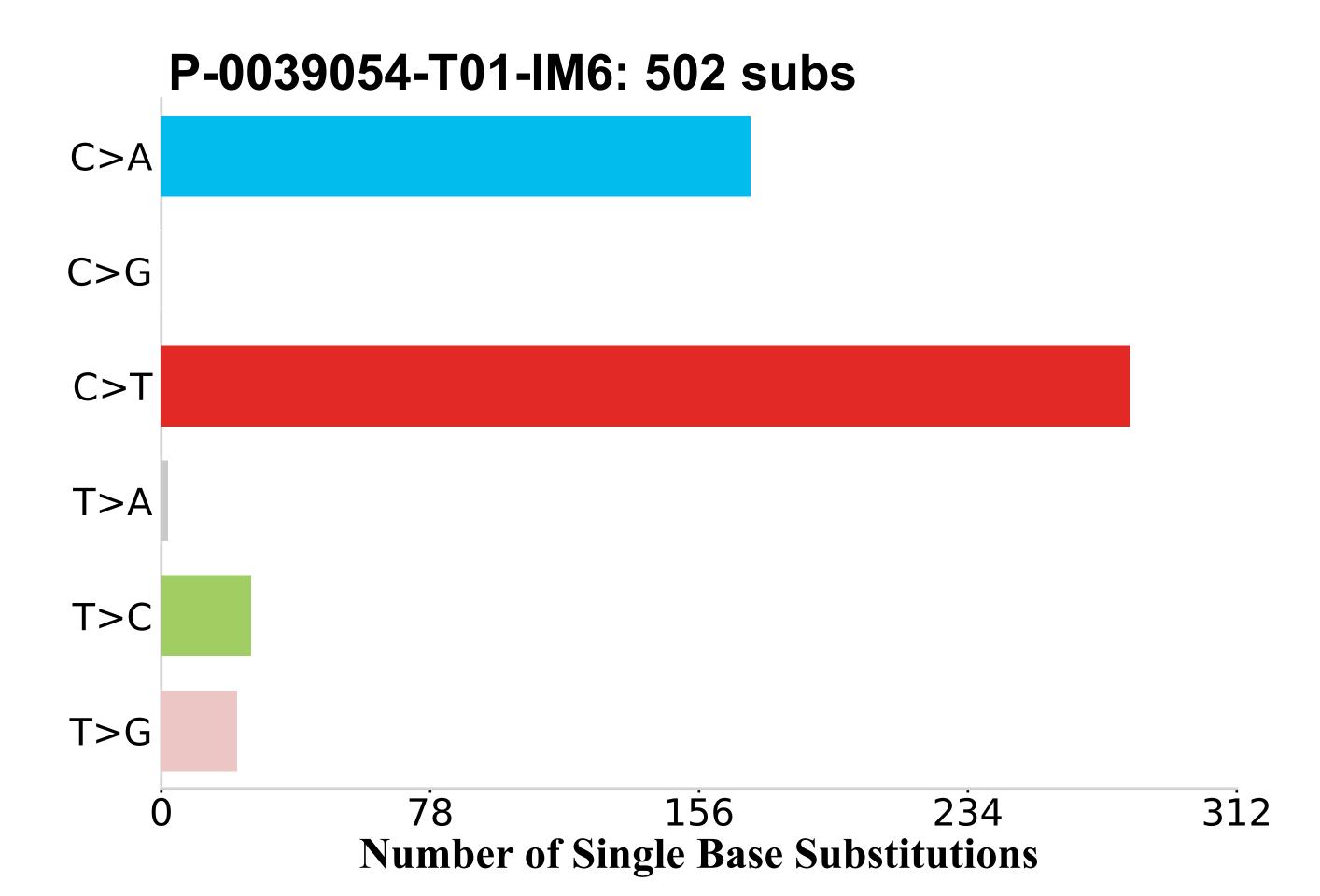


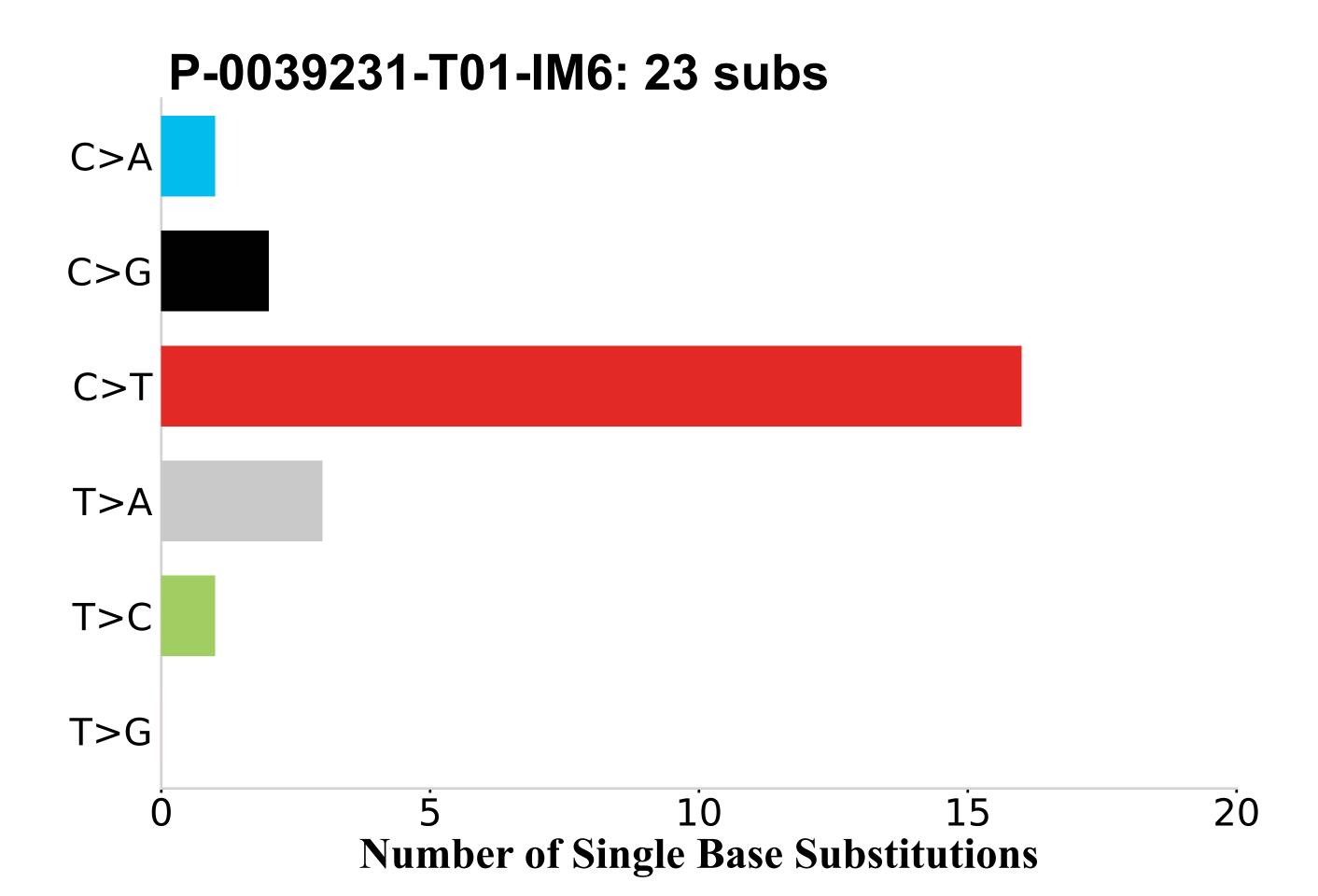


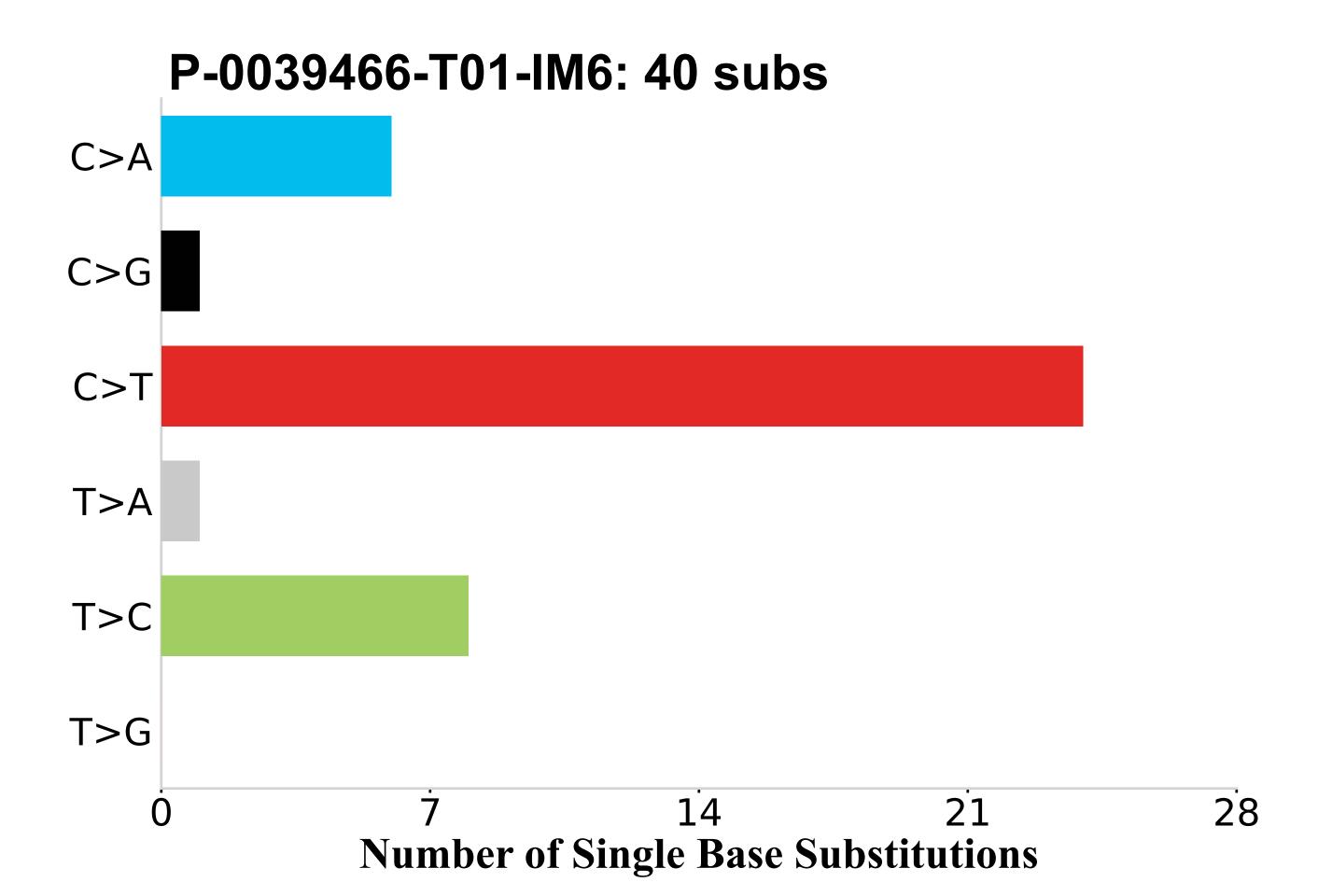


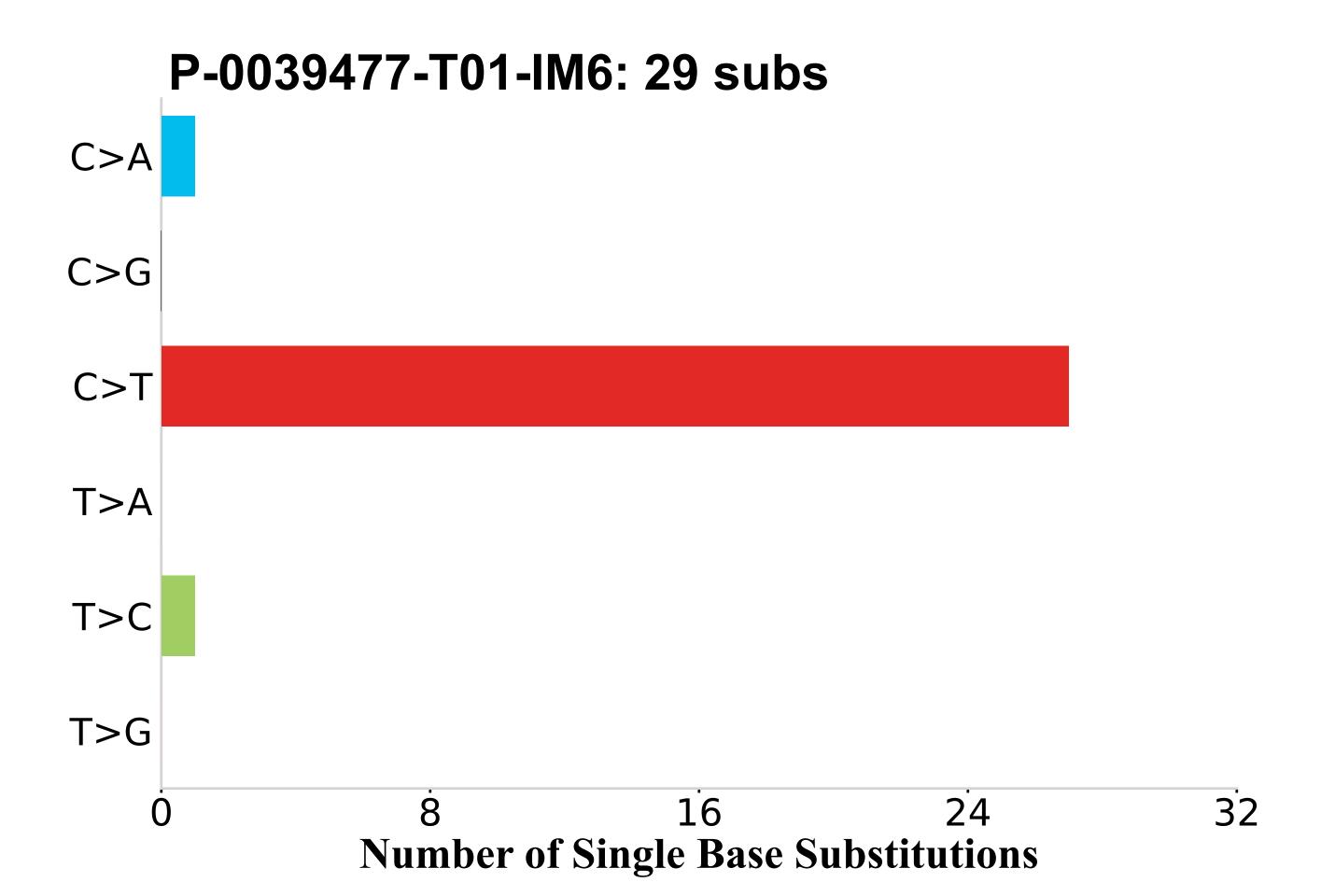


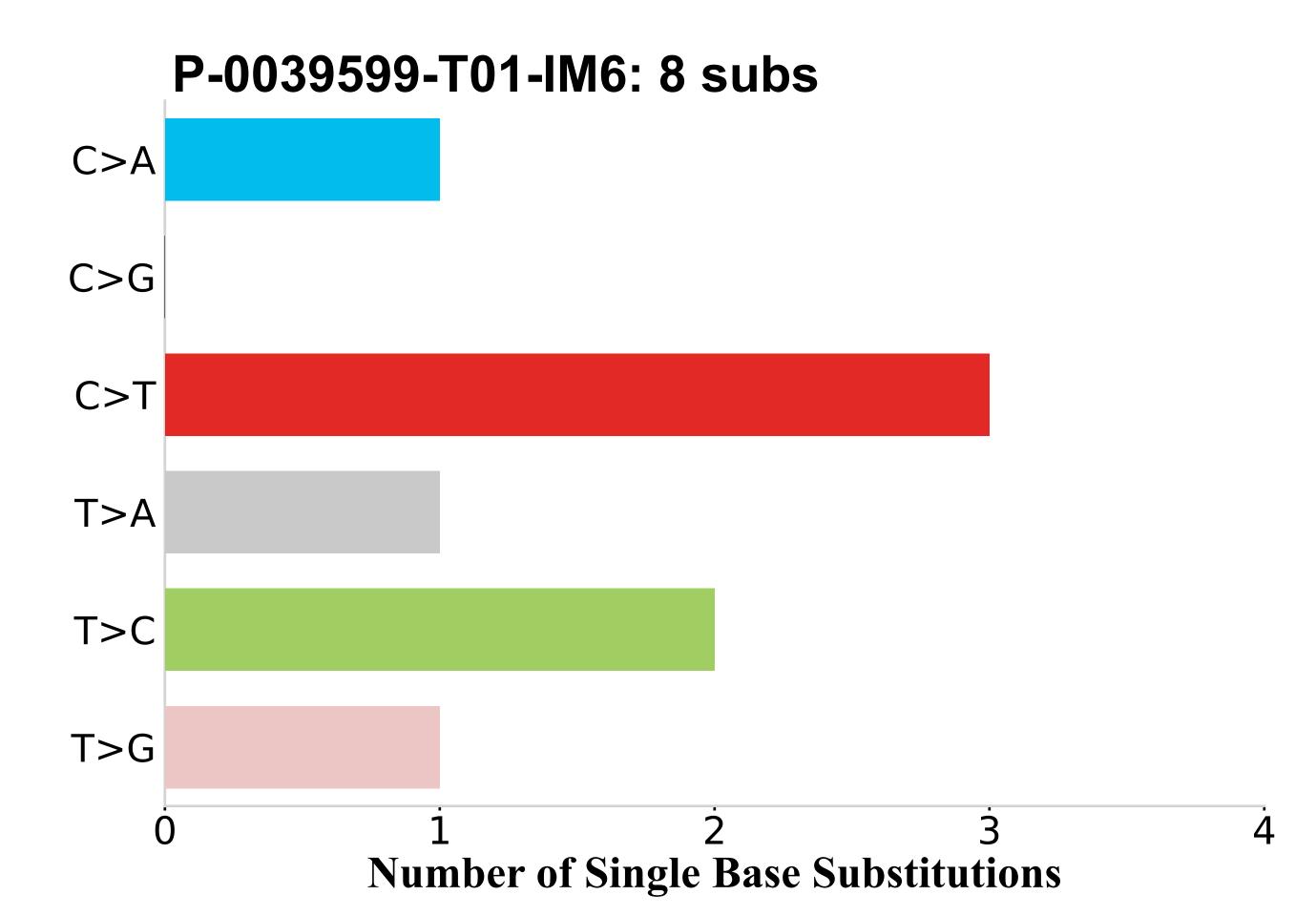


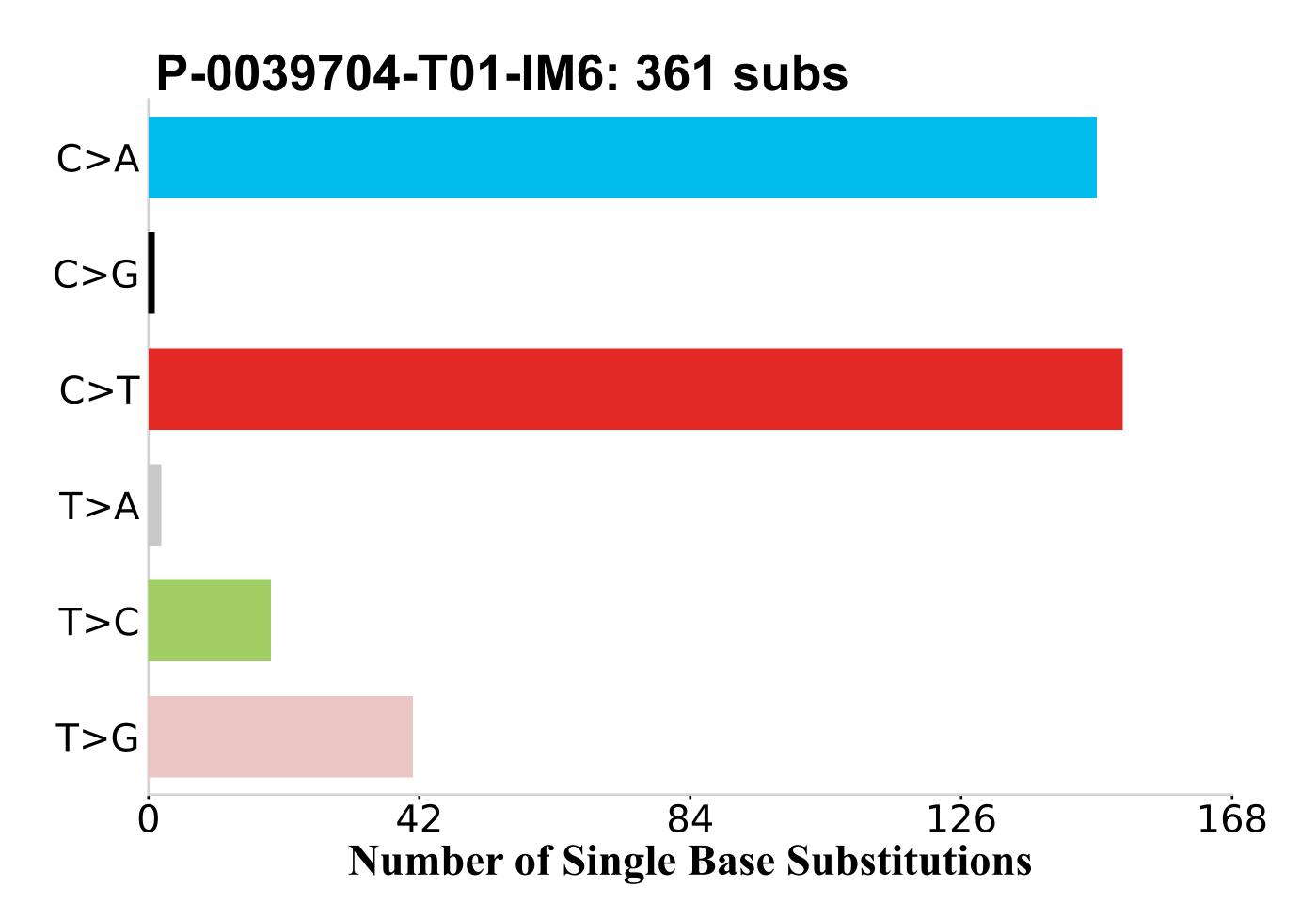


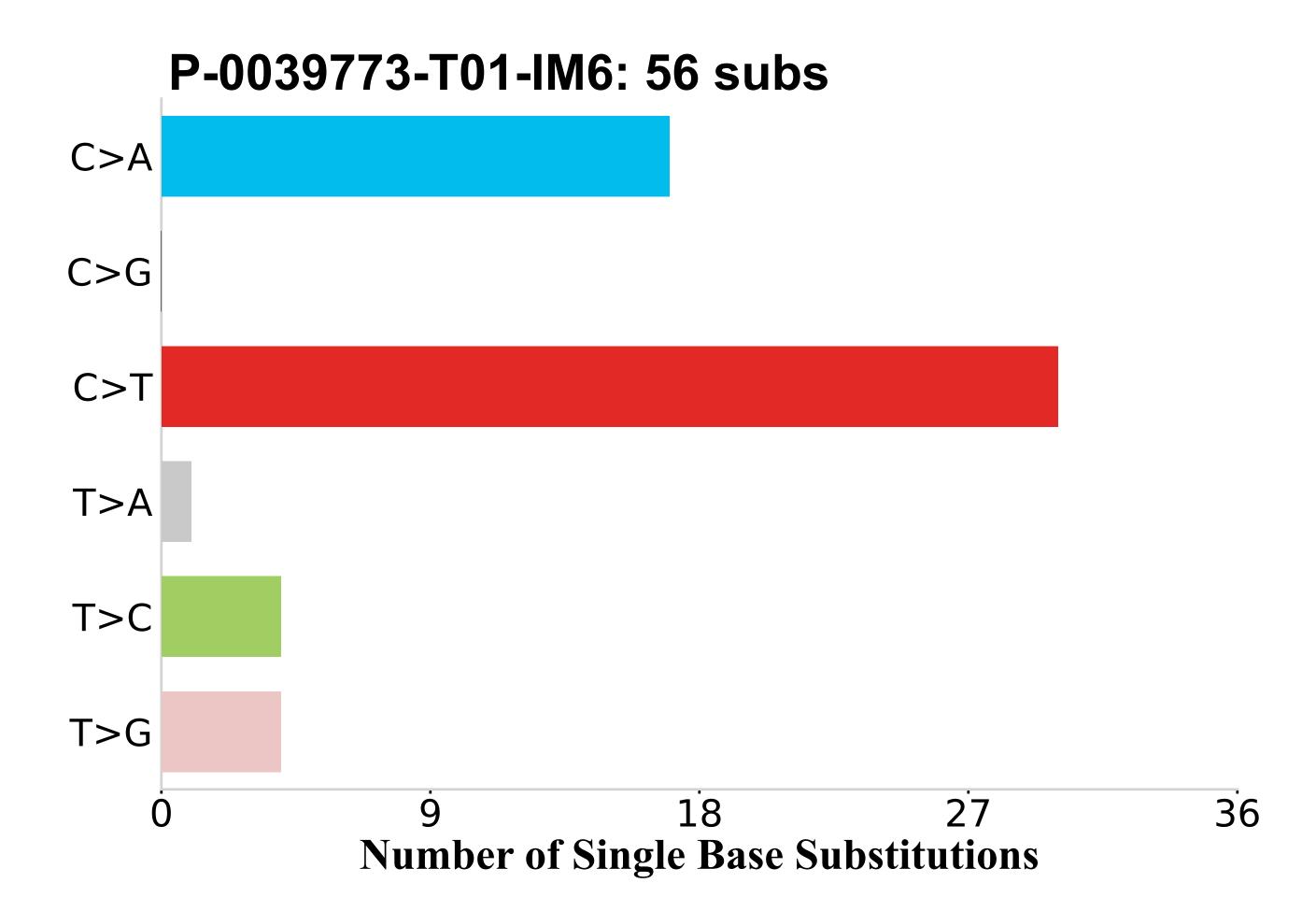


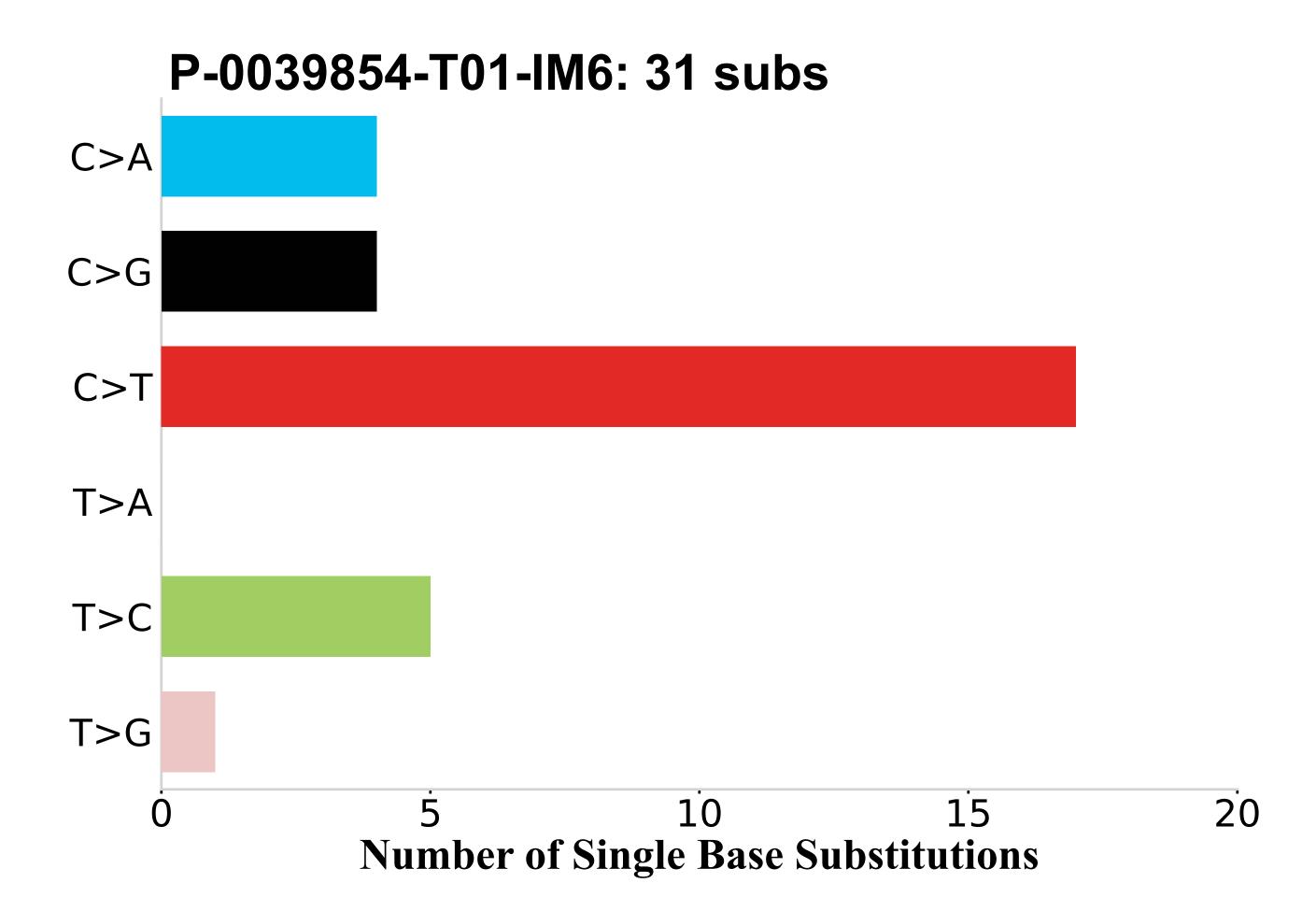


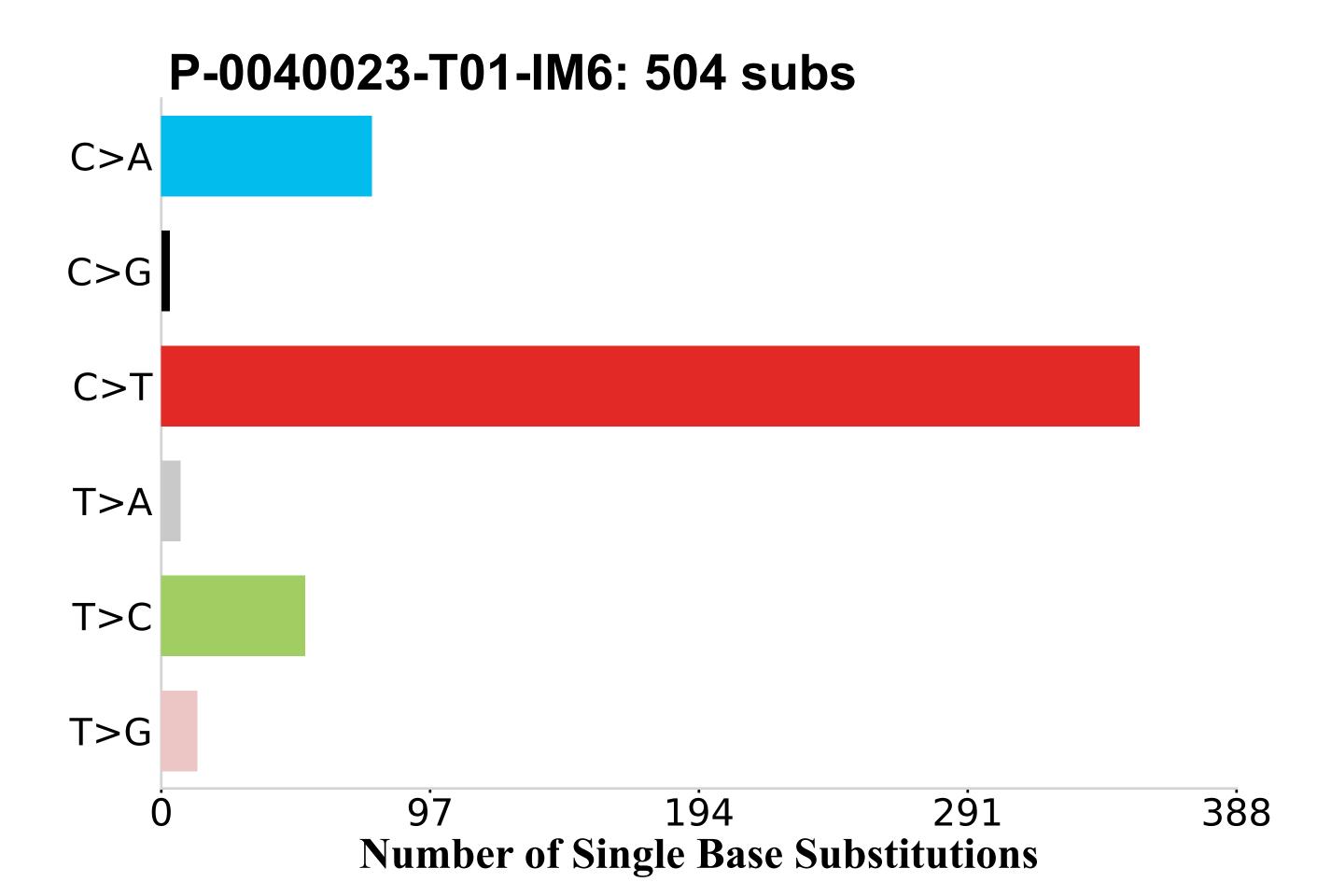


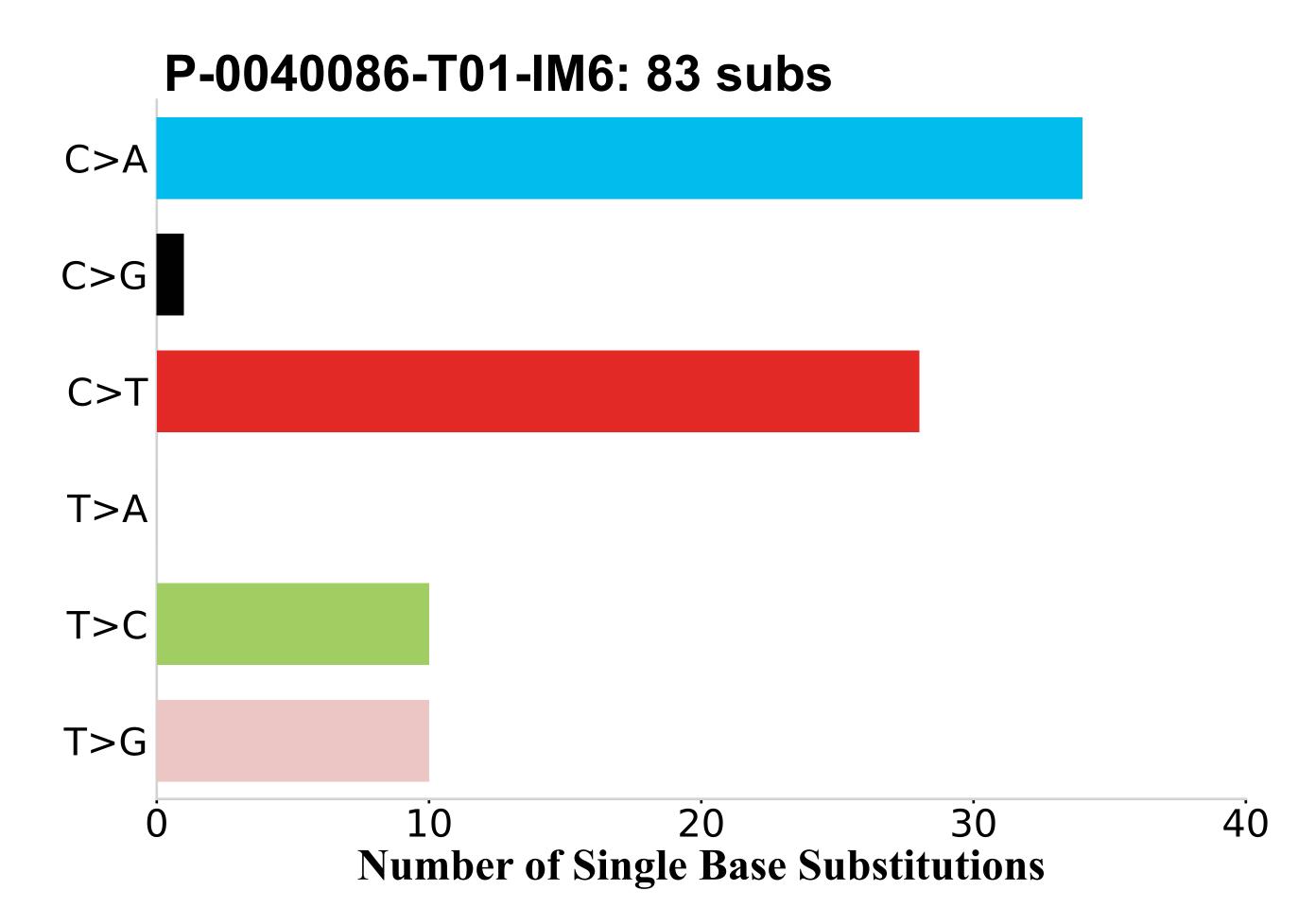


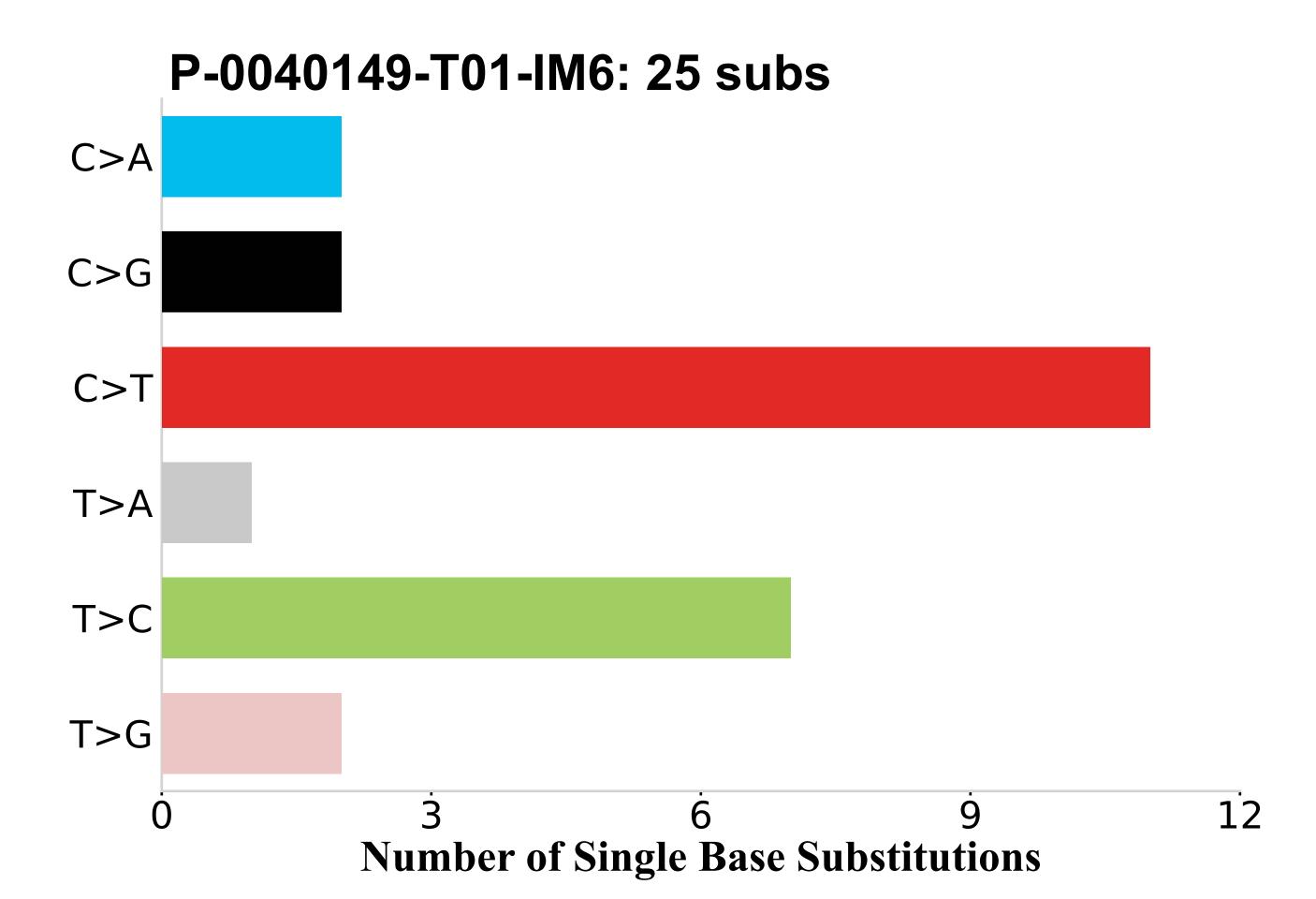




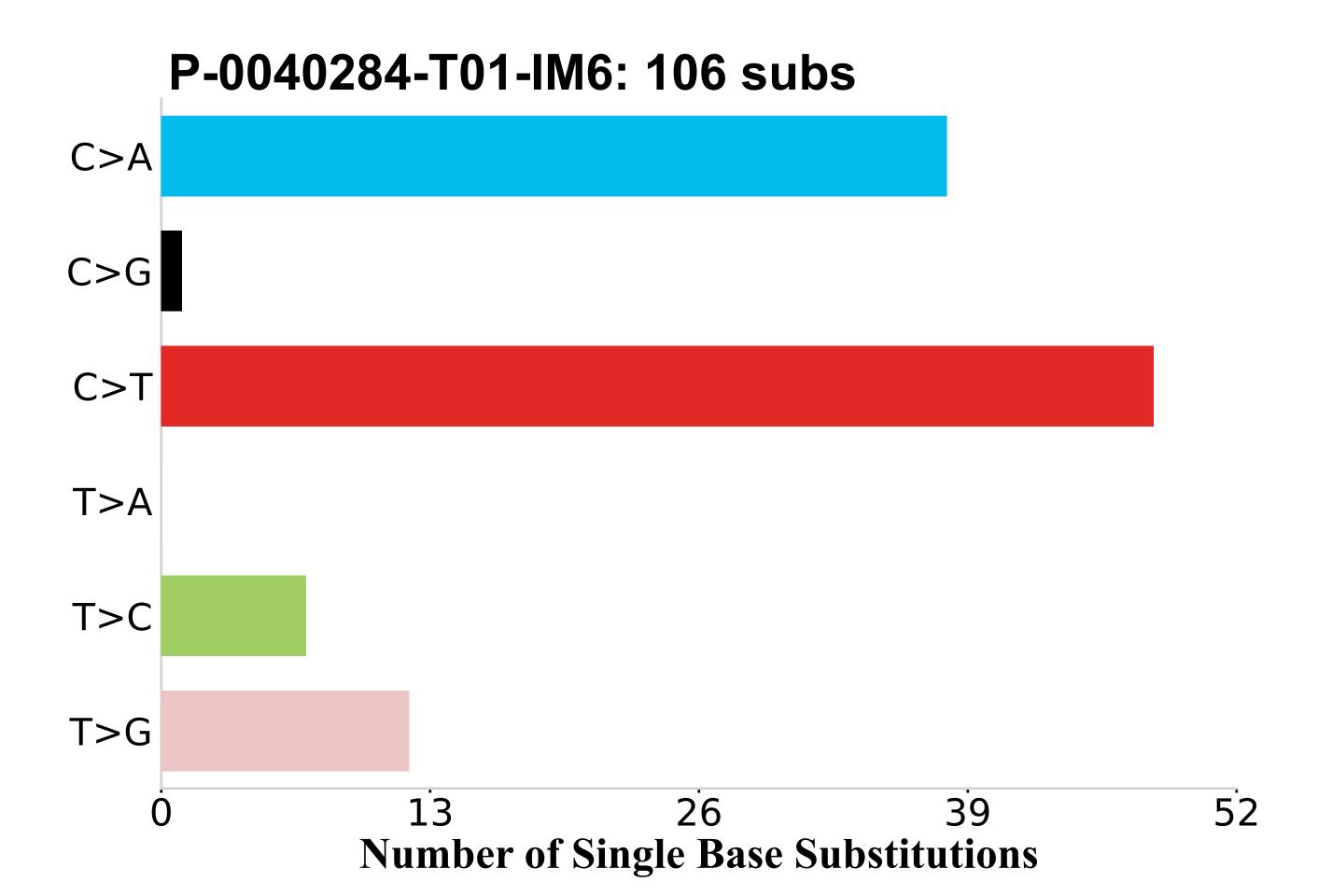


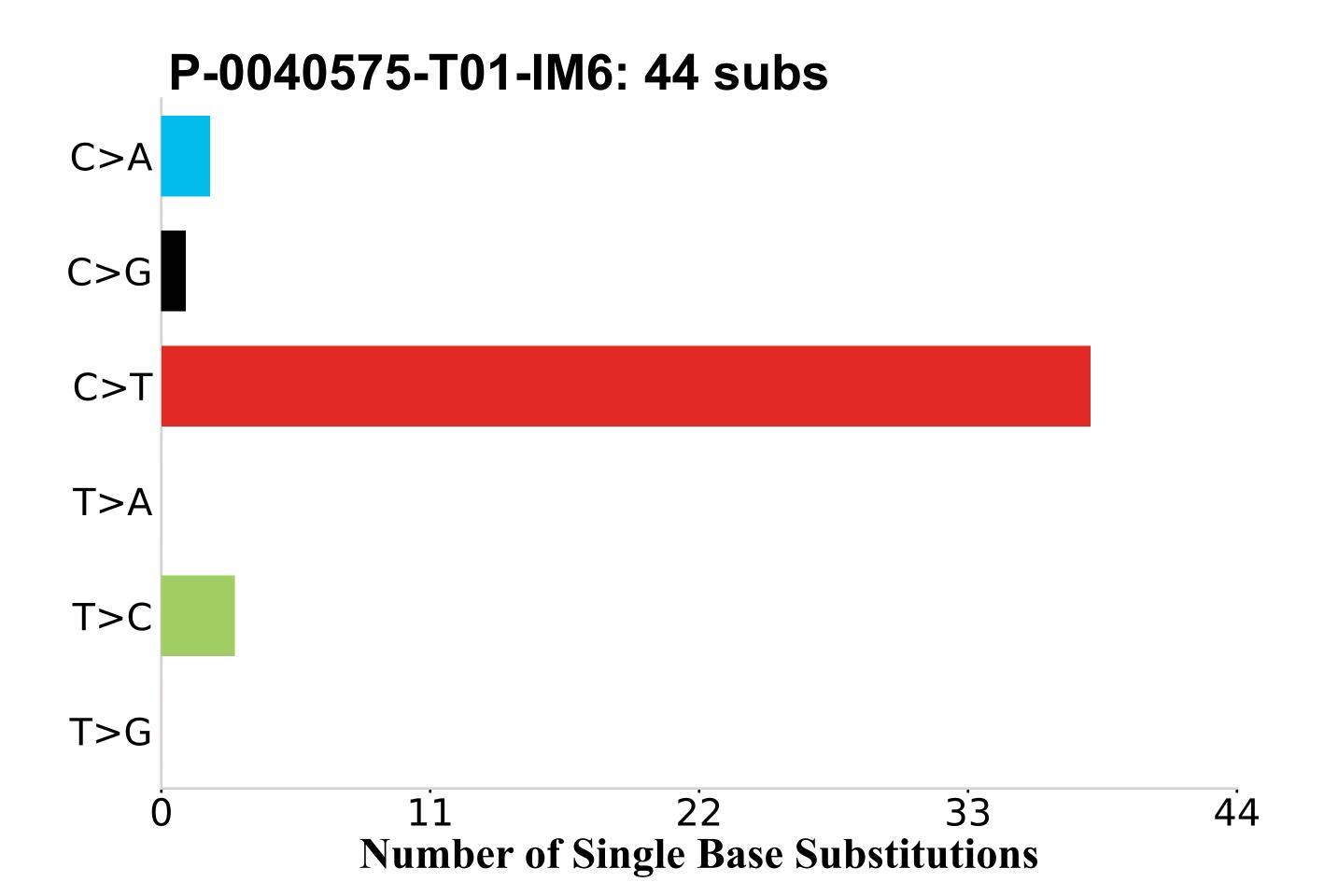


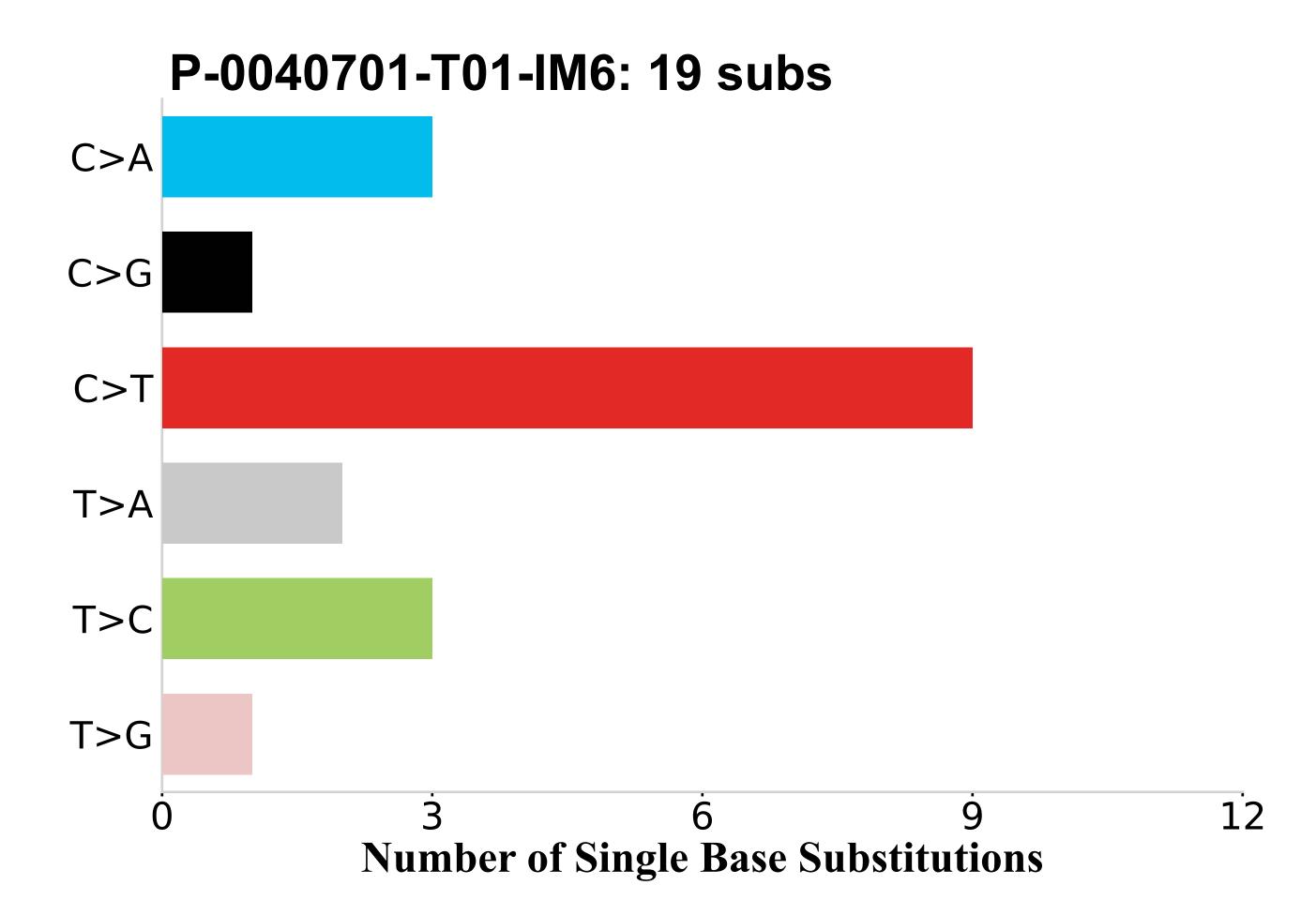


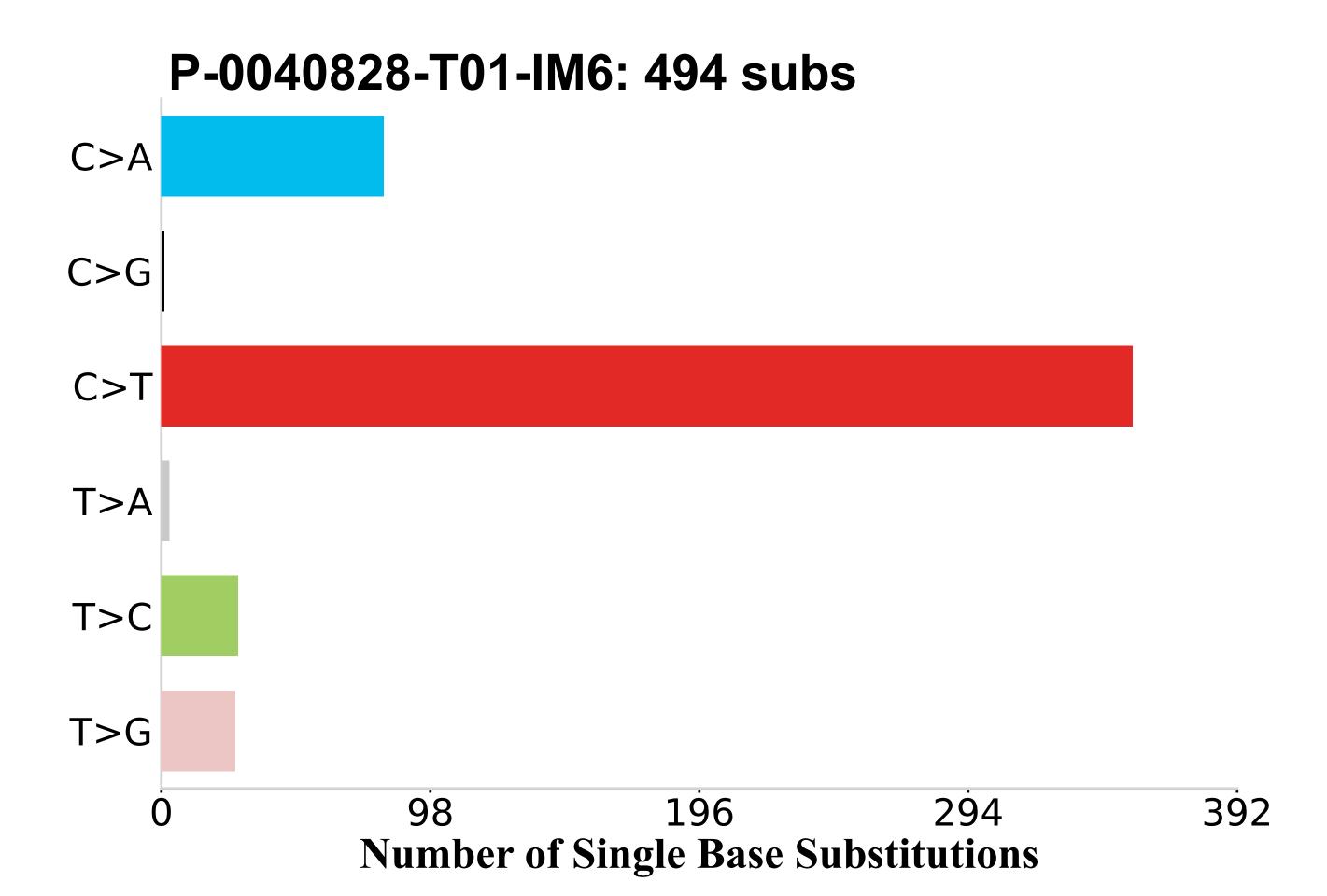


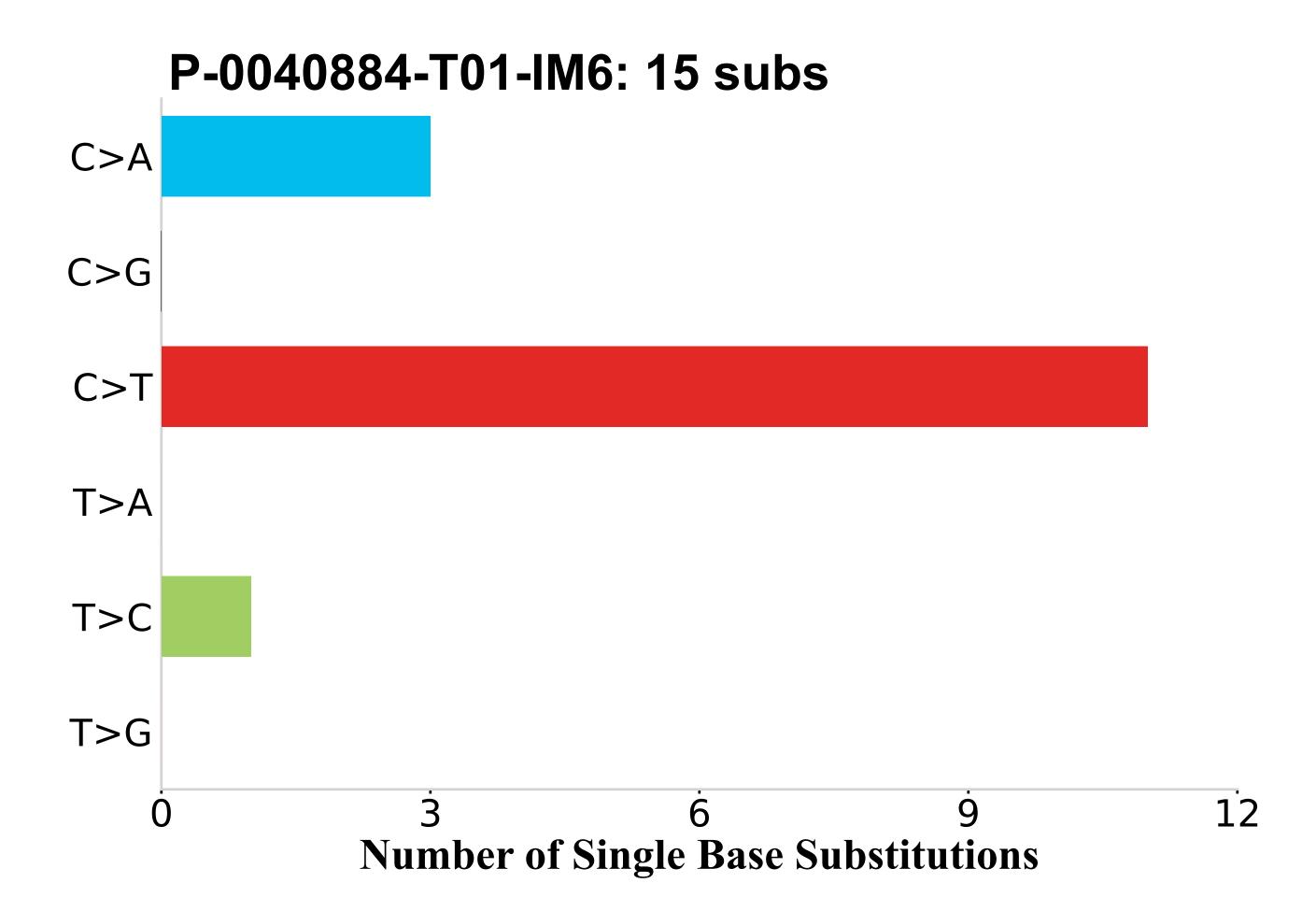
P-0040228-T01-IM6: 24 subs C>A C>G C>T T>A T>C T>G 16 12 **Number of Single Base Substitutions**

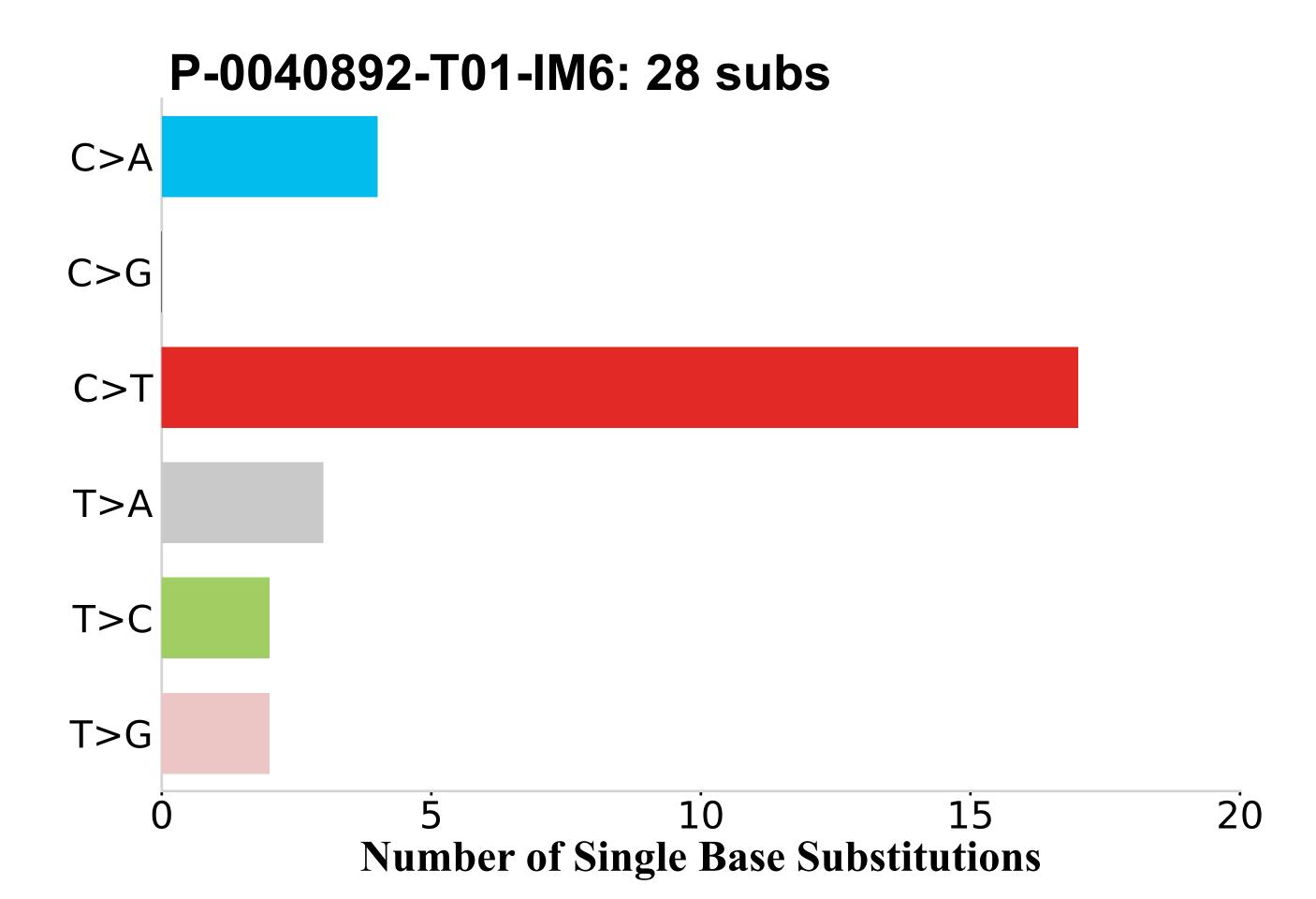


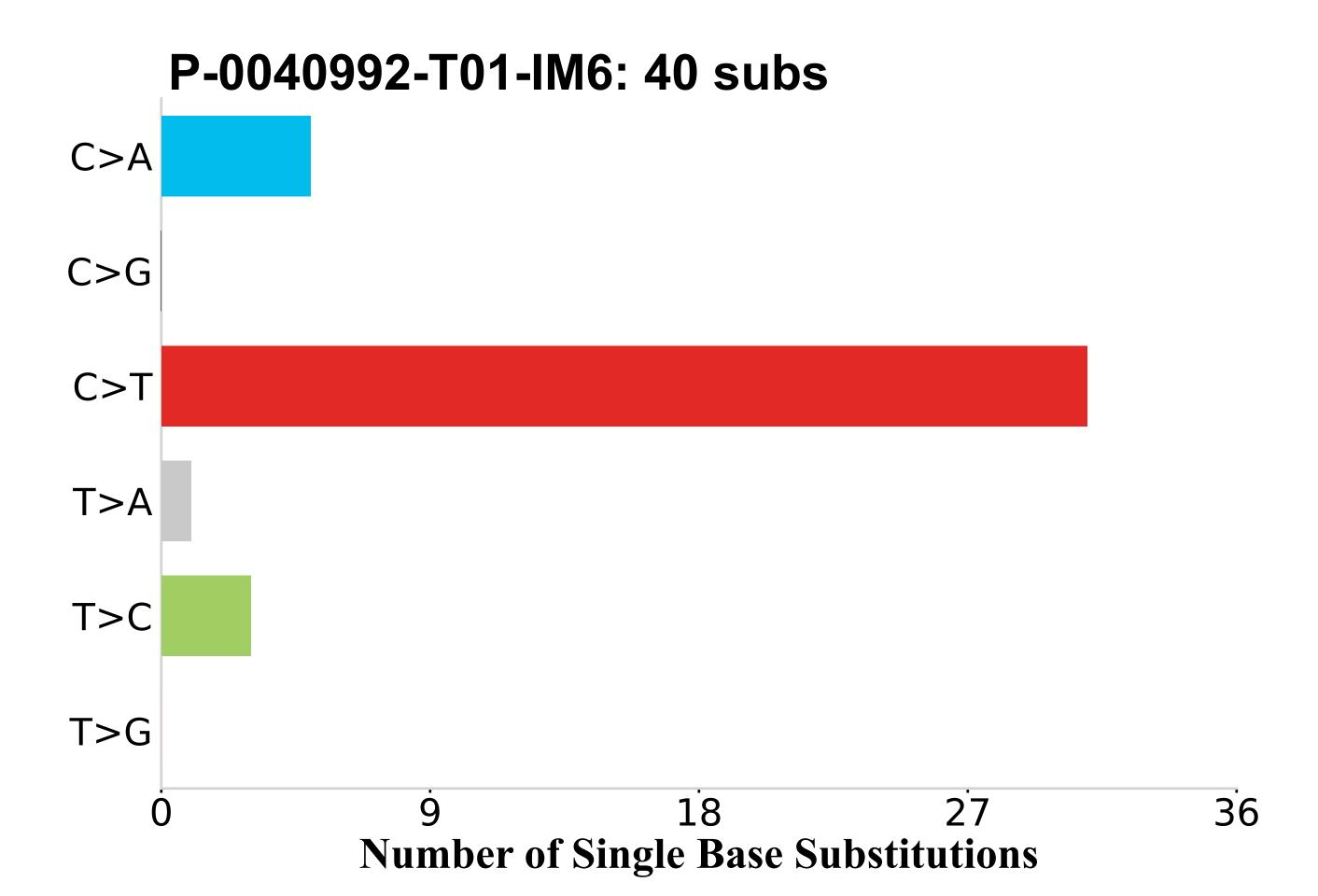




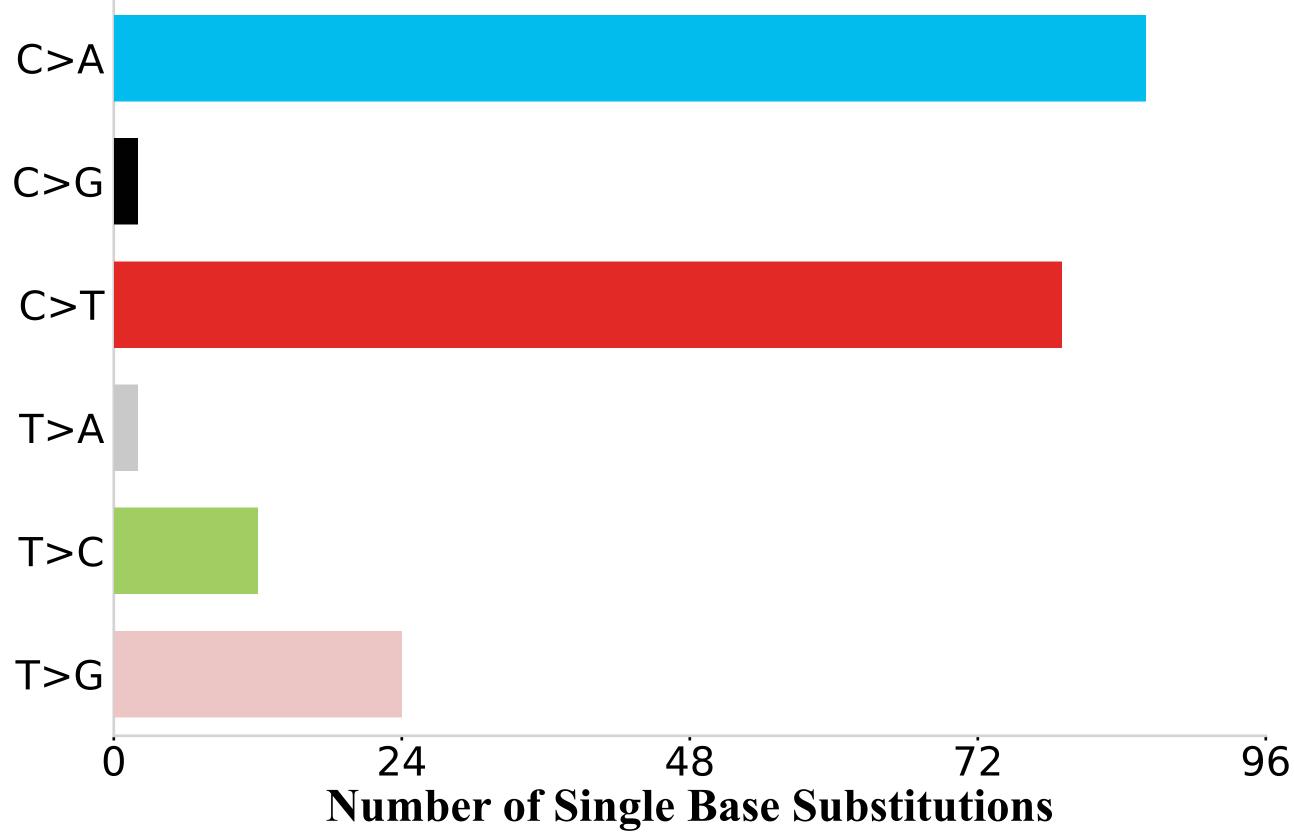


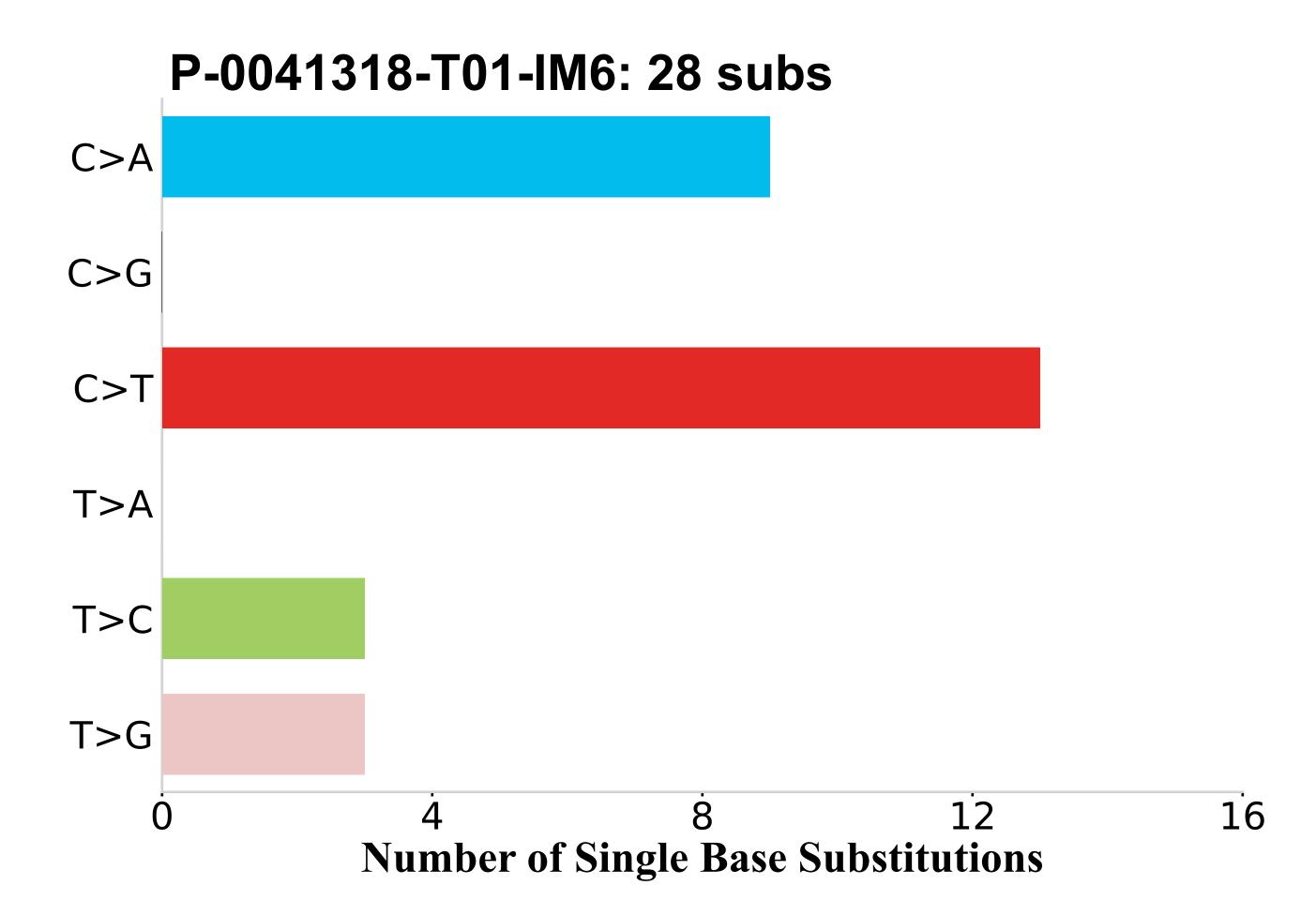




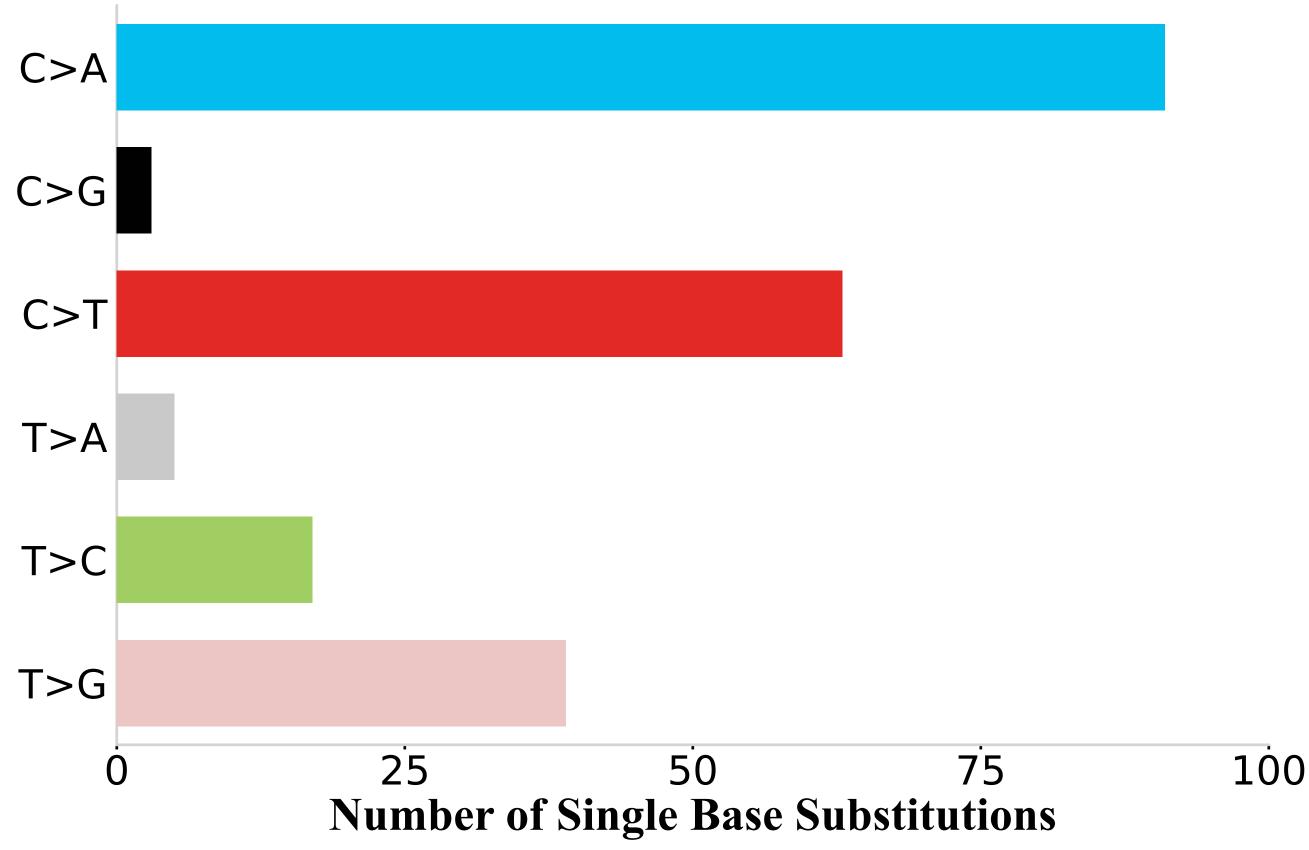


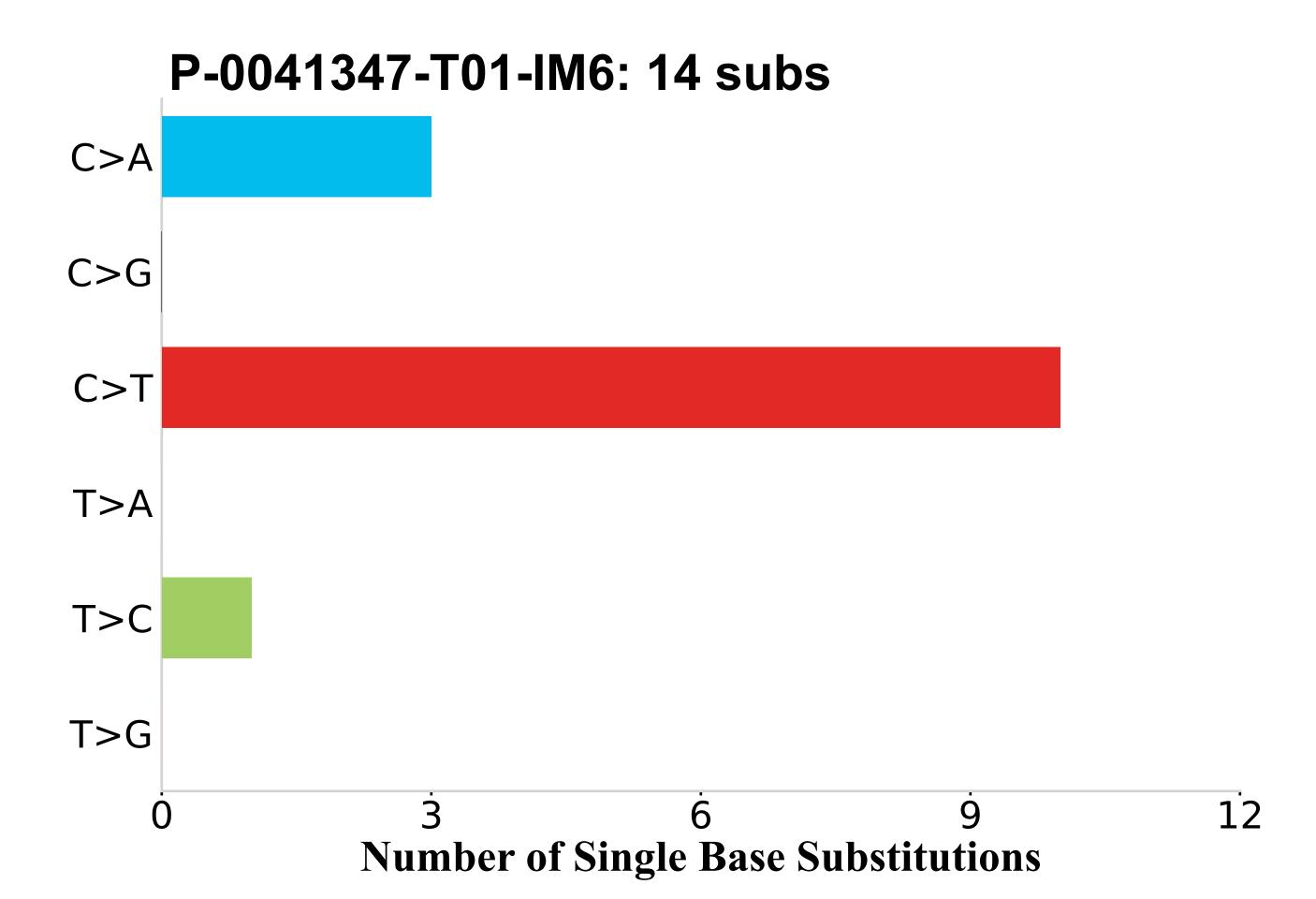
P-0041146-T01-IM6: 205 subs

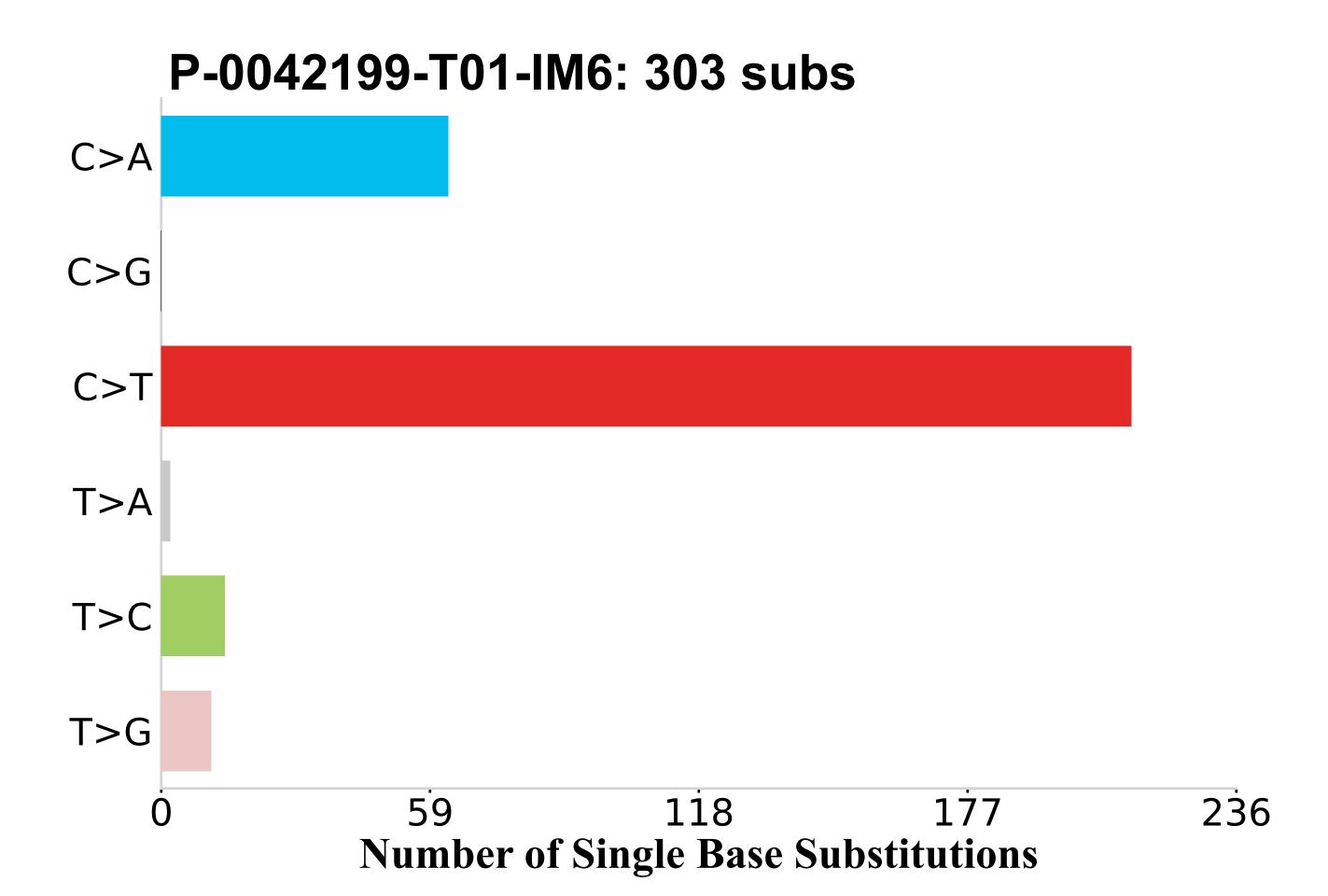


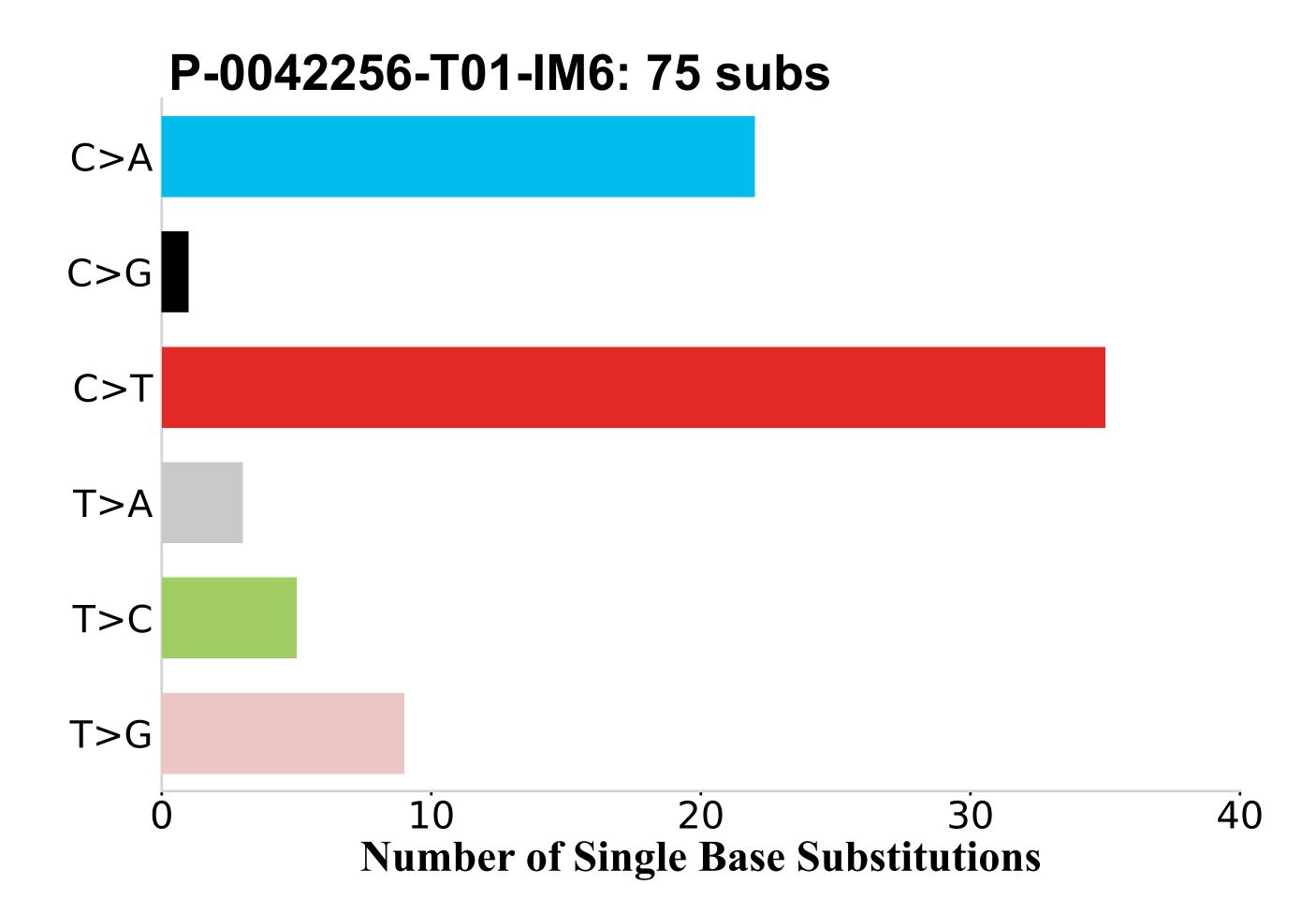


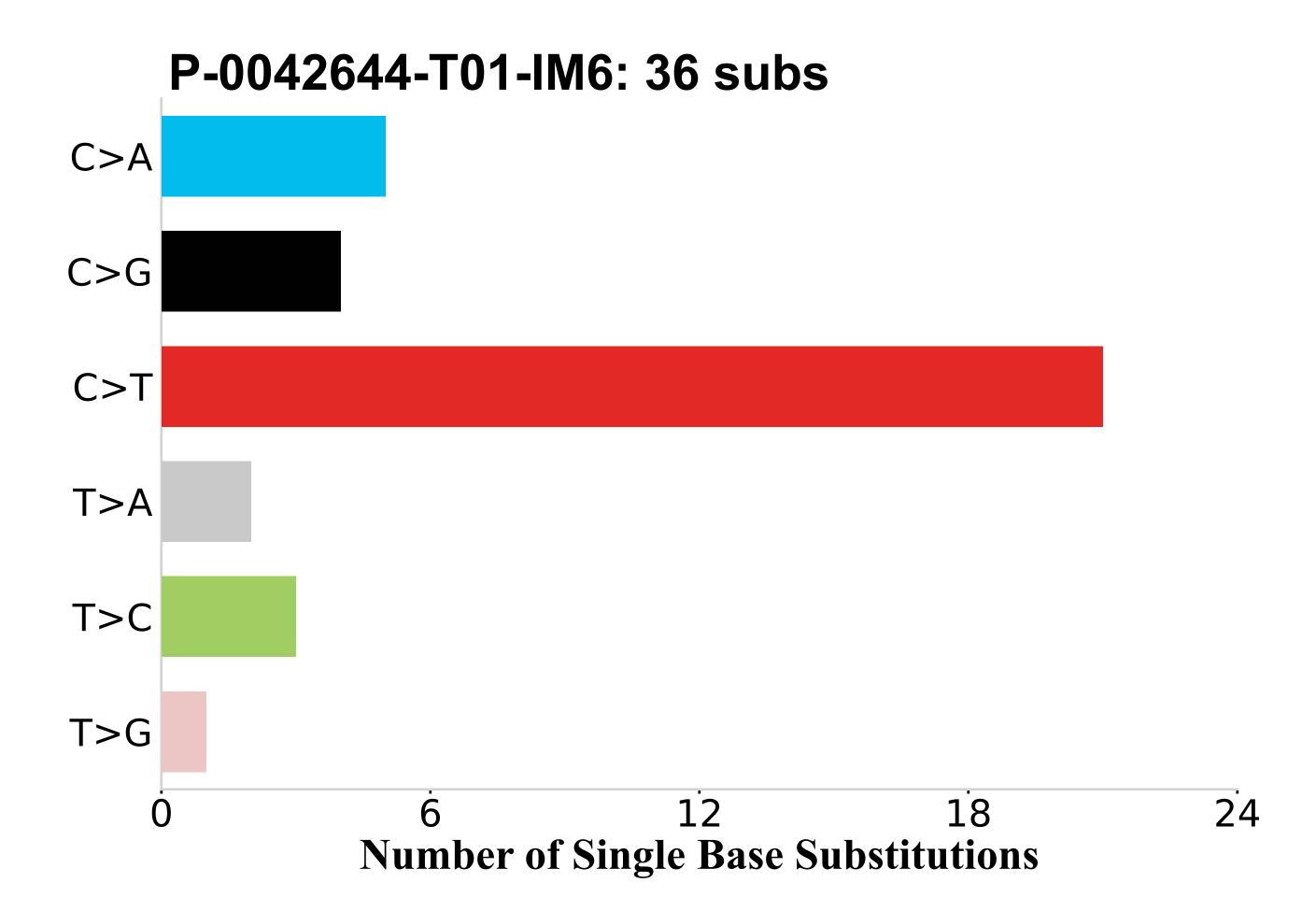


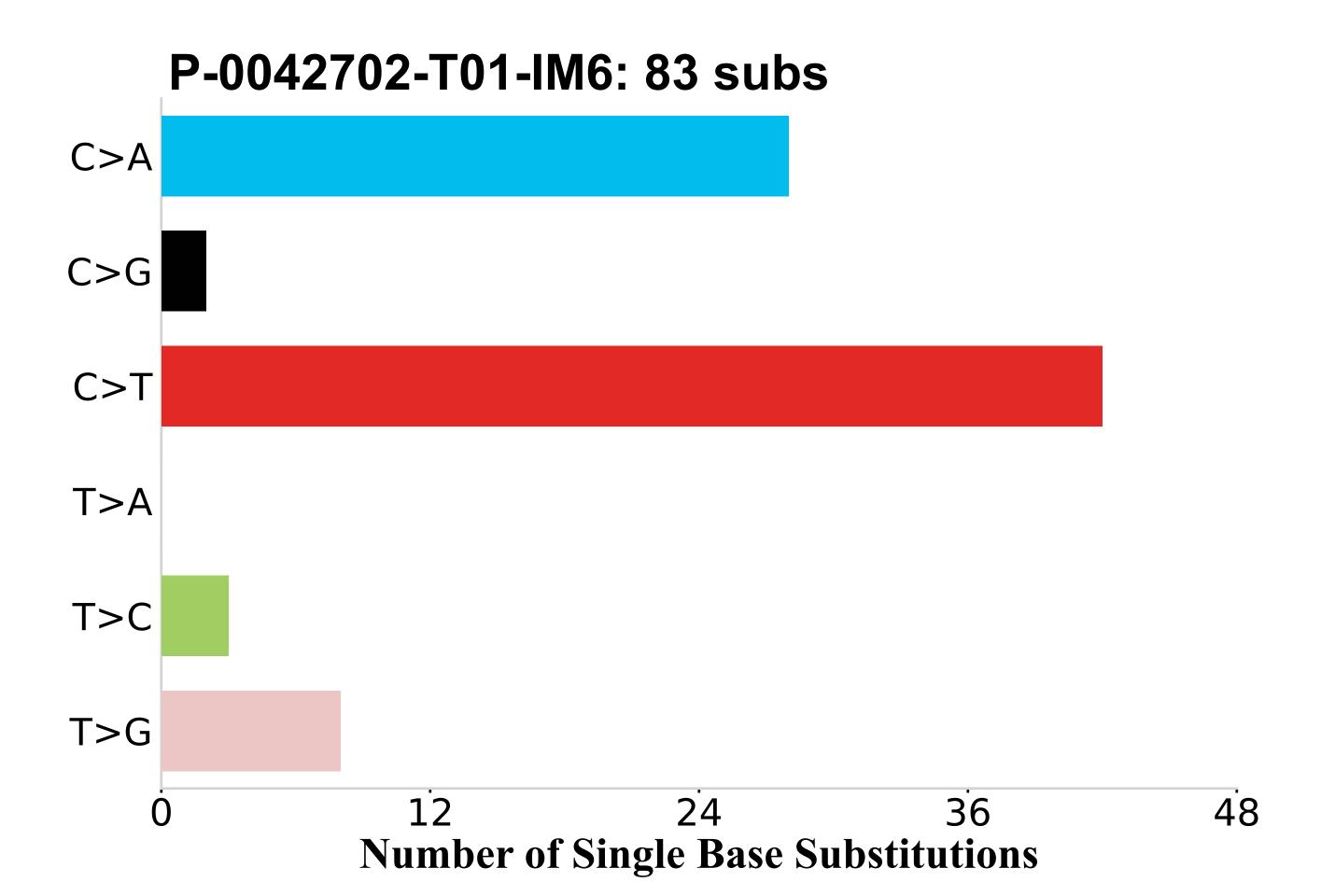


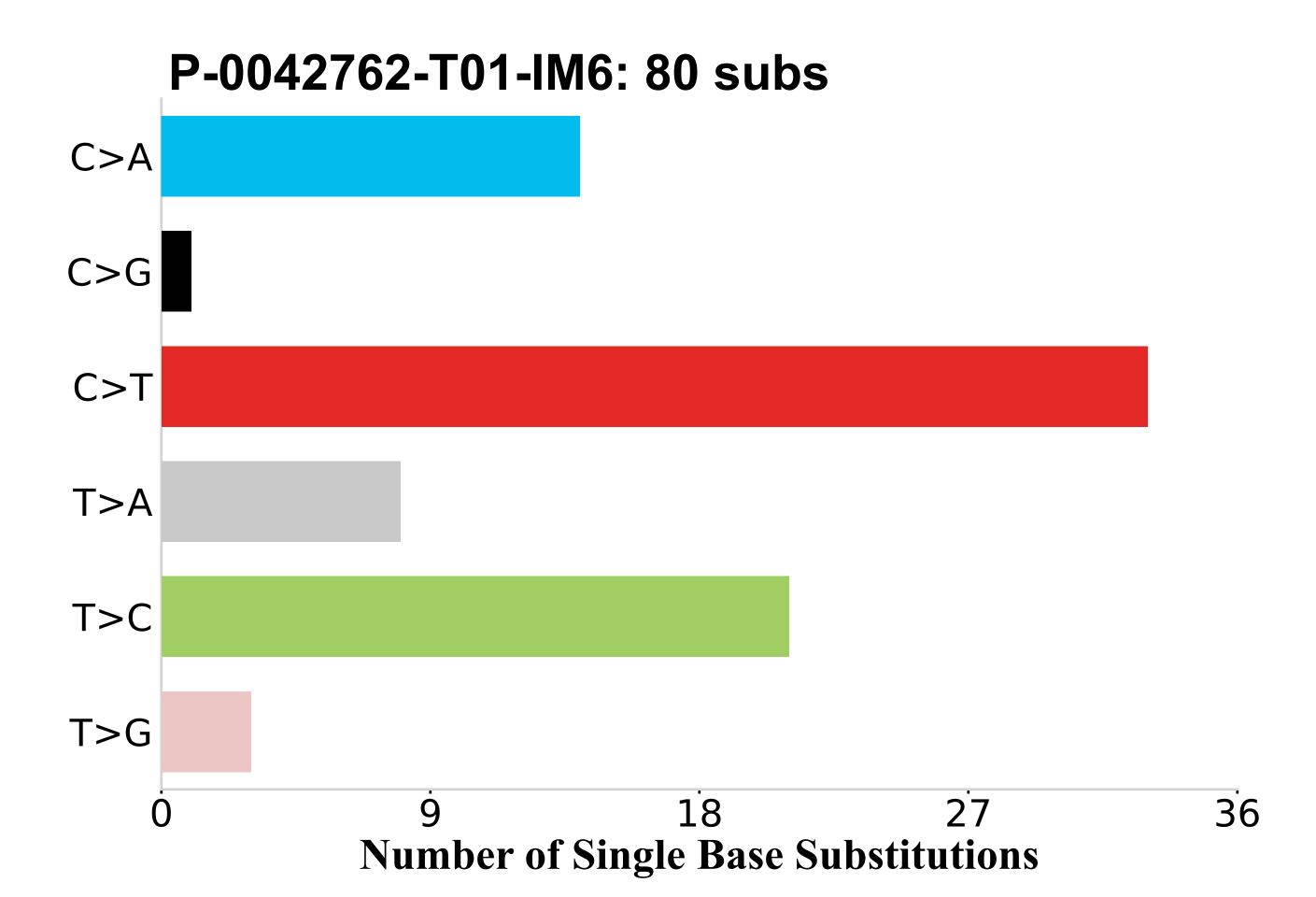


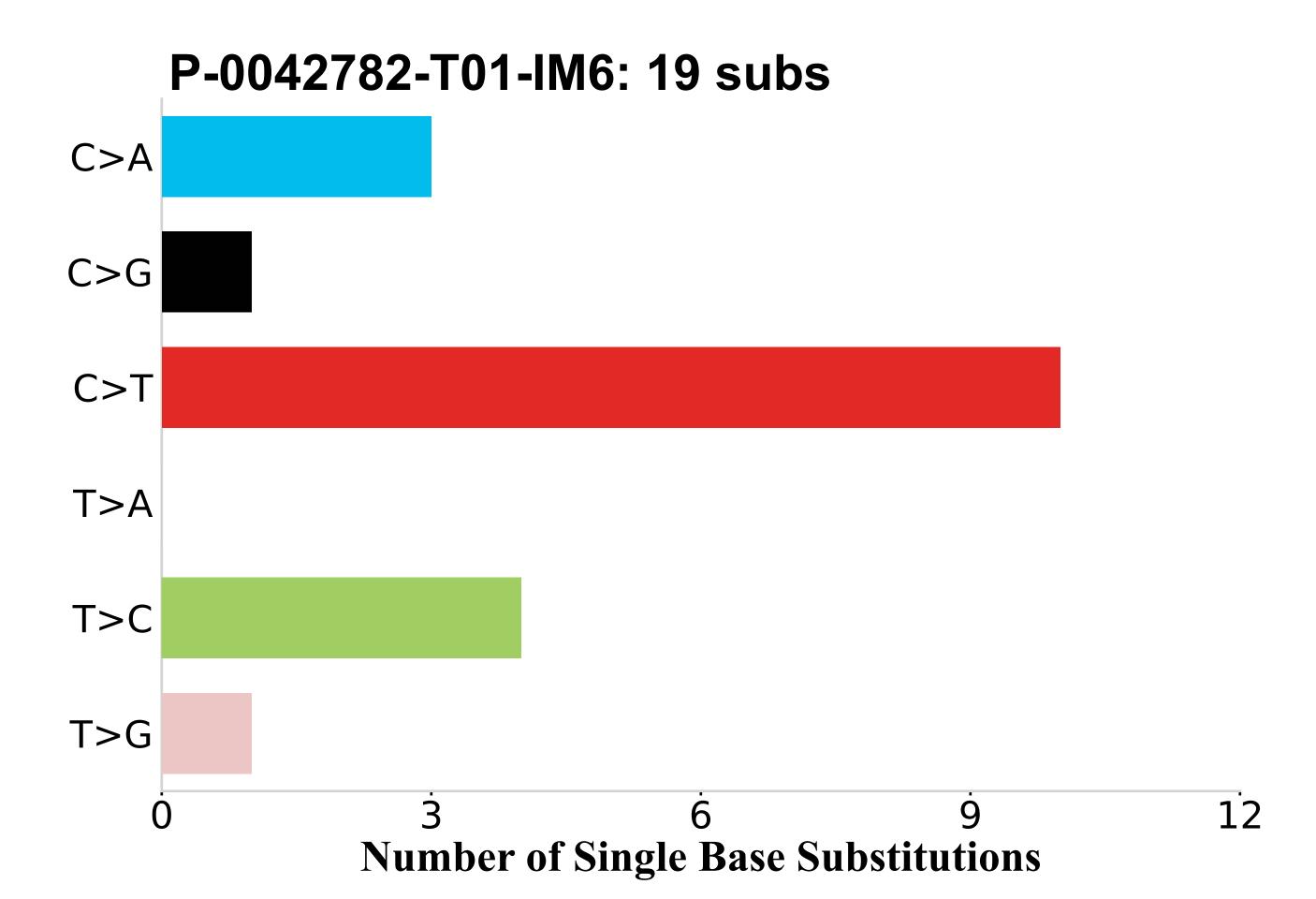


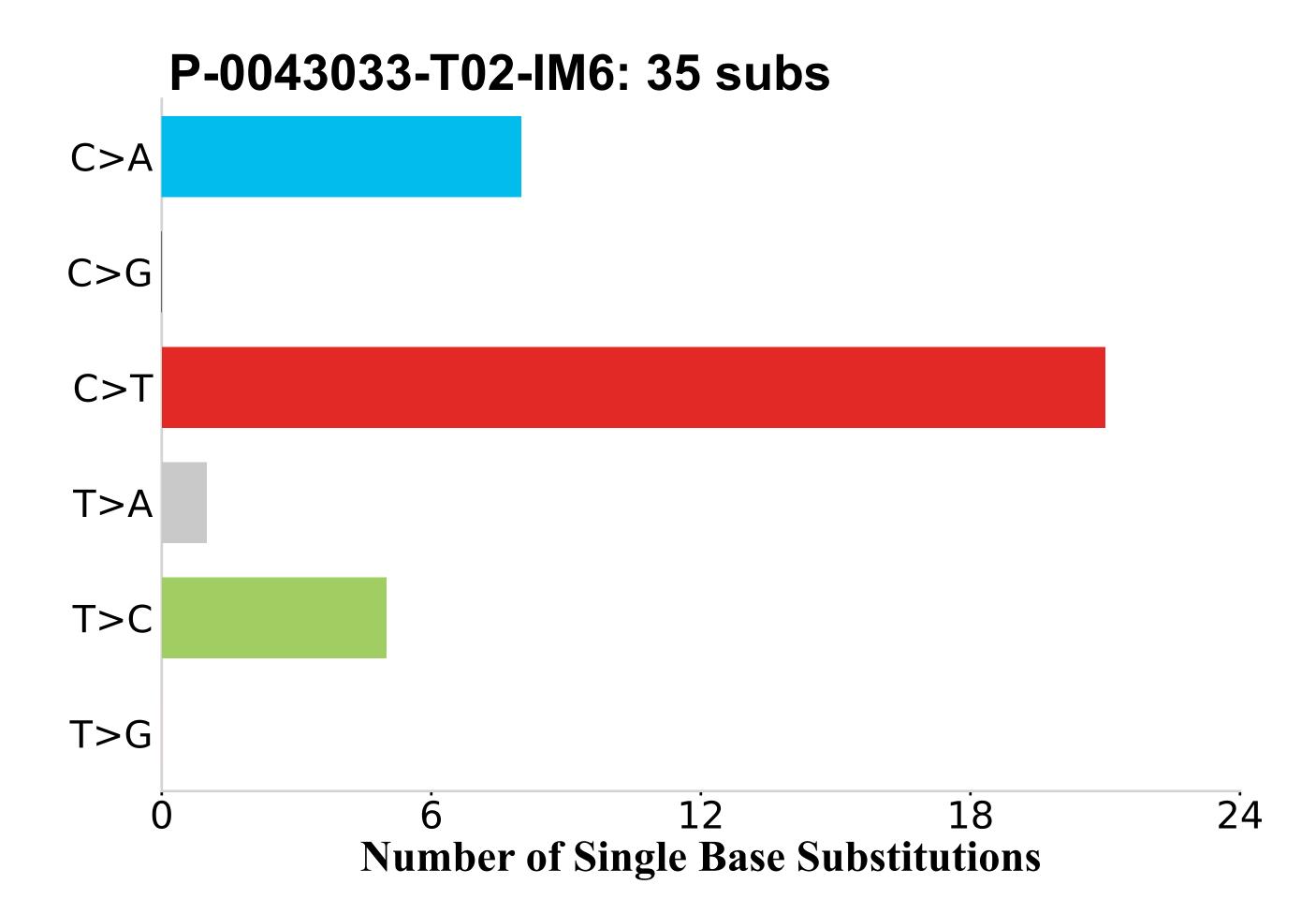


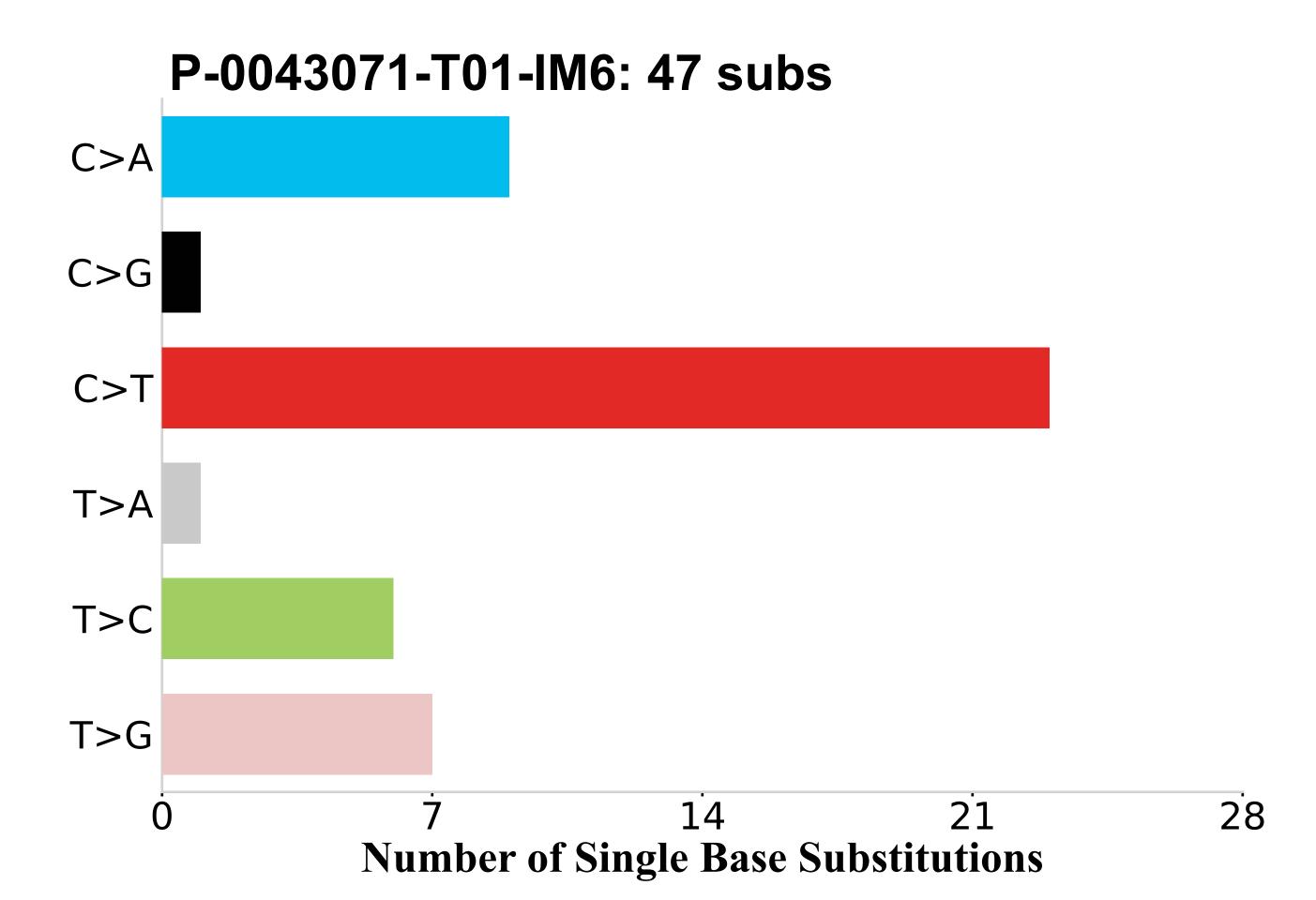


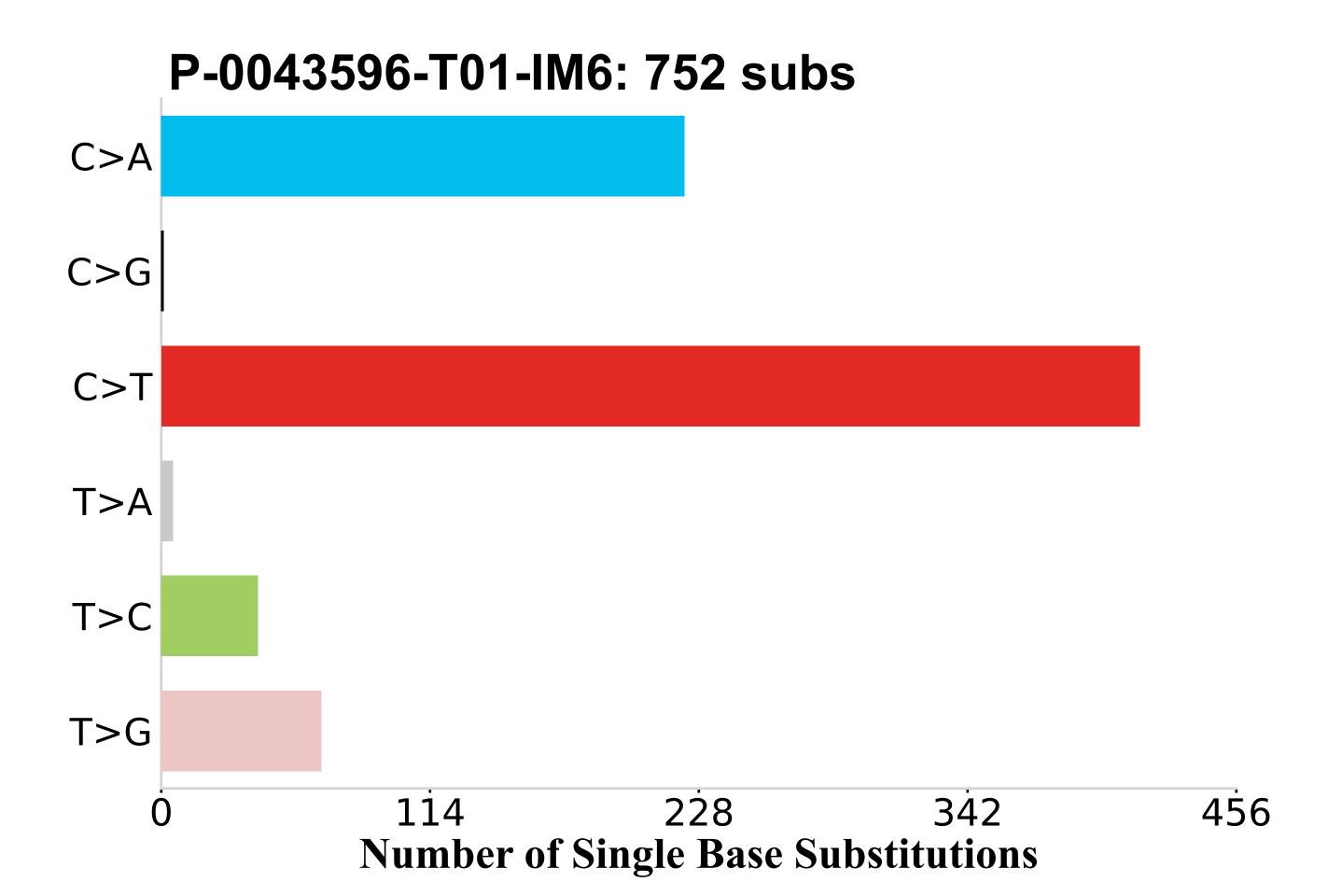


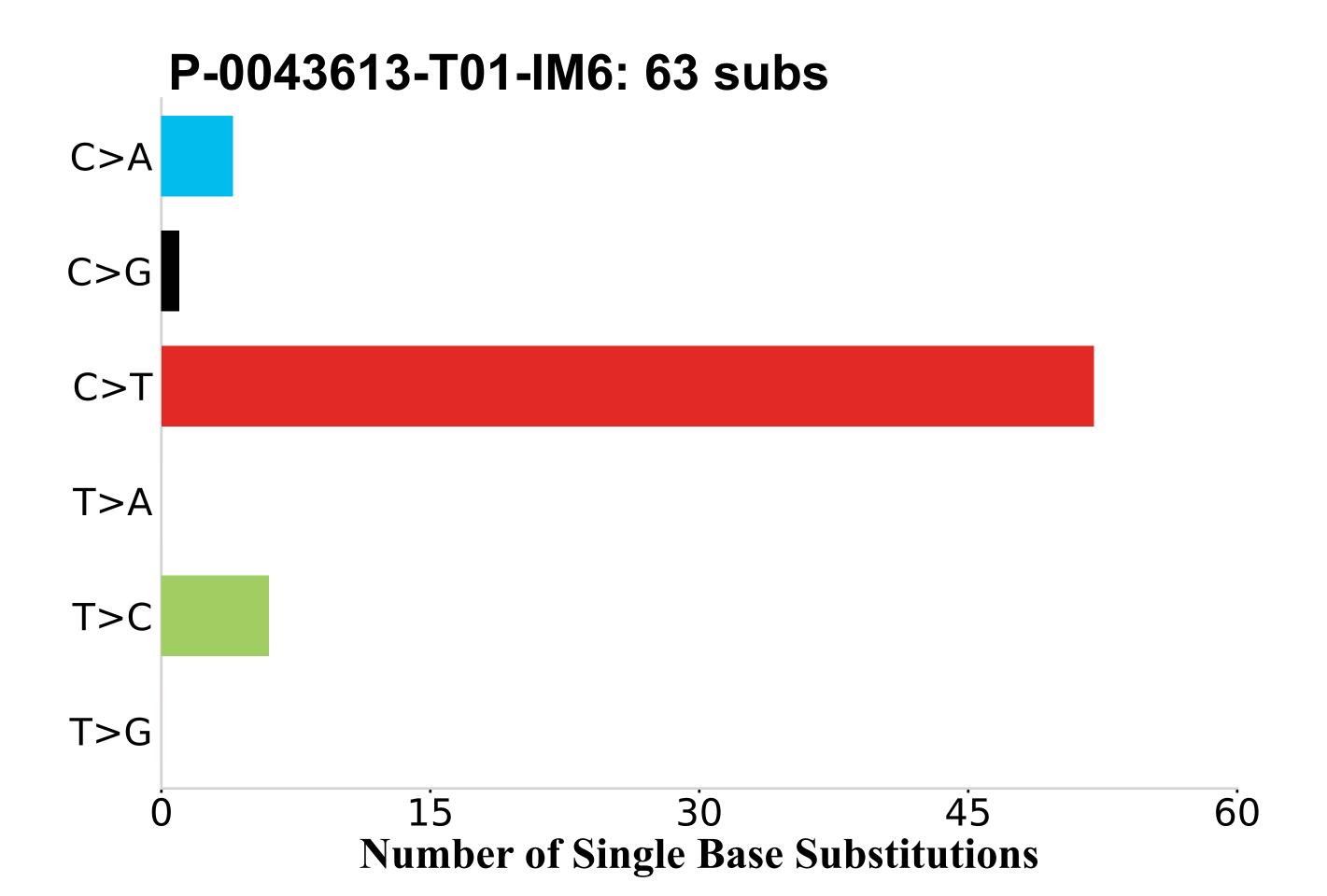


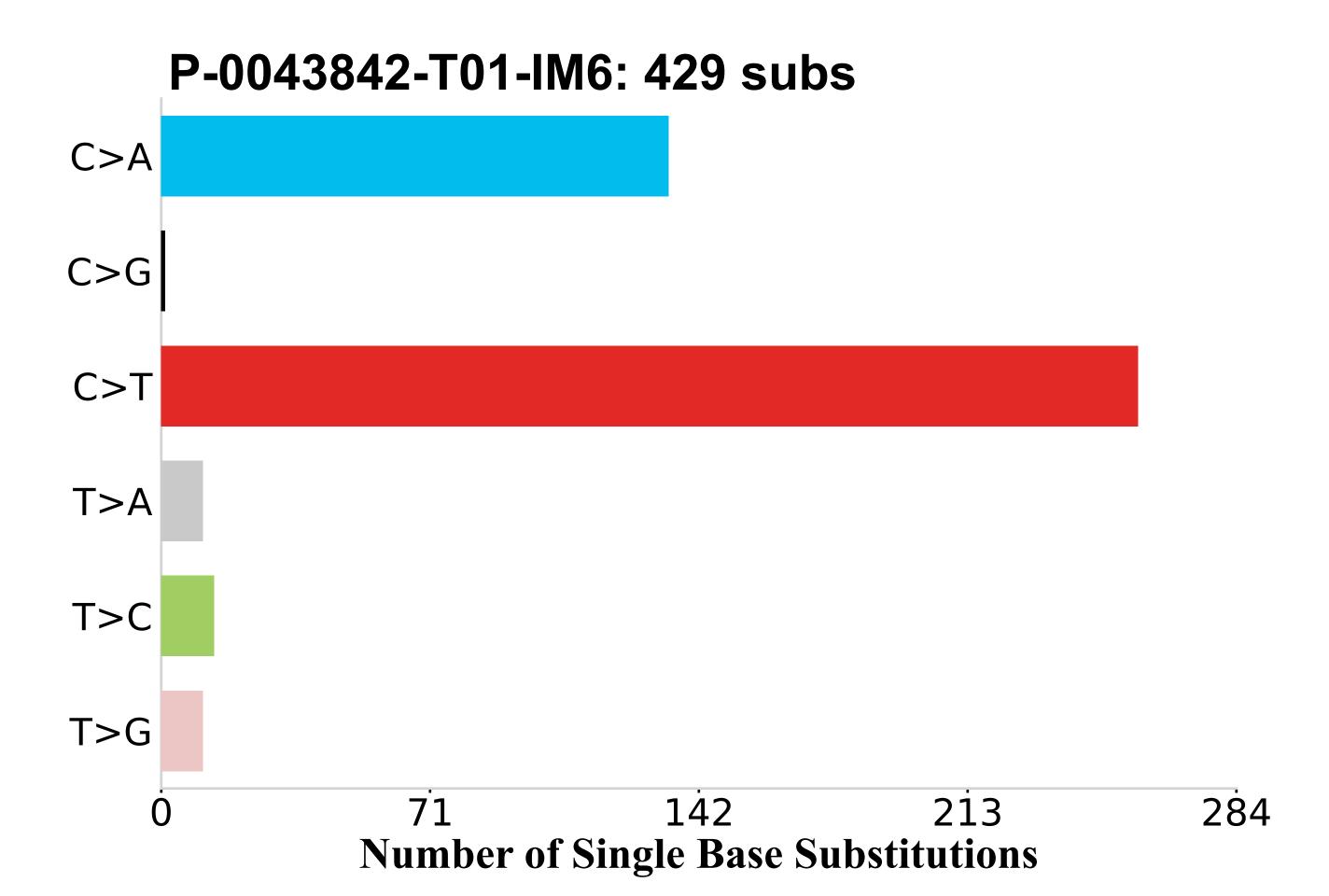


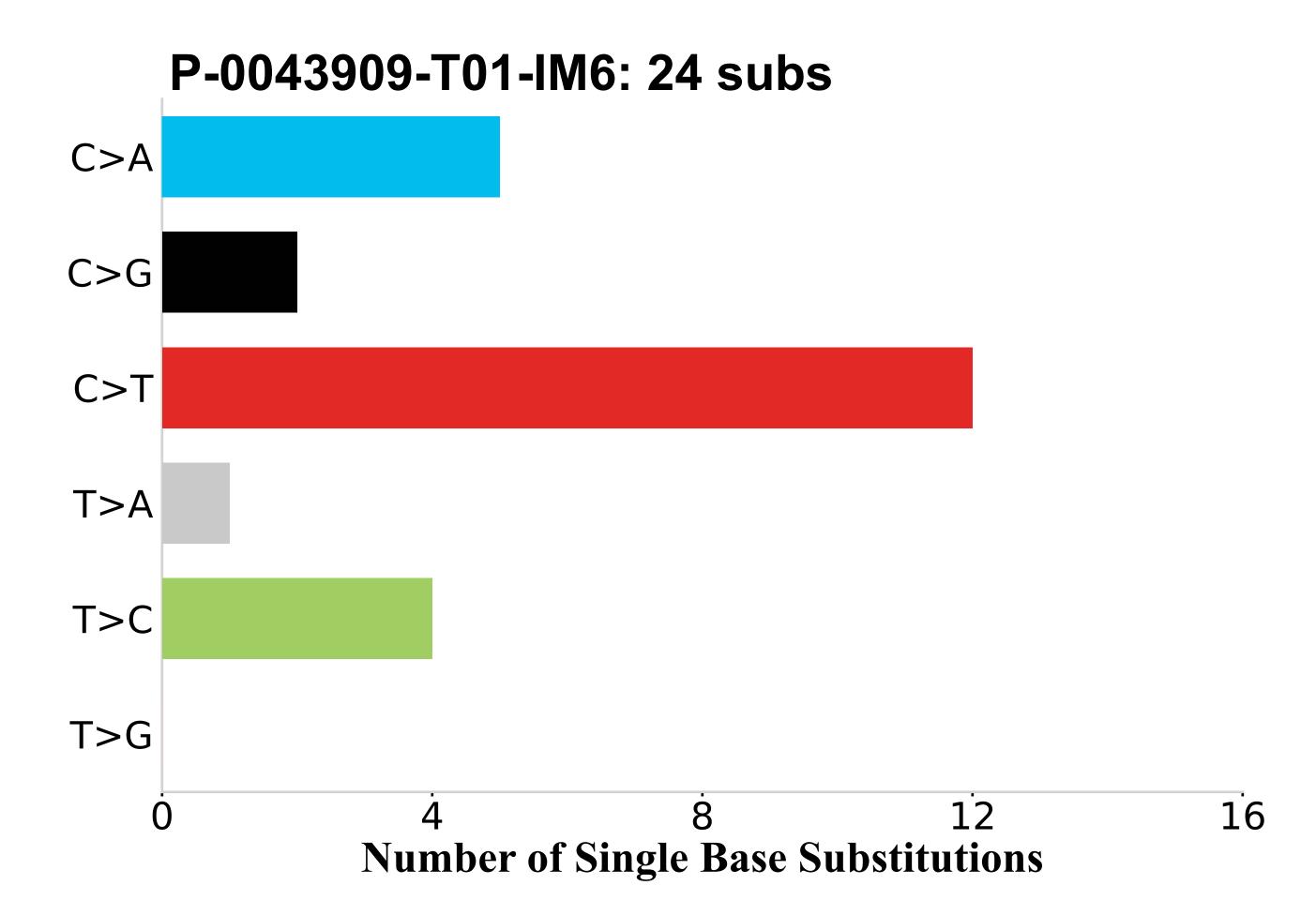


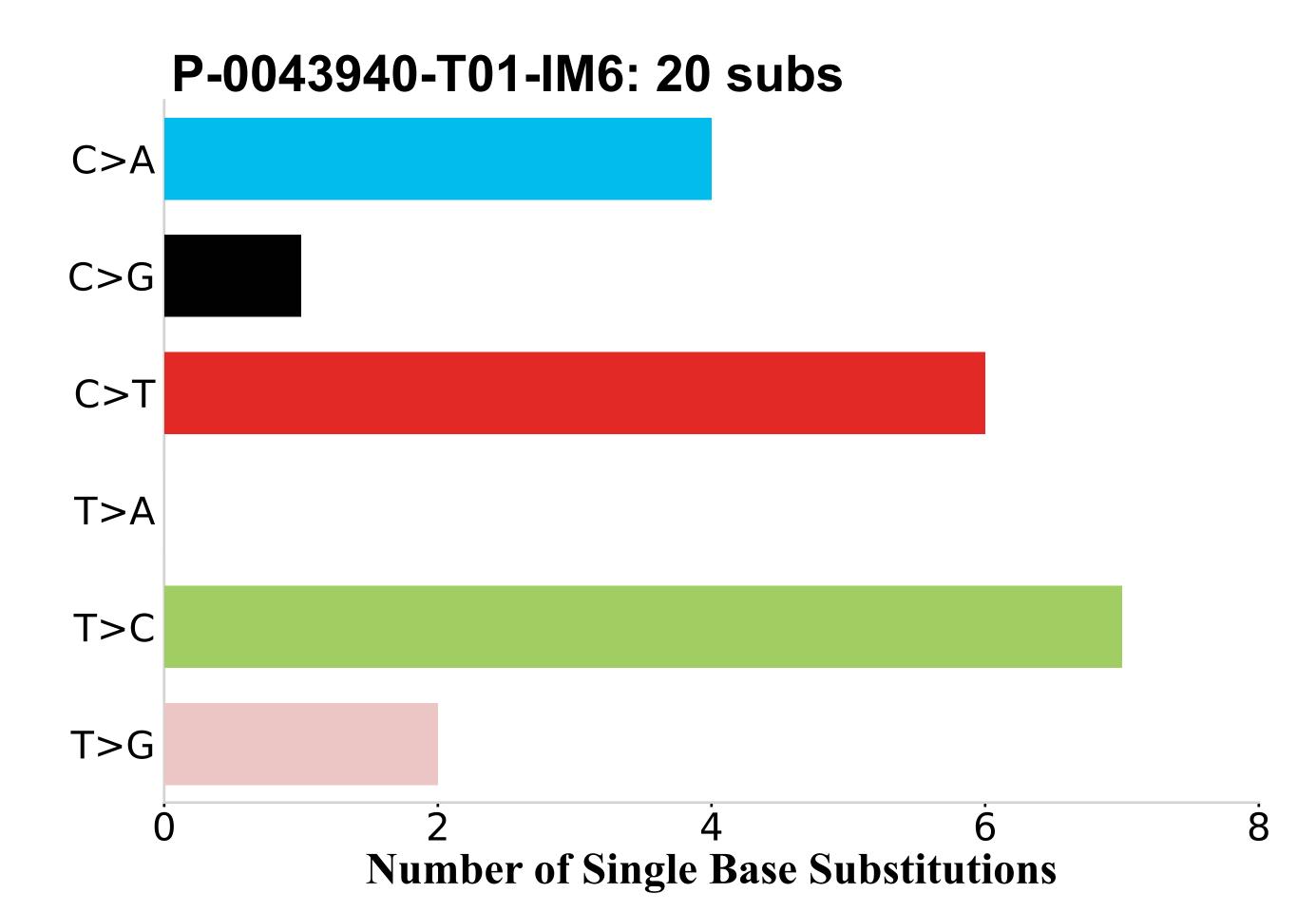


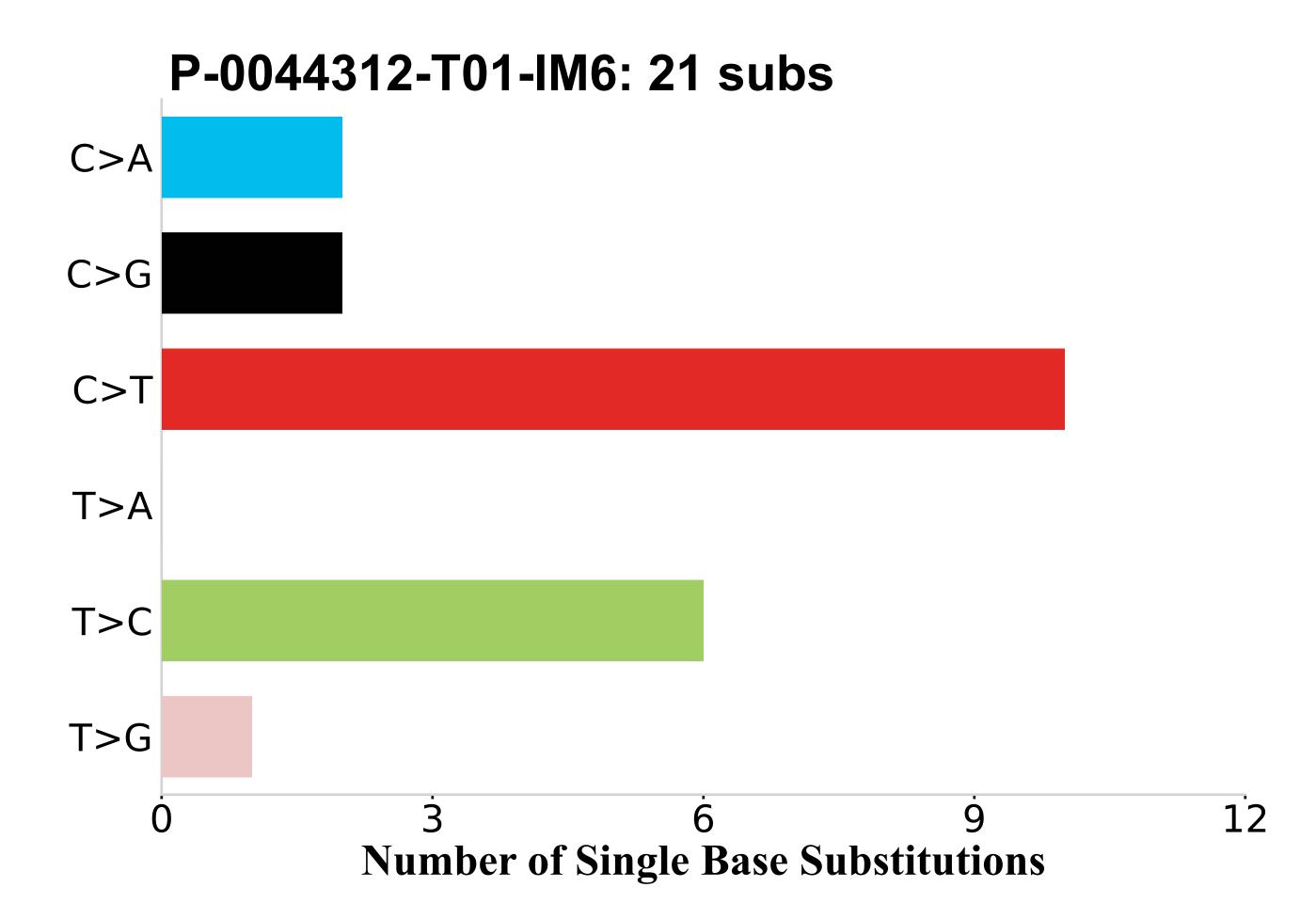


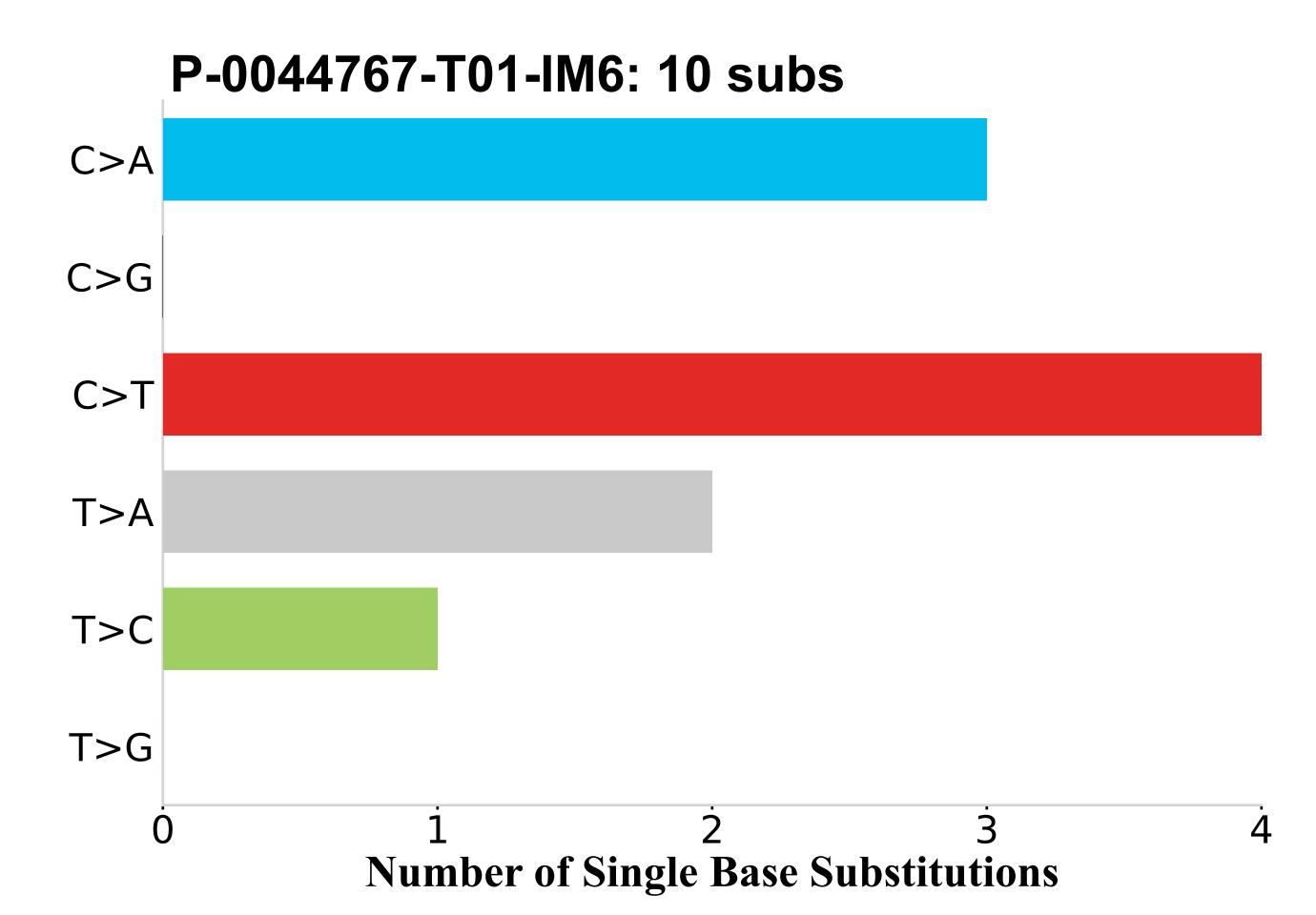


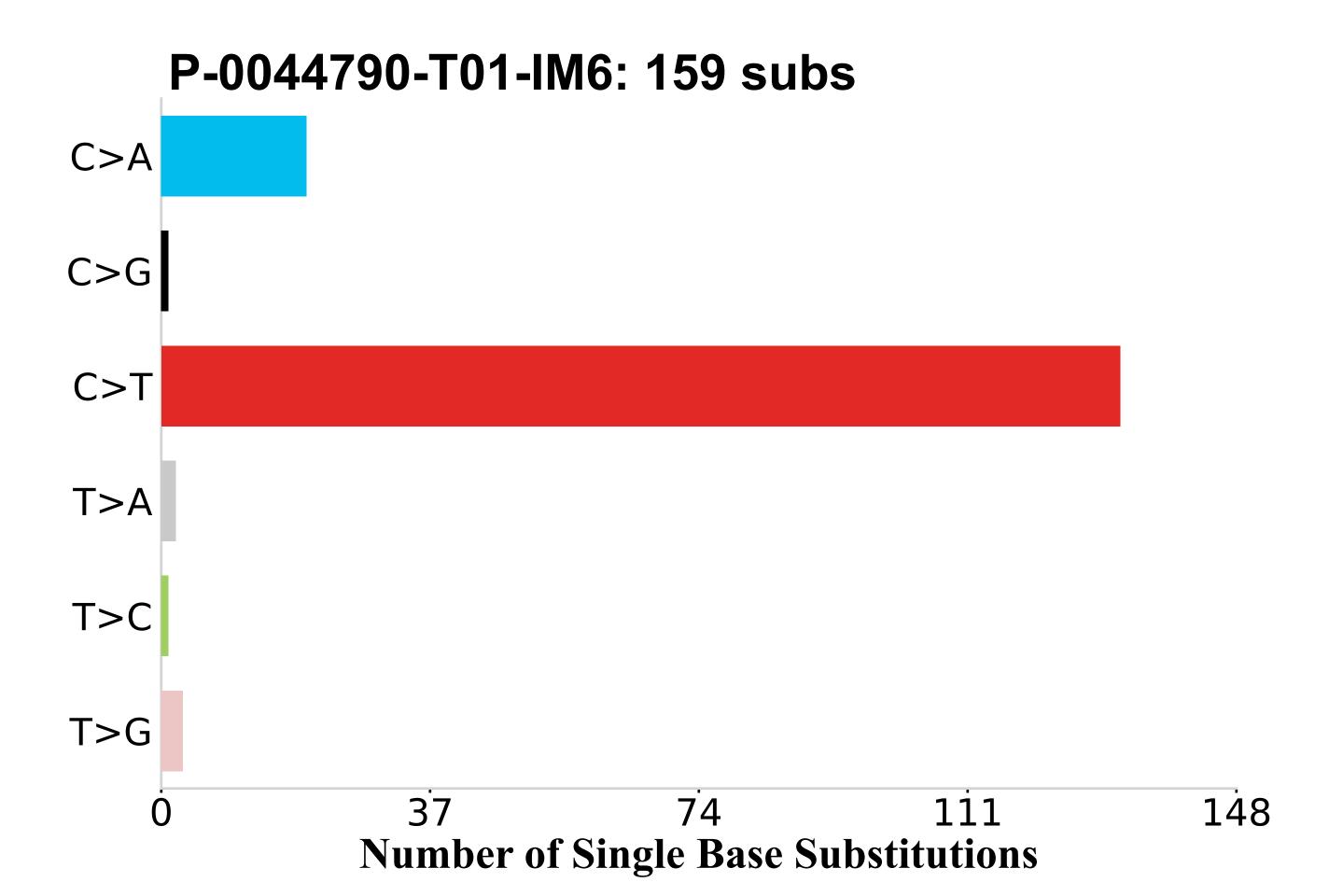


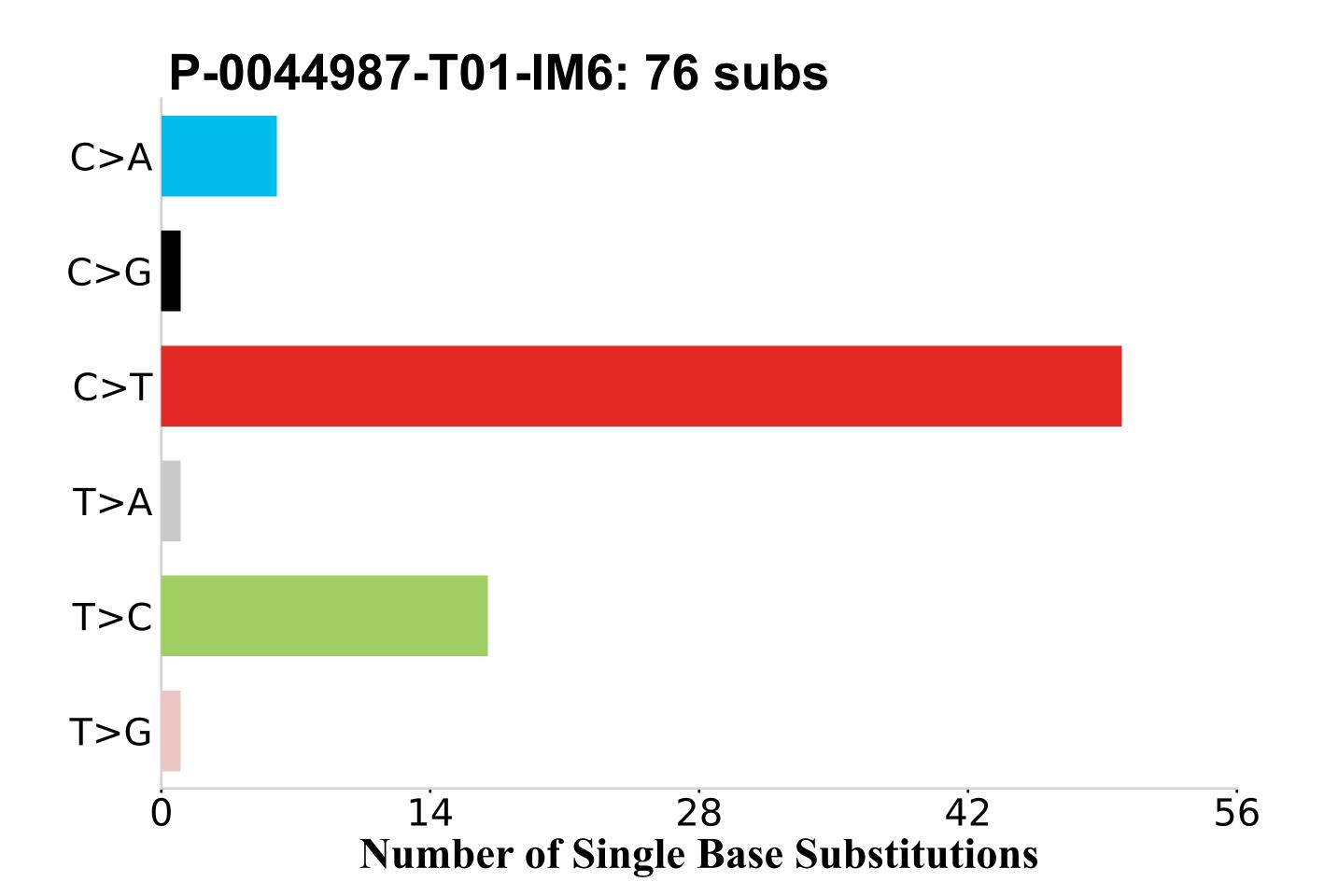


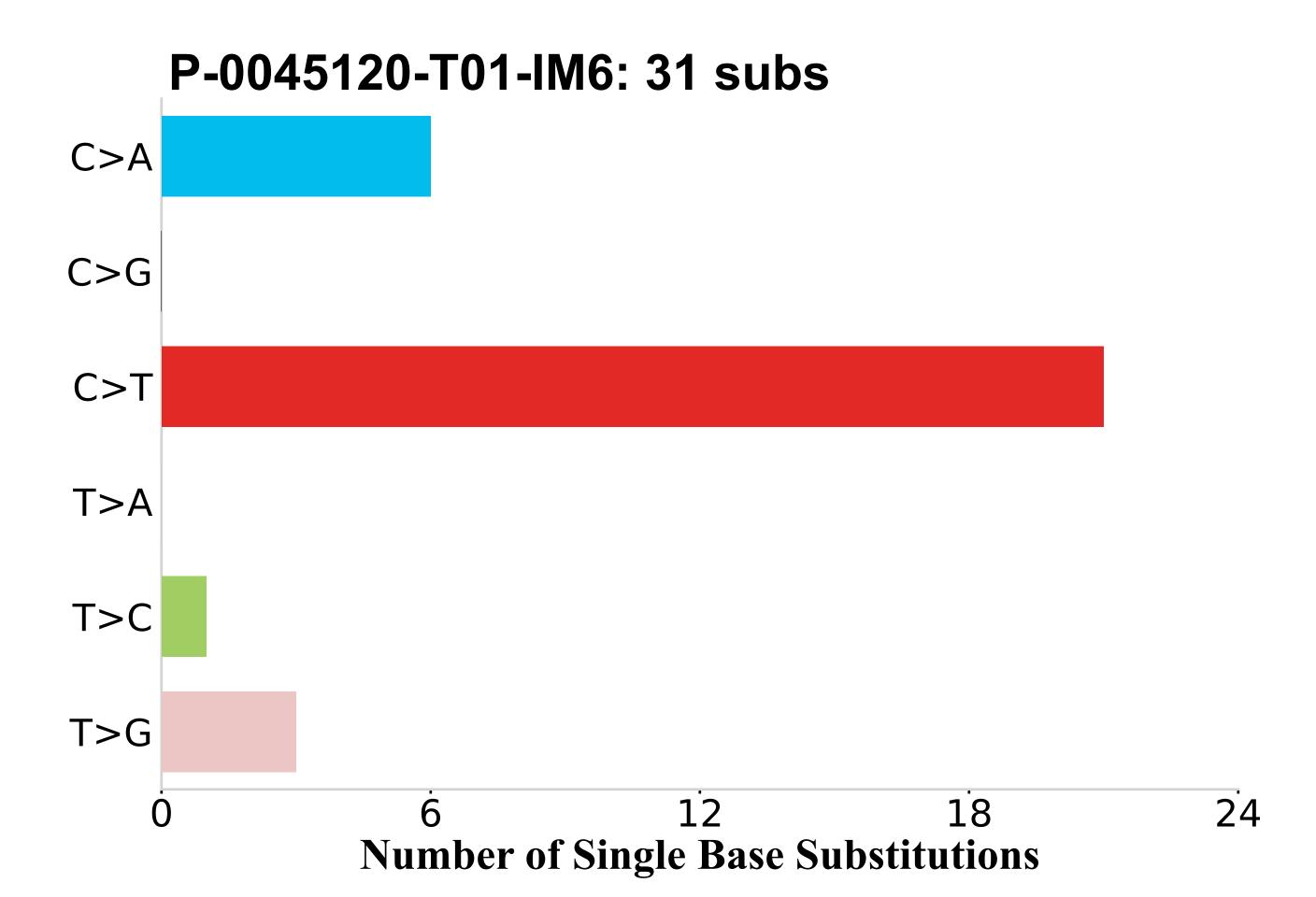


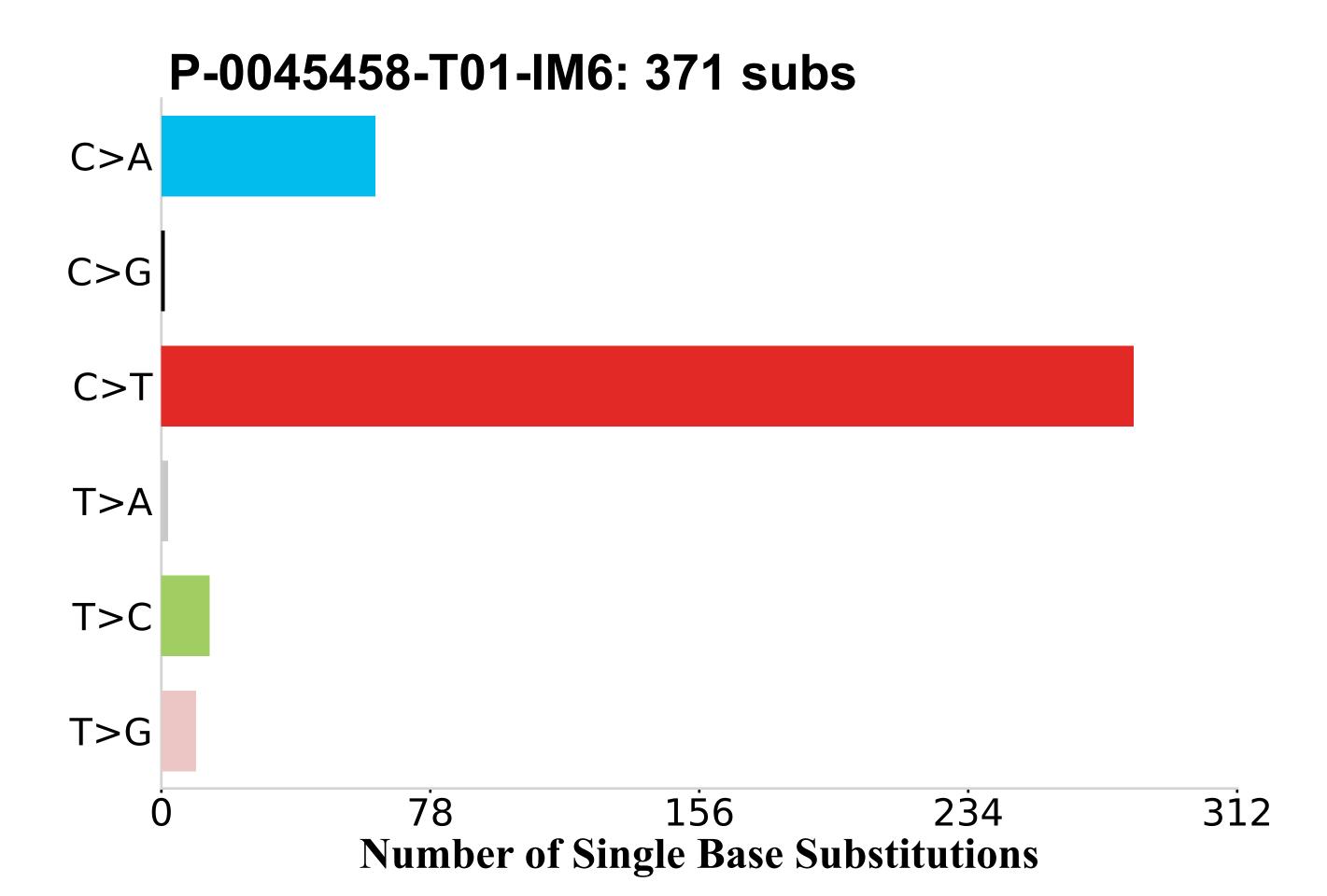


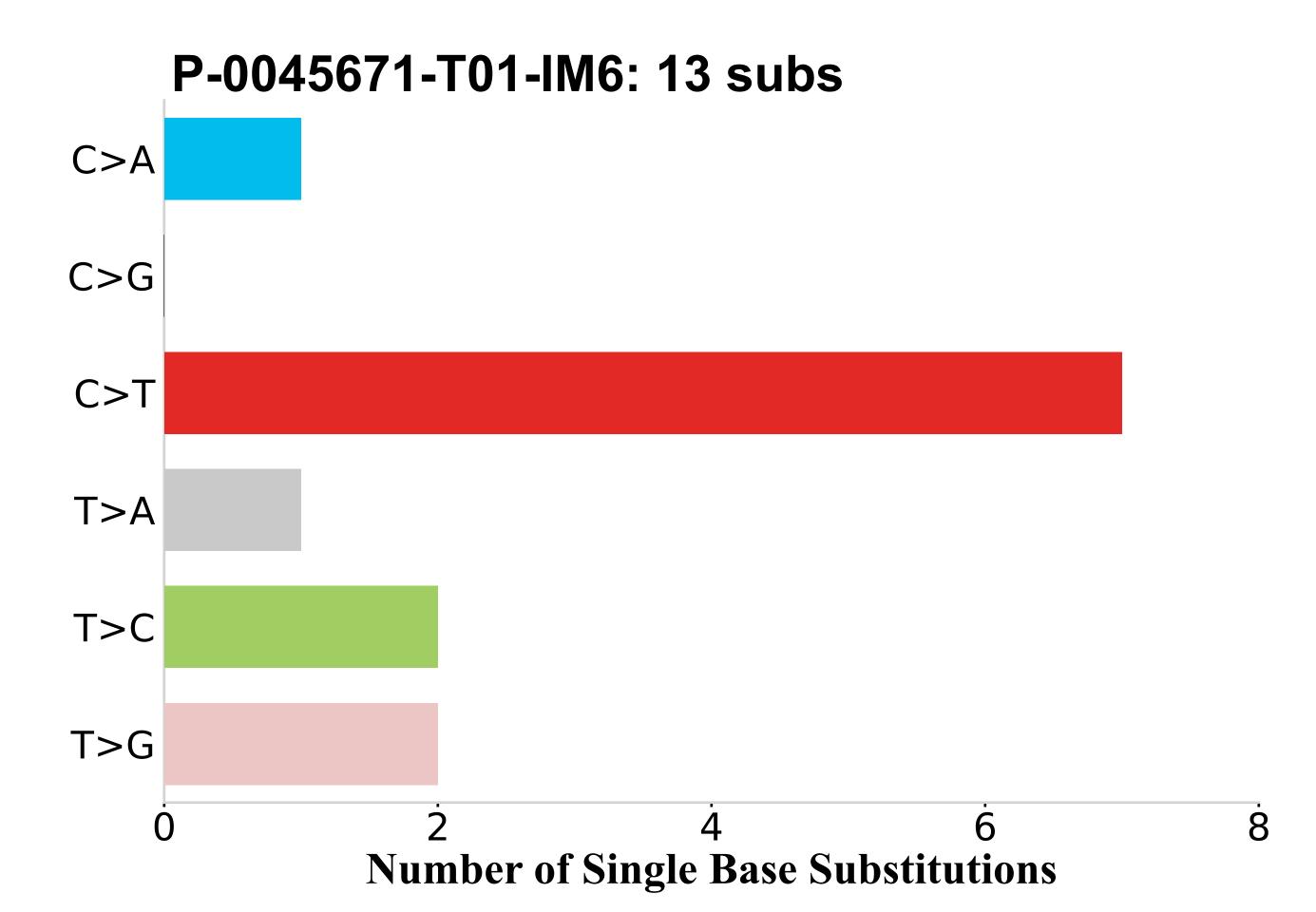


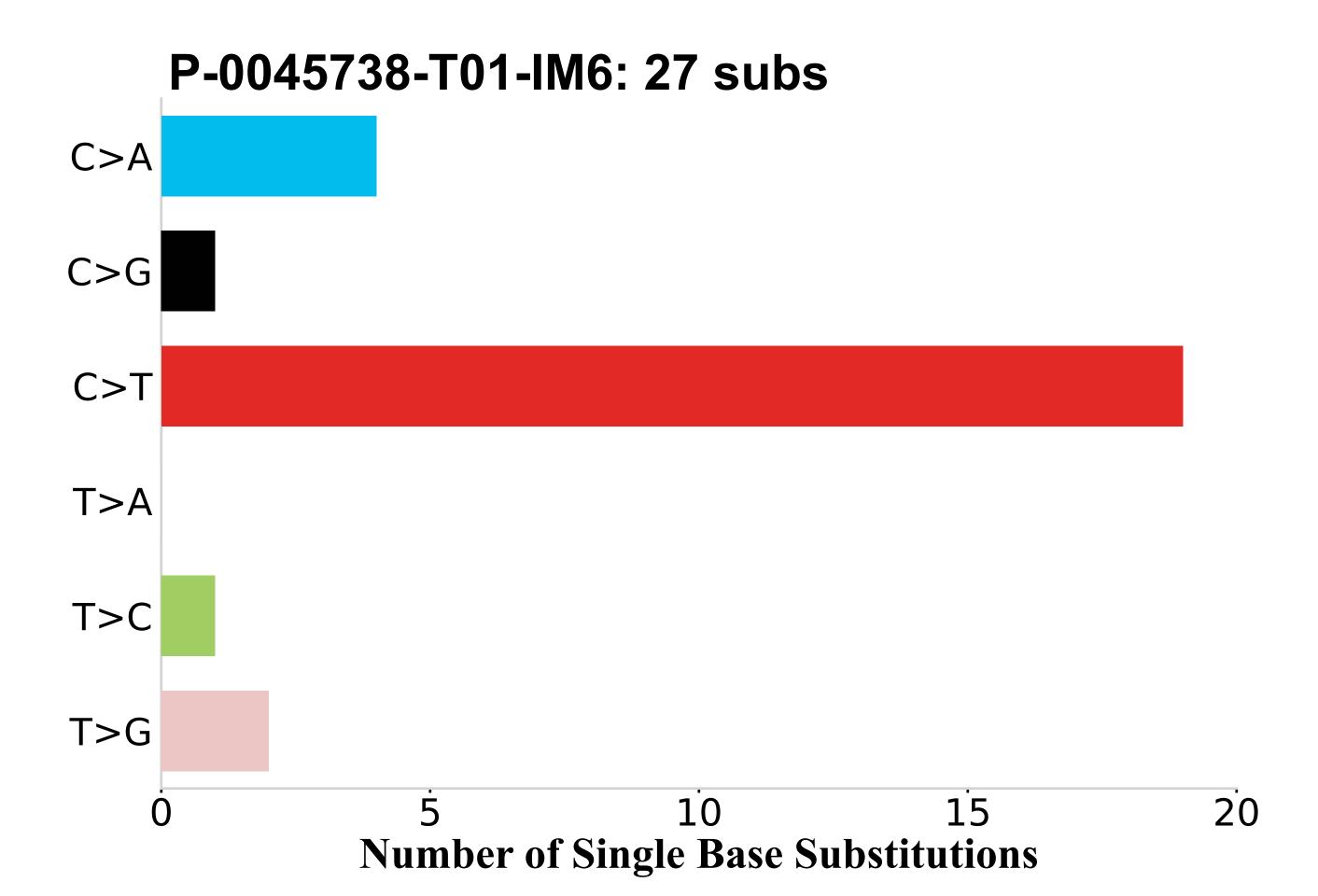


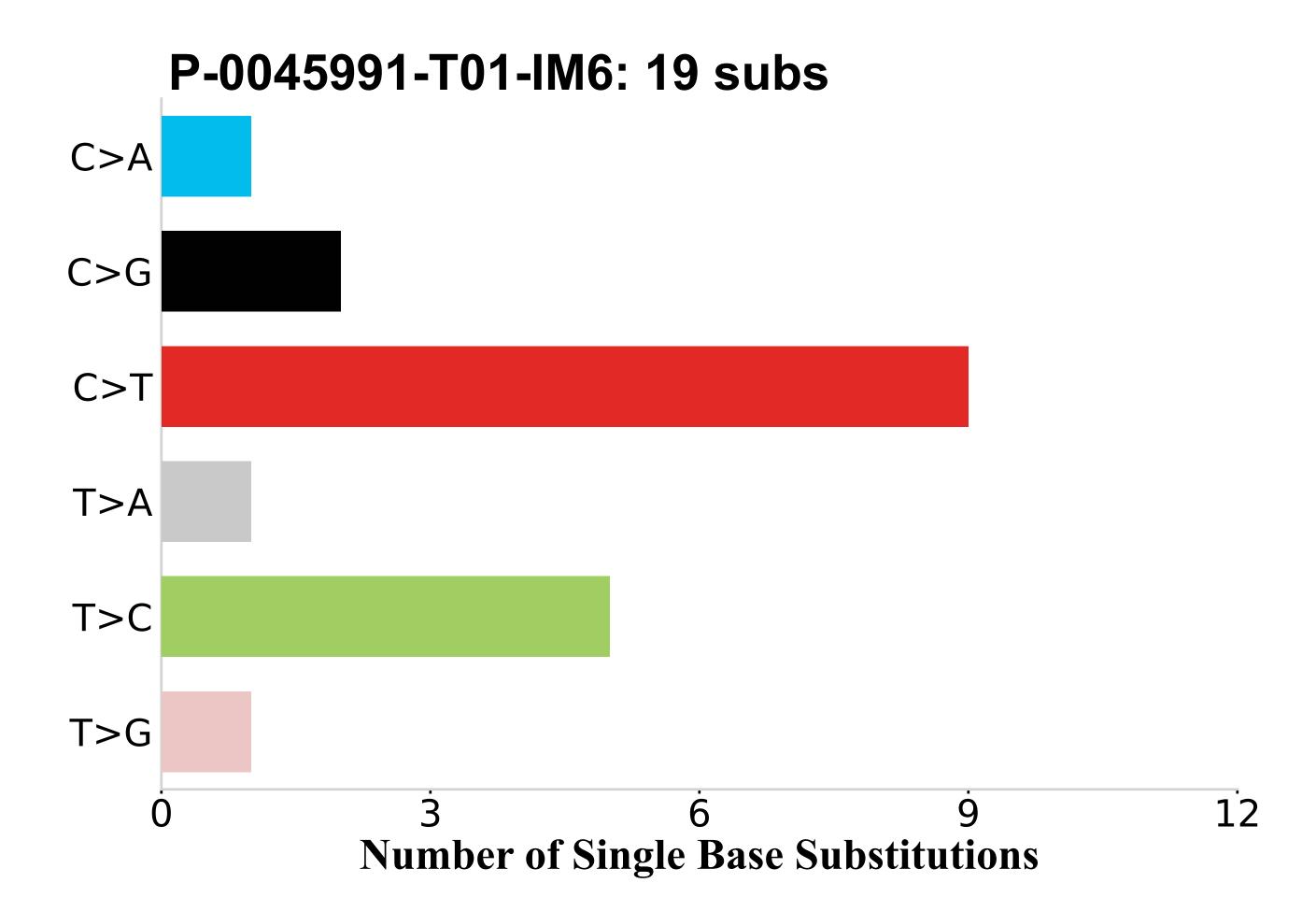


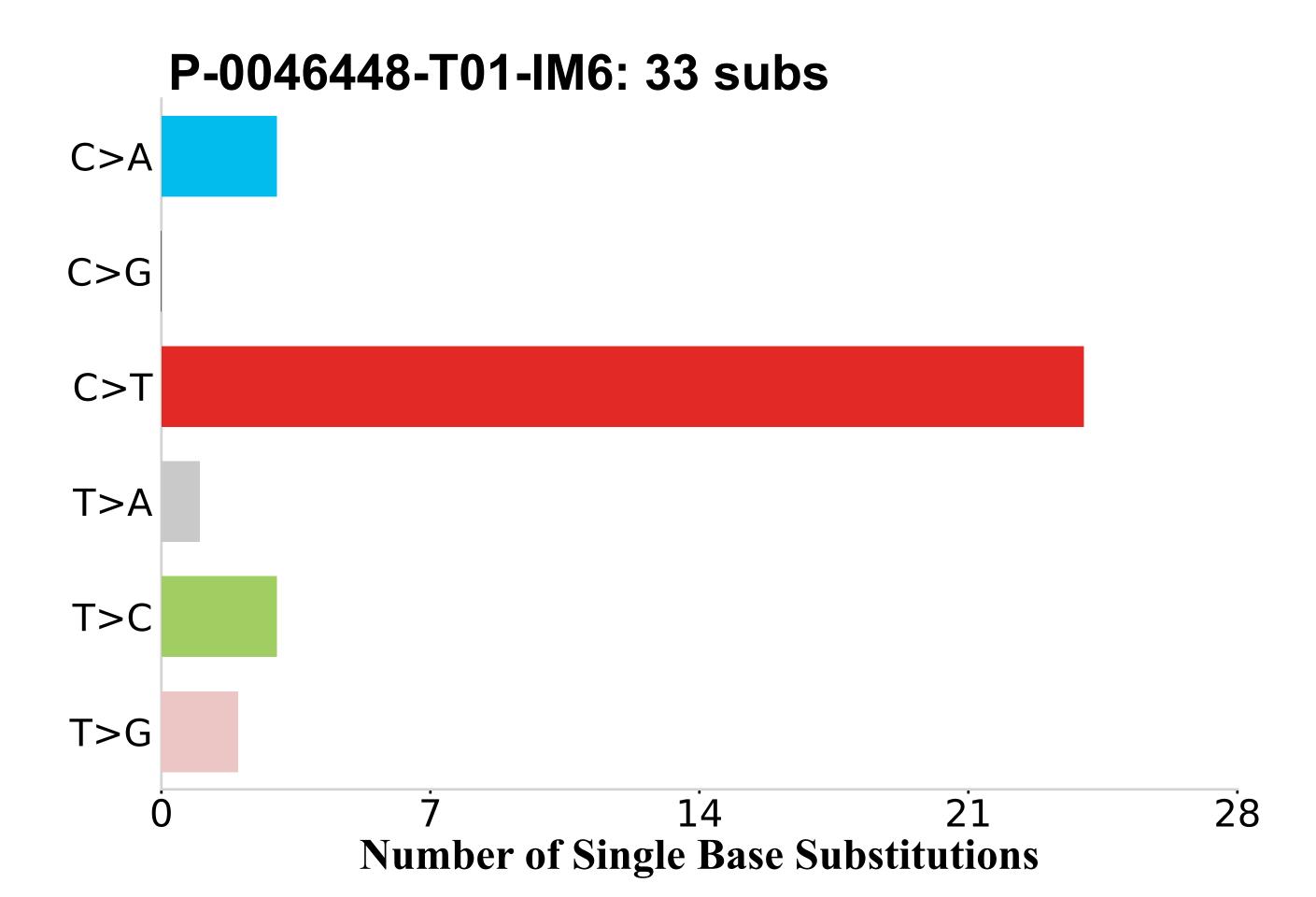


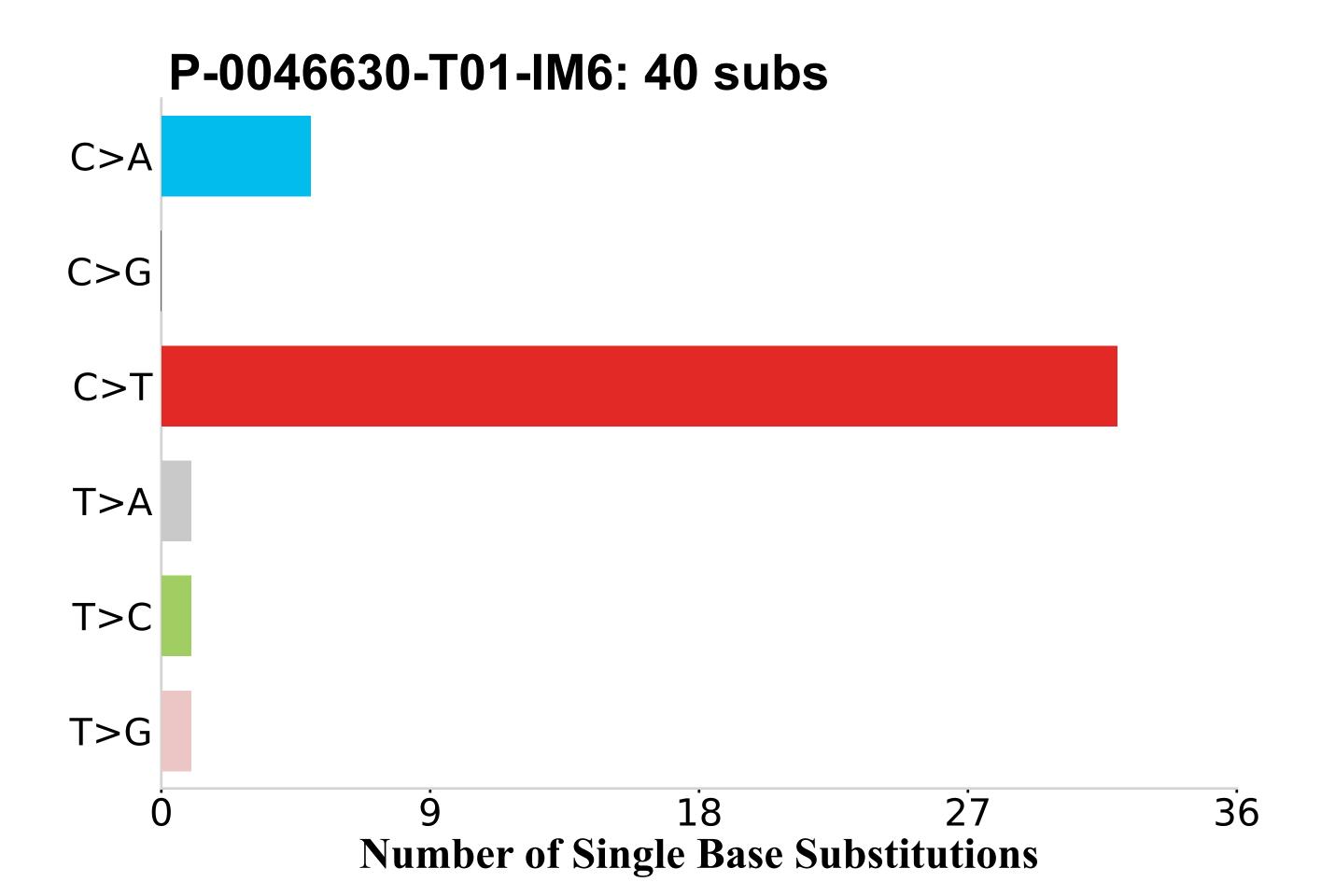


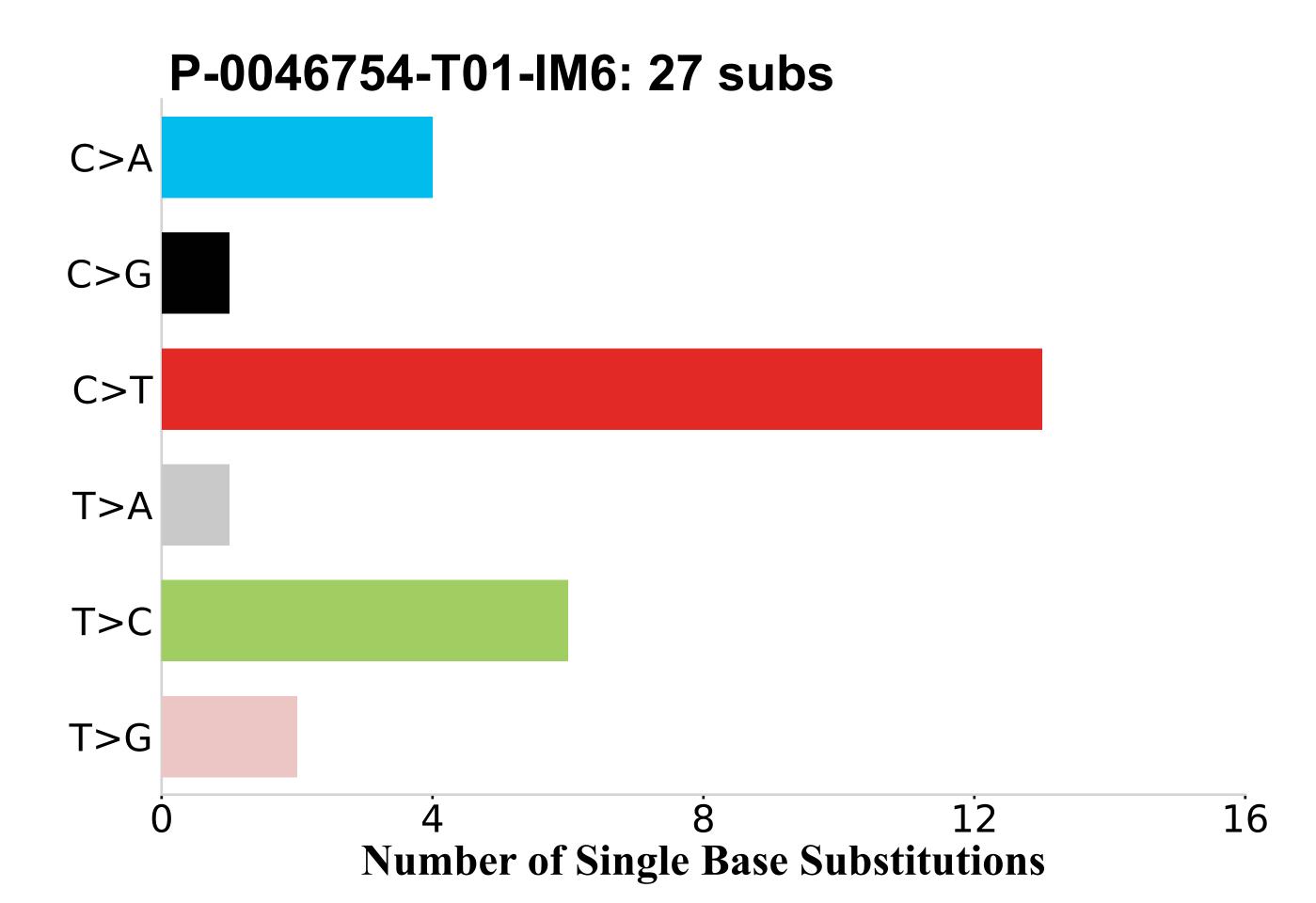












P-0046965-T01-IM6: 479 subs C>A C>G C>T T>A T>C T>G



