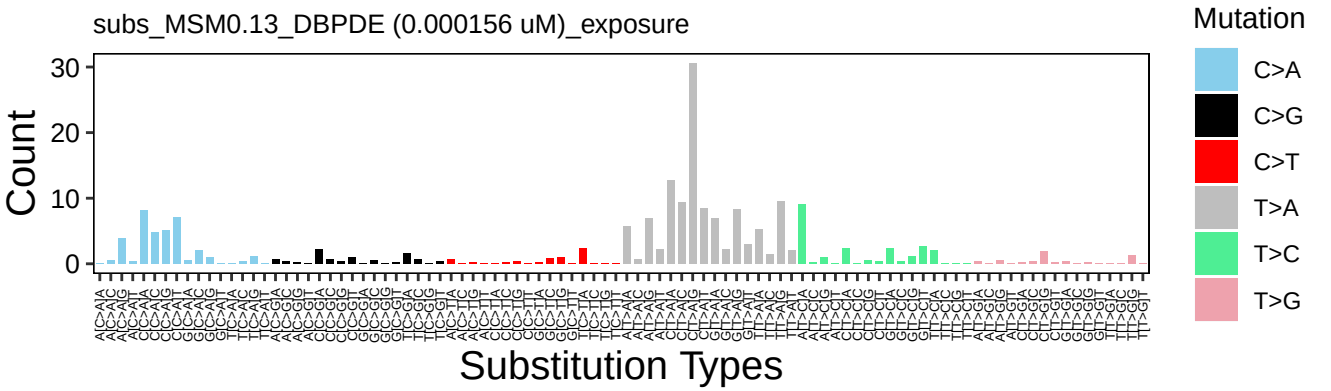
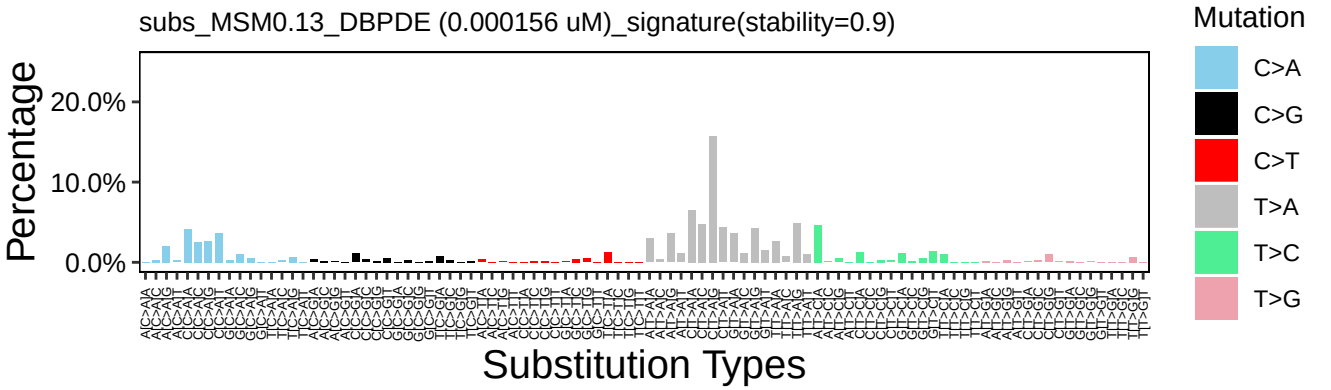

subs_MSM0.13_DBPDE (0.000156 uM)_exposure
Count
30
20
10
0
Substitution Types
Mutation
C>A
C>G
C>T
T>A
T>C
T>G


subs_MSM0.13_DBPDE (0.000156 uM)_signature(stability=0.9)
Percentage
20.0%
10.0%
0.0%
Substitution Types
Mutation
C>A
C>G
C>T
T>A
T>C
T>G