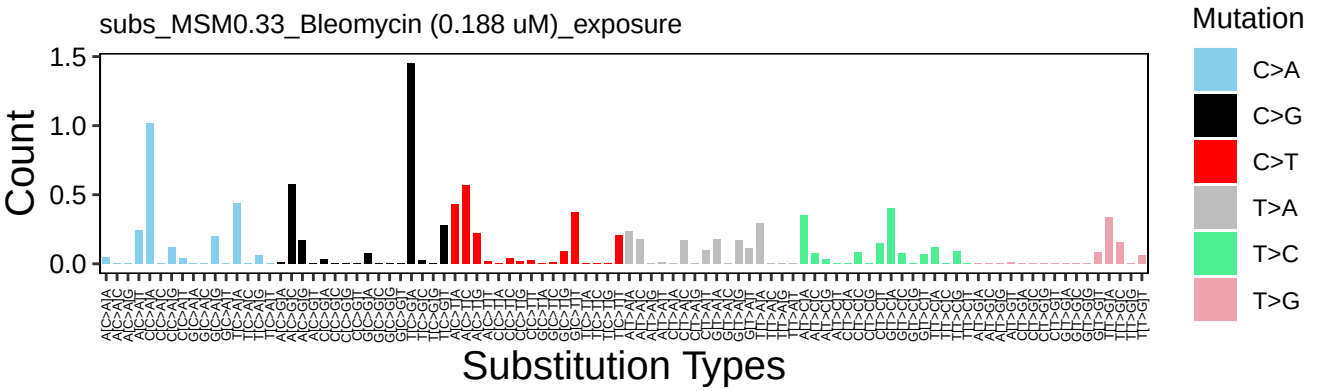
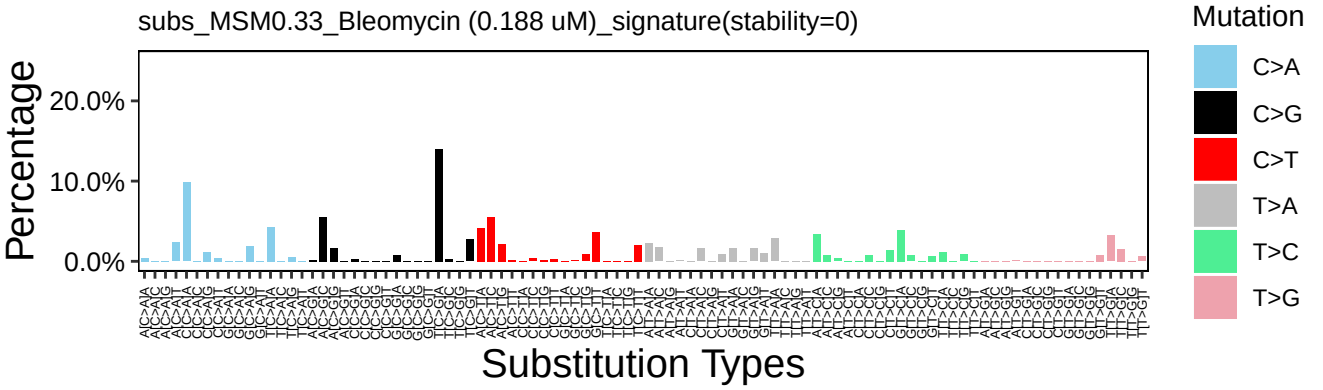

subs_MSM0.33_Bleomycin (0.188 uM)_exposure
Count
1.5
1.0
0.5
0.0
Substitution Types
Mutation
C>A
C>G
C>T
T>A
T>C
T>G


subs_MSM0.33_Bleomycin (0.188 uM)_signature(stability=0)
Percentage
20.0%
10.0%
0.0%
Substitution Types
Mutation
C>A
C>G
C>T
T>A
T>C
T>G