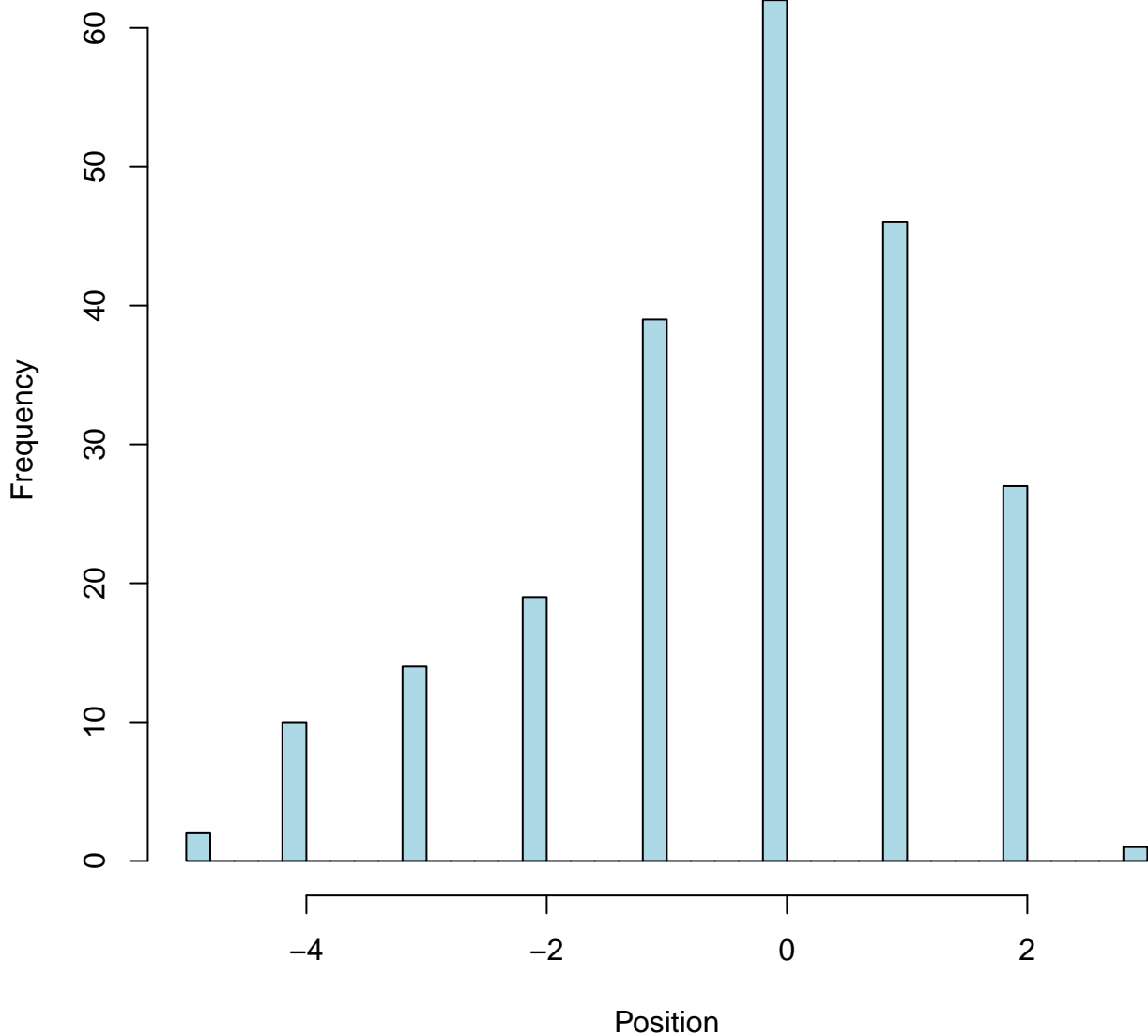
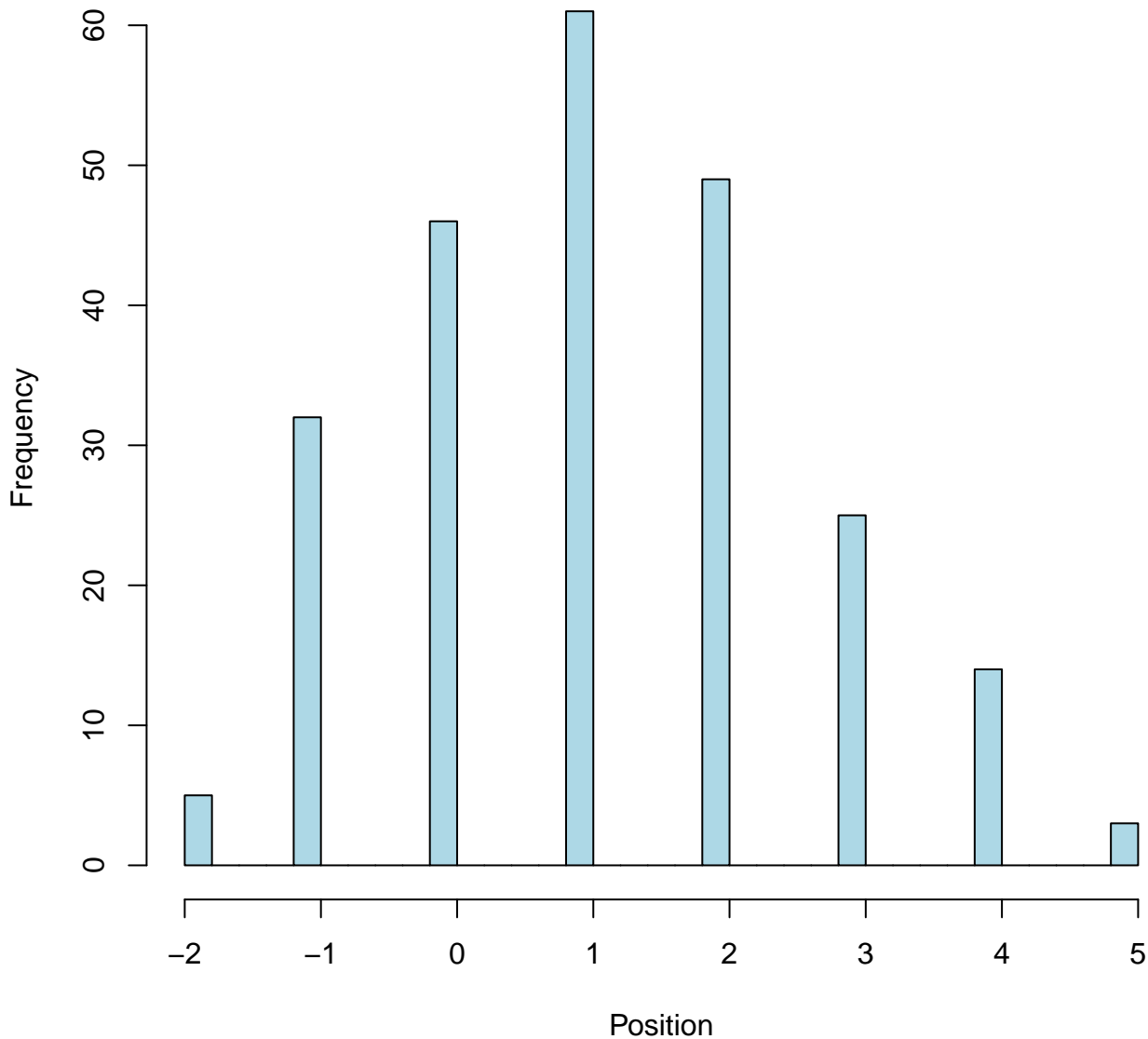


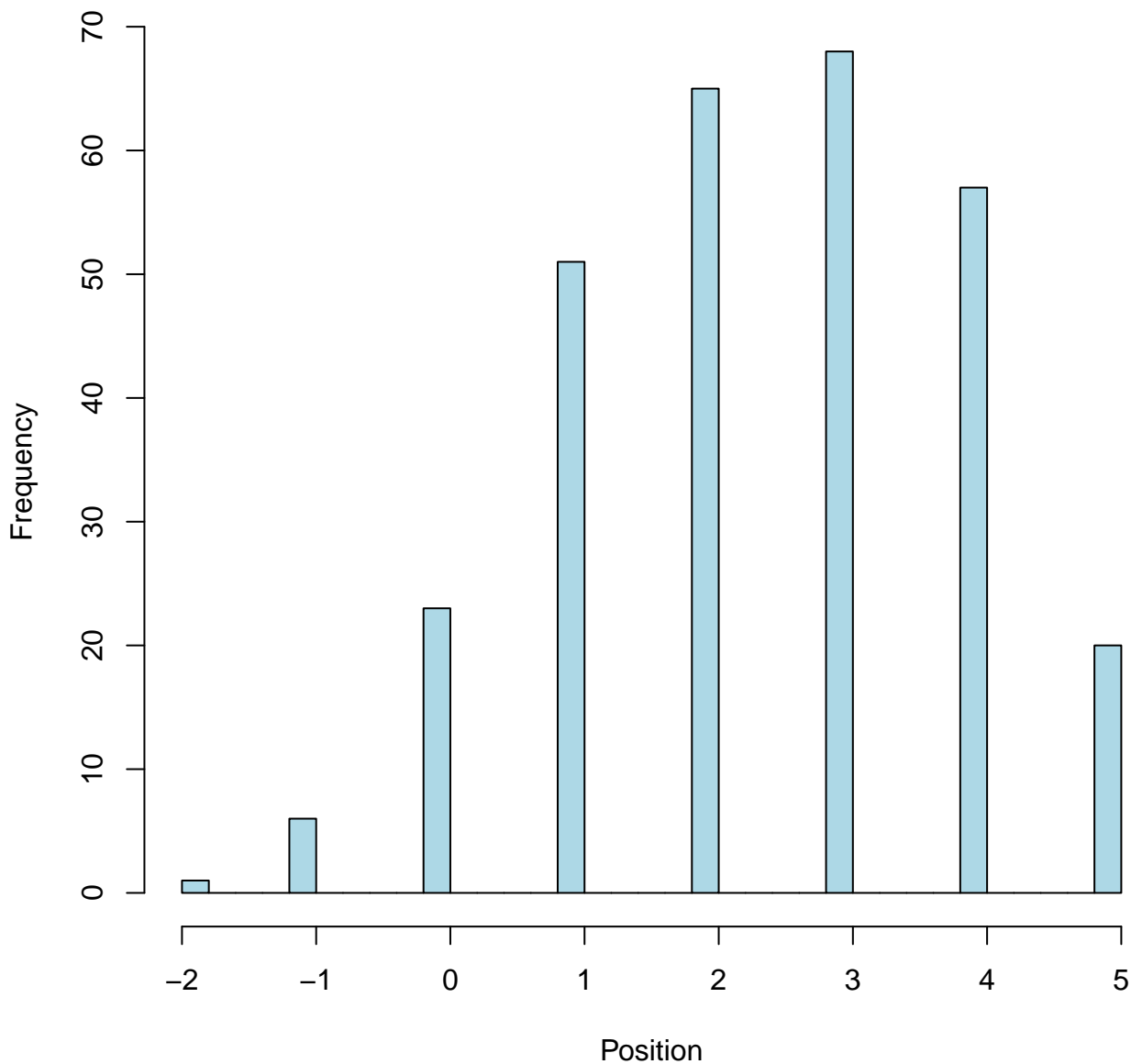
Interacting aa position relative to the center in peptide (alpha)



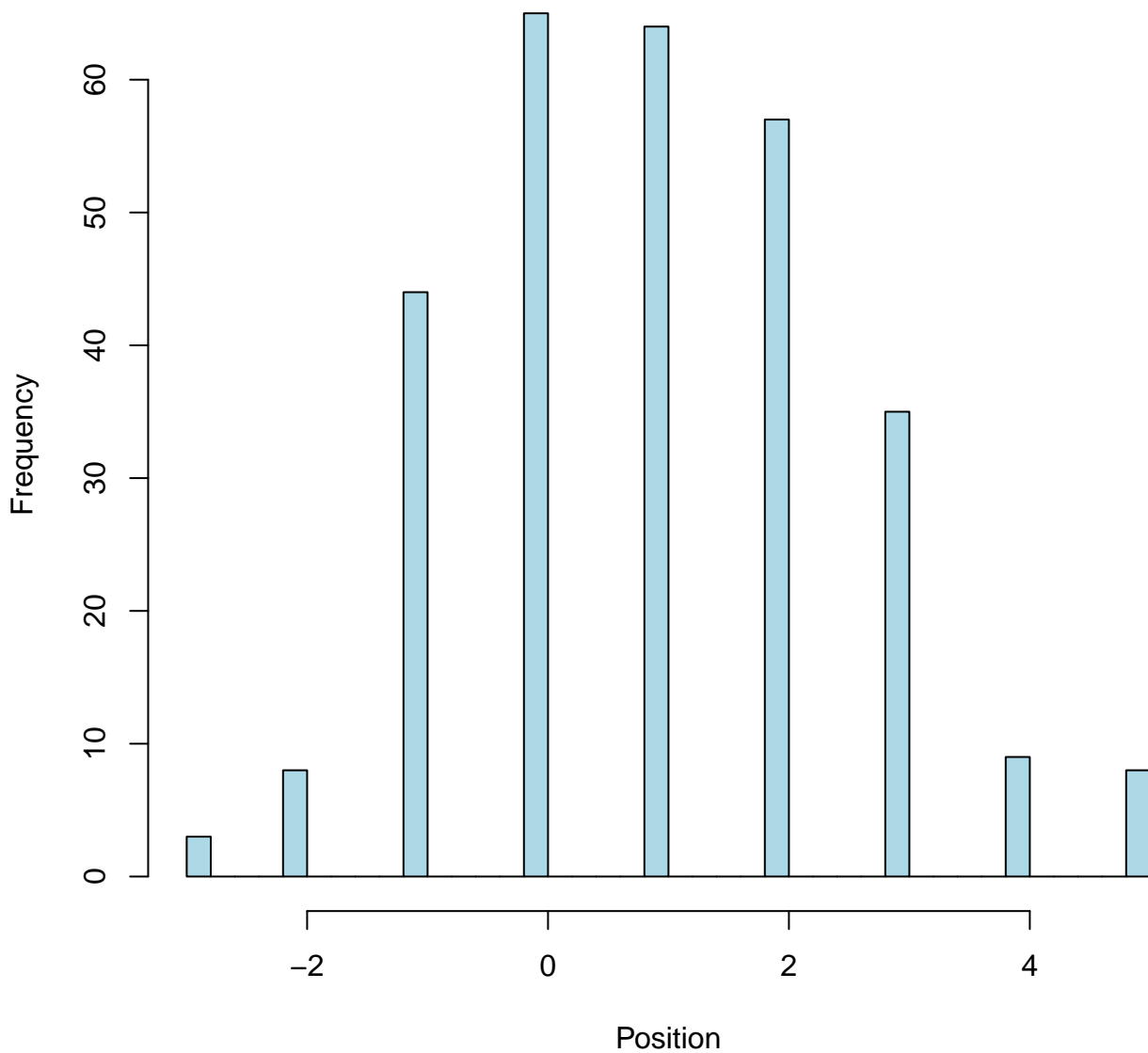
Interacting aa position relative to the center in cdr3 (alpha)

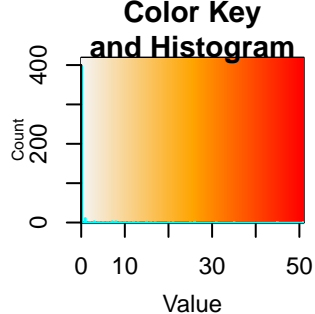


Interacting aa position relative to the center in peptide (beta)

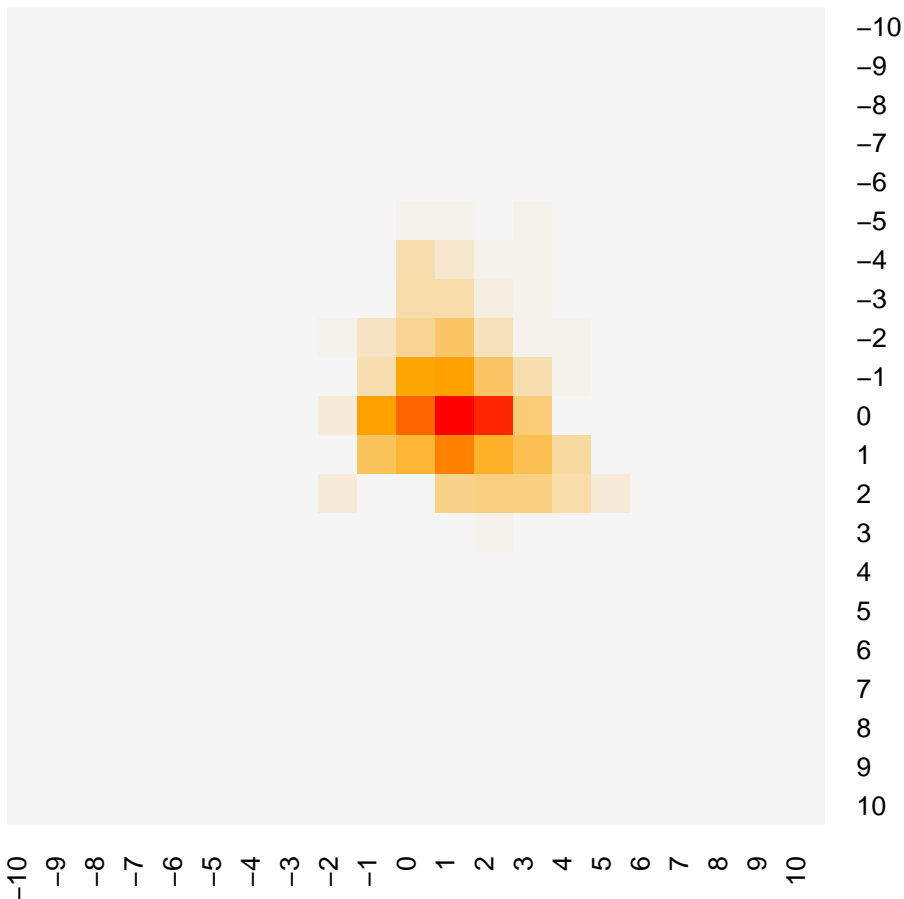


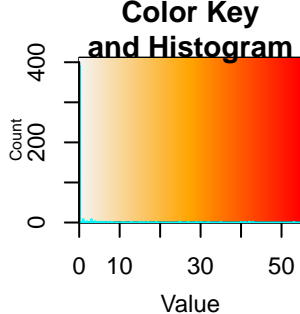
Interacting aa position relative to the center in cdr3 (beta)



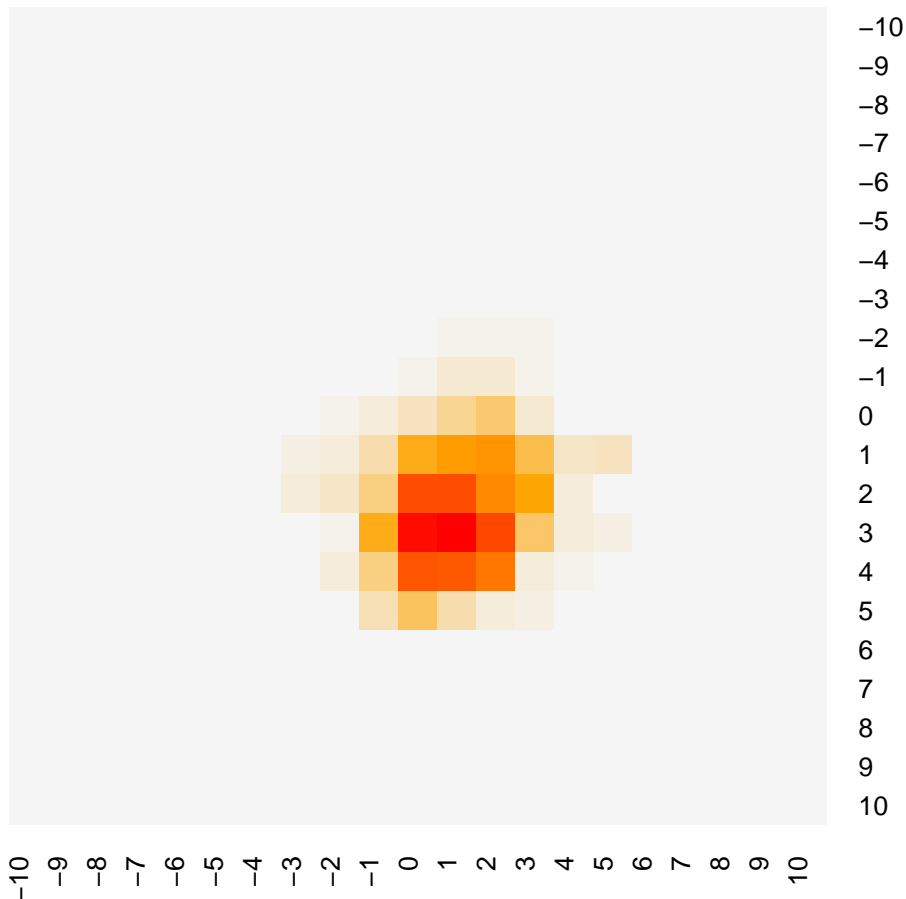


y – relative aa peptide position
x – relative aa CDR3 position (alpha)

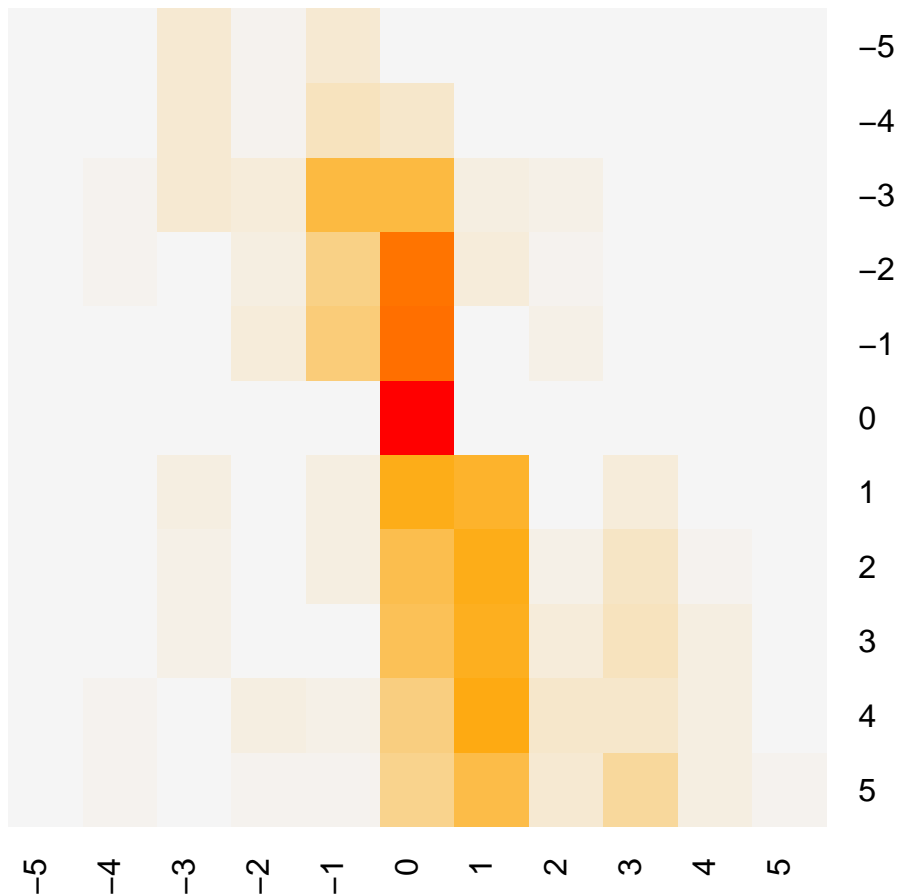
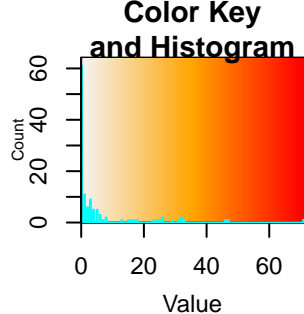


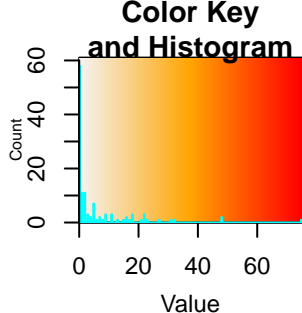


y – relative aa peptide position
x – relative aa CDR3 position (beta)

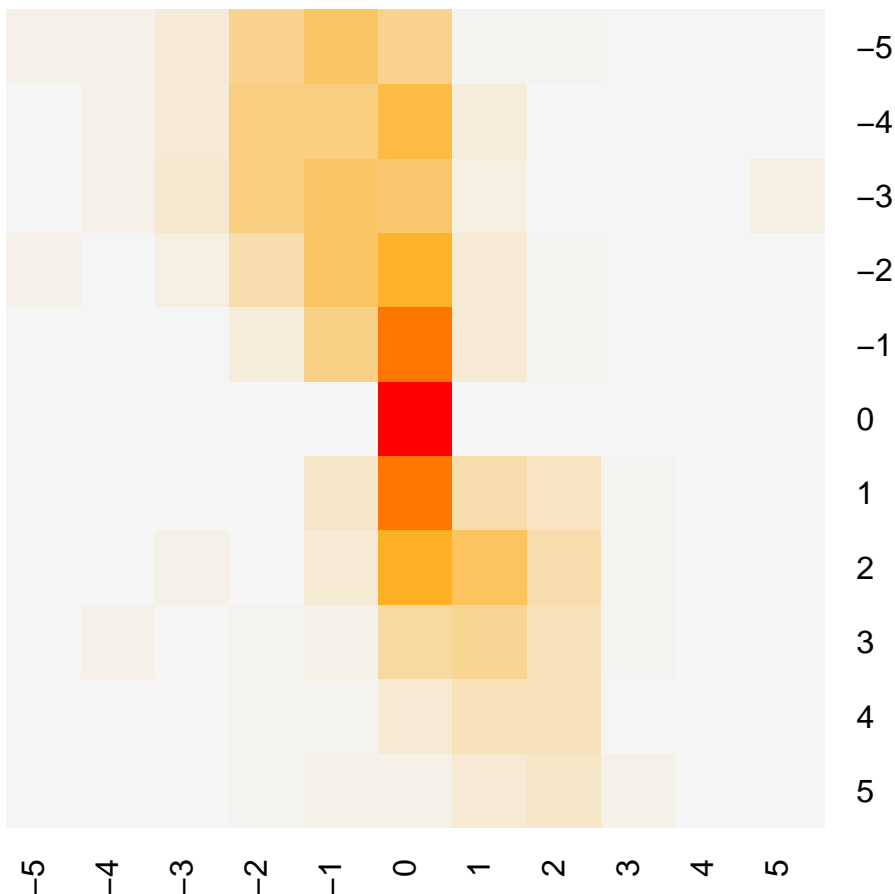


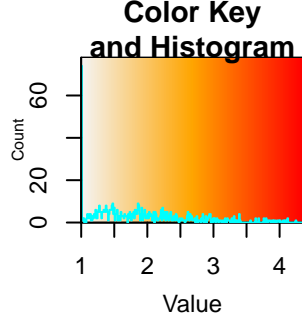
**Relative position from the
most interating CDR3 aa**
y – peptide aa pos; x – CDR3 (alpha)



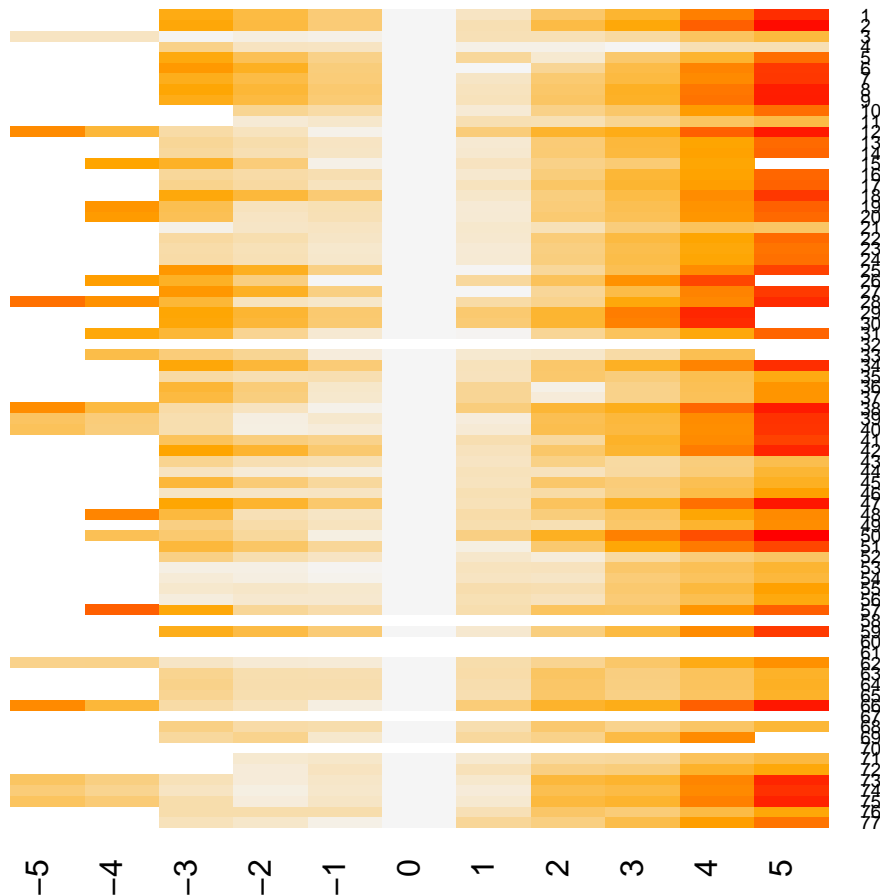


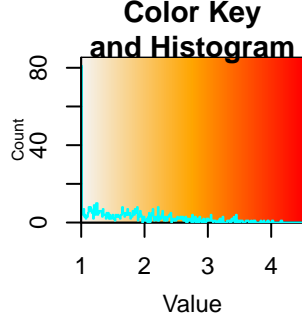
**Relative position from the
most interating CDR3 aa**
y – peptide aa pos; x – CDR3 (beta)



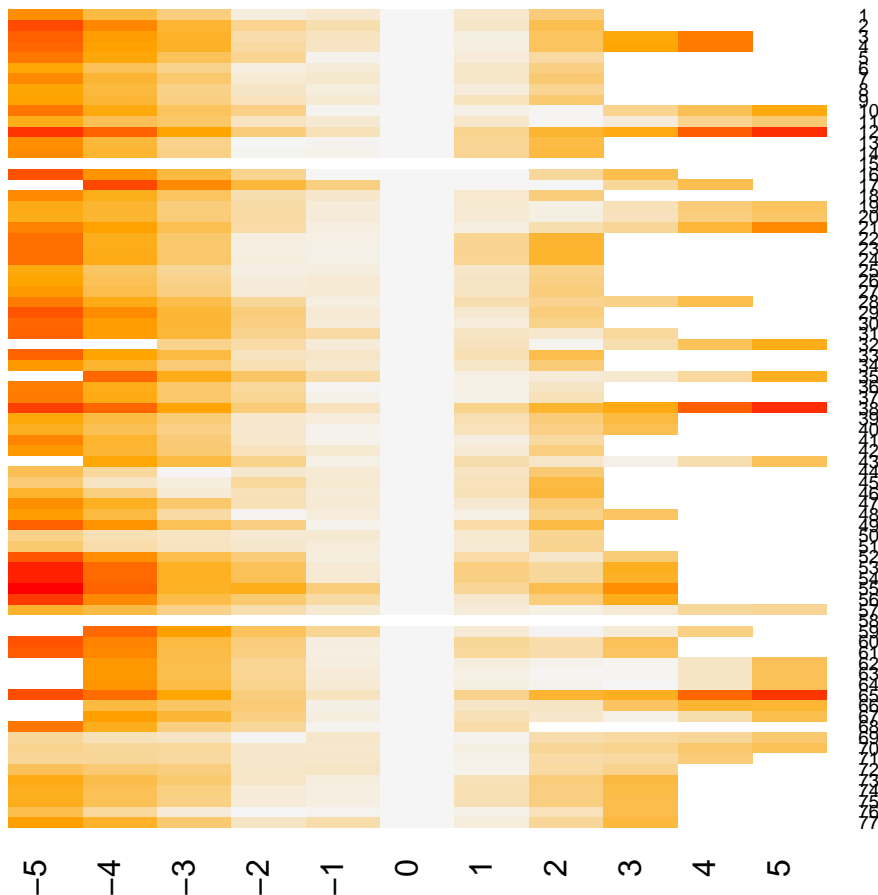


**y – Distance from peptide aa in
position 'x' form the center to
CDR3; x – Peptide aa position (alpha)**

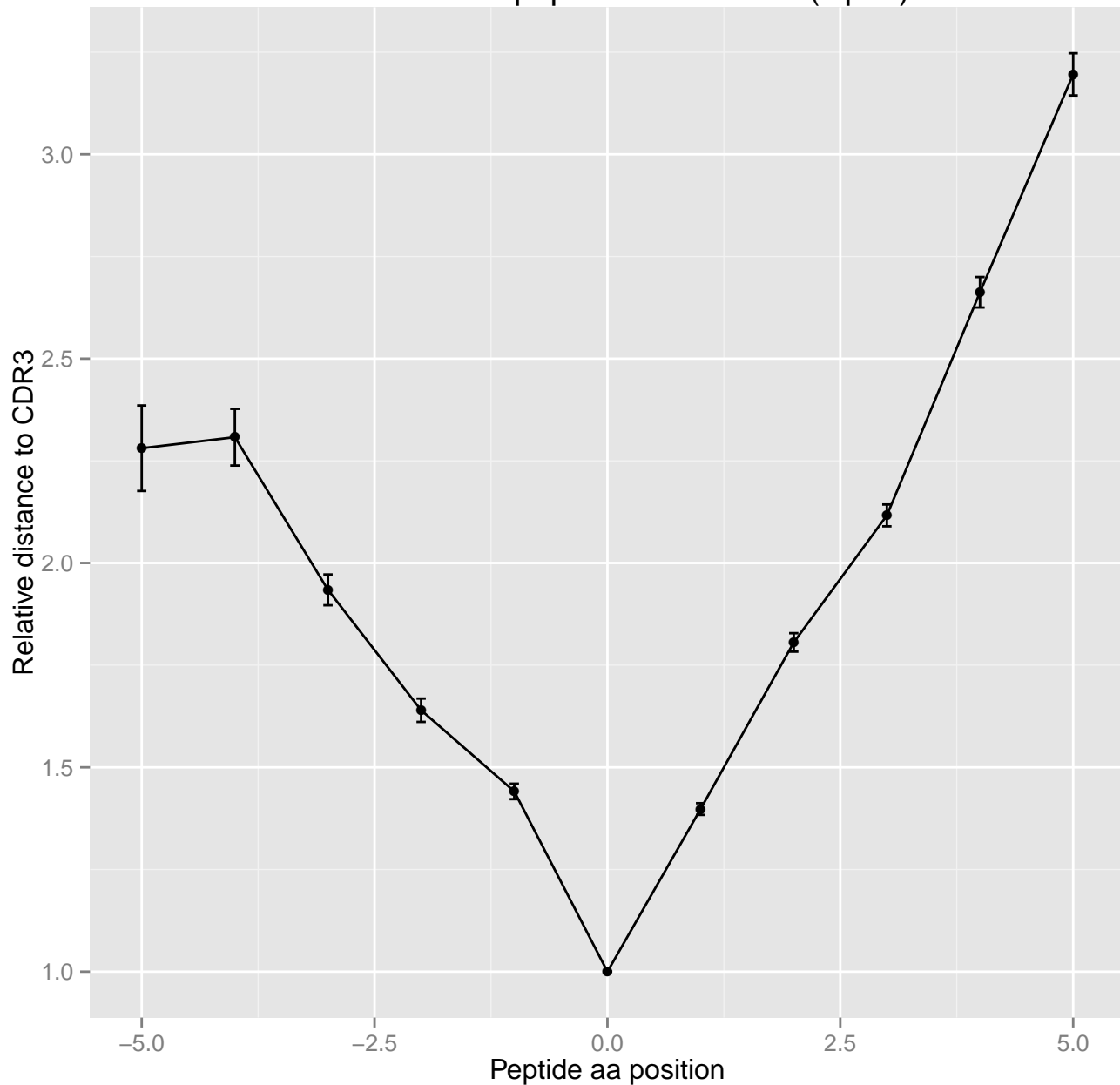




y – Distance from peptide aa in position 'x' from the center to CDR3; x – Peptide aa position (beta)



Distance from peptide aa to CDR3 (alpha)



Distance from peptide aa to CDR3 (beta)

