

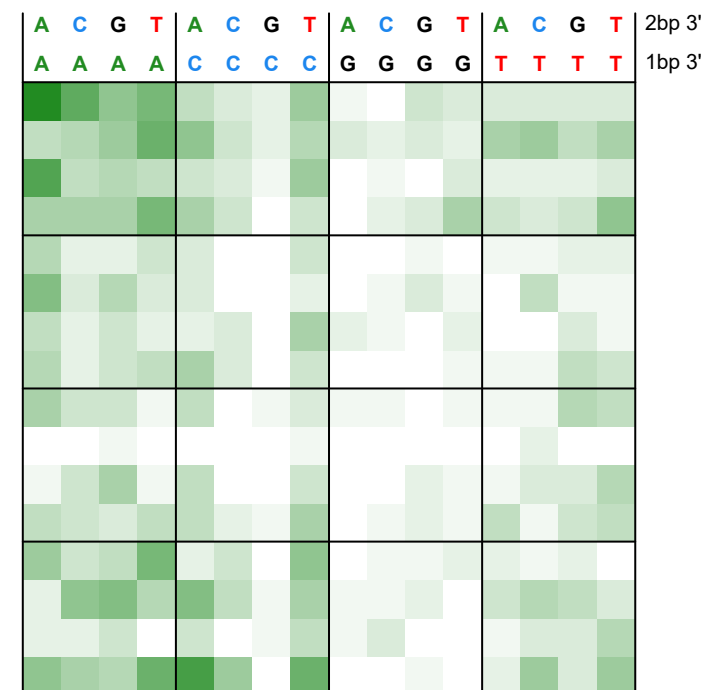
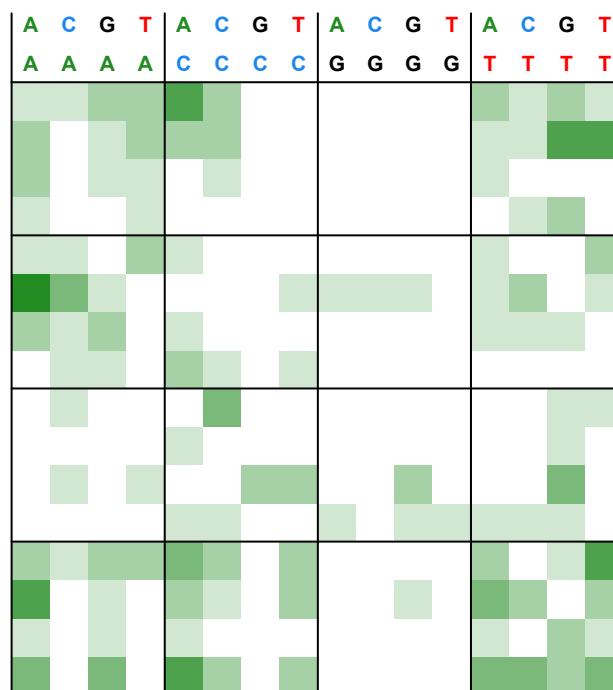
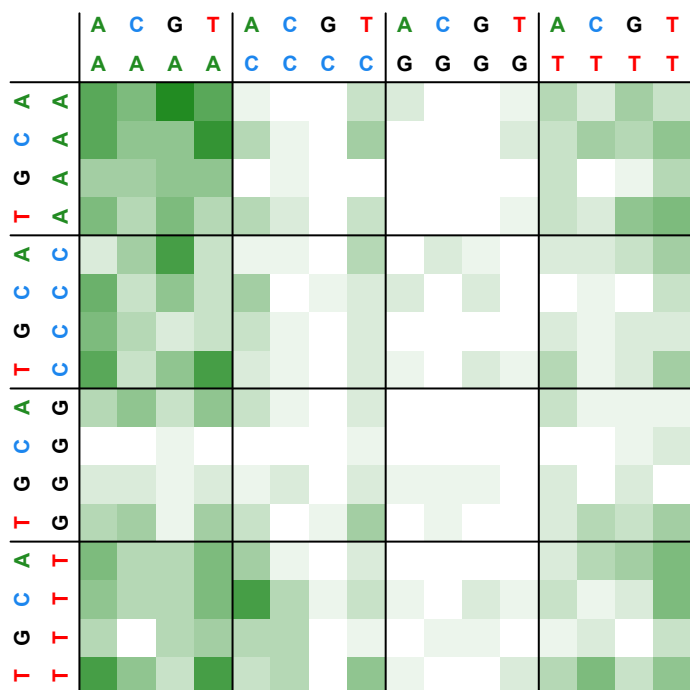
NDEA.cl1

C>A (N=649)

C>G (N=186)

C>T (N=844)

Preceding bases



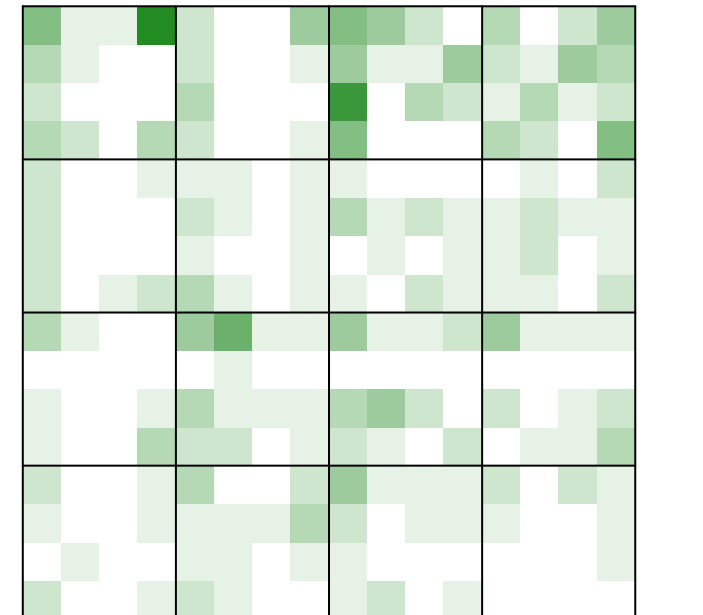
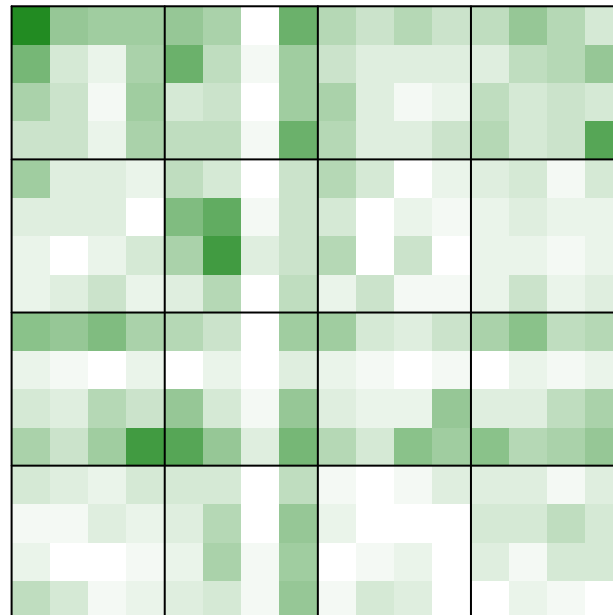
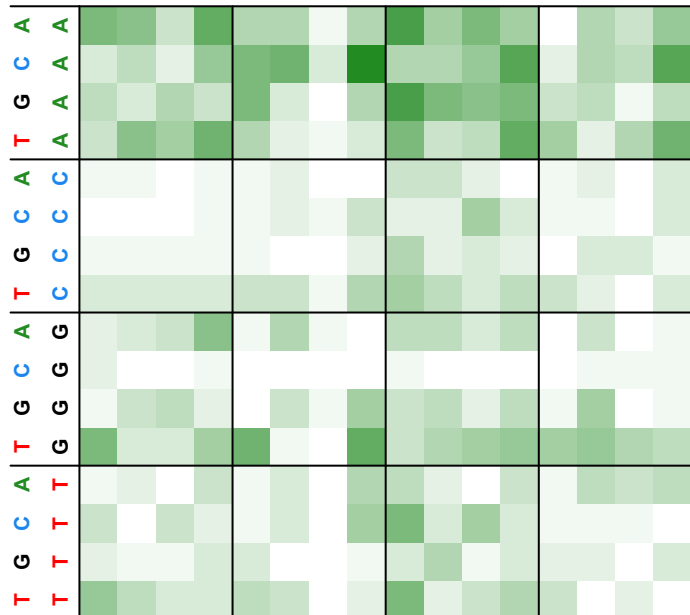
2bp 3'
1bp 3'

T>A (N=938)

T>C (N=1158)

T>G (N=299)

Preceding bases



2bp 5'
1bp 5'

NDEA.cl2

C>A (N=764)

C>G (N=255)

C>T (N=926)

T>A (N=1108)

T>C (N=1453)

T>G (N=333)

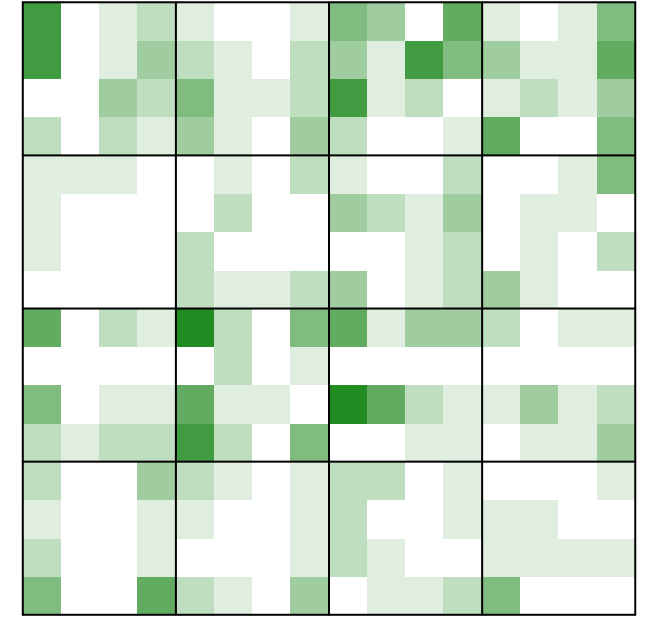
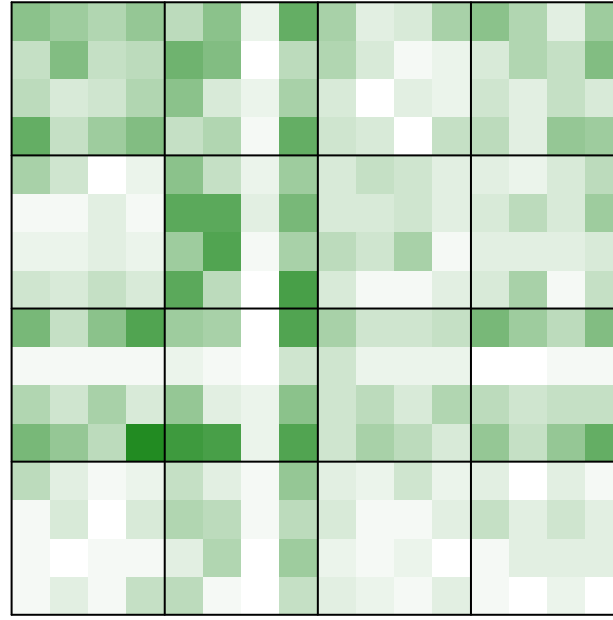
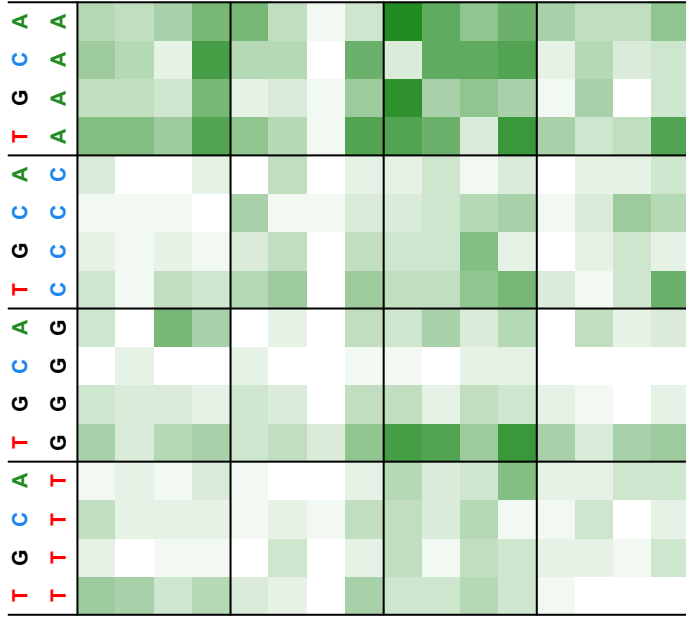
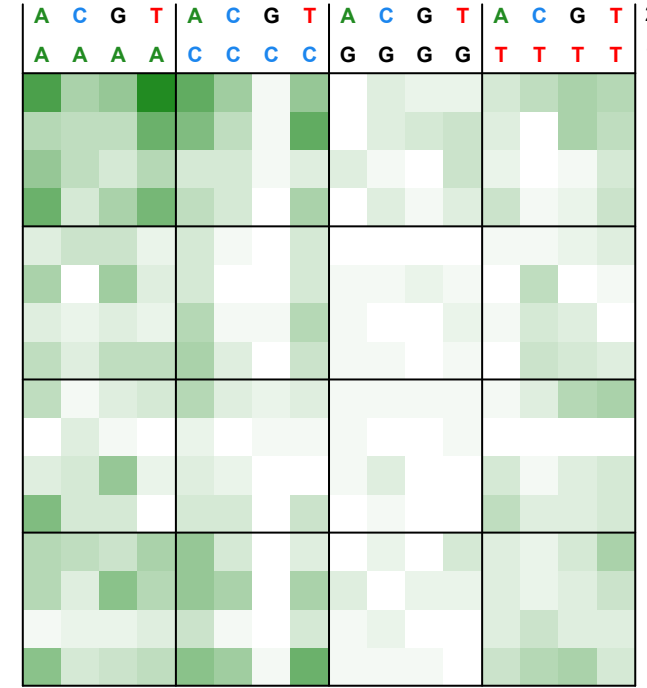
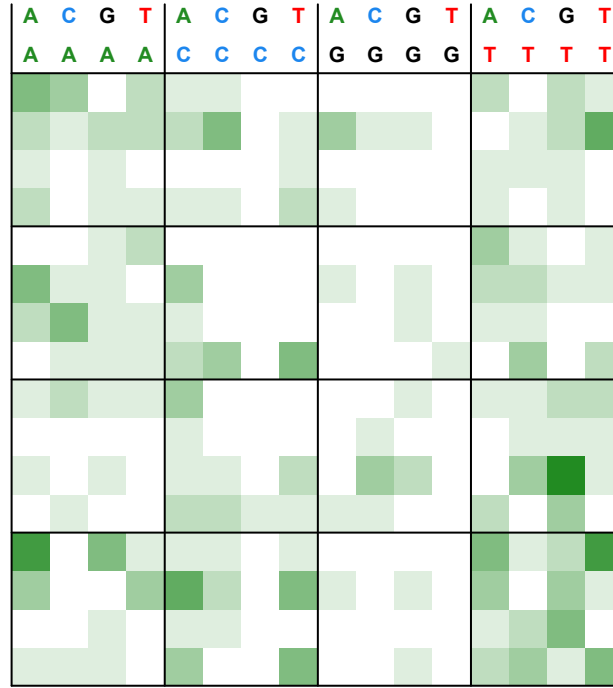
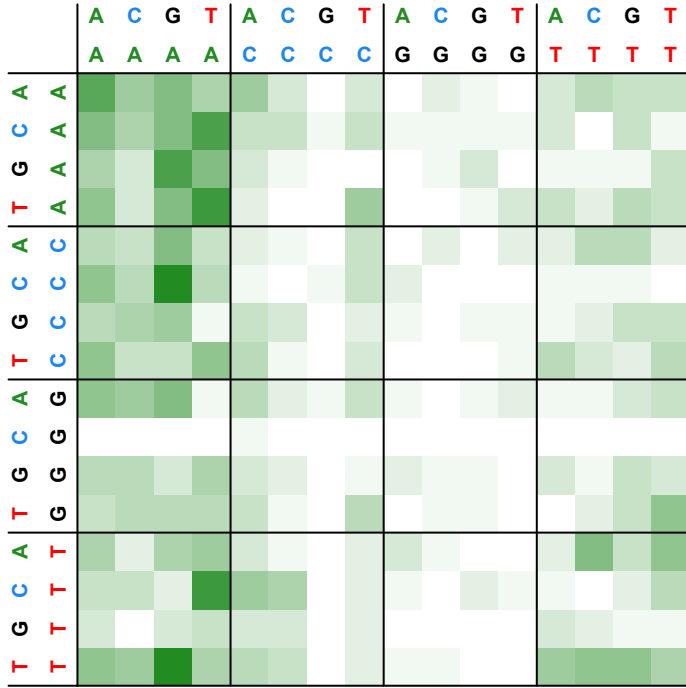
Preceding bases

Preceding bases

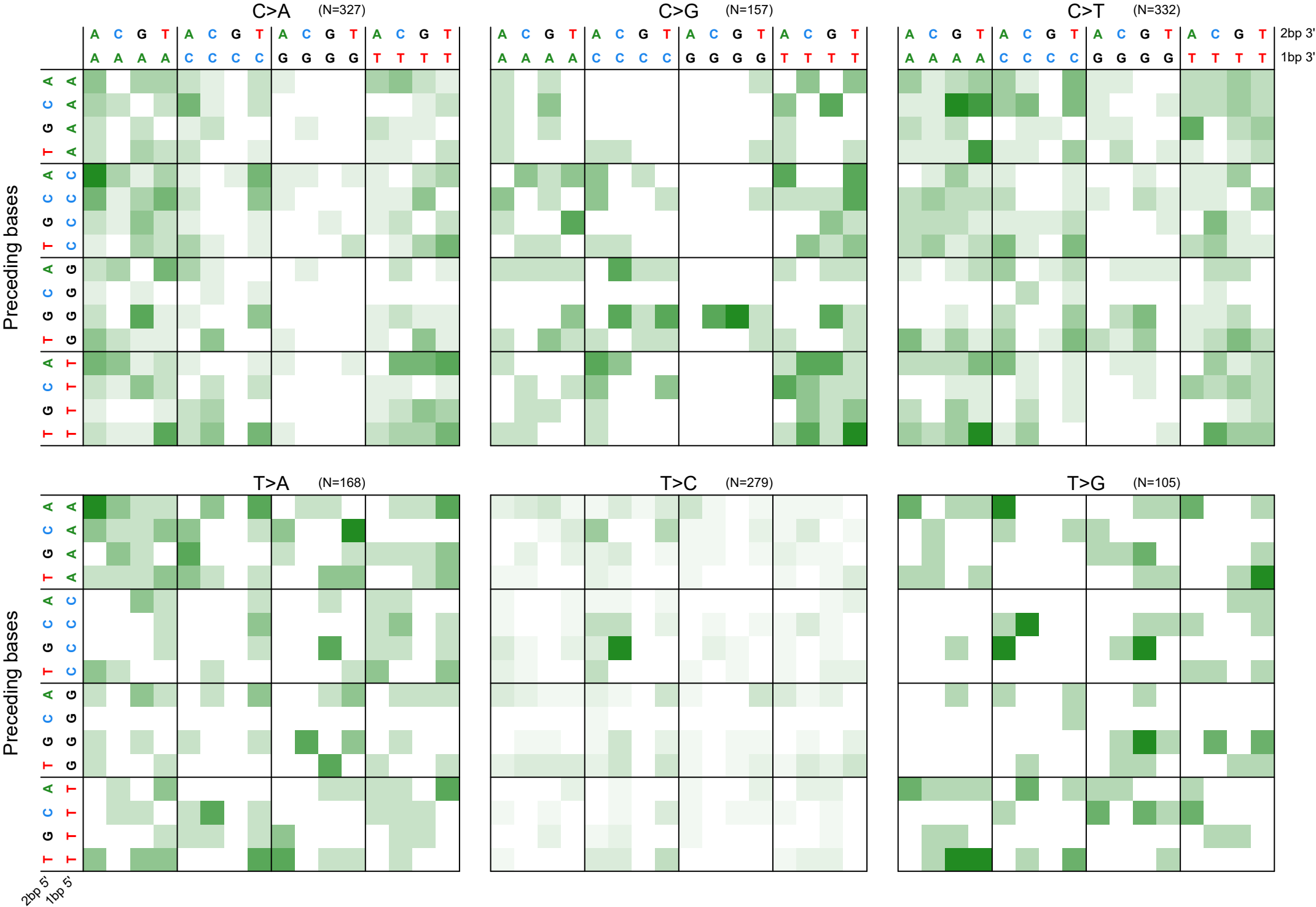
2bp 5'
1bp 5'

2bp 3'

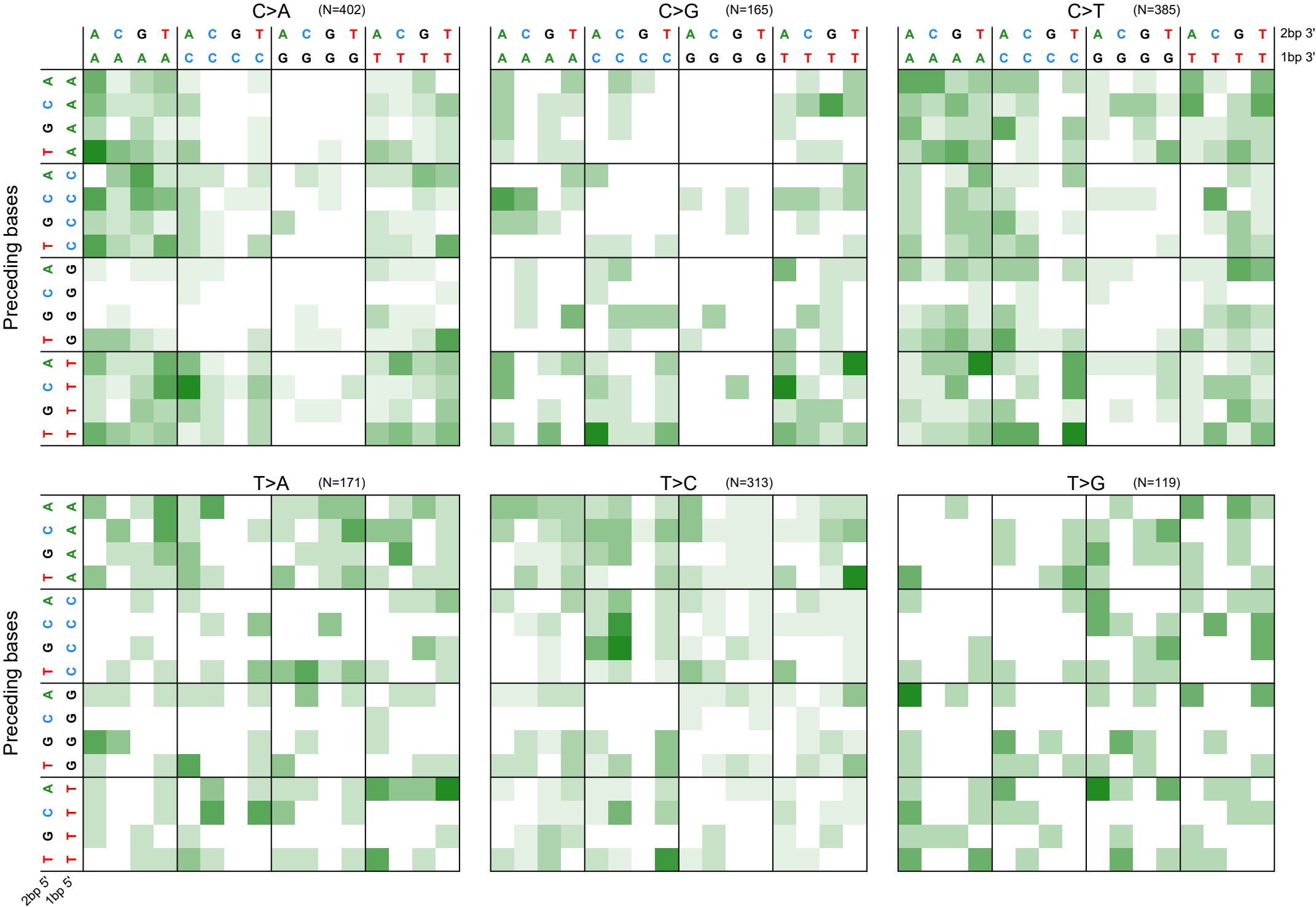
1bp 3'



NMDA.cl1



NMDA.cl2



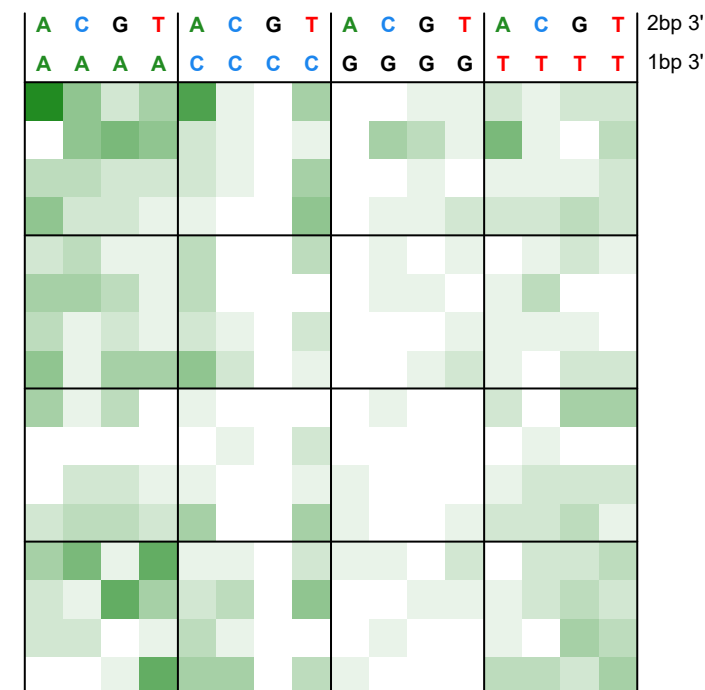
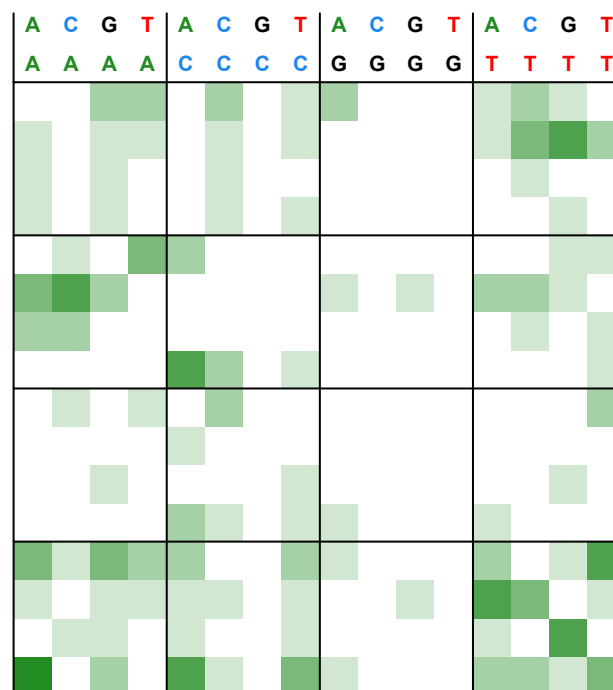
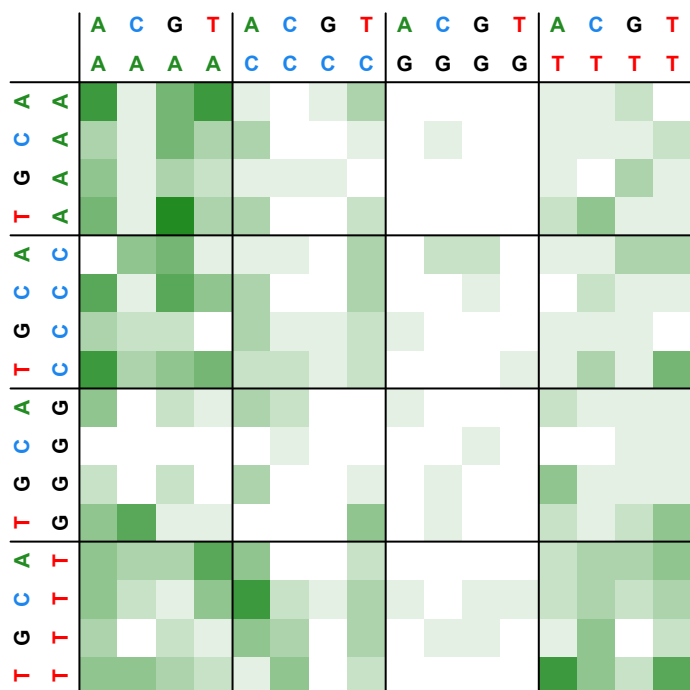
NPIP.cl1

C>A (N=408)

C>G (N=160)

C>T (N=403)

Preceding bases

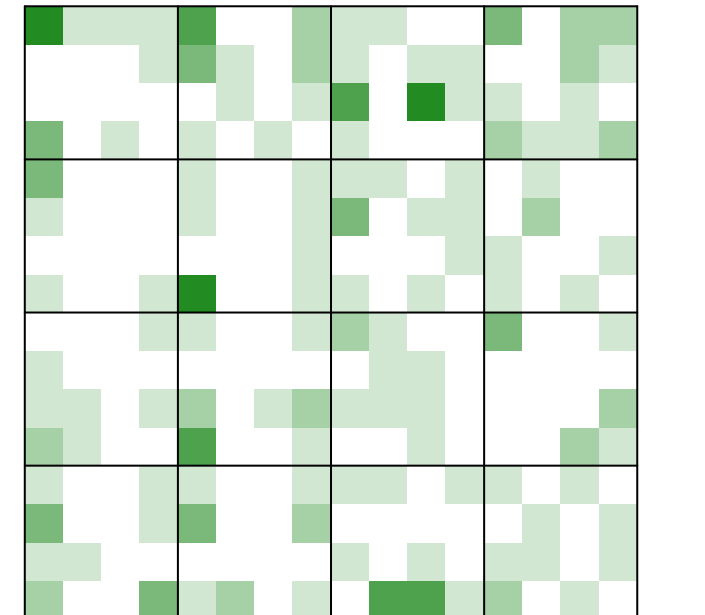
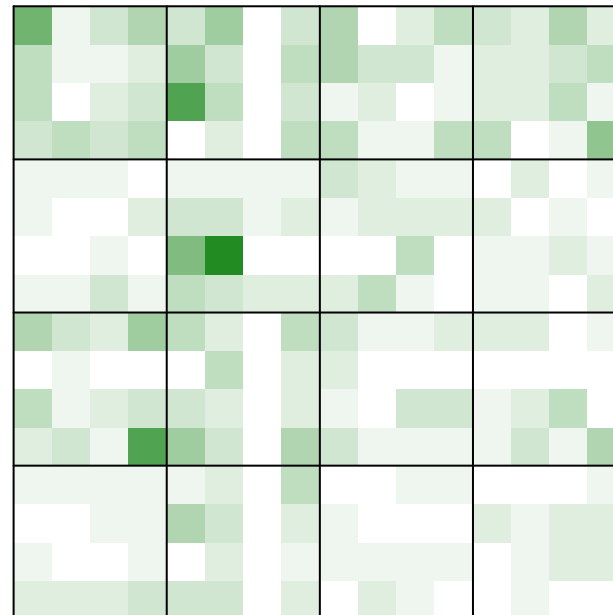
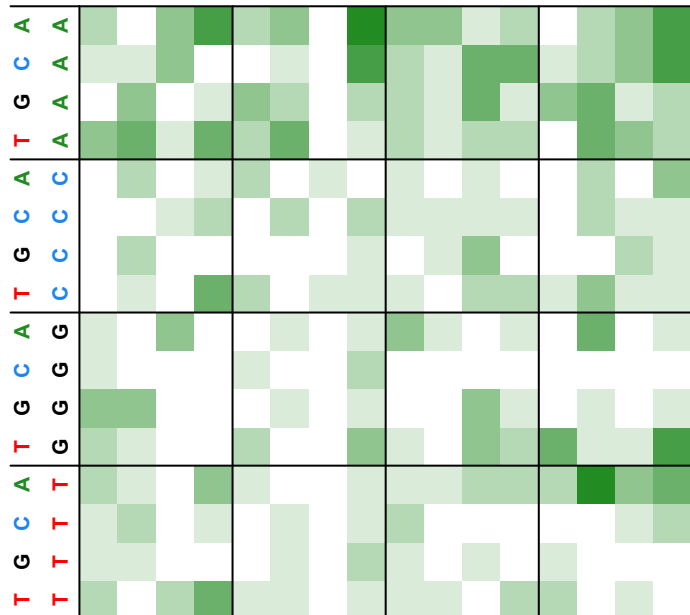


T>A (N=320)

T>C (N=472)

T>G (N=184)

Preceding bases



2bp 5'
1bp 5'

NPIP.cl2

C>A (N=419)

C>G (N=158)

C>T (N=422)

T>A (N=333)

T>C (N=480)

T>G (N=189)

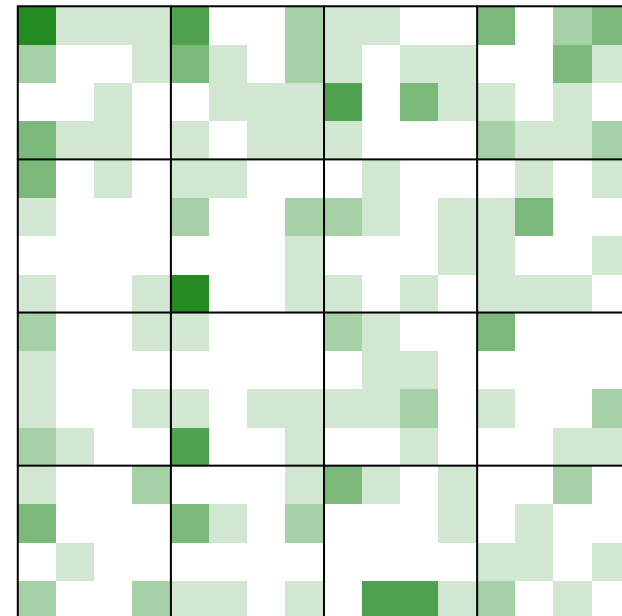
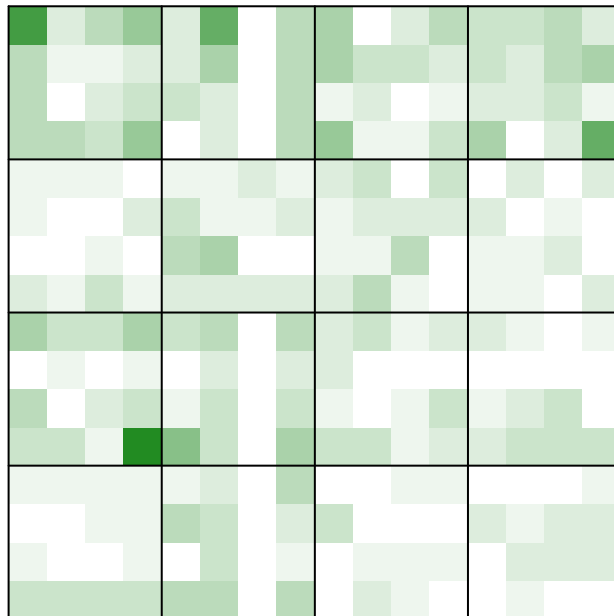
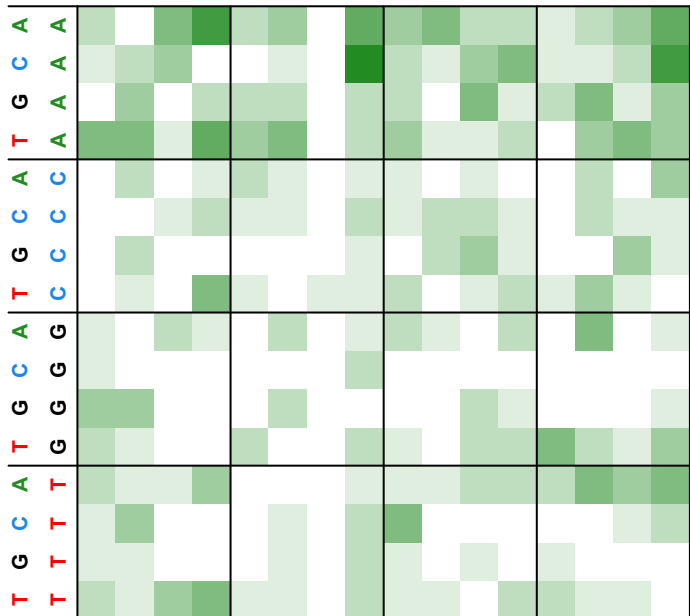
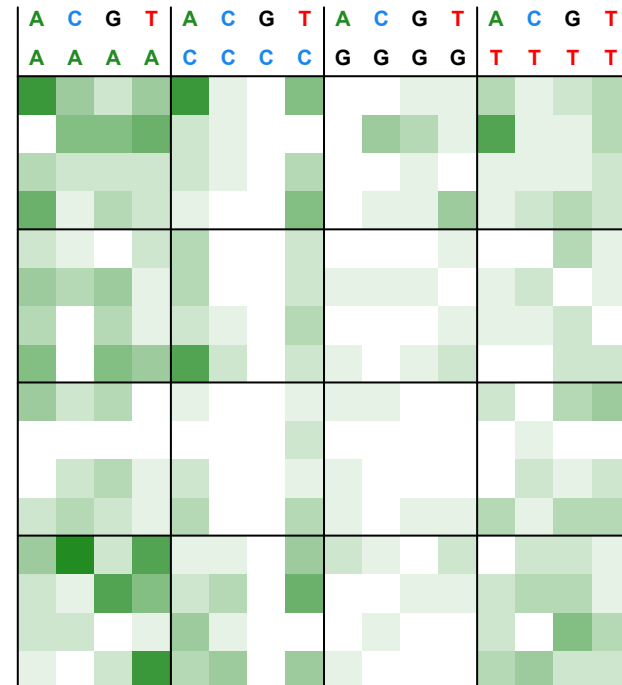
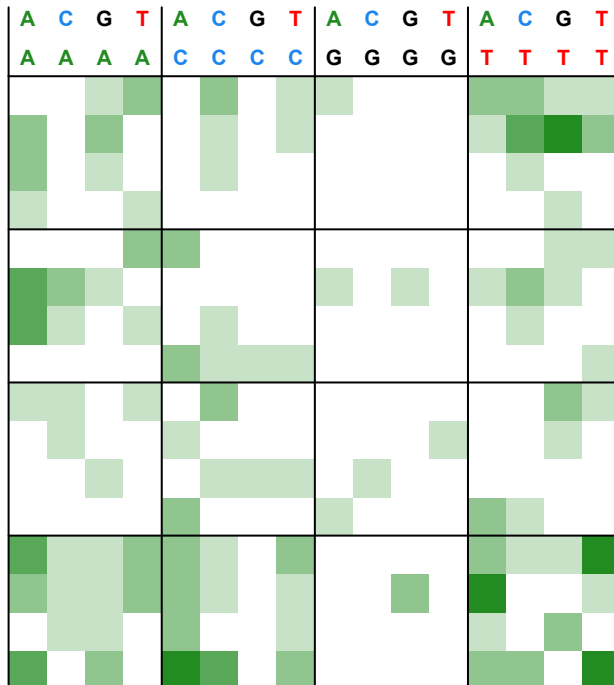
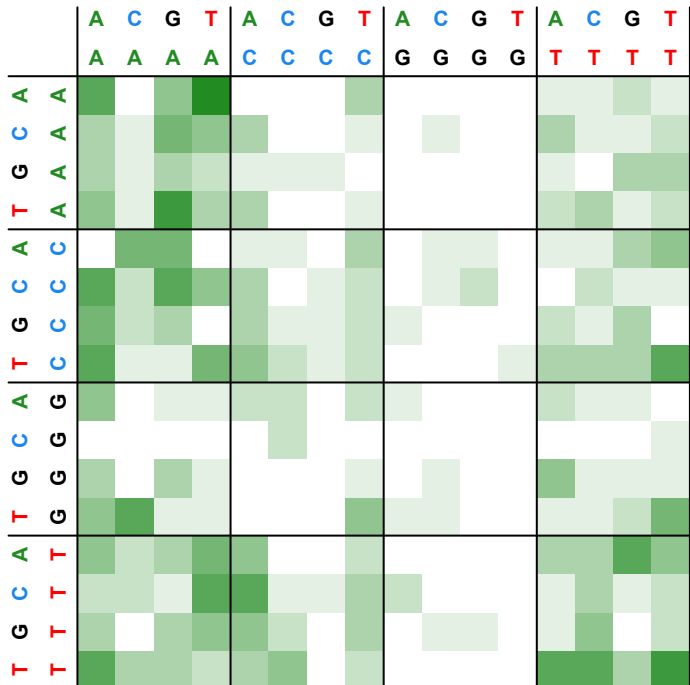
Preceding bases

Preceding bases

2bp 3'

1bp 3'

2bp 5'
1bp 5'



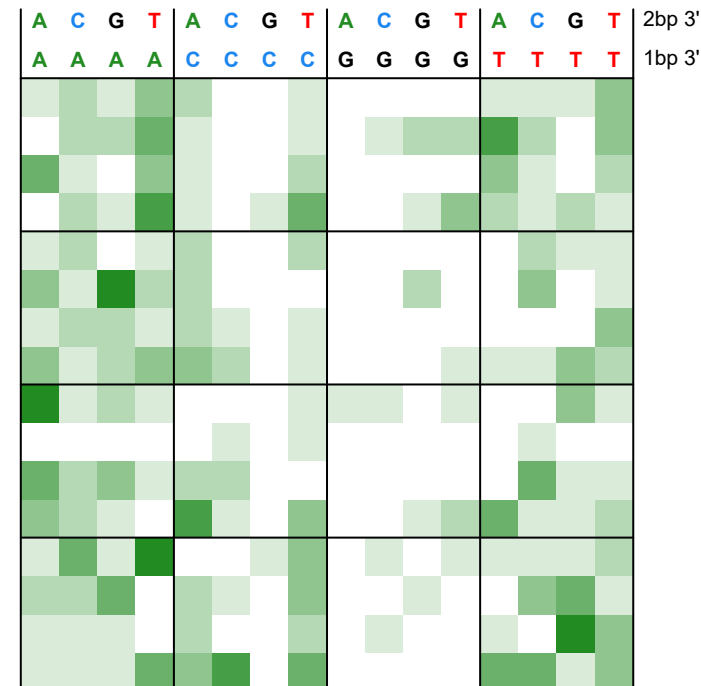
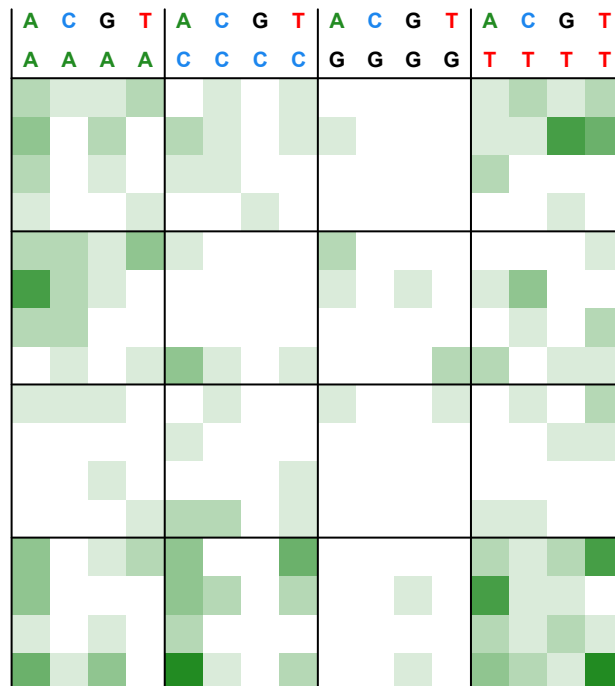
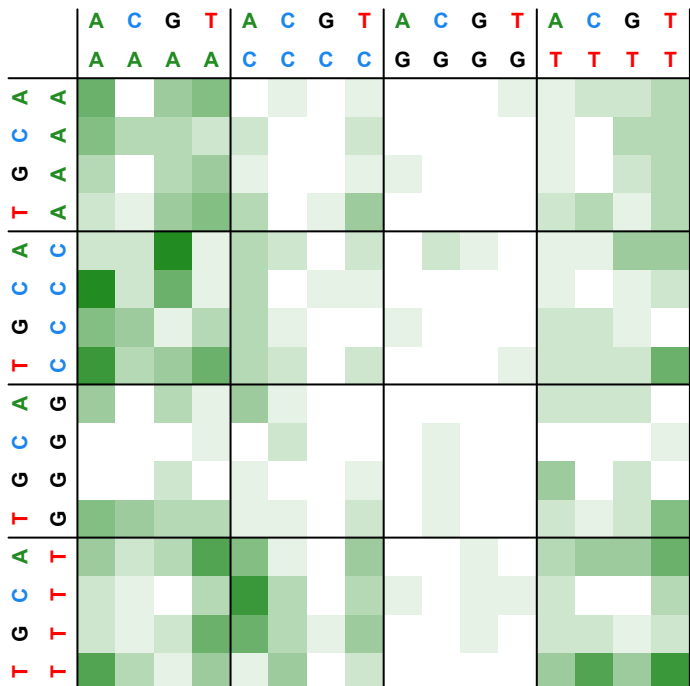
NPYR.cl1

C>A (N=438)

C>G (N=195)

C>T (N=312)

Preceding bases

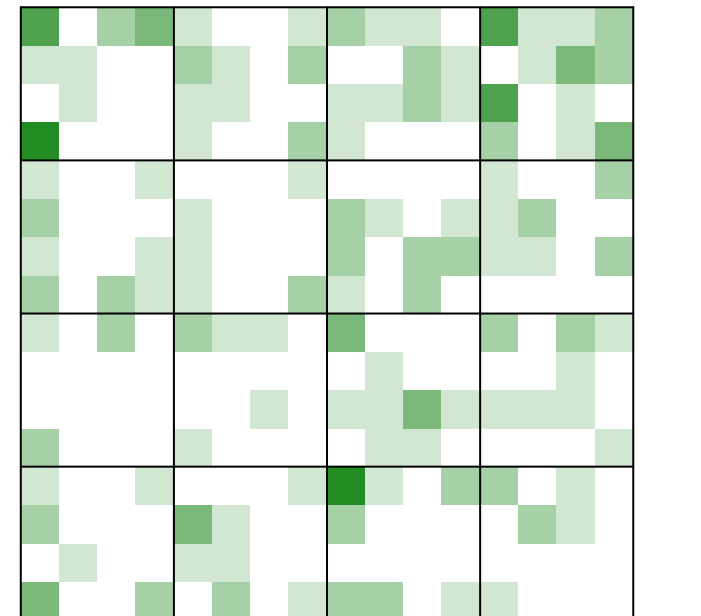
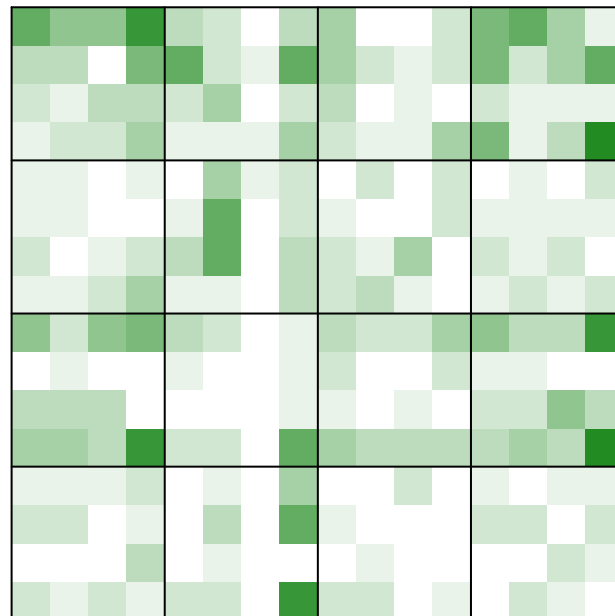
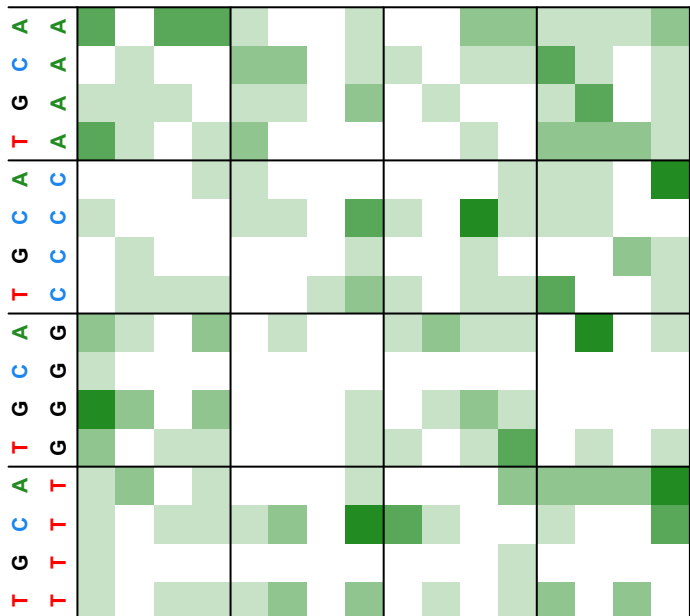


T>A (N=197)

T>C (N=500)

T>G (N=182)

Preceding bases



2bp 5'
1bp 5'

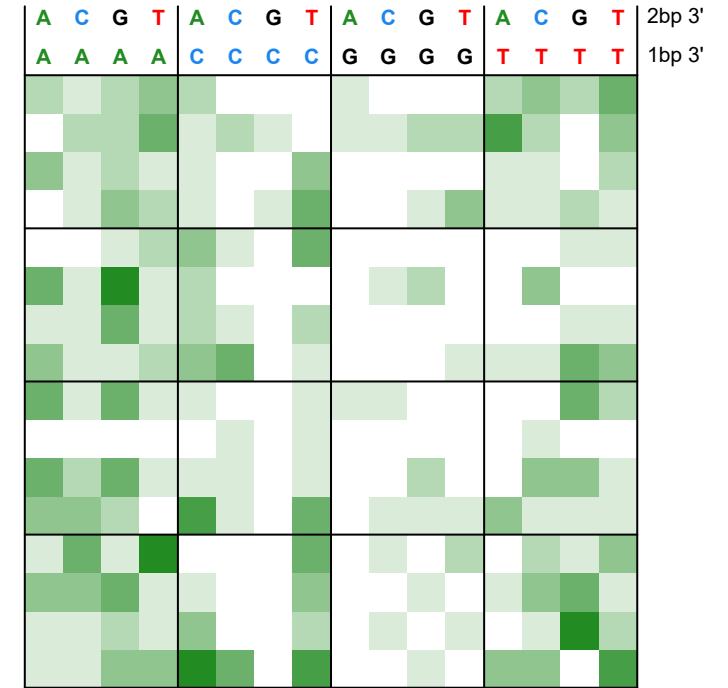
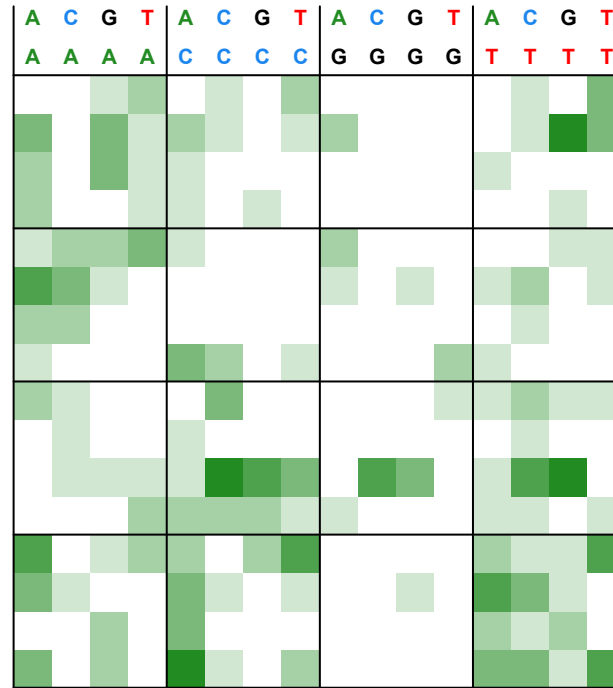
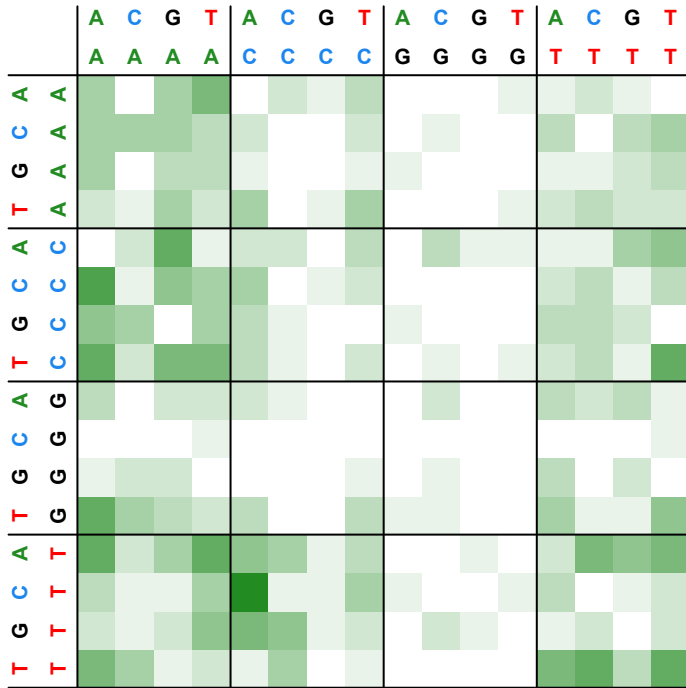
NPYR.cl2

C>A (N=468)

C>G (N=224)

C>T (N=337)

Preceding bases

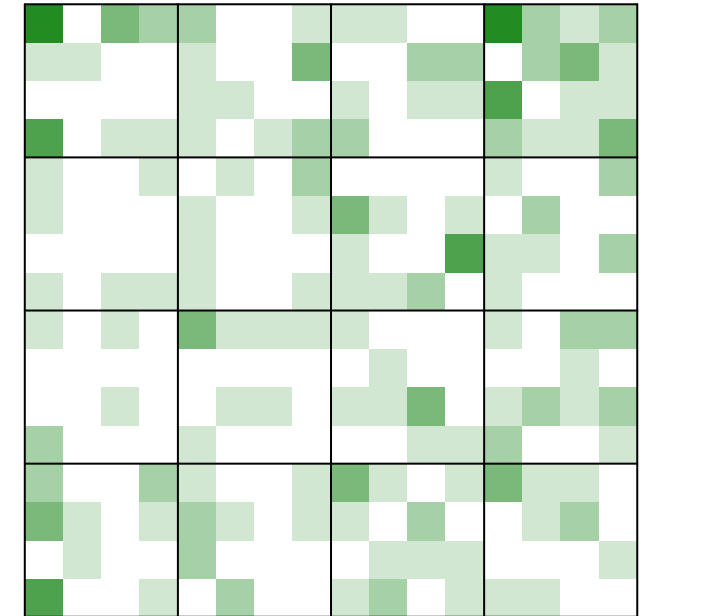
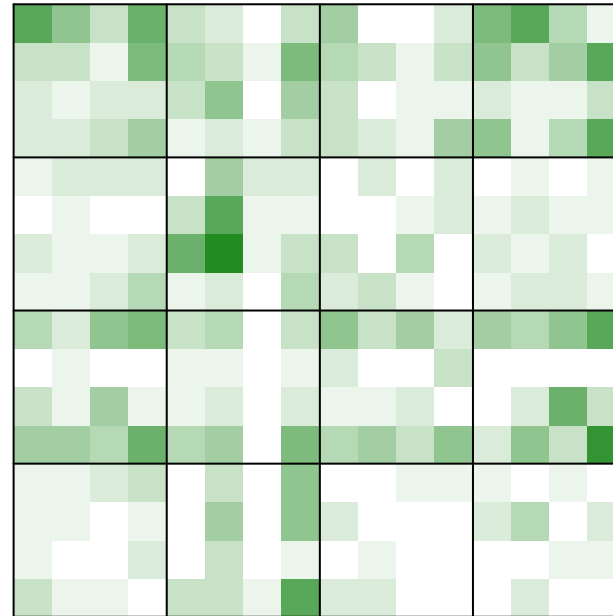
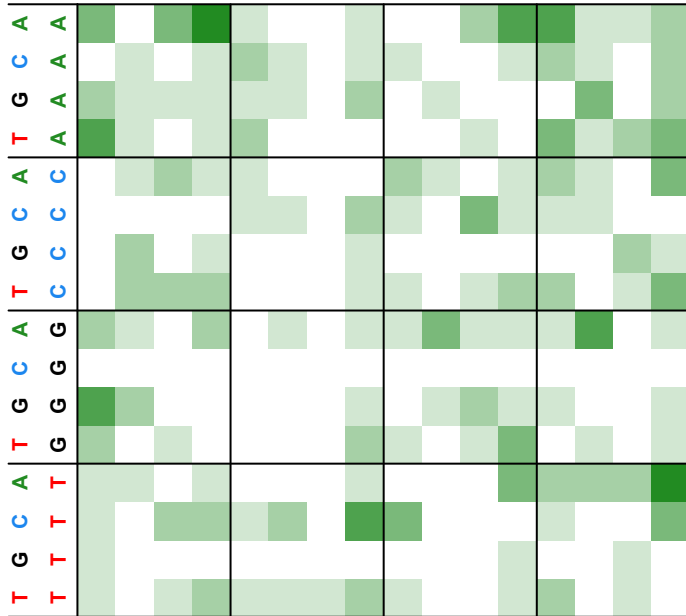


T>A (N=221)

T>C (N=591)

T>G (N=198)

Preceding bases



2bp 5'
1bp 5'

2bp 3'

1bp 3'