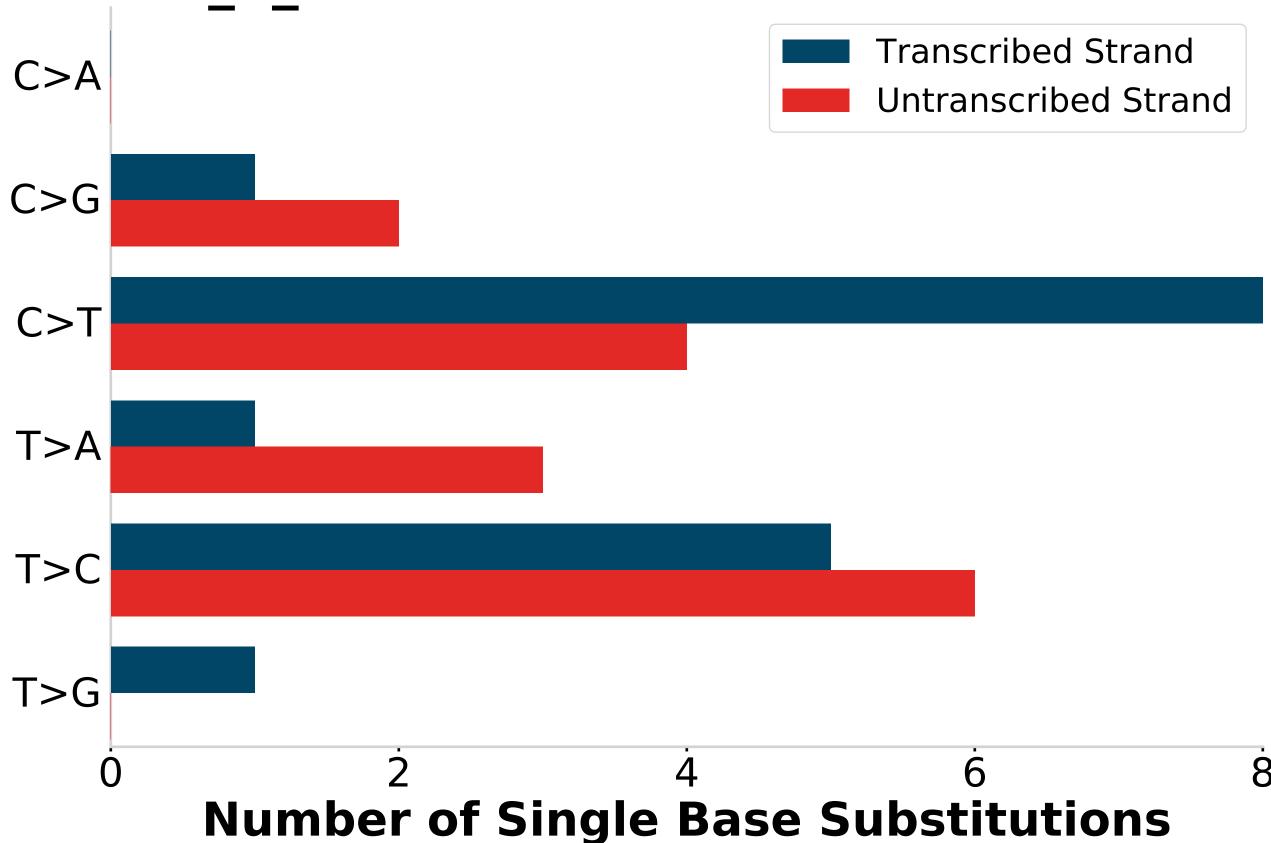
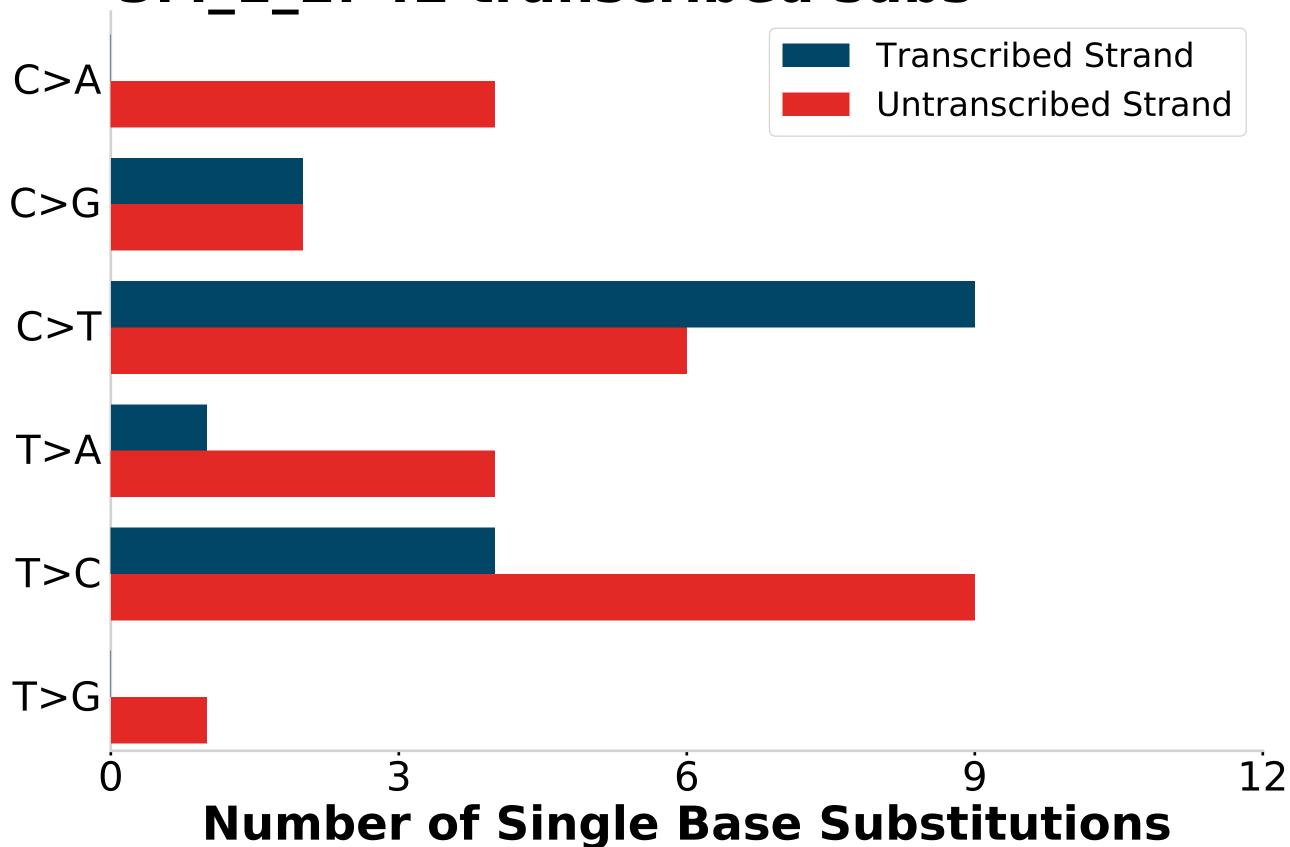
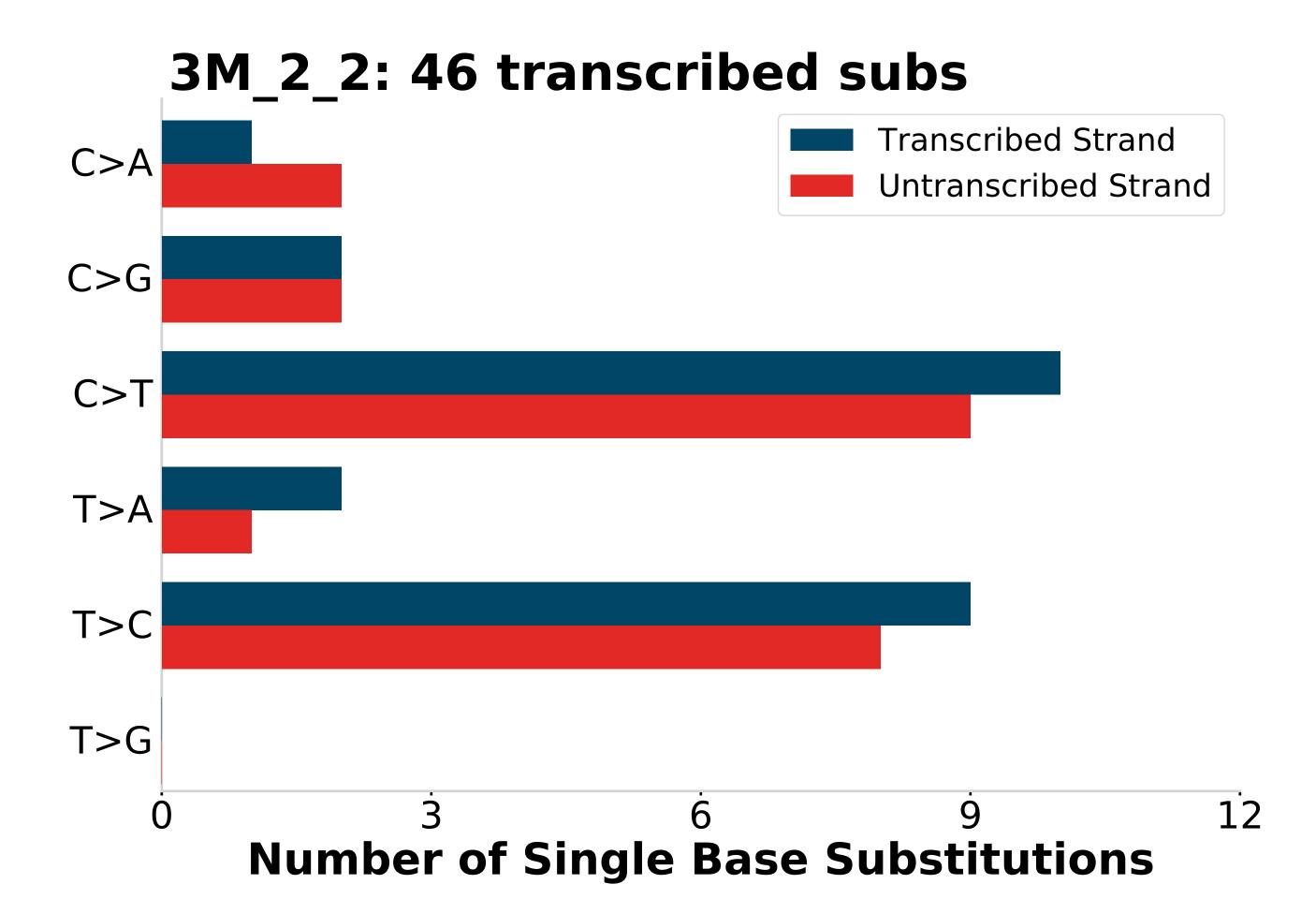
3M_1_1: 31 transcribed subs

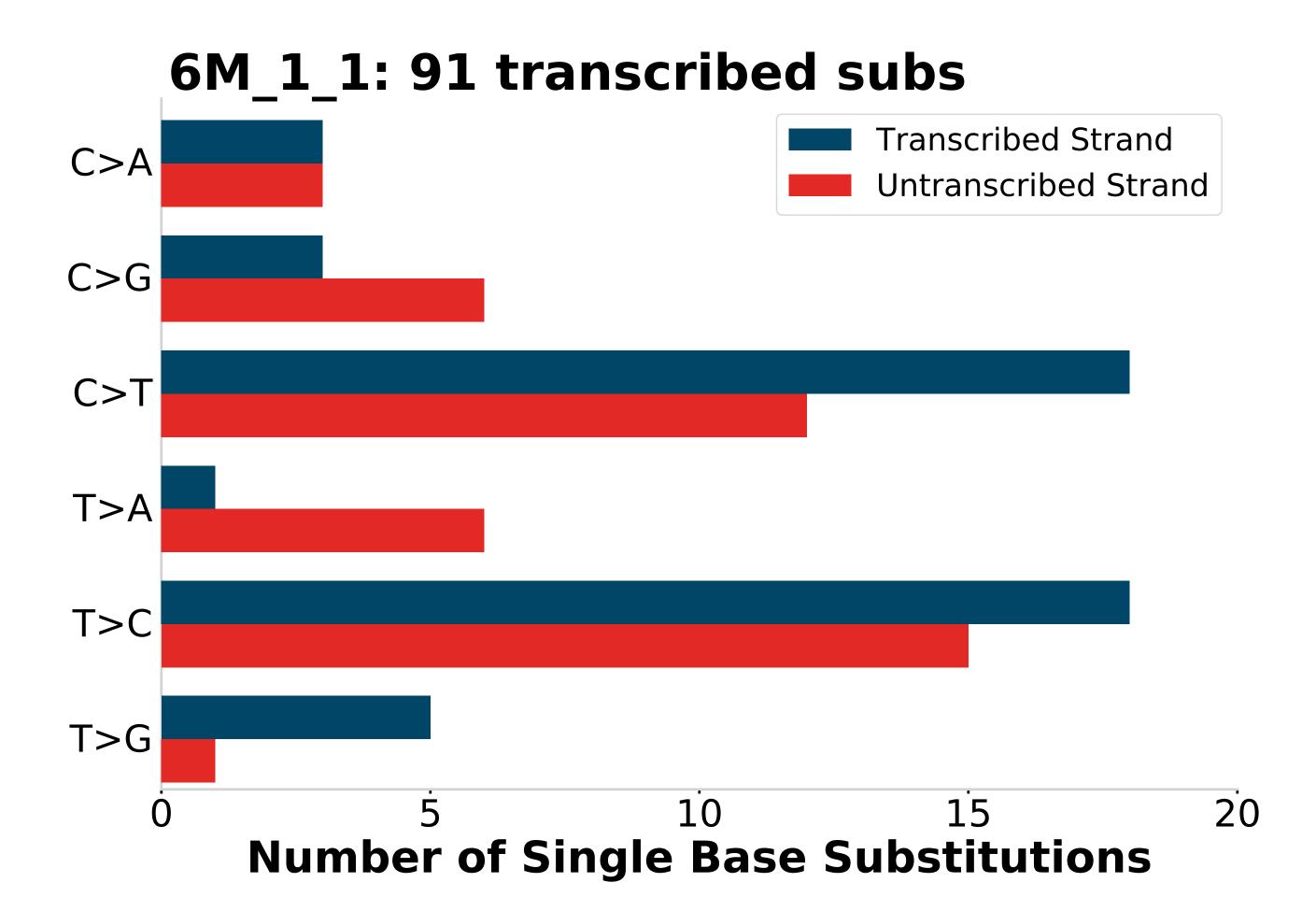


3M 1 2: 42 transcribed subs



3M 2 1: 45 transcribed subs **Transcribed Strand** C>A **Untranscribed Strand** C>G C>T T>A T>C T>G 12 9 **Number of Single Base Substitutions**





6M 1 2: 35 transcribed subs **Transcribed Strand** C>A **Untranscribed Strand** C>G C>T T>A T>C T>G **Number of Single Base Substitutions**

6M 2 1: 58 transcribed subs **Transcribed Strand** C>A **Untranscribed Strand** C>G C>T T>A T>C T>G 12 16 **Number of Single Base Substitutions**

6M_2_2: 46 transcribed subs

