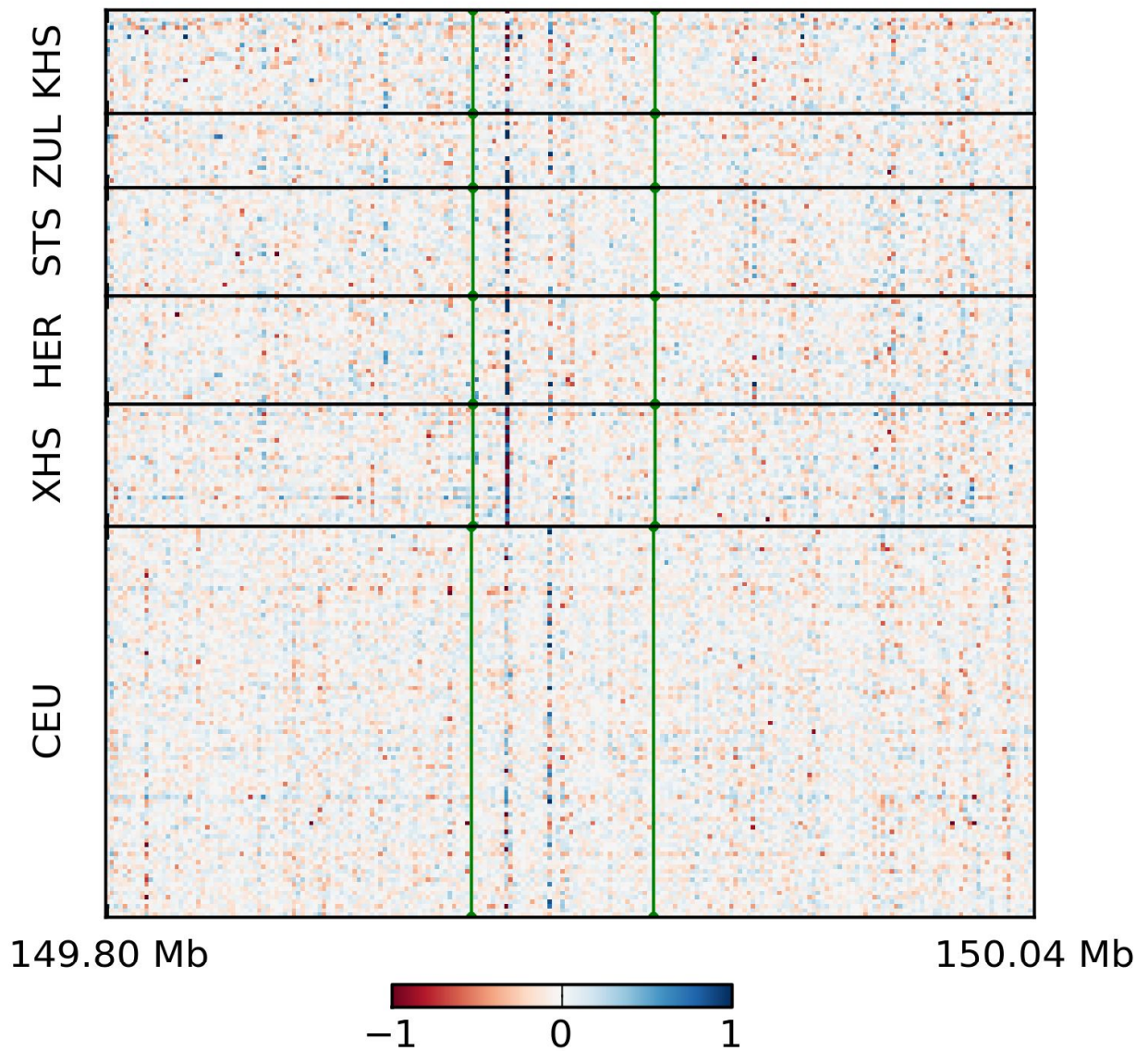
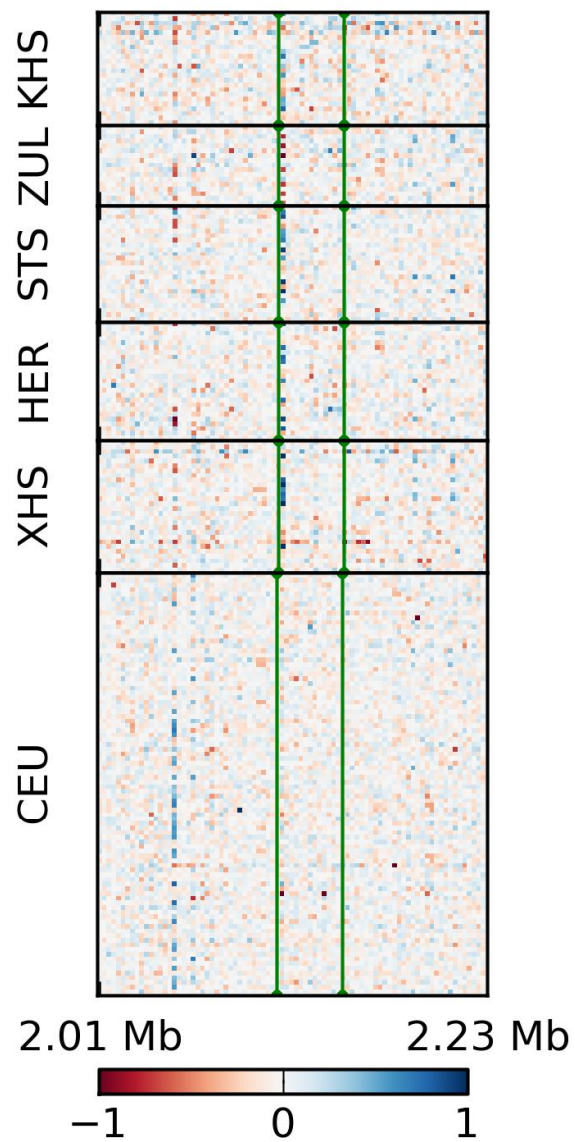


**Supplementary Data File S-6:** Heatmaps showing copy number variation analysis for candidate genes. Red shading shows a deletion at that position, blue shading shows a copy number gain at that position. HUGO gene symbol is shown. Chrm = chromosome. Green lines denote the gene start and end. Horizontal axis shows position on the chromosome. Vertical axis shows the individuals from each population: CEU = HapMap Central European population, XHS = Xhosa population, HER = Herero population, STS = Setswana population, ZUL = Zulu population, KHS = Khoisan population.

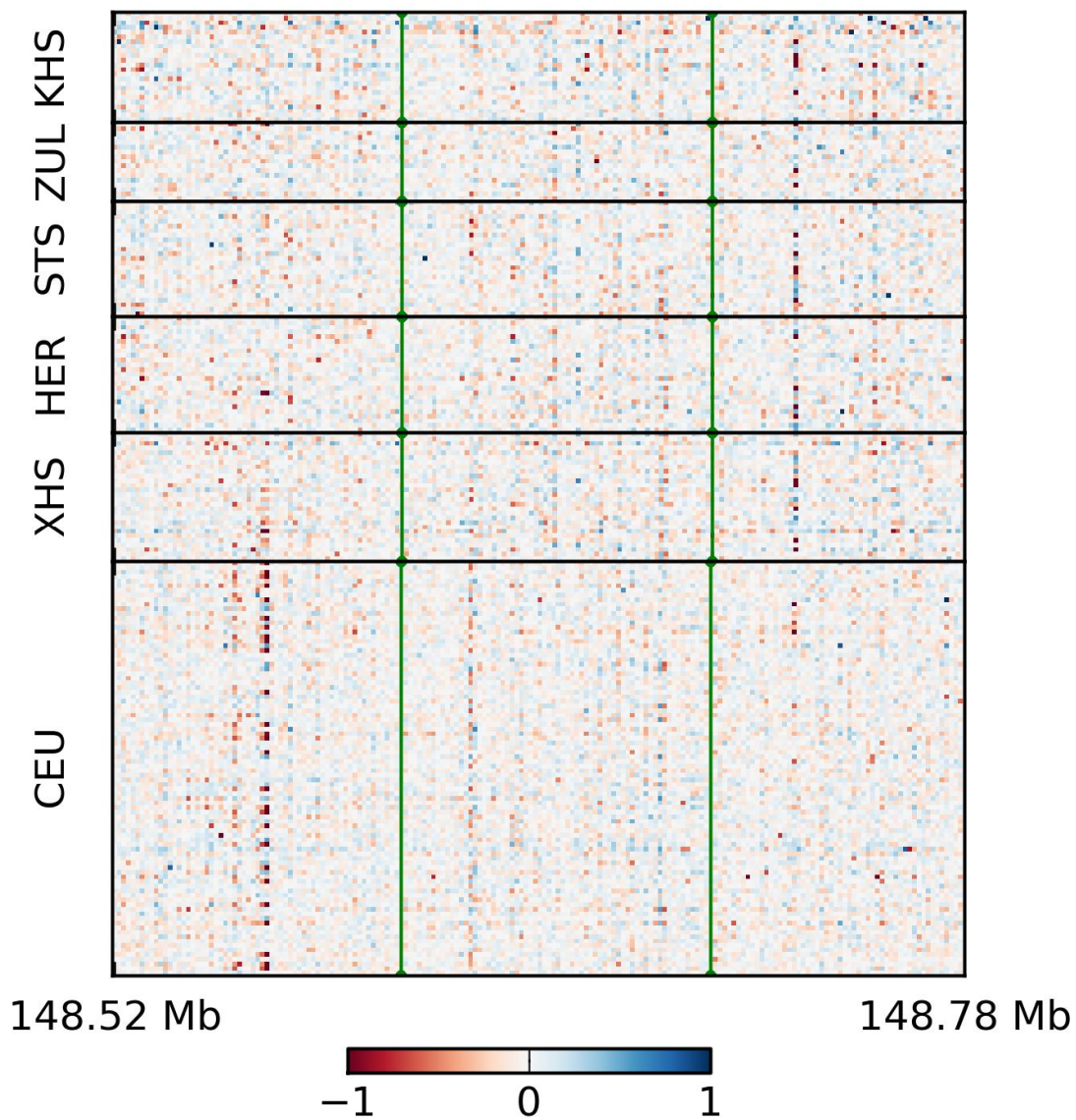
## AGTR1 (Chrm 3)



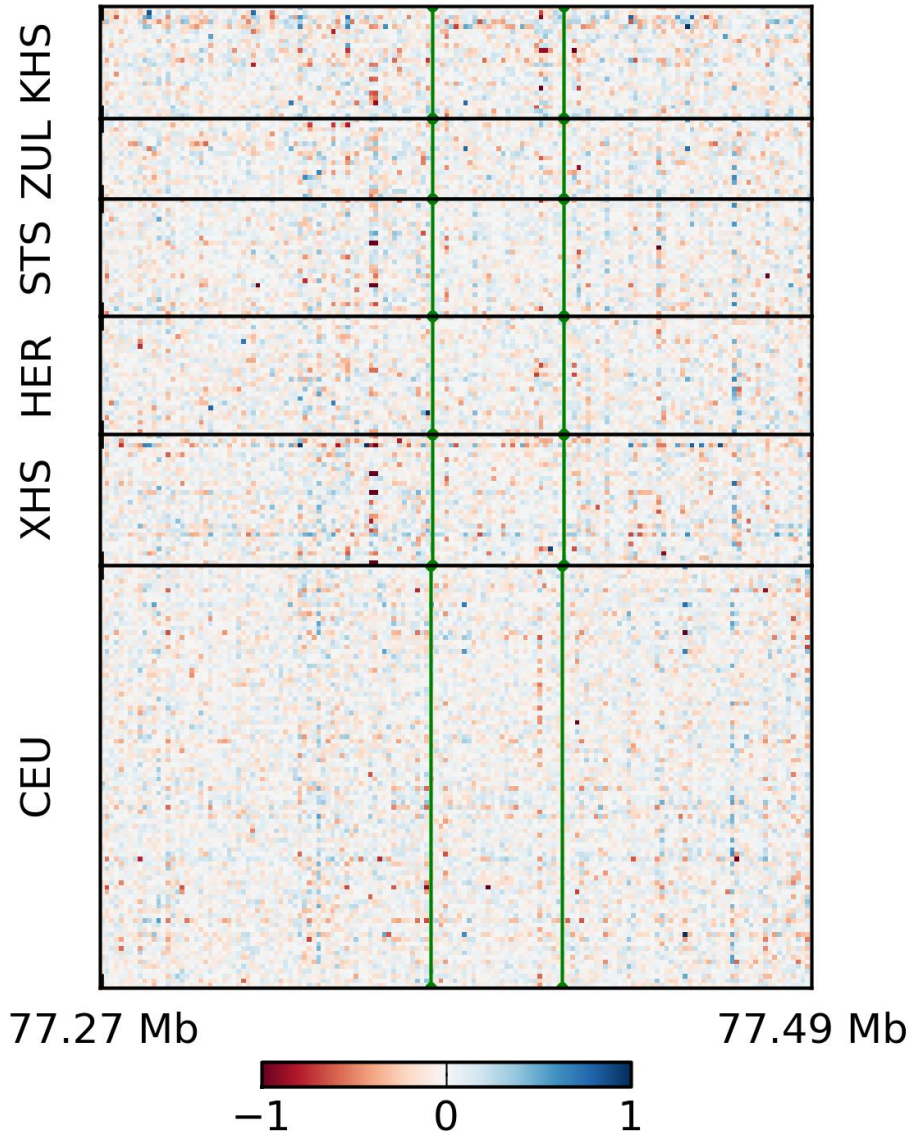
# INS (Chrm 11)



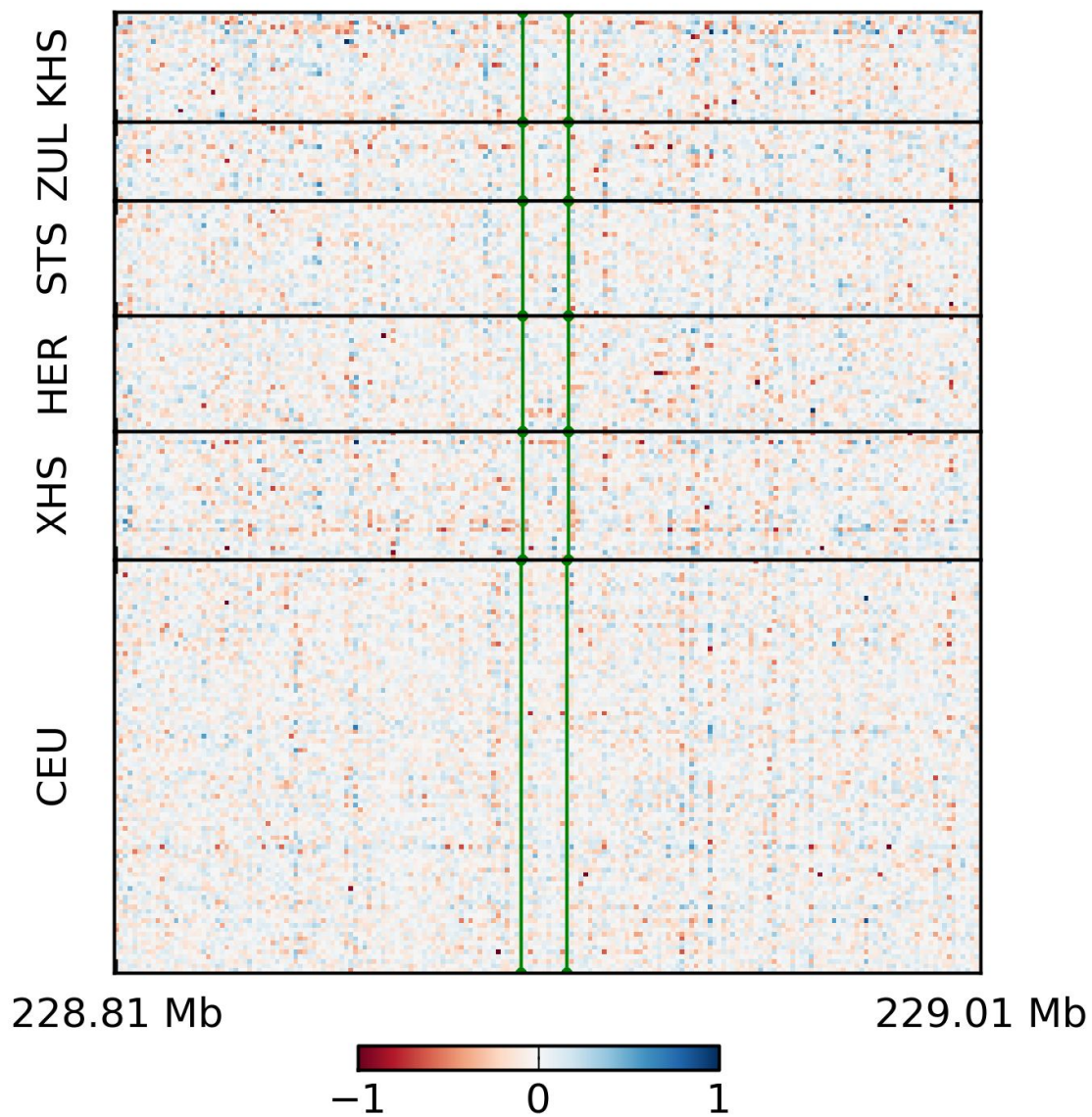
# EDNRA (Chrm 4)



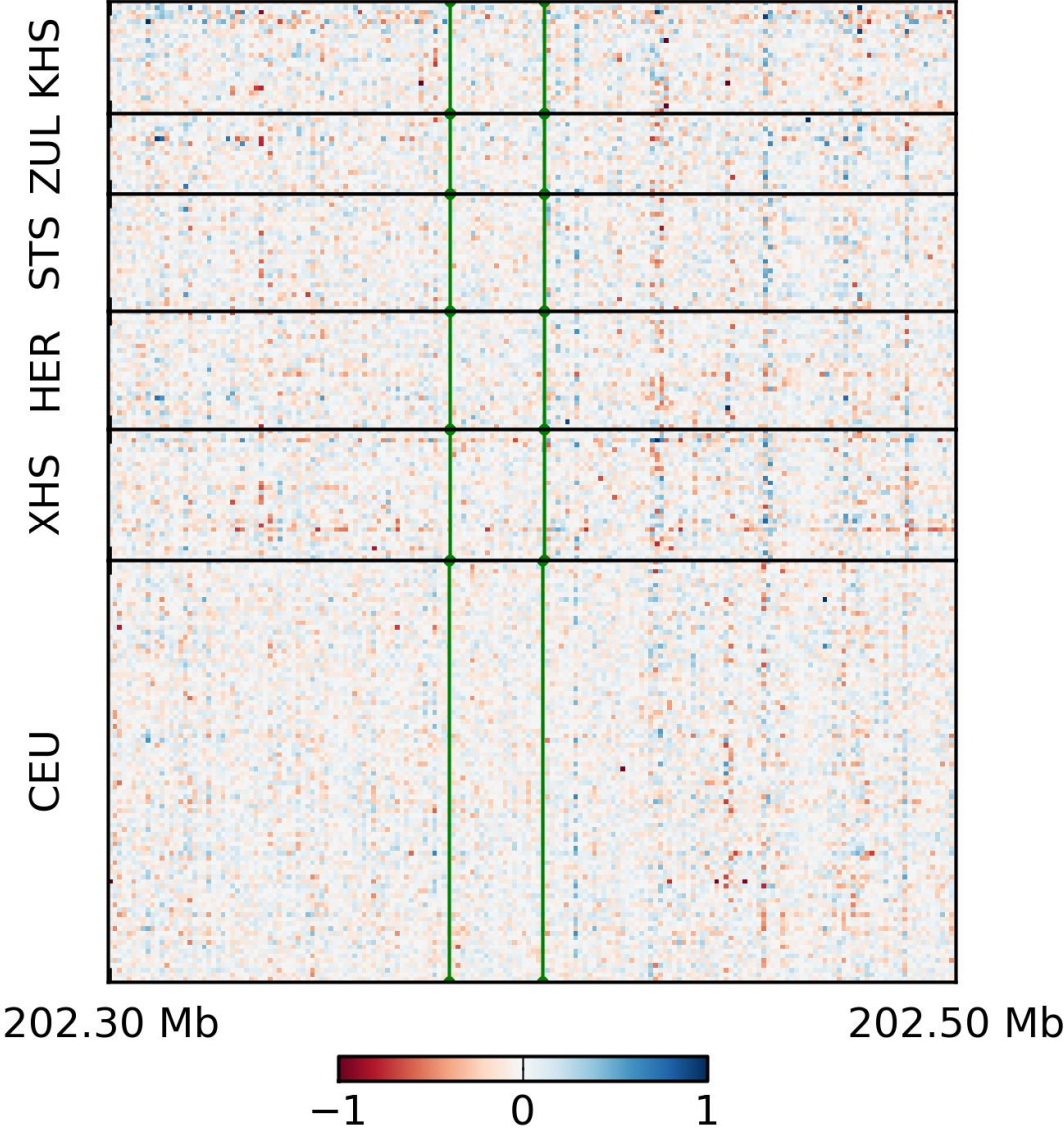
# EDNRB (Chrm 13)



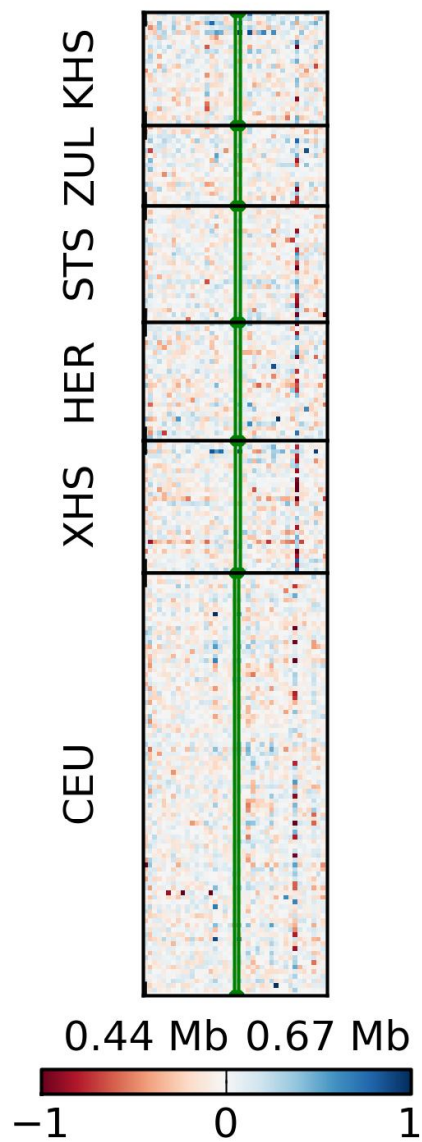
# AGT (Chrm 1)



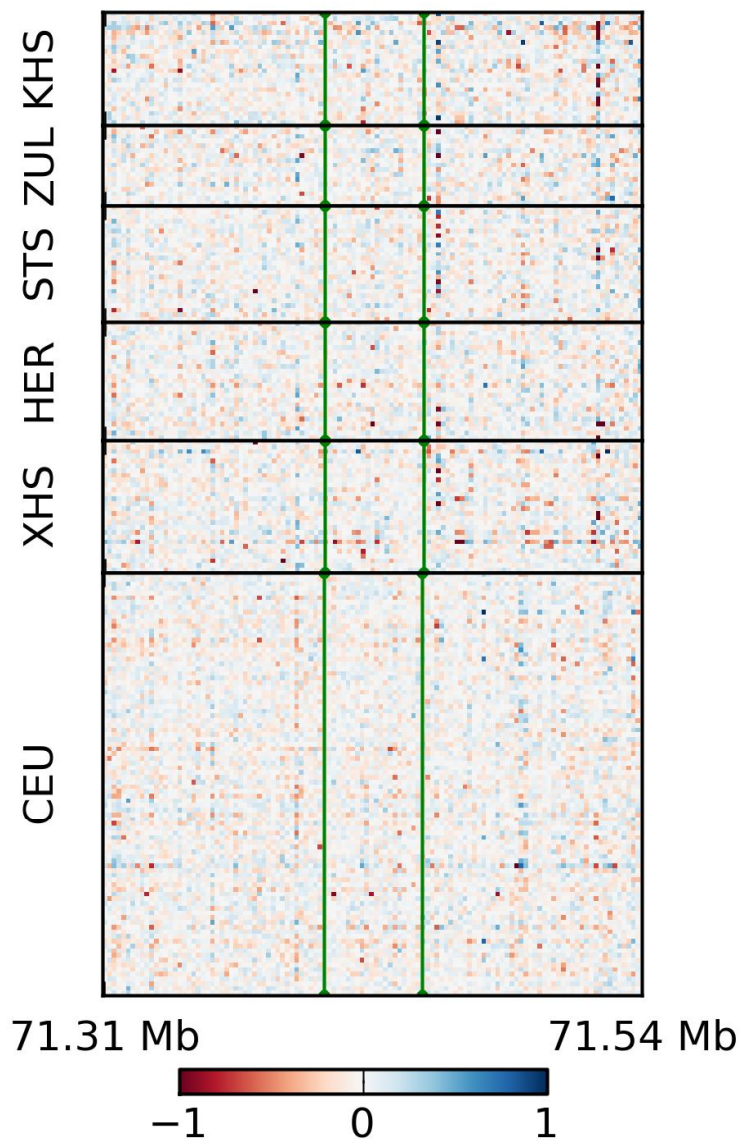
# REN (Chrm 1)



# HCN2 (Chrm 19)

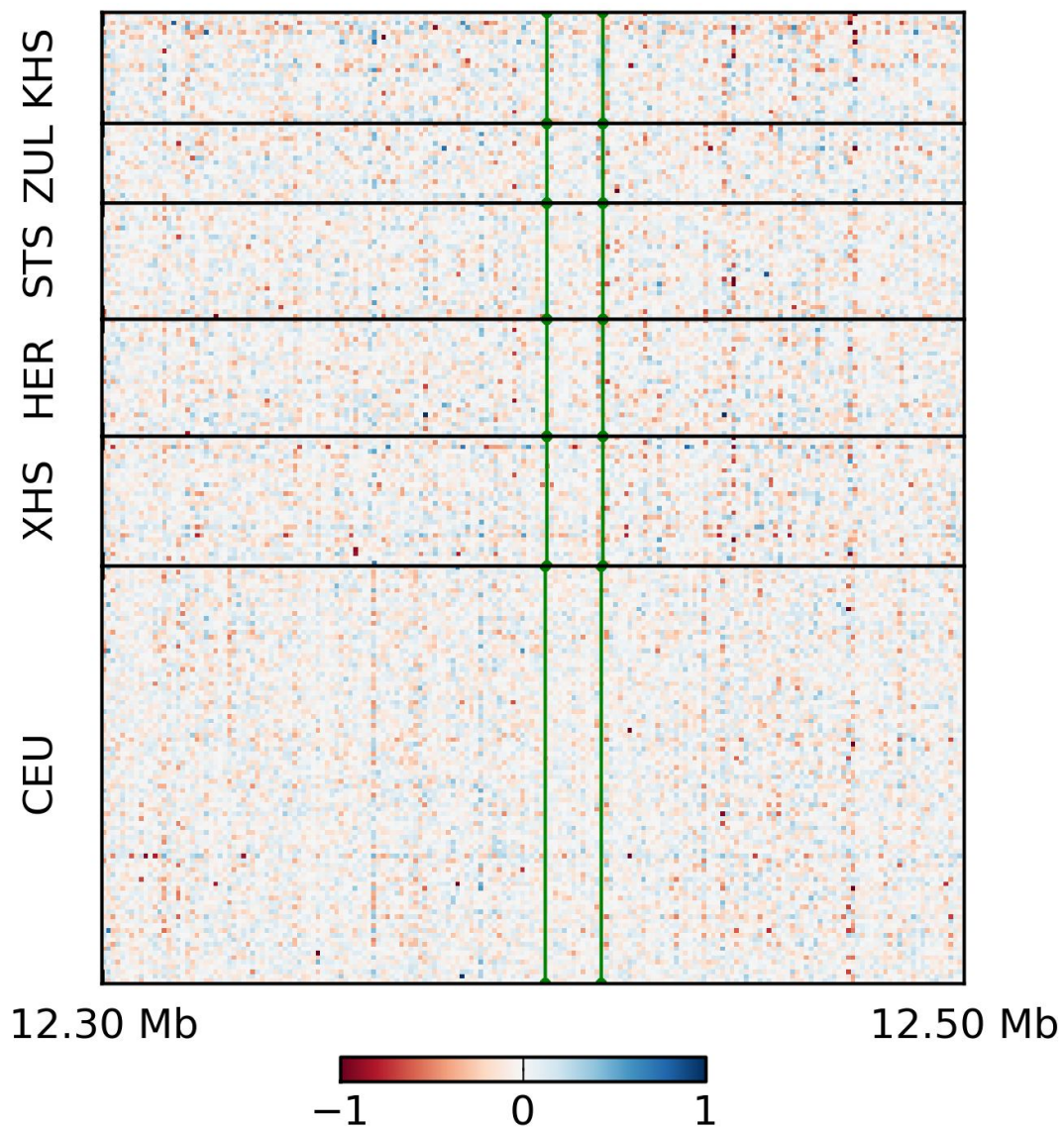


# HCN4 (Chrm 15)

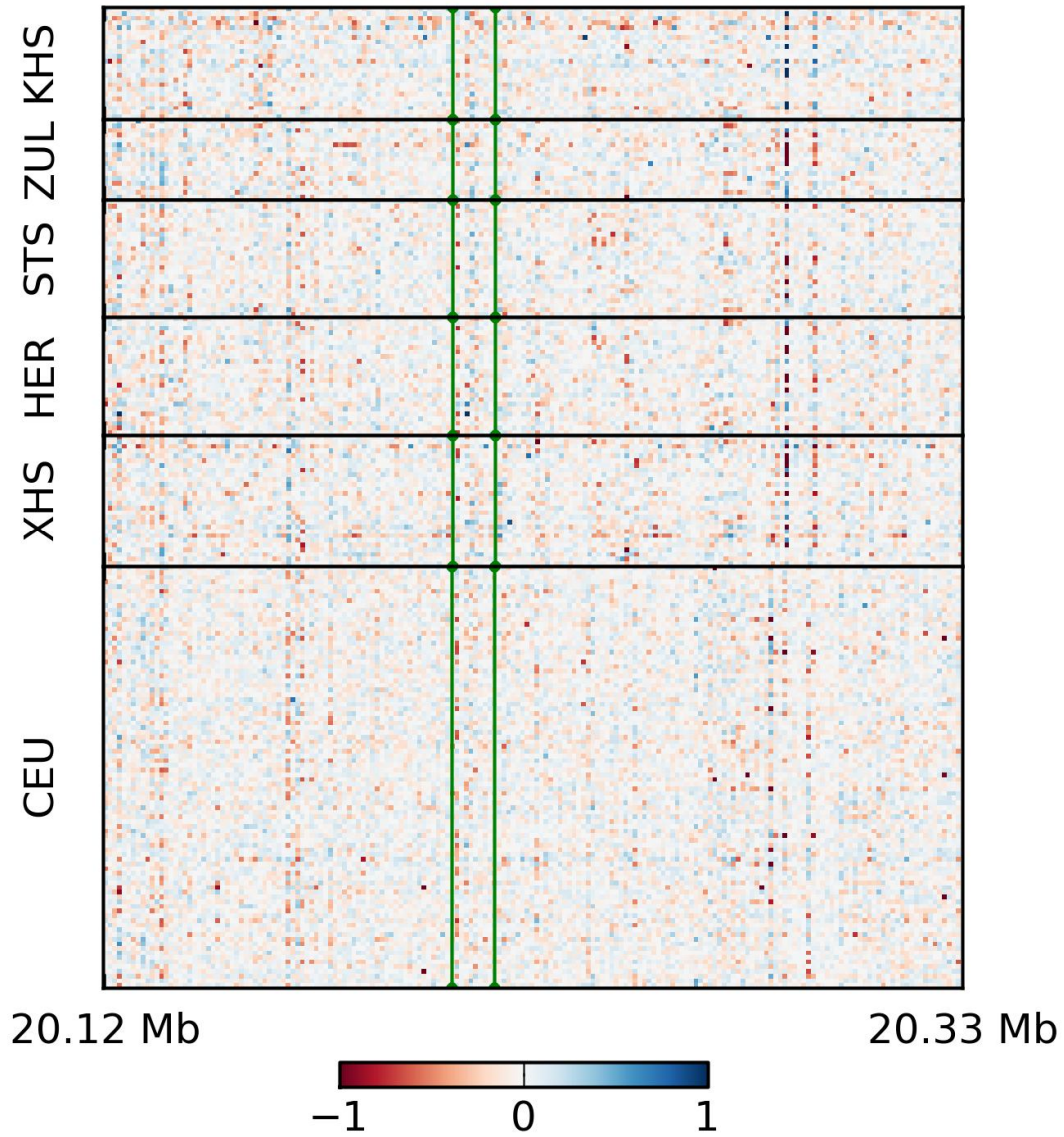




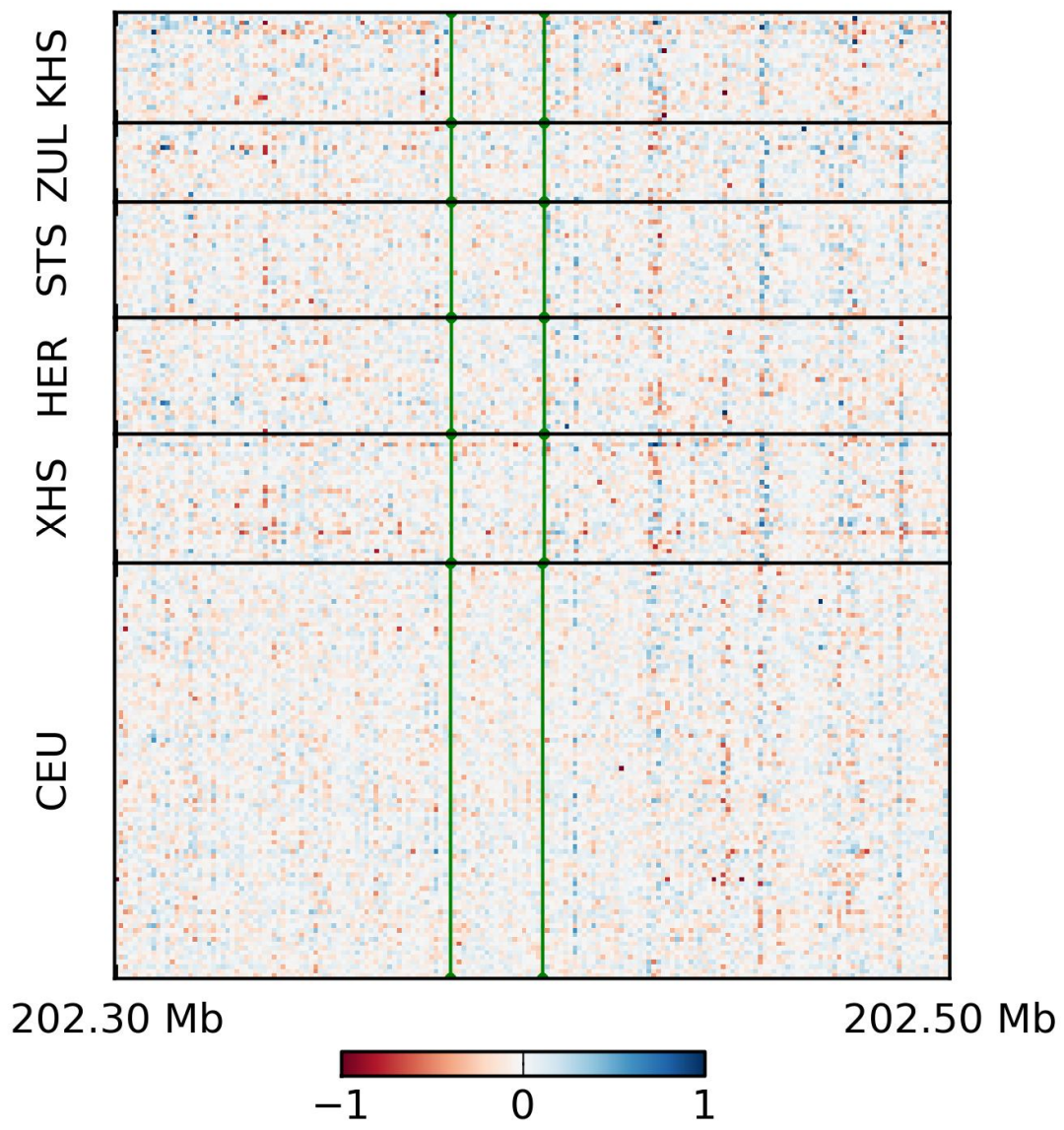
# EDN1 (Chrm 6)



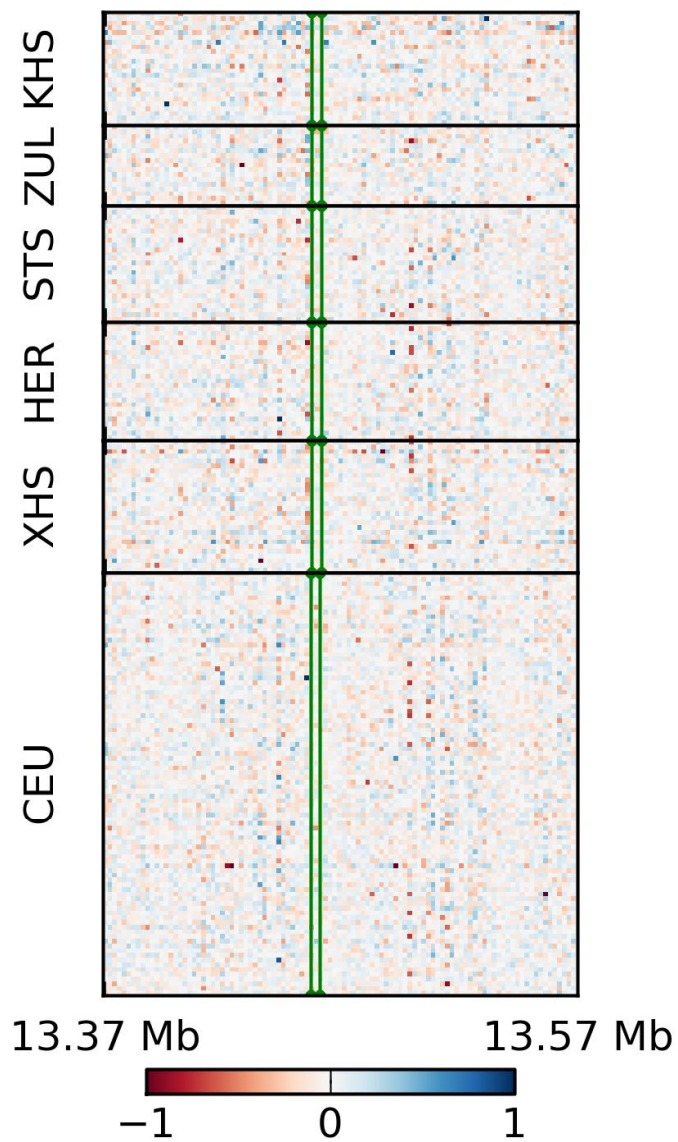
# ANG (Chrm 14)



# REN (Chrm 1)



# PTH (Chrm 11)



# NPPA (Chrm 1)

