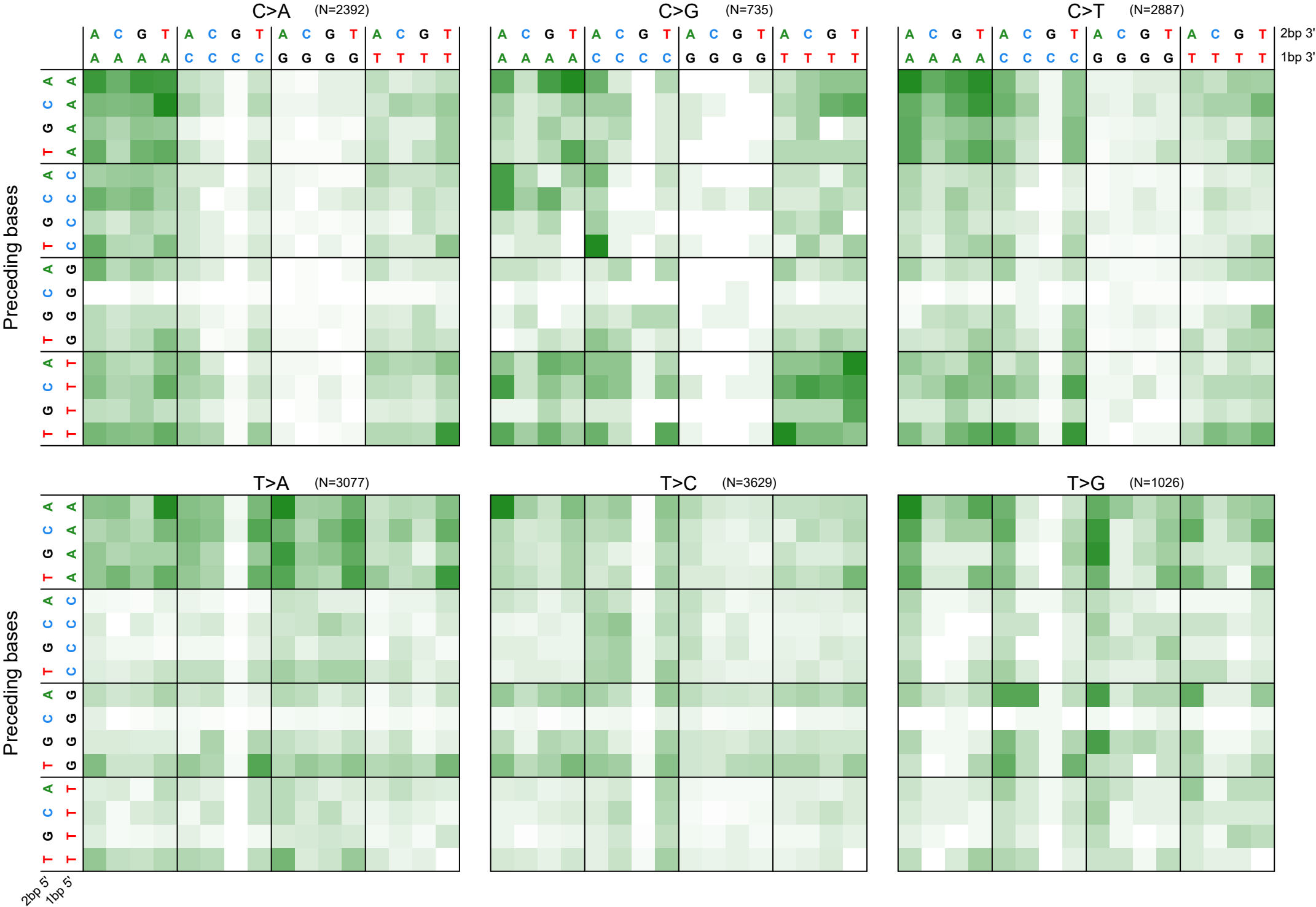


NDEA_cl1



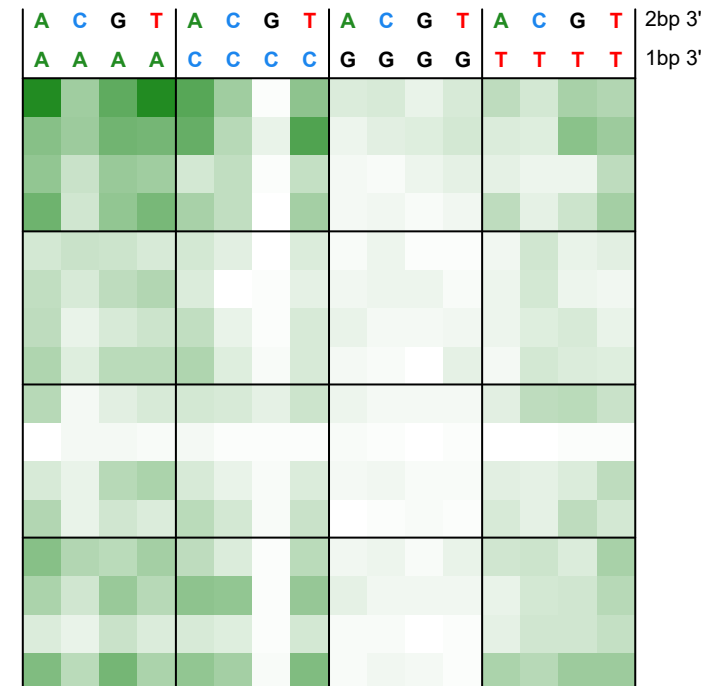
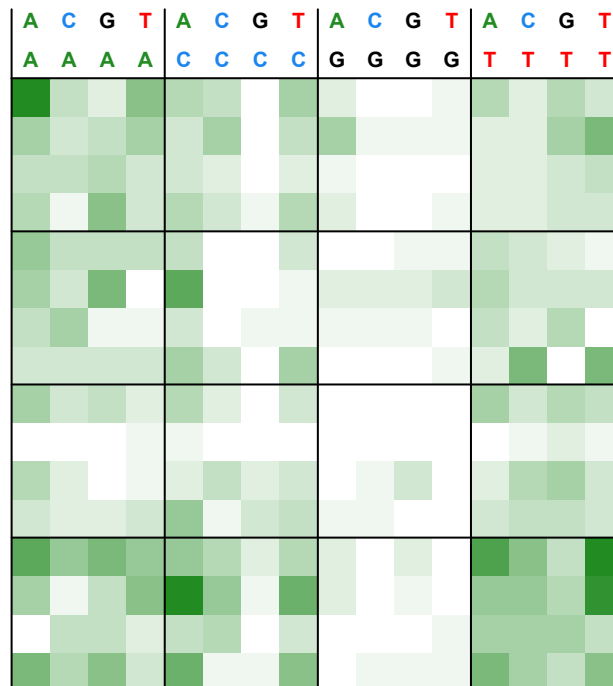
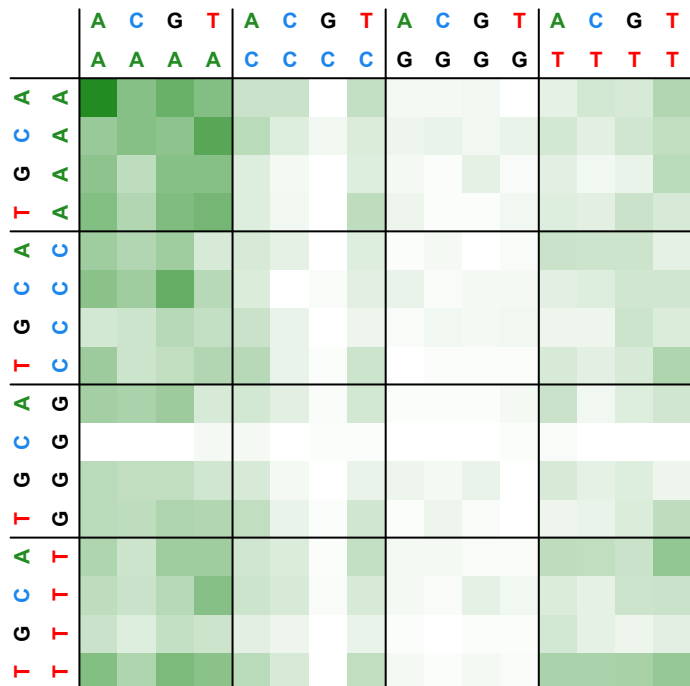
NDEA_cl2

C>A (N=2865)

C>G (N=804)

C>T (N=3157)

Preceding bases

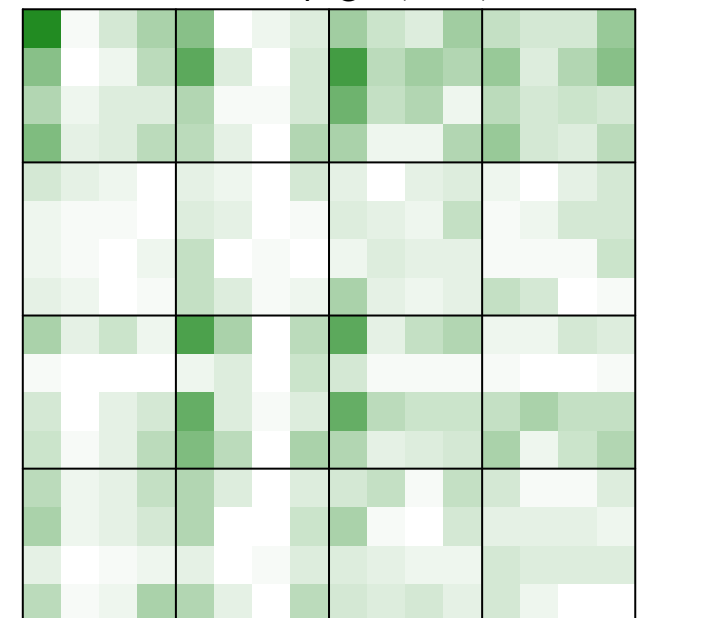
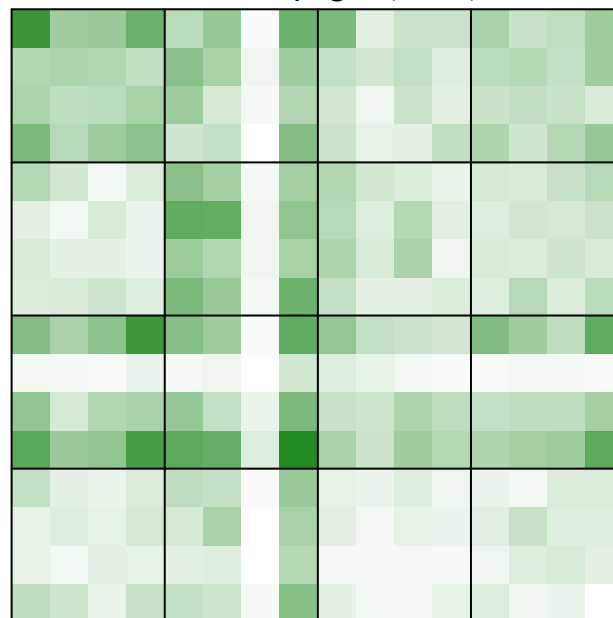
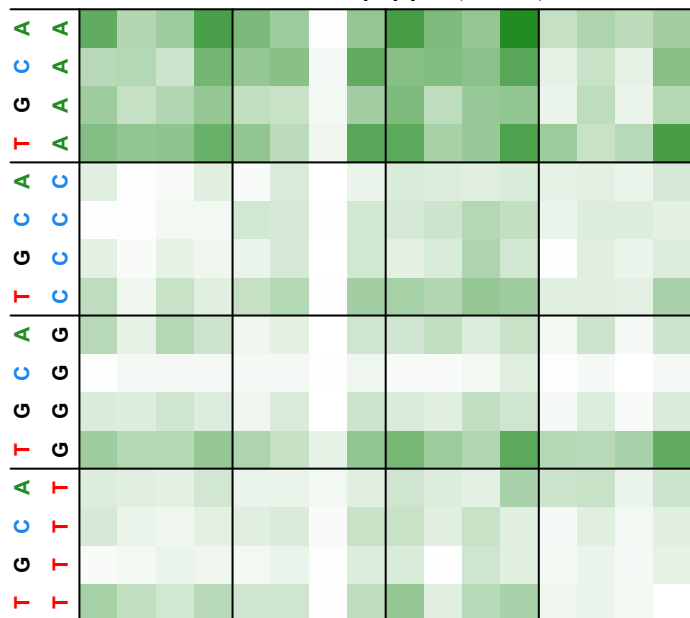


Preceding bases

T>A (N=3958)

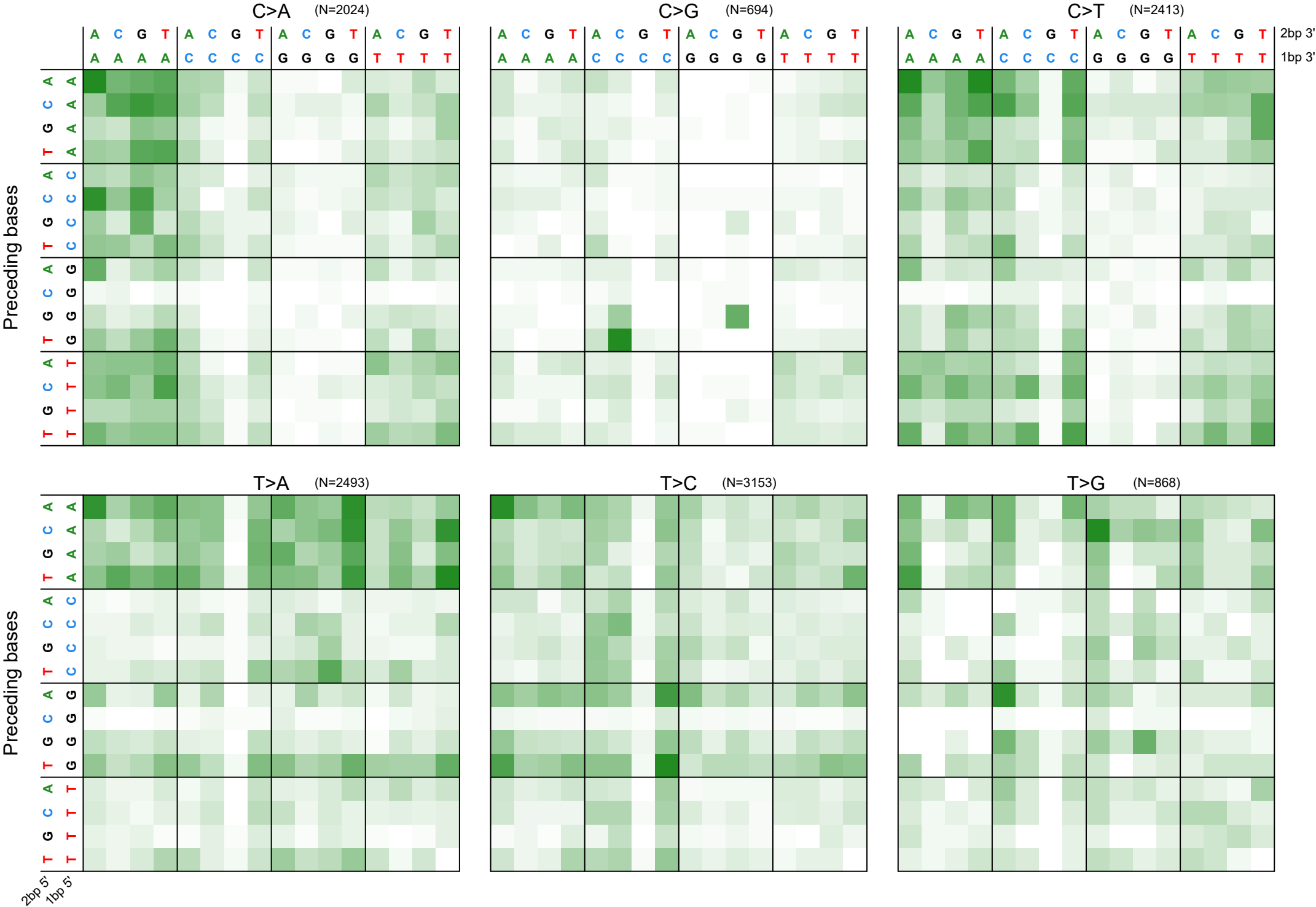
T>C (N=4596)

T>G (N=1205)



2bp 5'
1bp 5'

NDEA_cl3



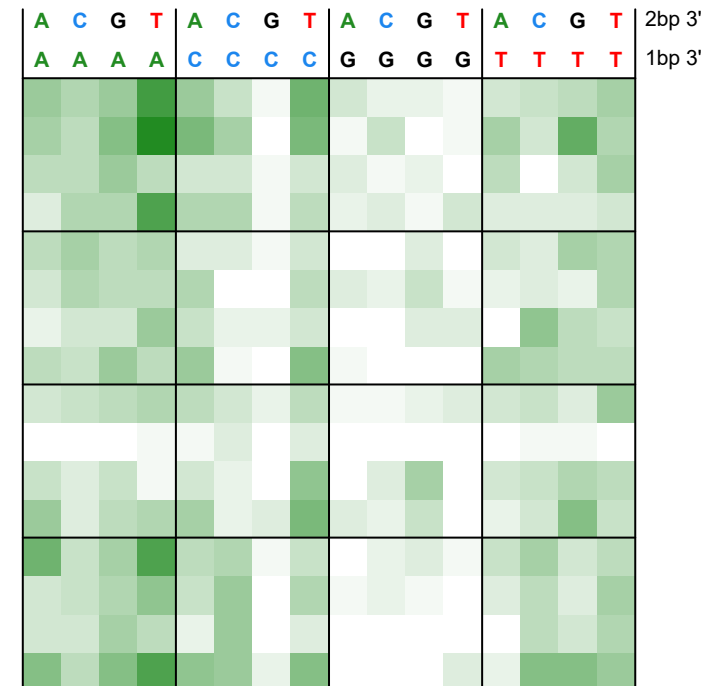
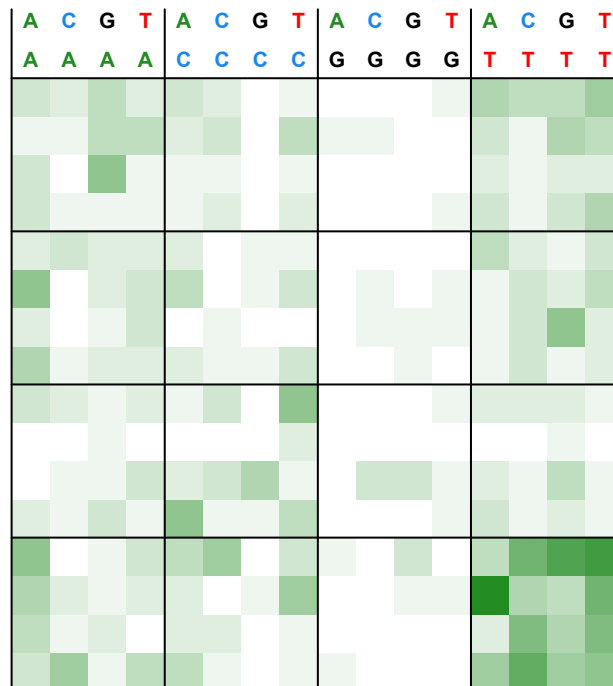
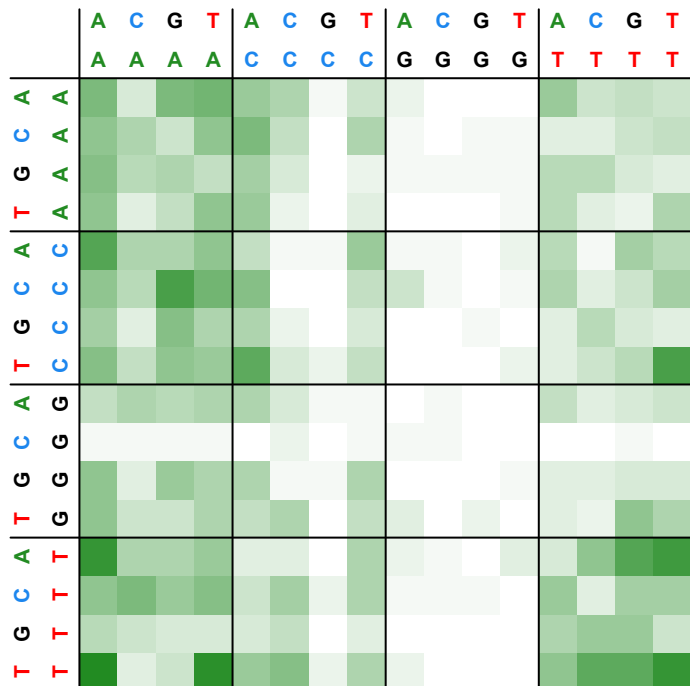
NDMA_cl1

C>A (N=1324)

C>G (N=511)

C>T (N=1167)

Preceding bases

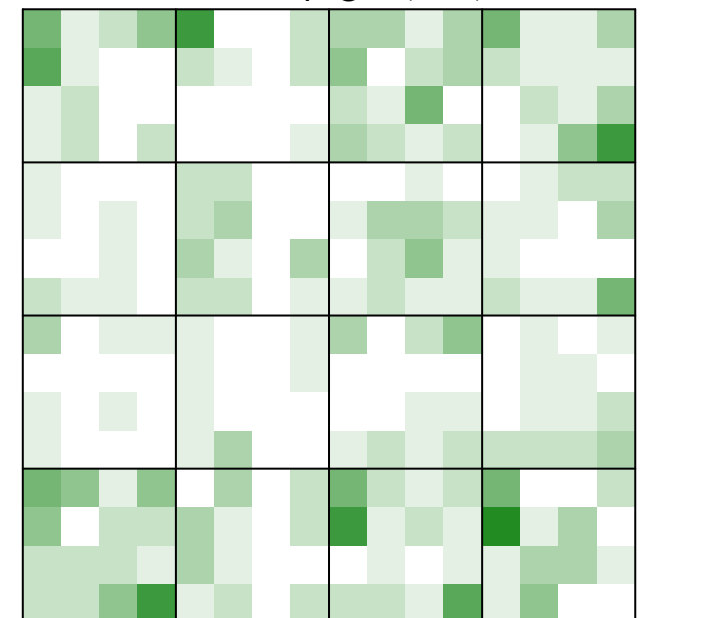
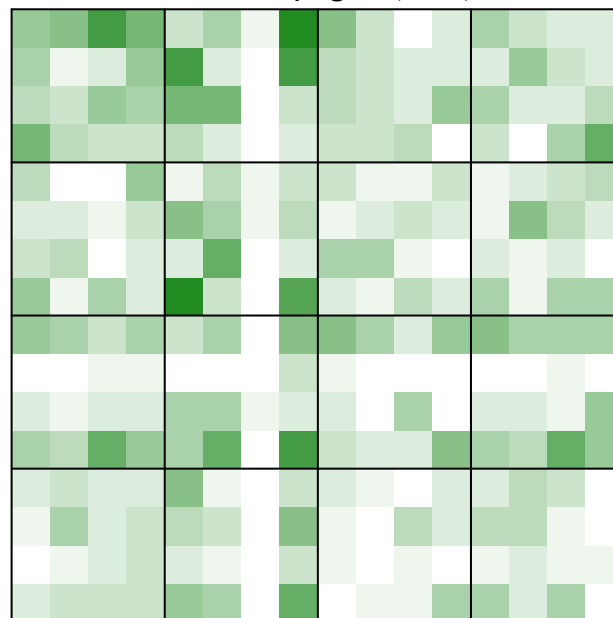
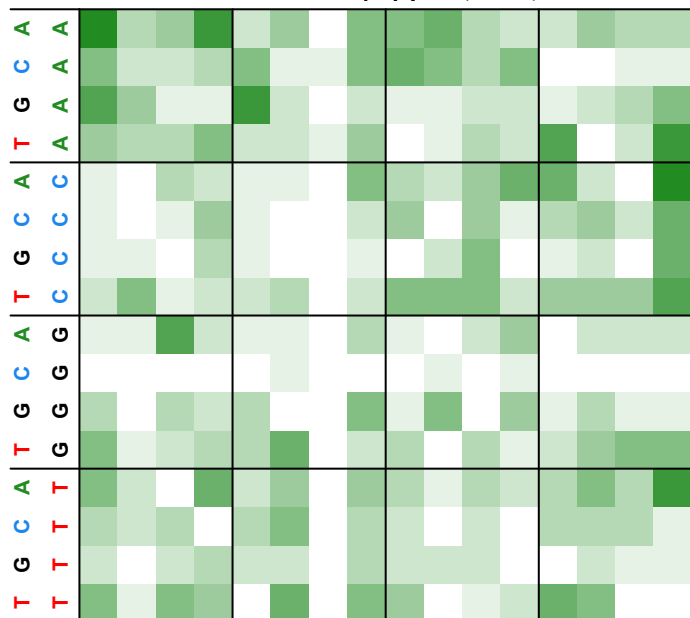


T>A (N=612)

T>C (N=797)

T>G (N=362)

Preceding bases



2bp 5'
1bp 5'

2bp 3'
1bp 3'

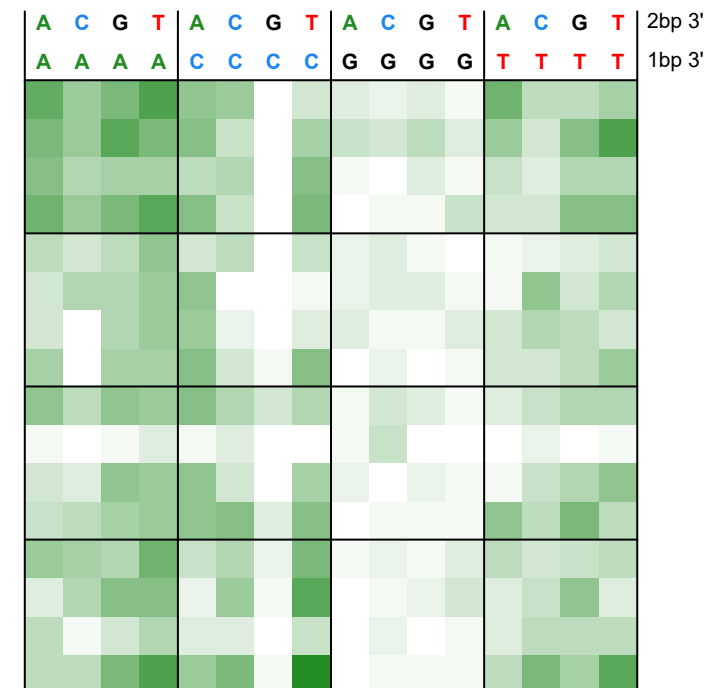
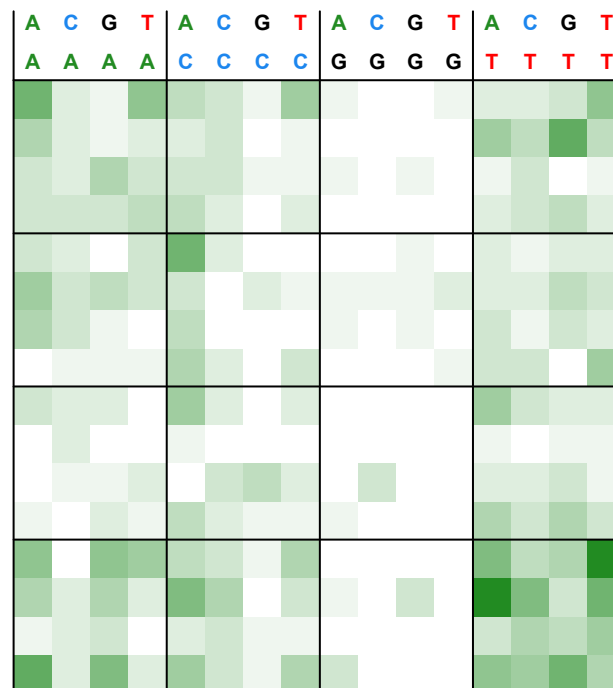
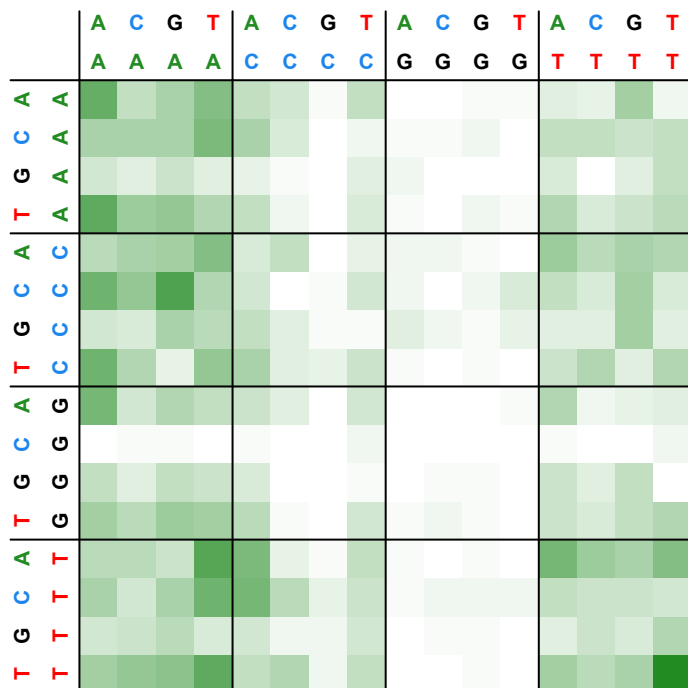
NDMA_cl2

C>A (N=1488)

C>G (N=583)

C>T (N=1357)

Preceding bases

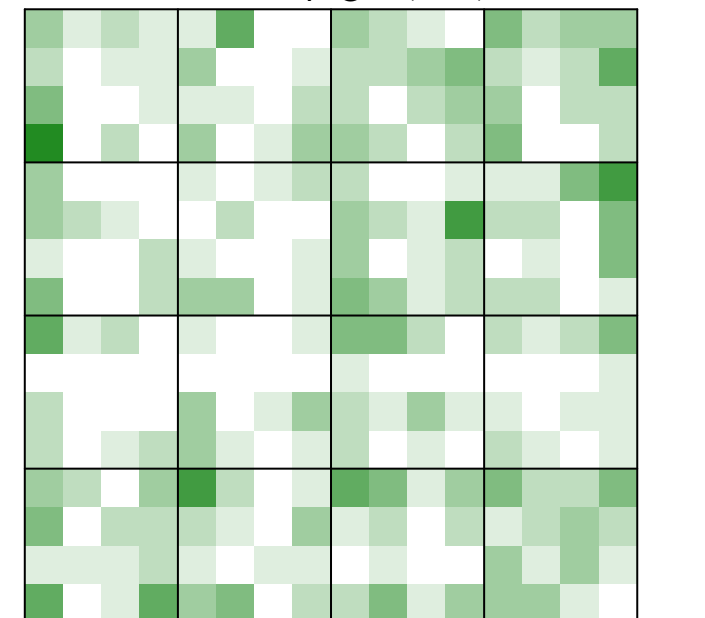
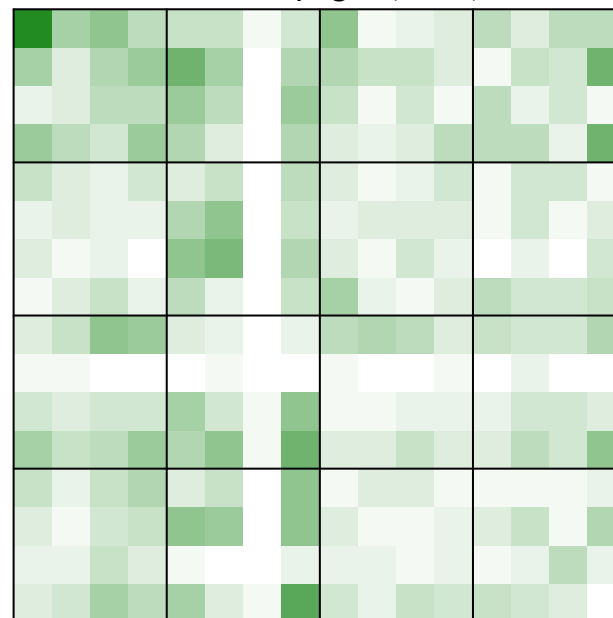
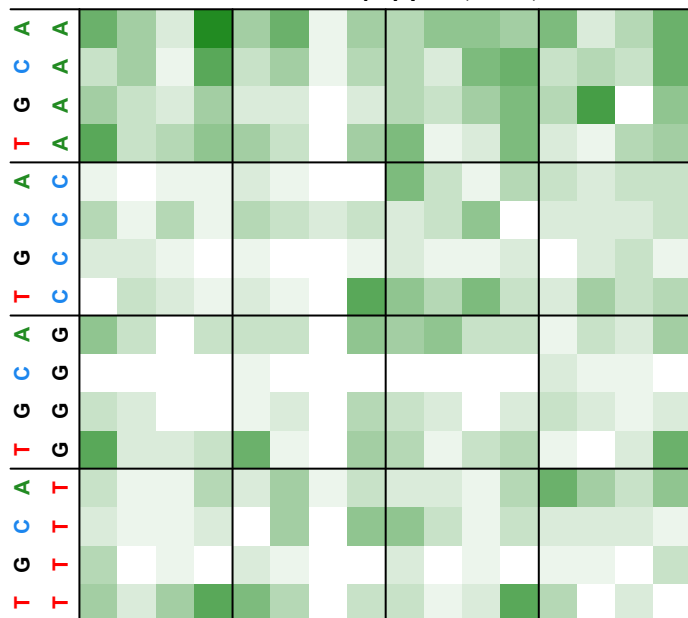


T>A (N=724)

T>C (N=1011)

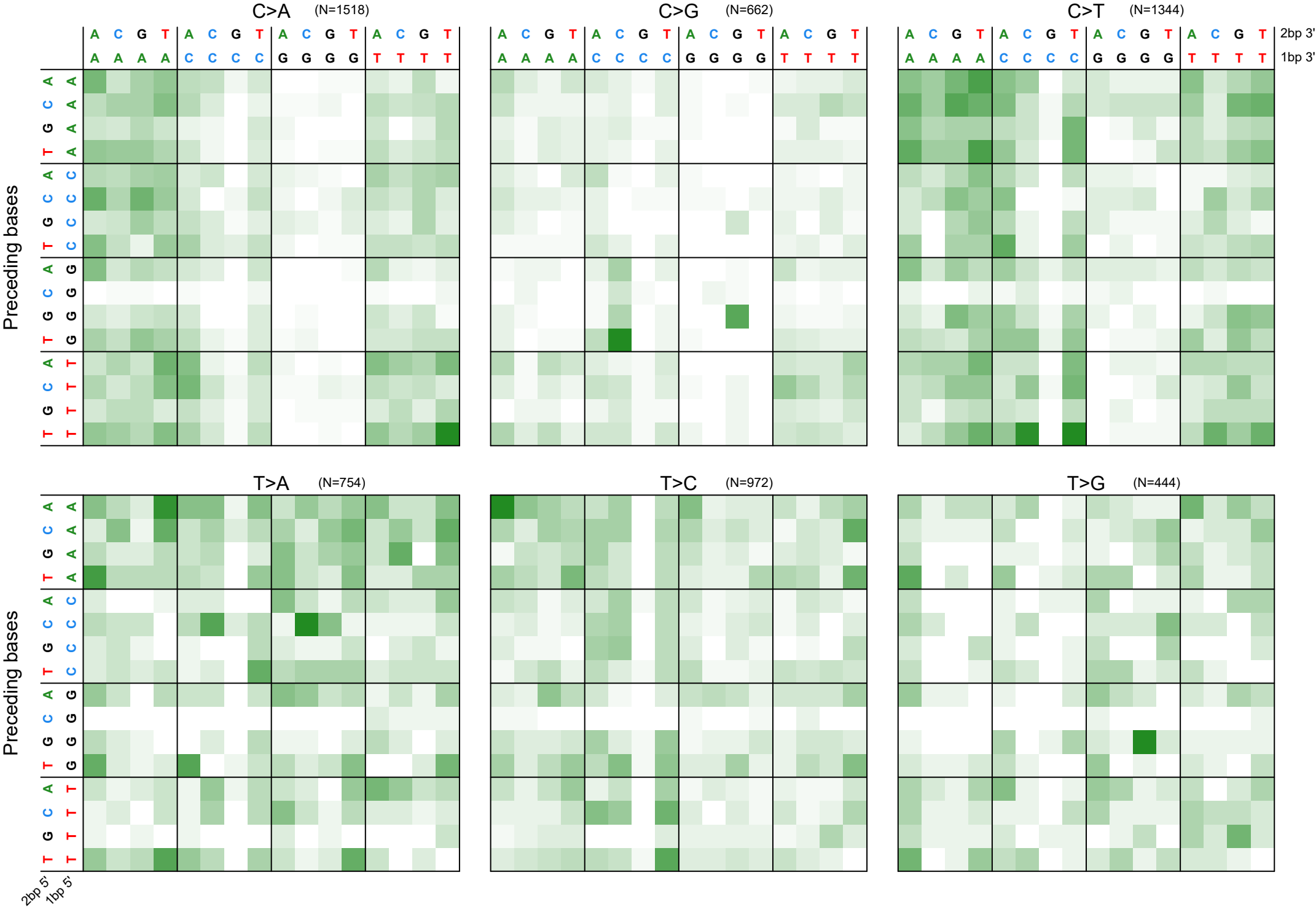
T>G (N=396)

Preceding bases



2bp 5'
1bp 5'

NDMA_cl3



NPIP_cl1

C>A (N=1599)

C>G (N=590)

C>T (N=1505)

T>A (N=1083)

T>C (N=1391)

T>G (N=583)

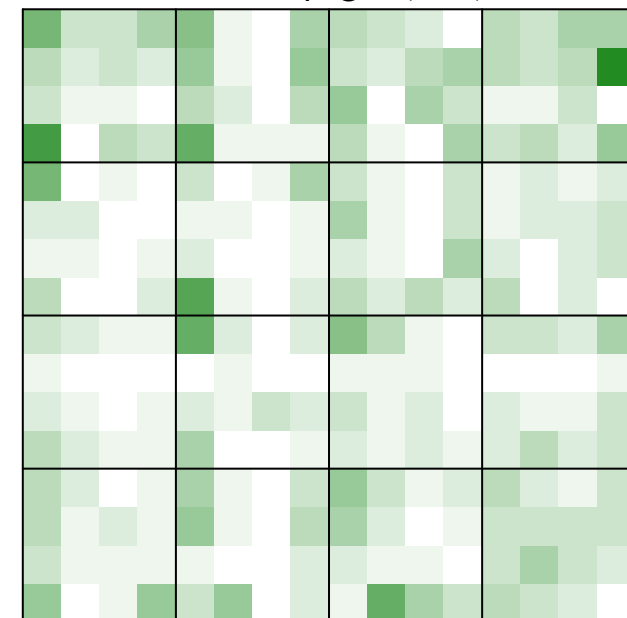
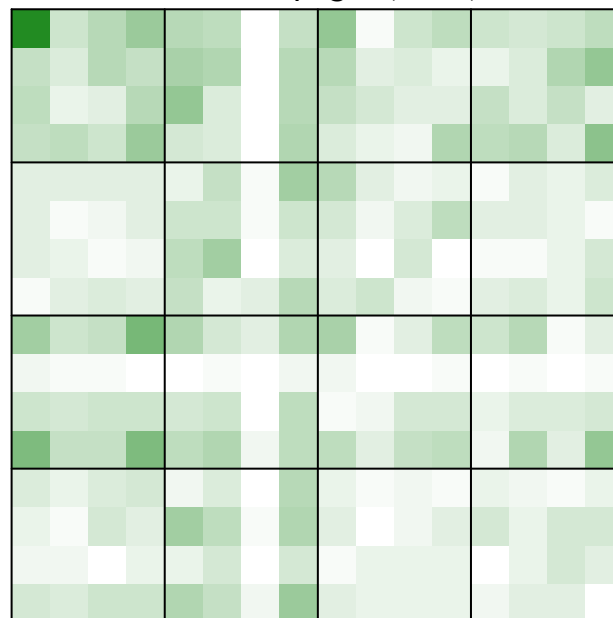
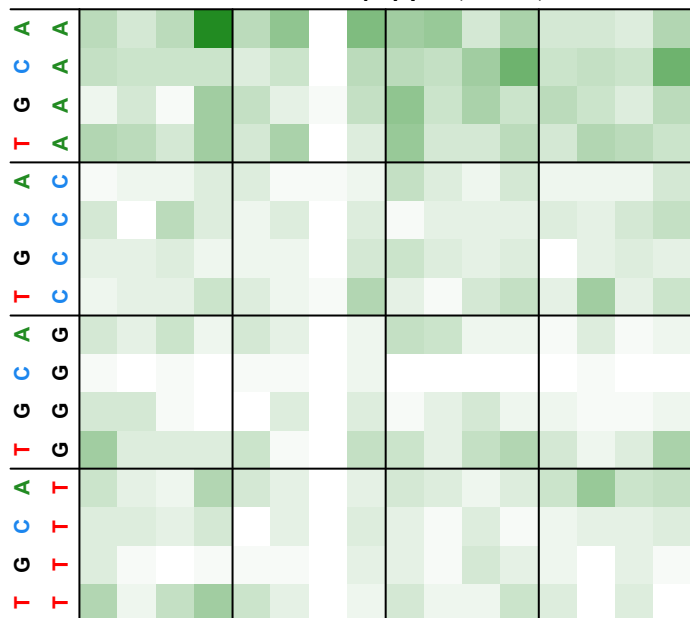
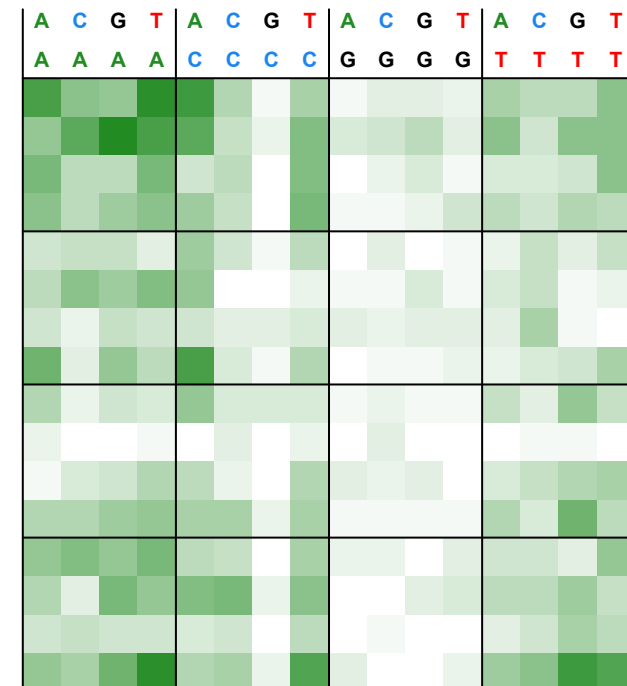
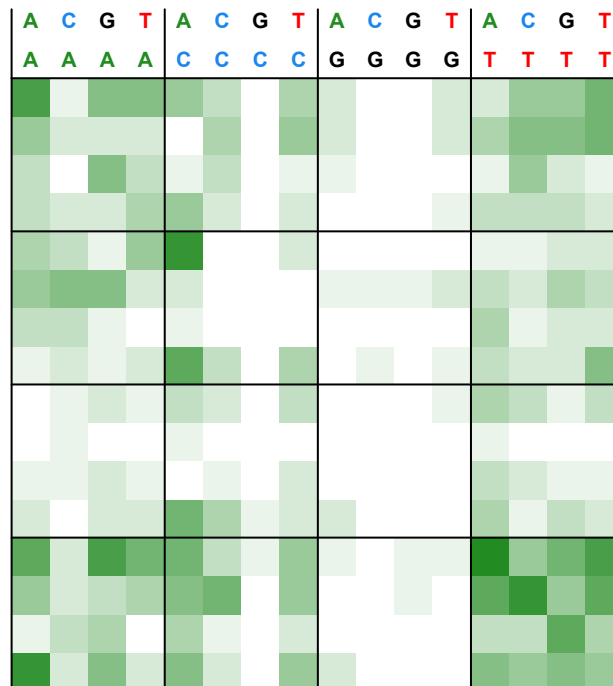
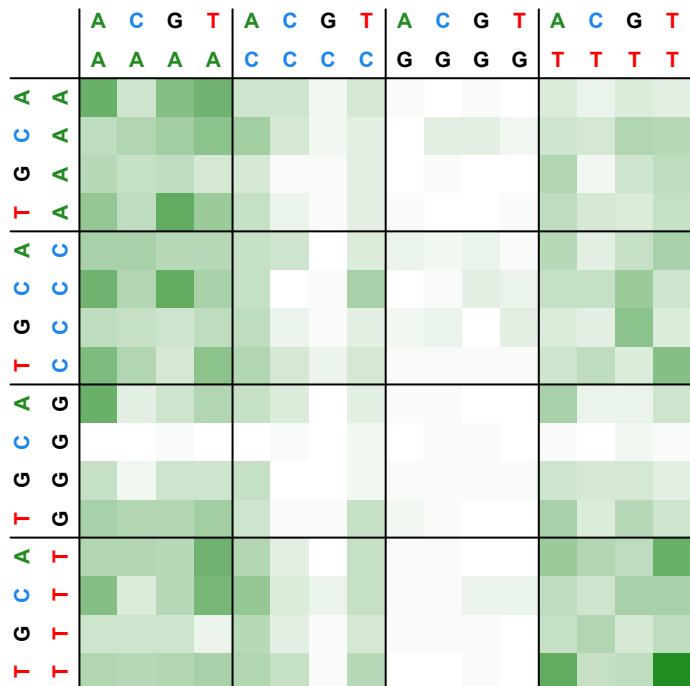
Preceding bases

Preceding bases

2bp 3'

1bp 3'

2bp 5'
1bp 5'



NPIP_cl2

C>A (N=1604)

C>G (N=610)

C>T (N=1505)

T>A (N=1153)

T>C (N=1425)

T>G (N=599)

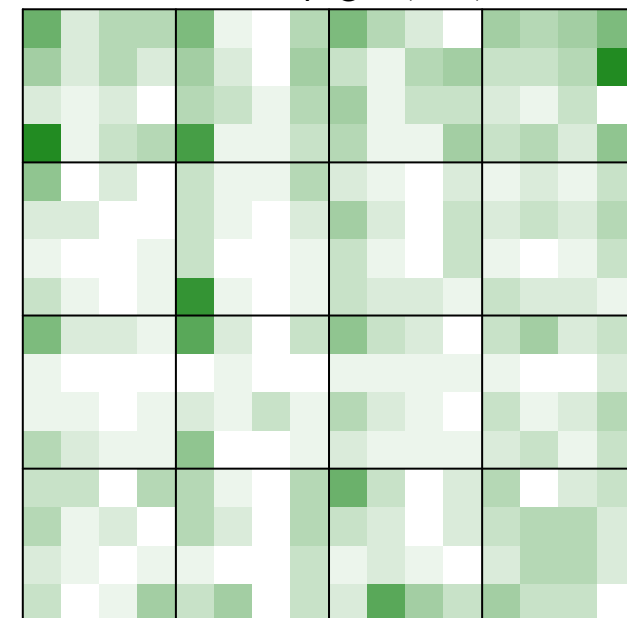
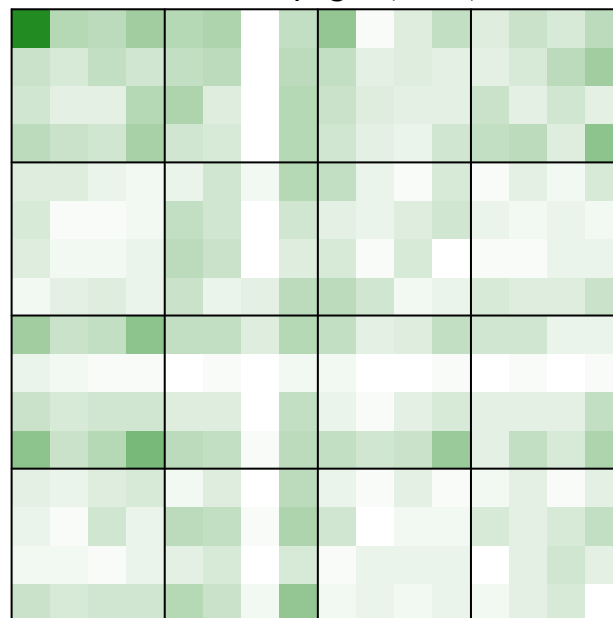
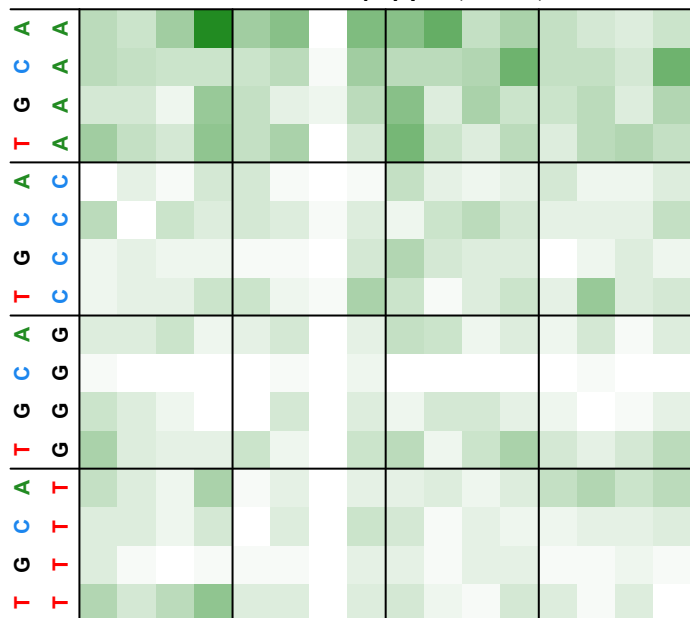
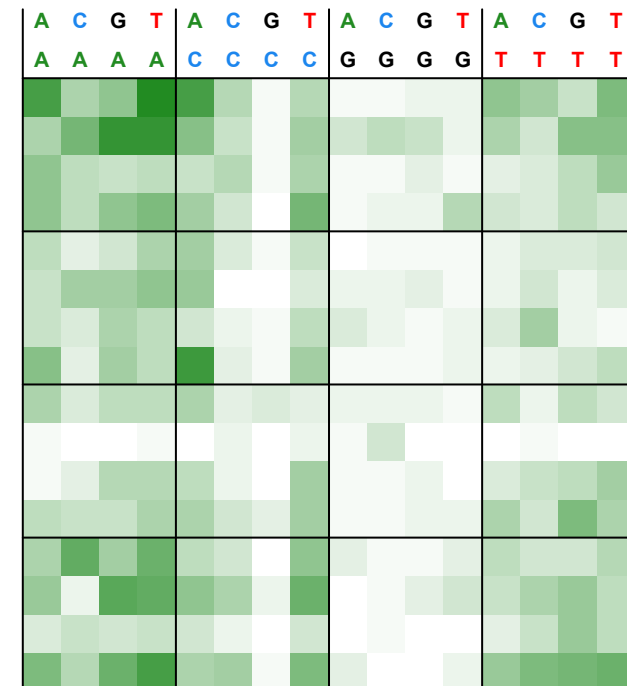
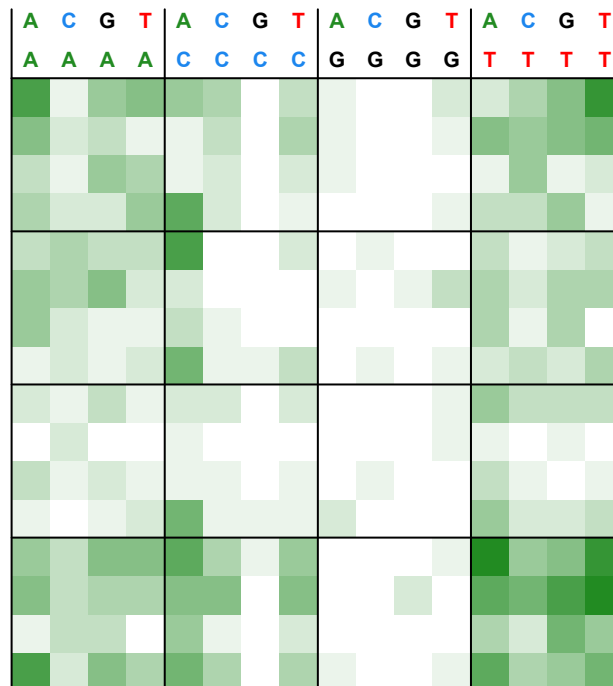
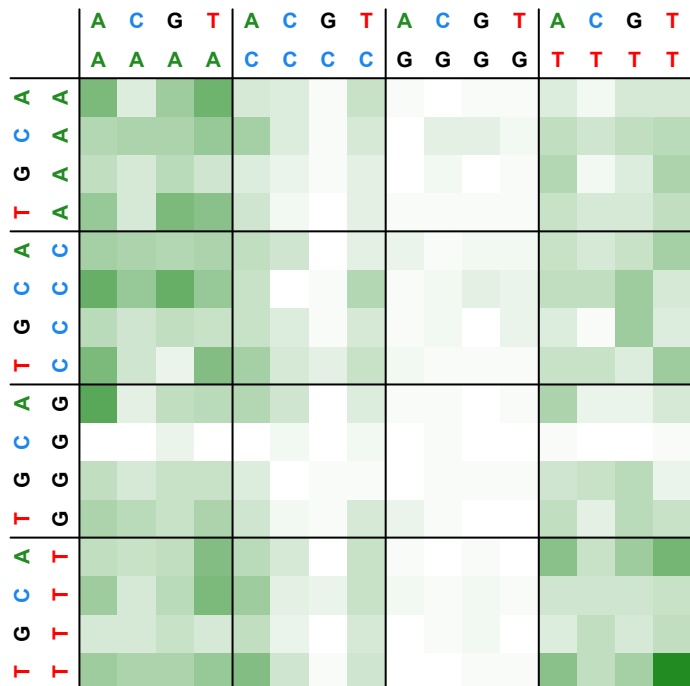
Preceding bases

Preceding bases

2bp 3'

1bp 3'

2bp 5'
1bp 5'



NPIP_cl3

C>A (N=1430)

C>G (N=623)

C>T (N=1236)

T>A (N=761)

T>C (N=976)

T>G (N=500)

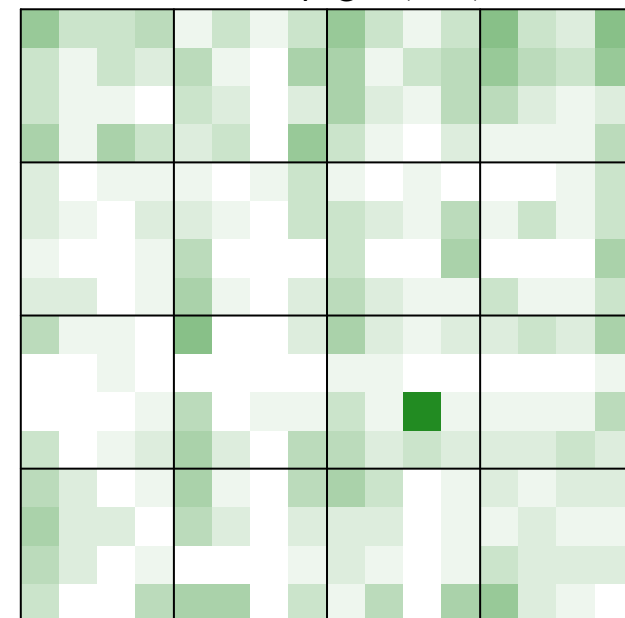
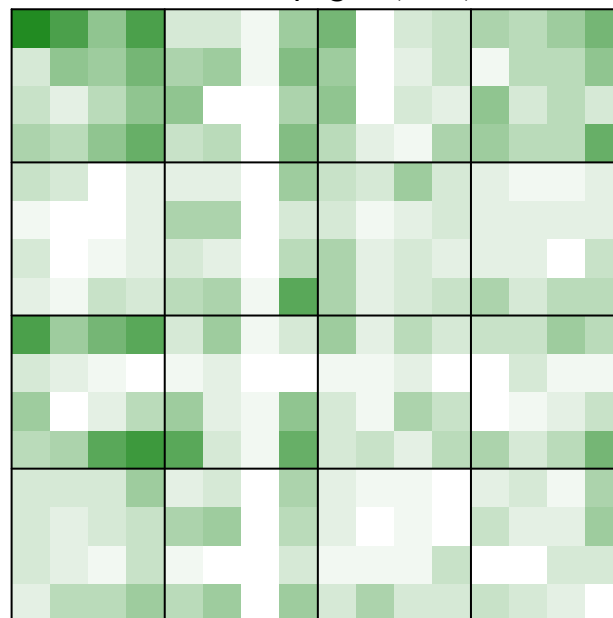
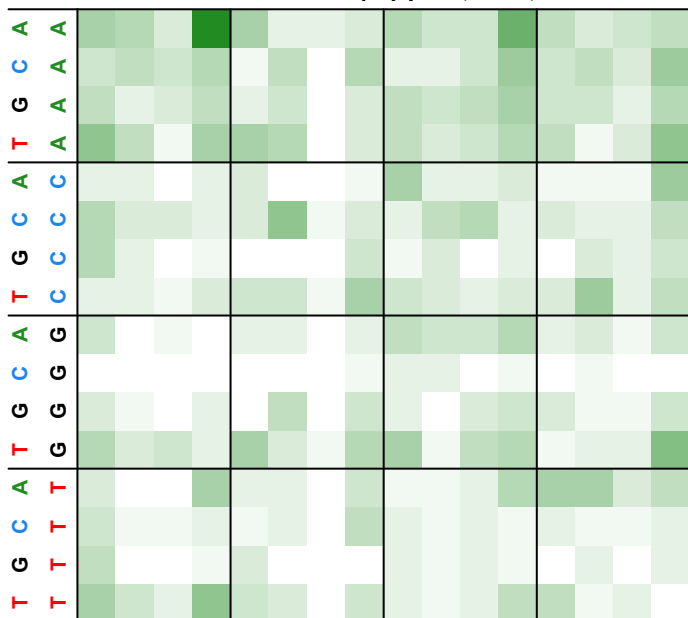
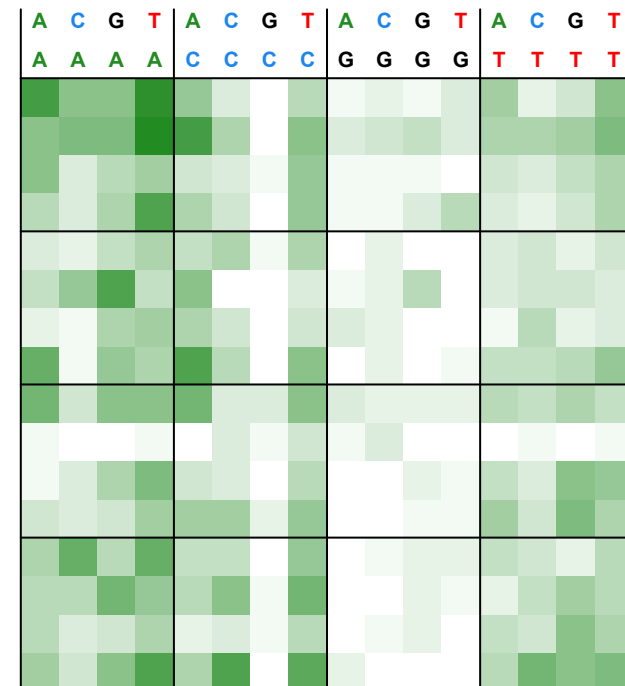
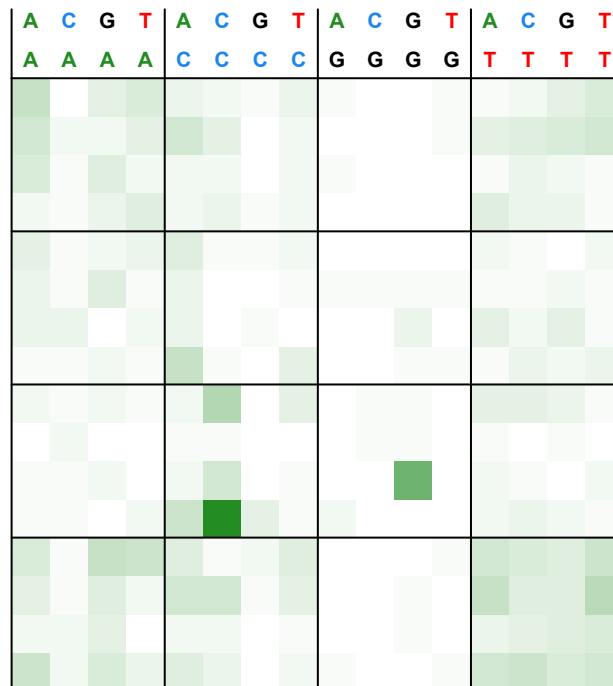
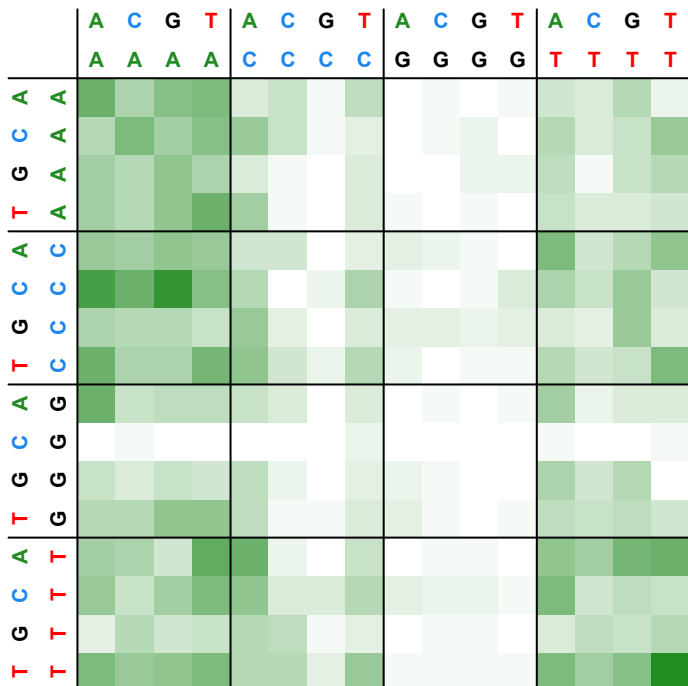
Preceding bases

Preceding bases

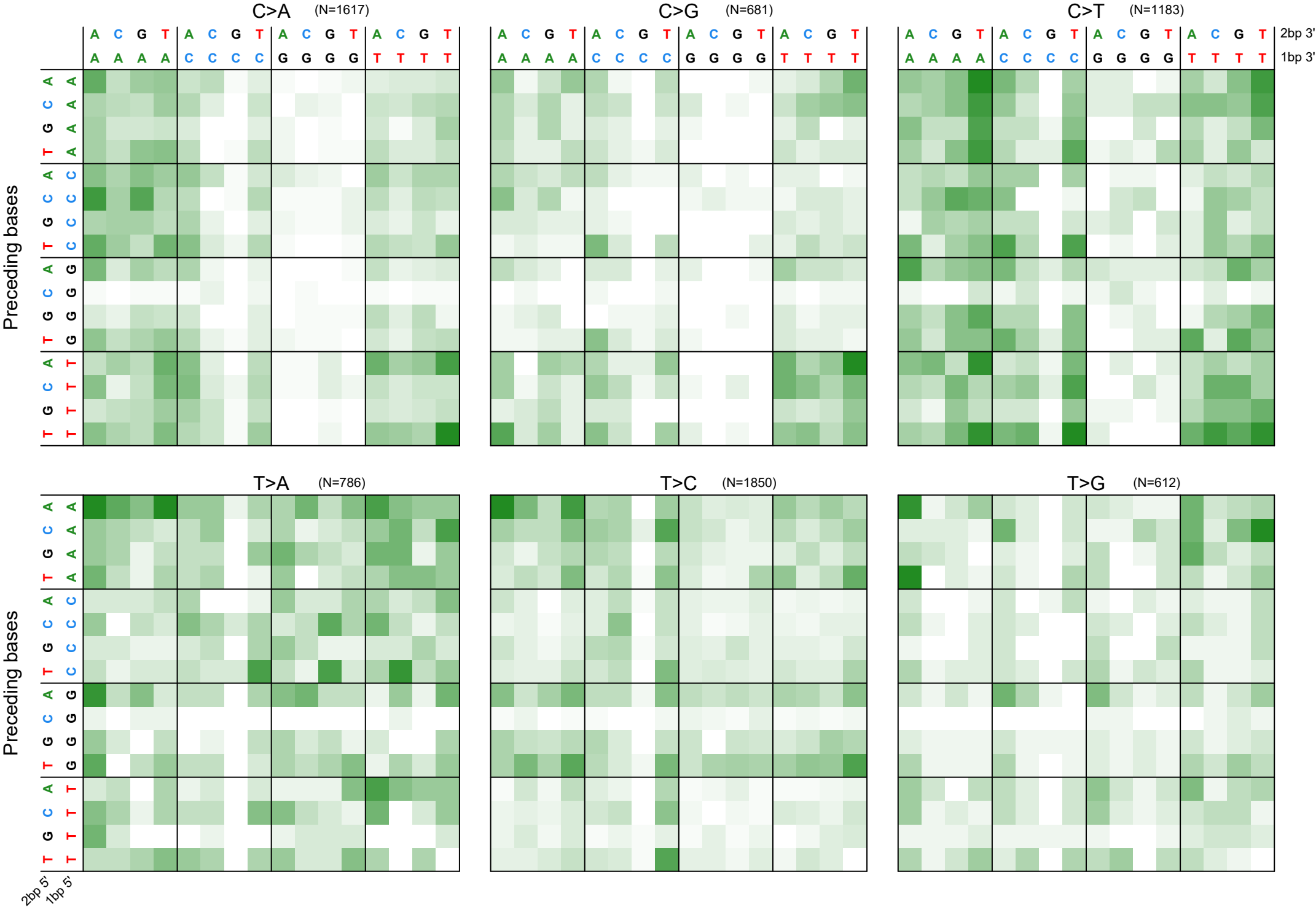
2bp 3'

1bp 3'

2bp 5'
1bp 5'



NPYR_cl1



NPYR_cl2

