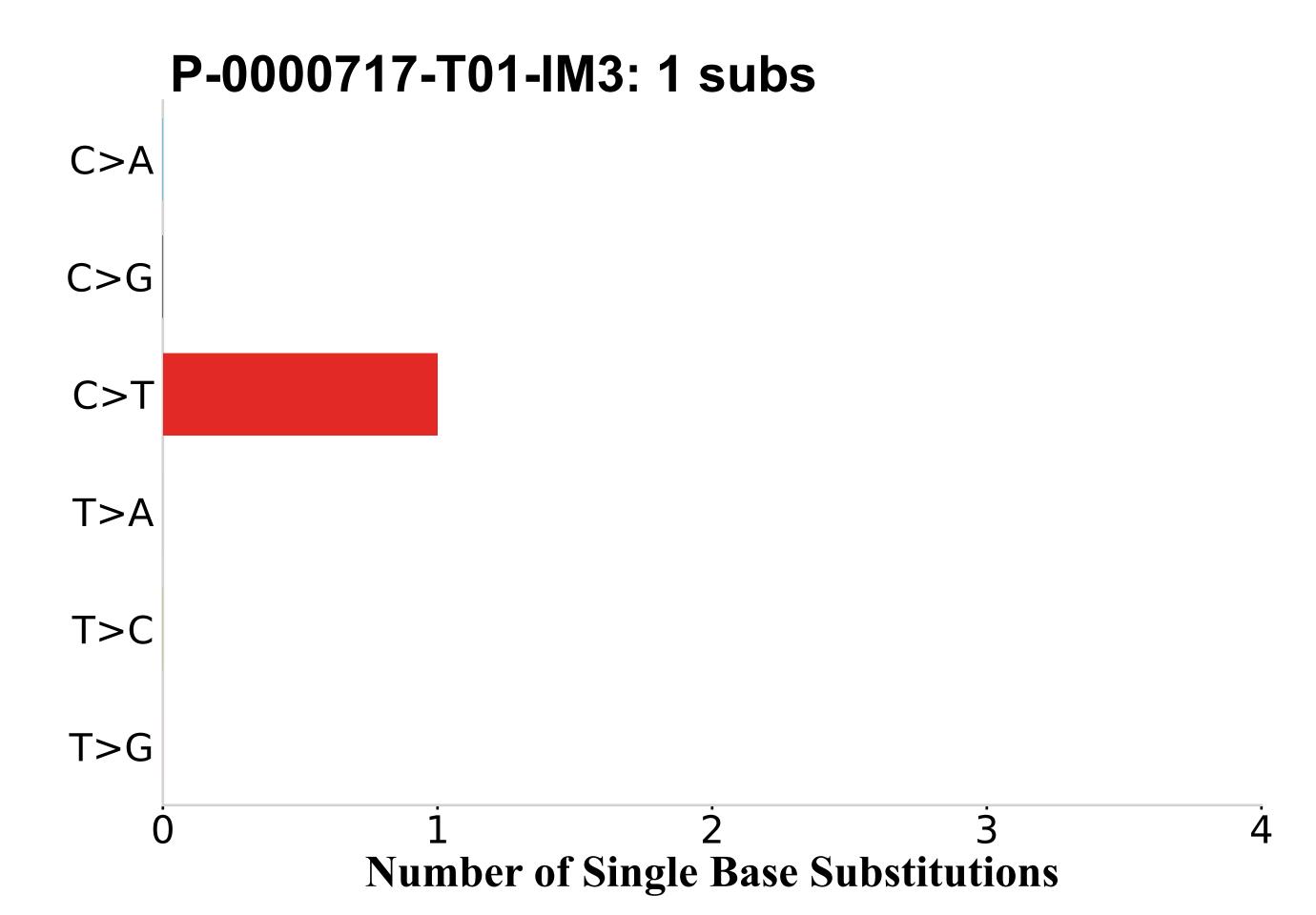
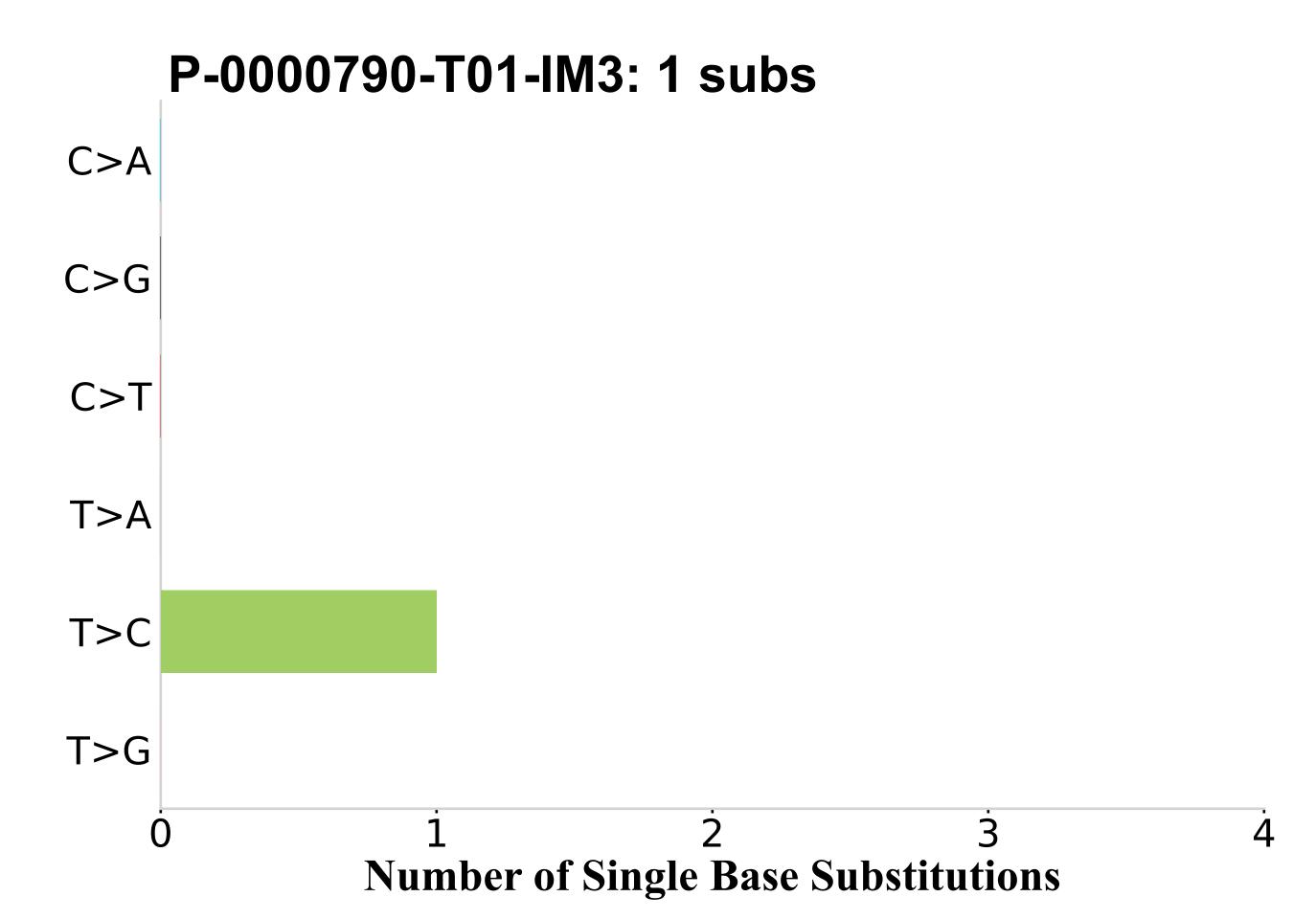


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Number of Single Base Substitutions



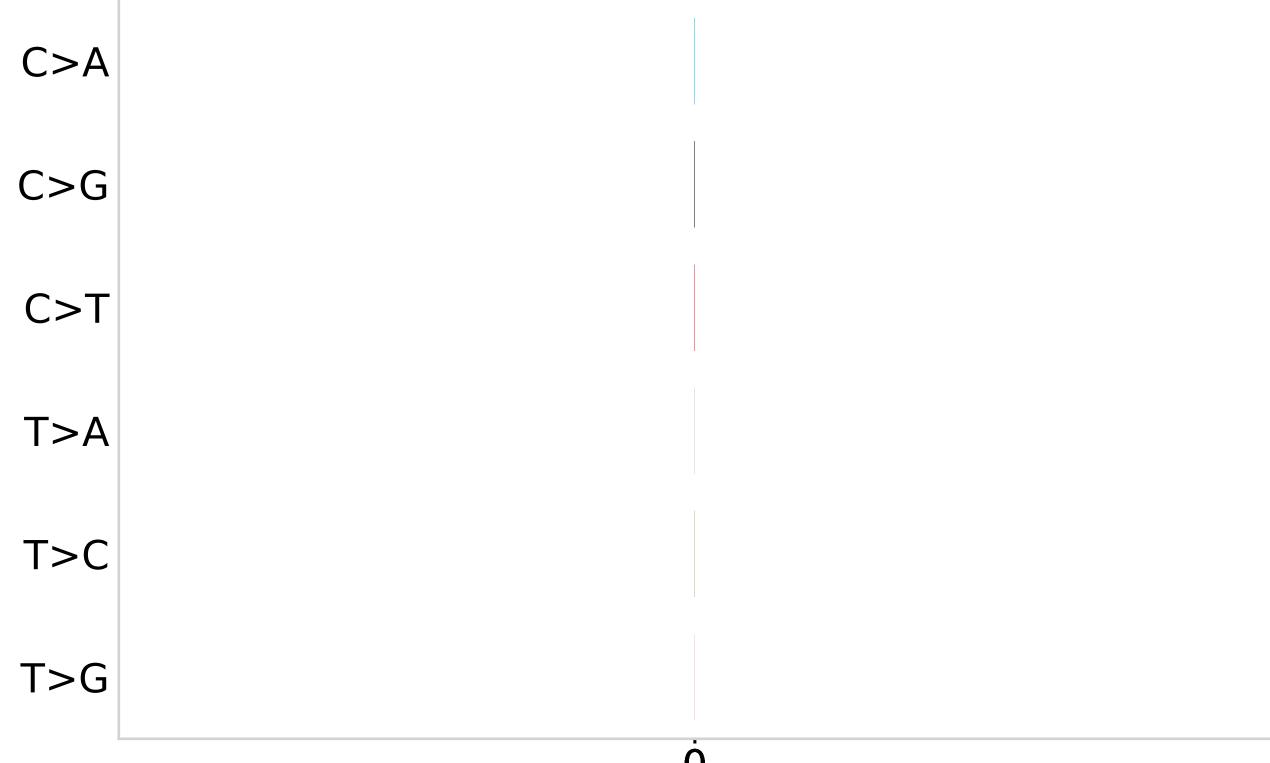


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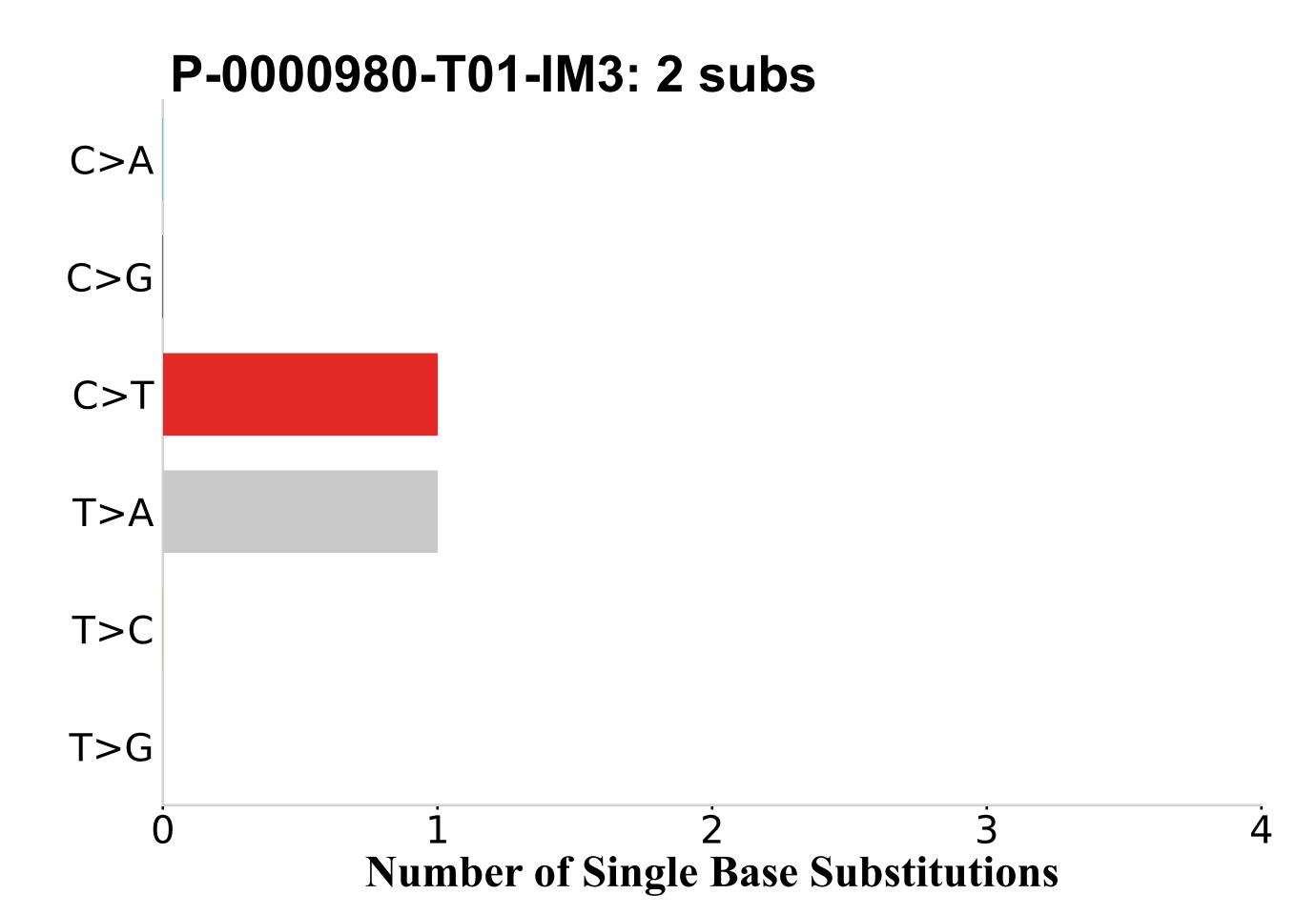


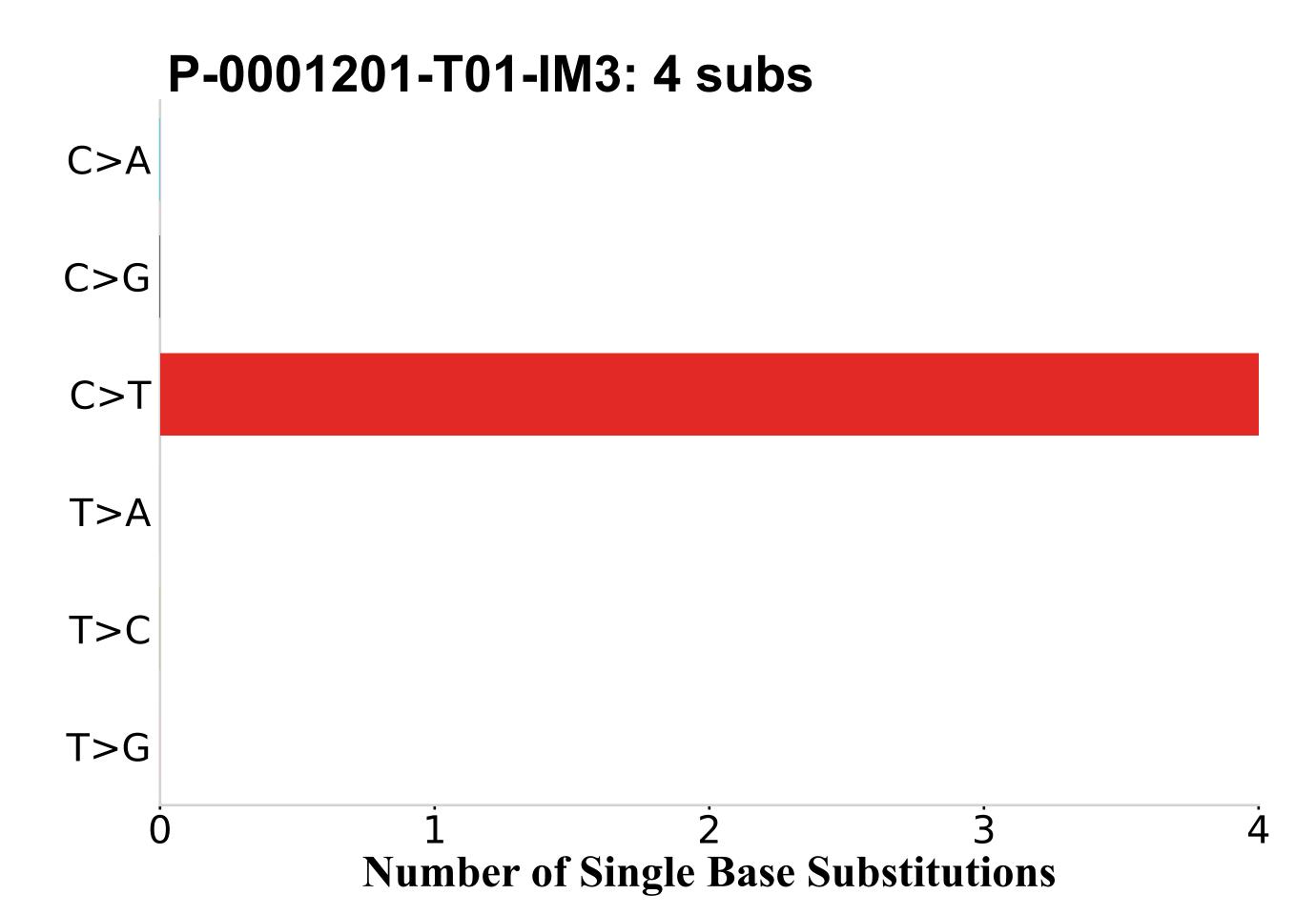
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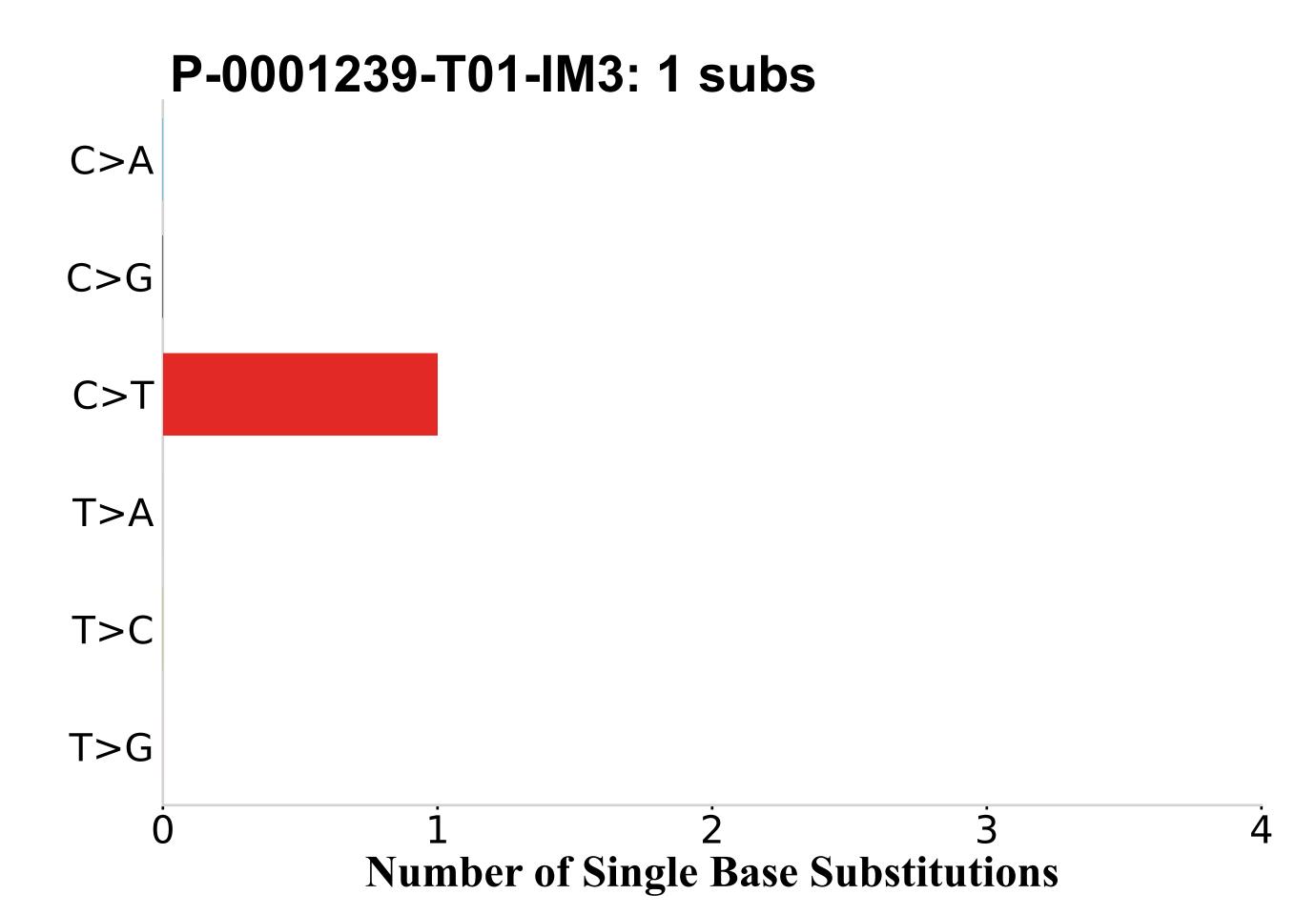
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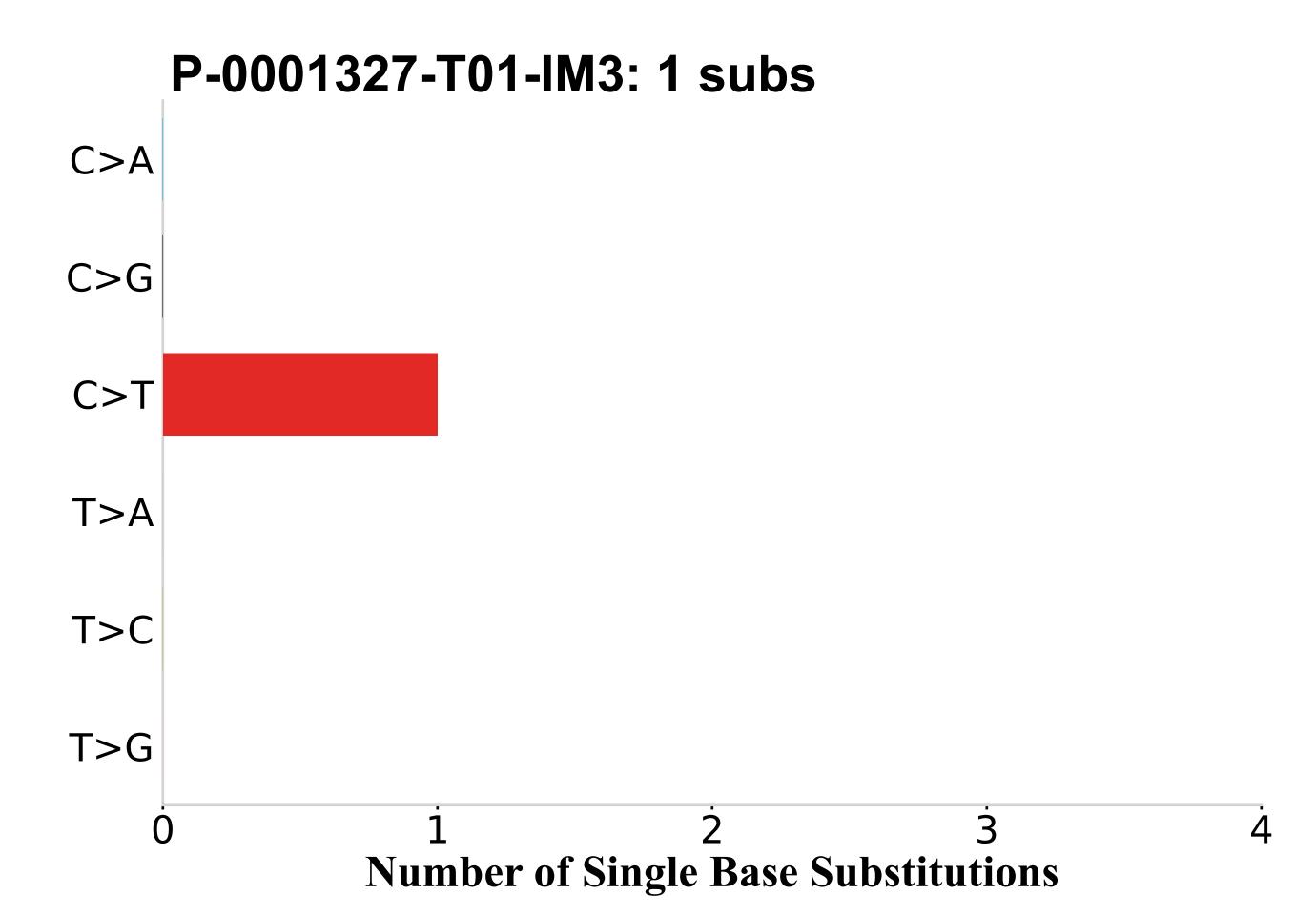


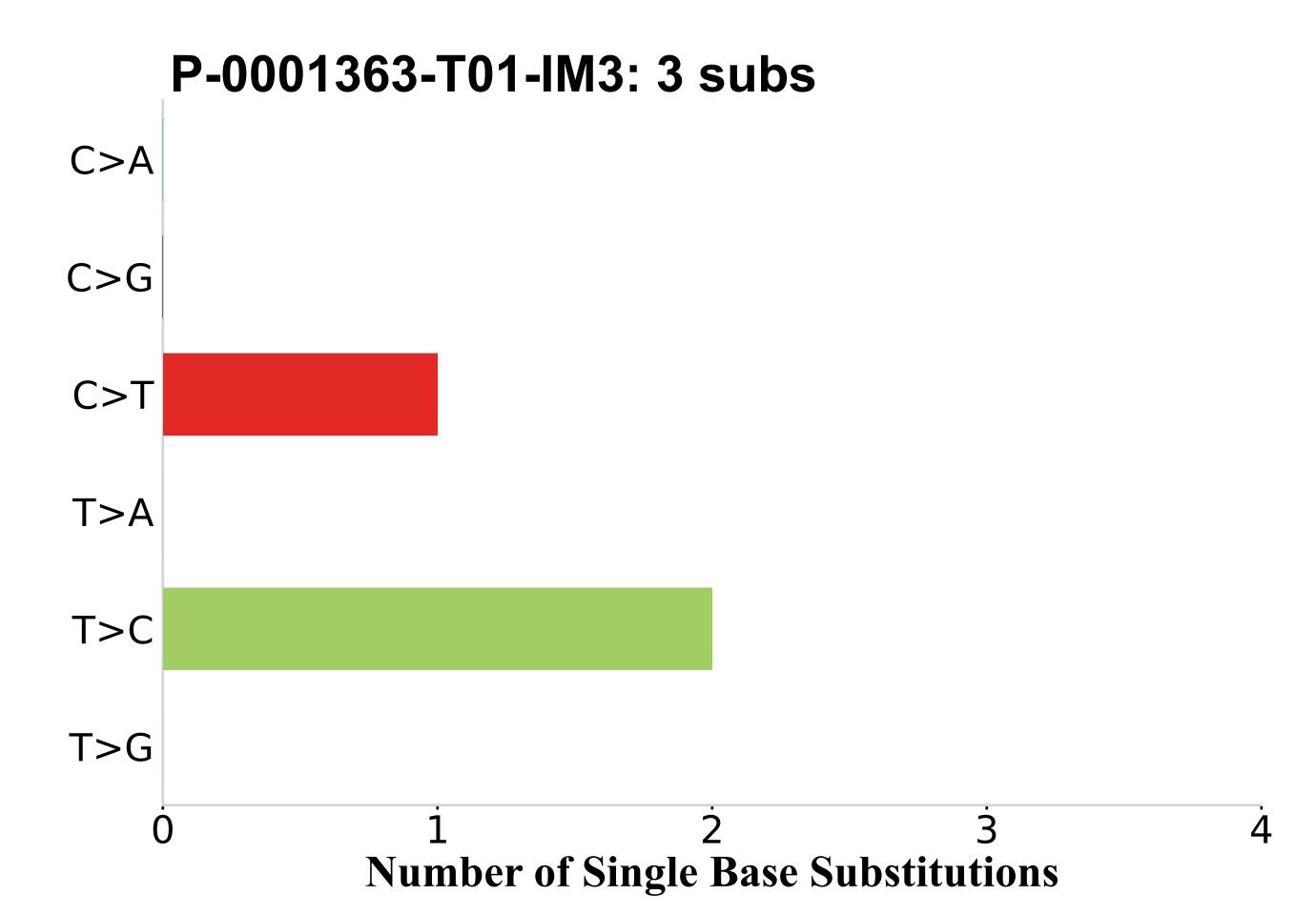
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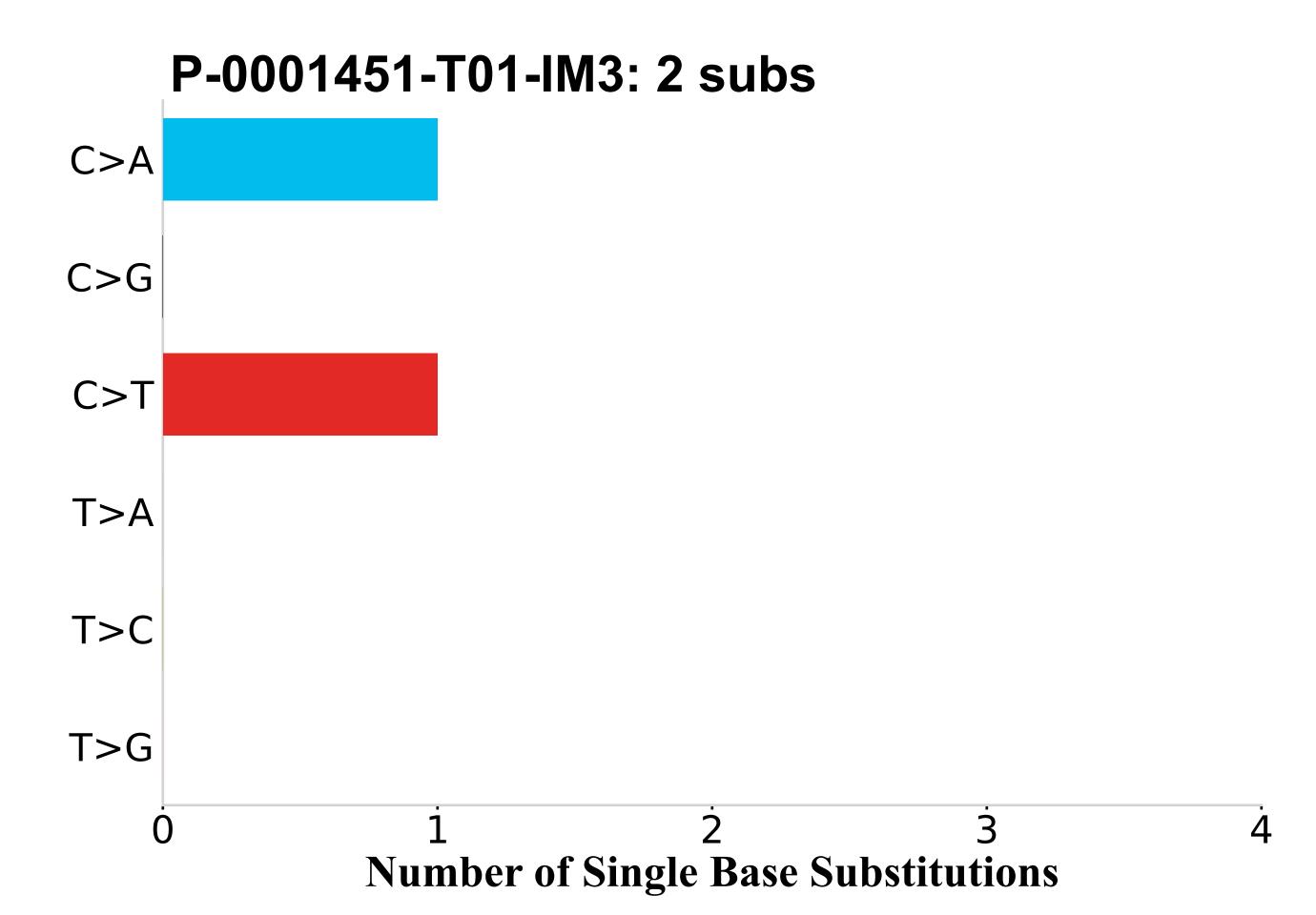








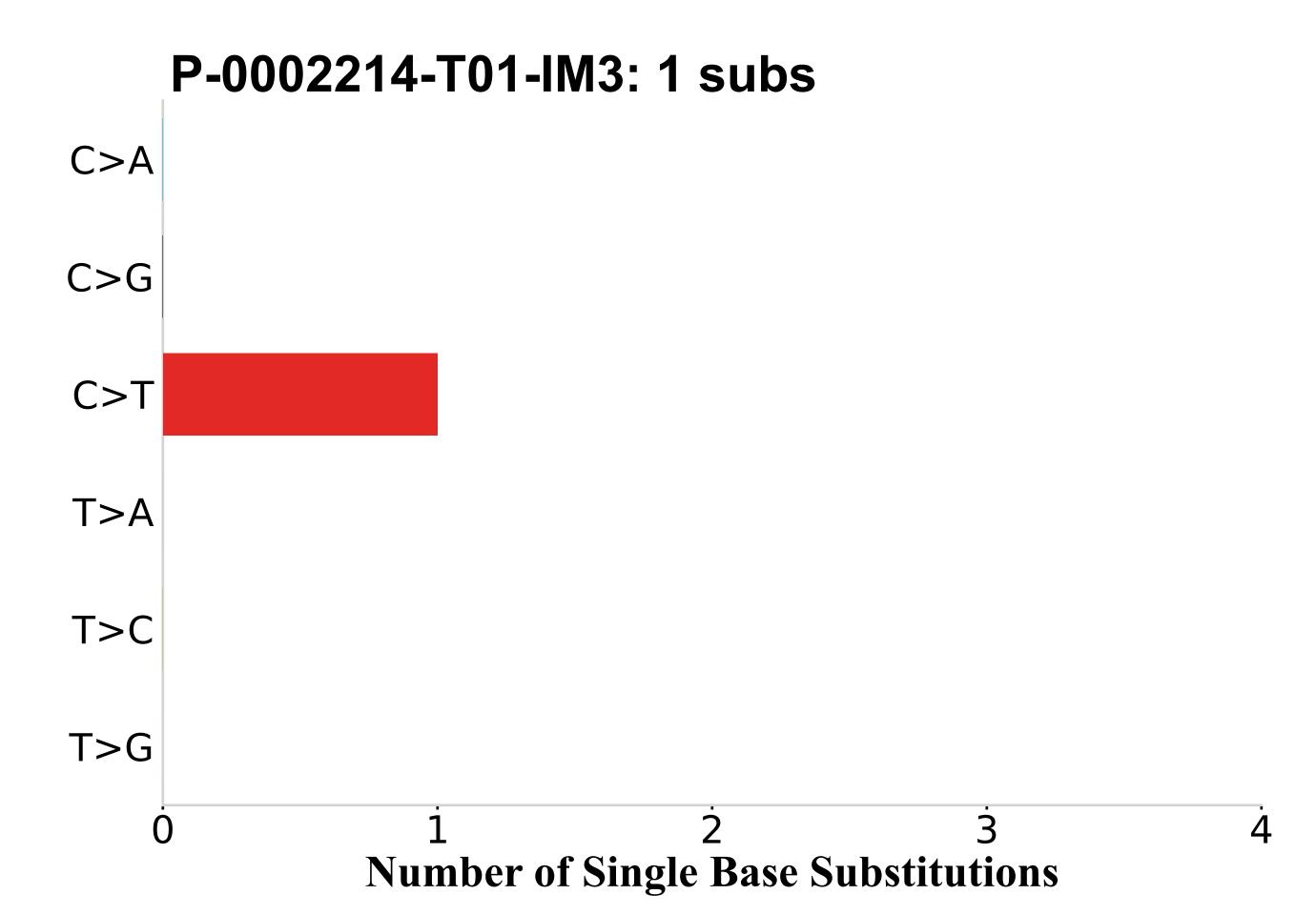




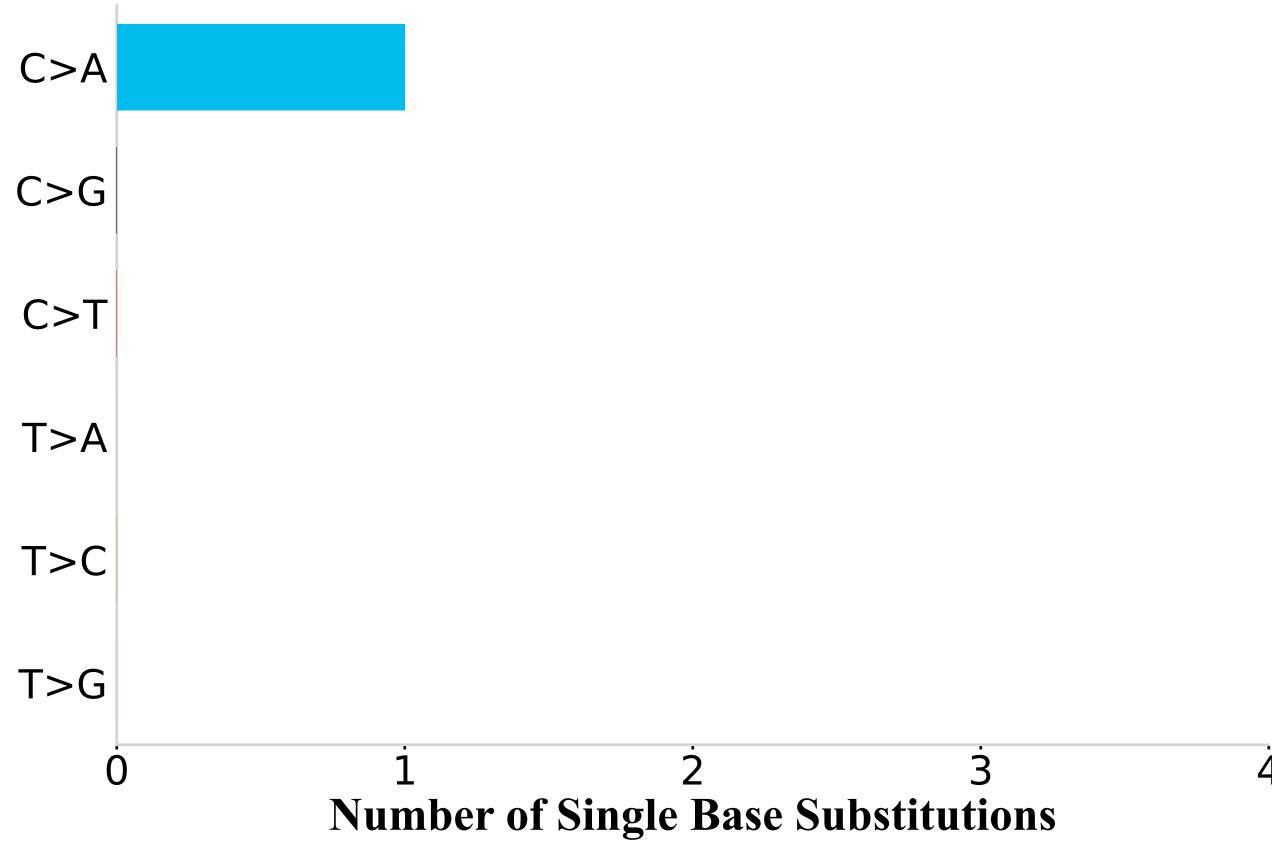
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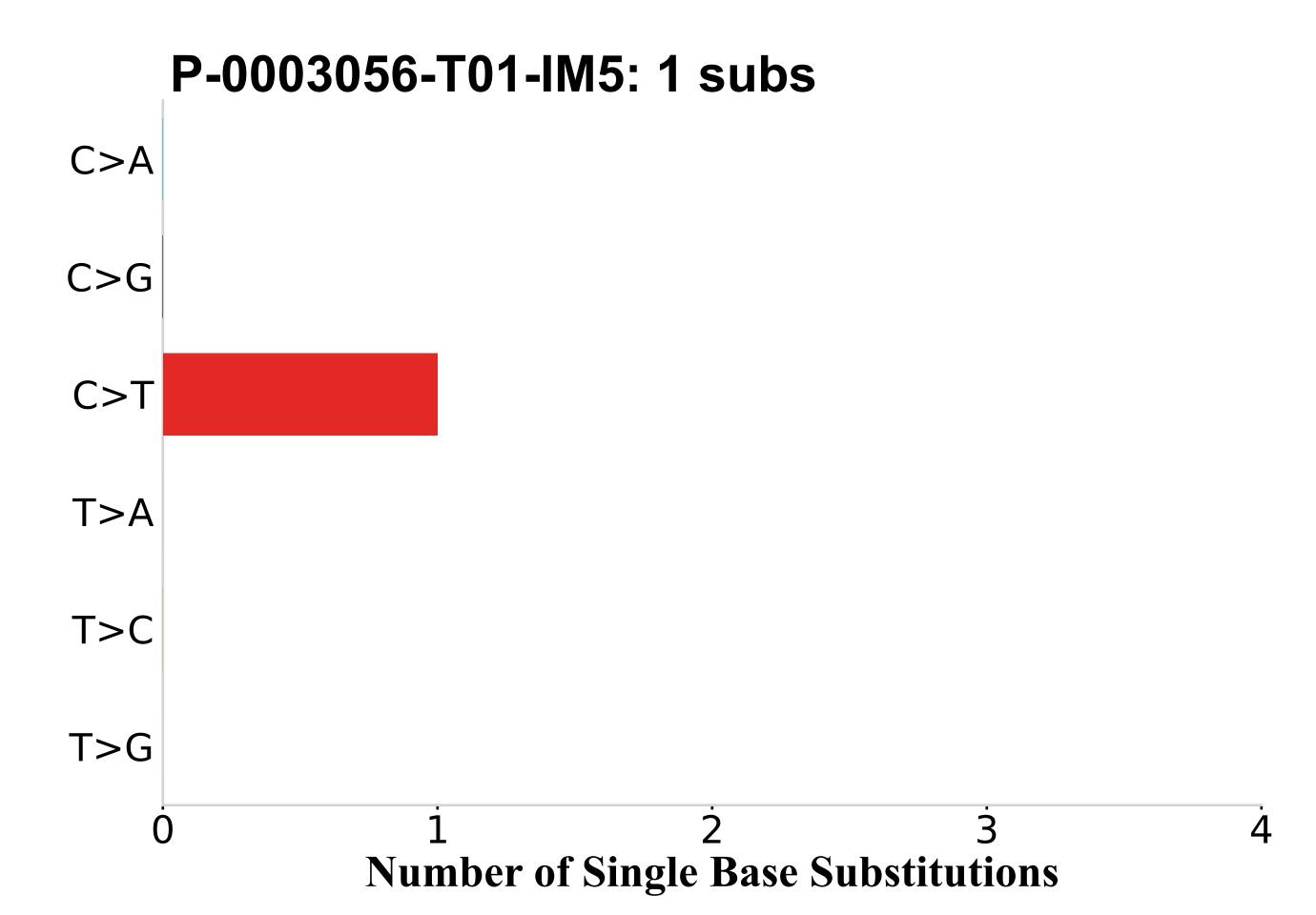


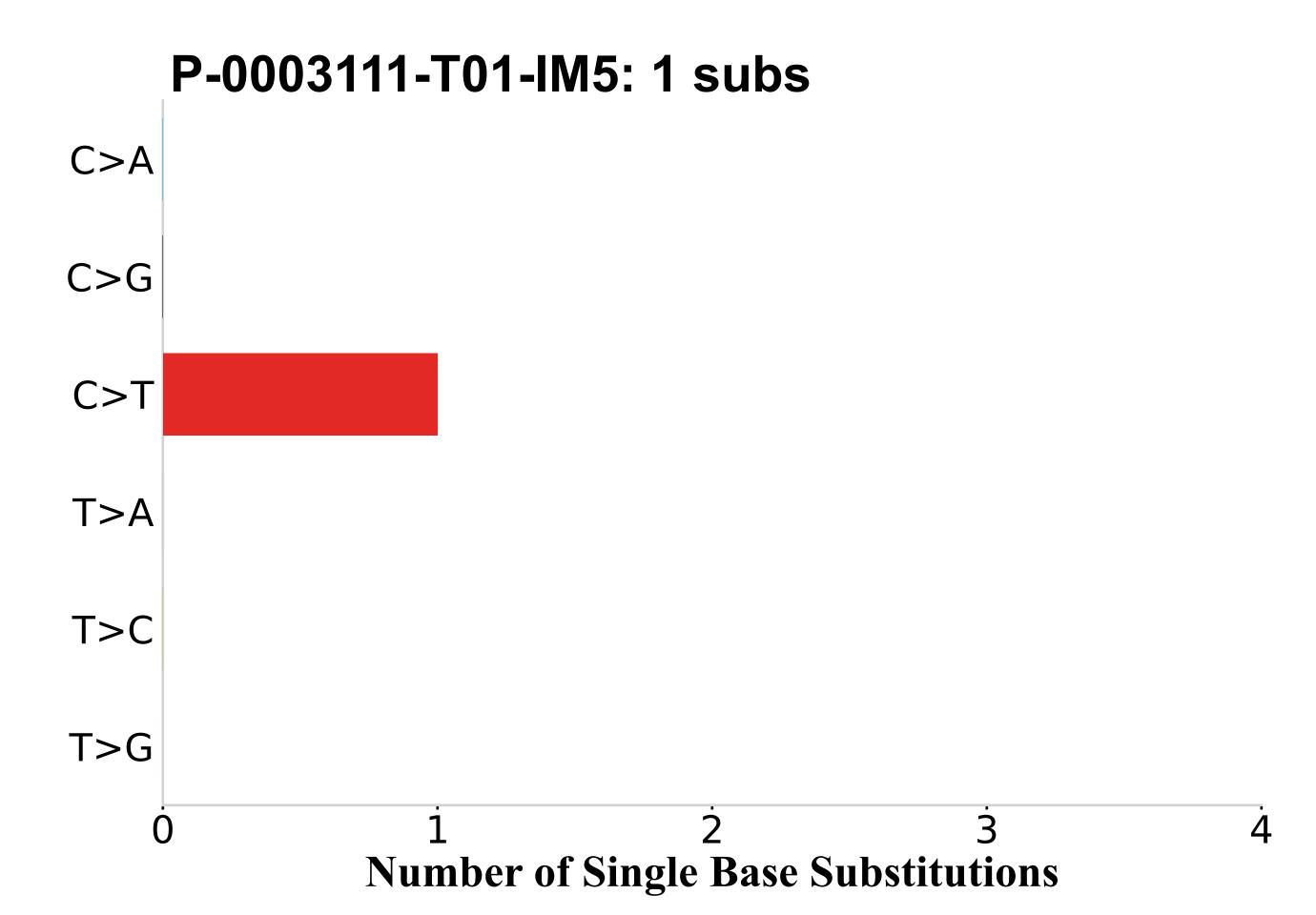
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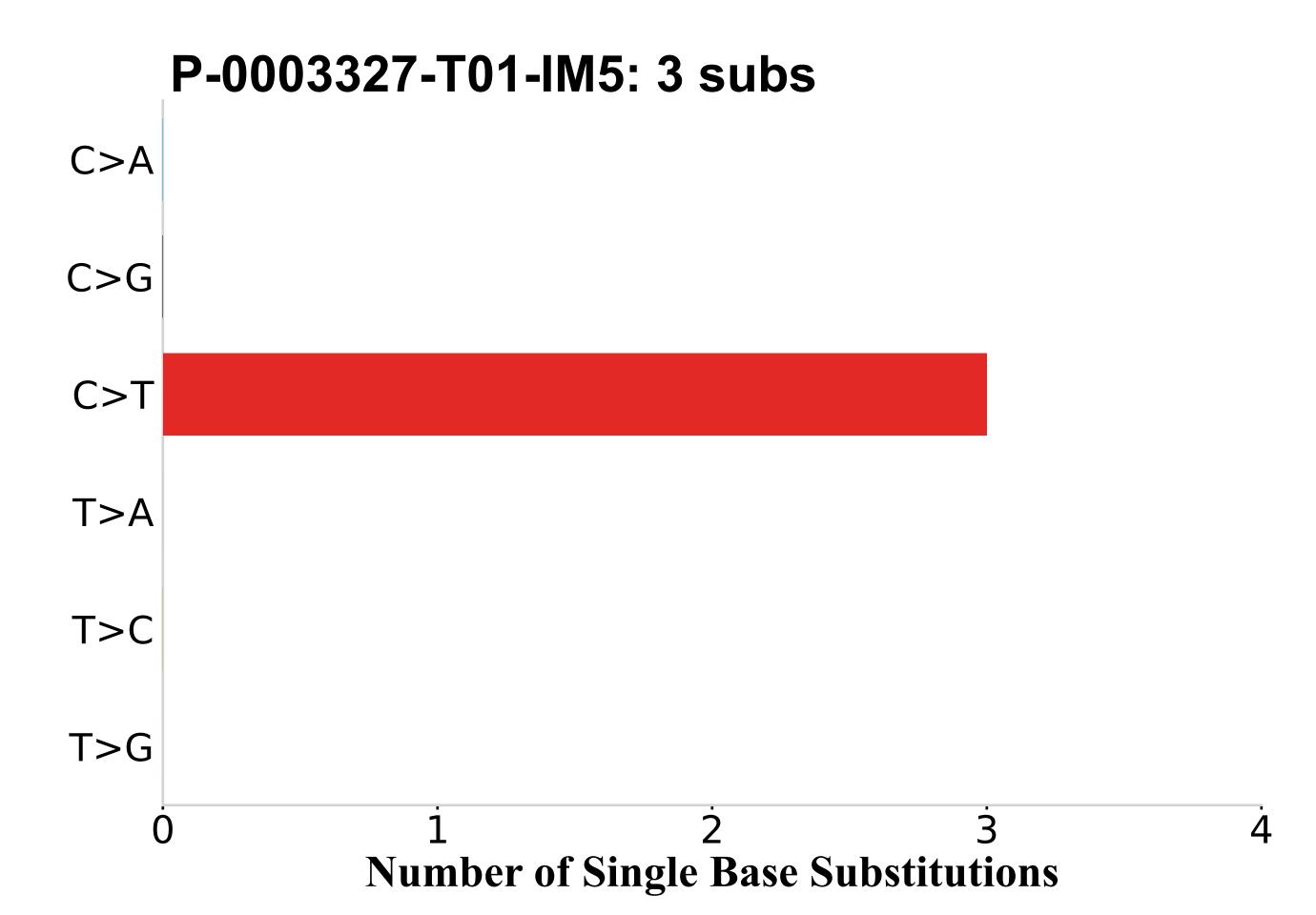


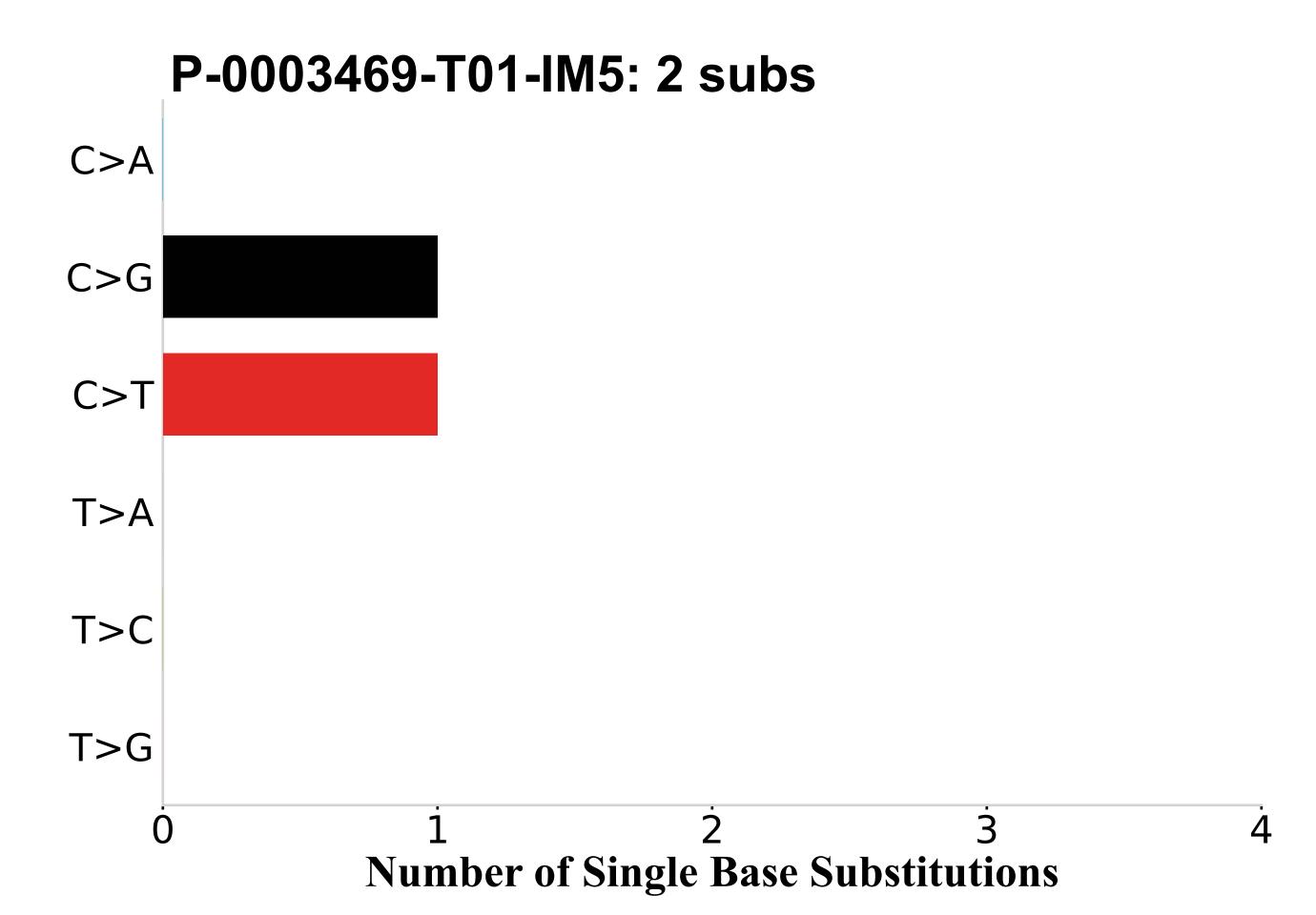


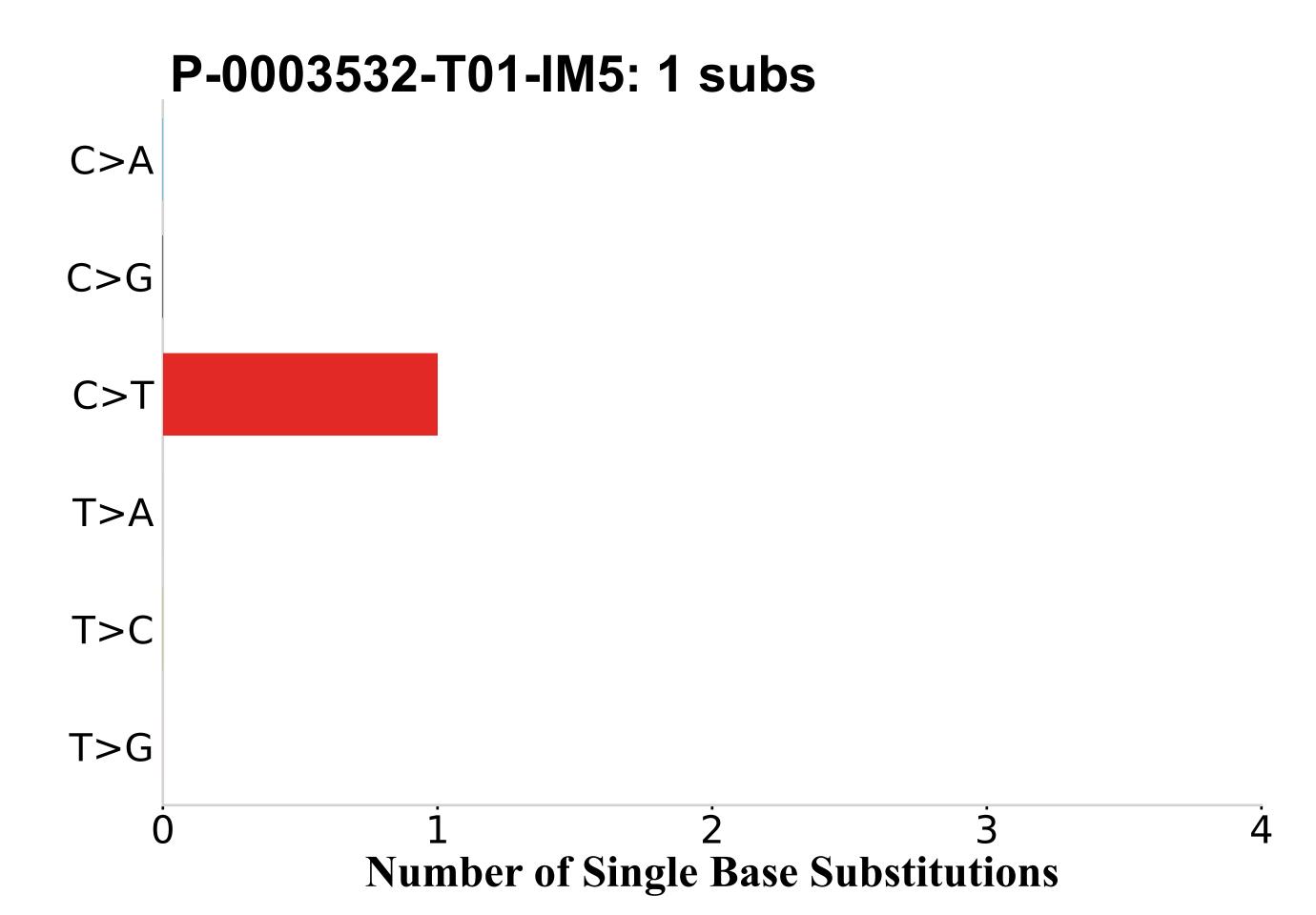




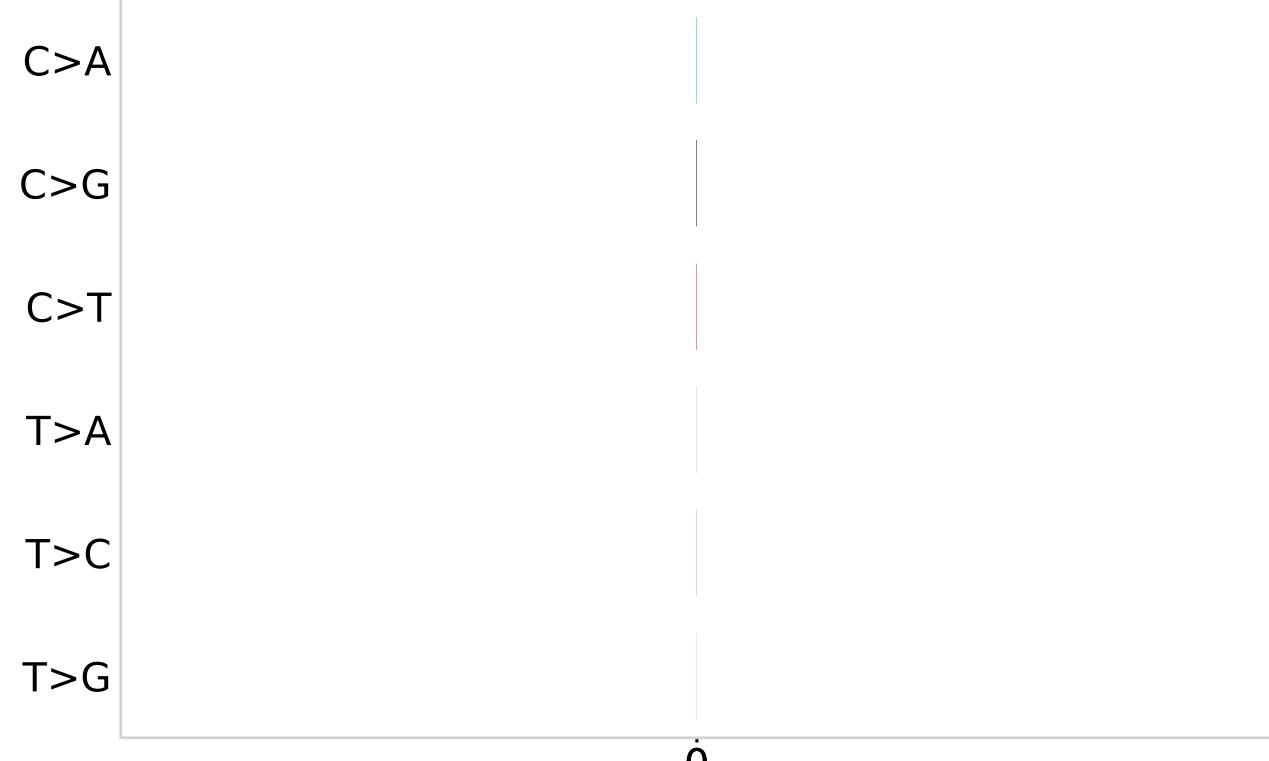




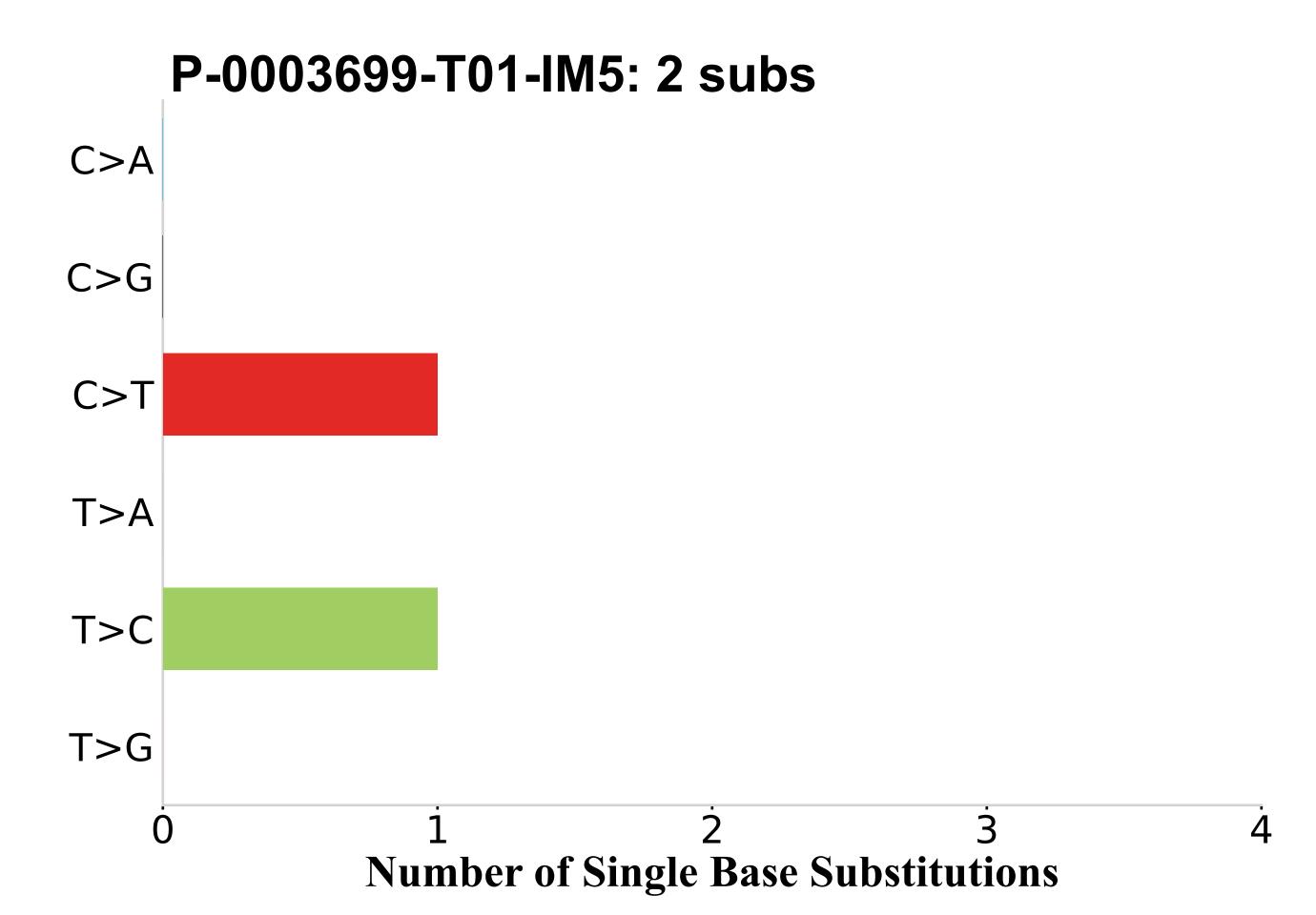


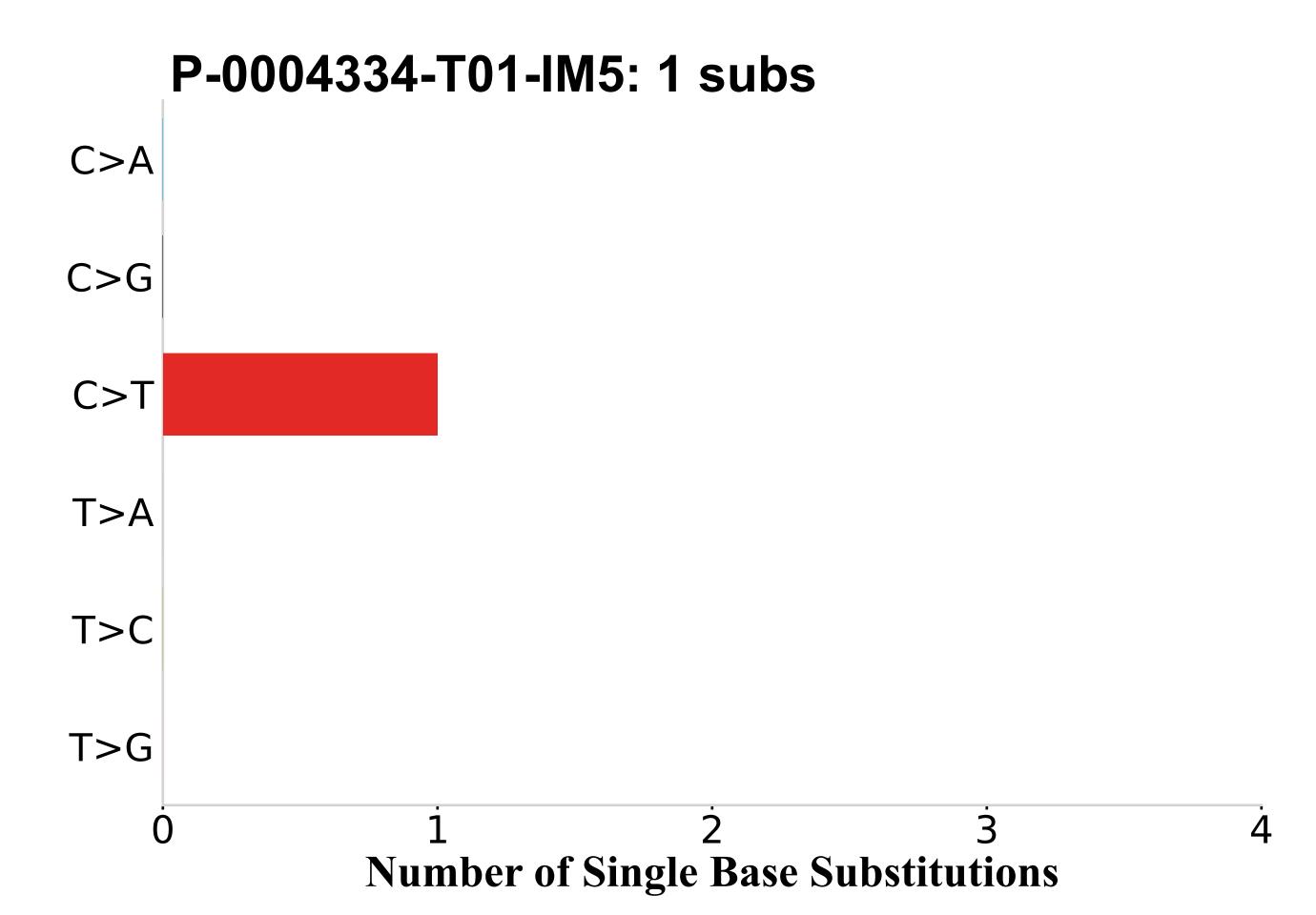


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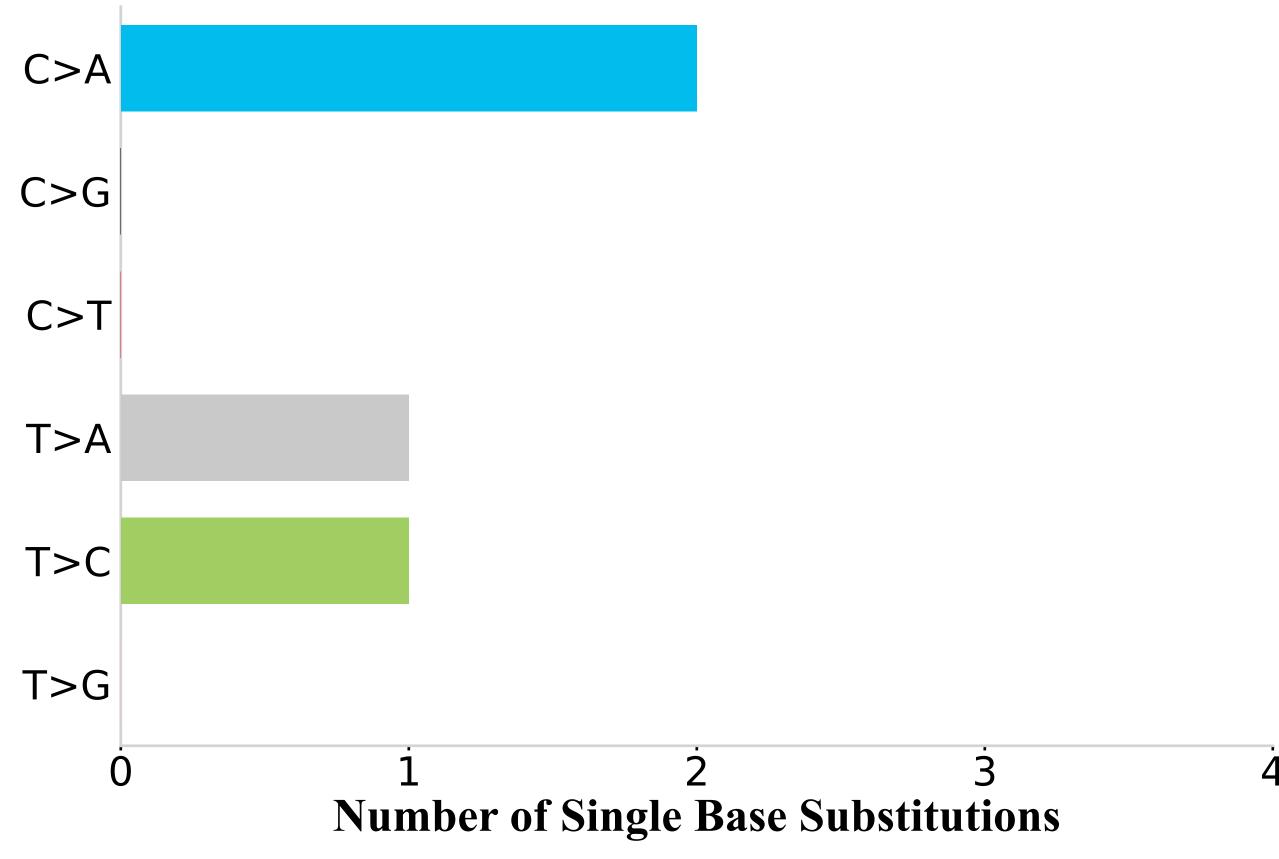


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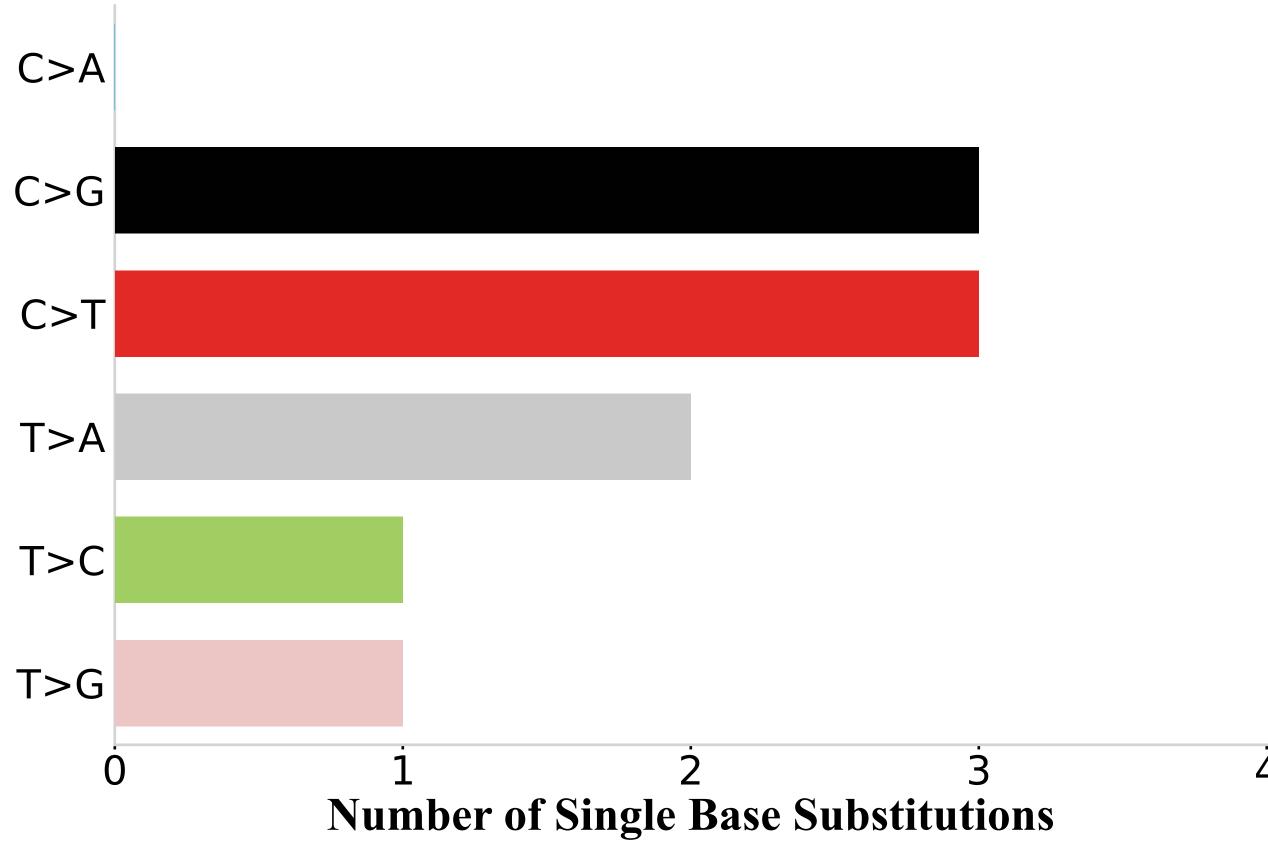


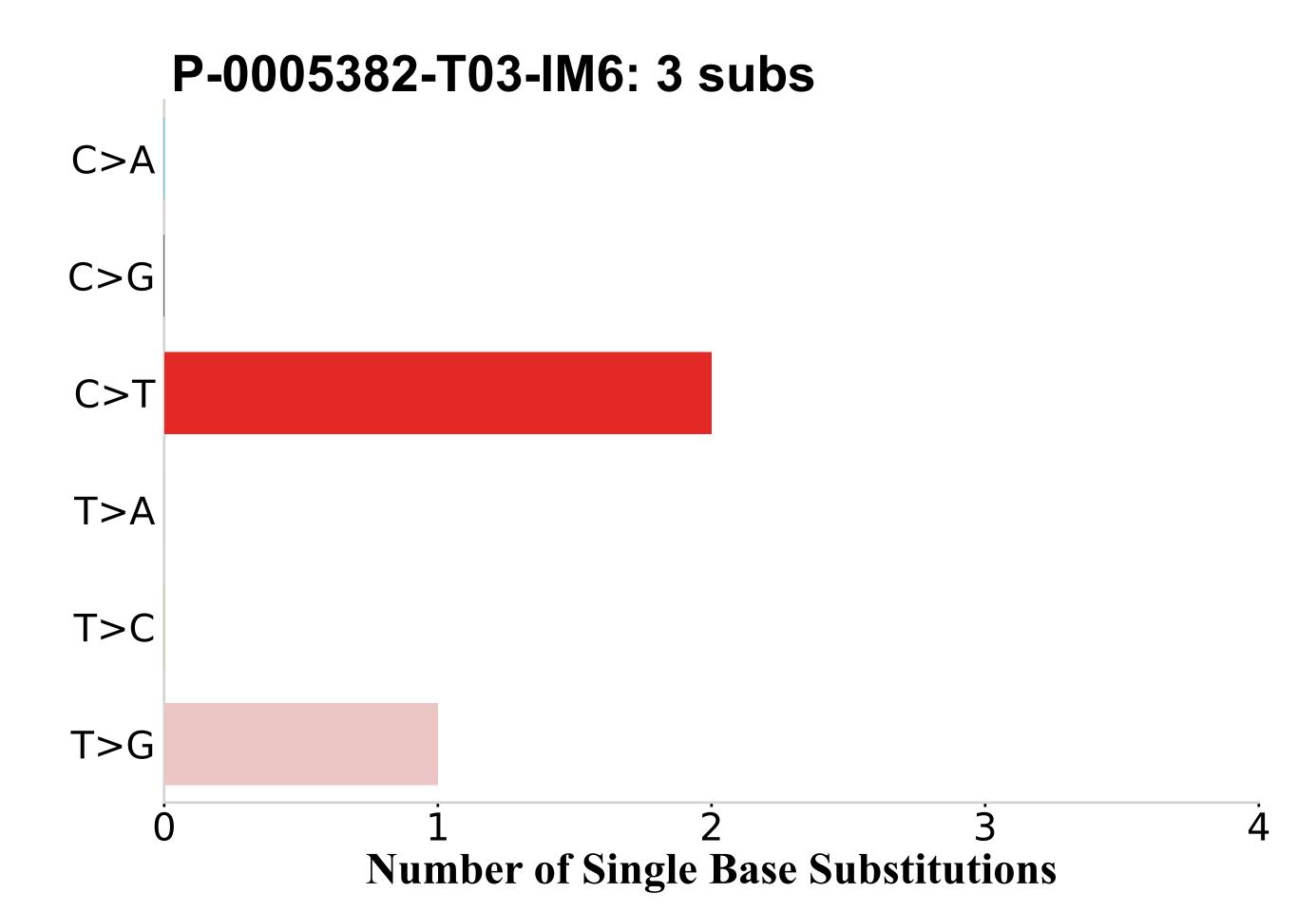




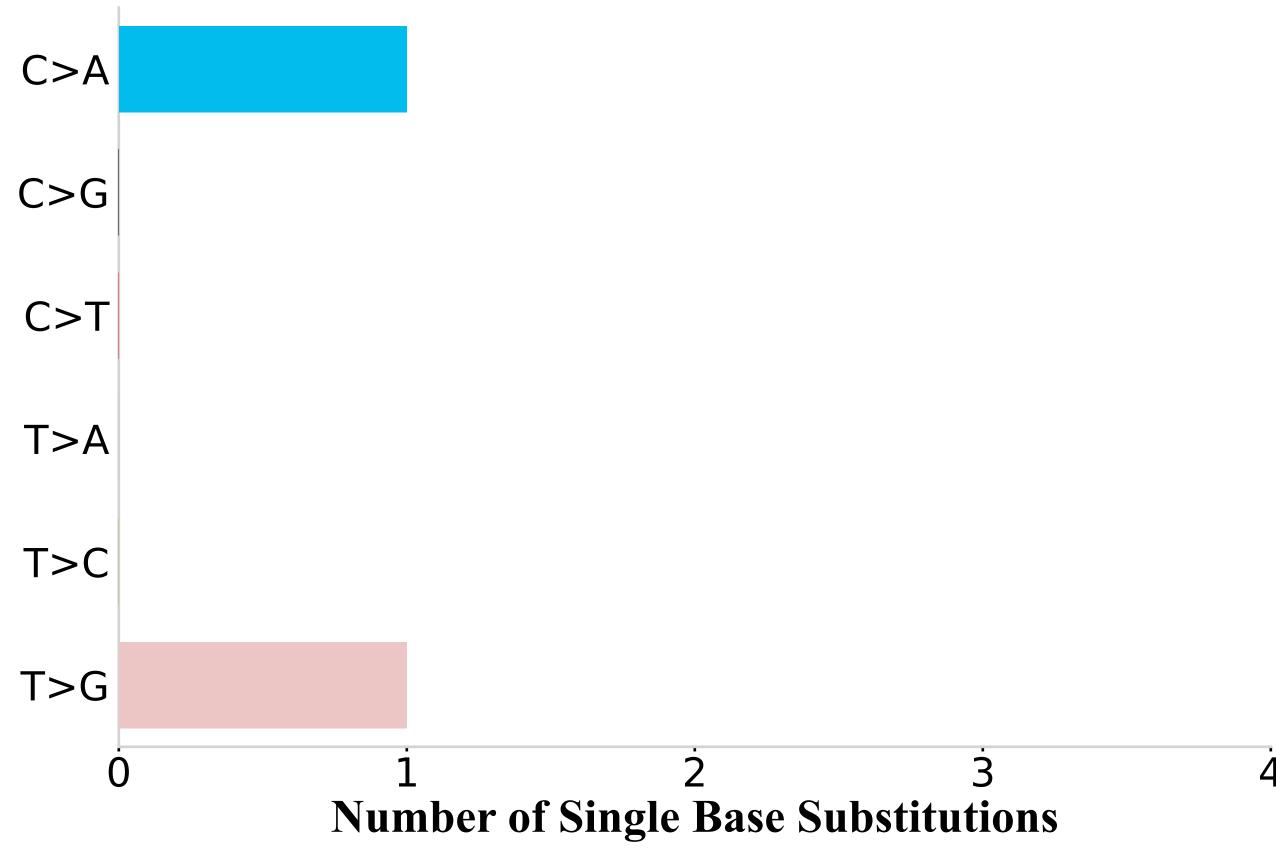


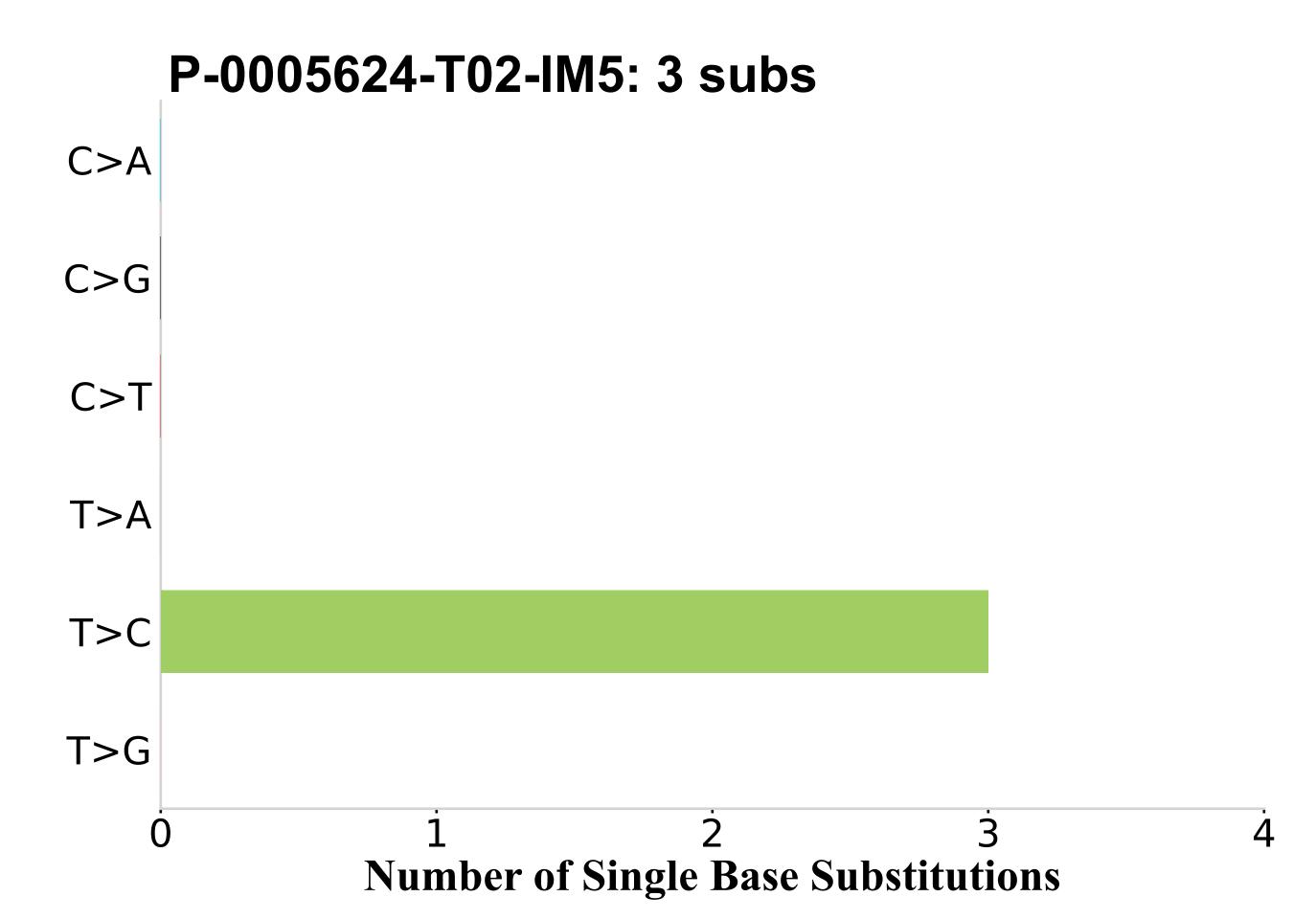


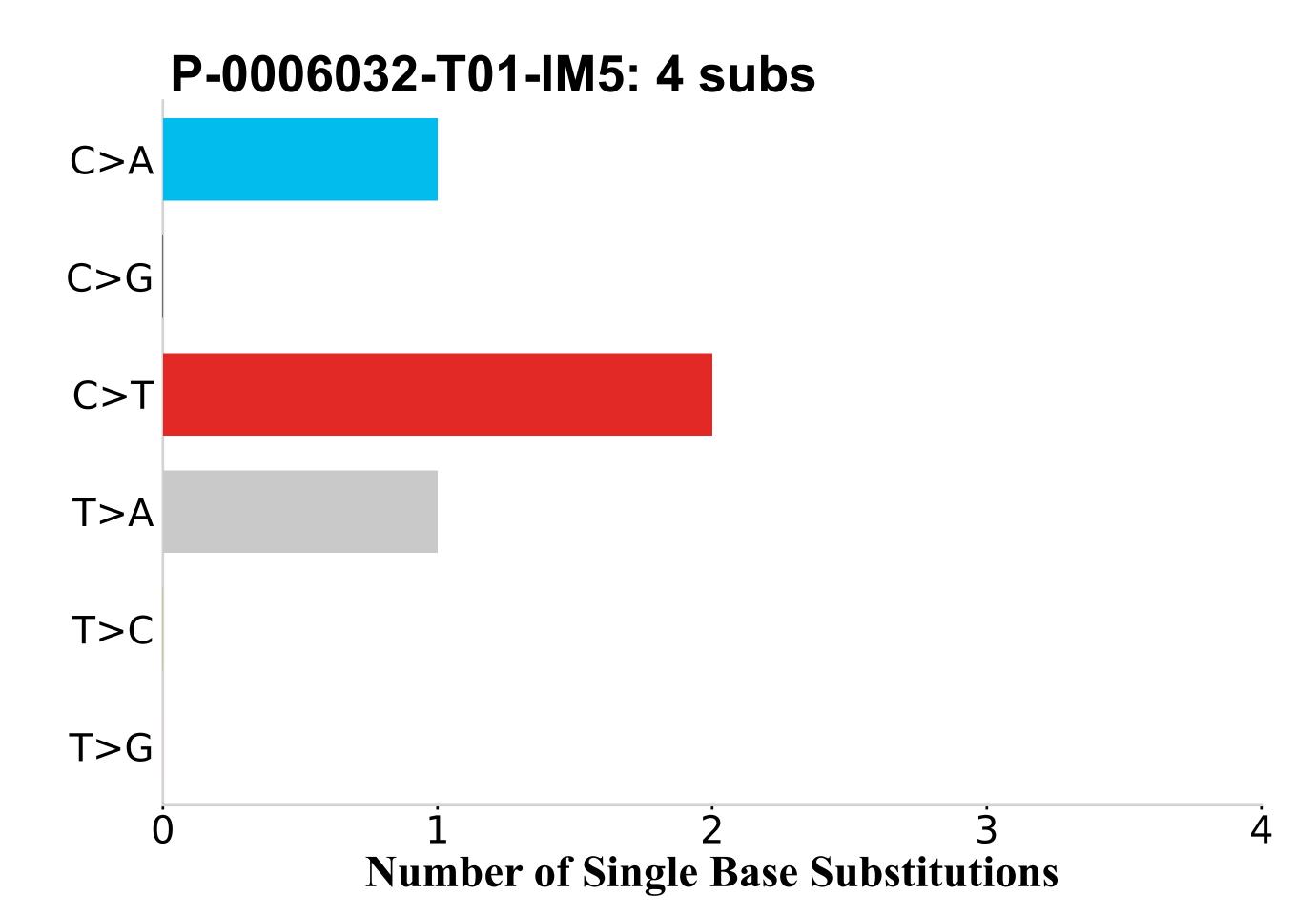


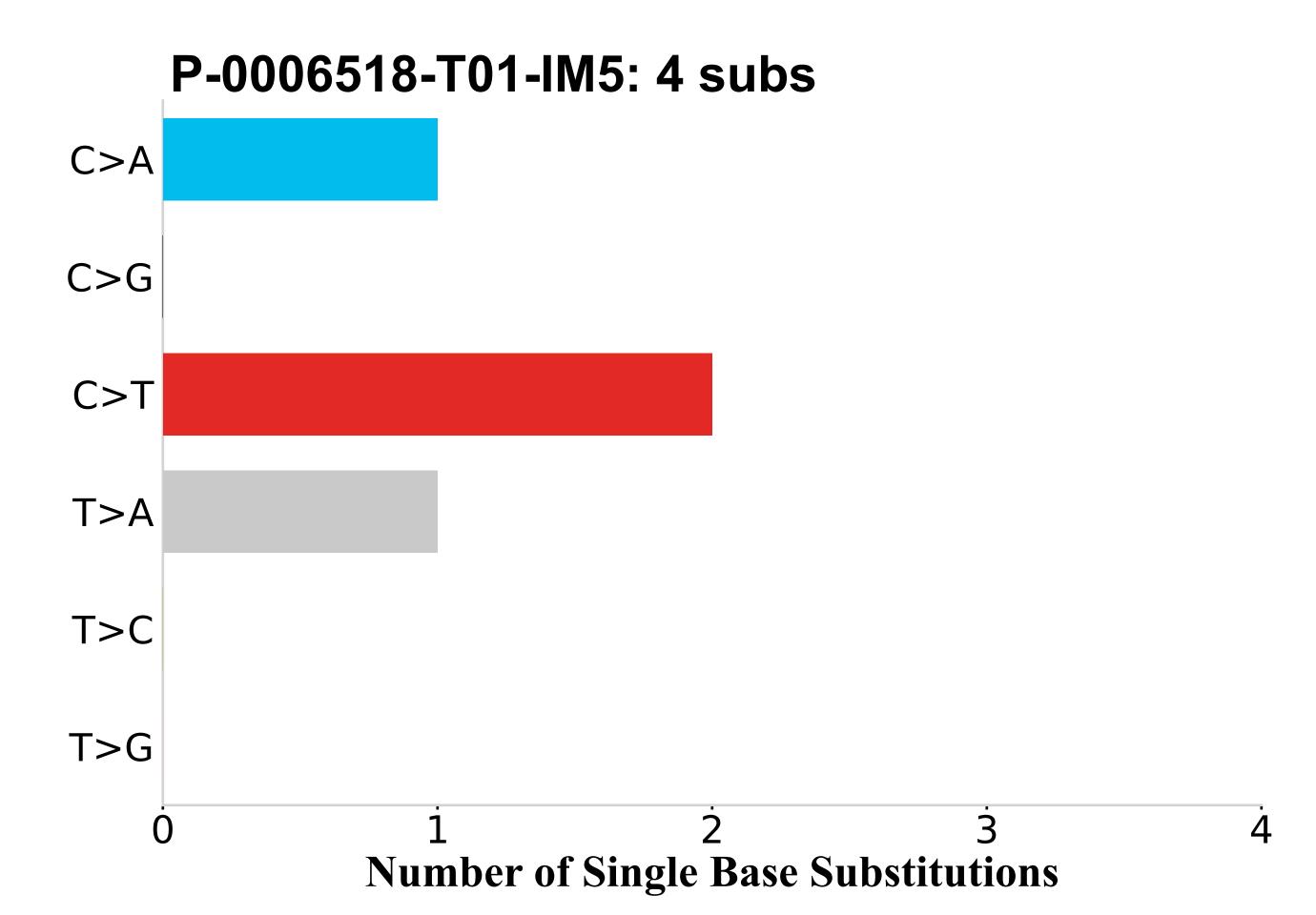


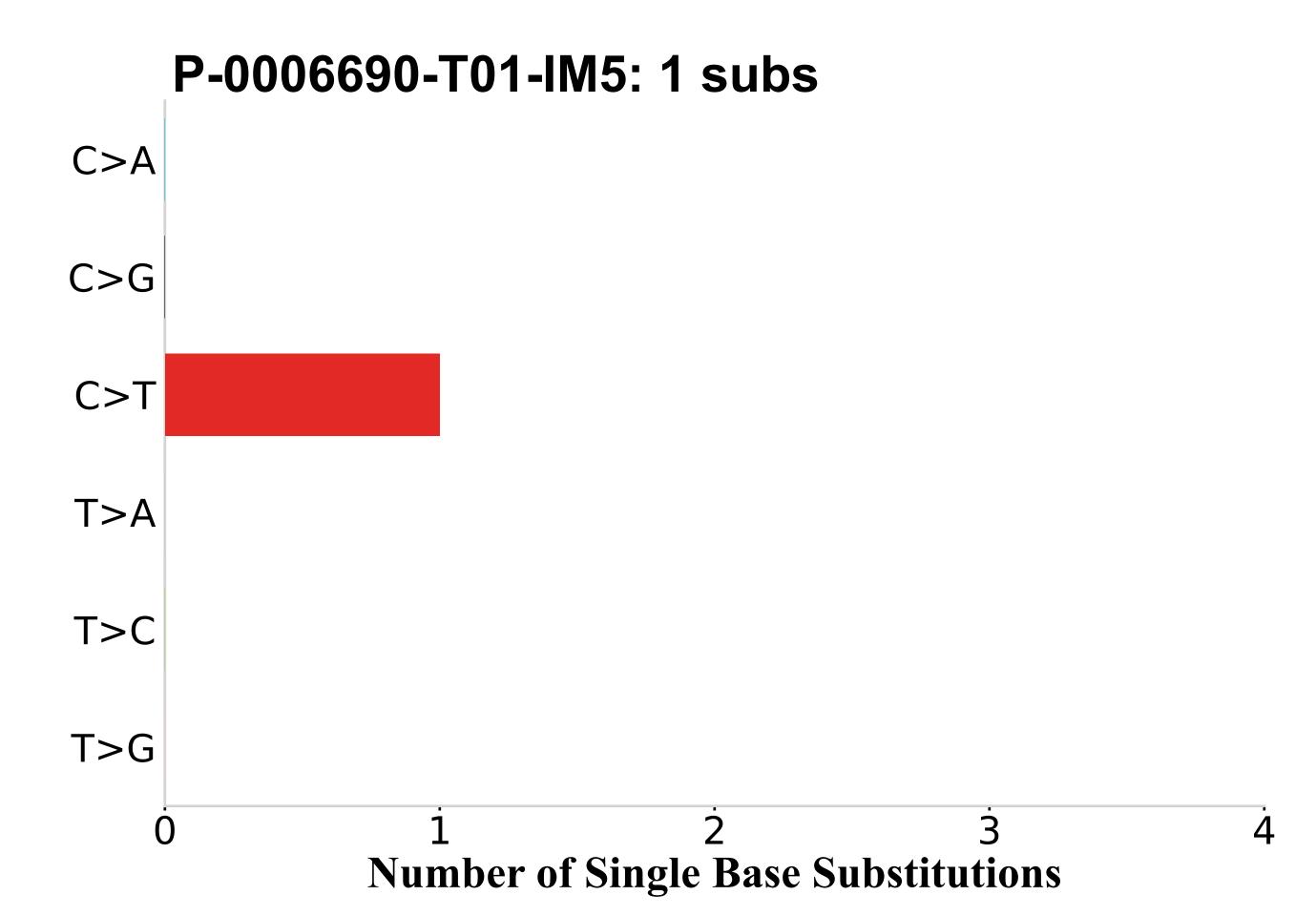
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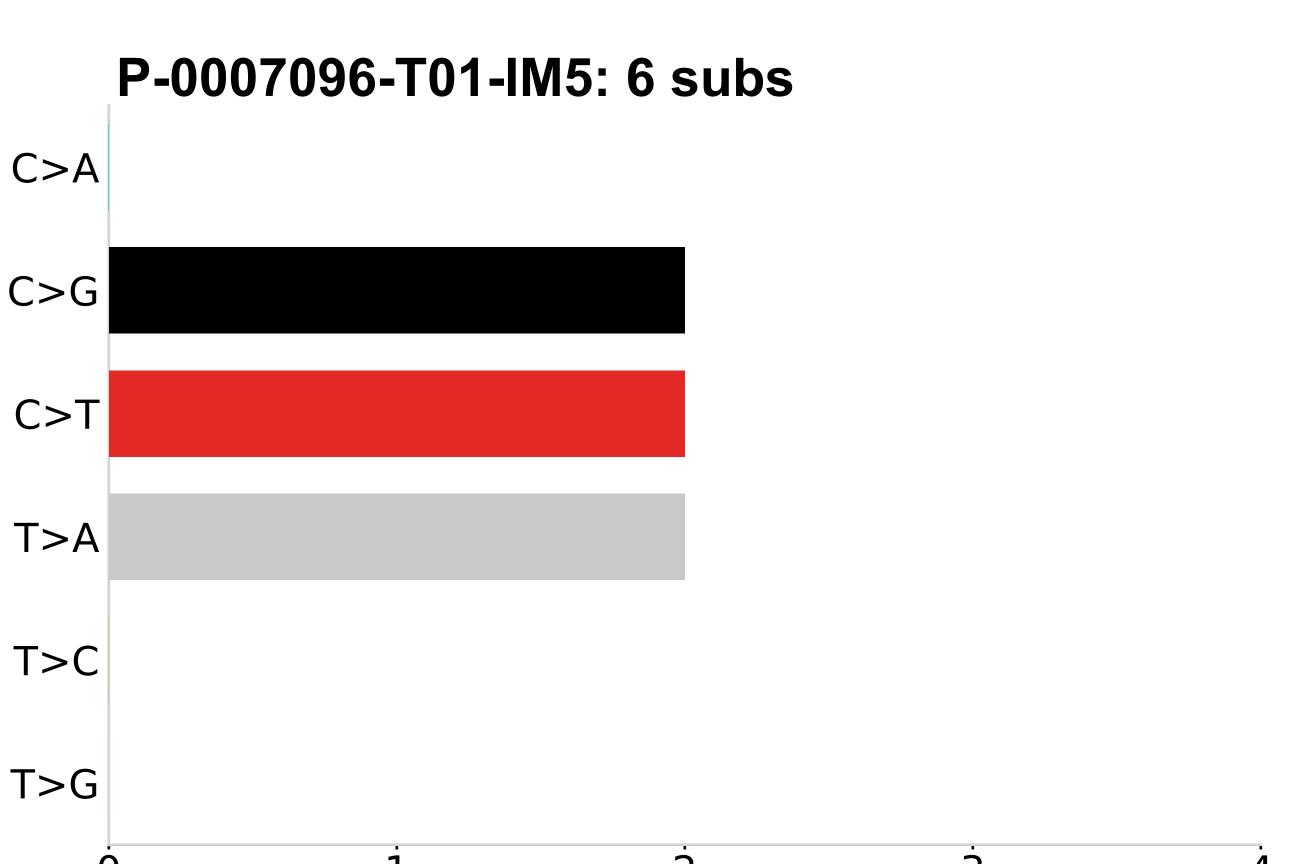




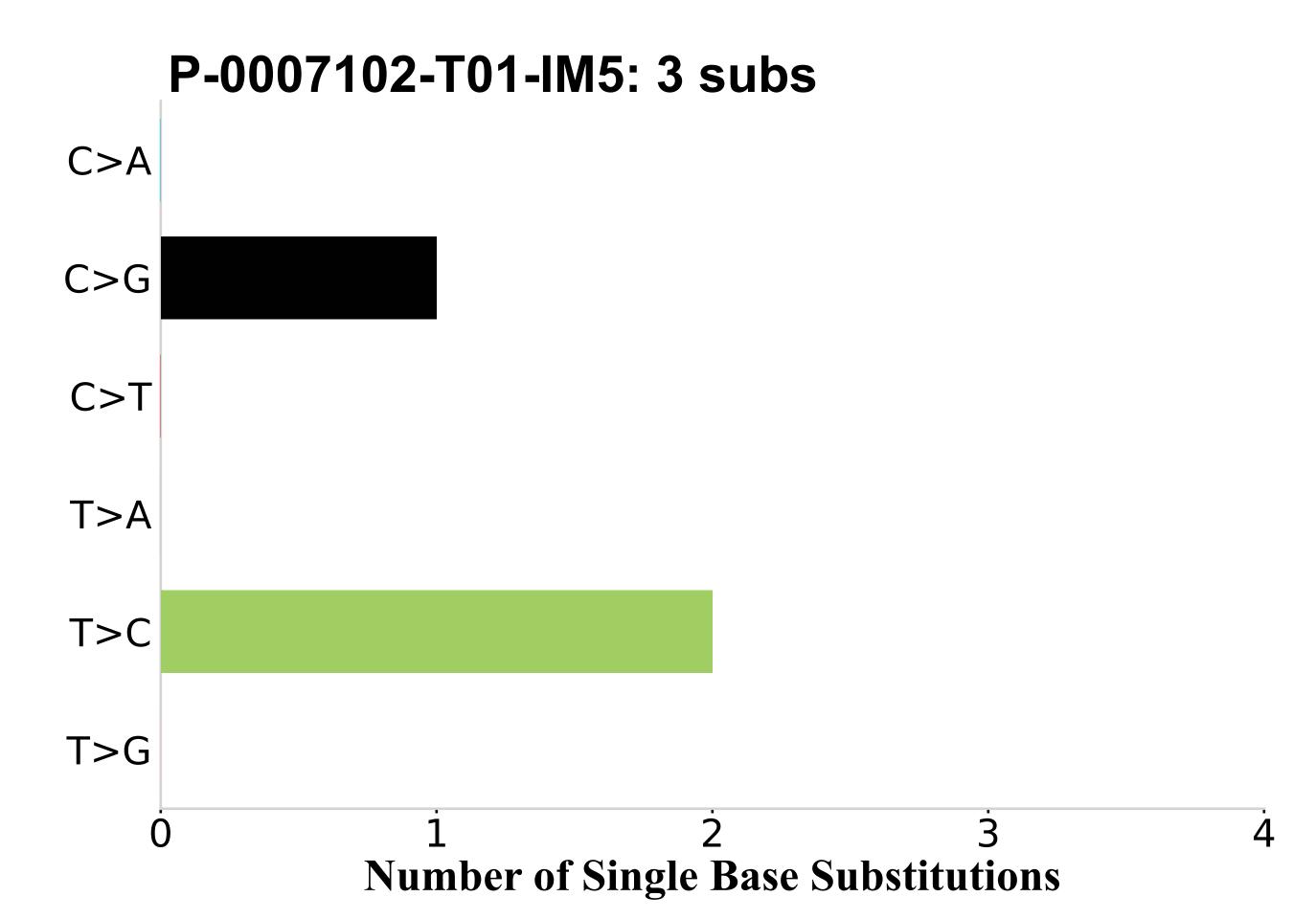


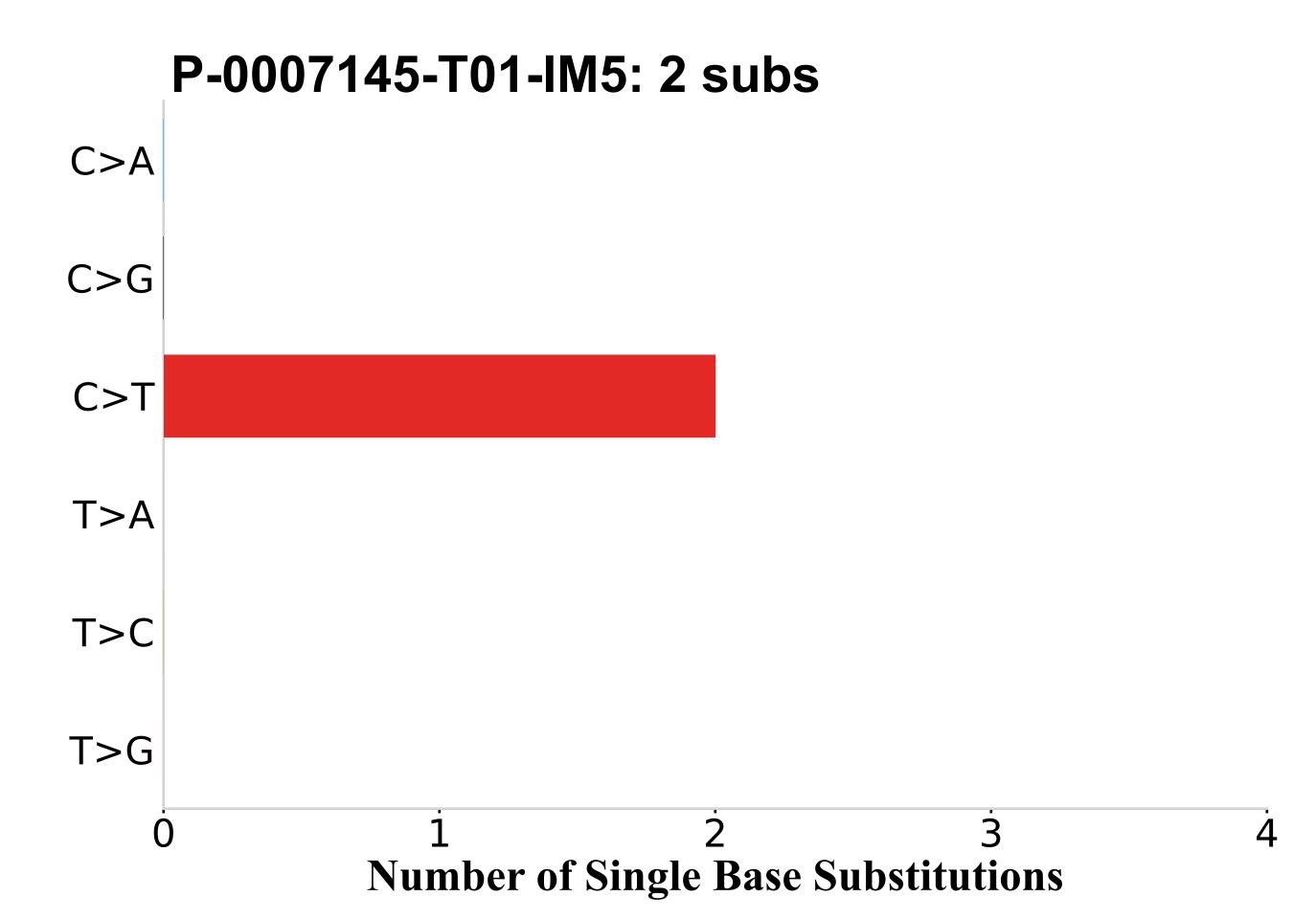




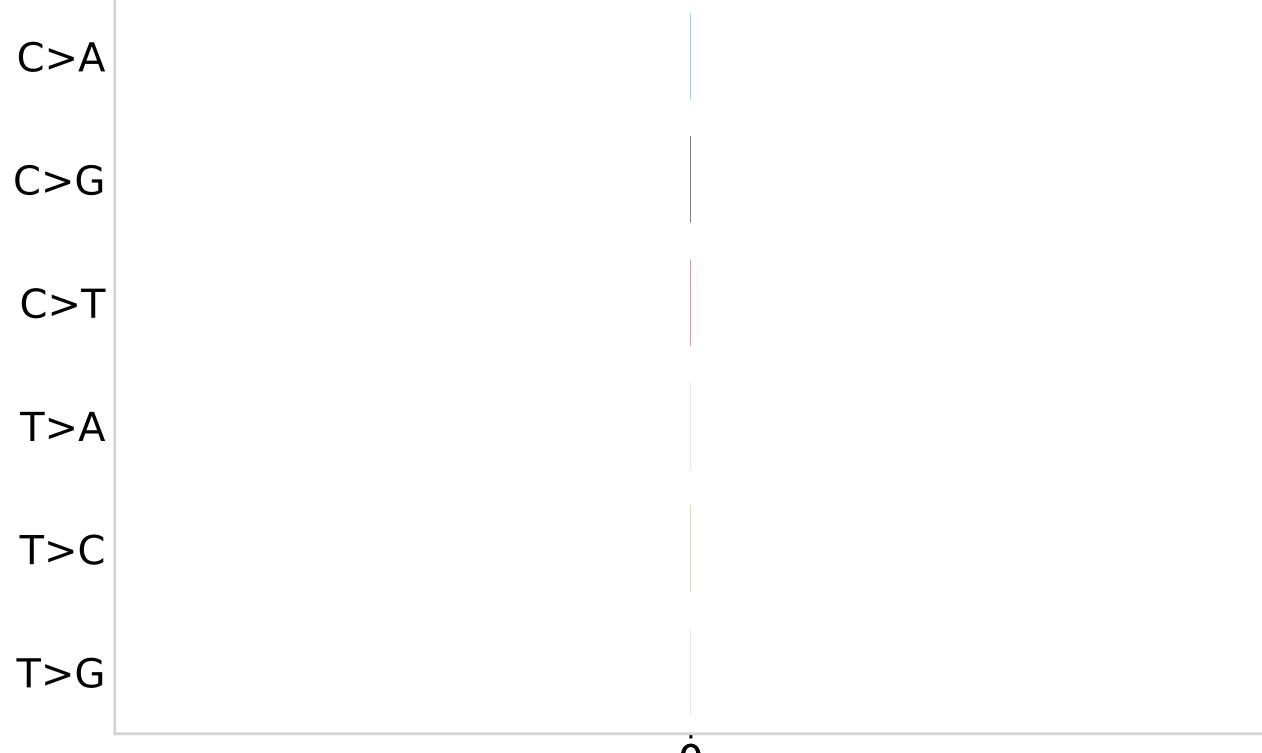


Number of Single Base Substitutions

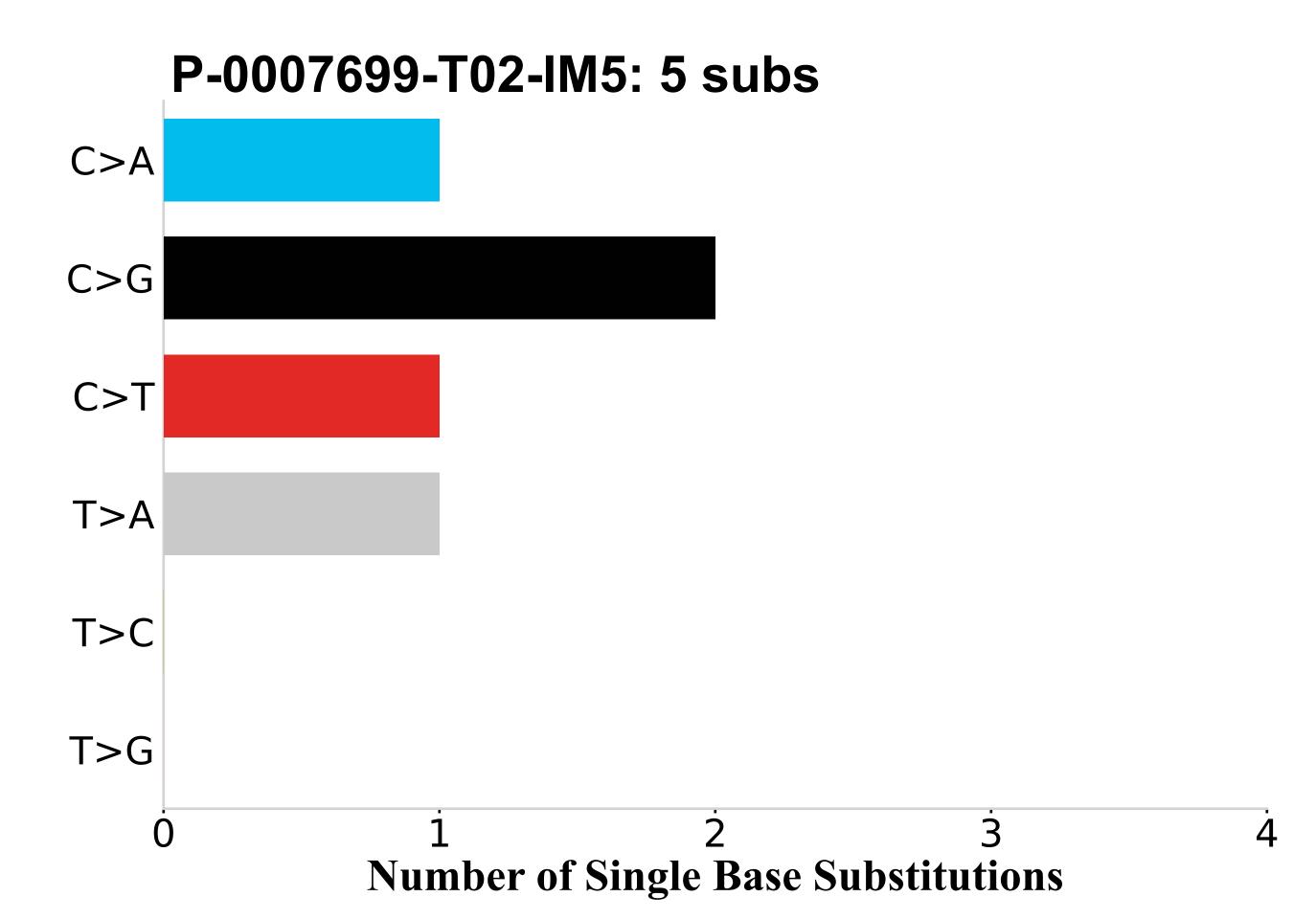


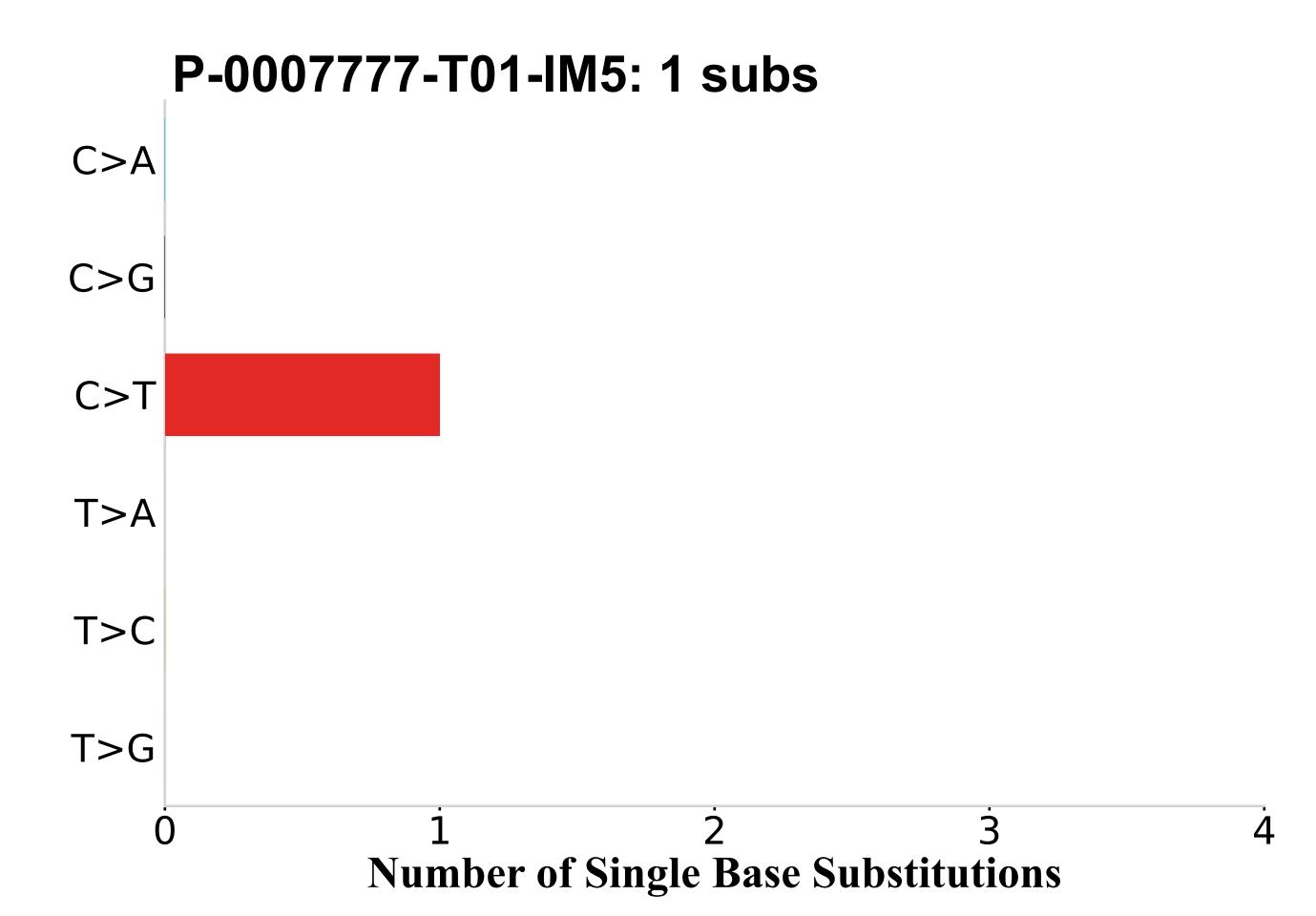


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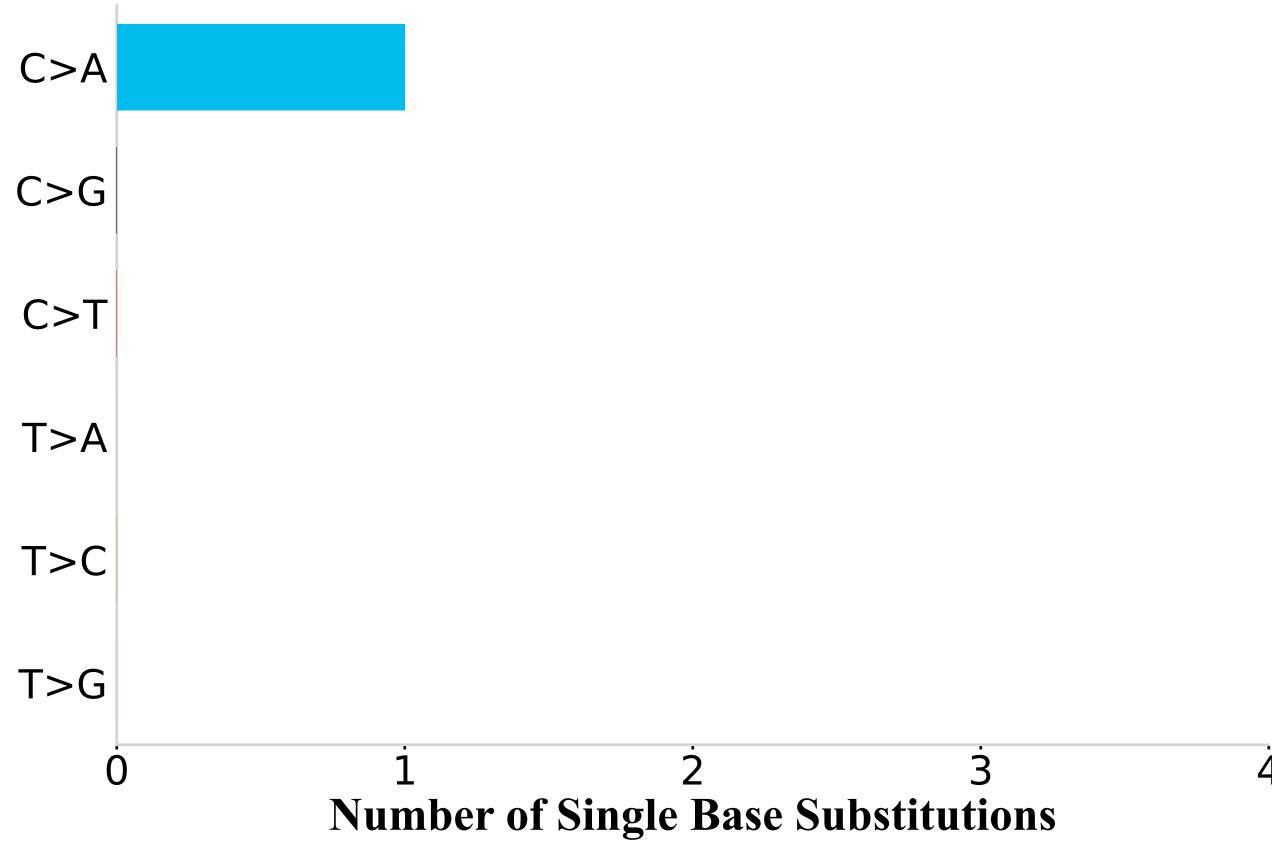


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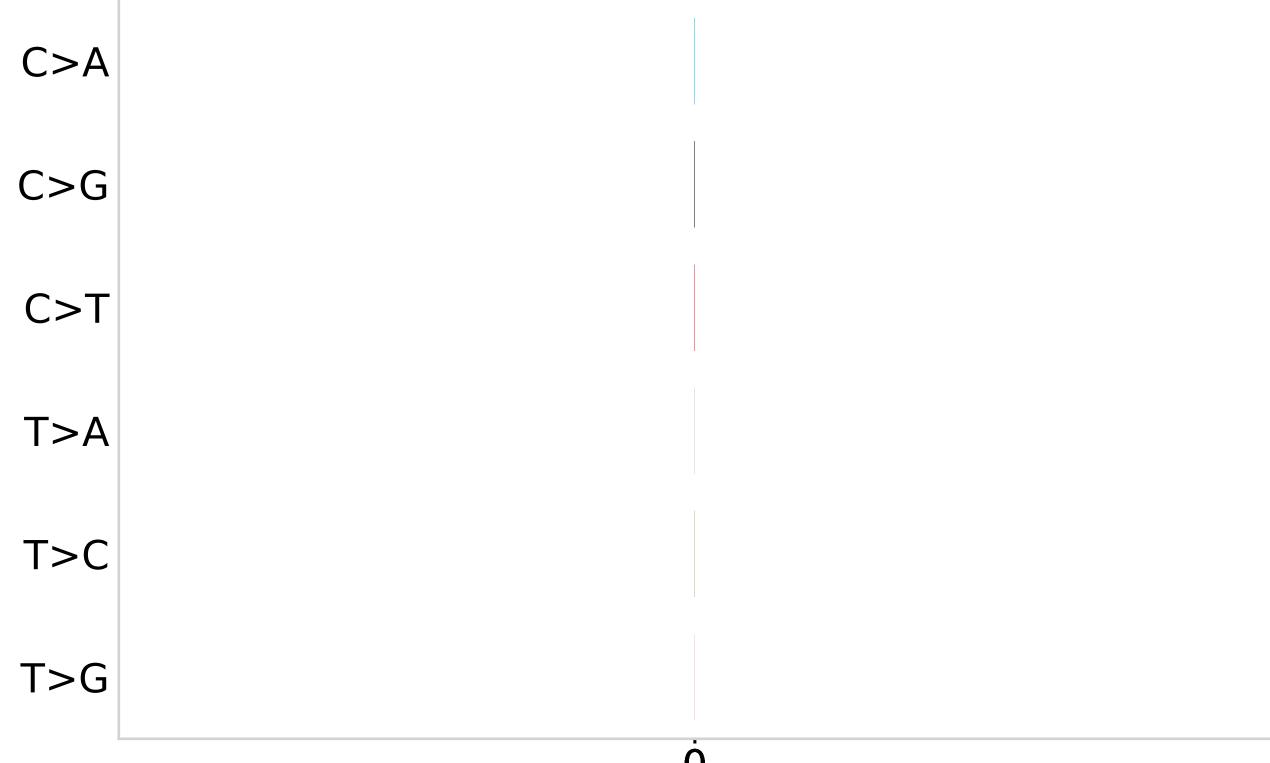






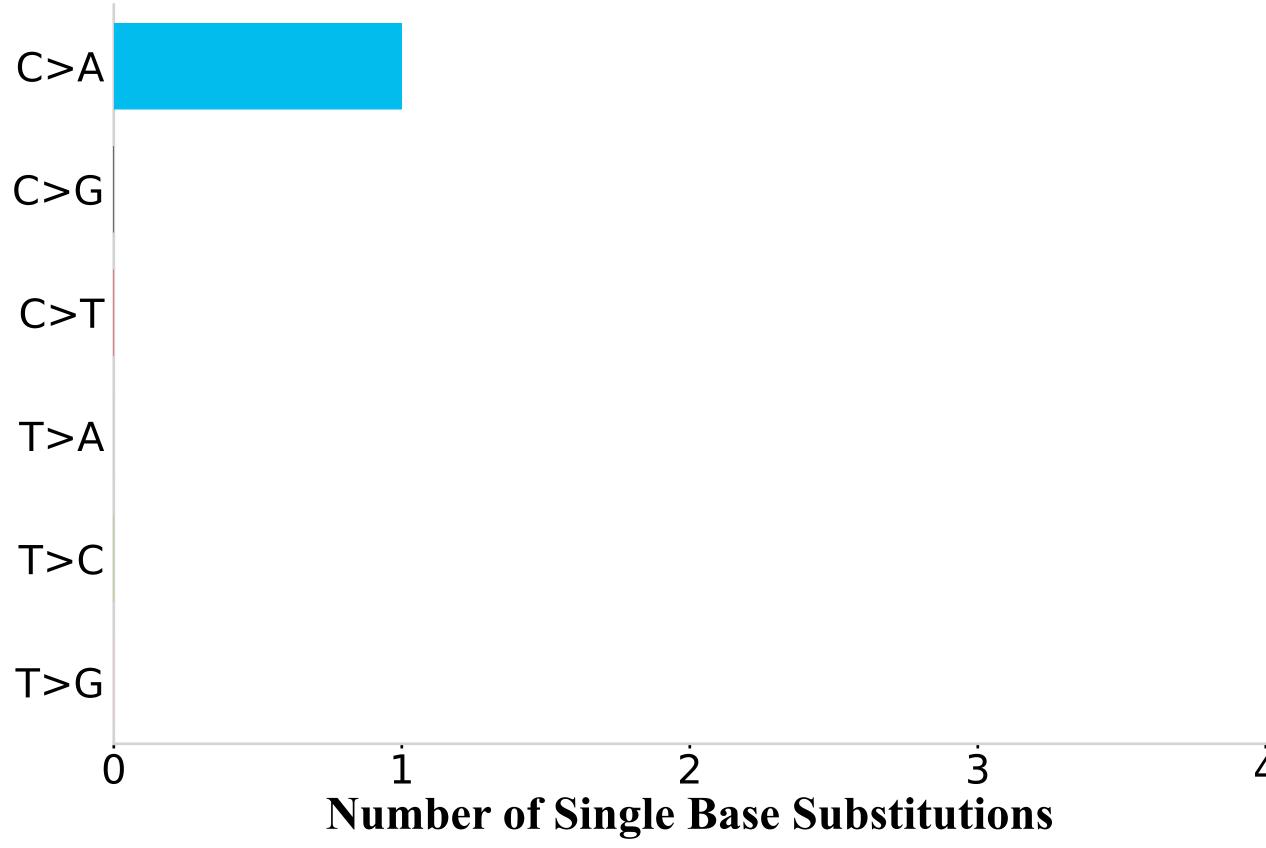


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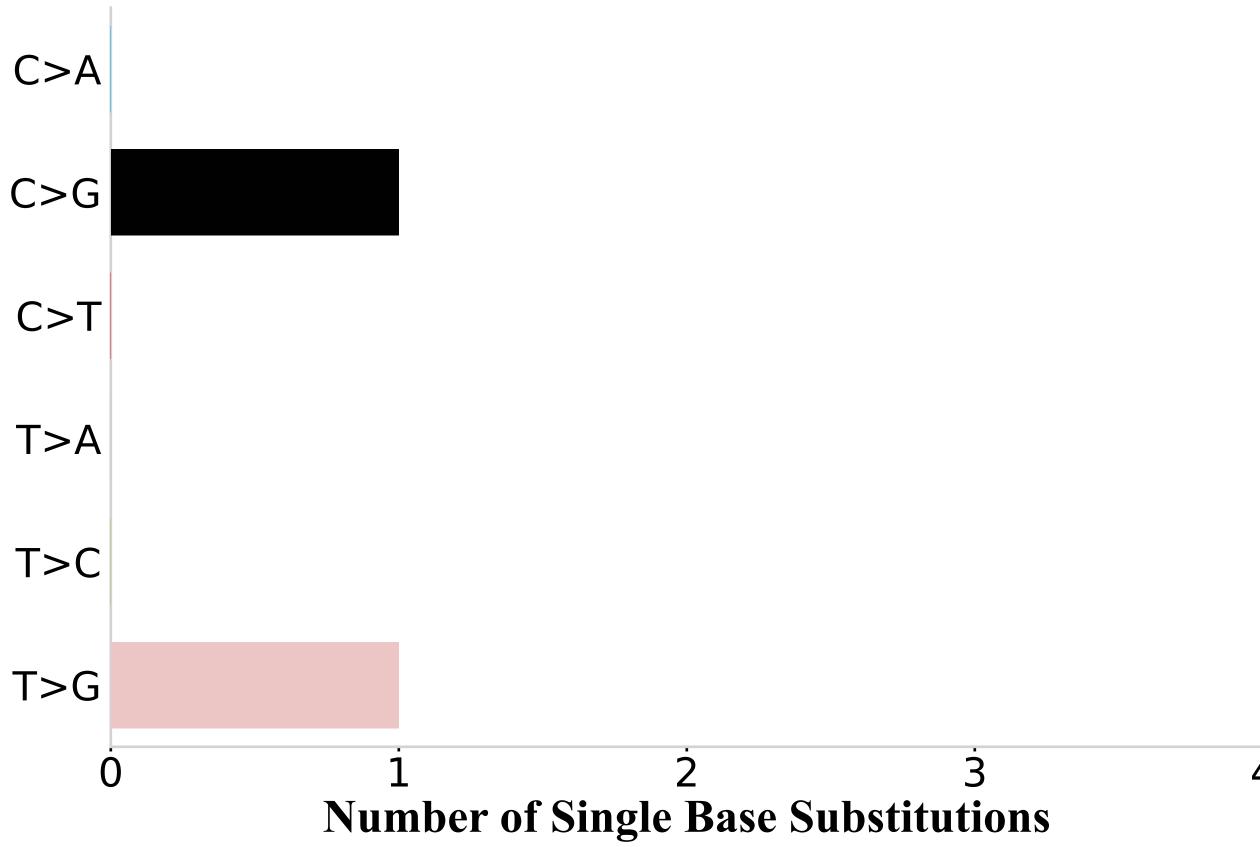


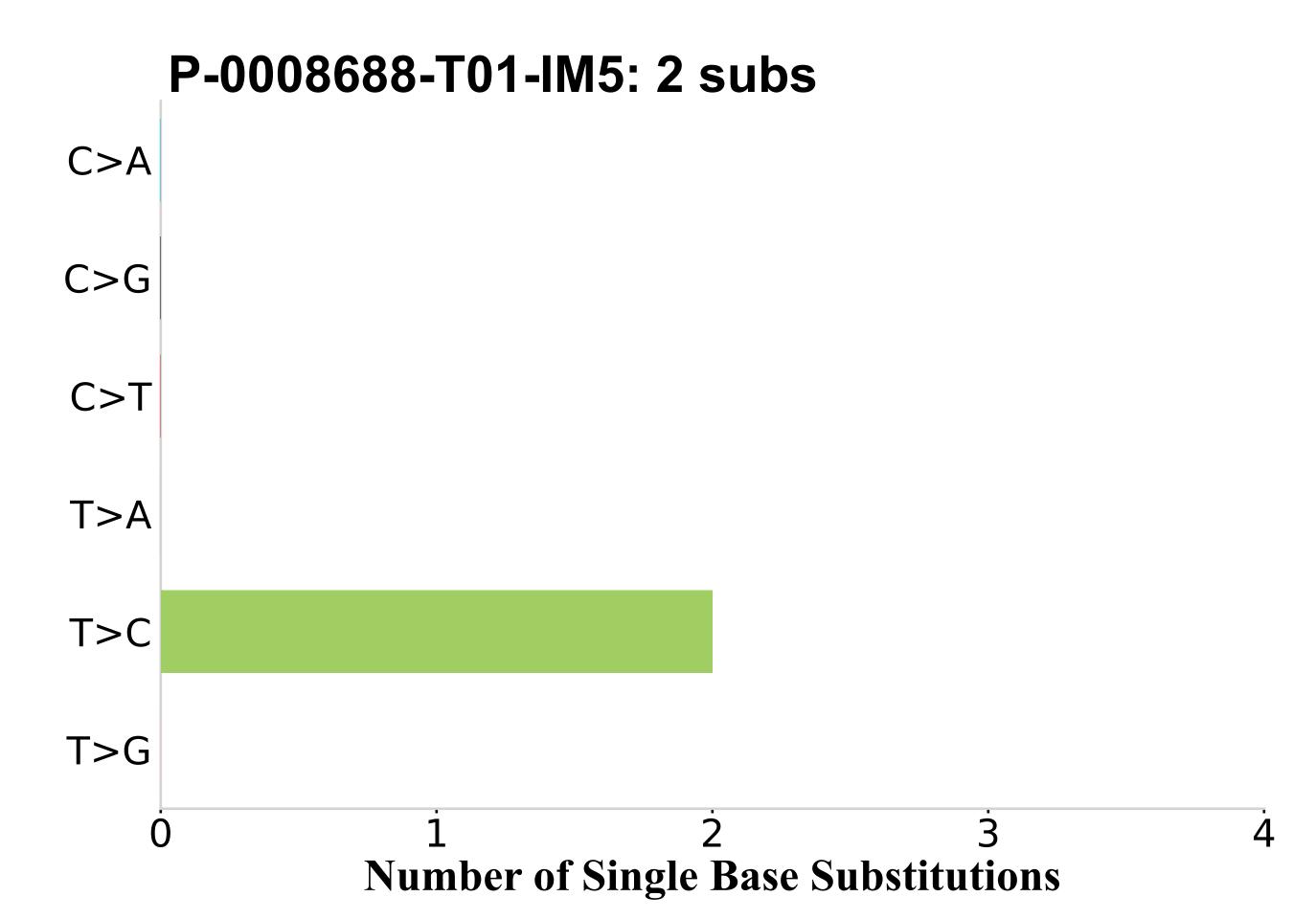
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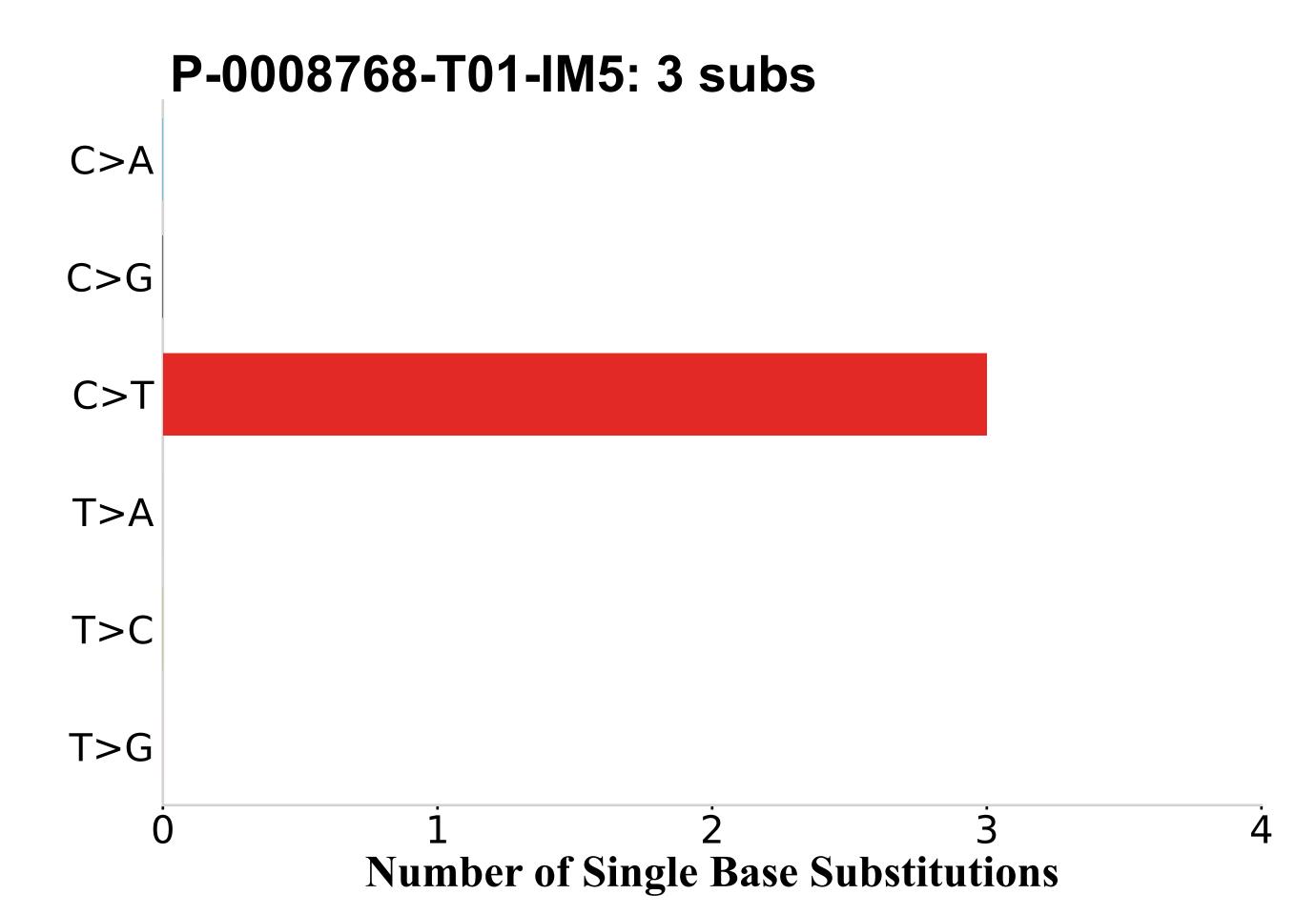


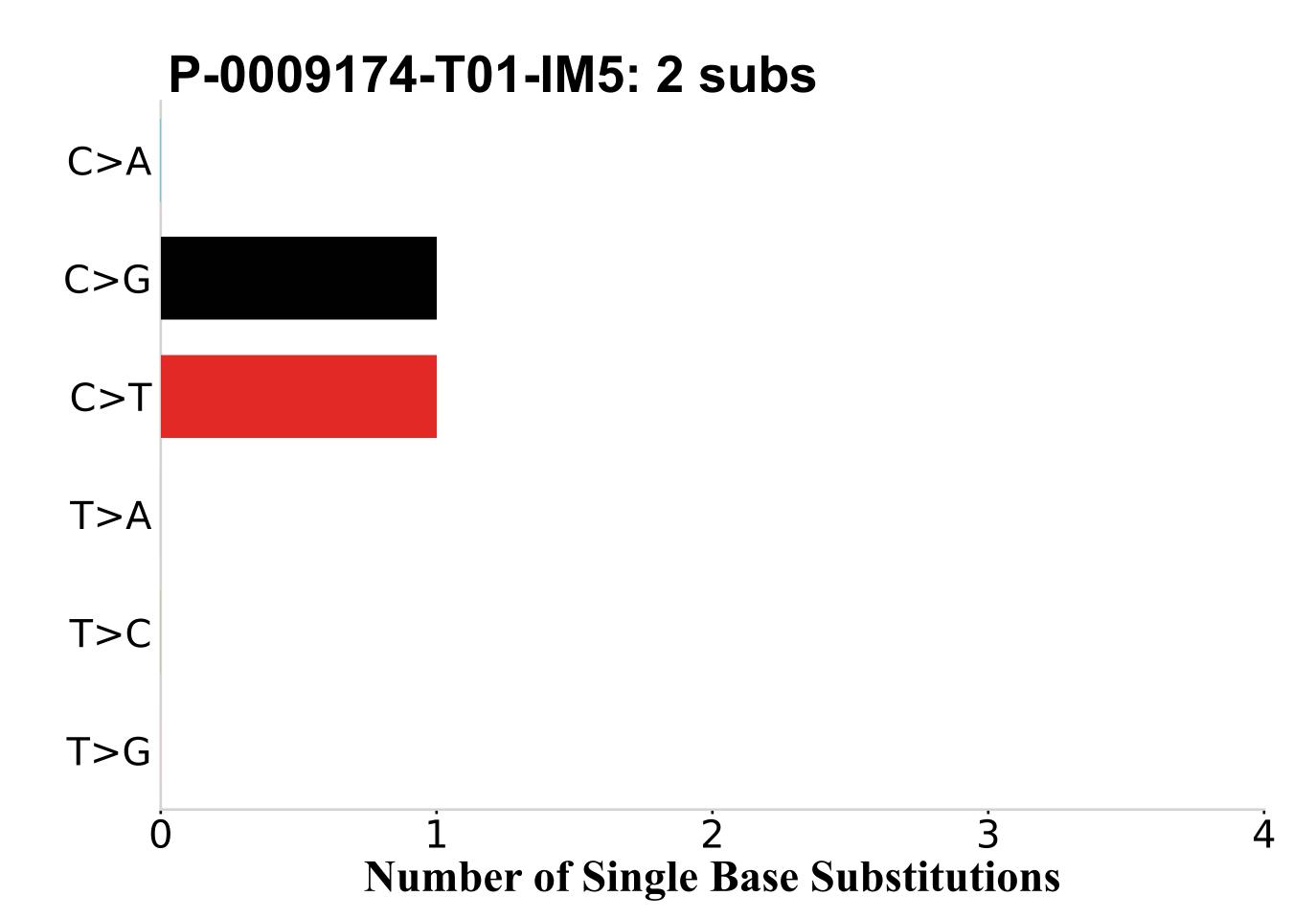


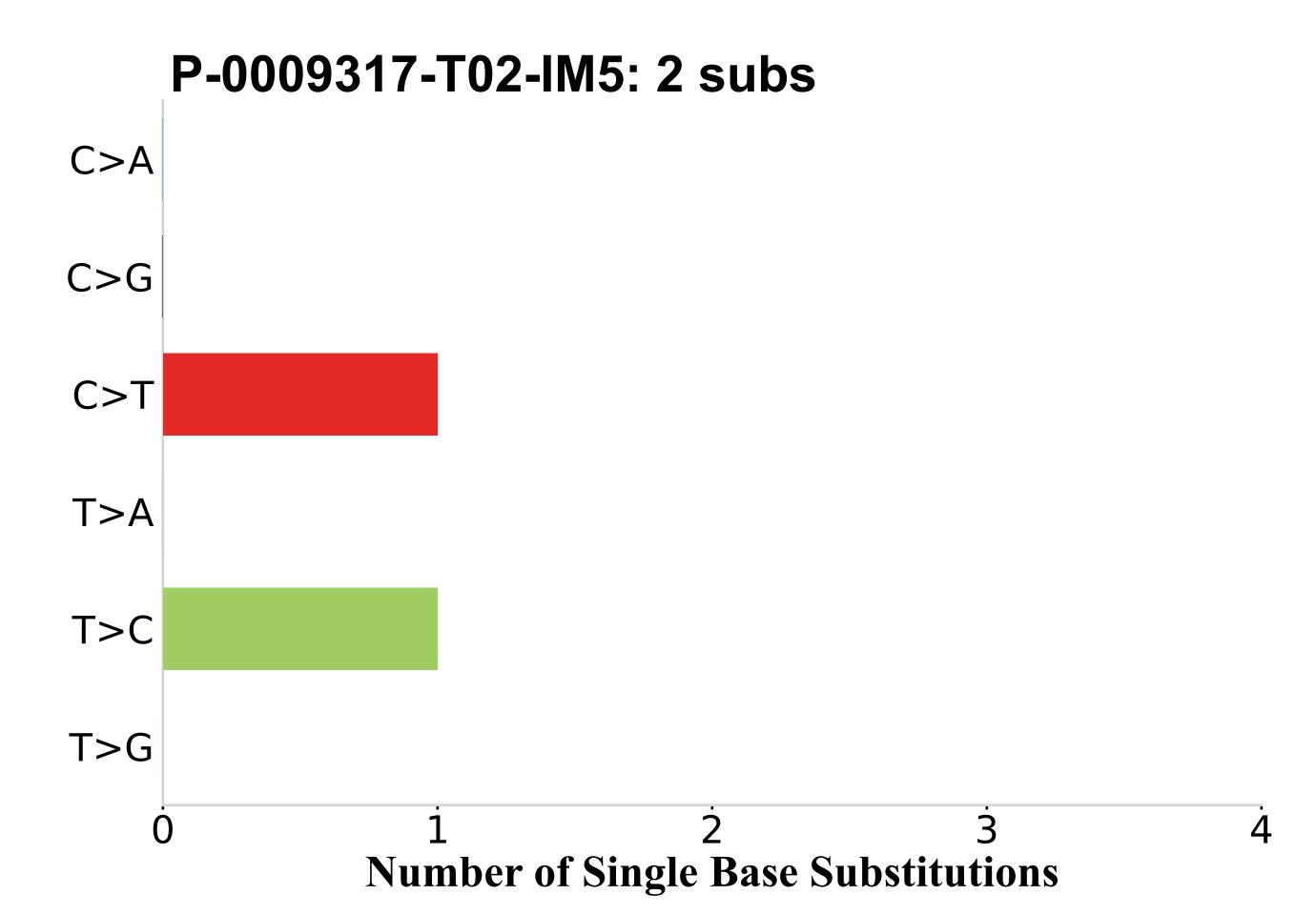


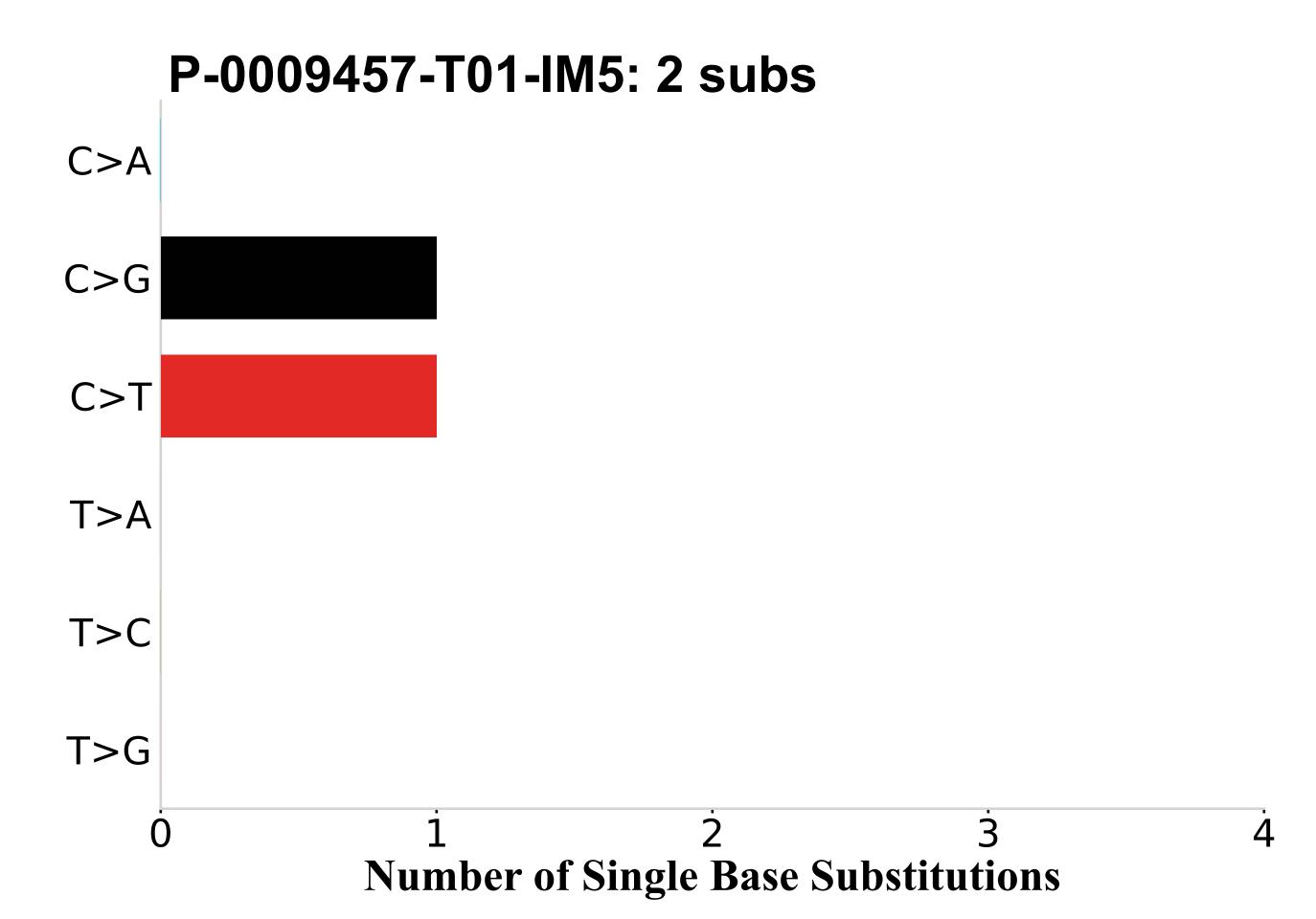




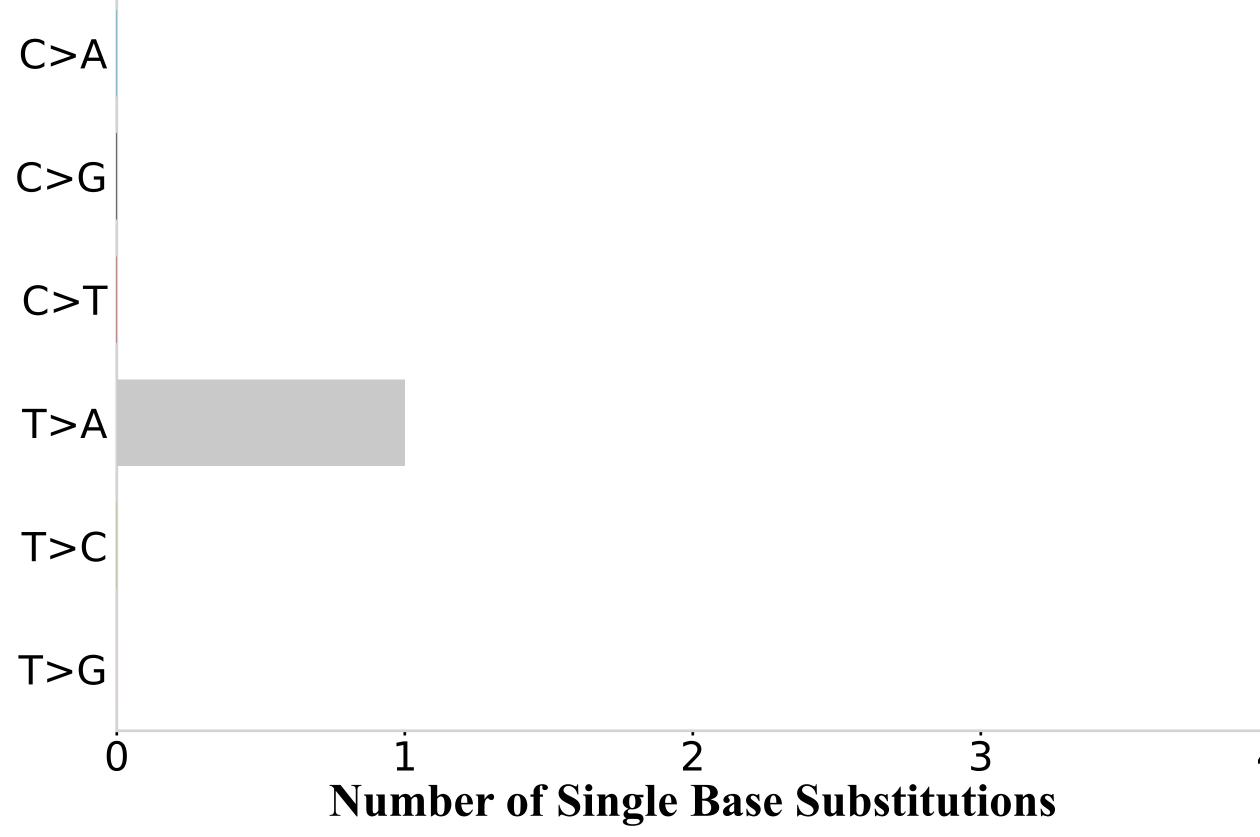


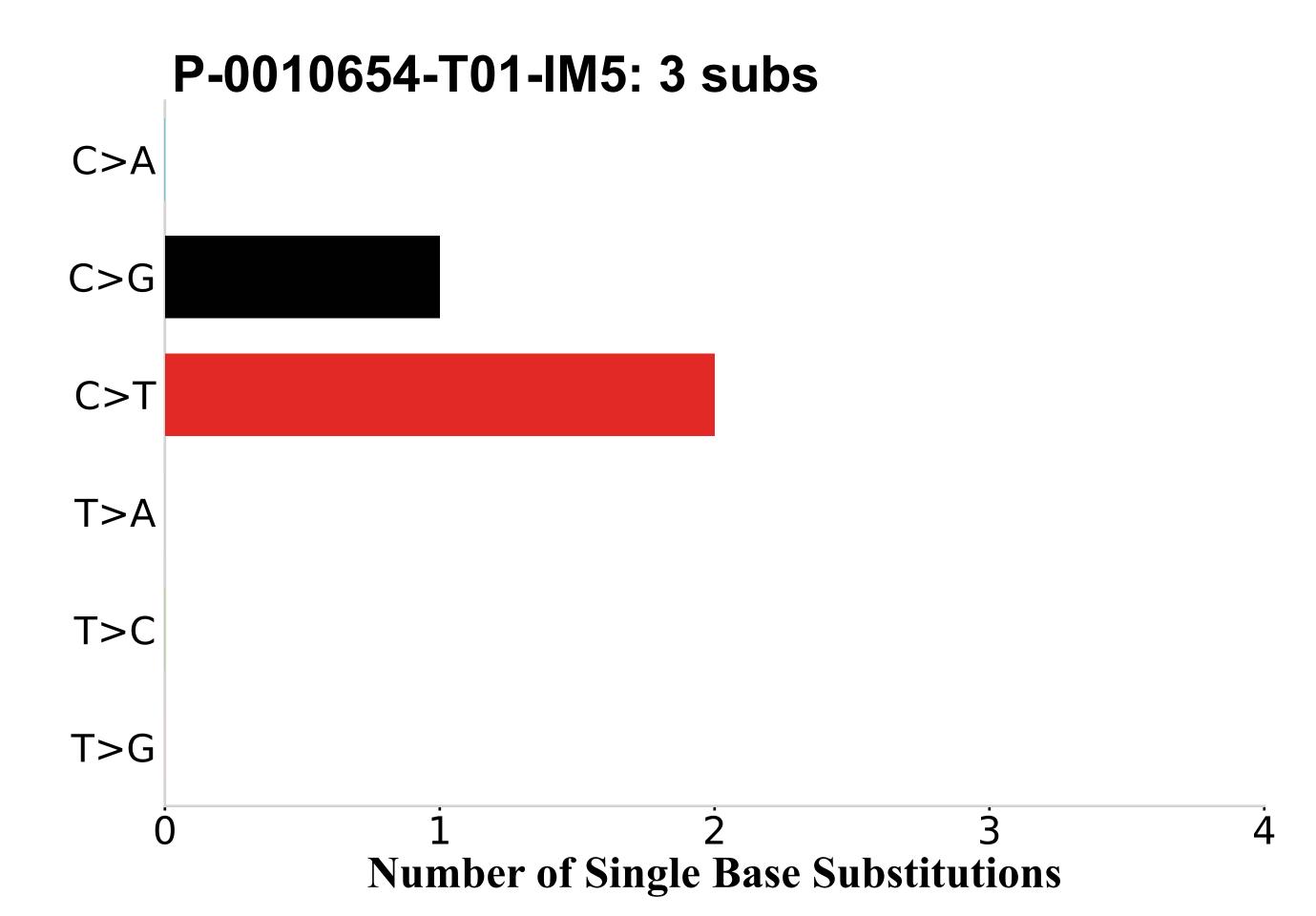


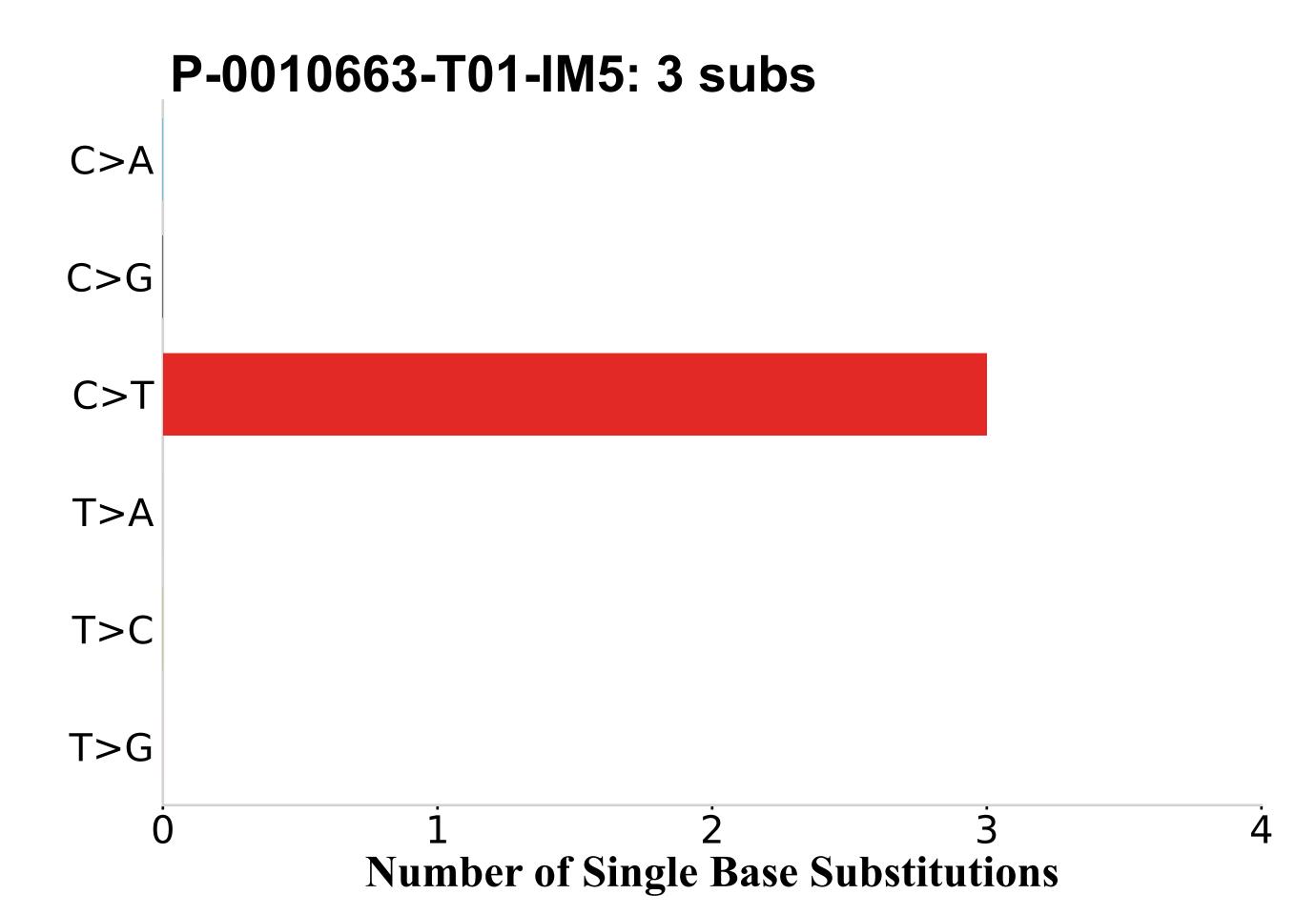


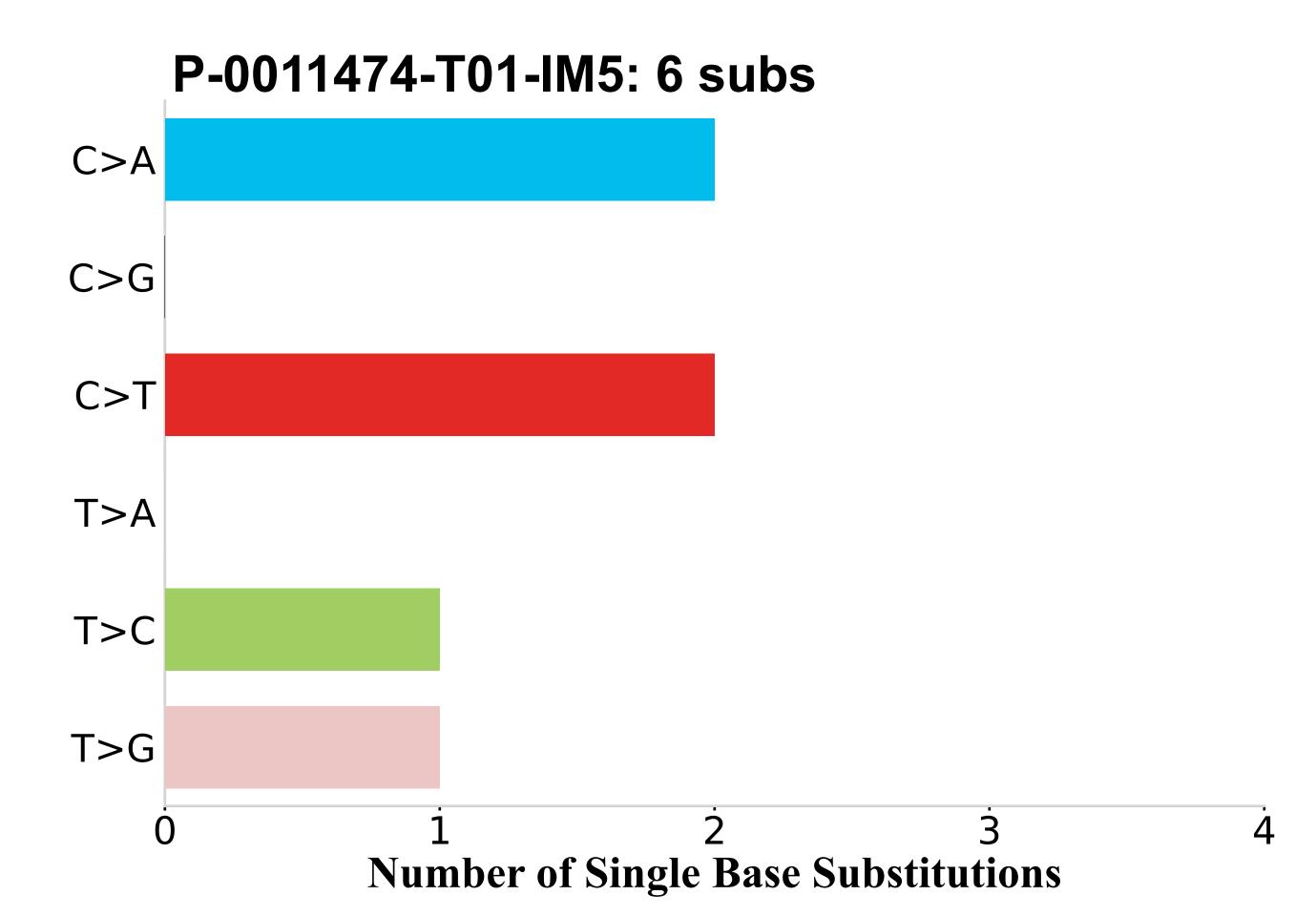


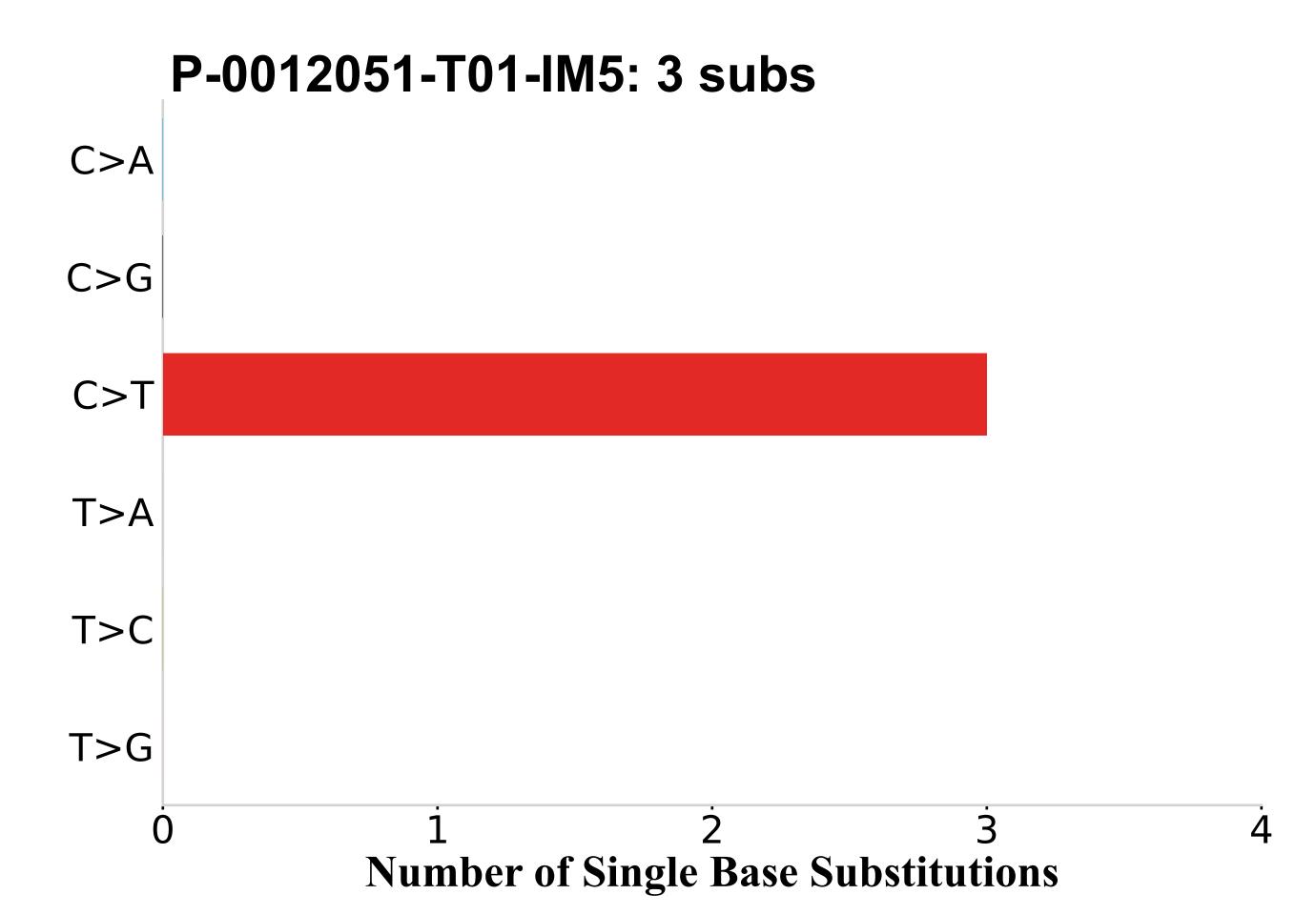
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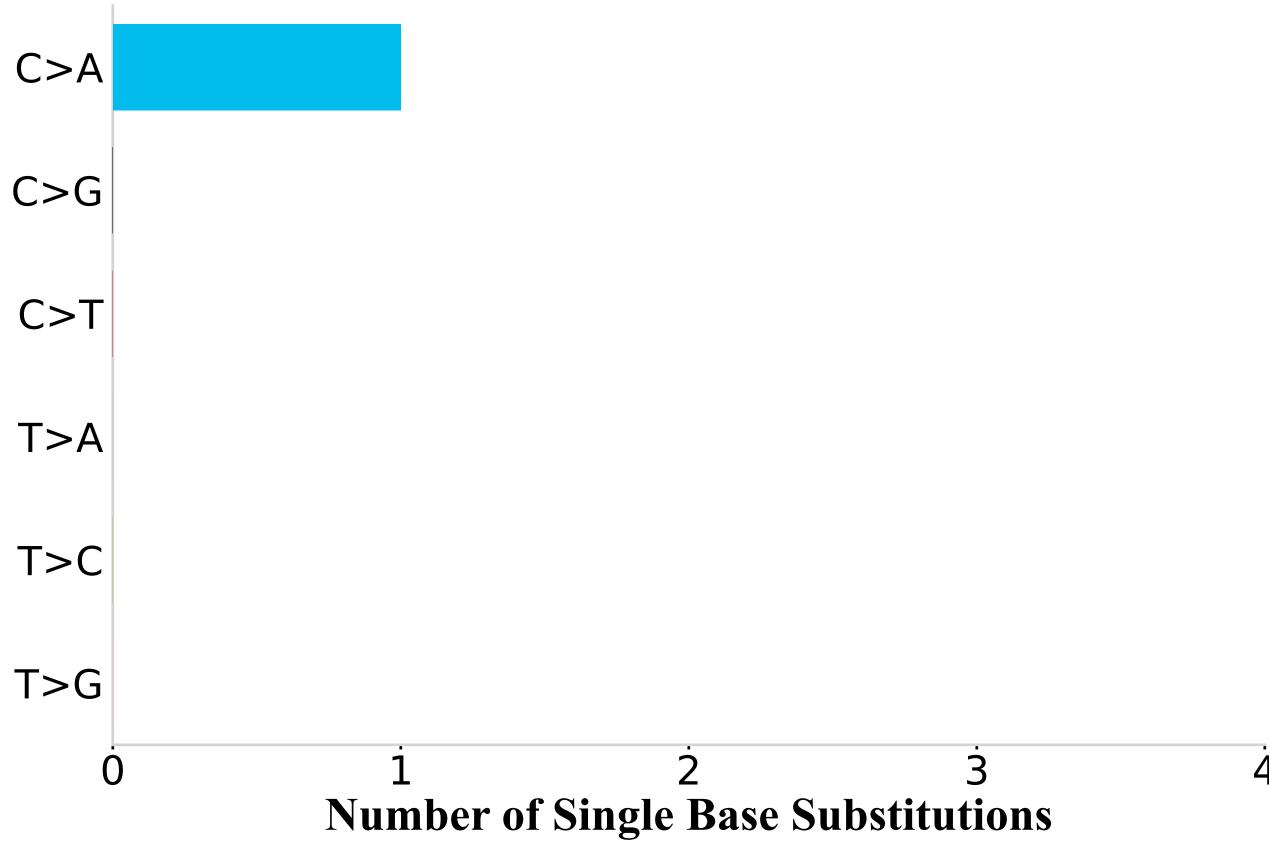




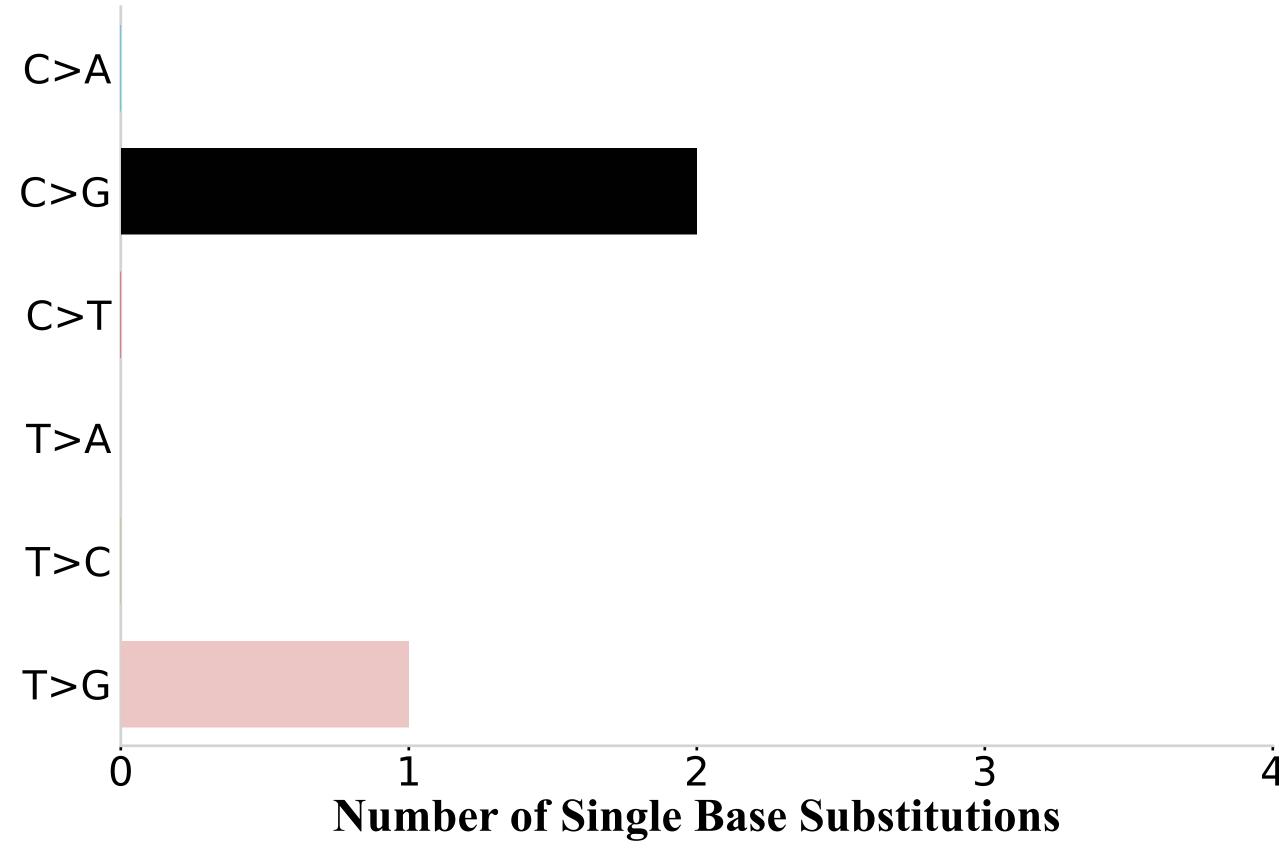




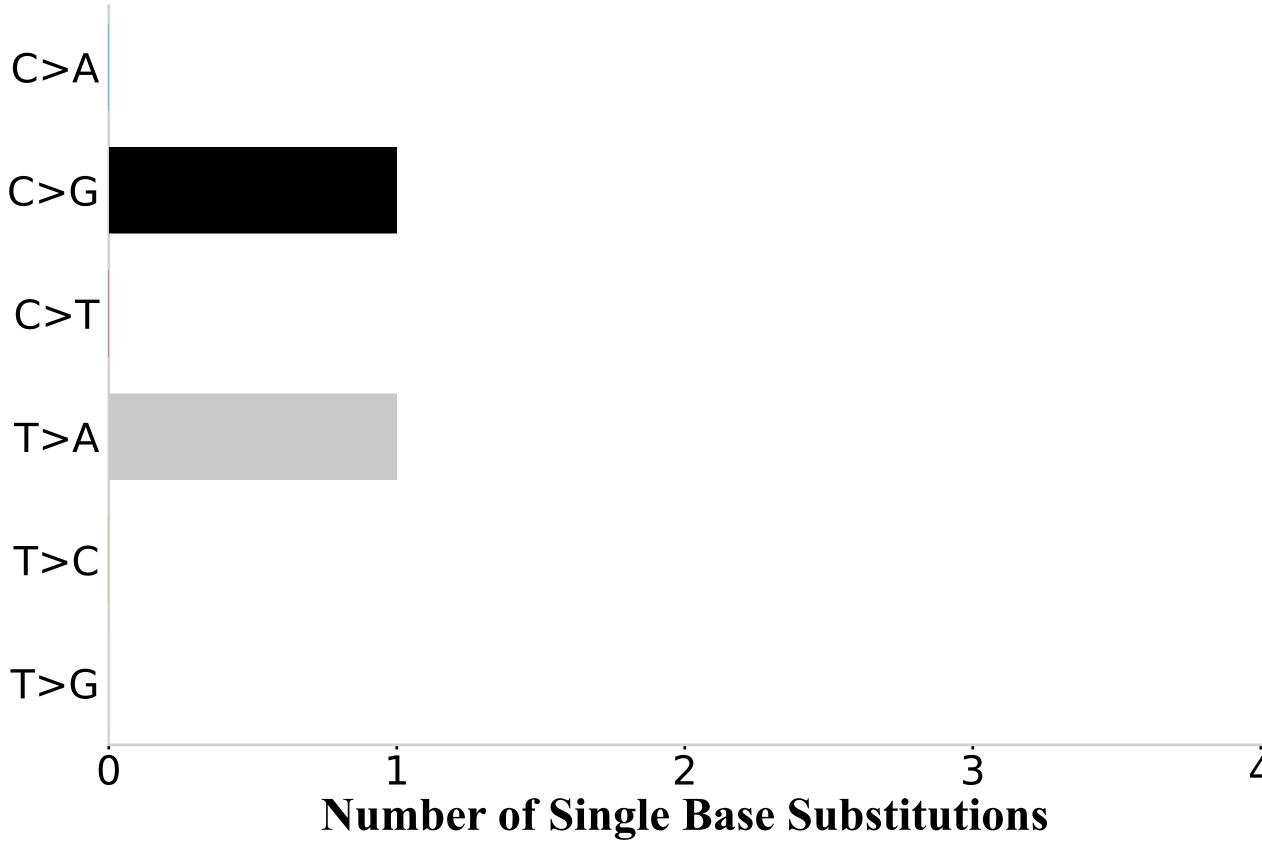


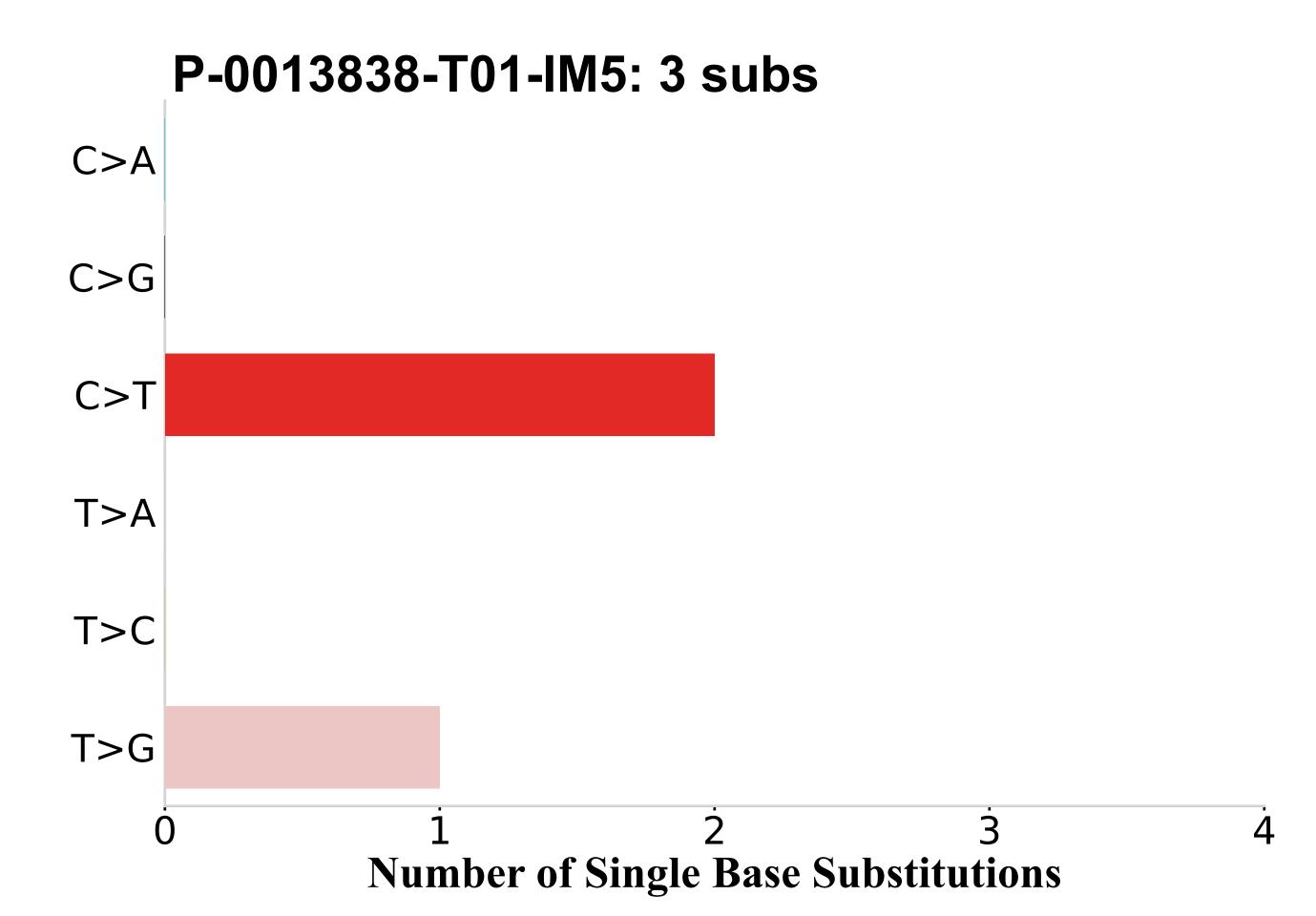


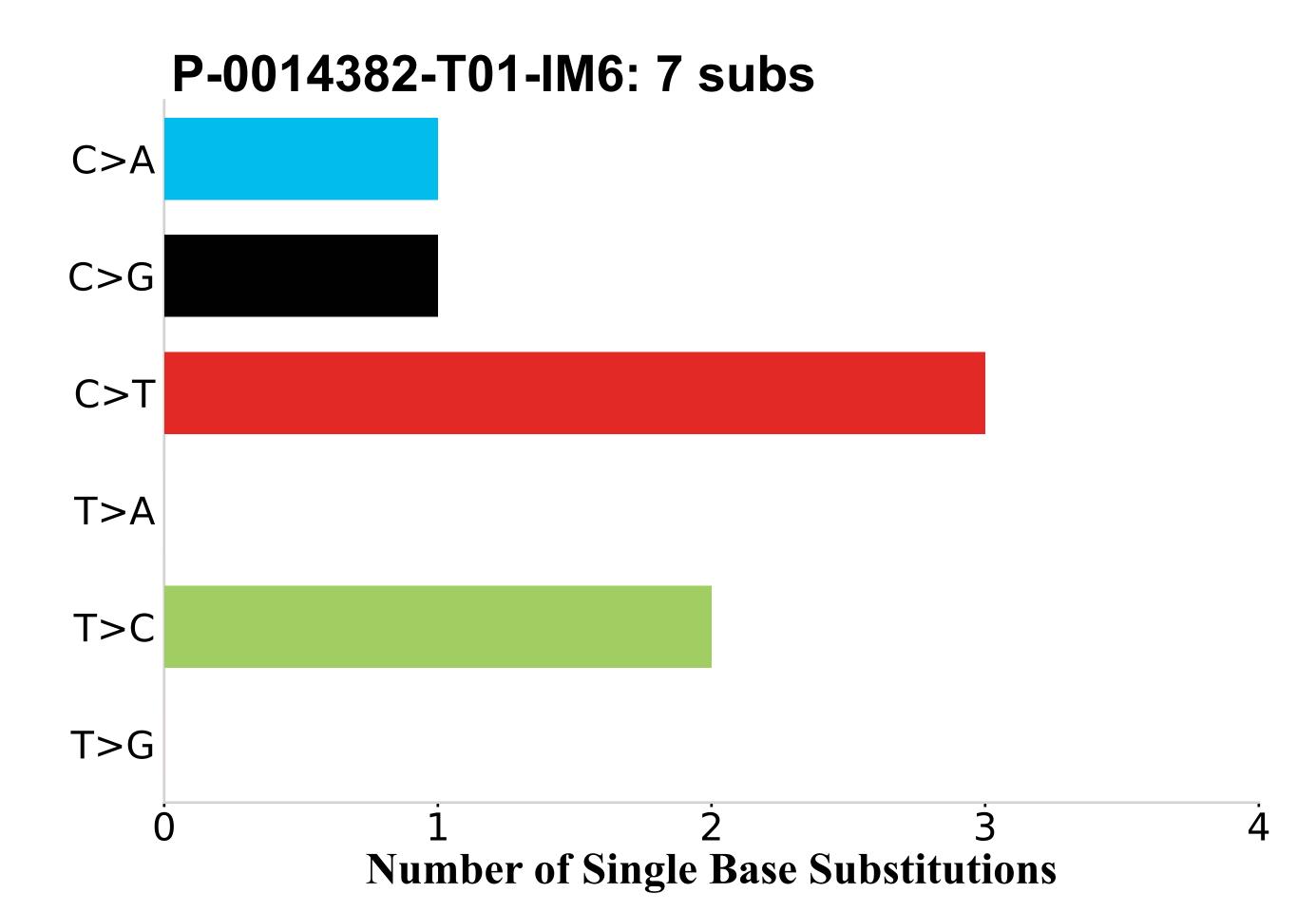




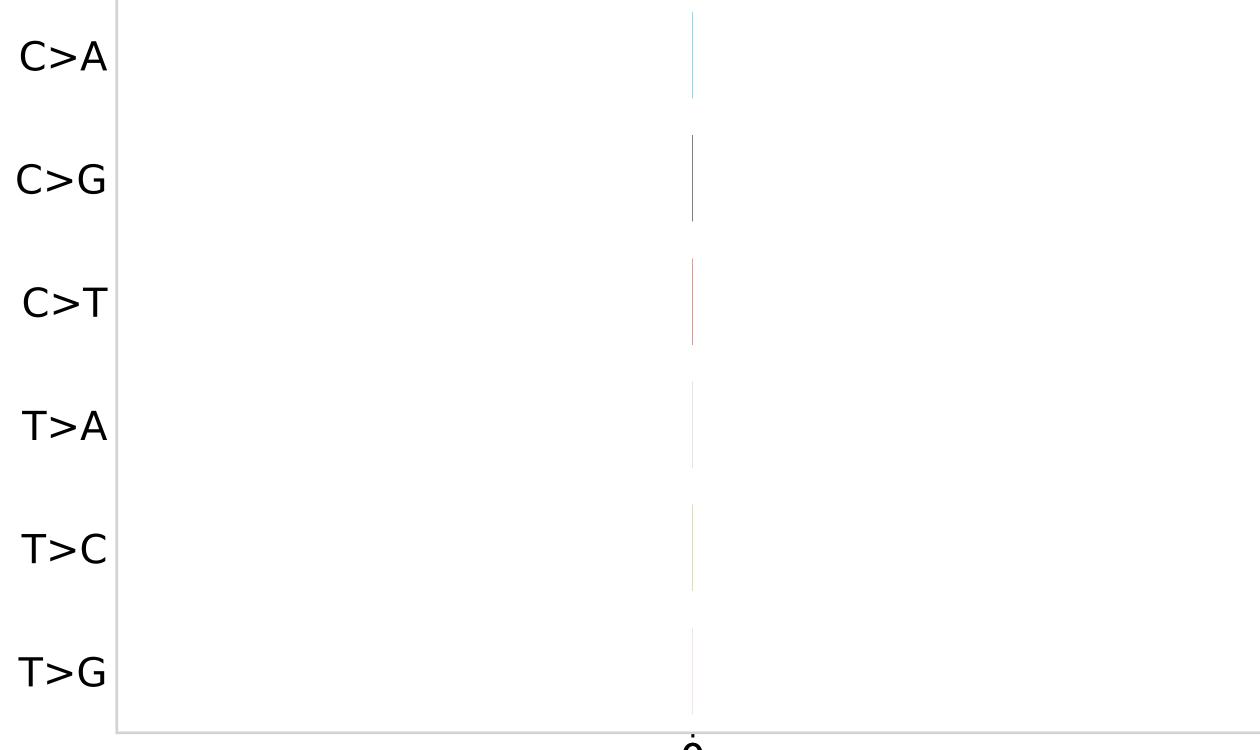






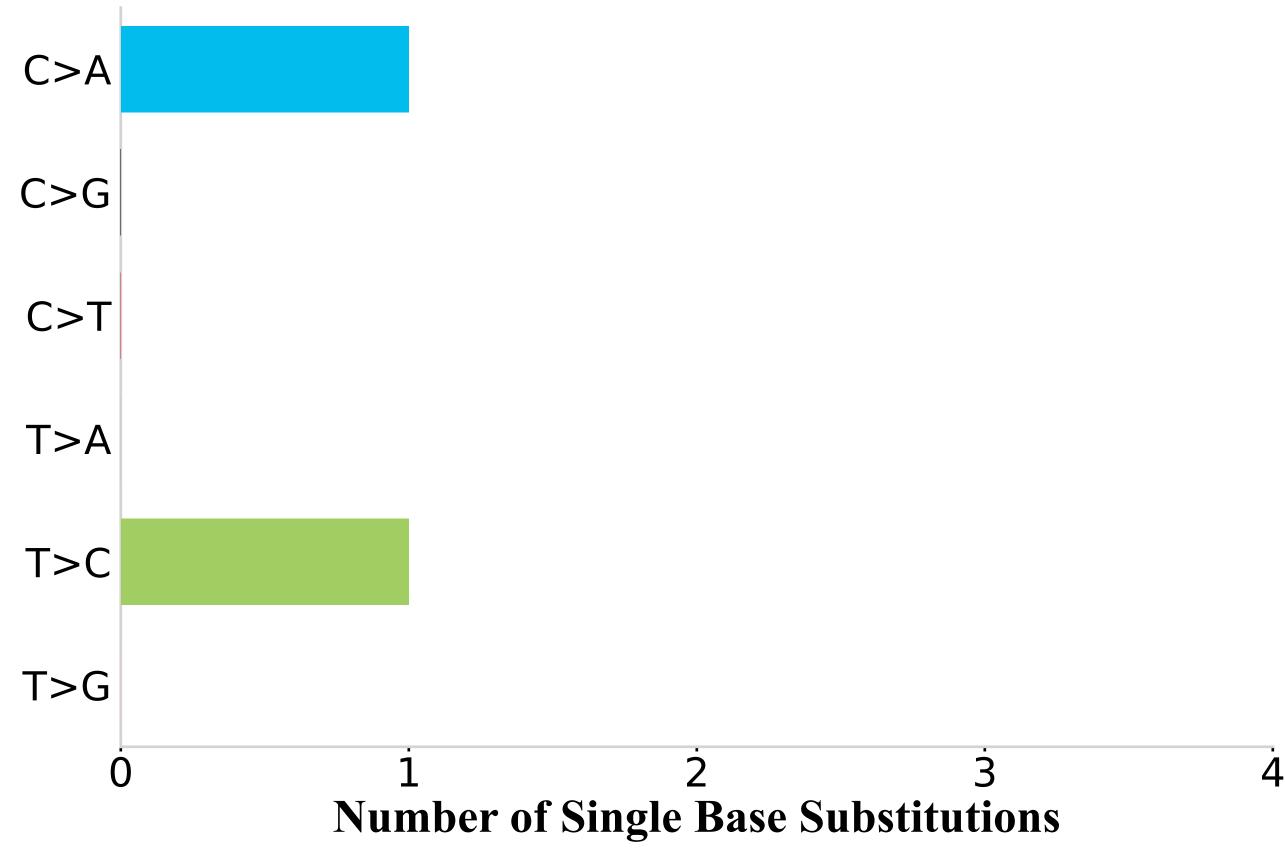


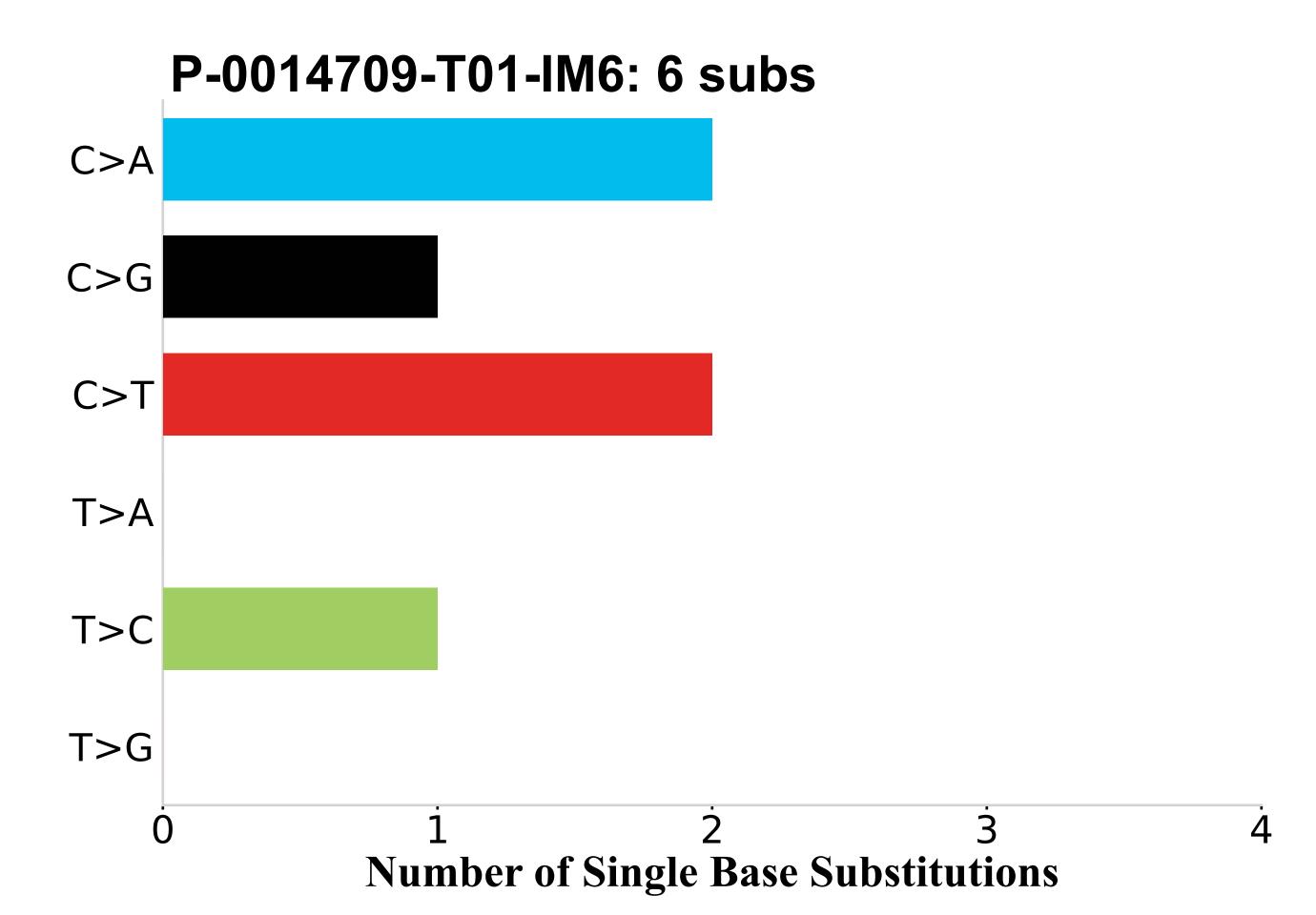
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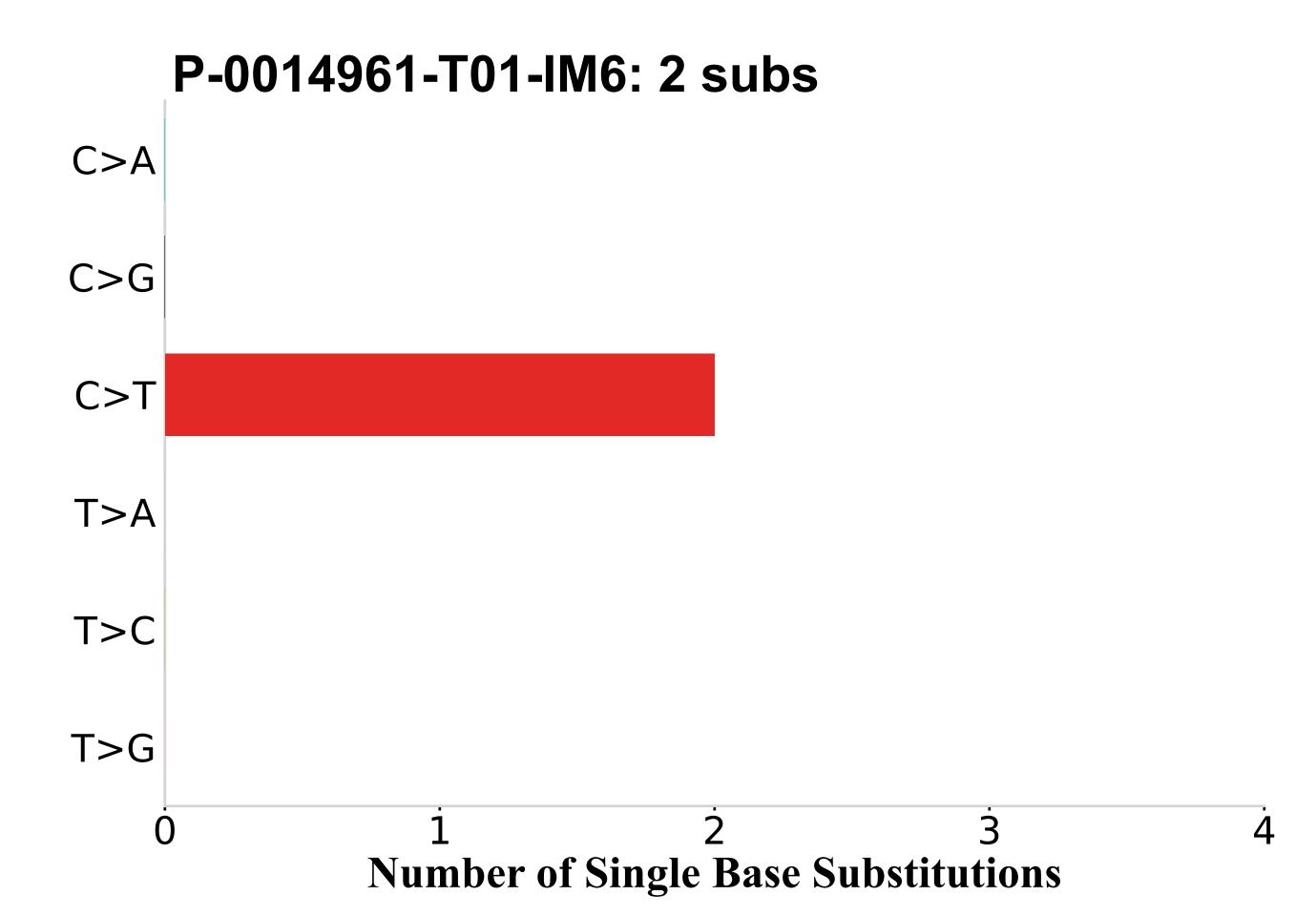


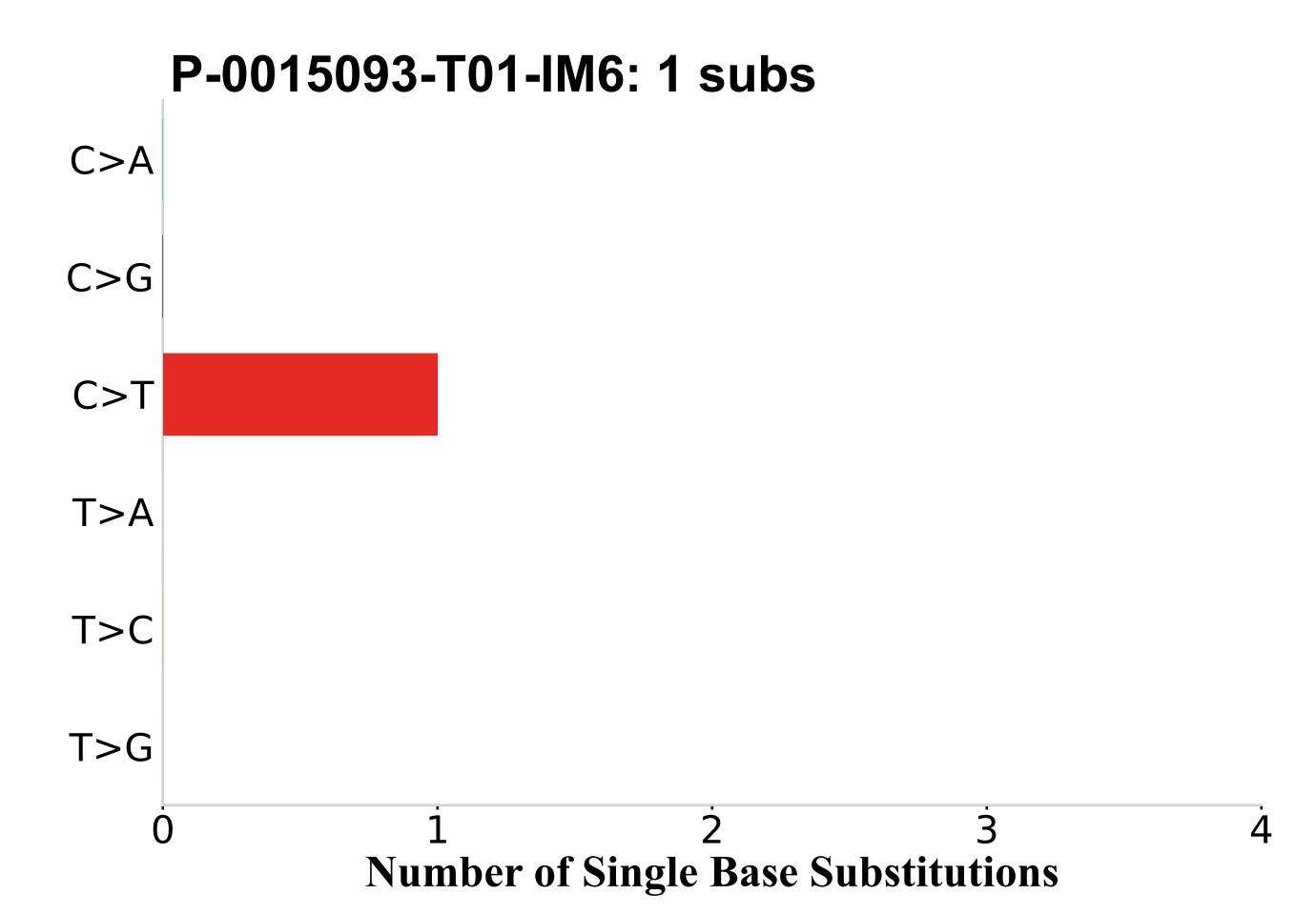
Number of Single Base Substitutions

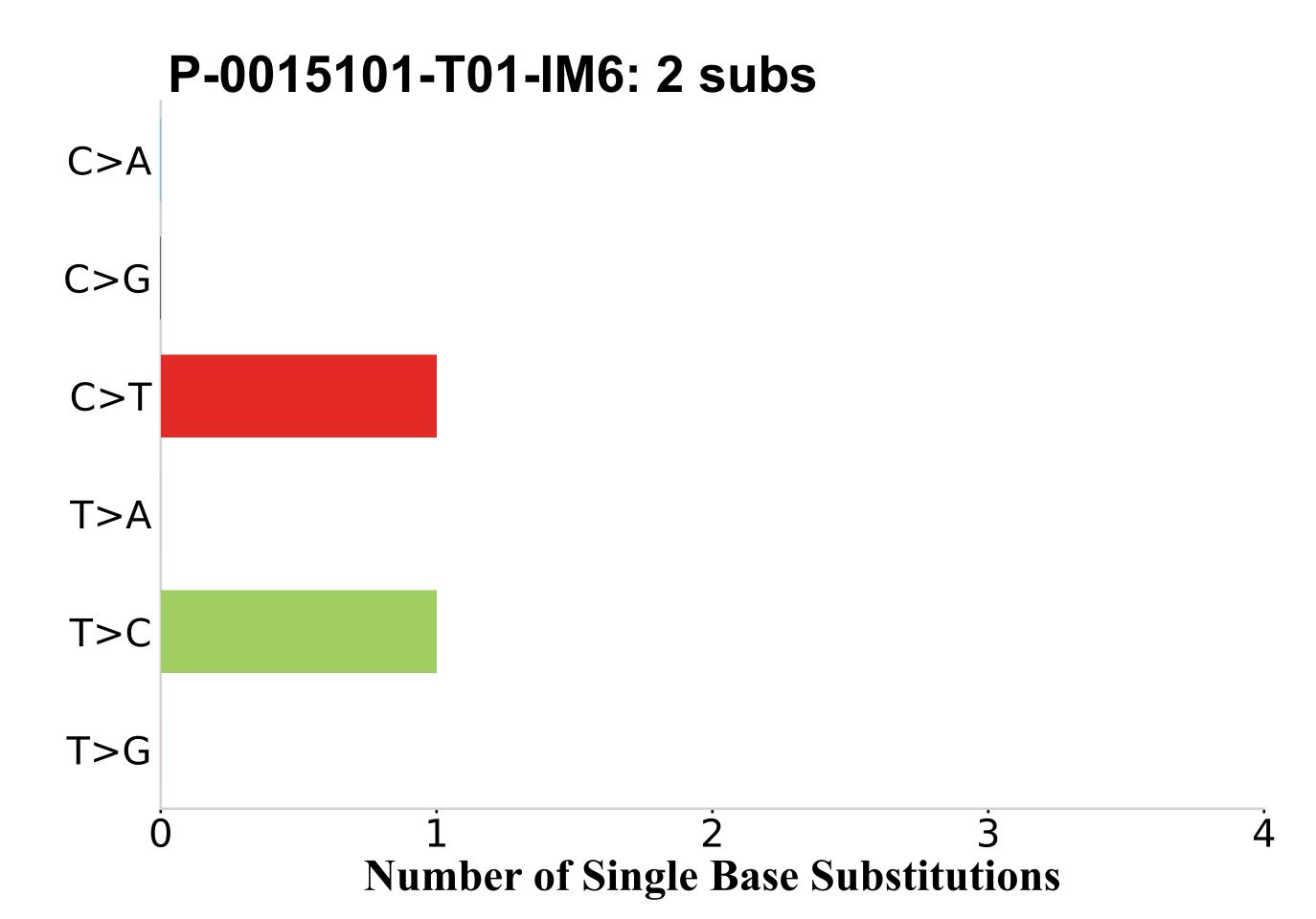
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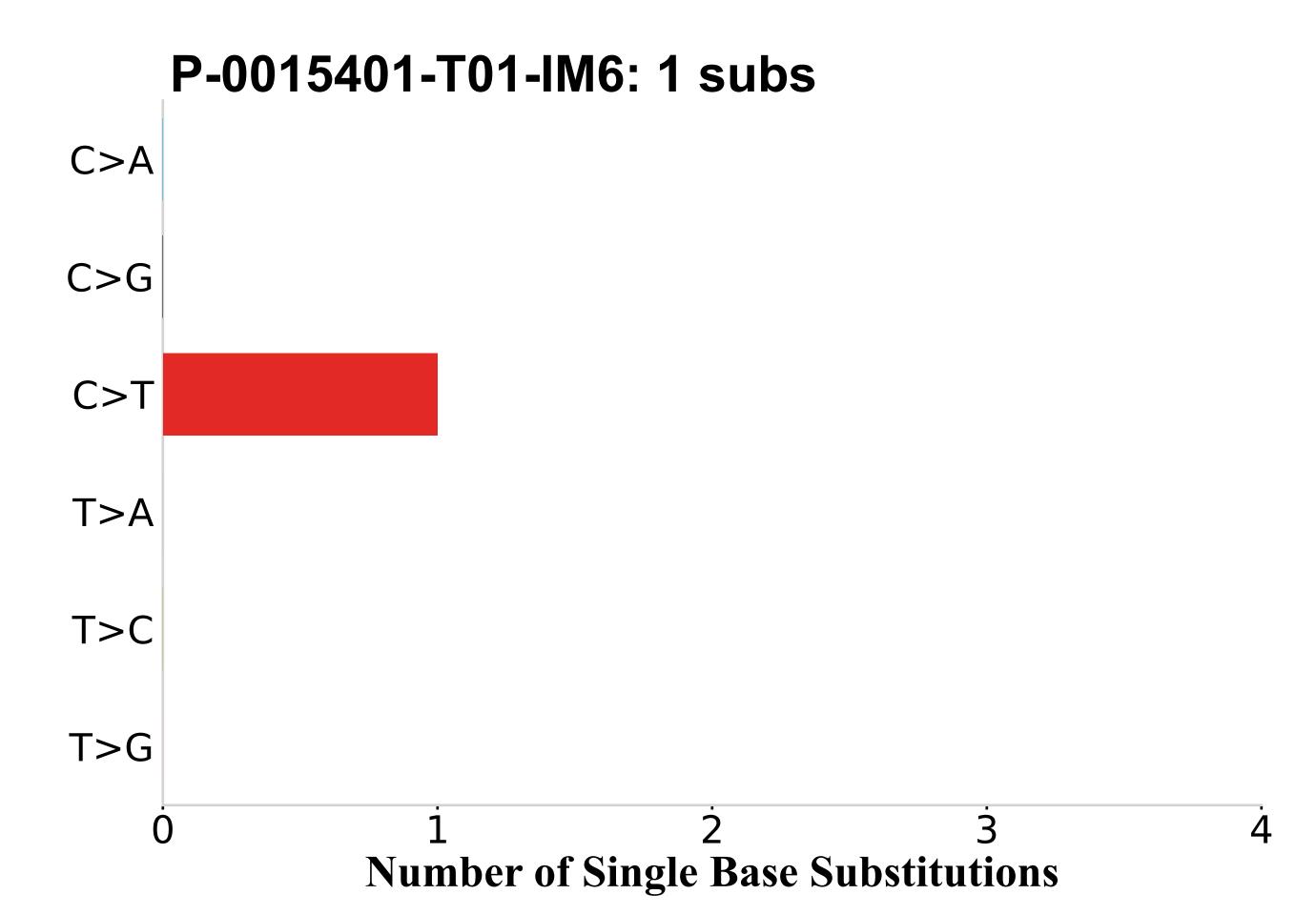








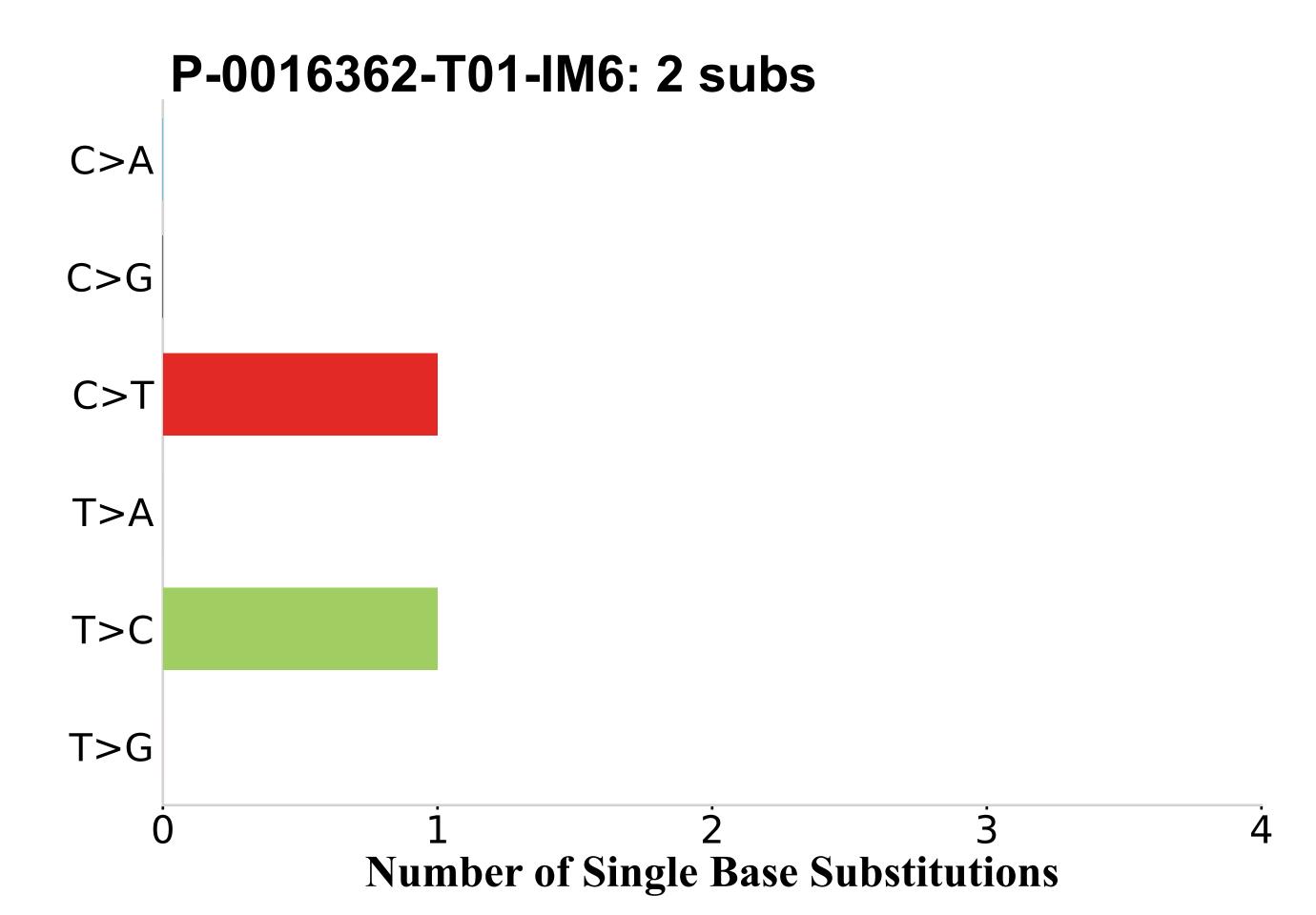




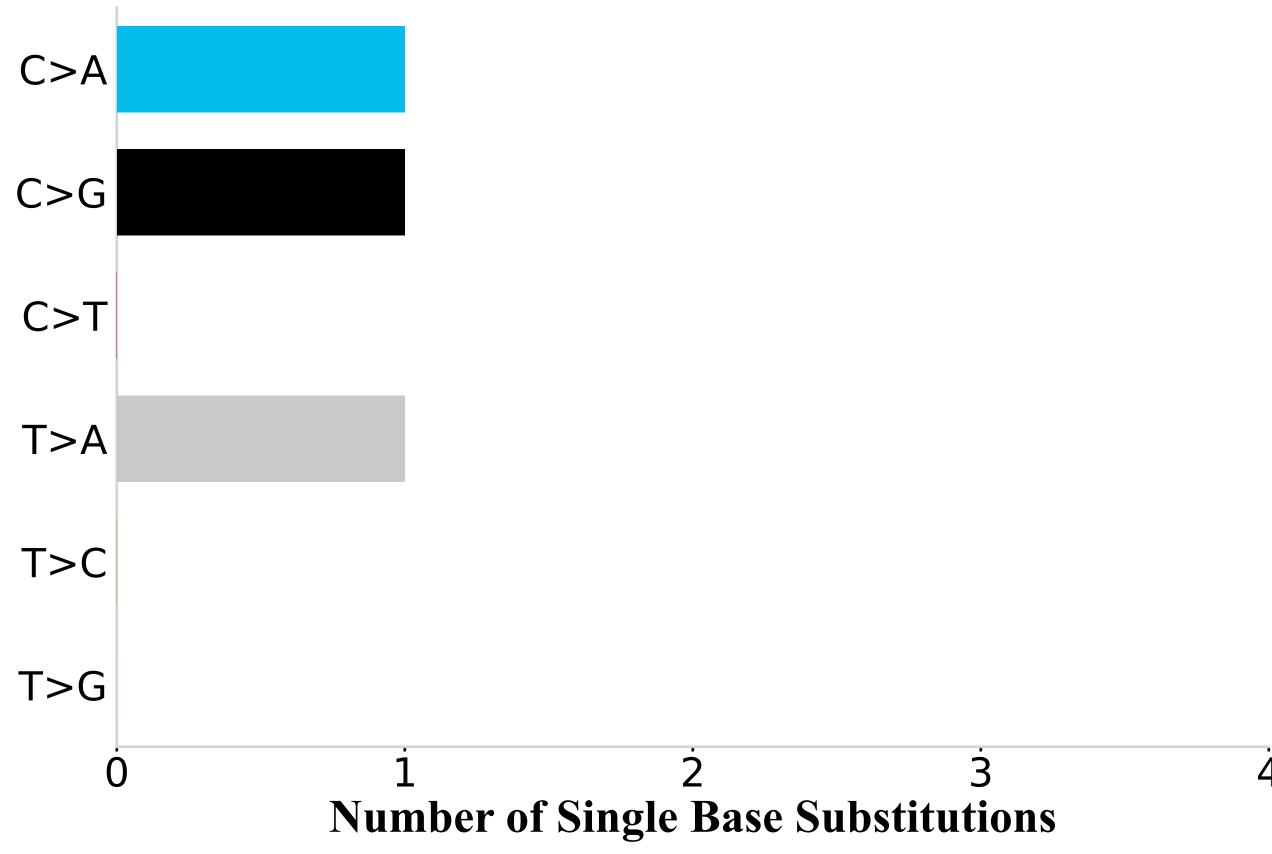
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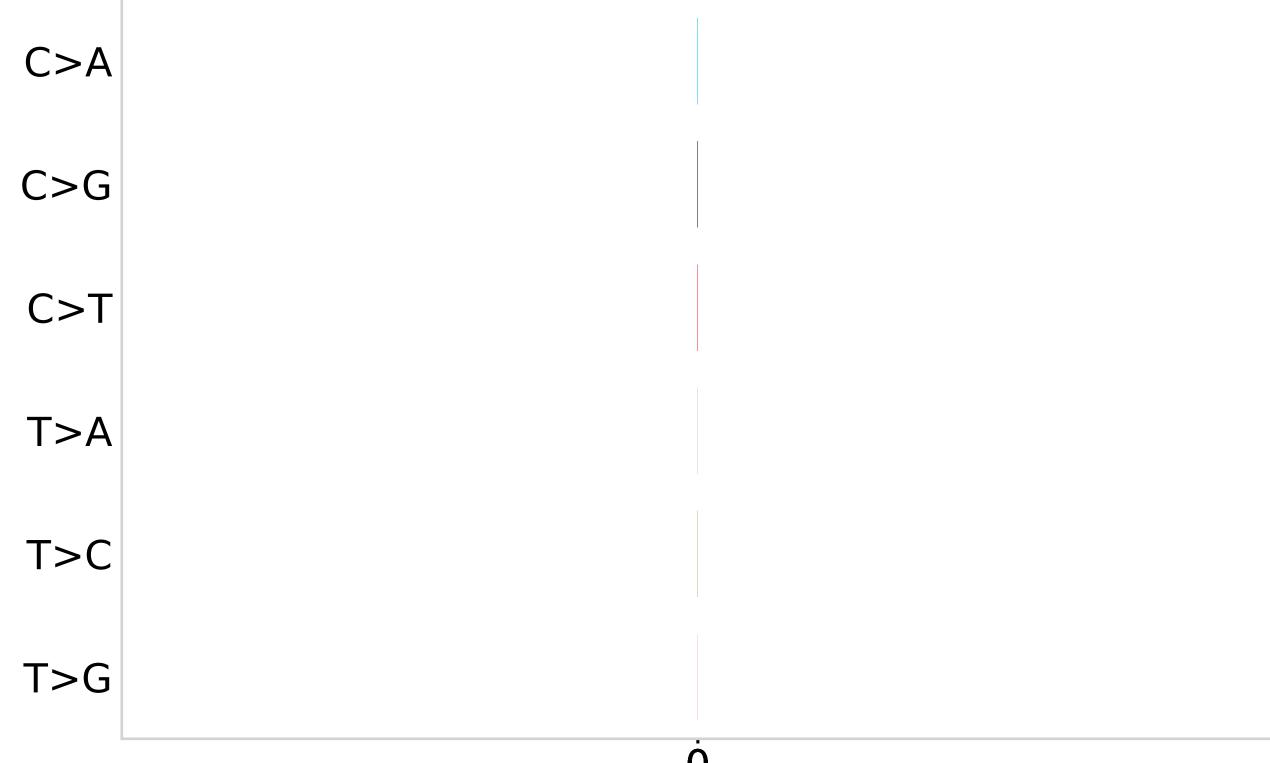
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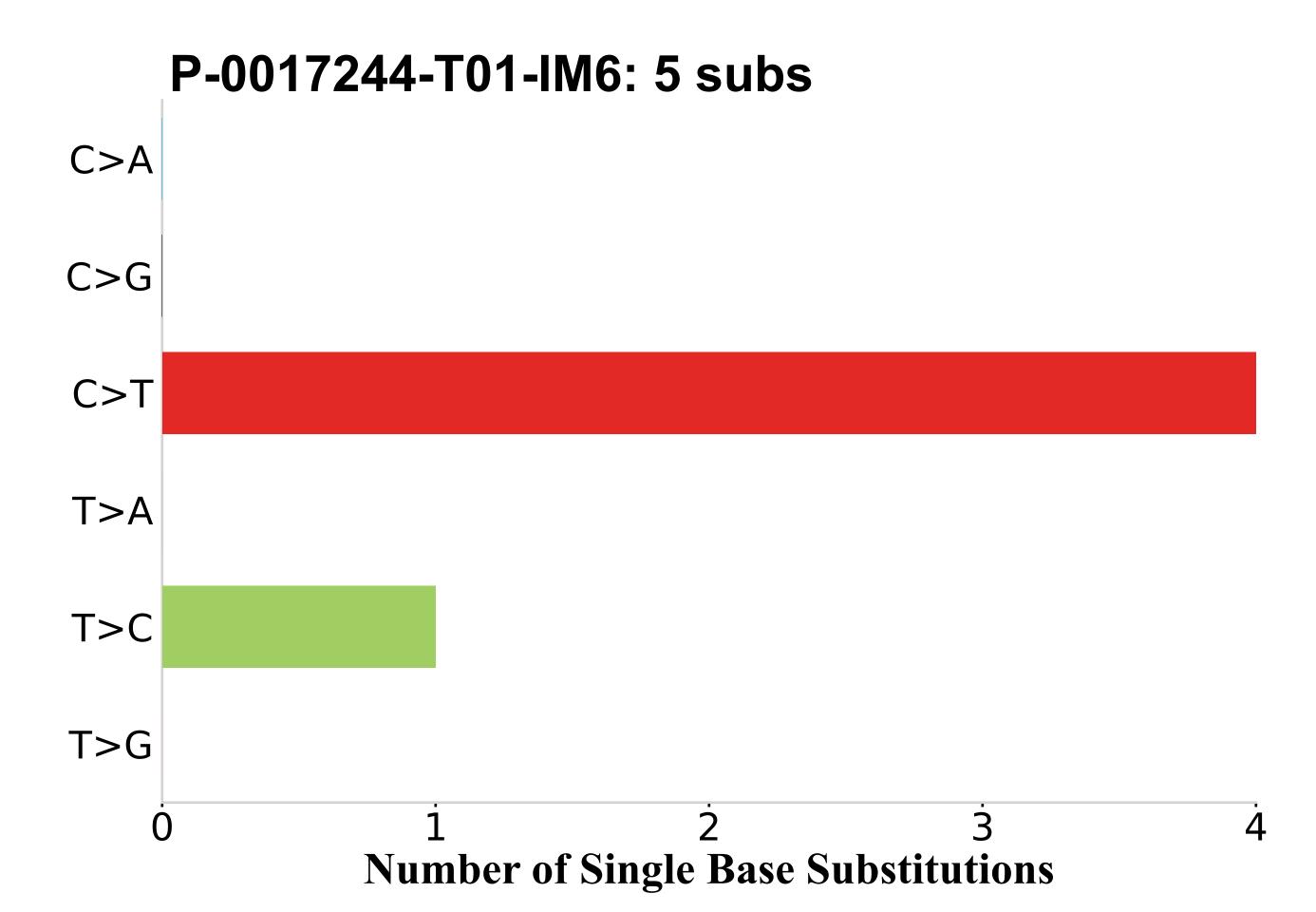
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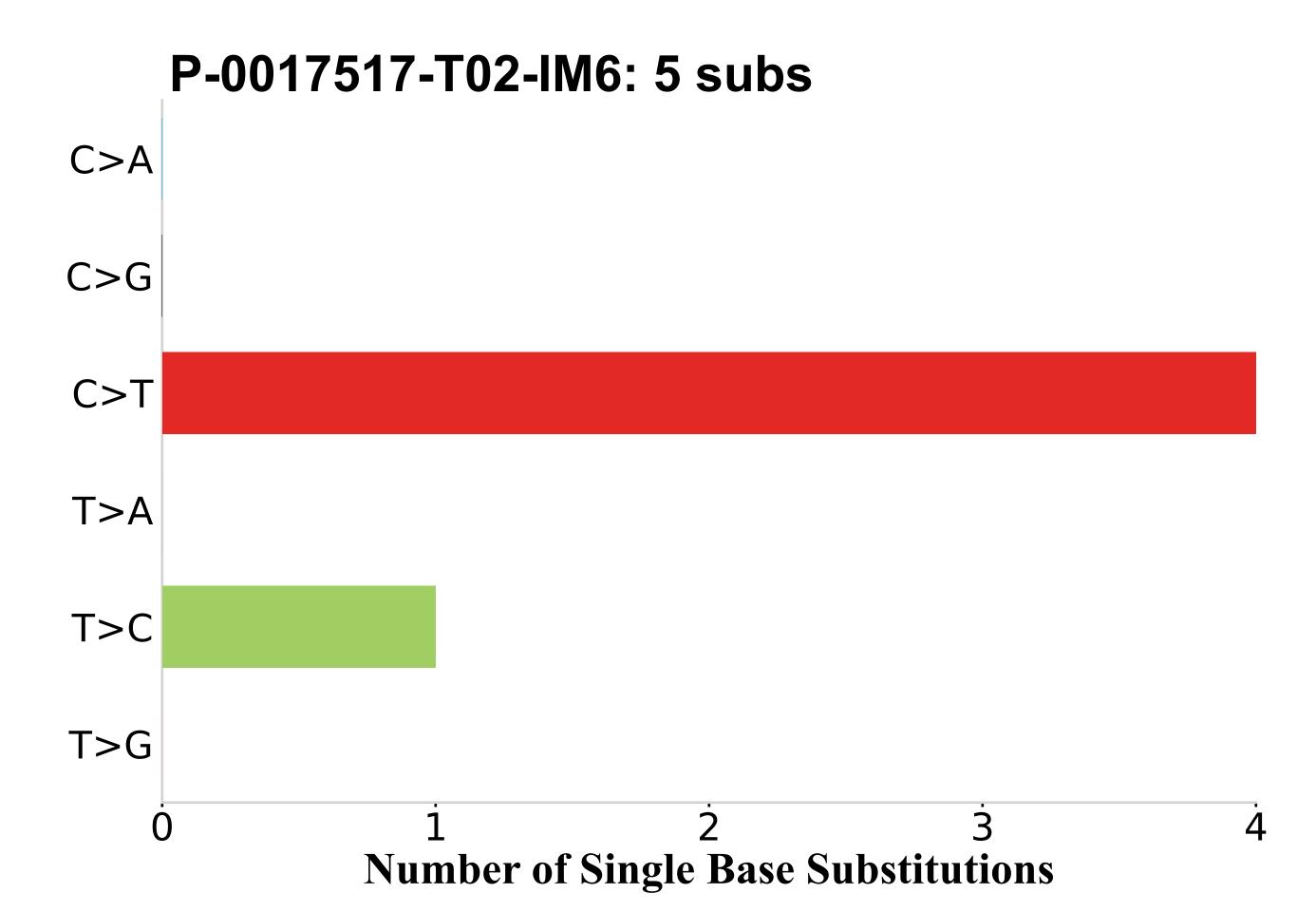


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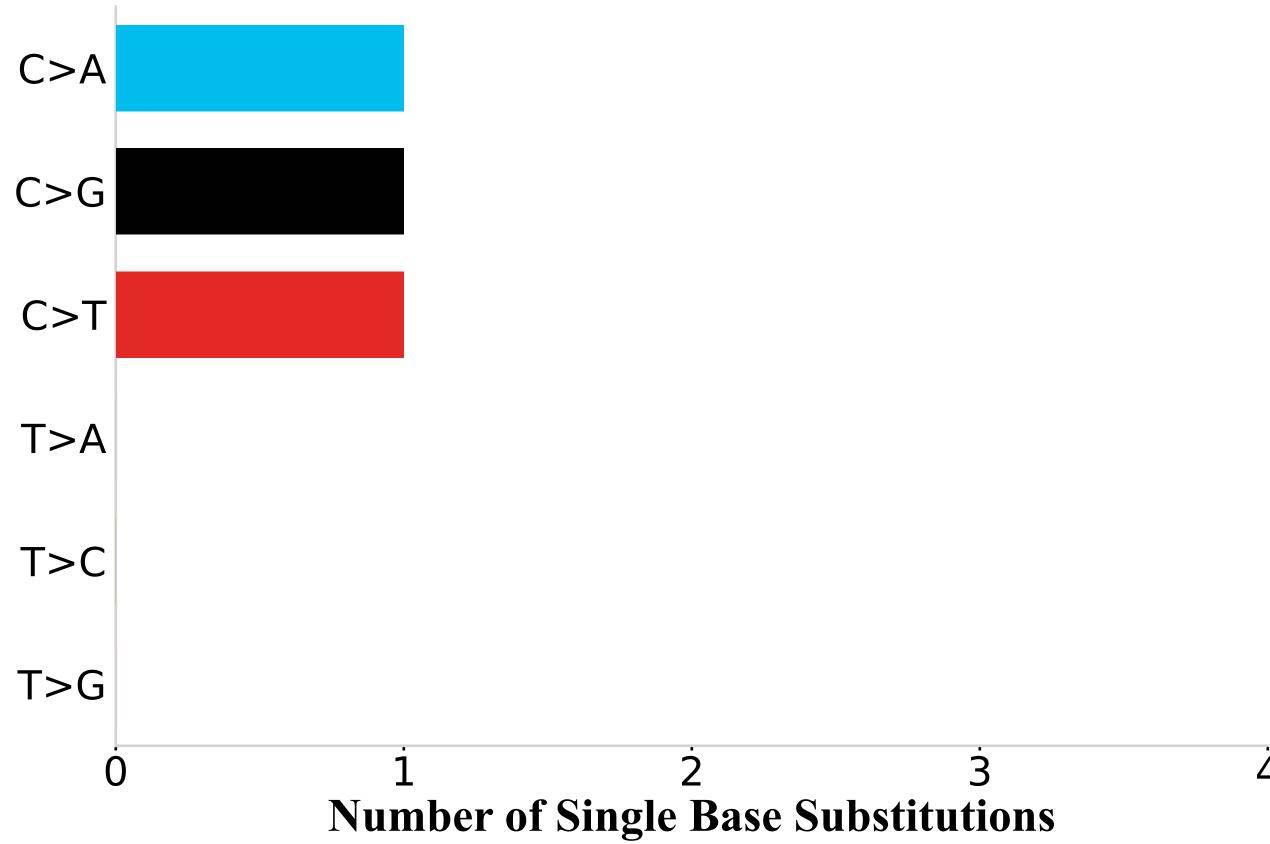


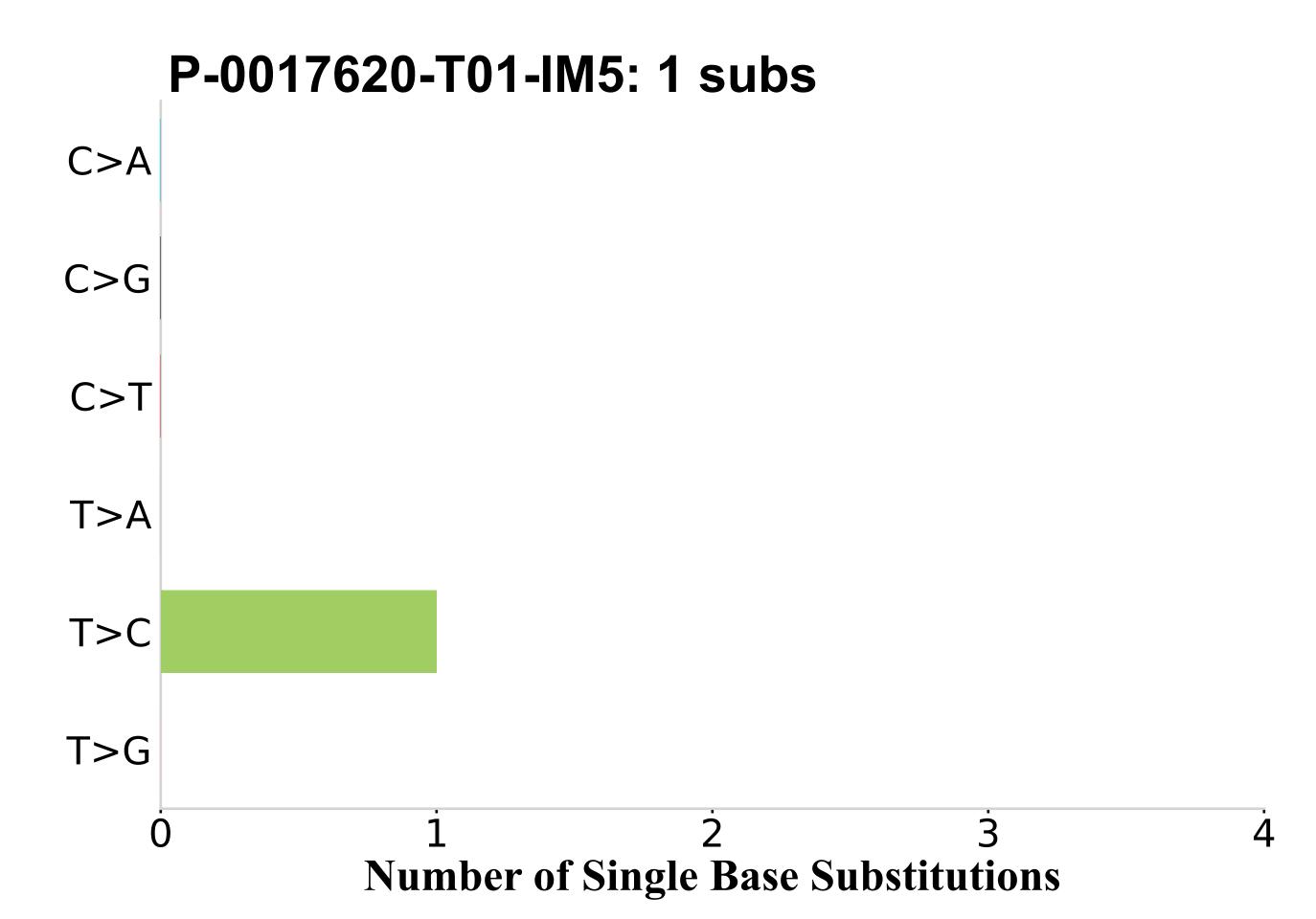
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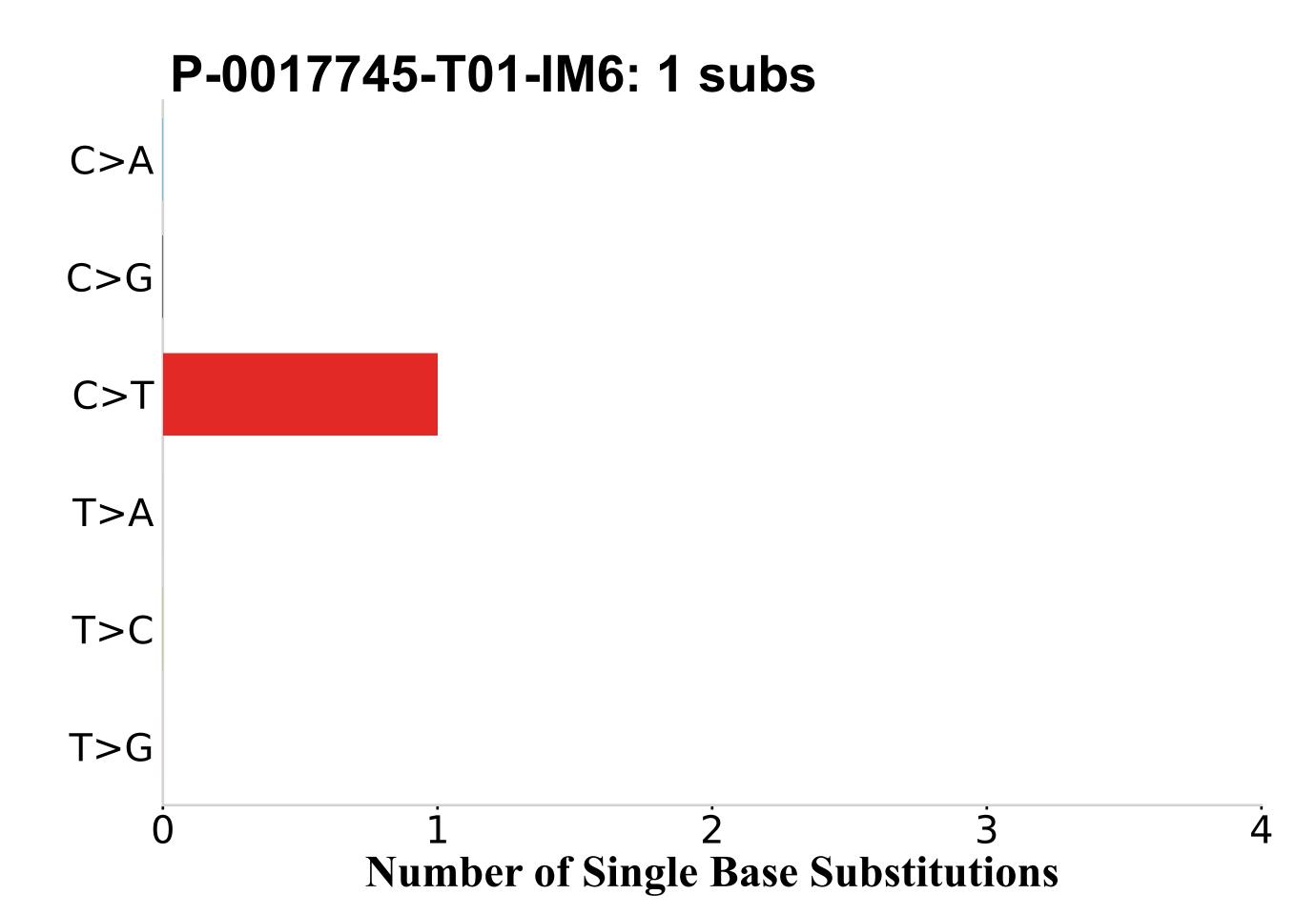


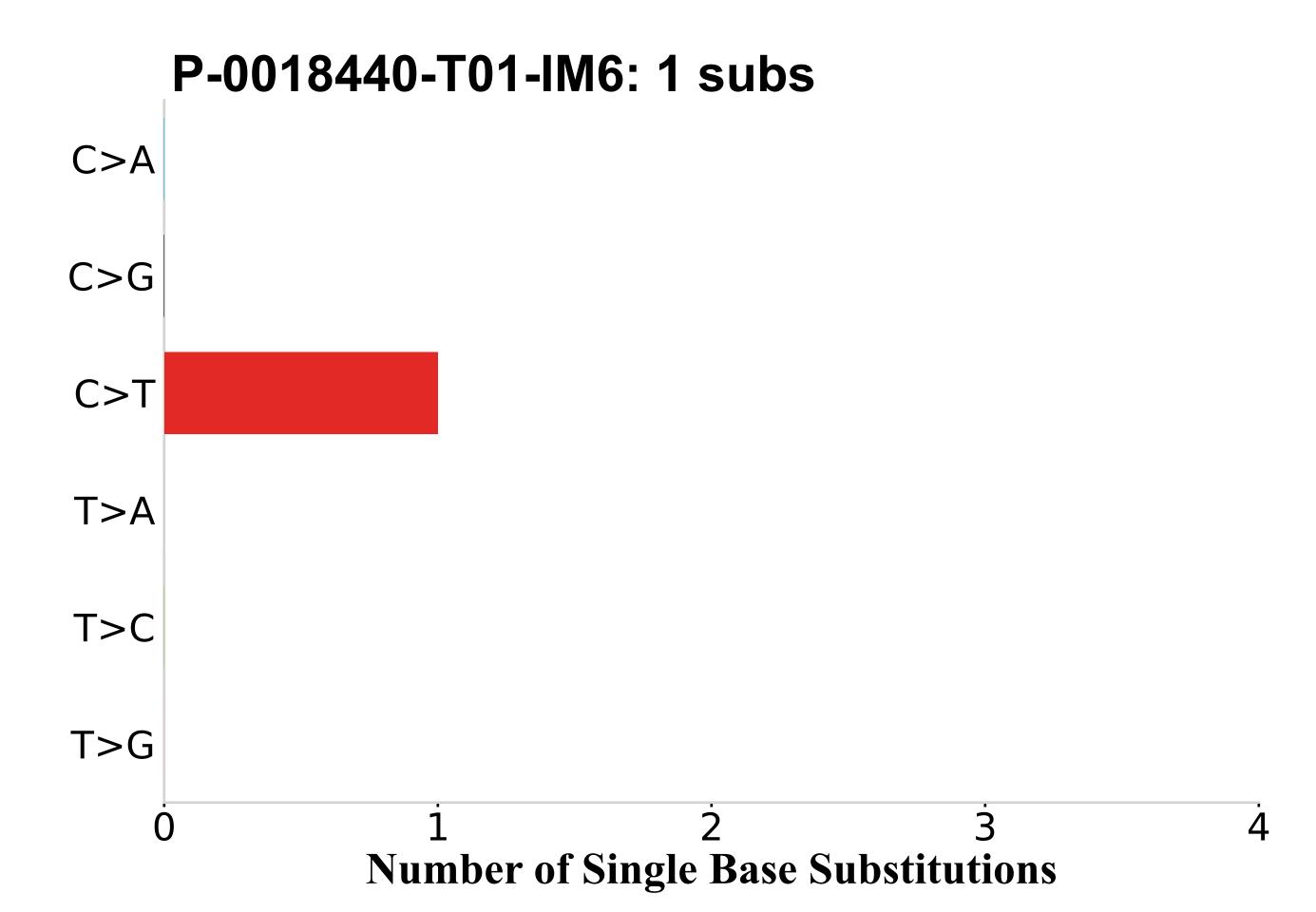


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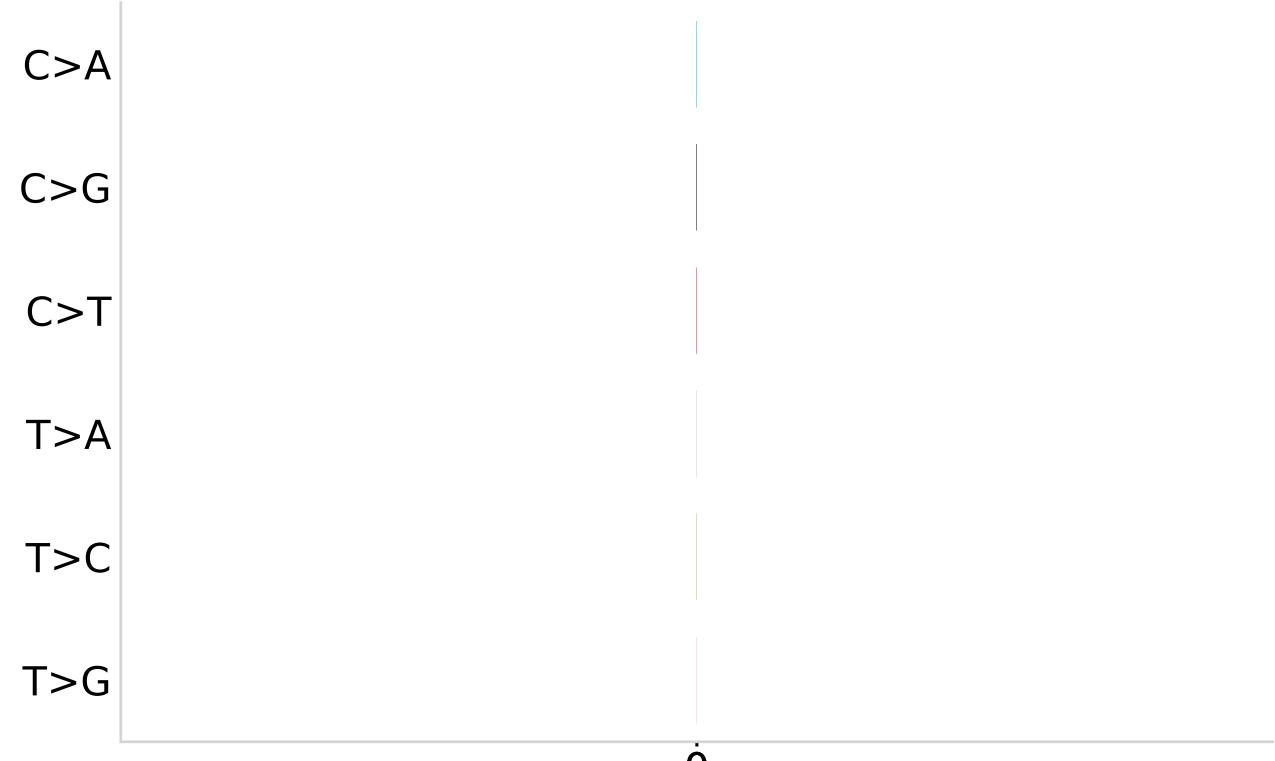




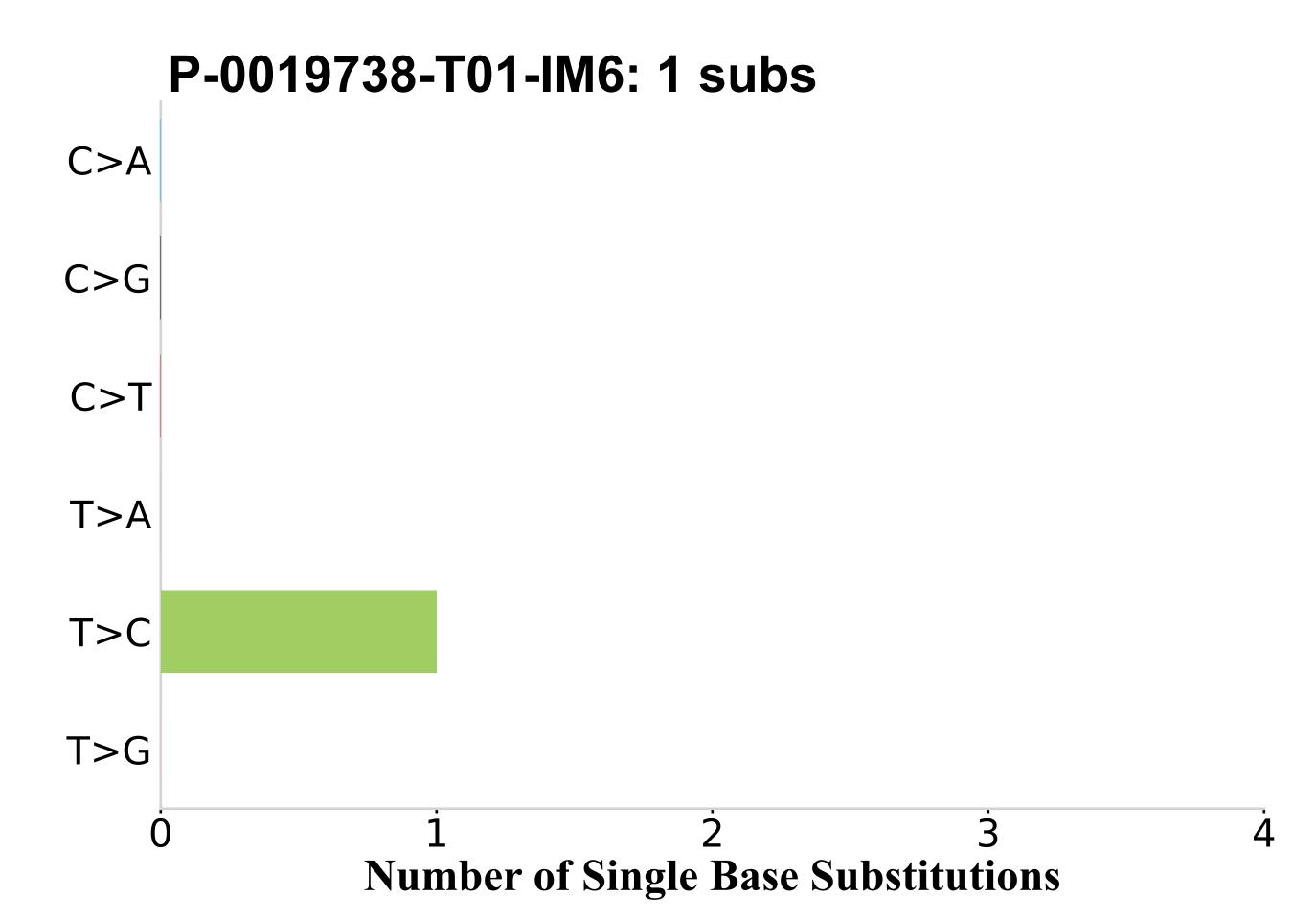




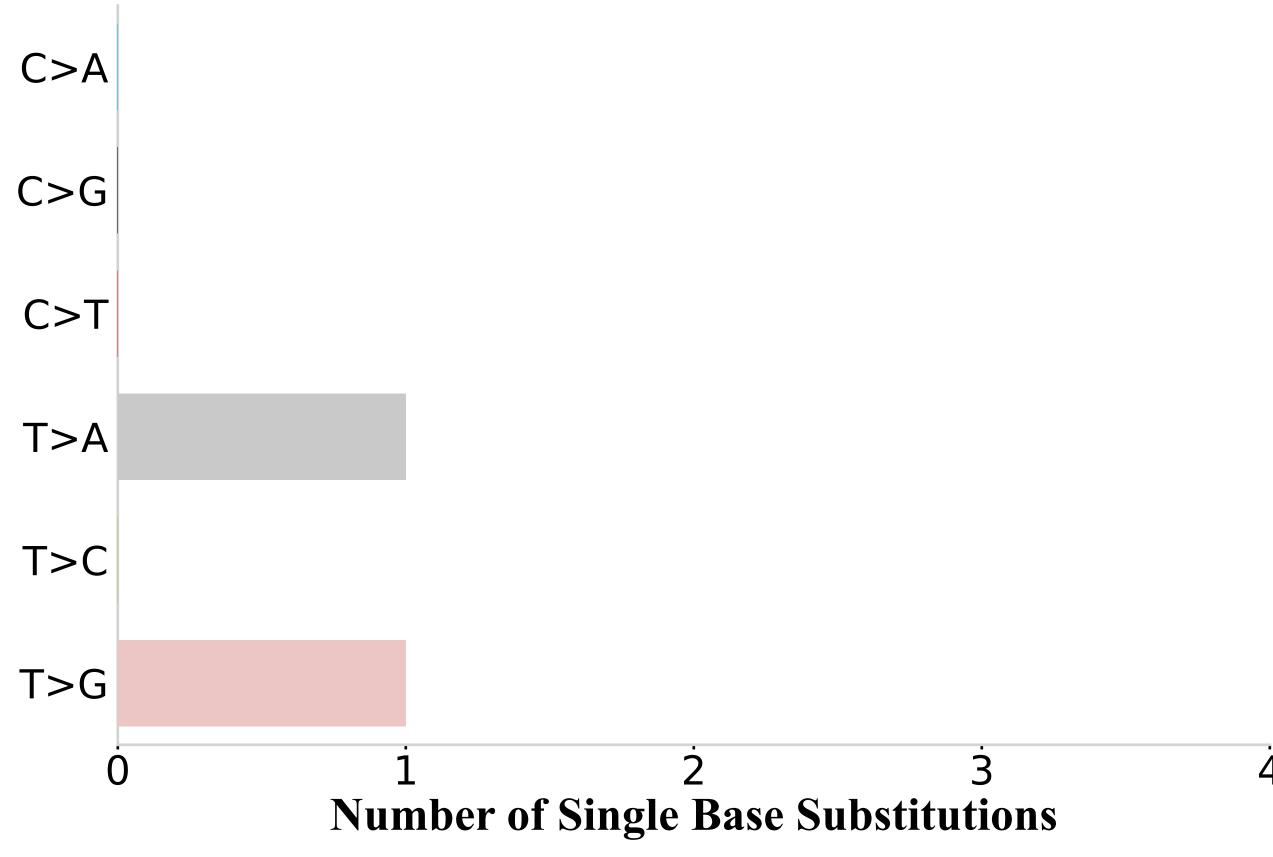
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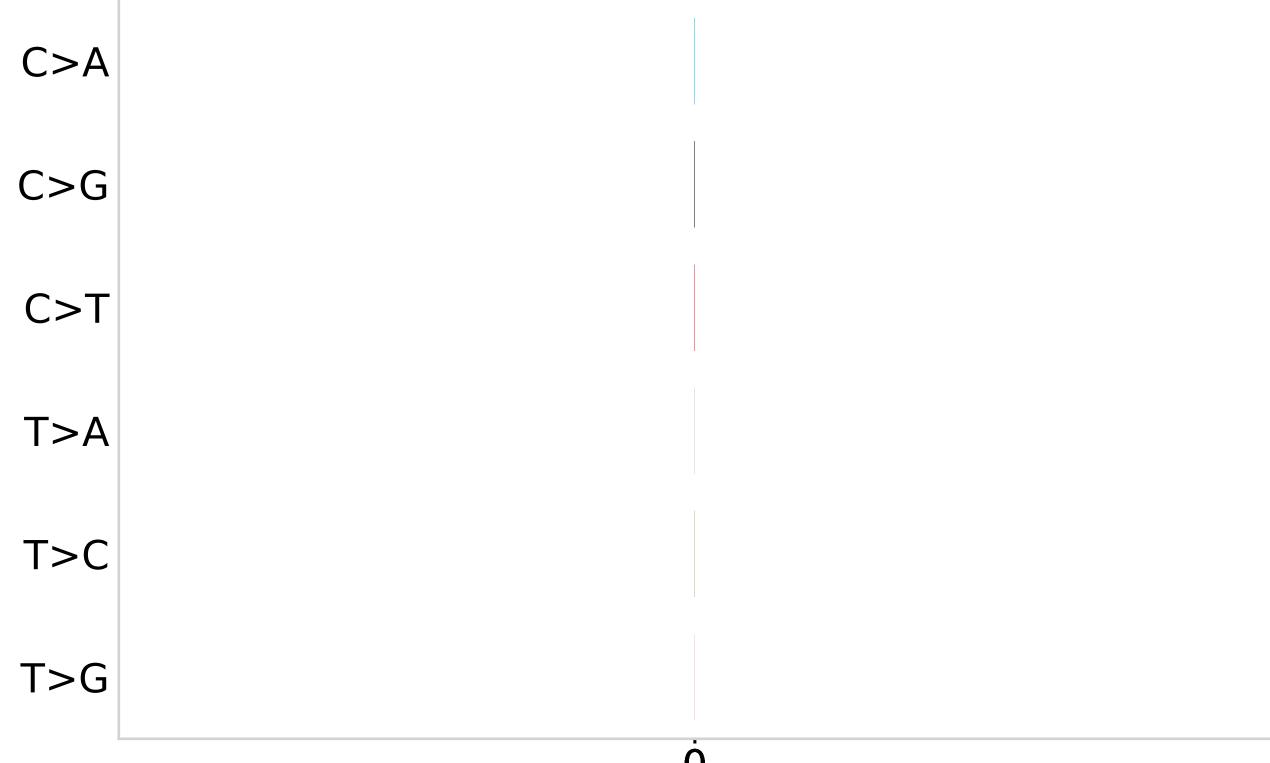
Number of Single Base Substitutions



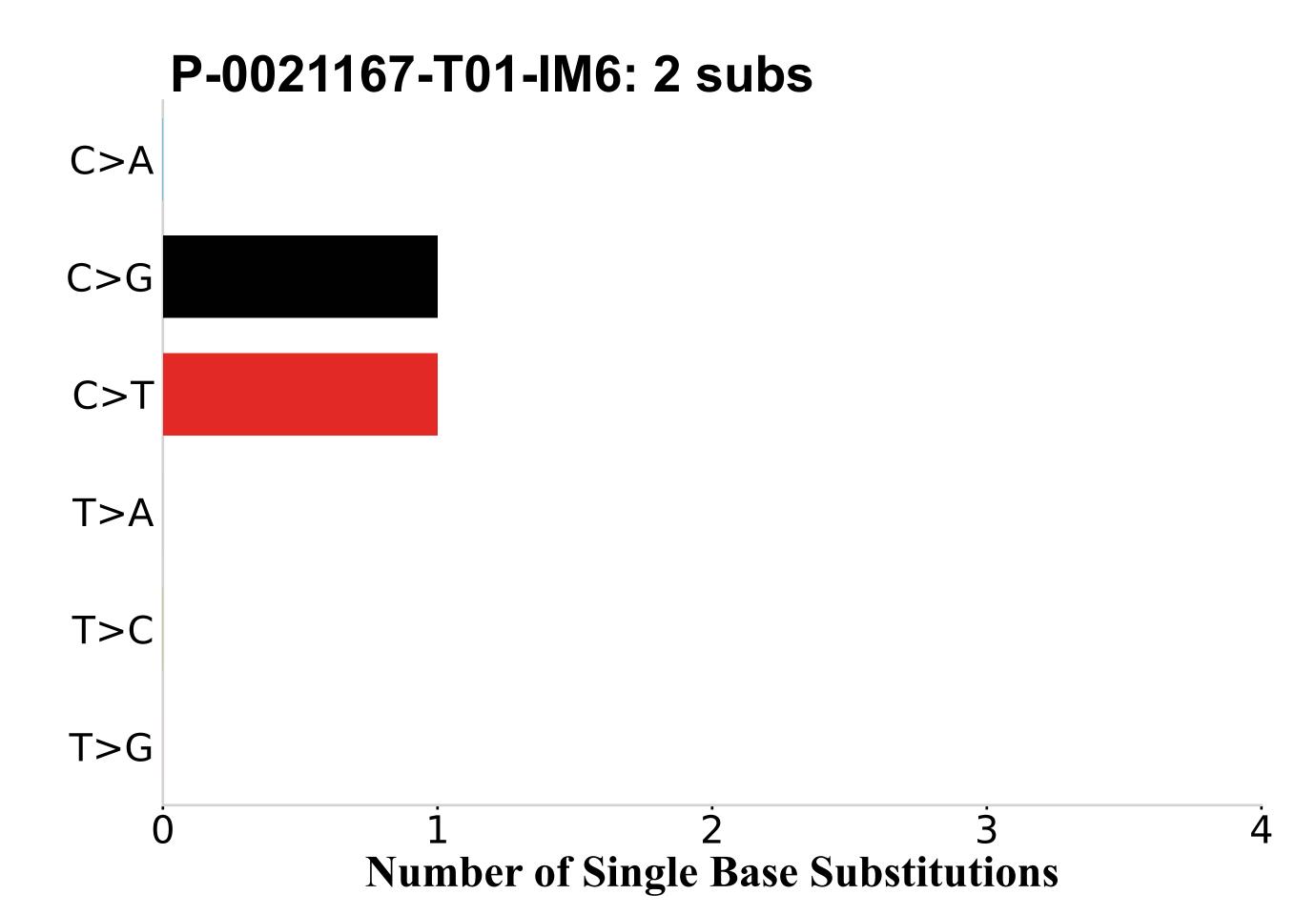


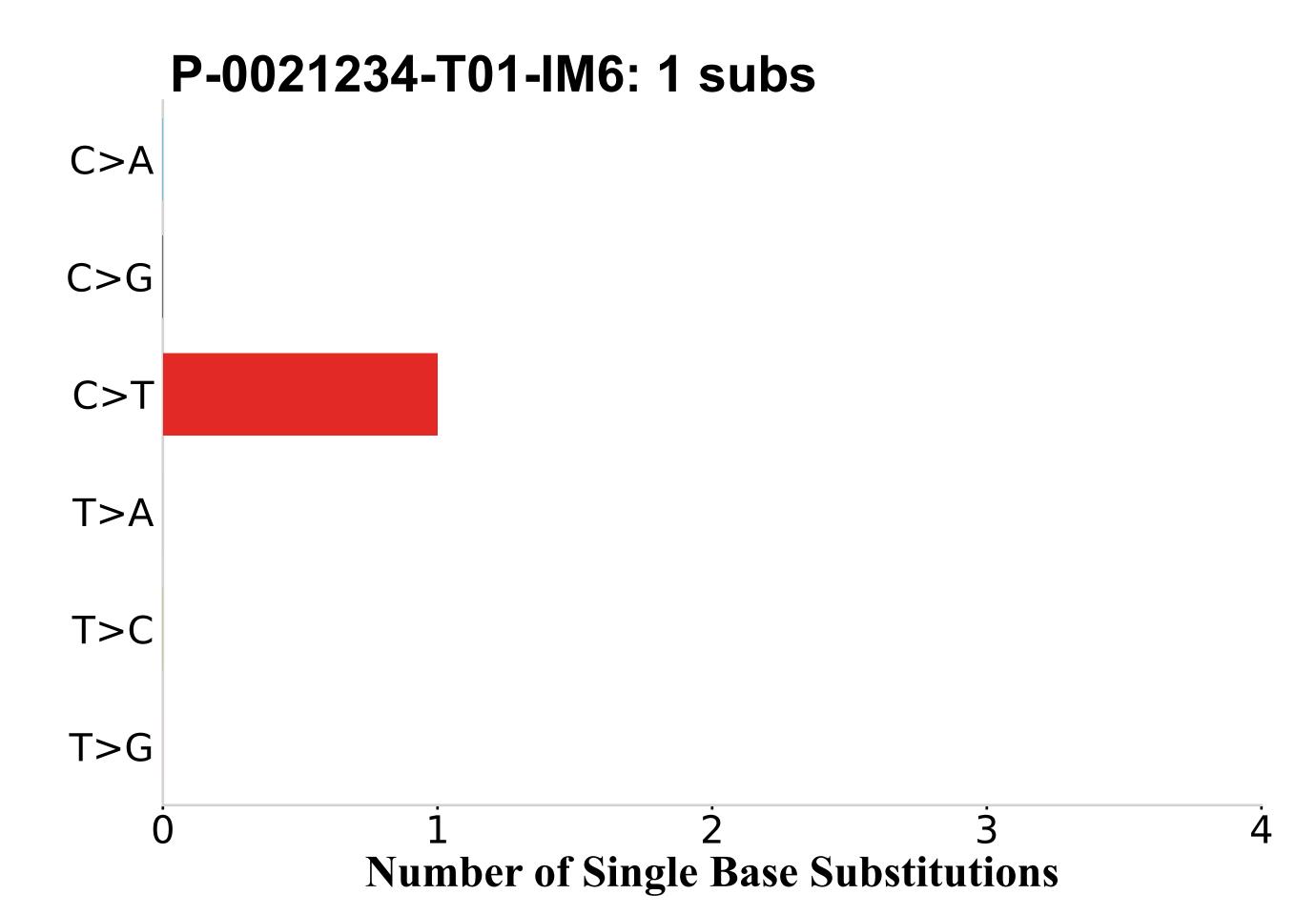


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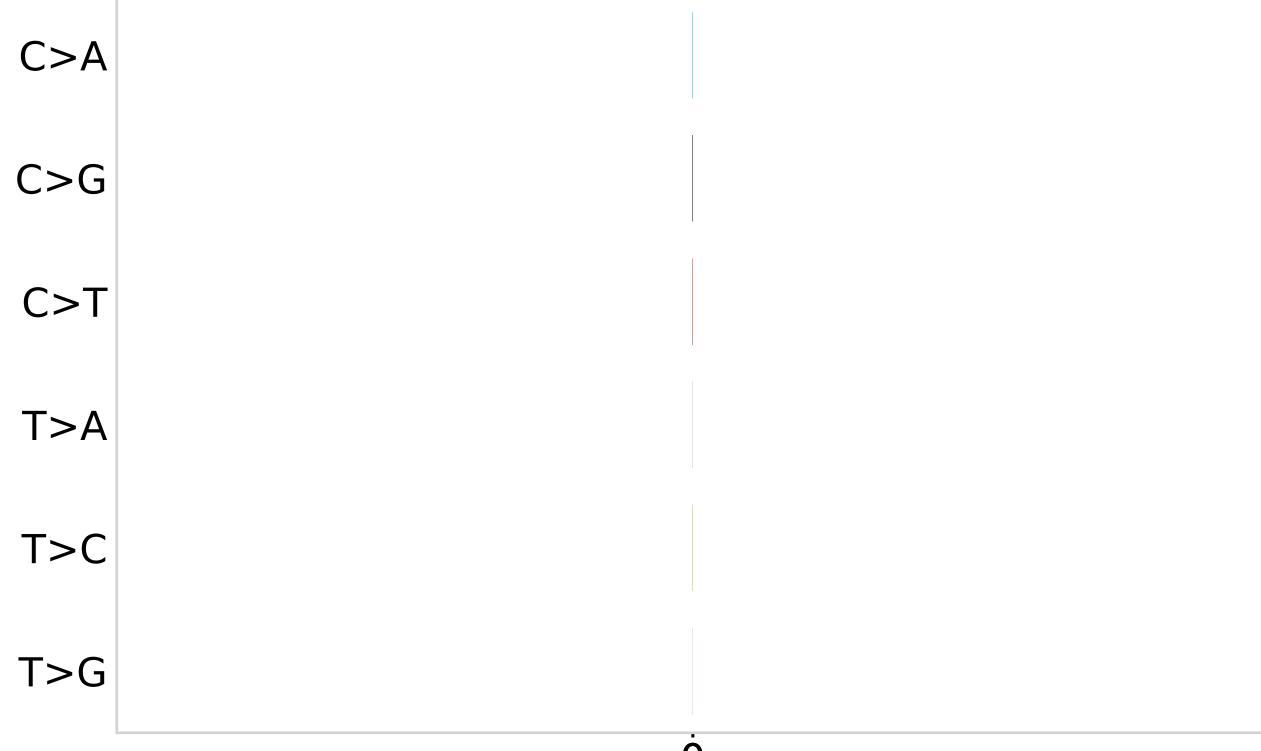


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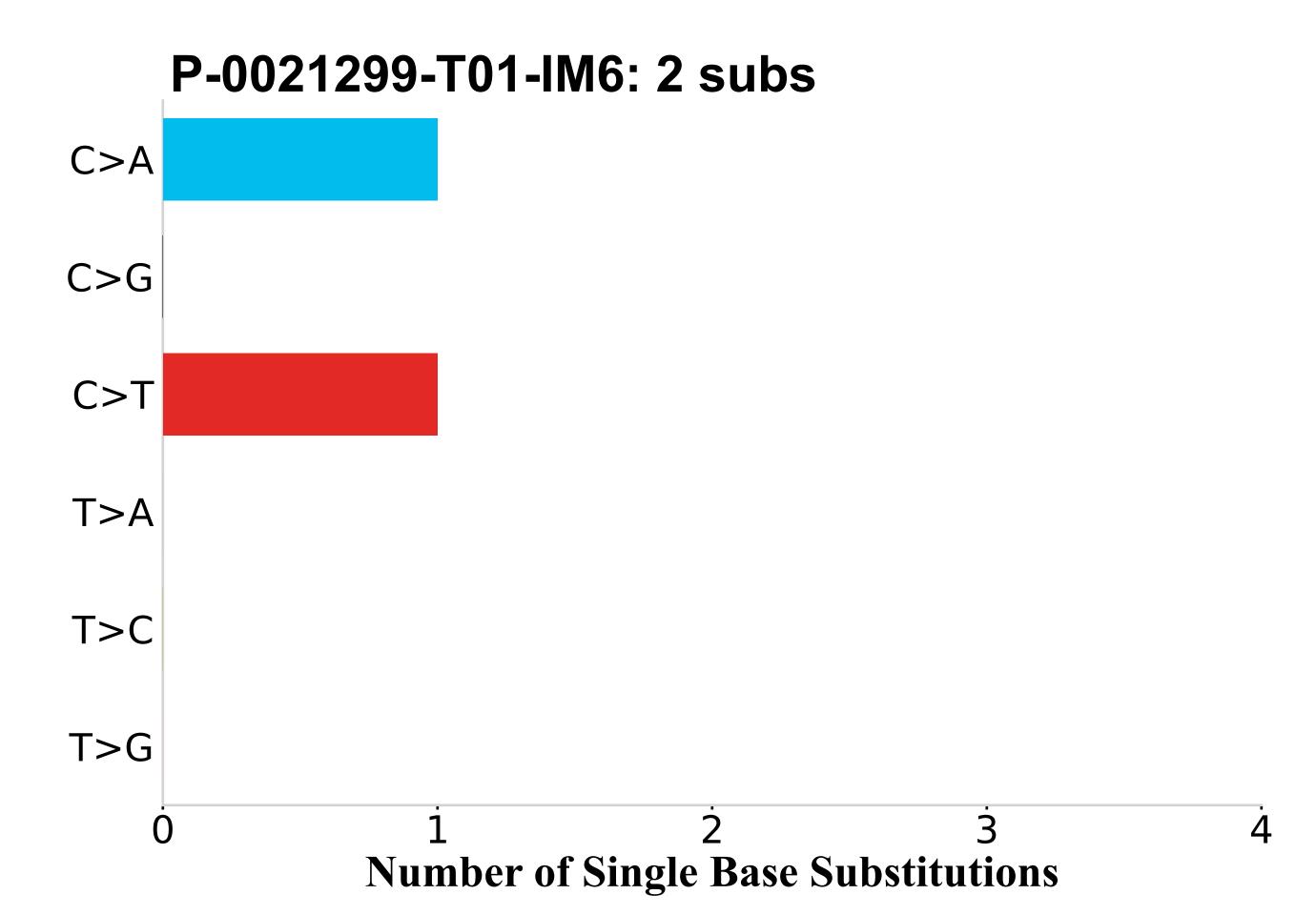


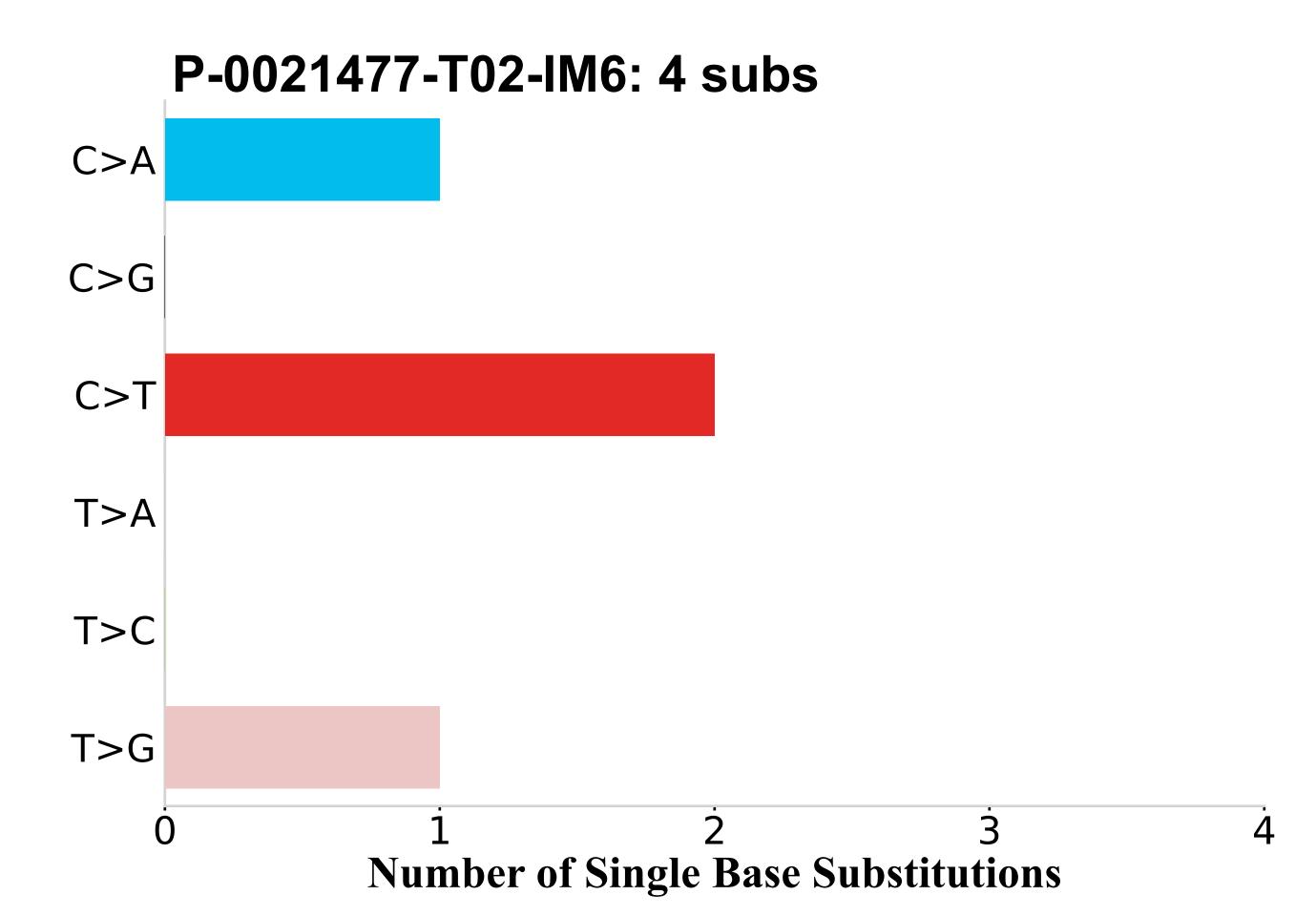


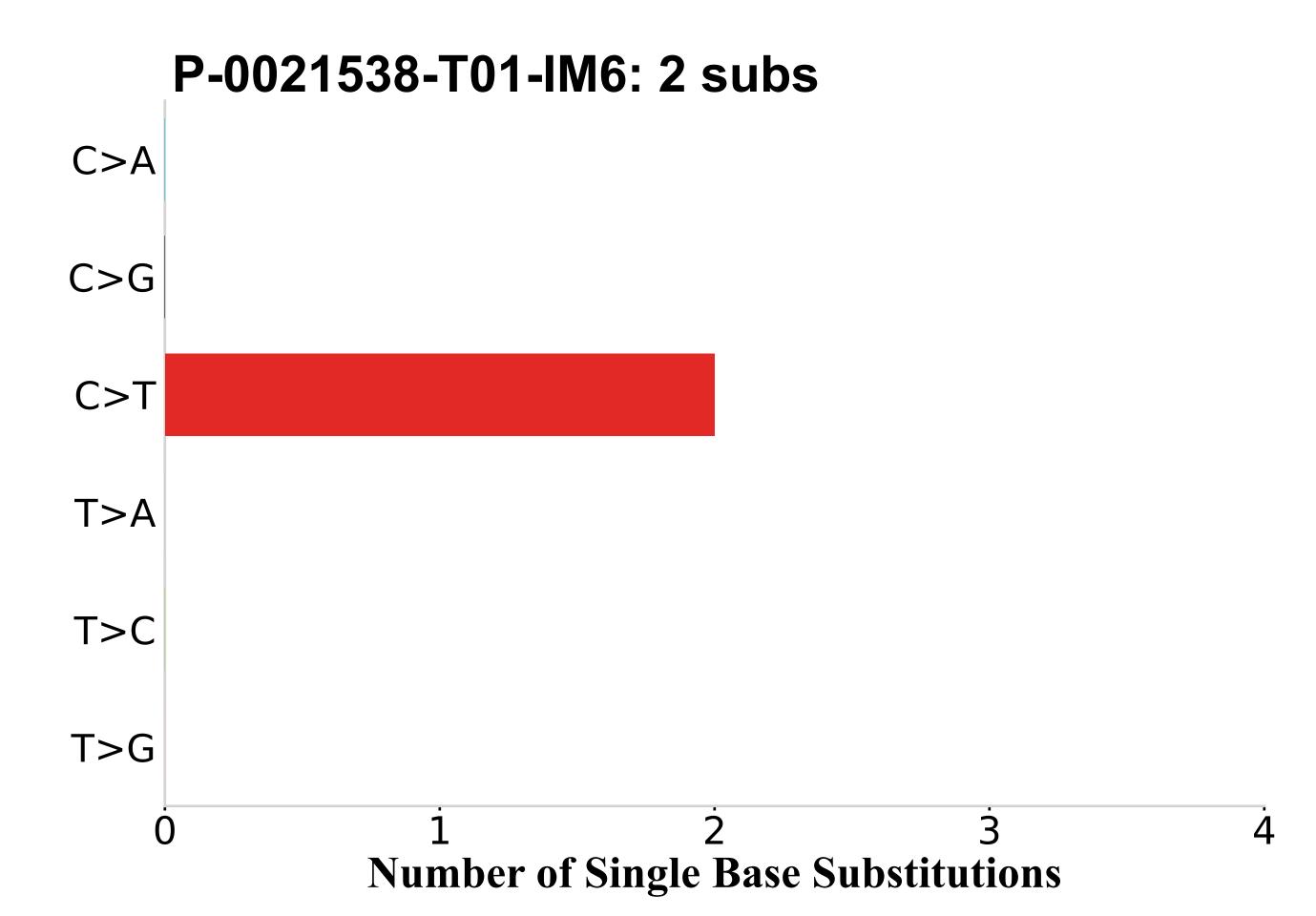
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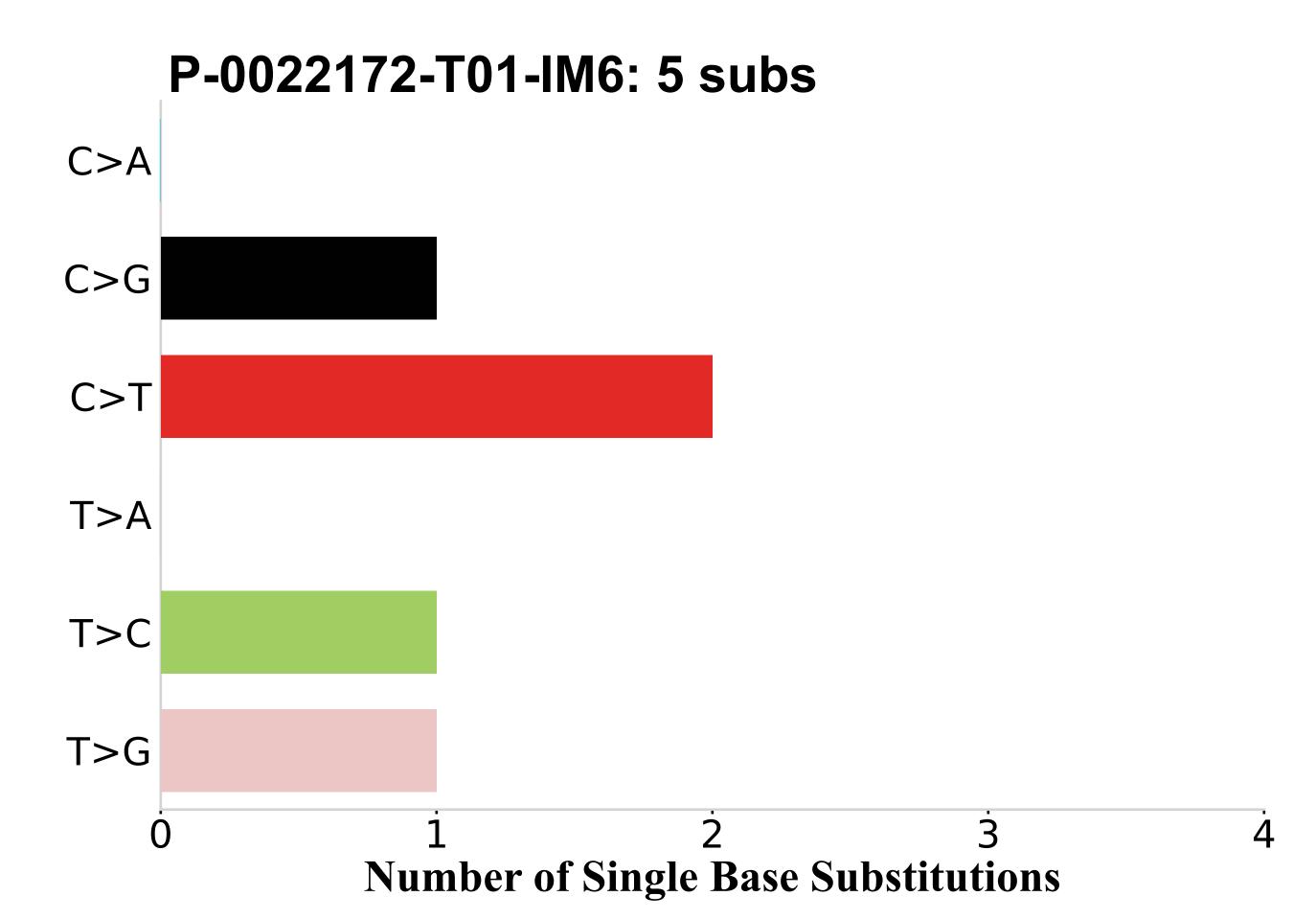


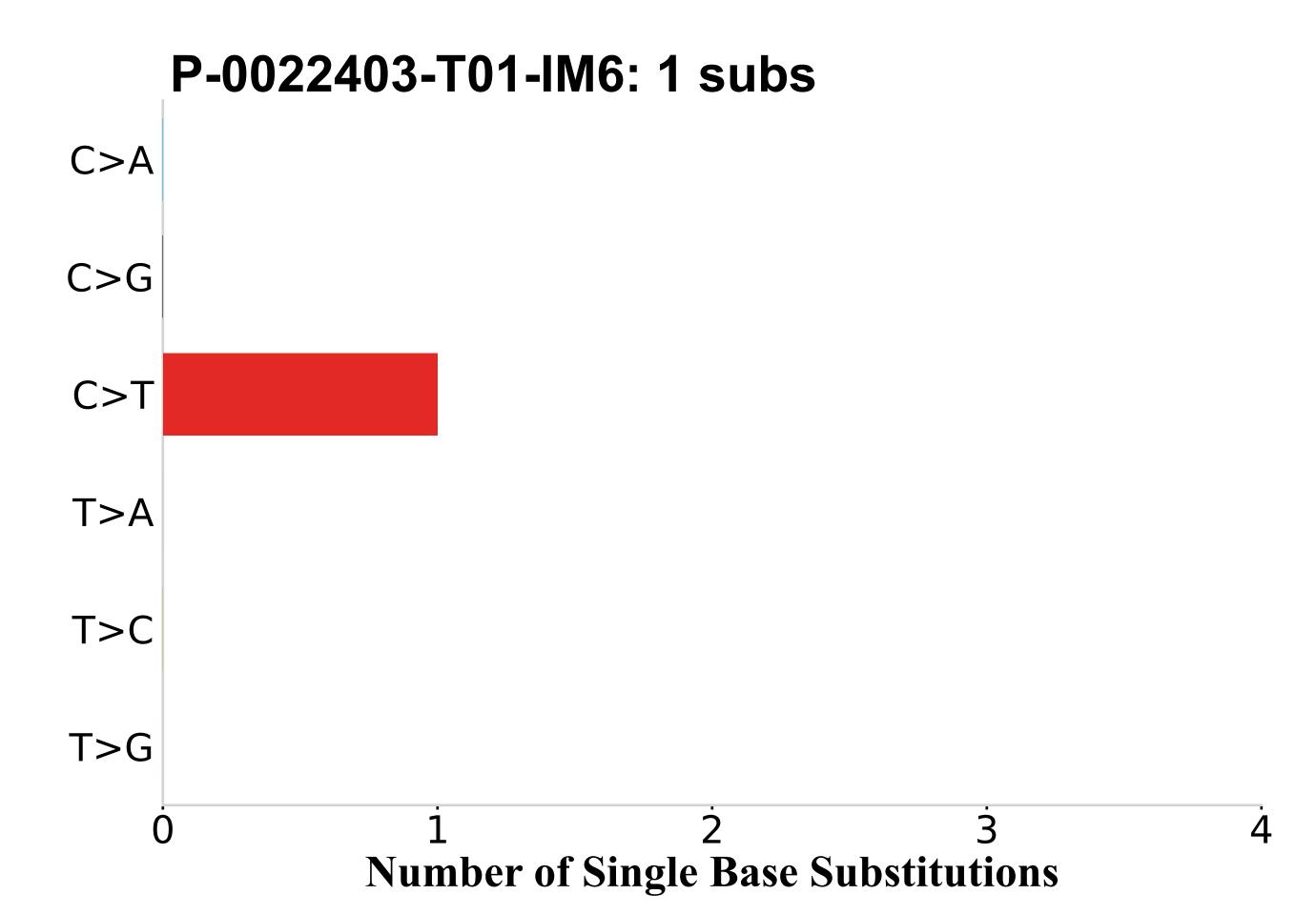
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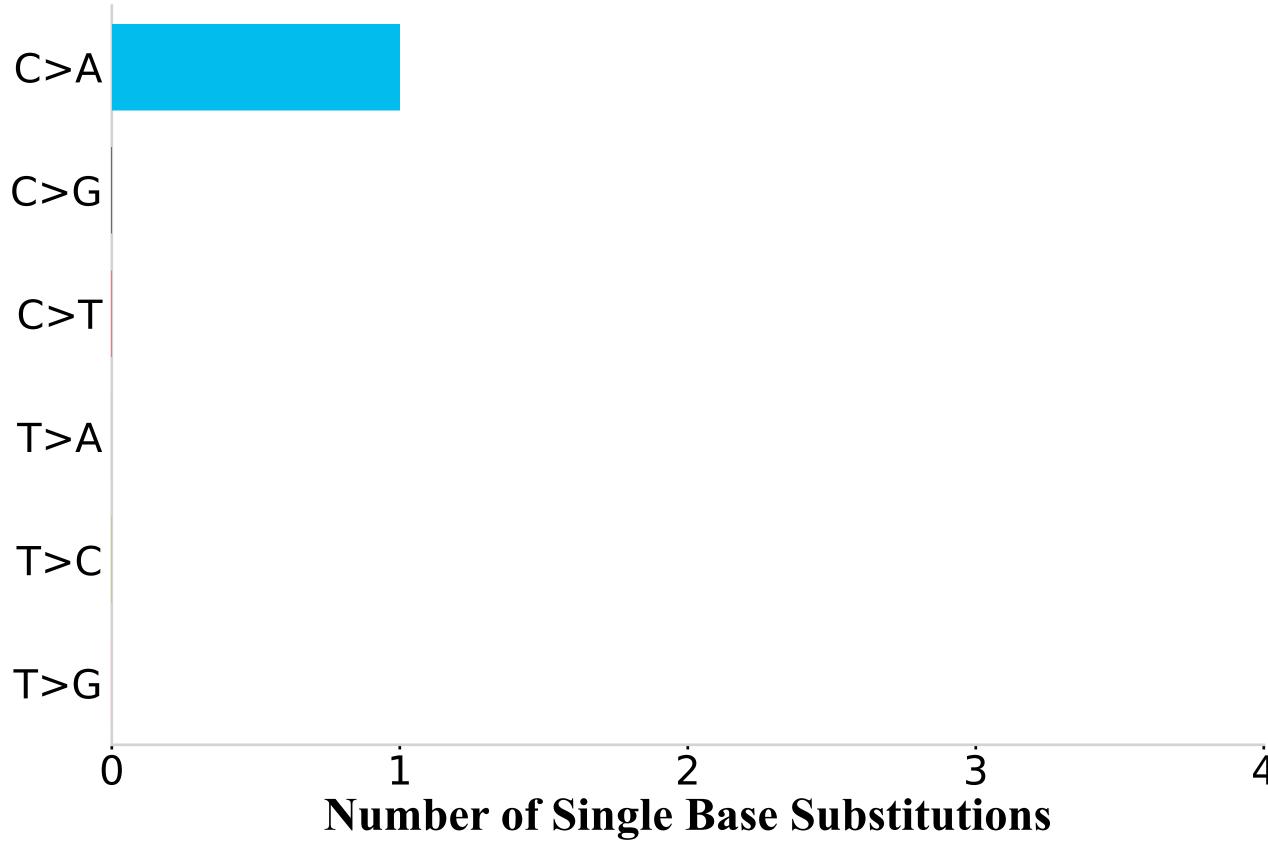














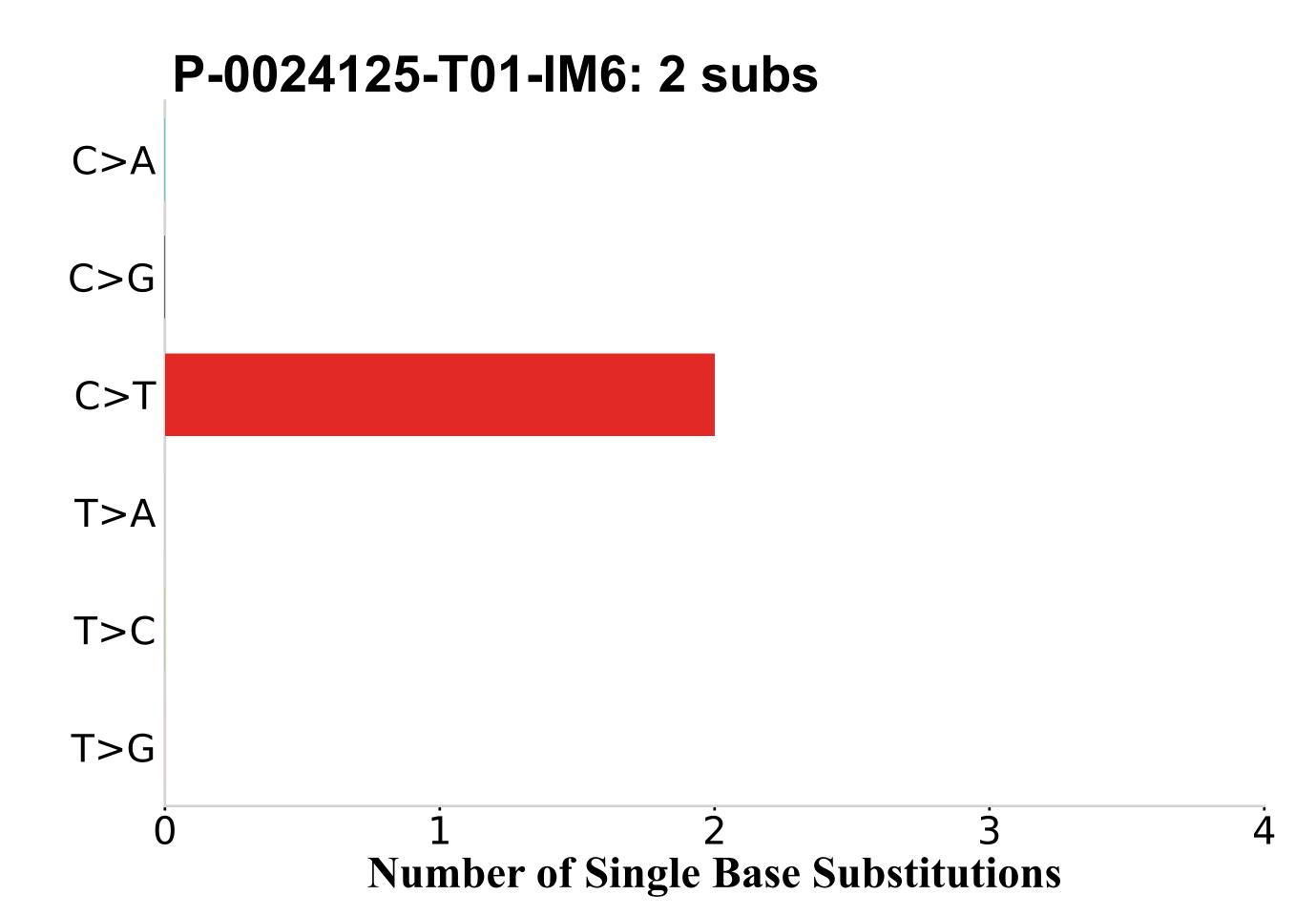


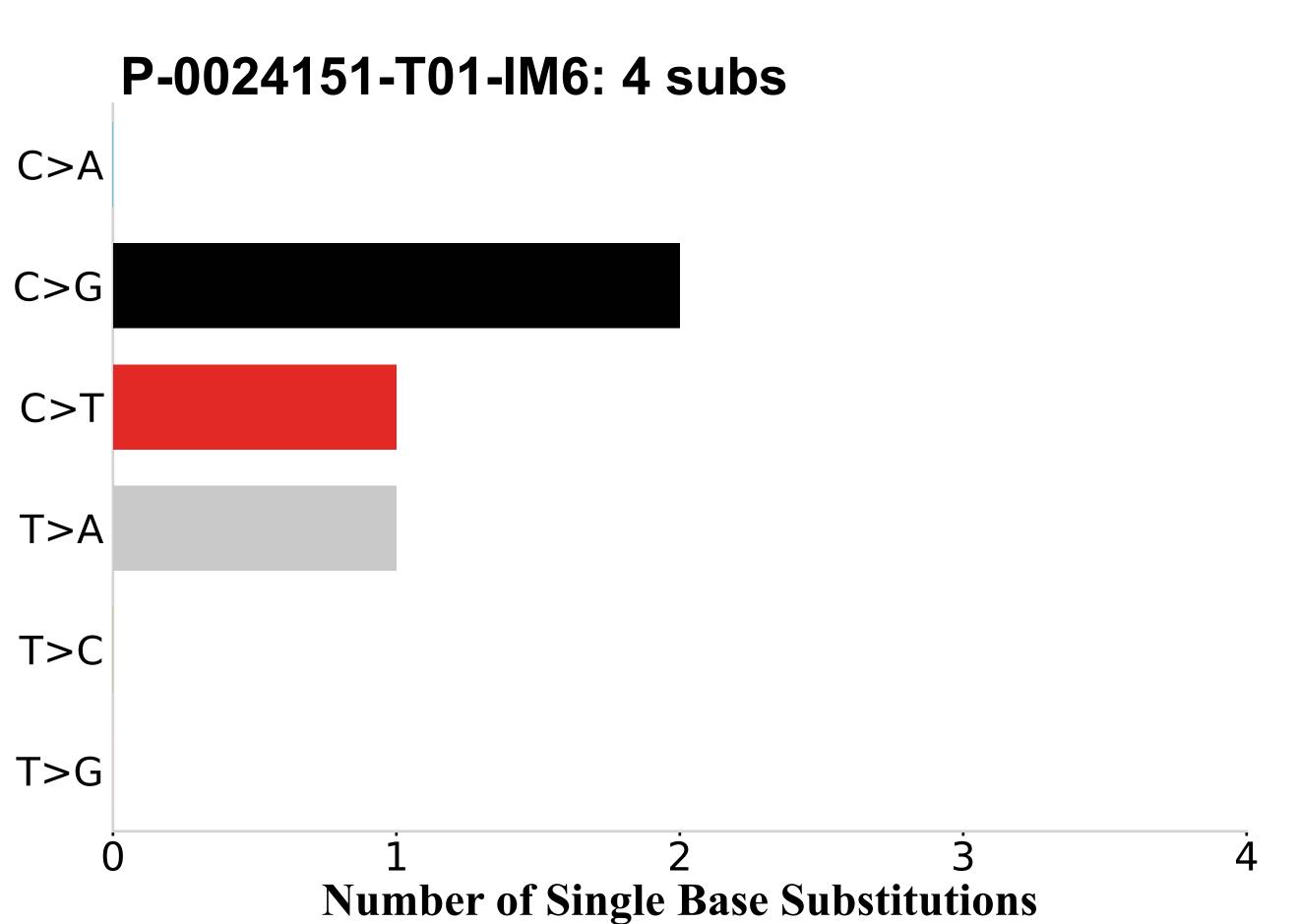
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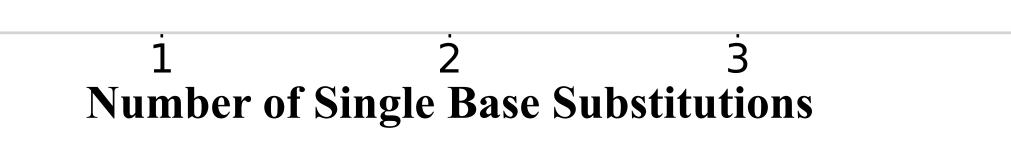


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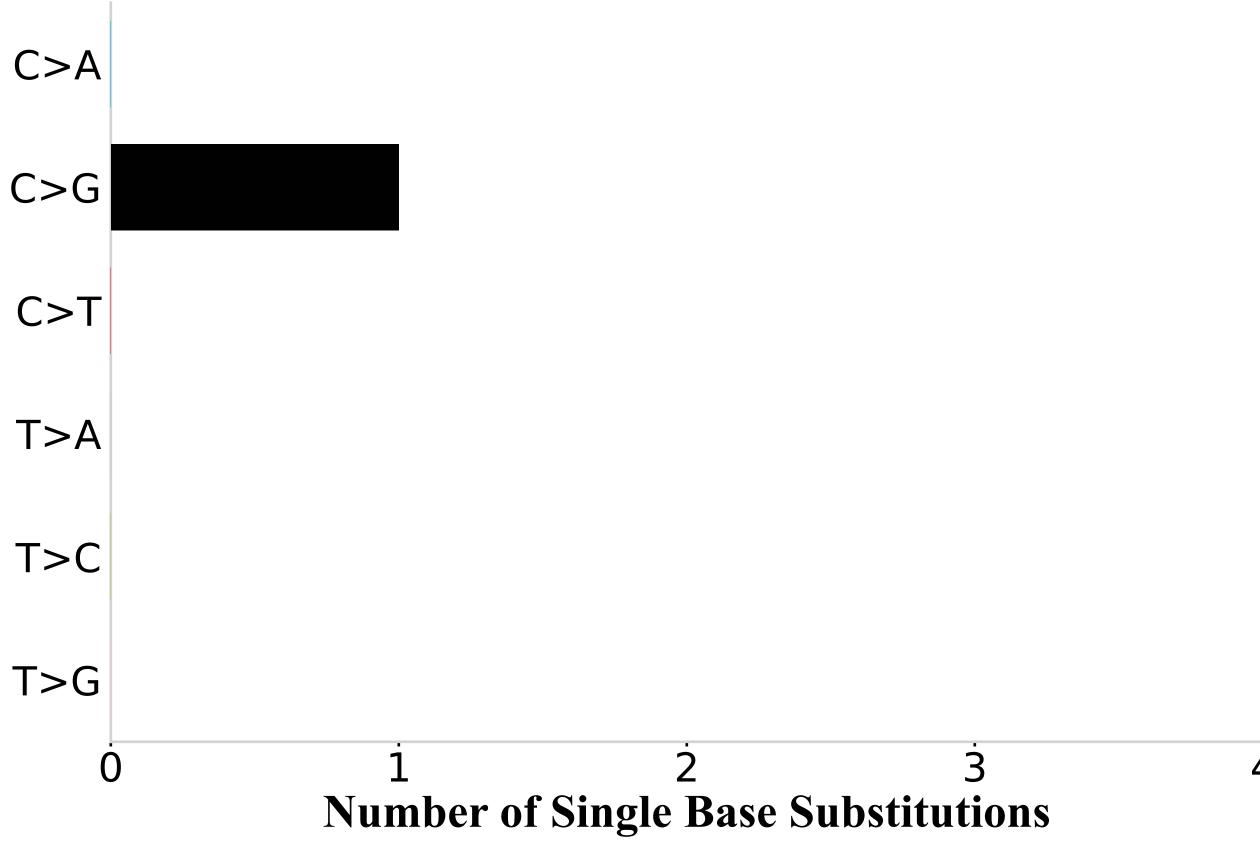


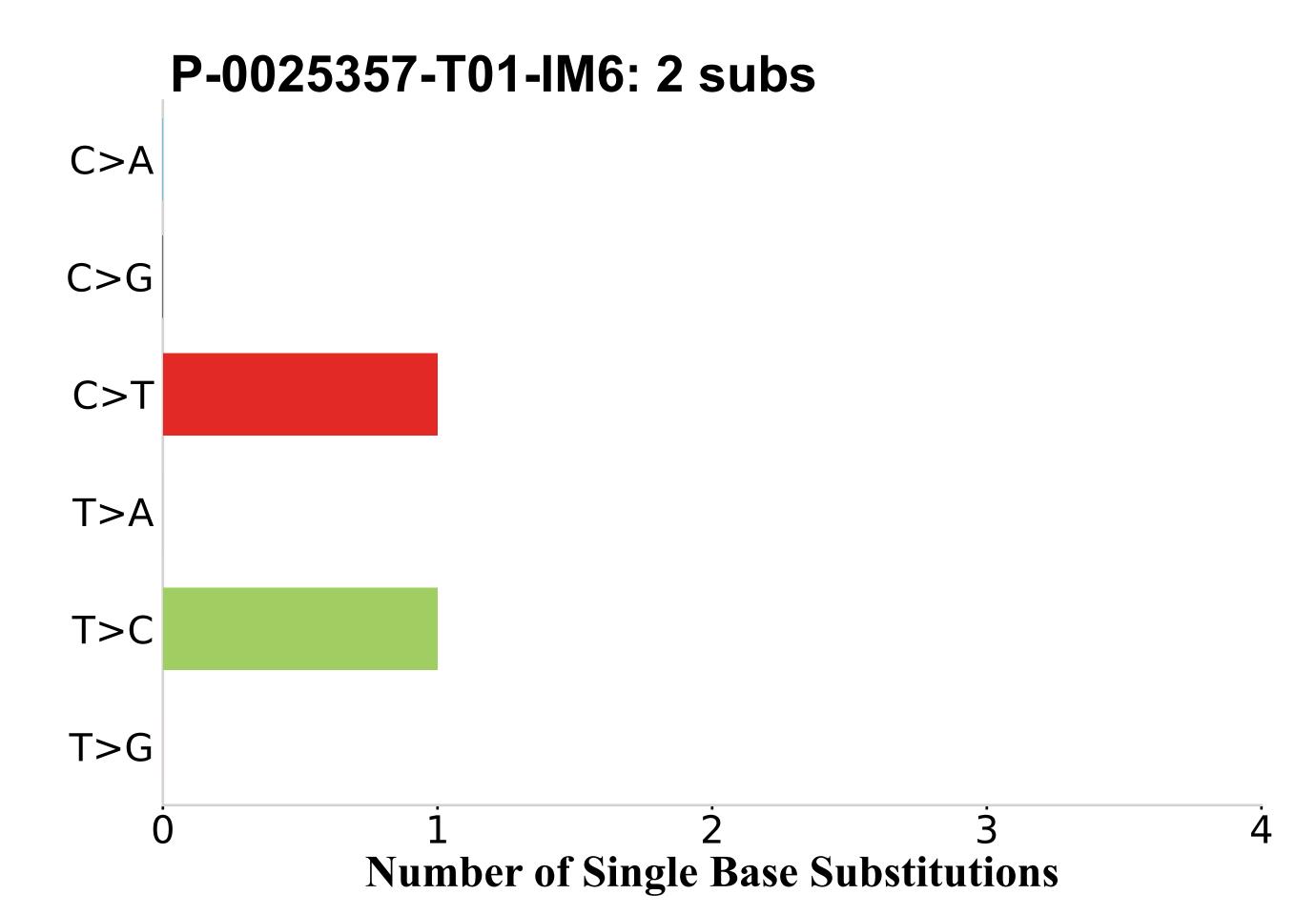


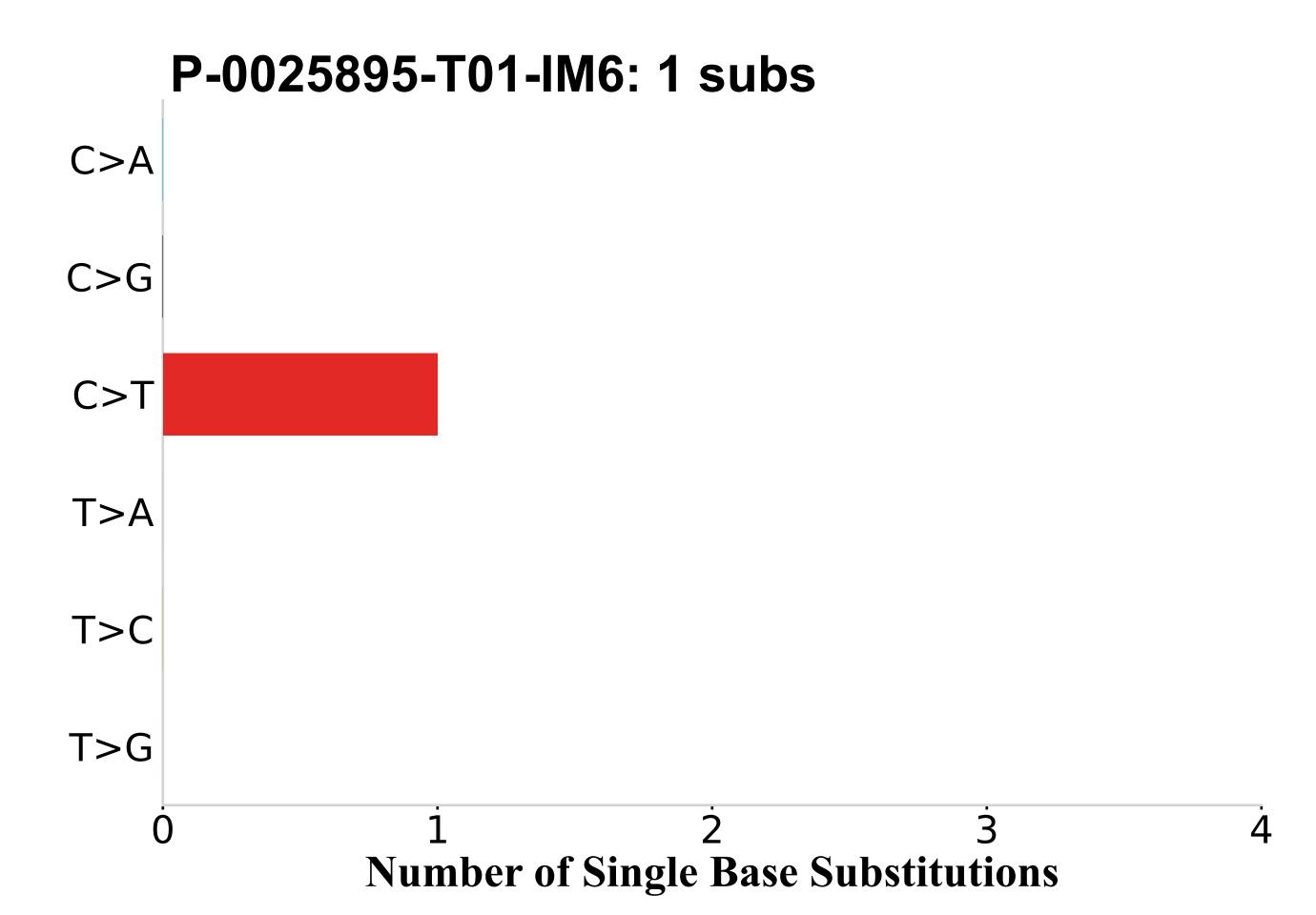
P-0024612-T01-IM6: 2 subs C>A C>G C>T T>A T>C T>G



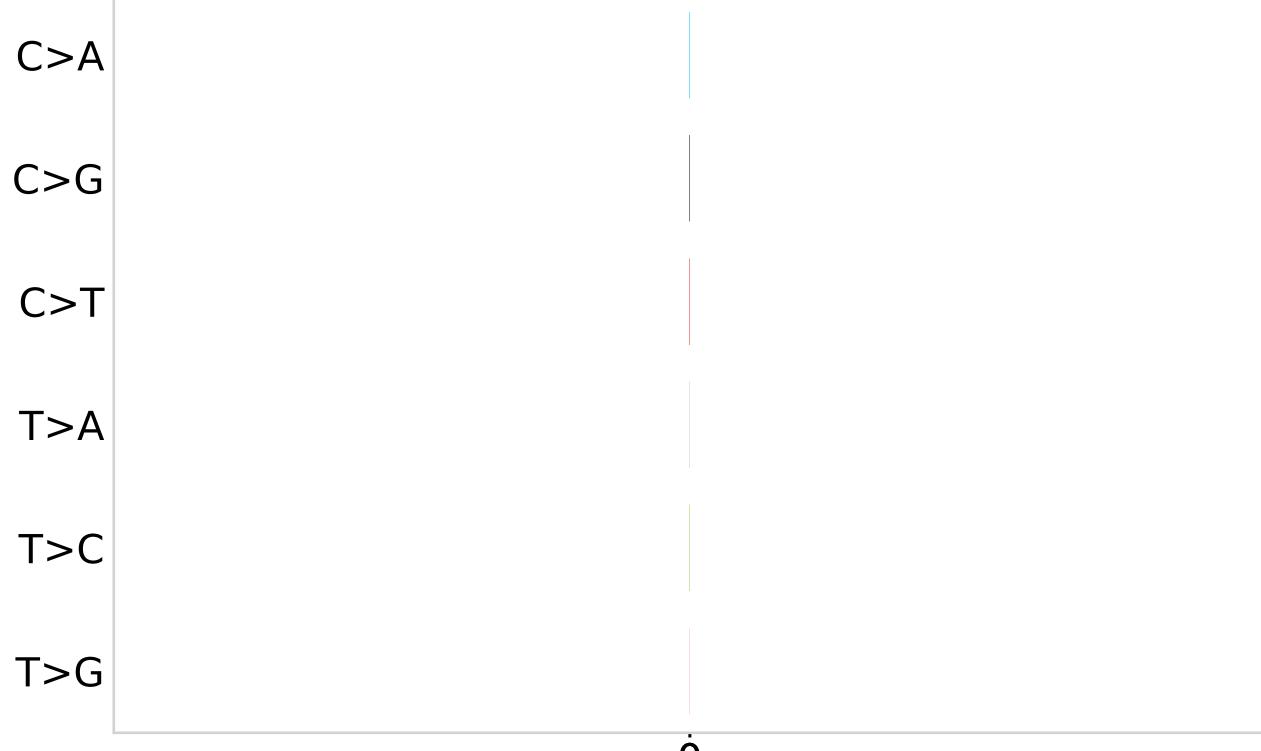




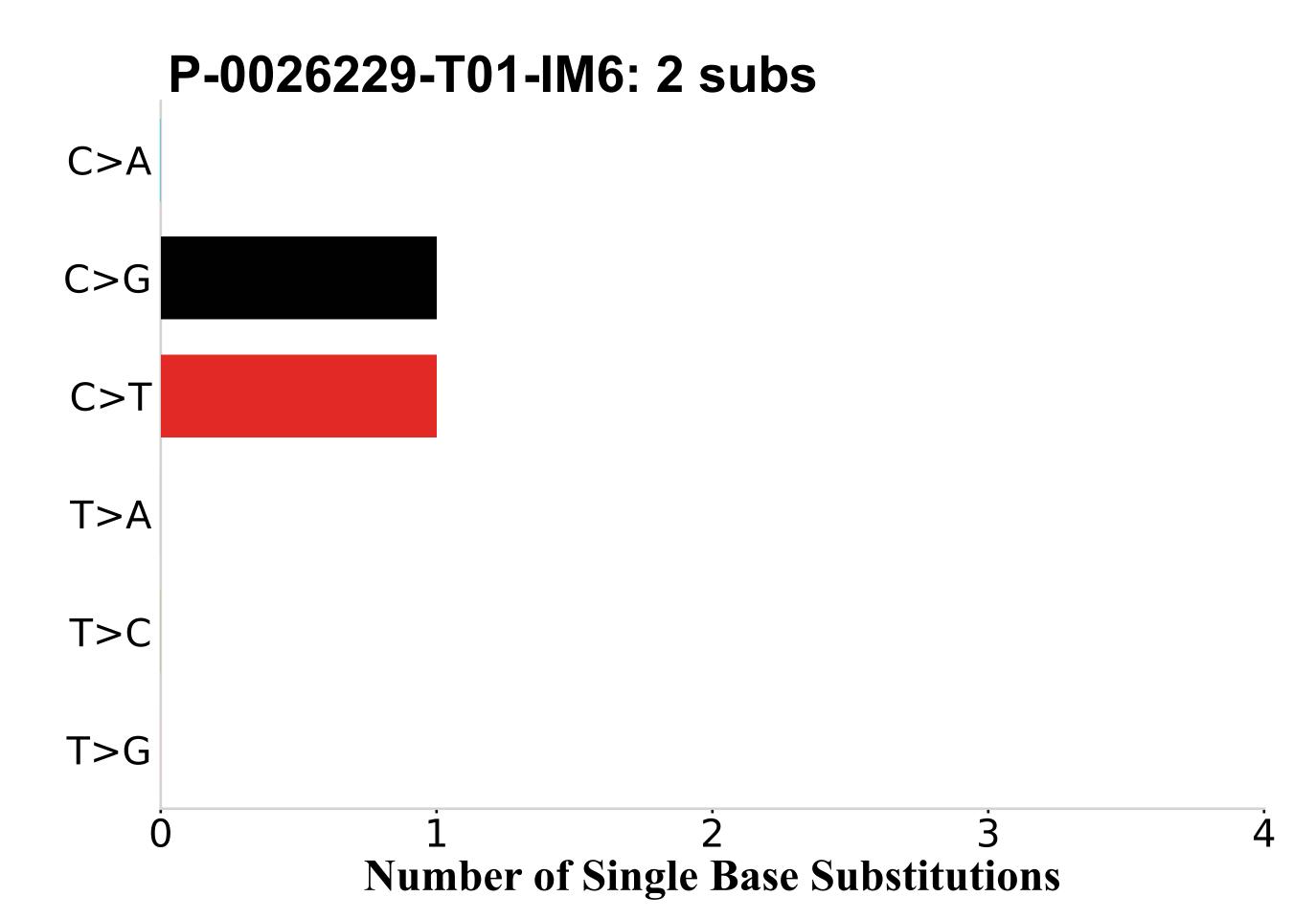


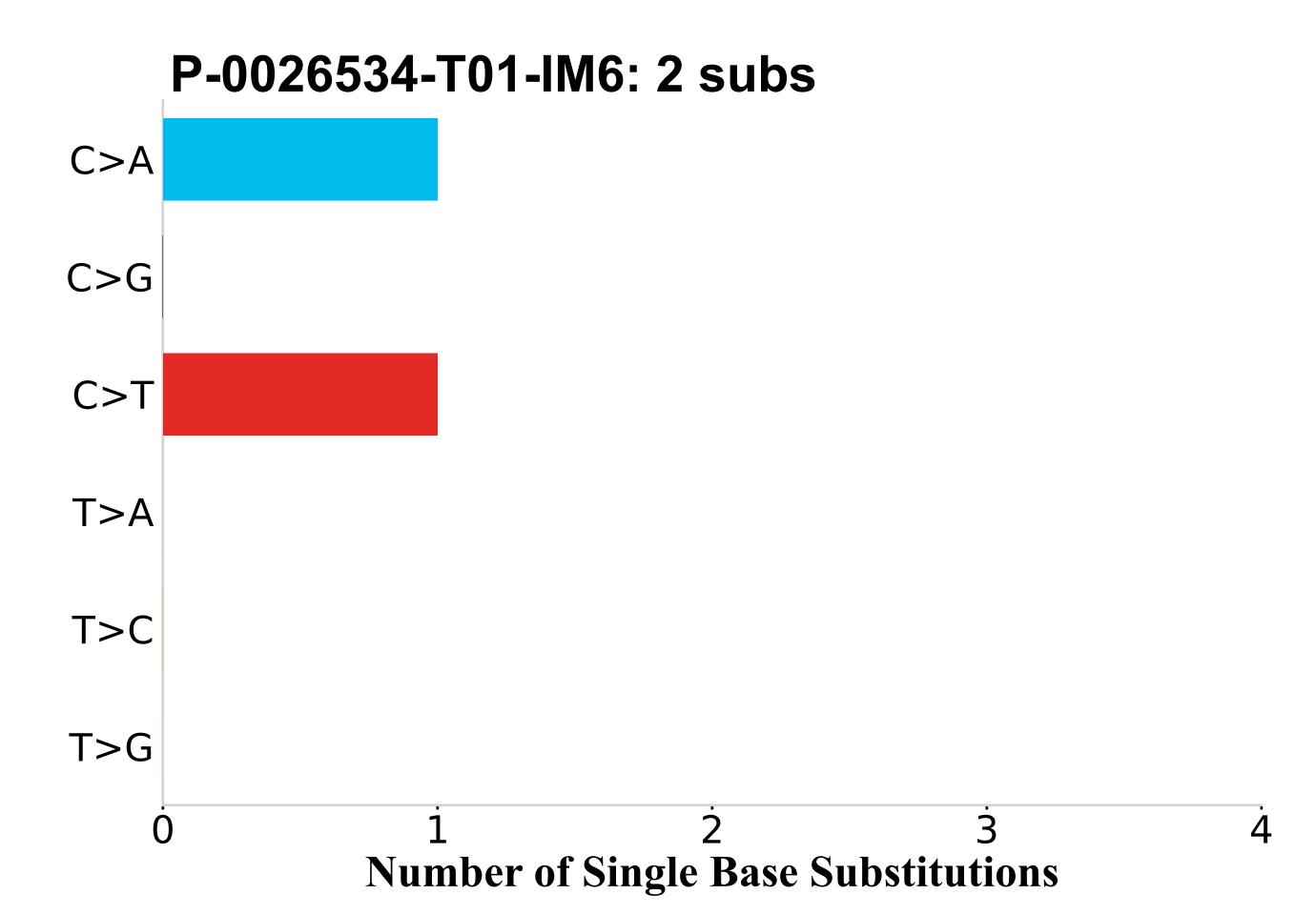


P-0025944-T02-IM6: 0 subs

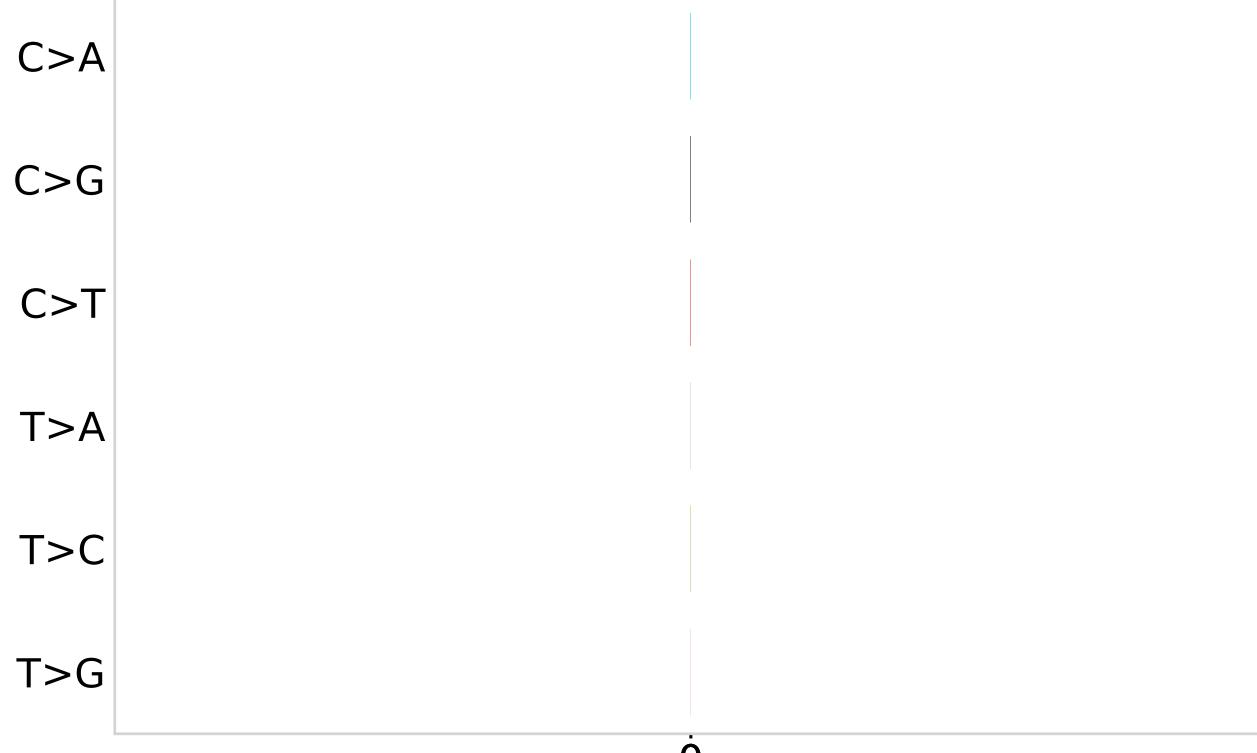


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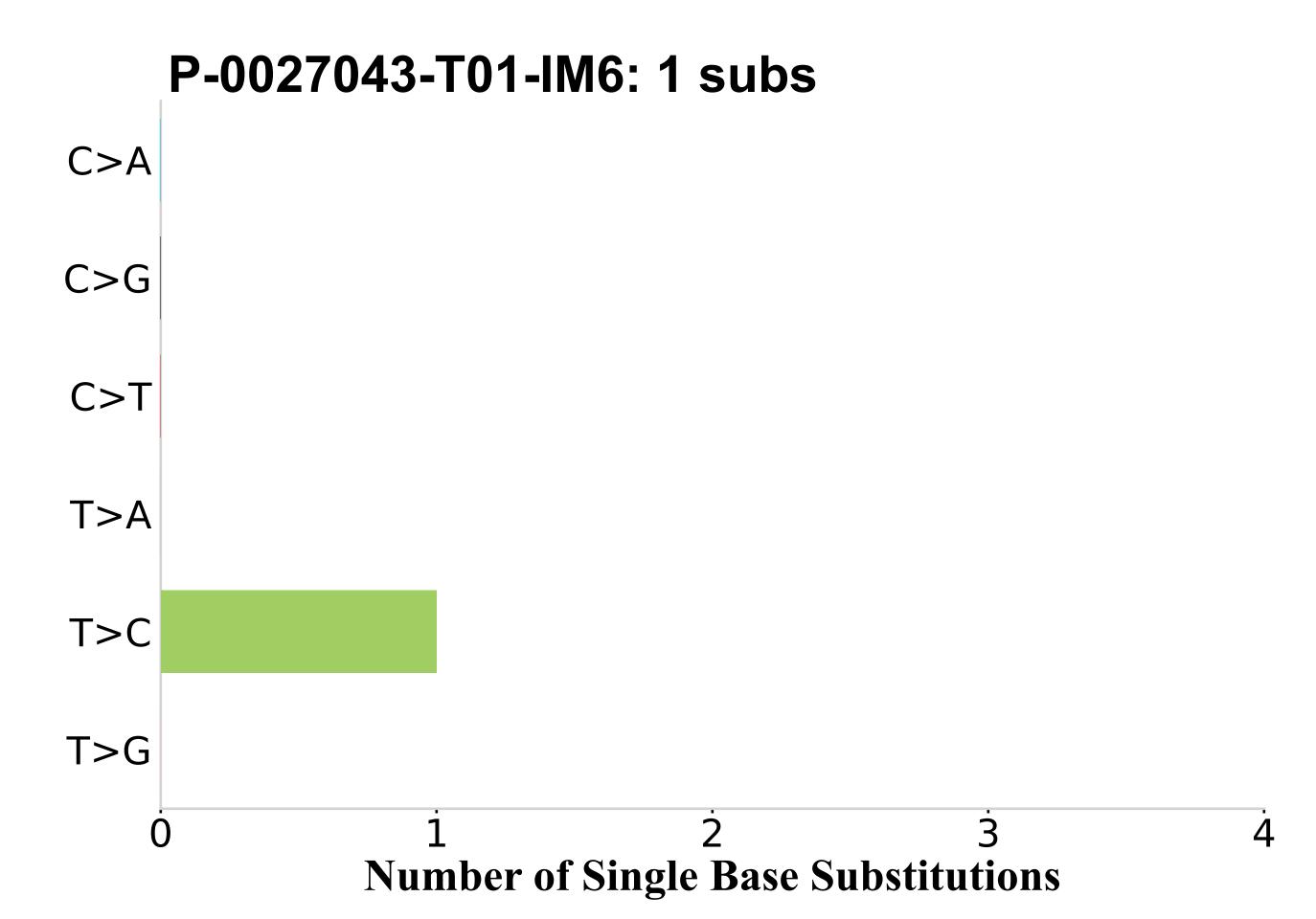


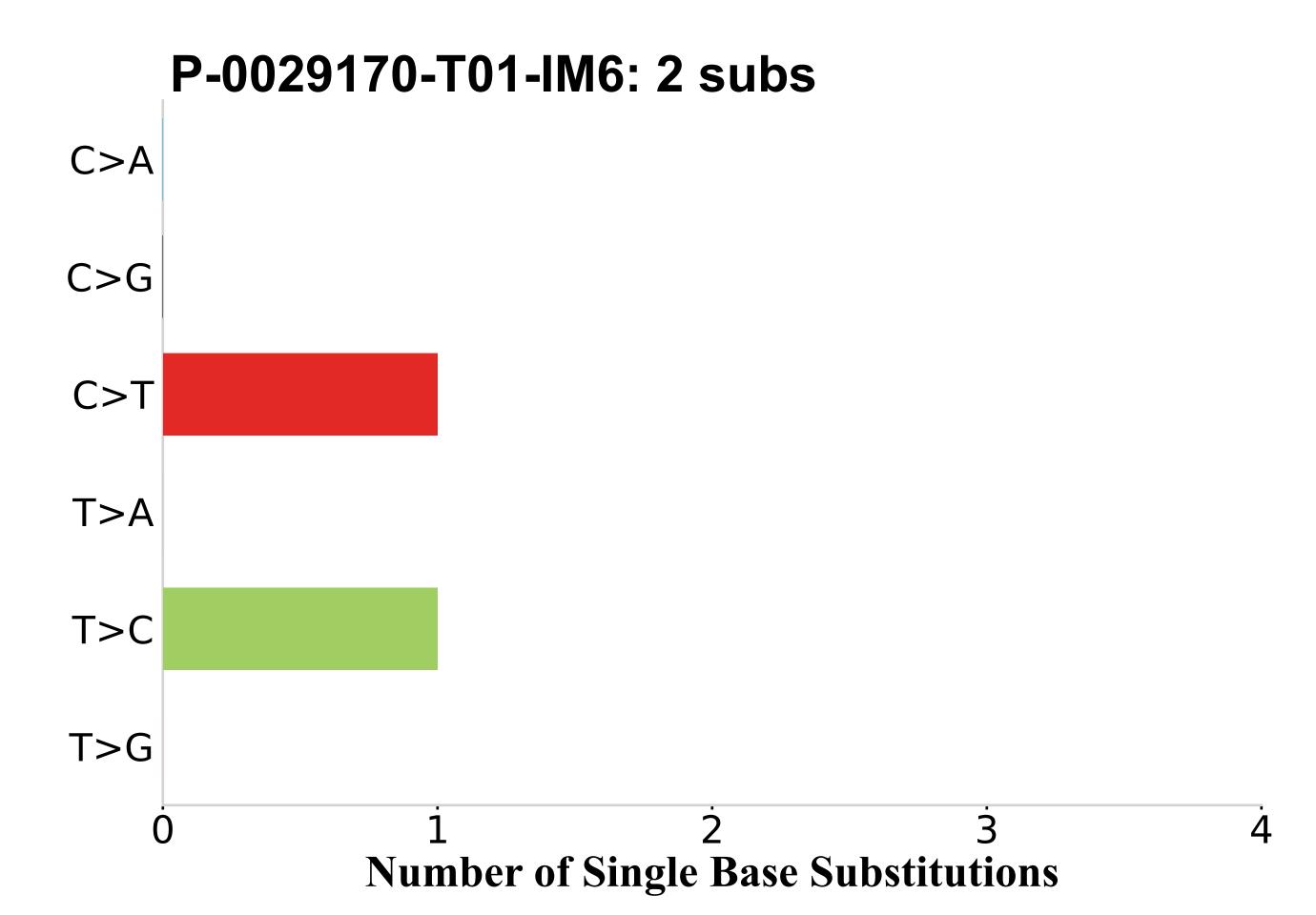


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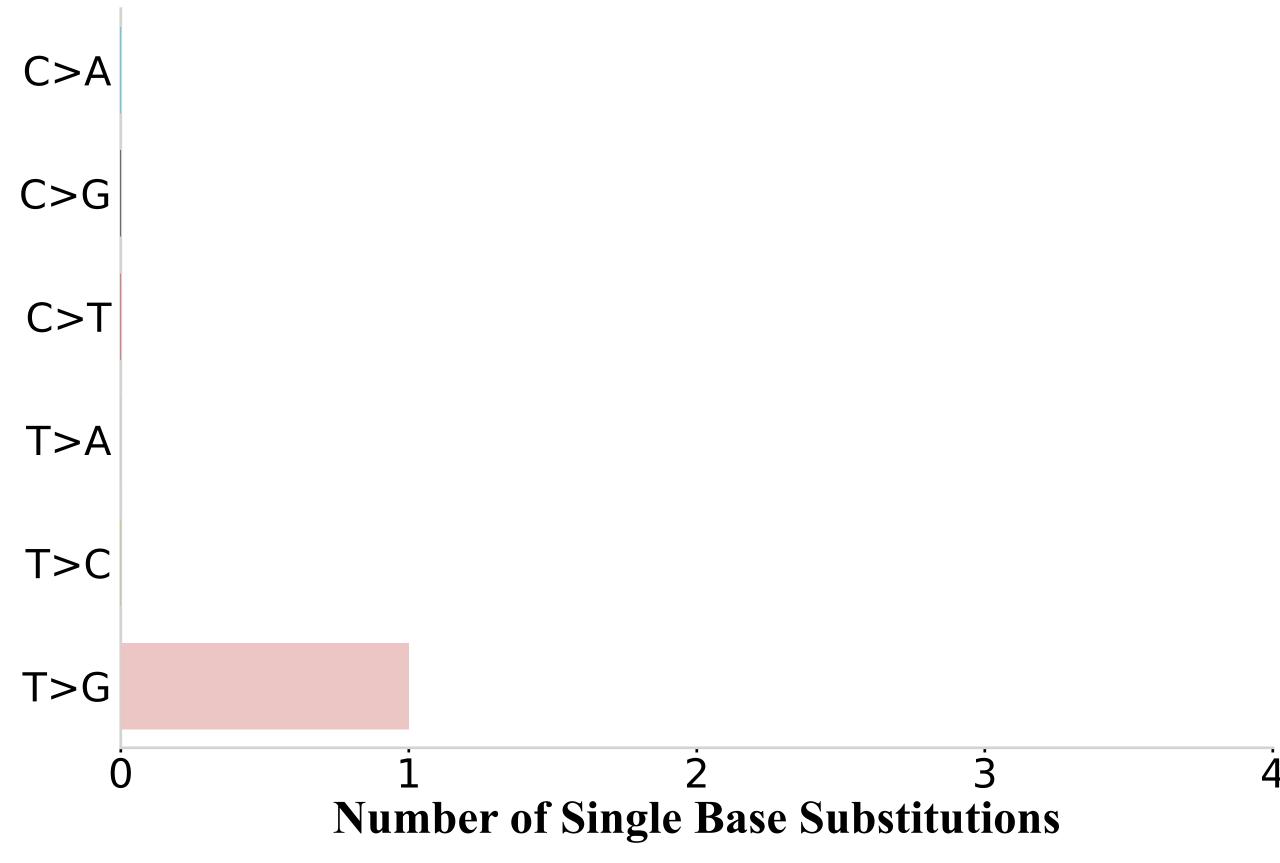


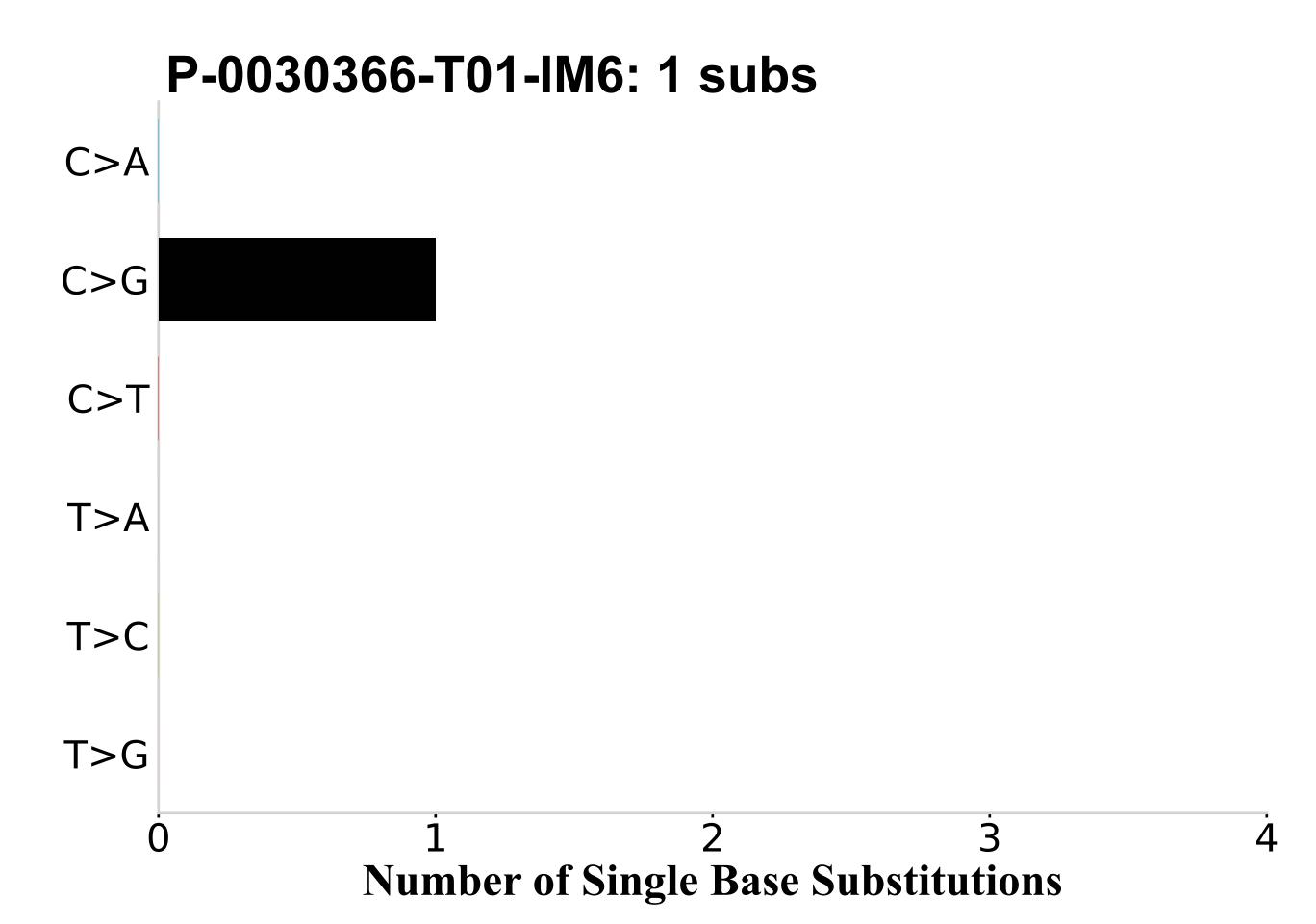
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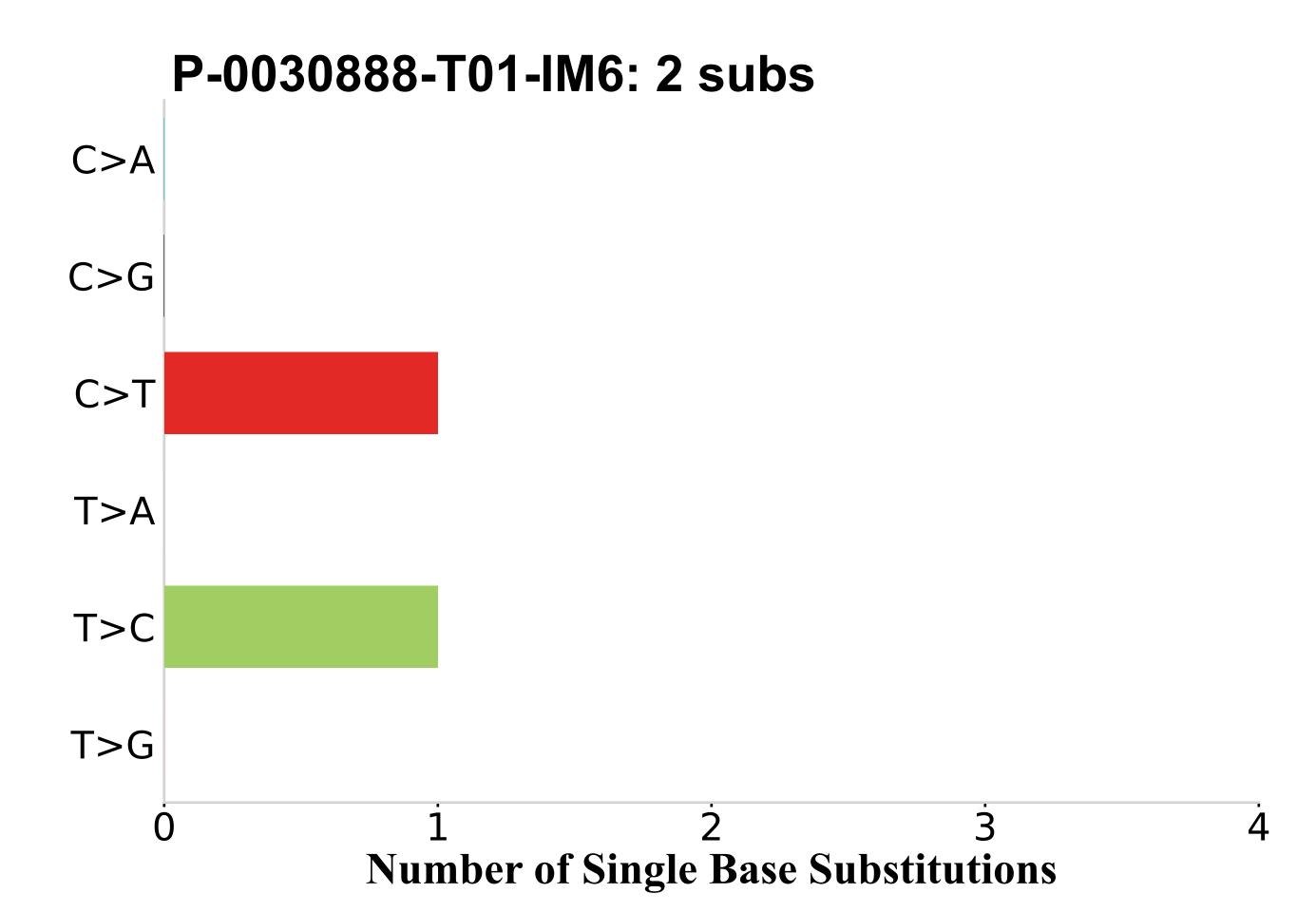


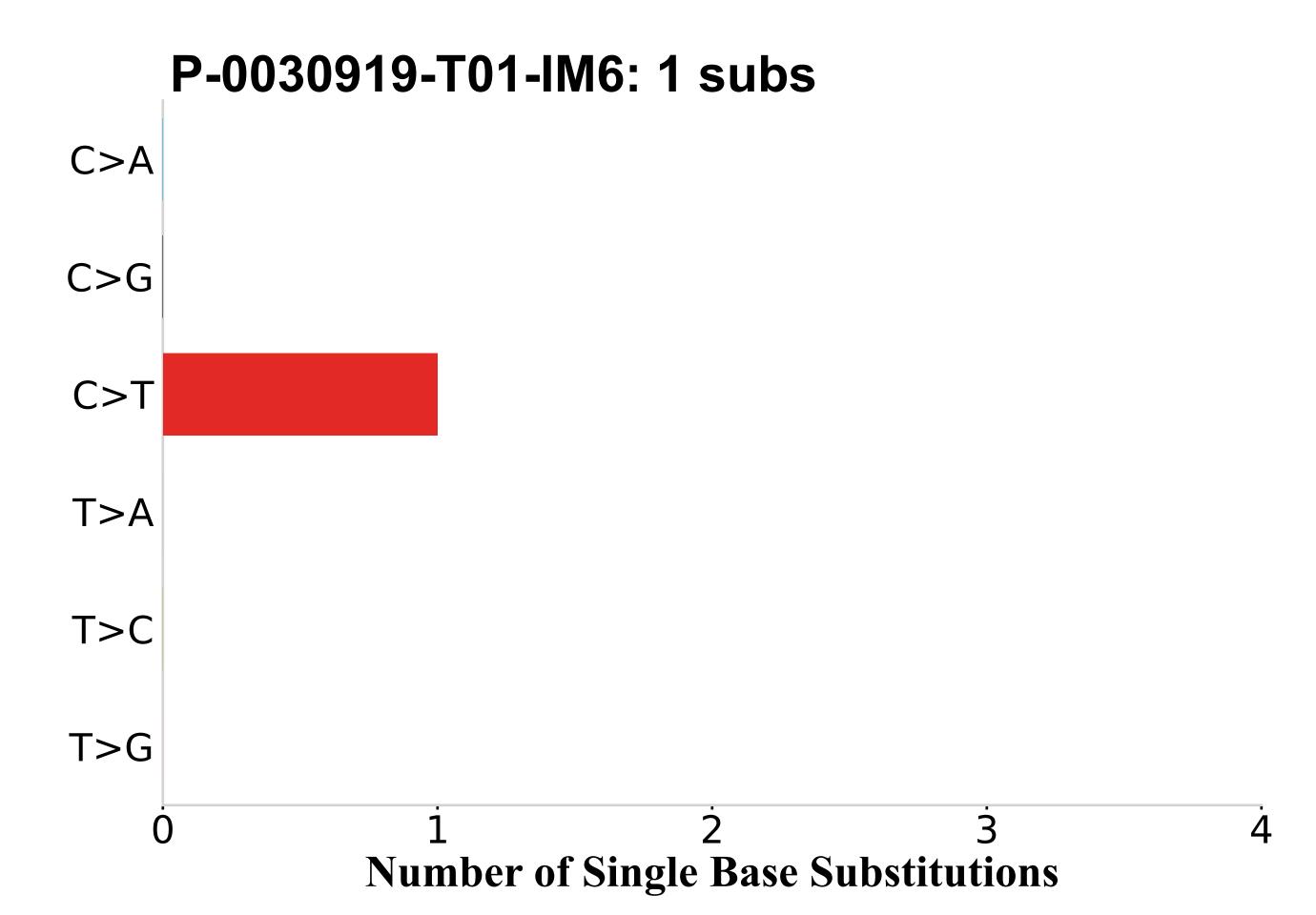


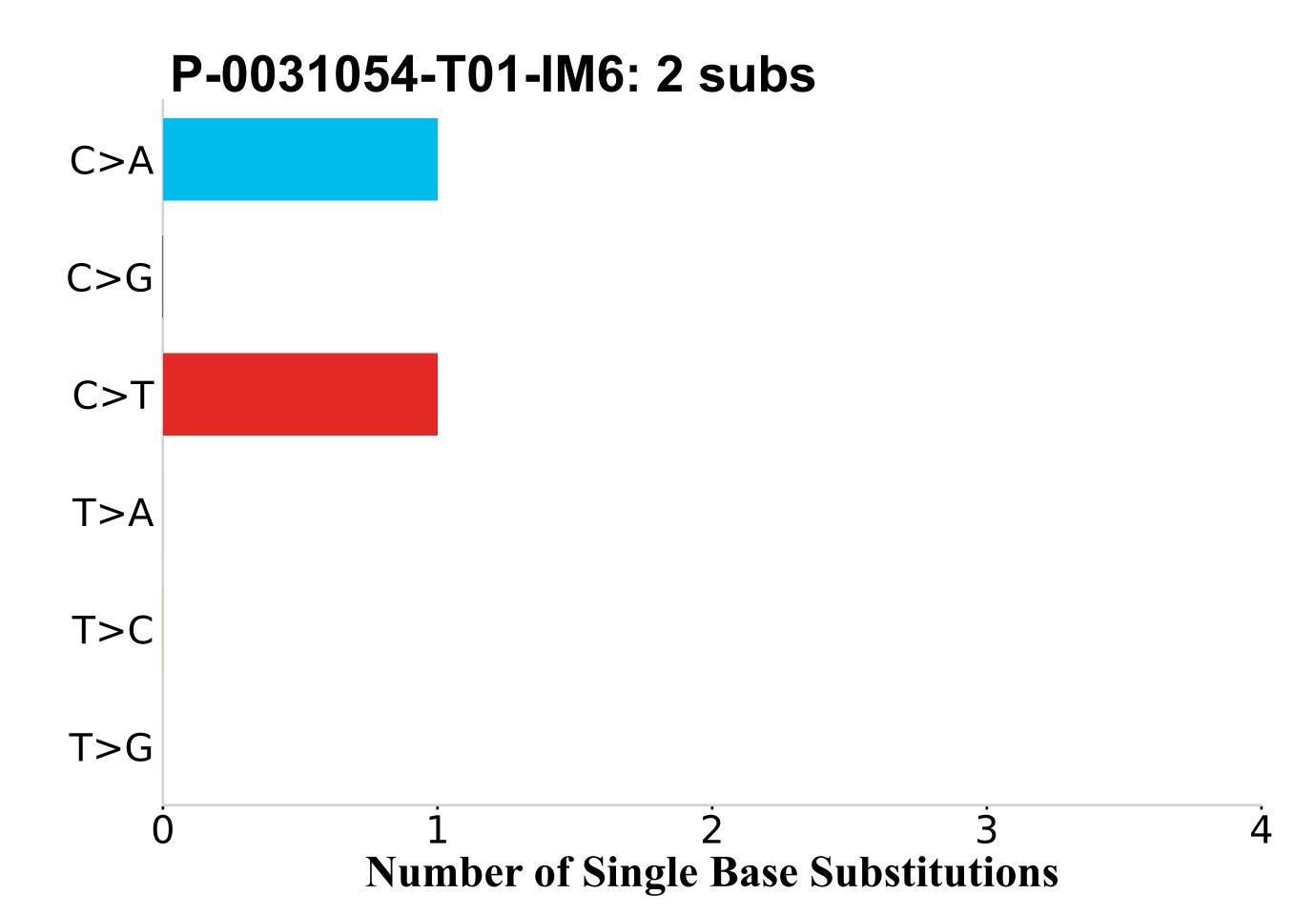
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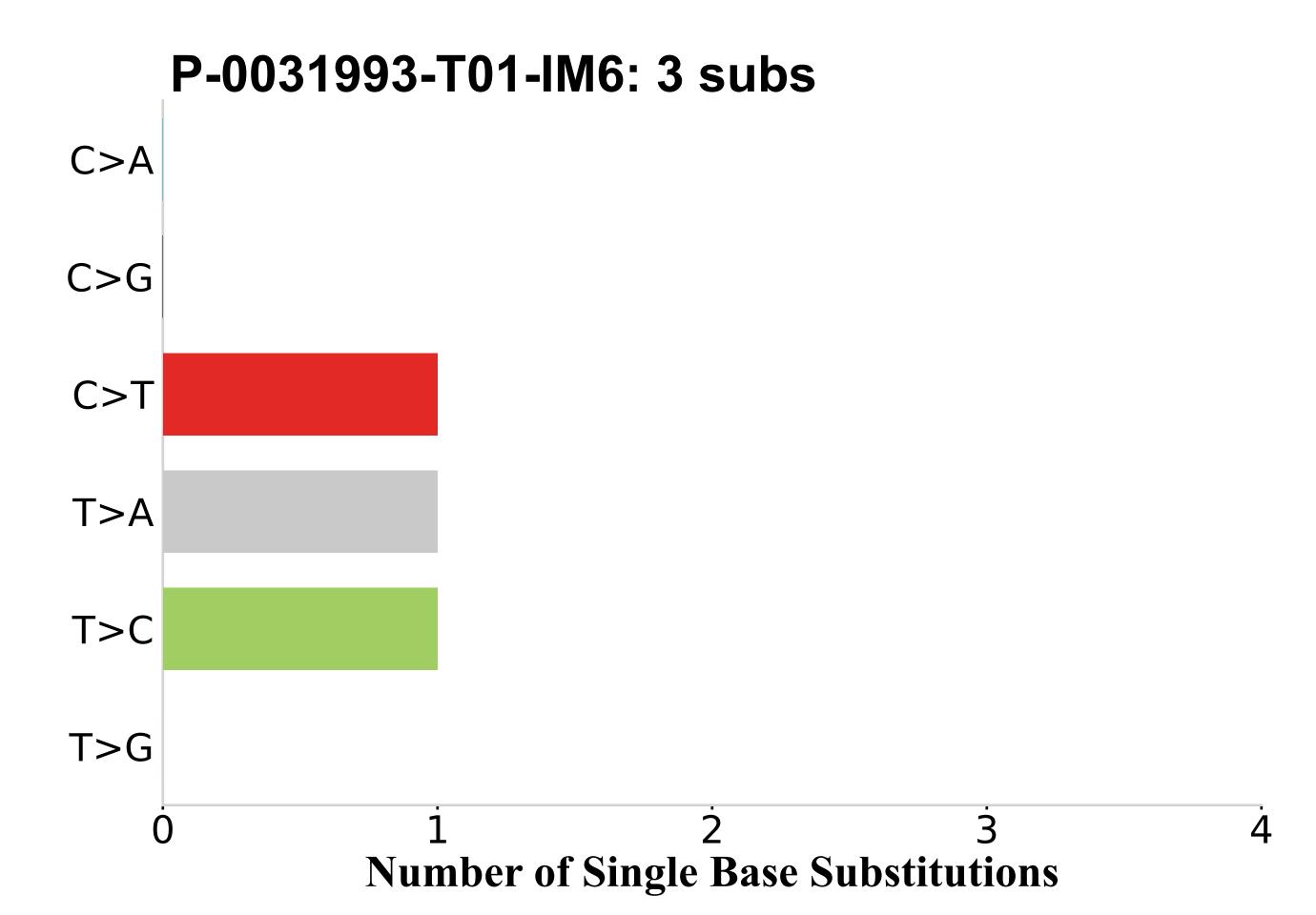








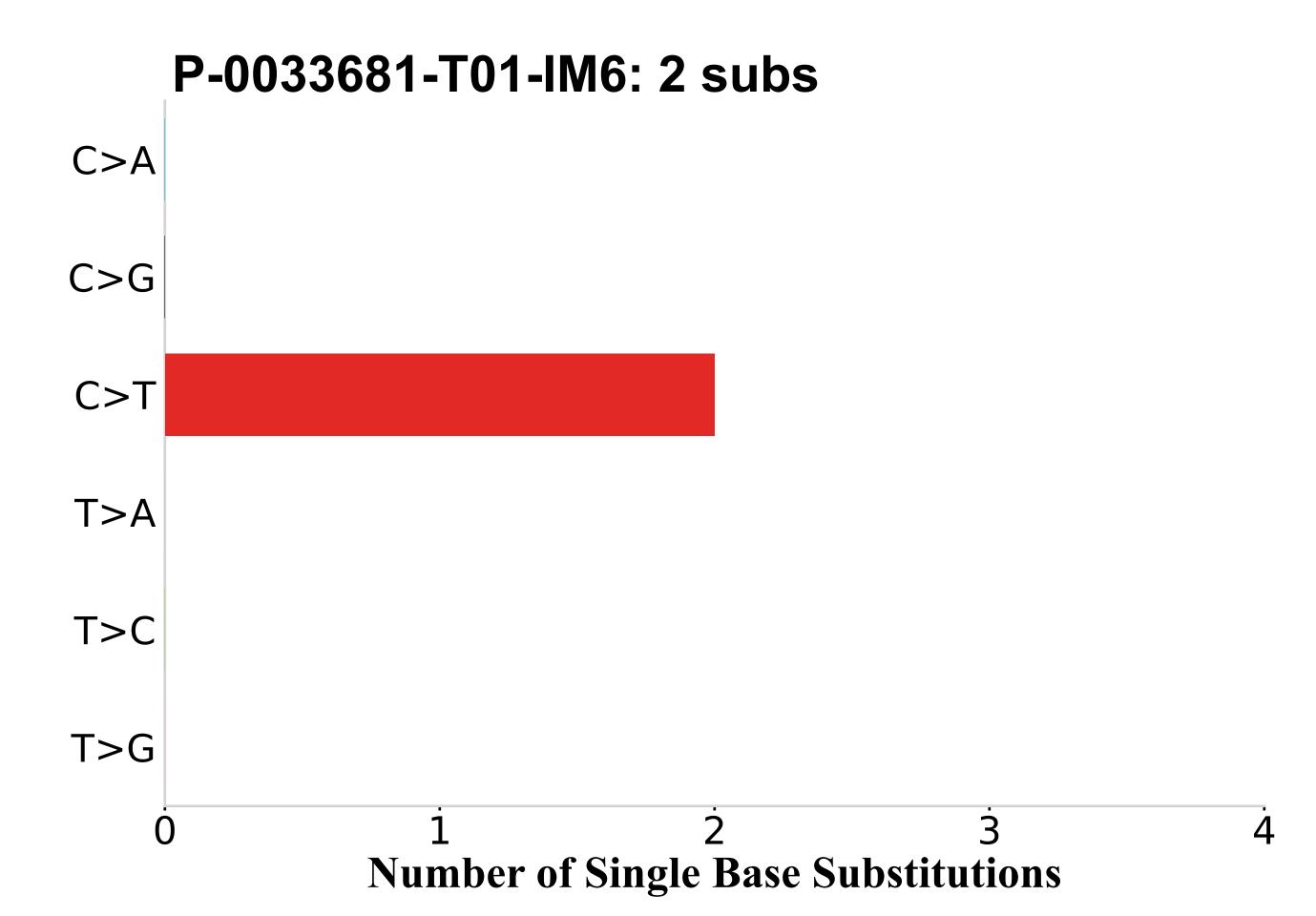


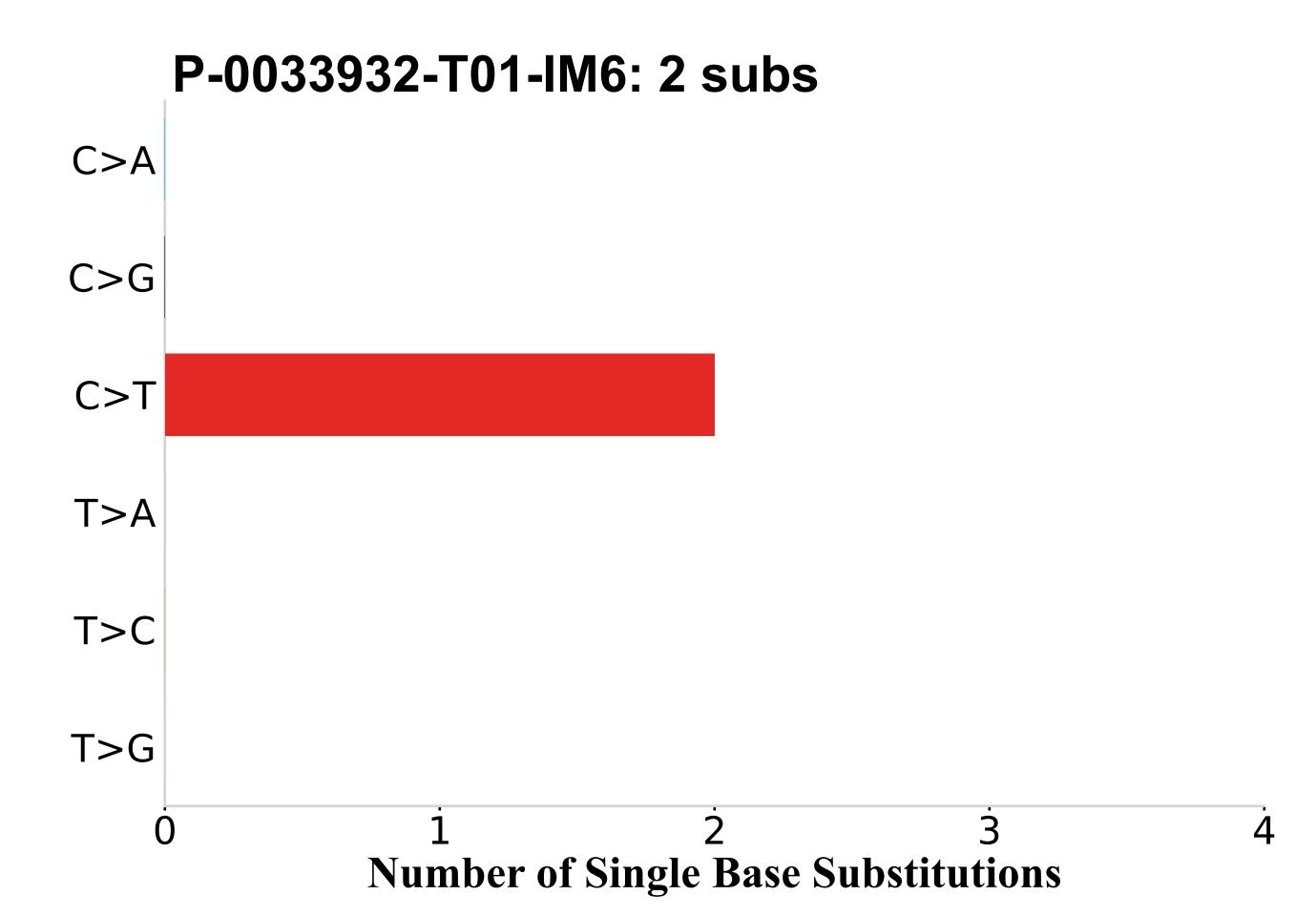


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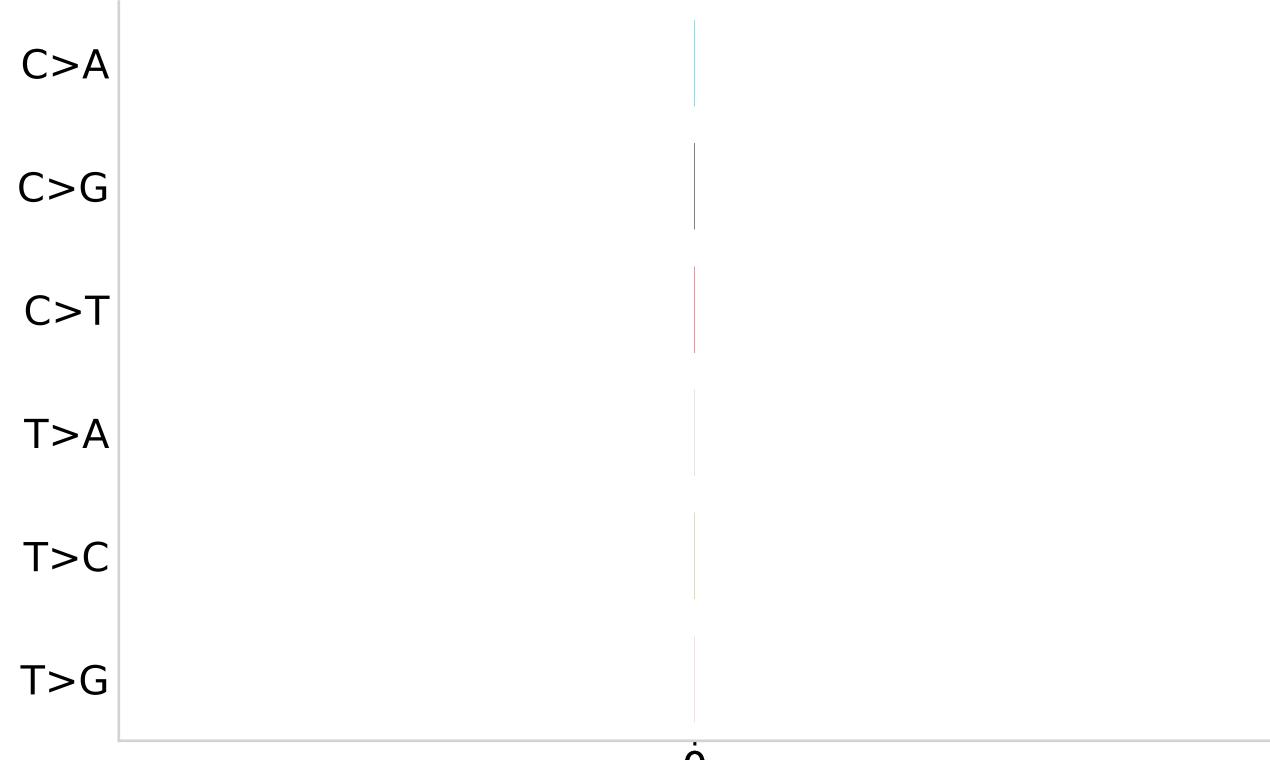


Number of Single Base Substitutions





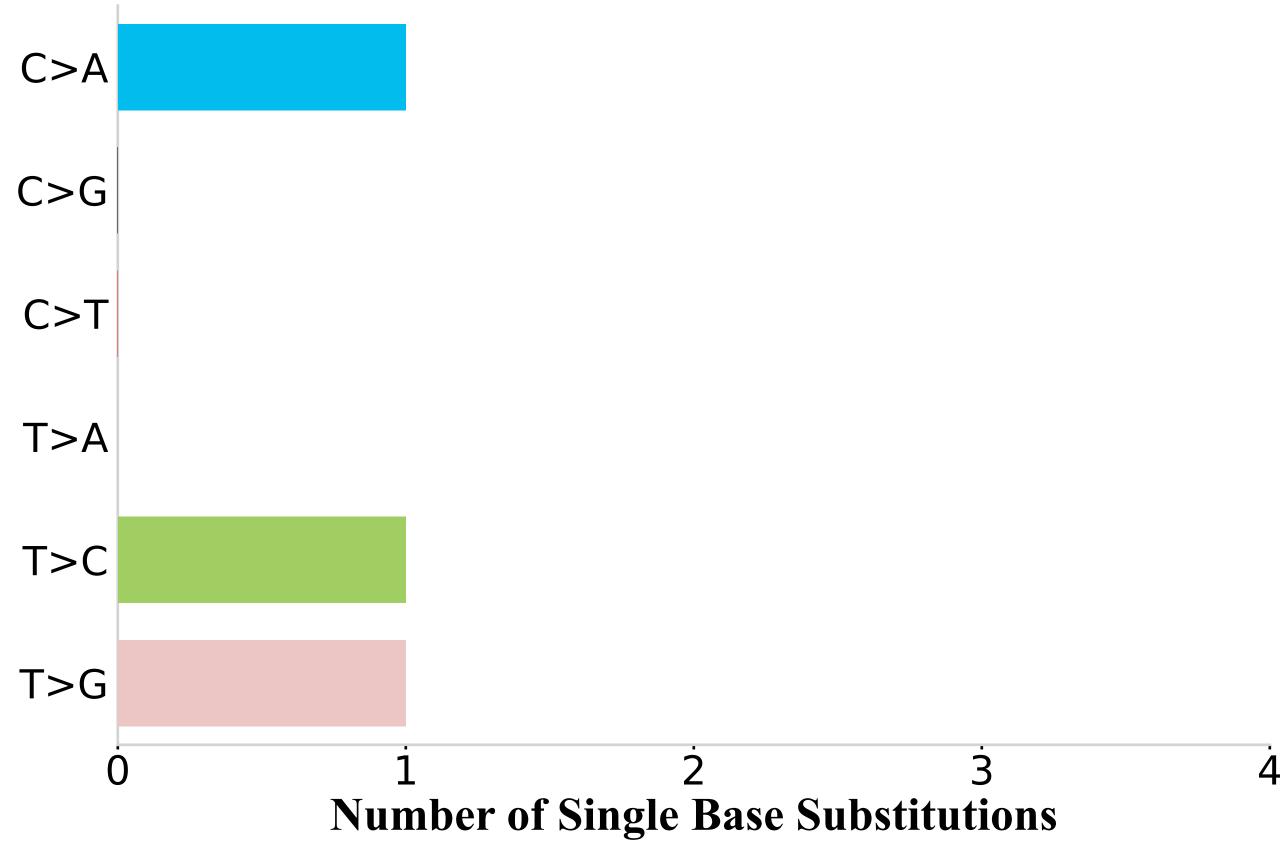
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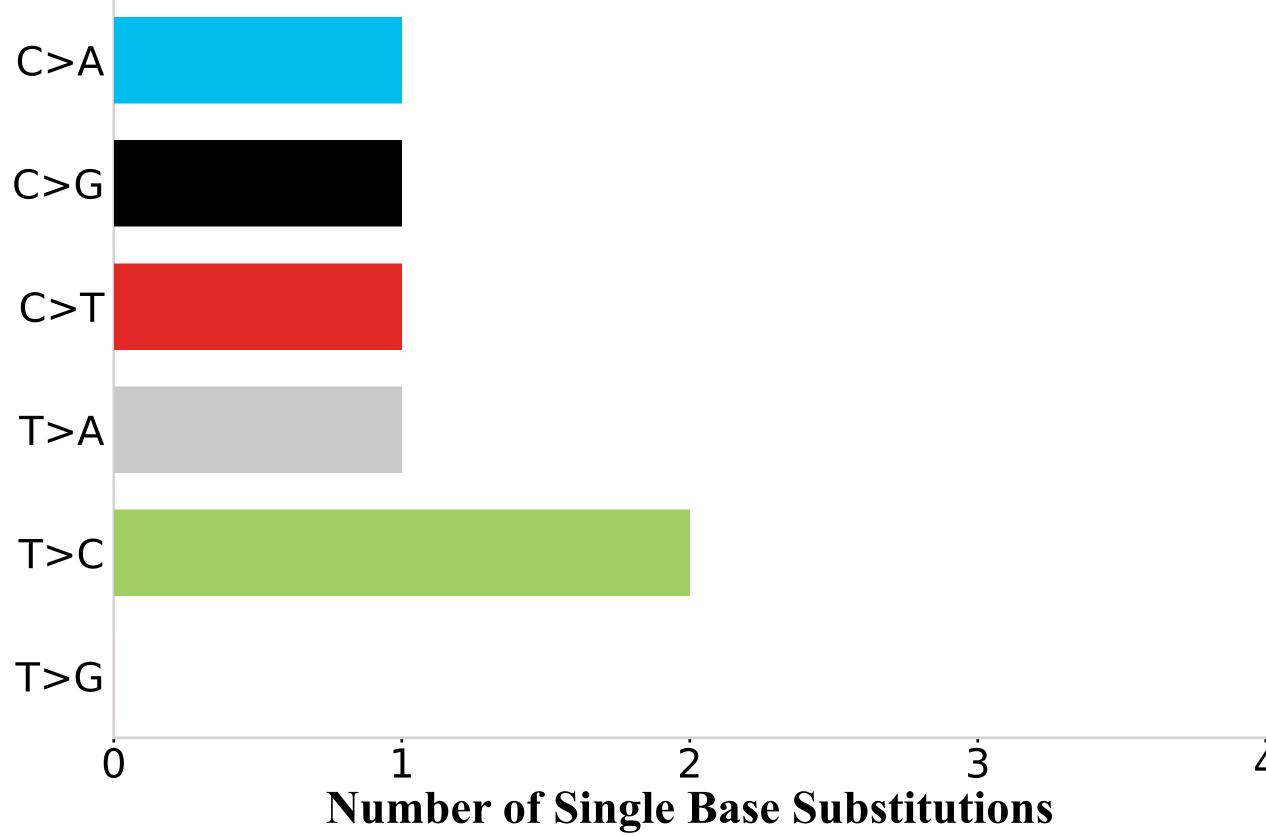
Number of Single Base Substitutions

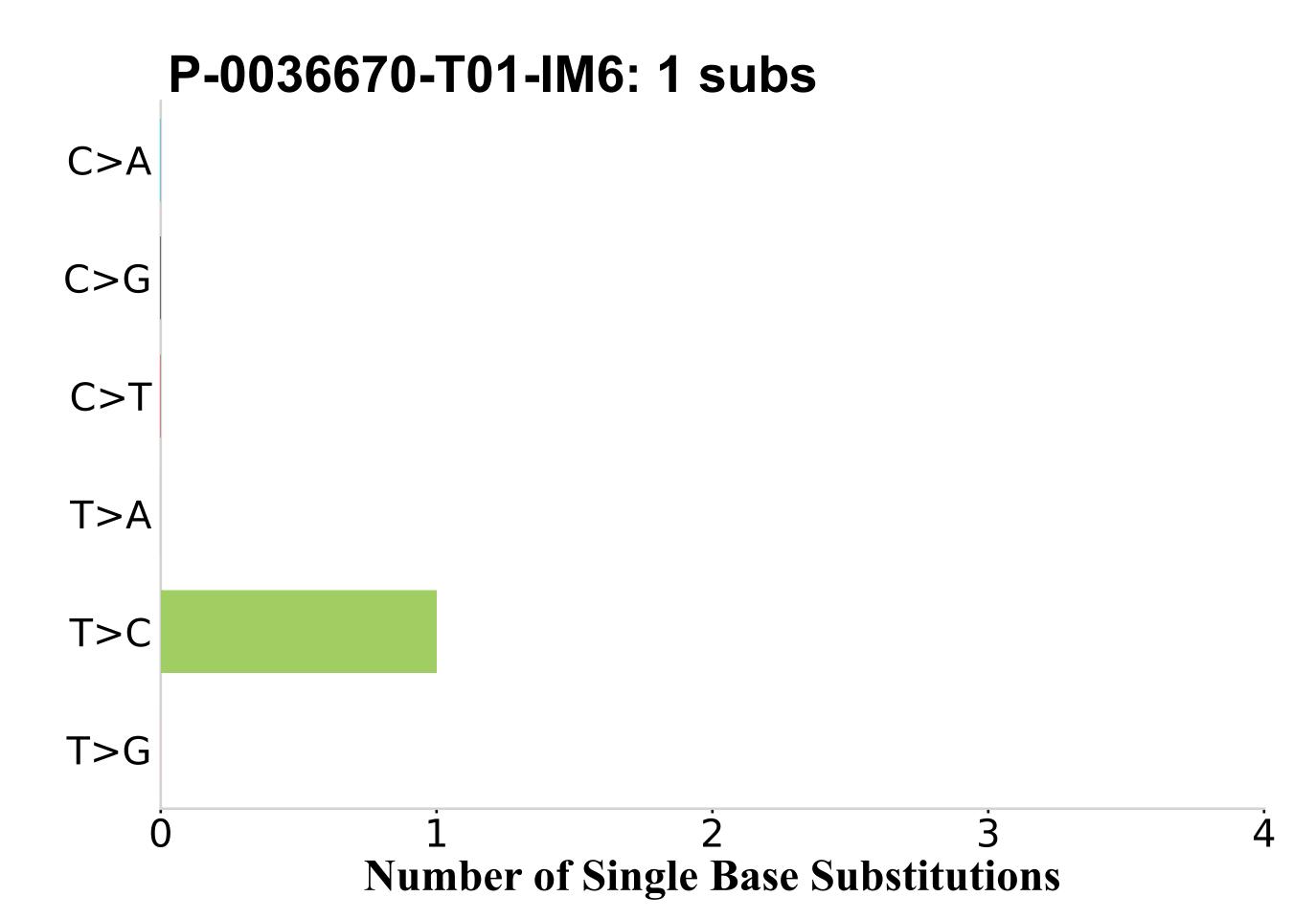
P-0035775-T01-IM6: 3 subs C>A C>G C>T T>A T>C T>G **Number of Single Base Substitutions**

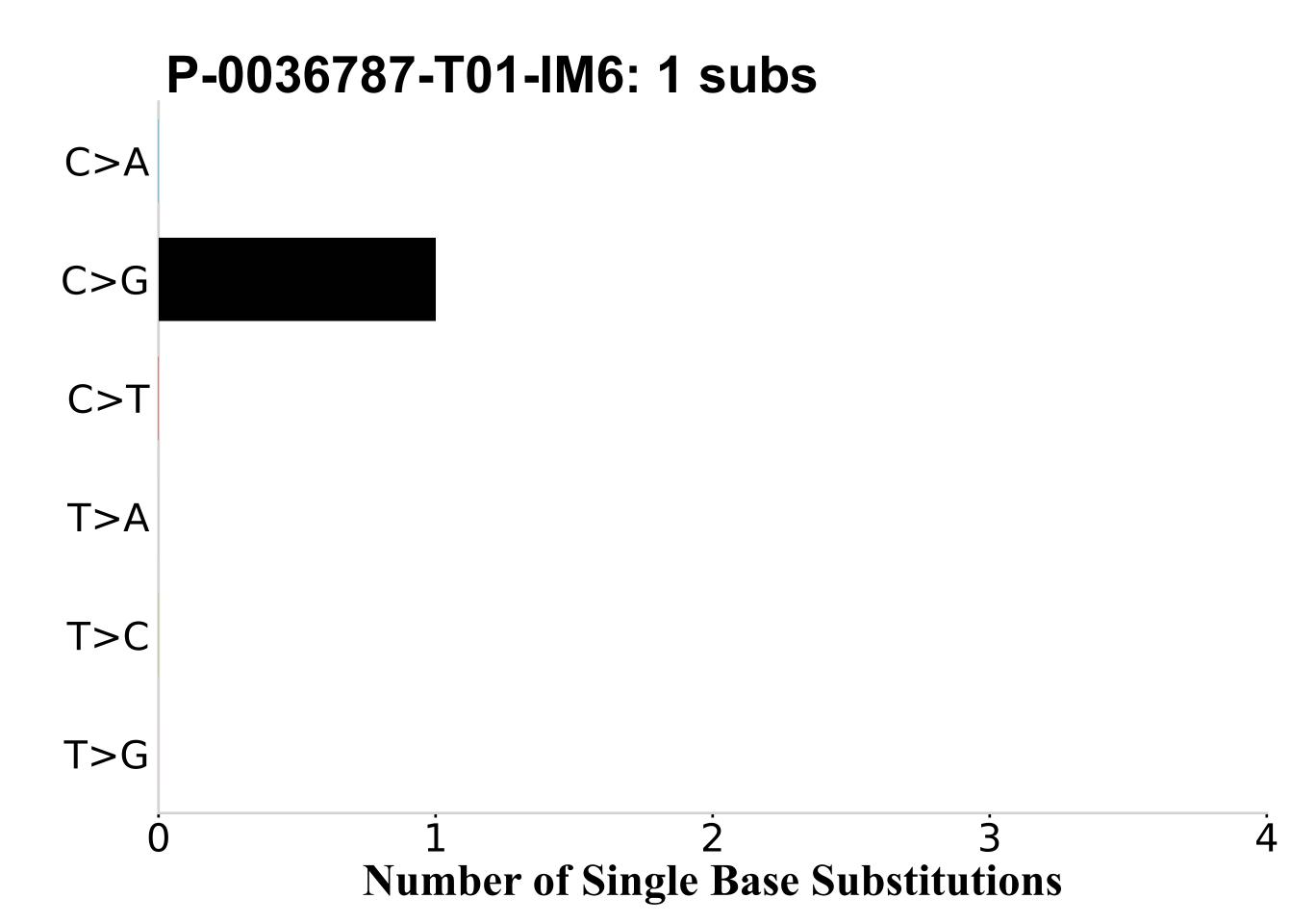


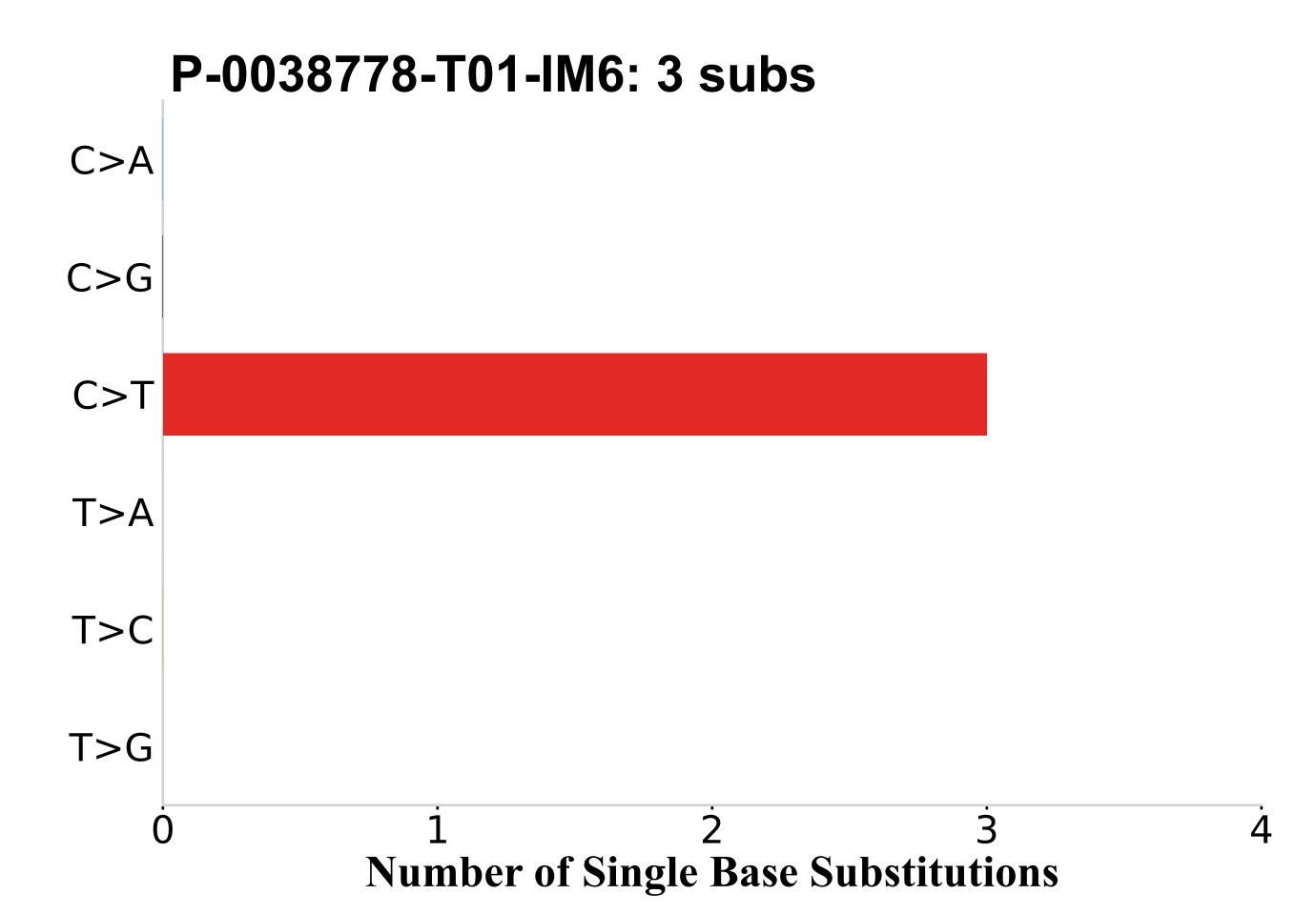


P-0036669-T01-IM6: 6 subs





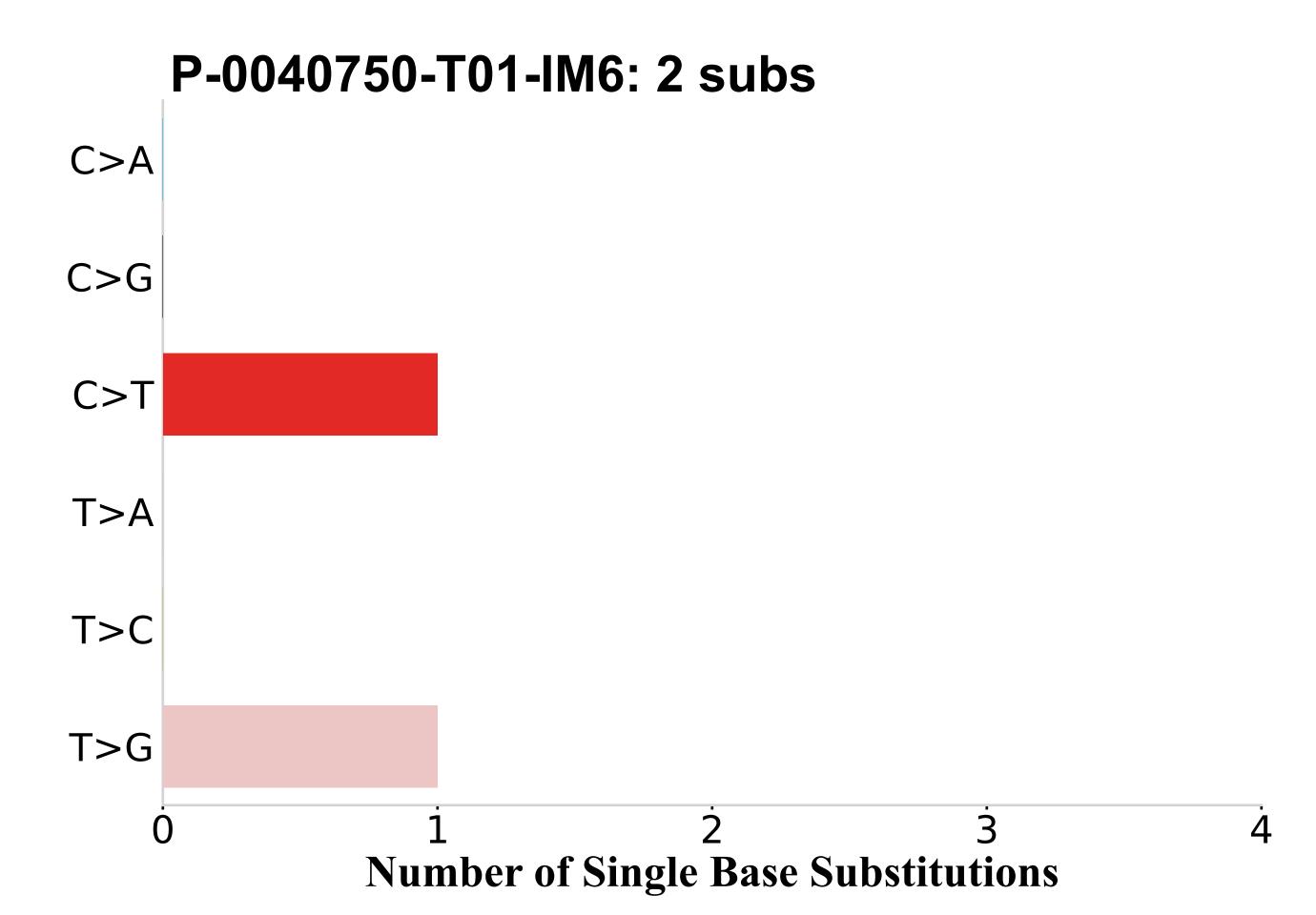


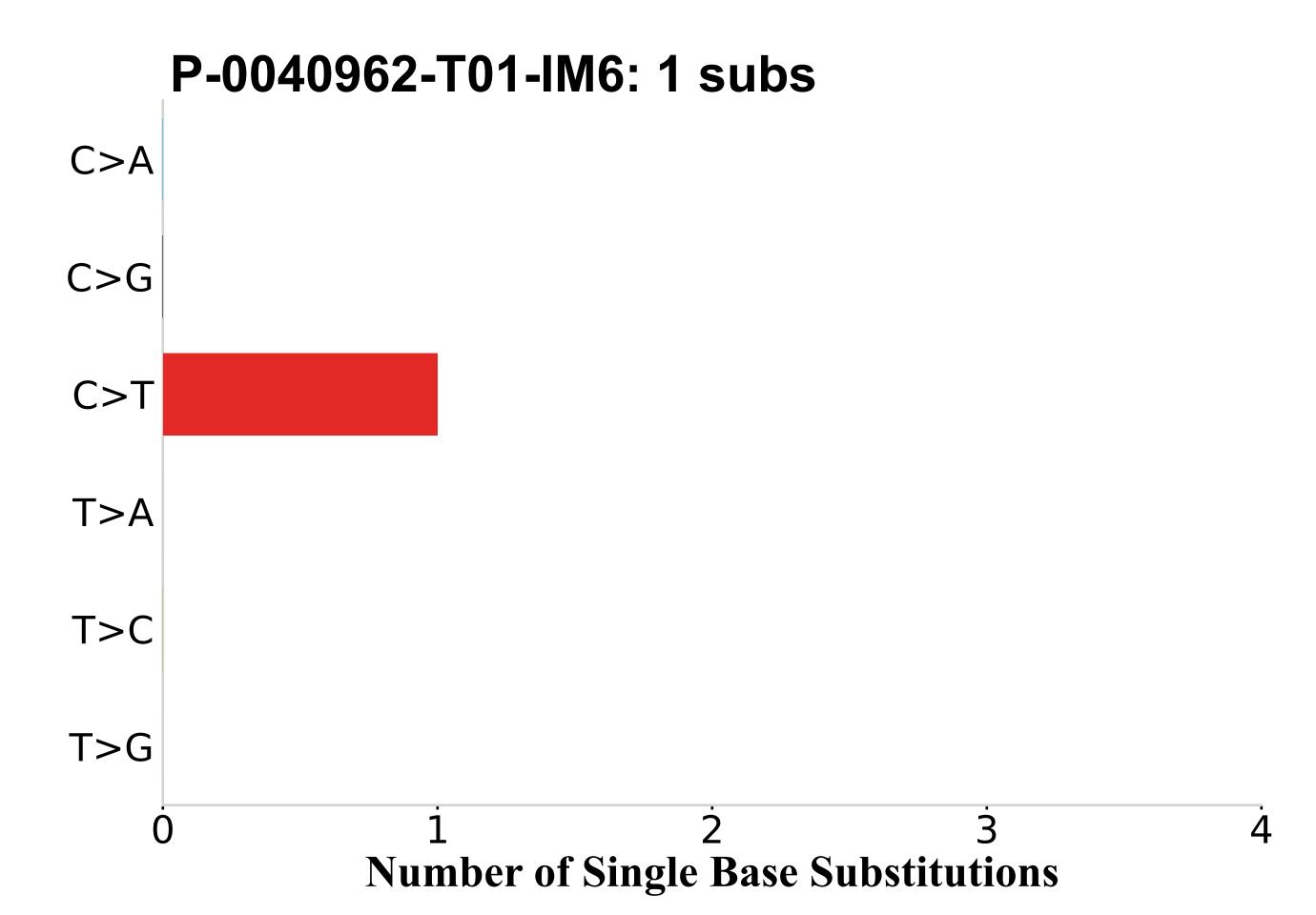


P-0040422-T02-IM6: 0 subs

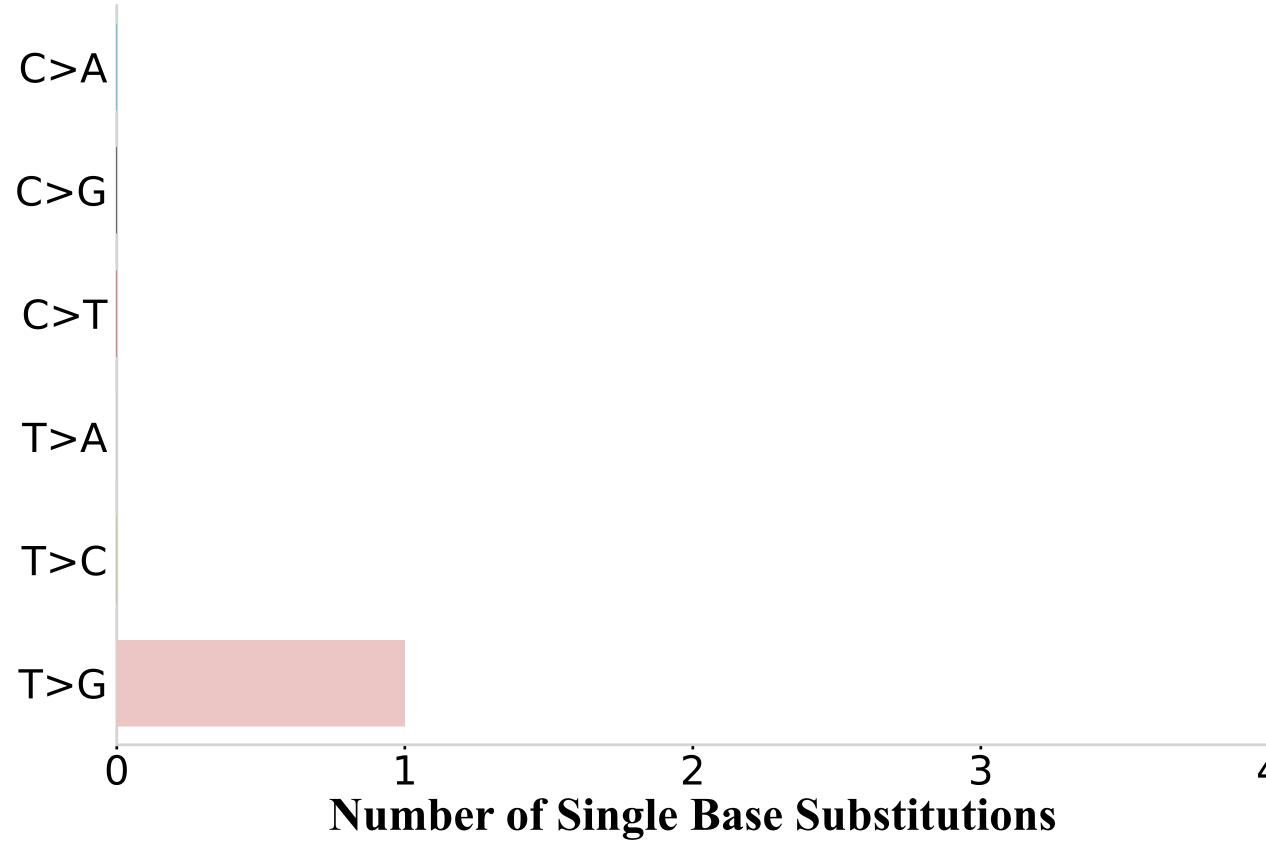


Number of Single Base Substitutions

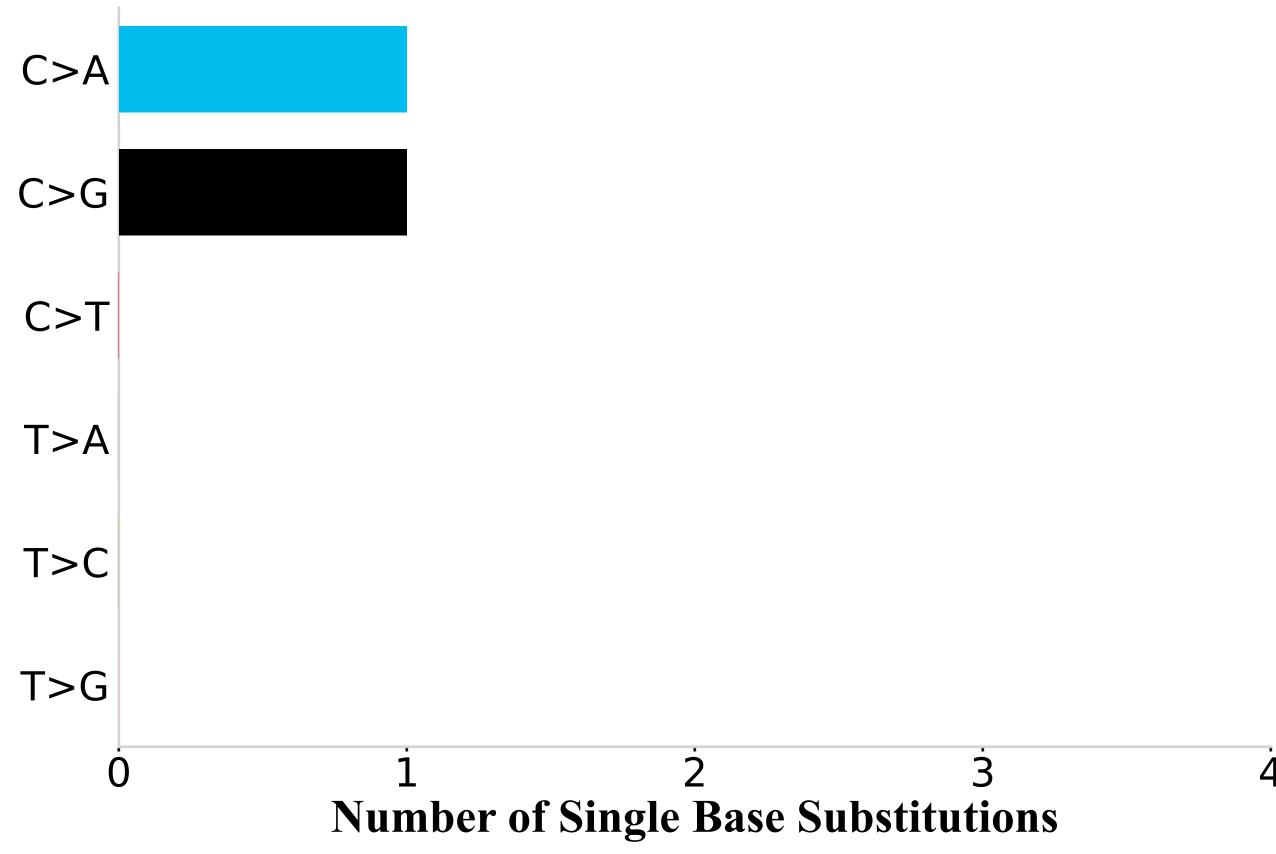


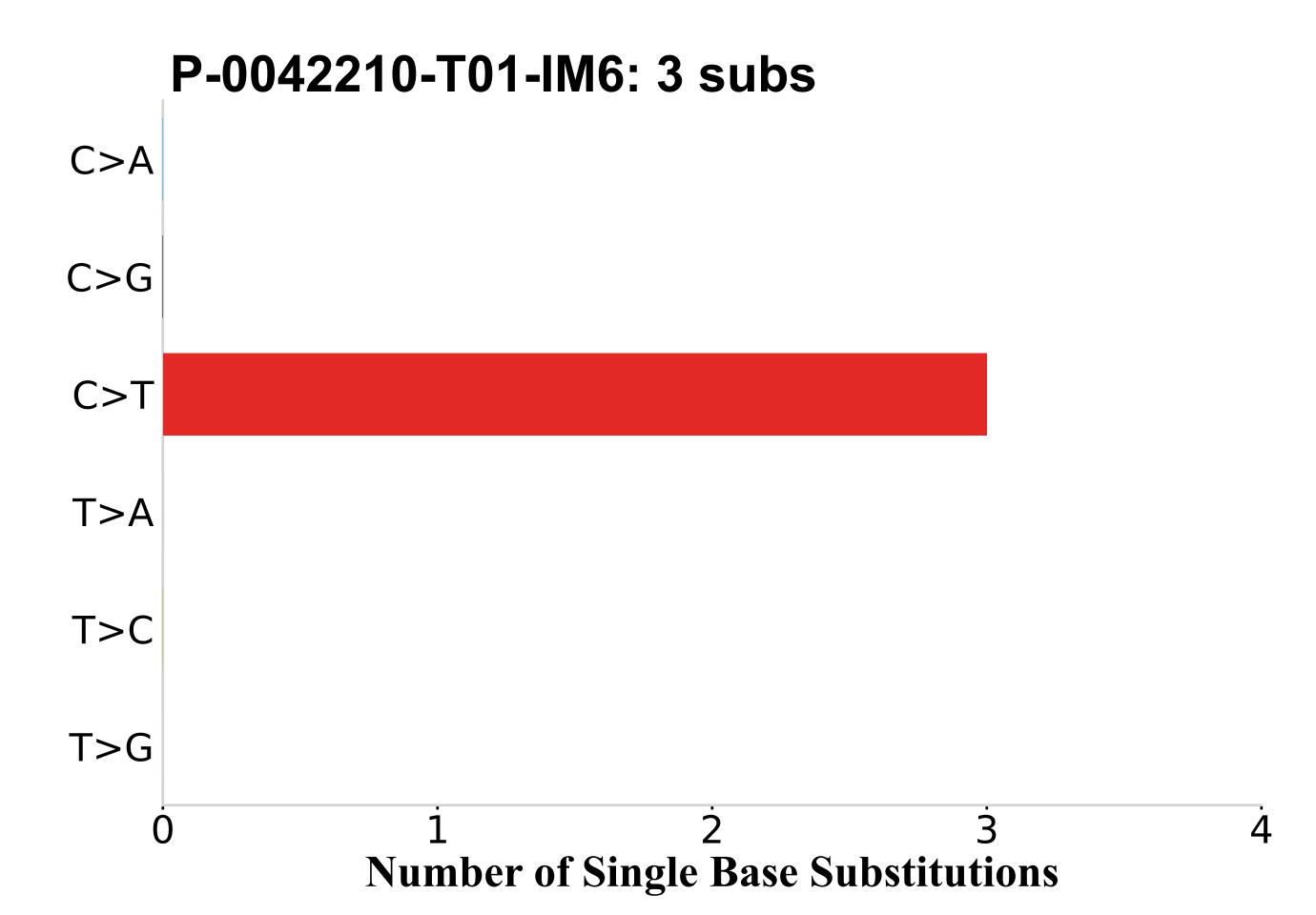


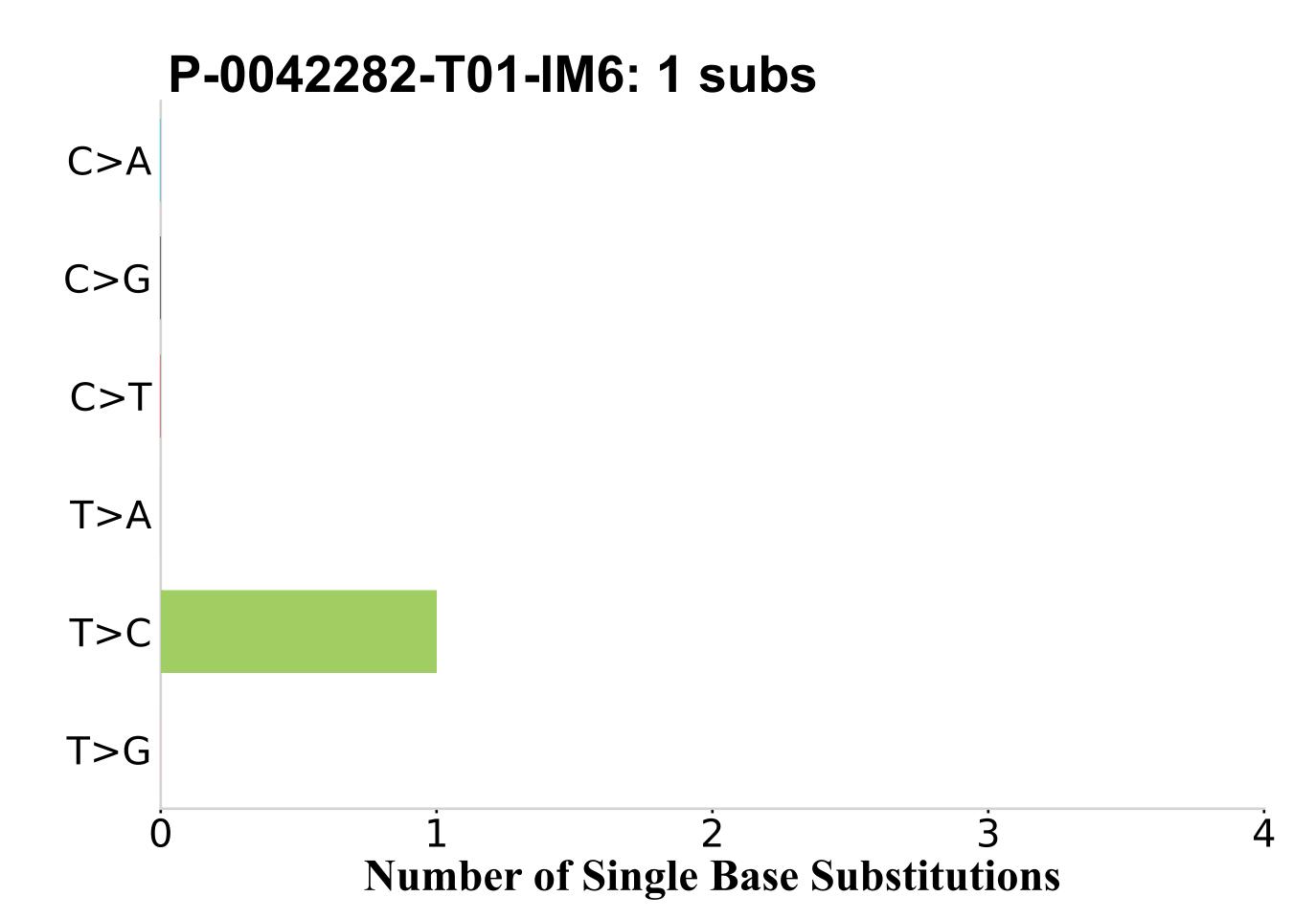




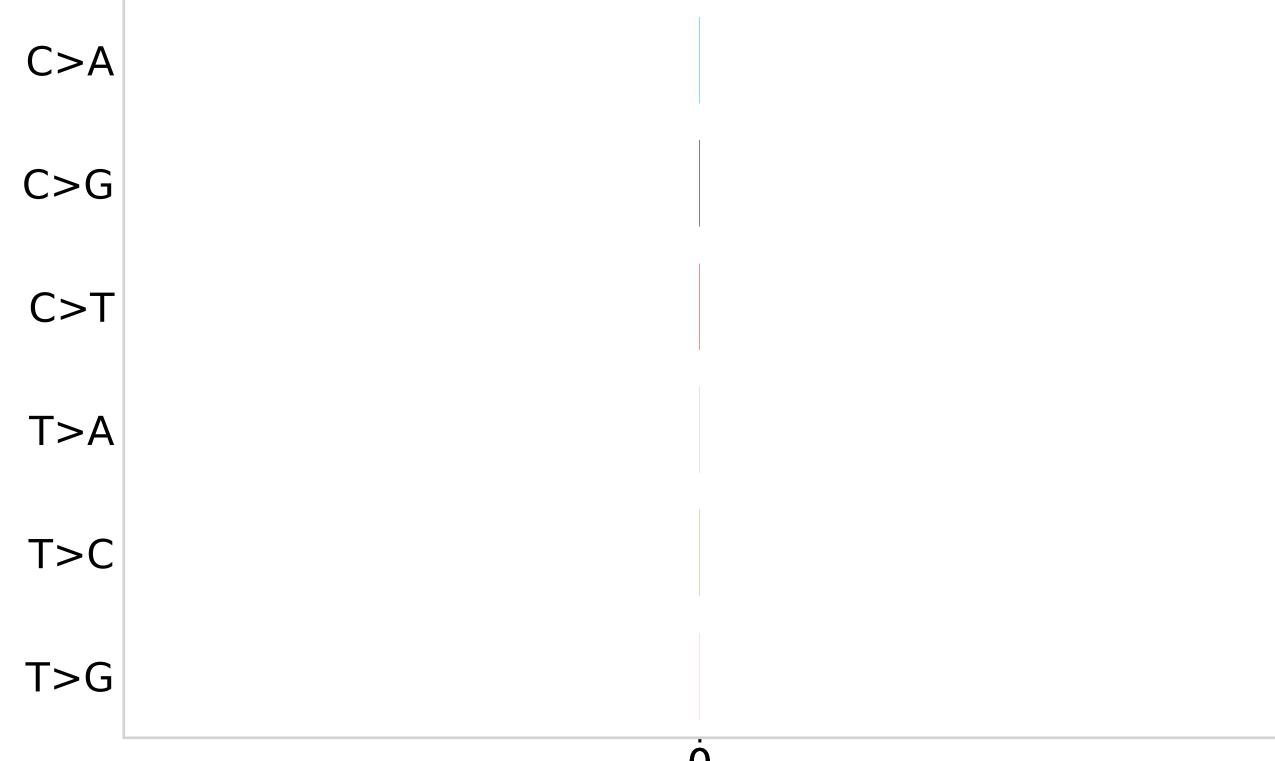
P-0041646-T01-IM6: 2 subs



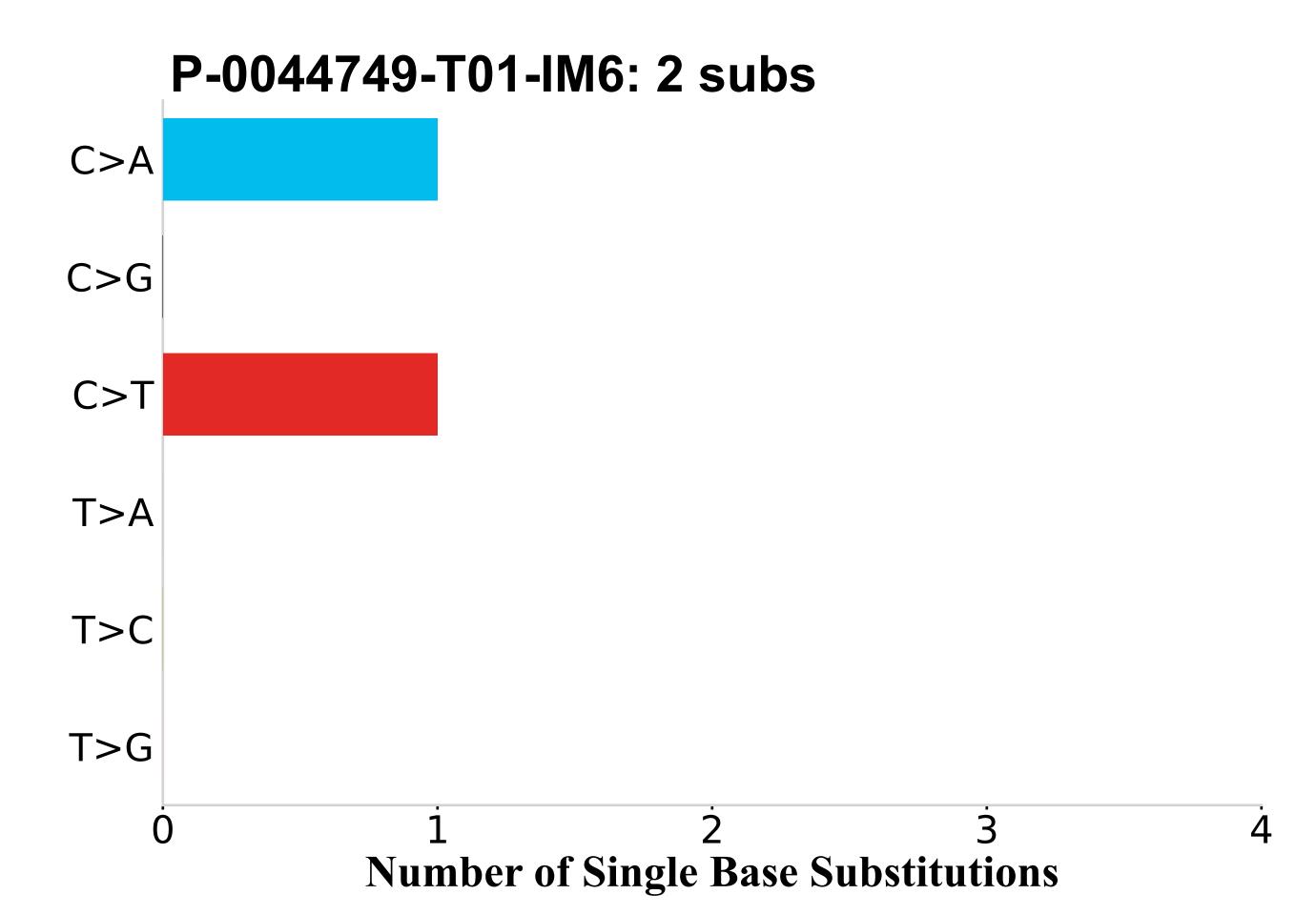




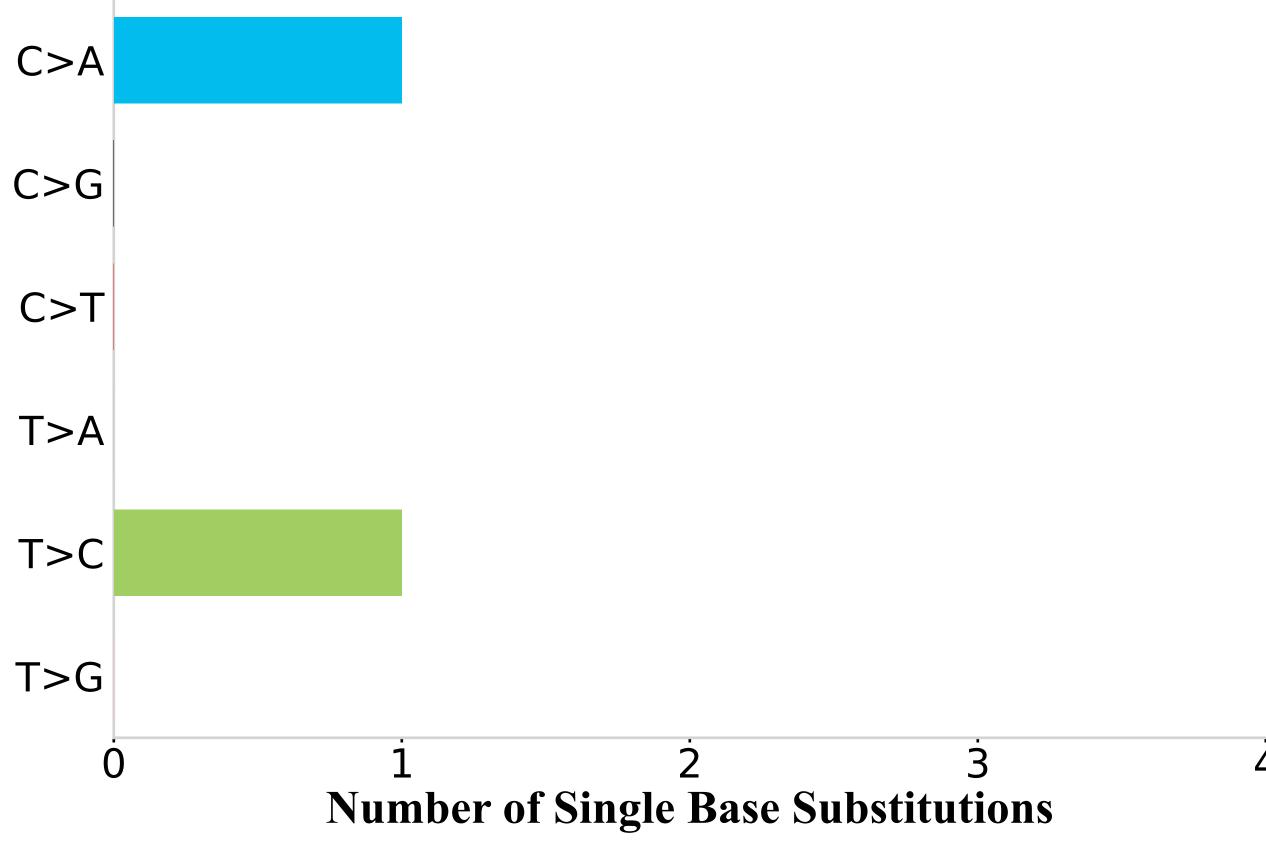
P-0043104-T01-IM6: 0 subs

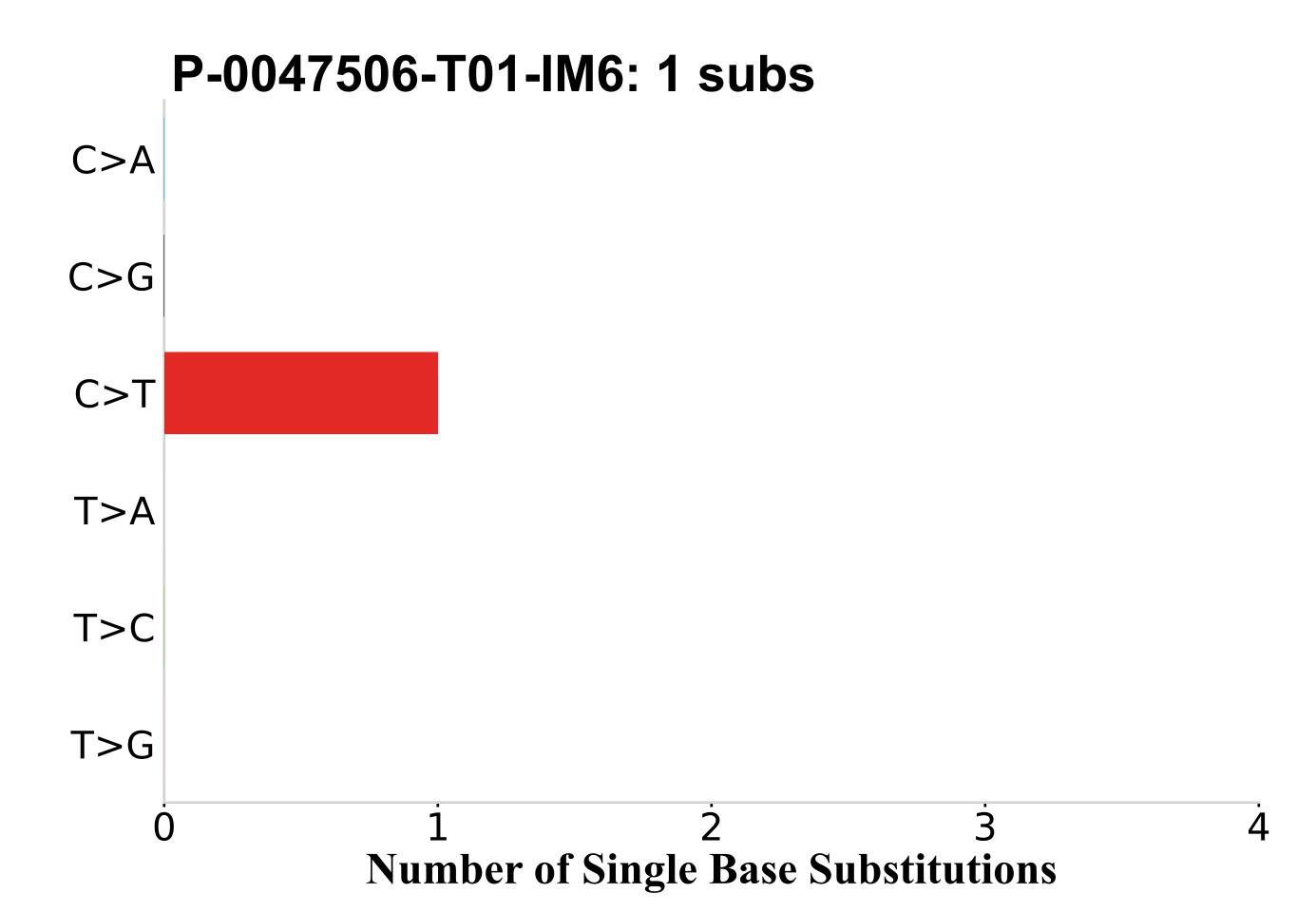


Number of Single Base Substitutions



P-0046481-T01-IM6: 2 subs





P-0049196-T01-IM6: 3 subs C>A C>G C>T T>A T>C T>G **Number of Single Base Substitutions**

