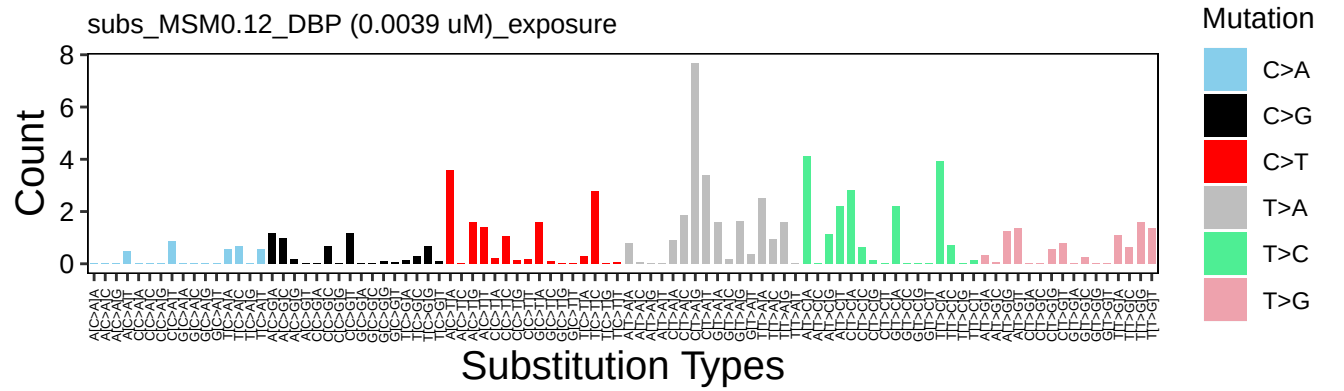
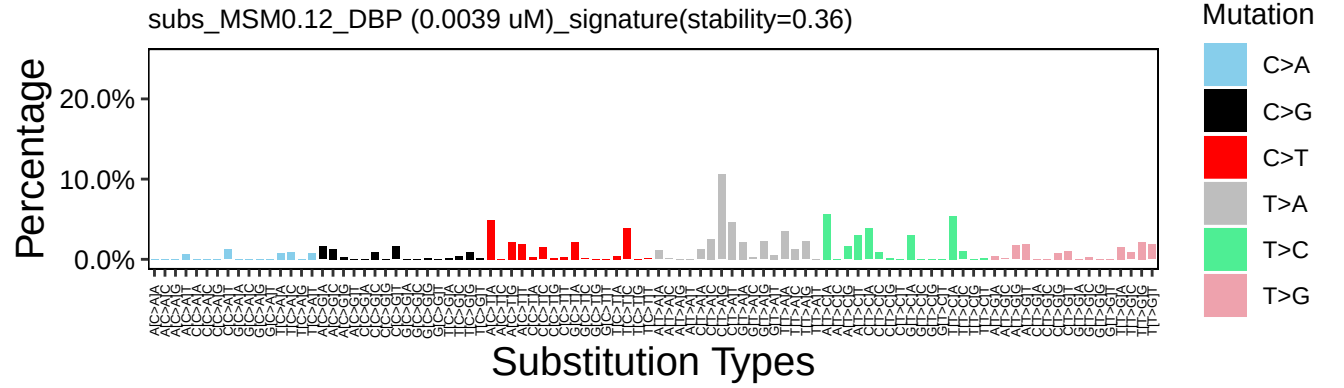

subs_MSM0.12_DBP (0.0039 uM)_exposure
Count
Substitution Types
Mutation
C>A
C>G
C>T
T>A
T>C
T>G


subs_MSM0.12_DBP (0.0039 uM)_signature(stability=0.36)
Percentage
Substitution Types
Mutation
C>A
C>G
C>T
T>A
T>C
T>G