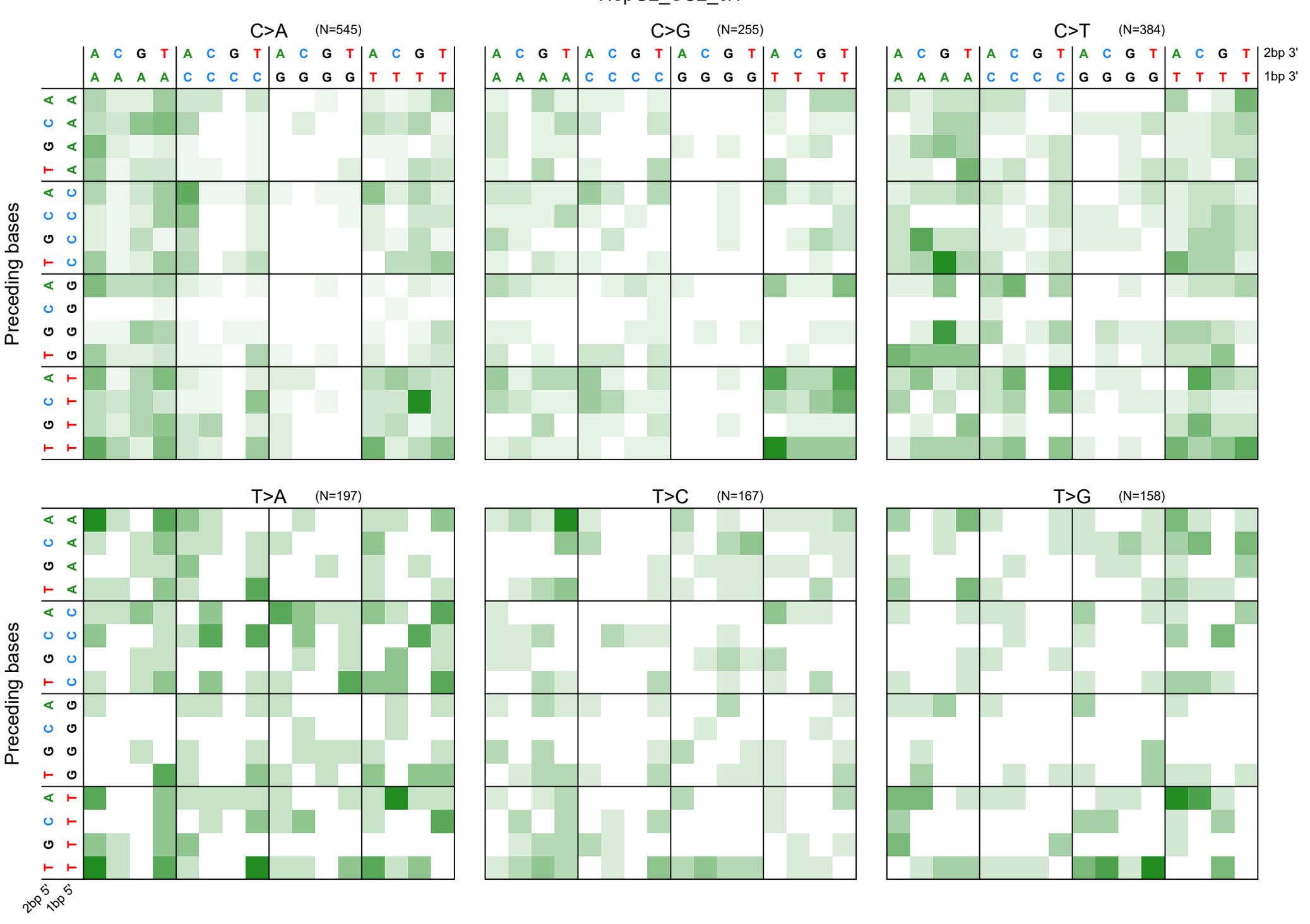
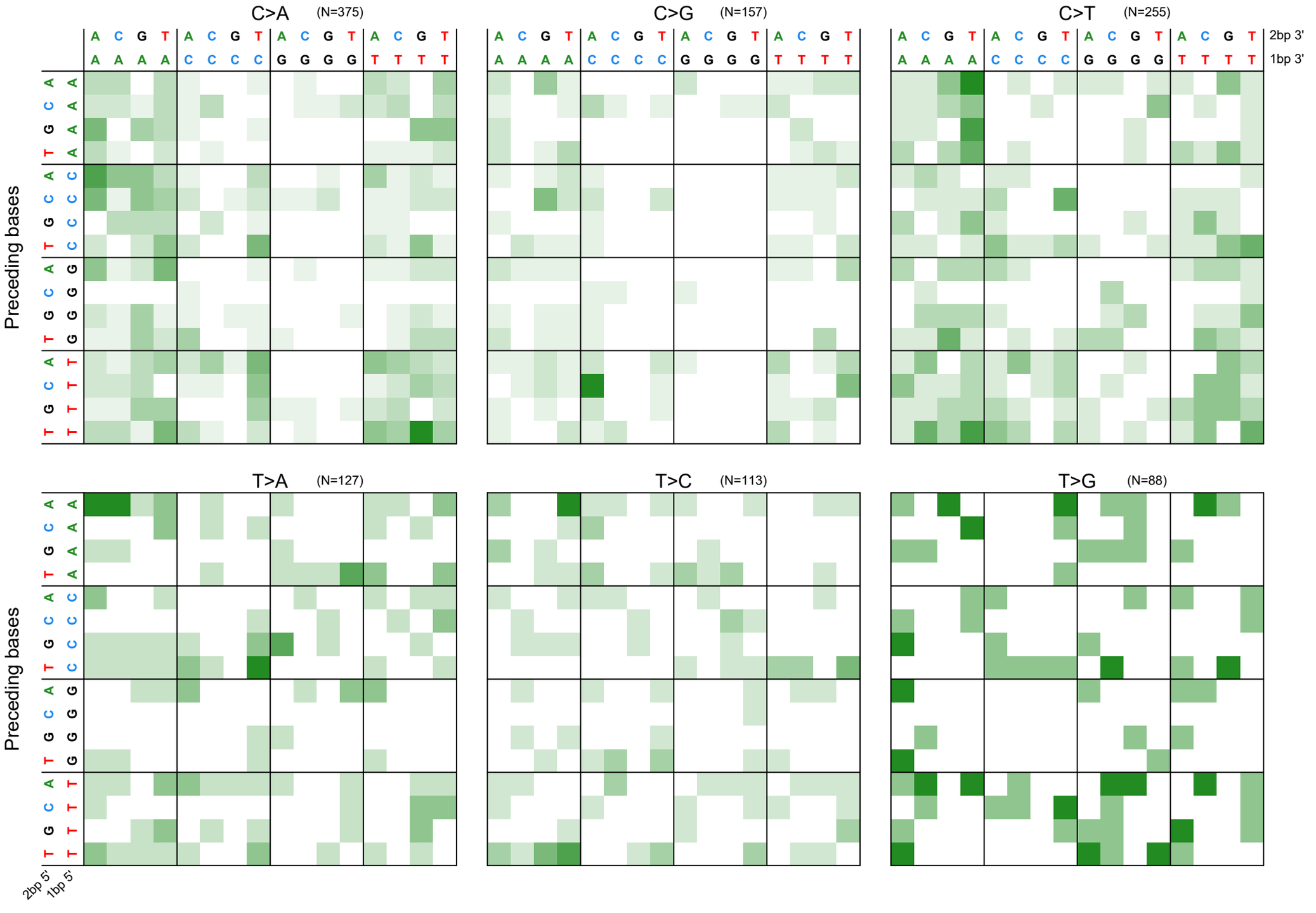
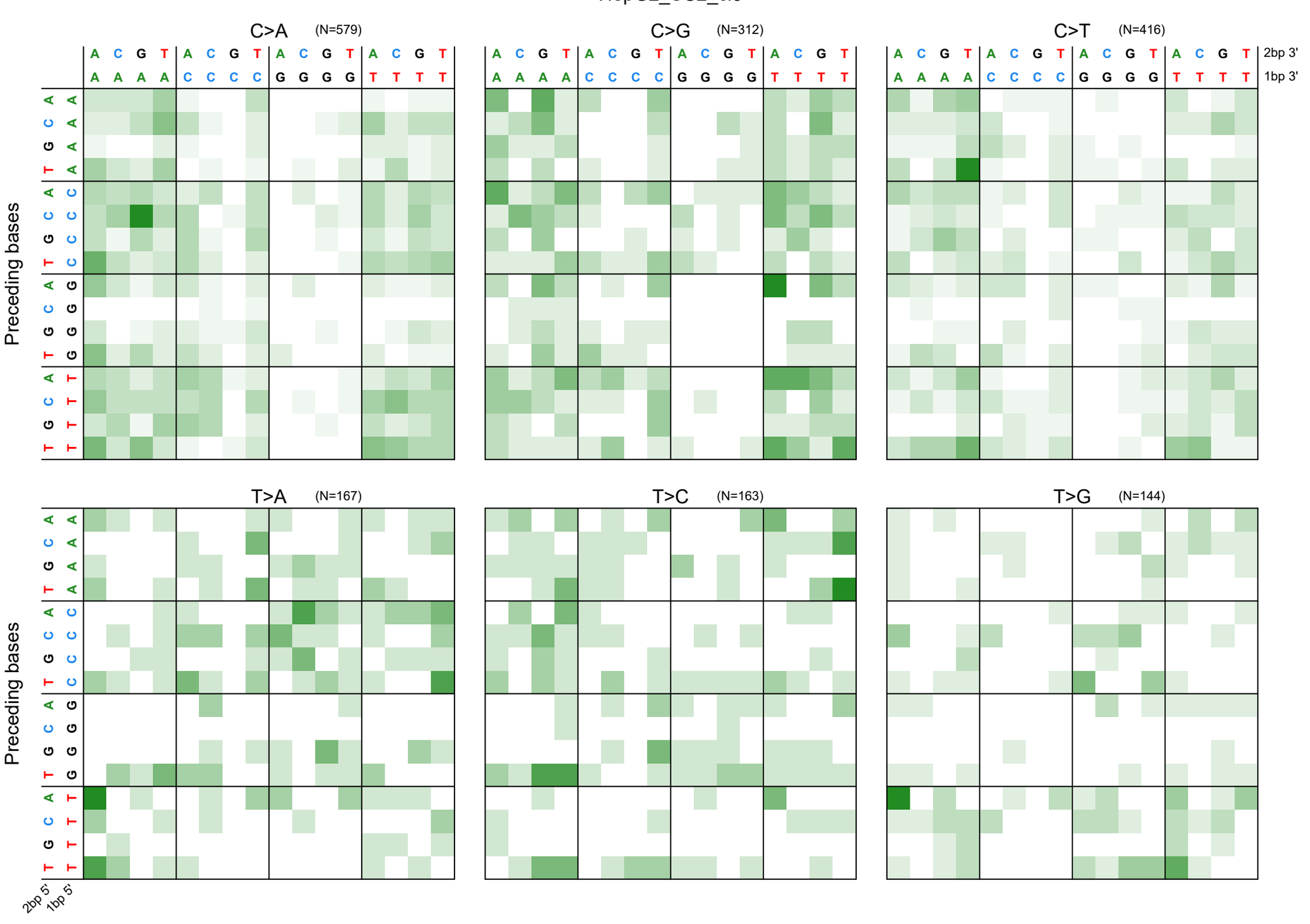
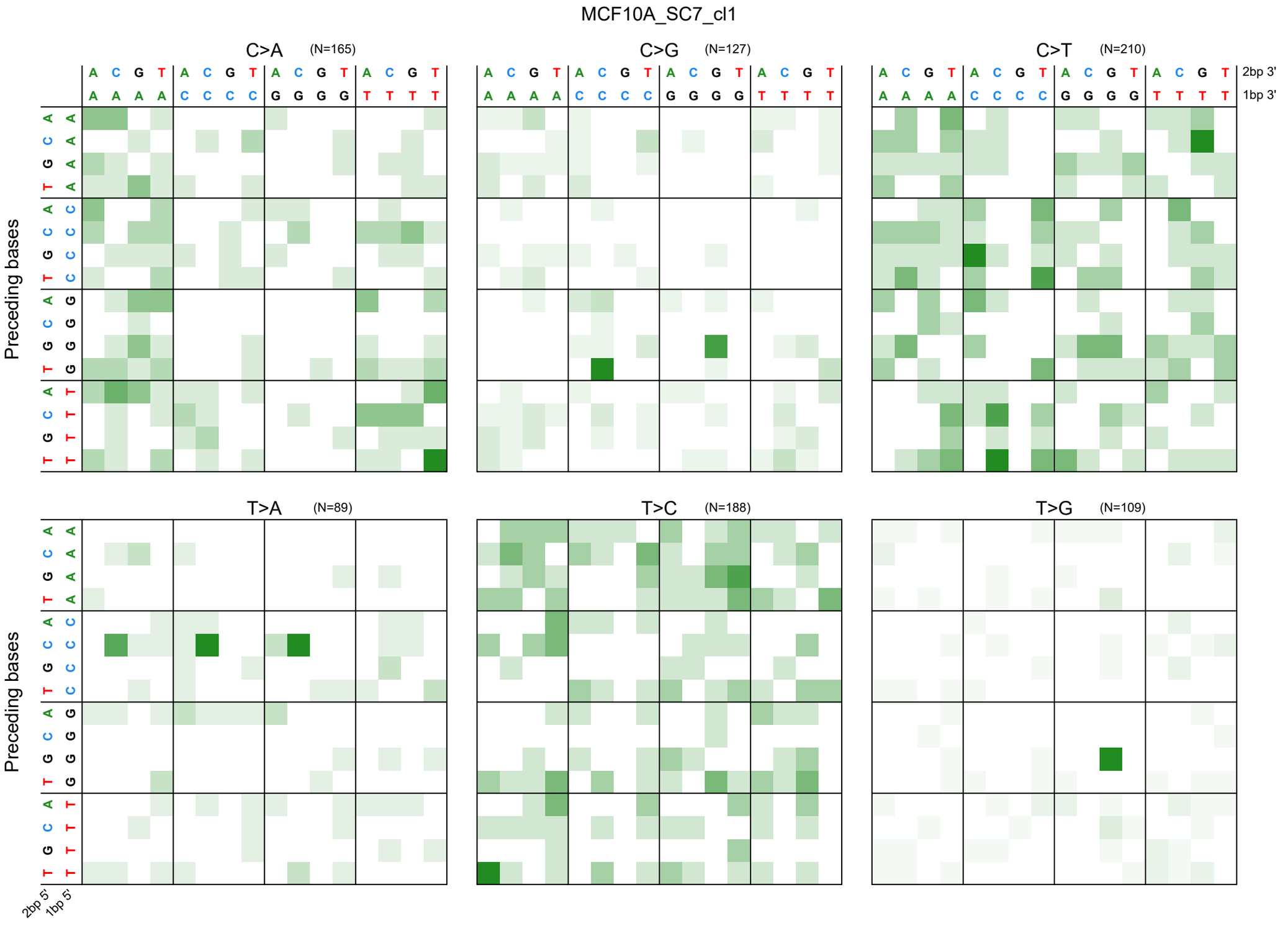
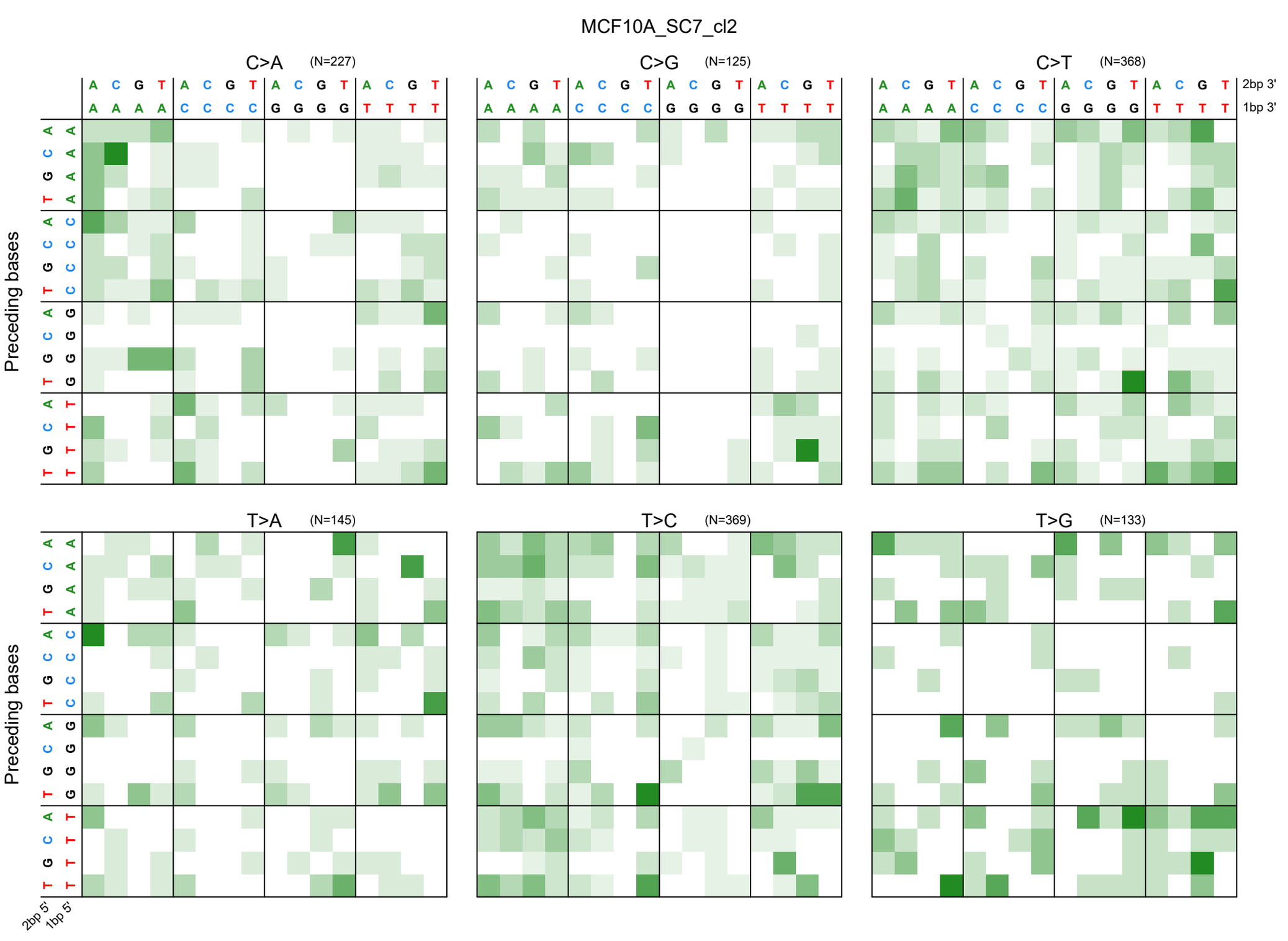
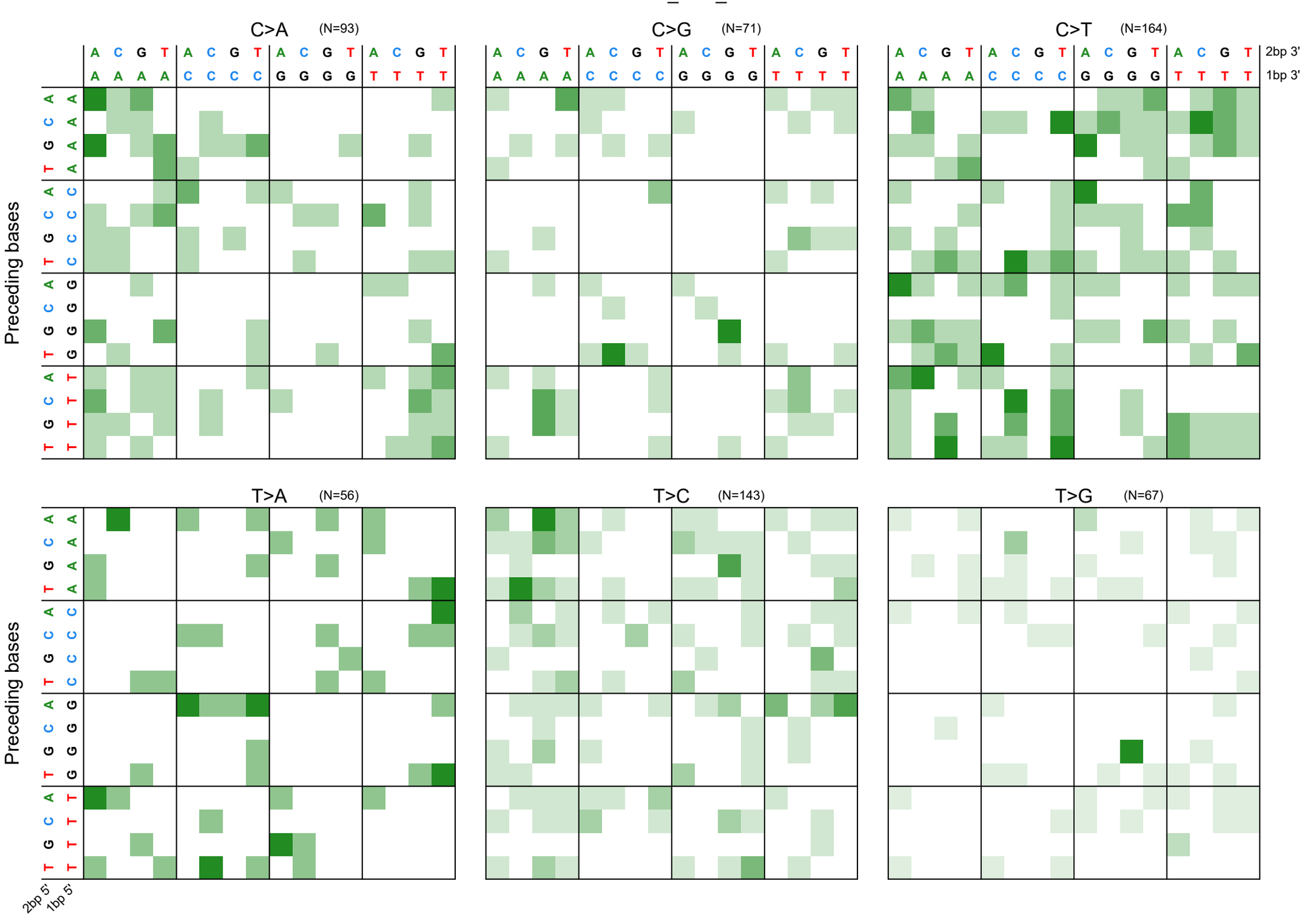

HepG2_SC2_cl1
C>A (N=545)
C>G (N=255)
C>T (N=384)
T>A (N=197)
T>C (N=167)
T>G (N=158)
Preceding bases
2bp 3'
1bp 3'
2bp 5'
1bp 5'


HepG2_SC2_cl2
C>A (N=375)
C>G (N=157)
C>T (N=255)
T>A (N=127)
T>C (N=113)
T>G (N=88)
2bp 3'
1bp 3'
Preceding bases
2bp 5'
1bp 5'


HepG2_SC2_cl3
C>A (N=579)
C>G (N=312)
C>T (N=416)
T>A (N=167)
T>C (N=163)
T>G (N=144)
Preceding bases
2bp 3'
1bp 3'
2bp 5'
1bp 5'


MCF10A_SC7_cl1
C>A (N=165)
C>G (N=127)
C>T (N=210)
T>A (N=89)
T>C (N=188)
T>G (N=109)
Preceding bases
2bp 3'
1bp 3'
2bp 5'
1bp 5'


MCF10A_SC7_cl2
C>A (N=227)
C>G (N=125)
C>T (N=368)
T>A (N=145)
T>C (N=369)
T>G (N=133)
Preceding bases
2bp 3'
1bp 3'
2bp 5'
1bp 5'


MCF10A_SC7_cl3
C>A (N=93)
C>G (N=71)
C>T (N=164)
T>A (N=56)
T>C (N=143)
T>G (N=67)
Preceding bases
2bp 3'
1bp 3'
2bp 5'
1bp 5'