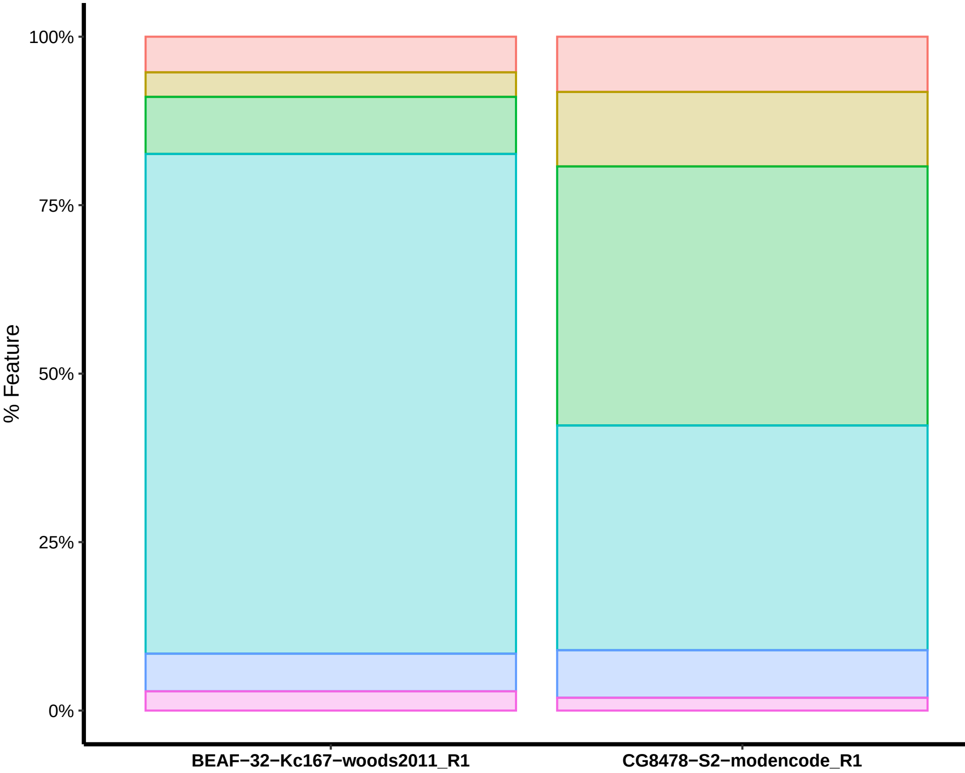
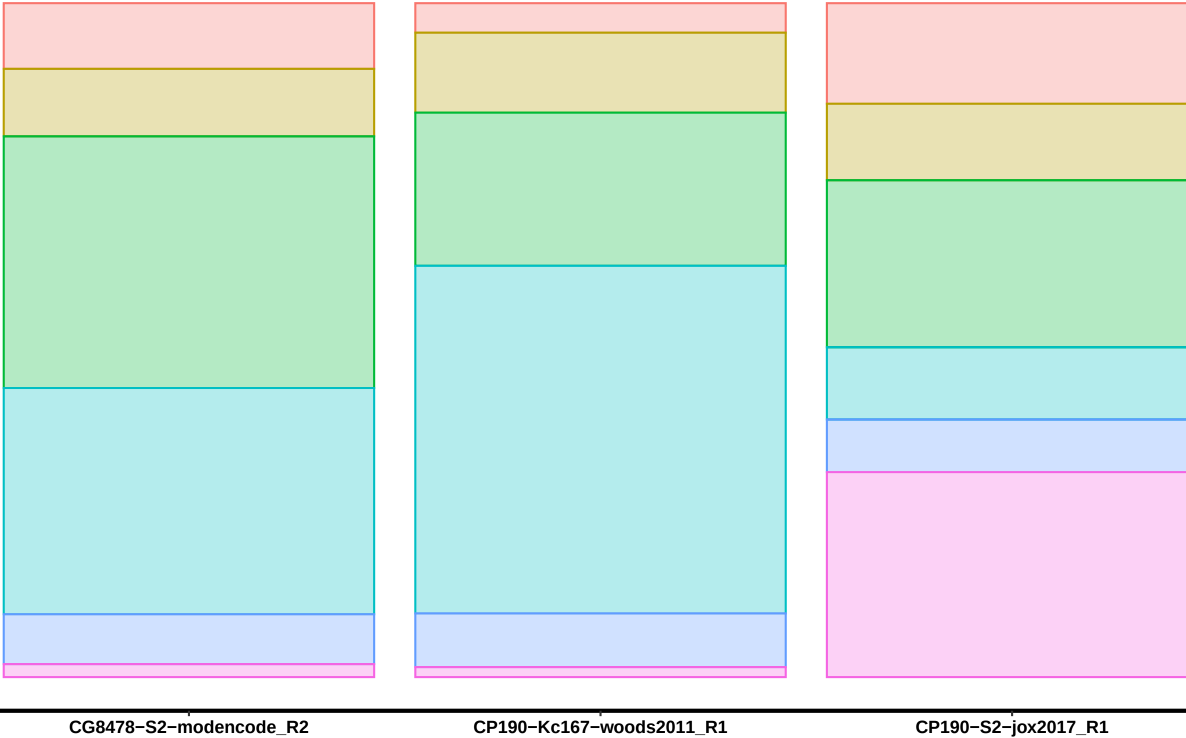
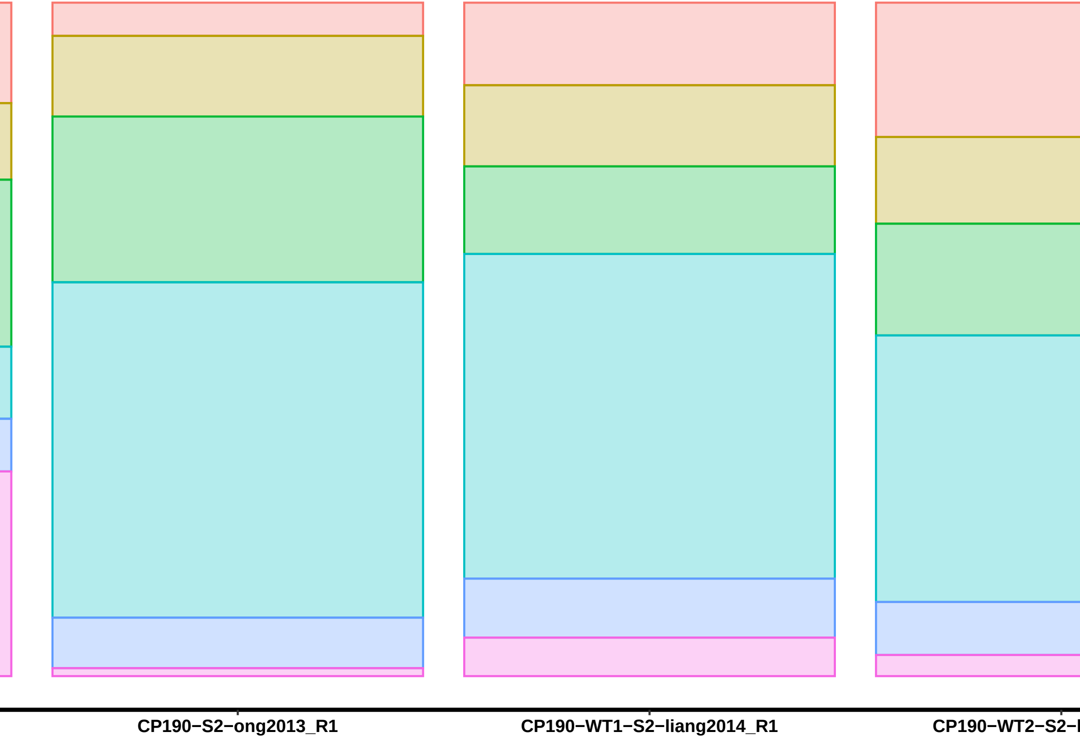
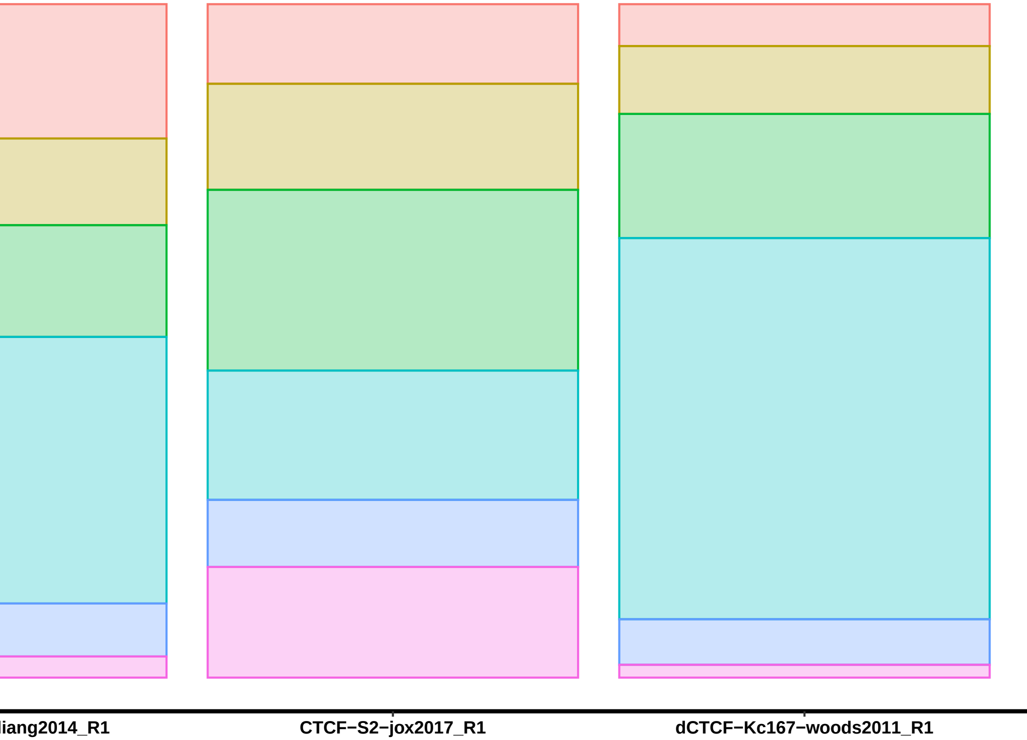
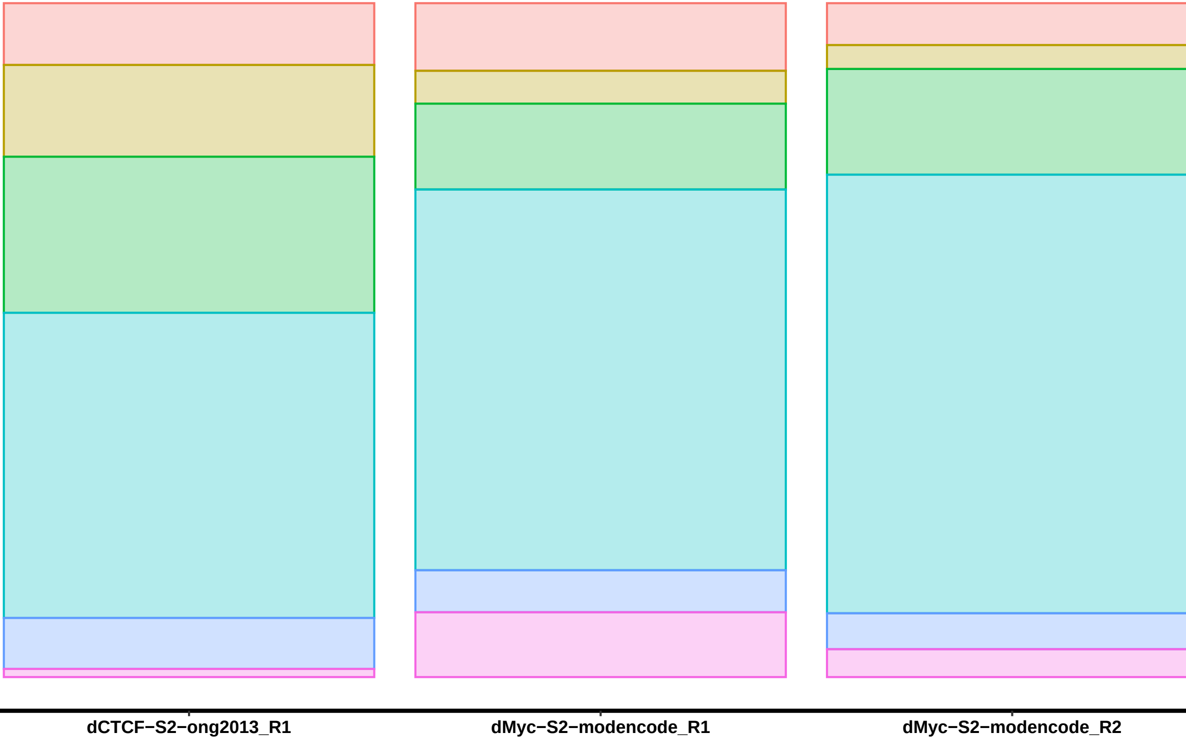
Peak Location Relative to Annotation

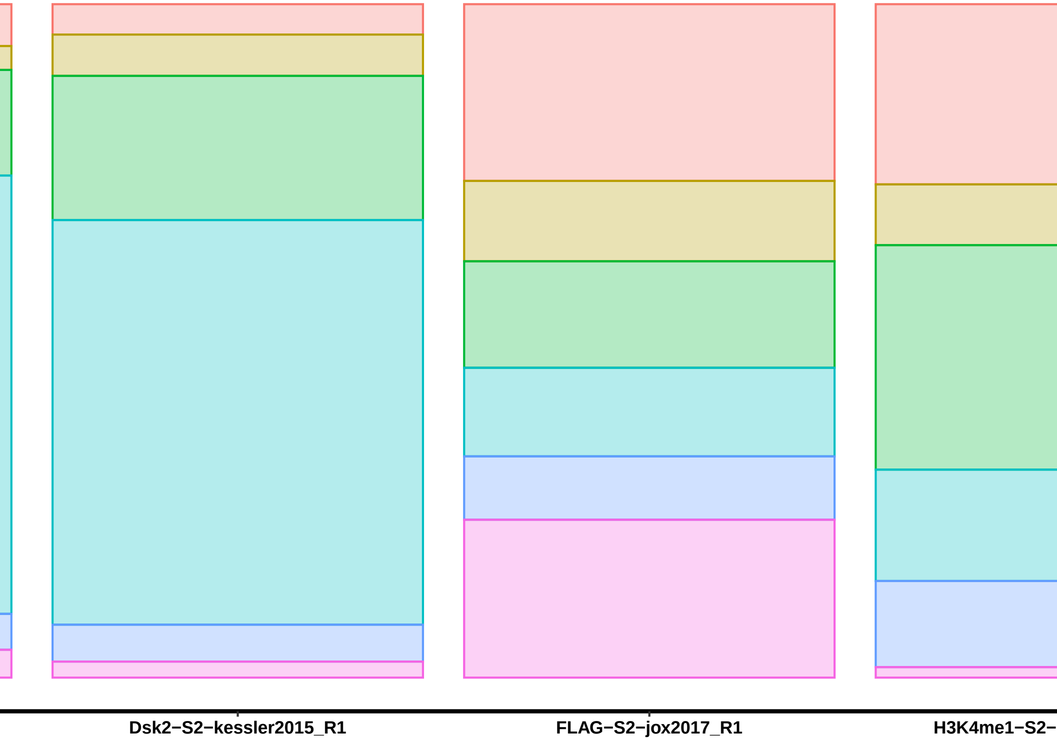


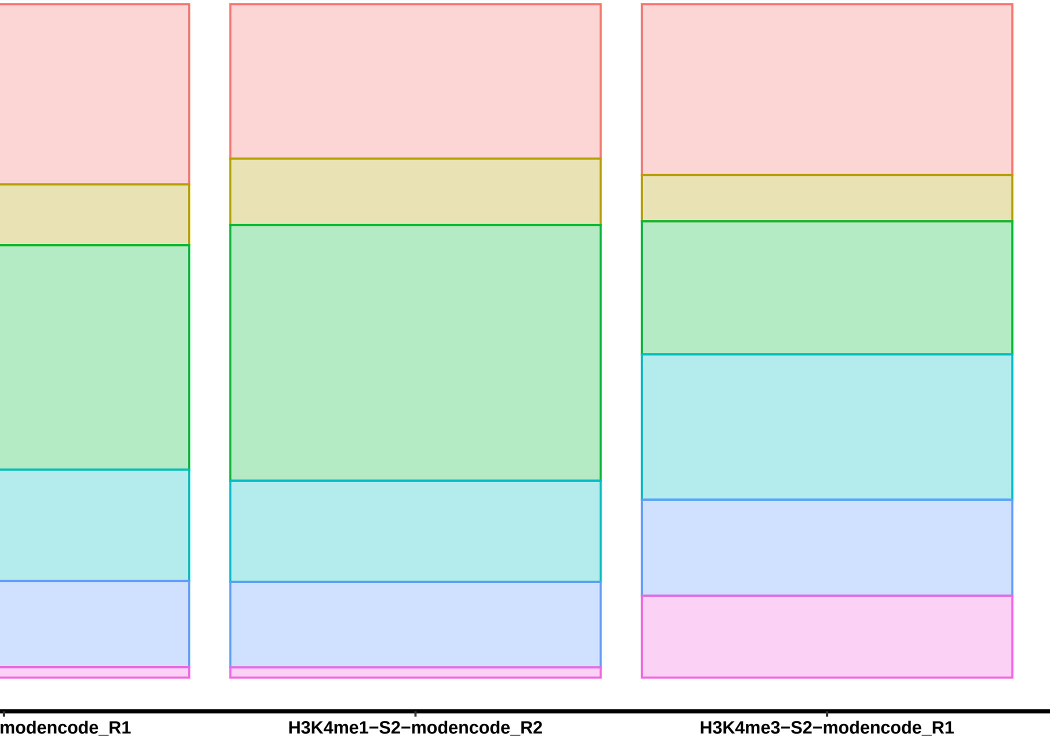


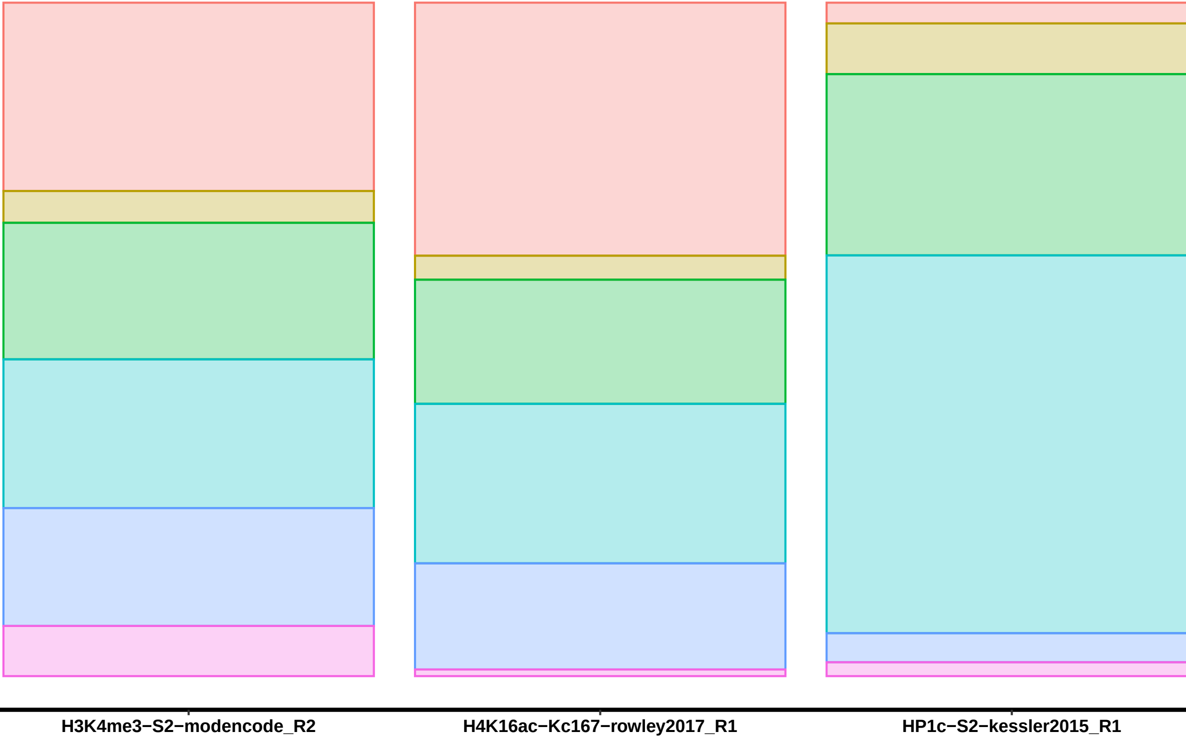


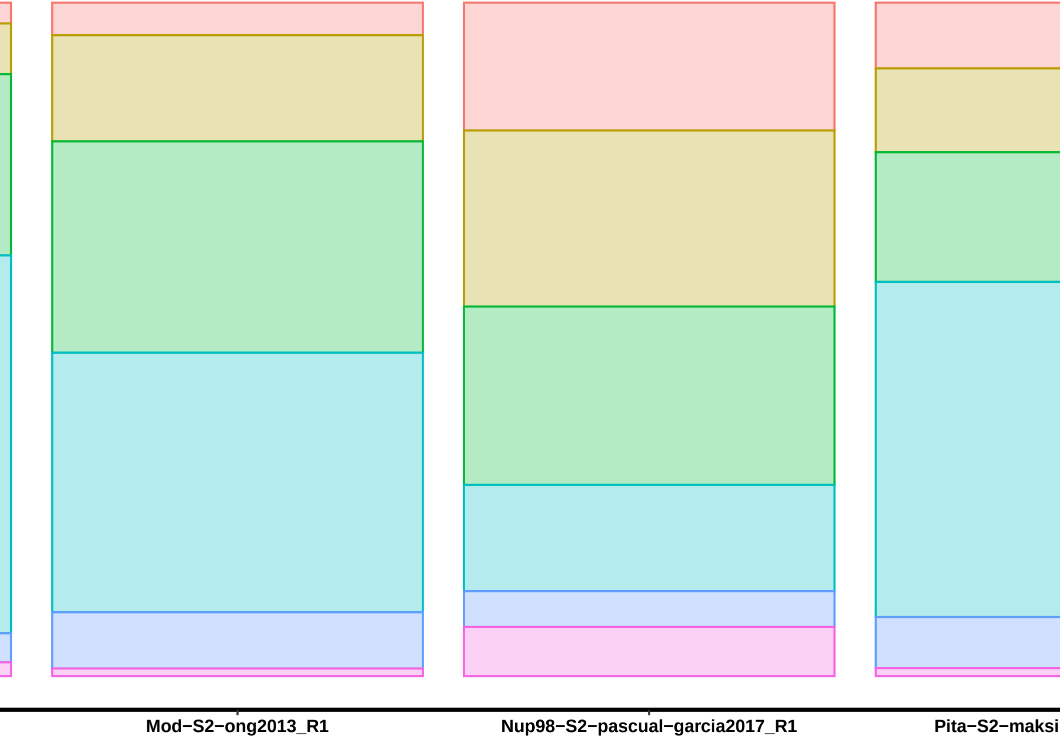


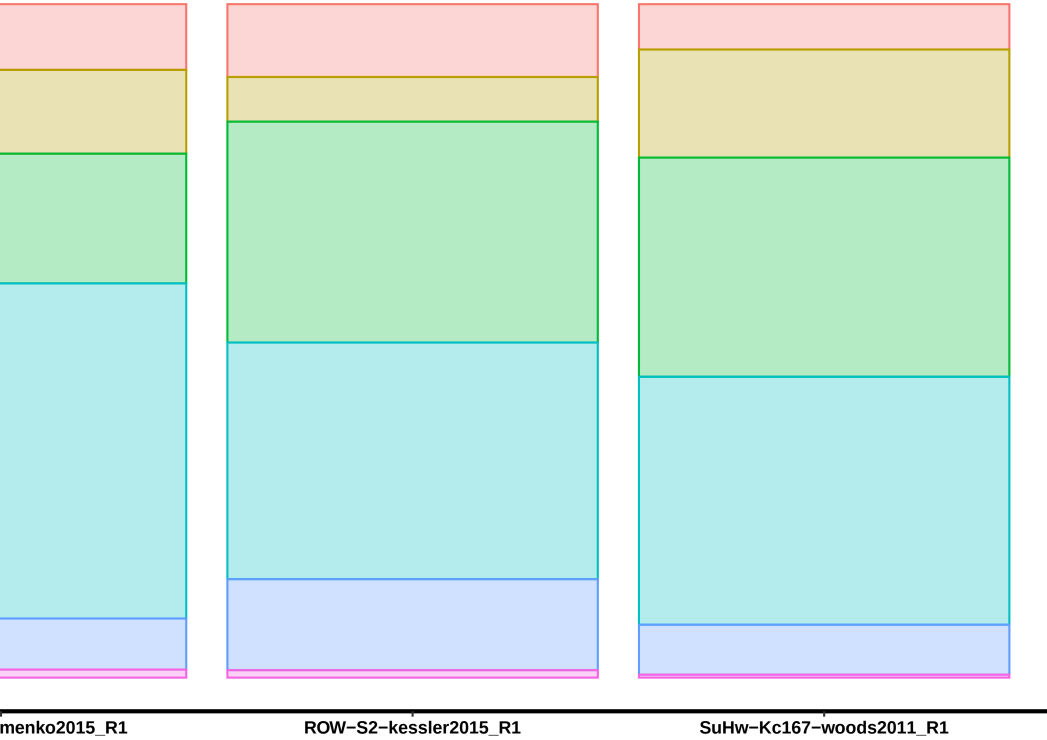


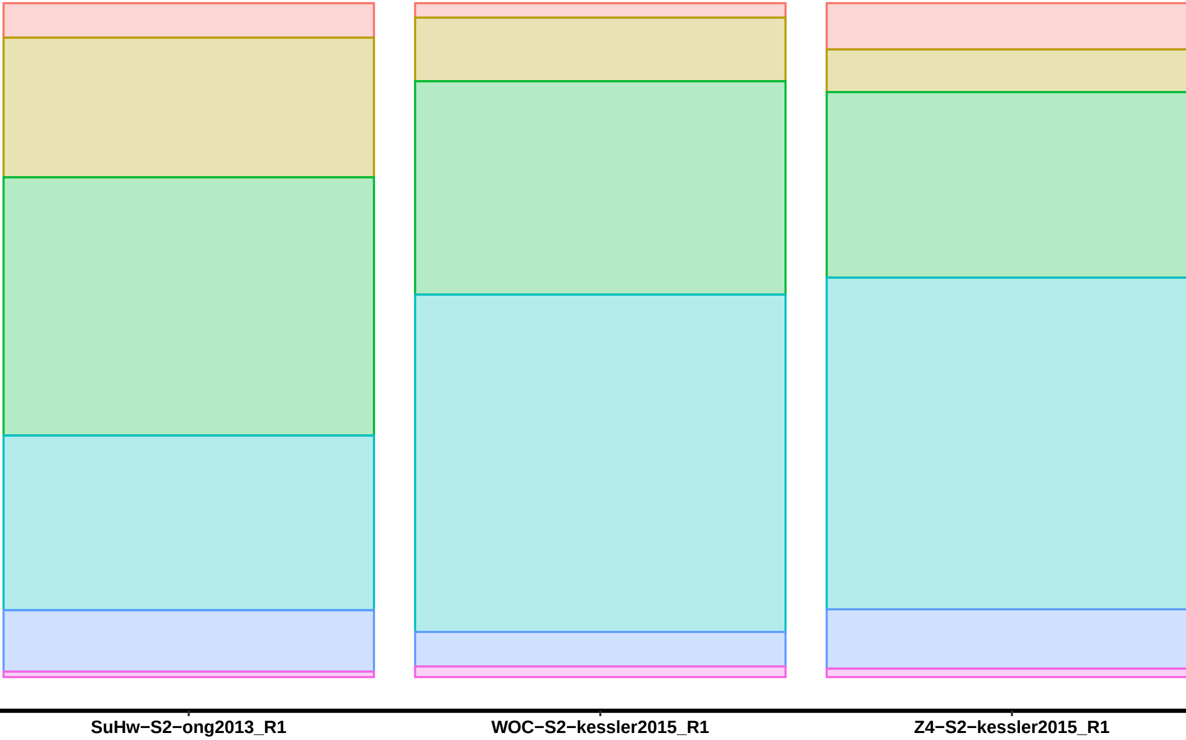


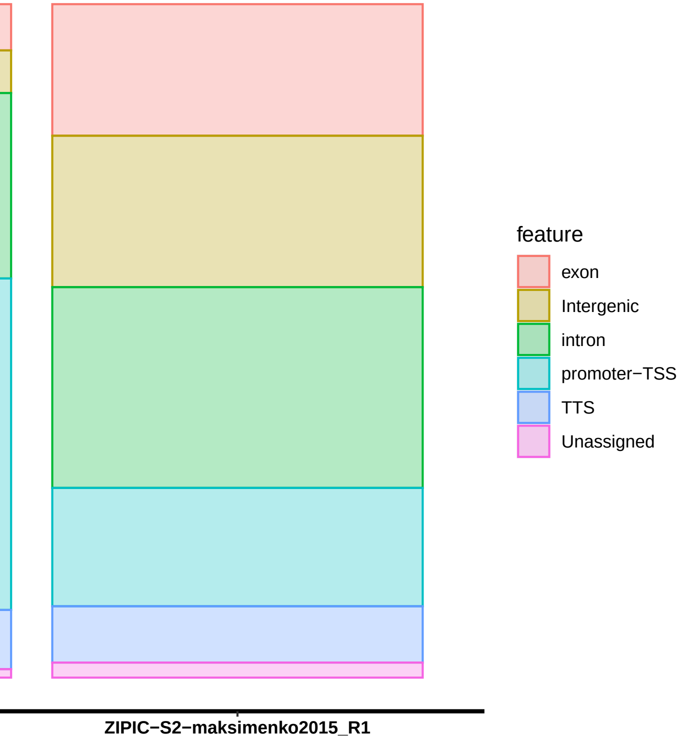


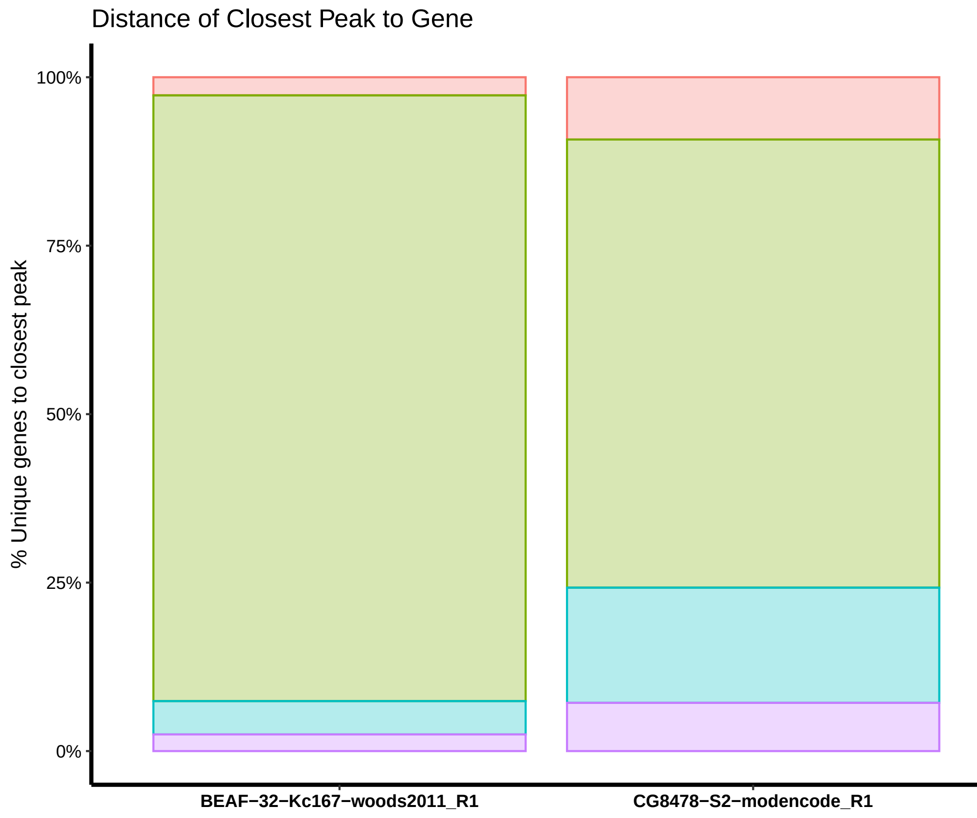


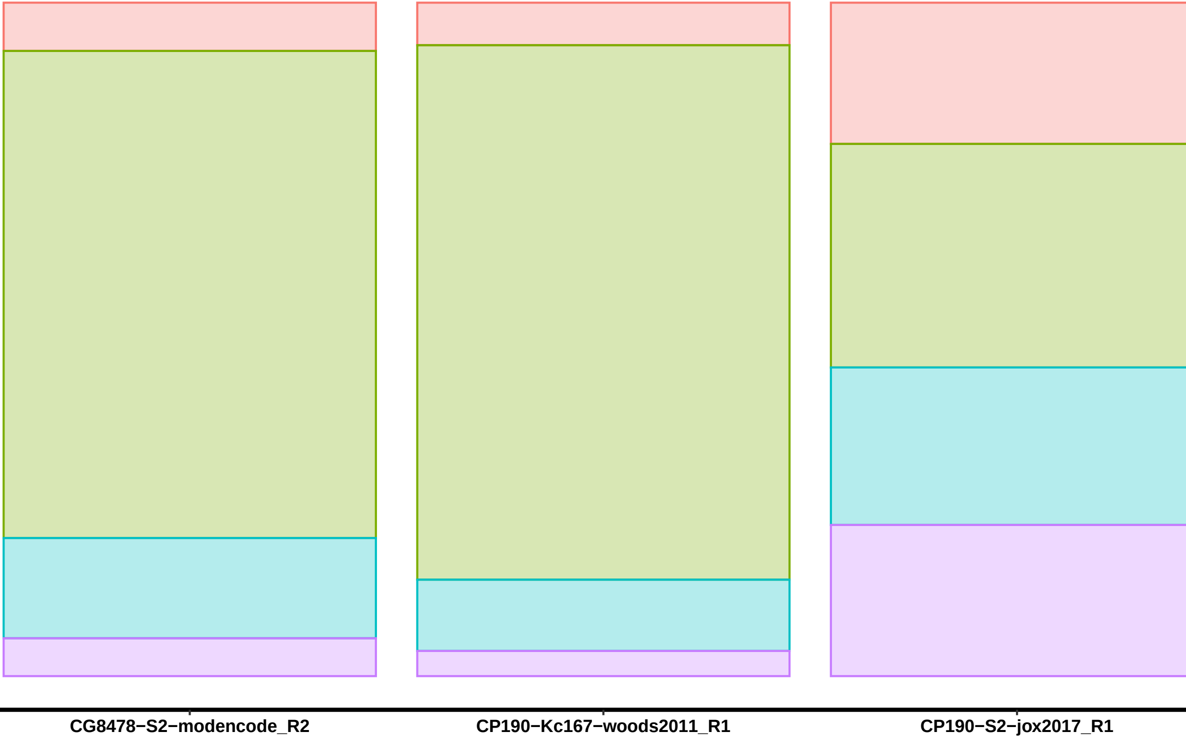


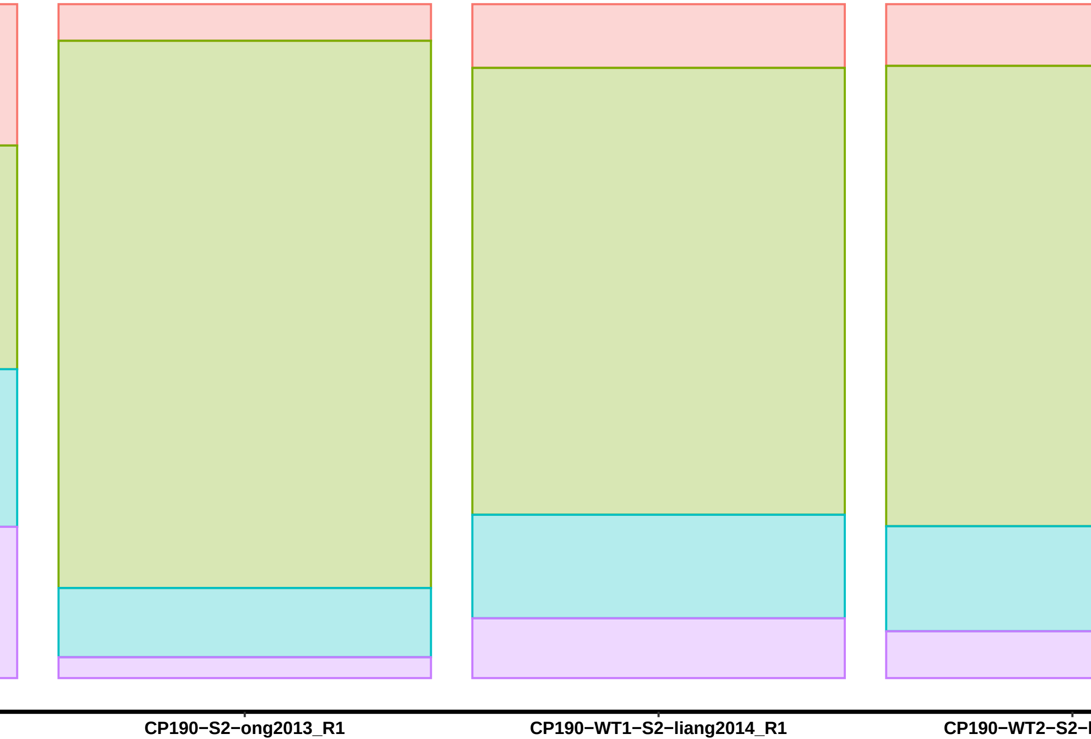


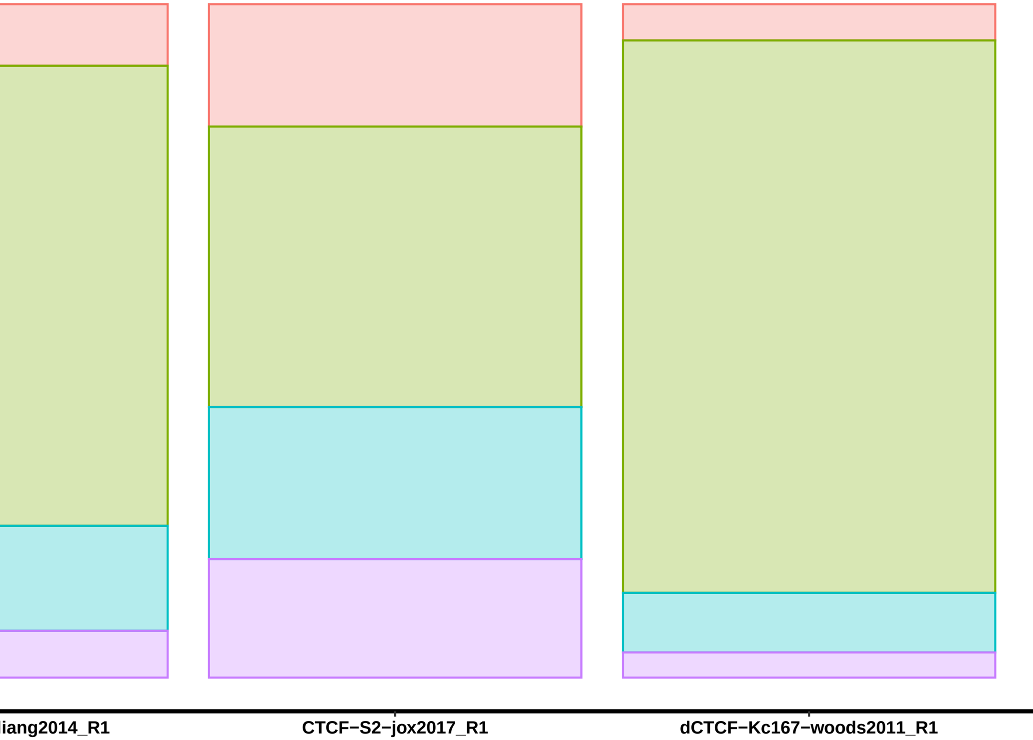


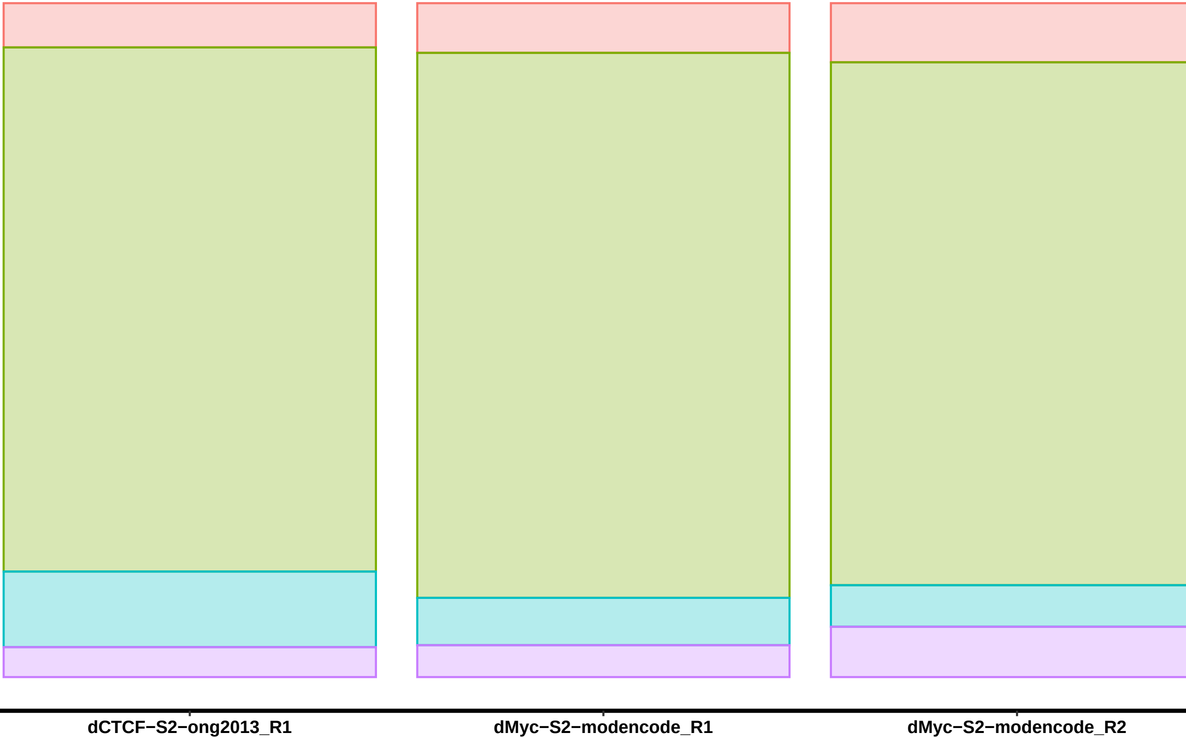



Distance of Closest Peak to Gene
% Unique genes to closest peak
100%
75%
50%
25%
0%
BEAF−32−Kc167−woods2011_R1
CG8478−S2−modencode_R1


CG8478−S2−modencode_R2
CP190−Kc167−woods2011_R1
CP190−S2−jox2017_R1


CP190−S2−ong2013_R1
CP190−WT1−S2−liang2014_R1
CP190−WT2−S2−l


iang2014_R1
CTCF−S2−jox2017_R1
dCTCF−Kc167−woods2011_R1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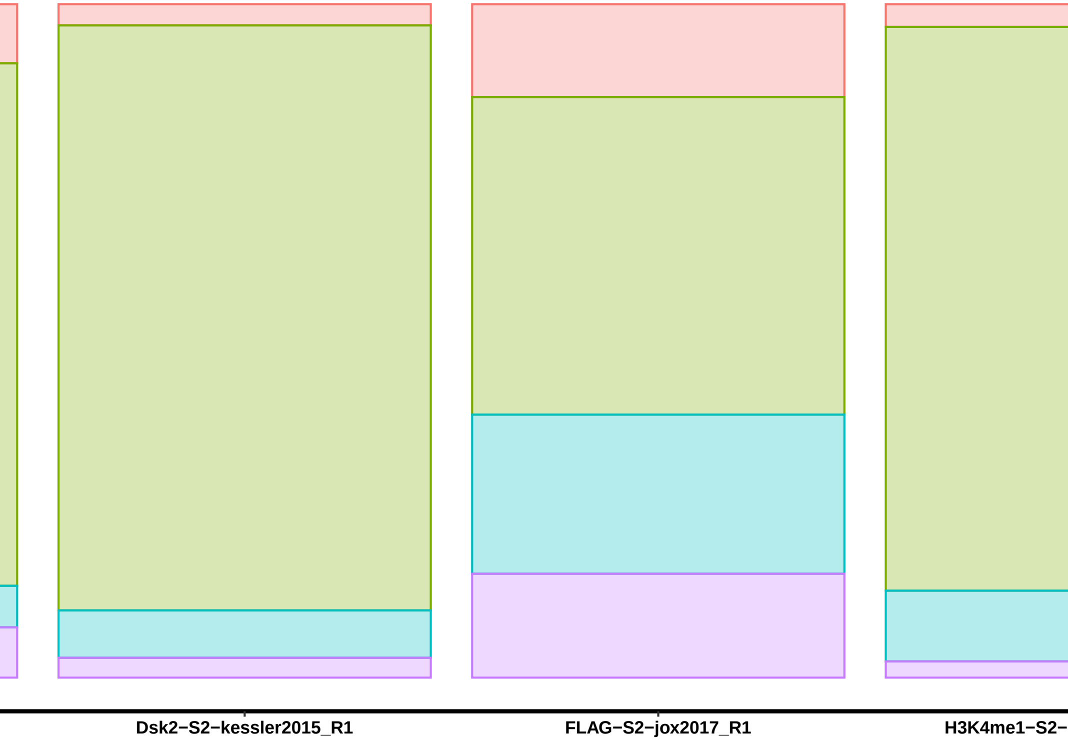
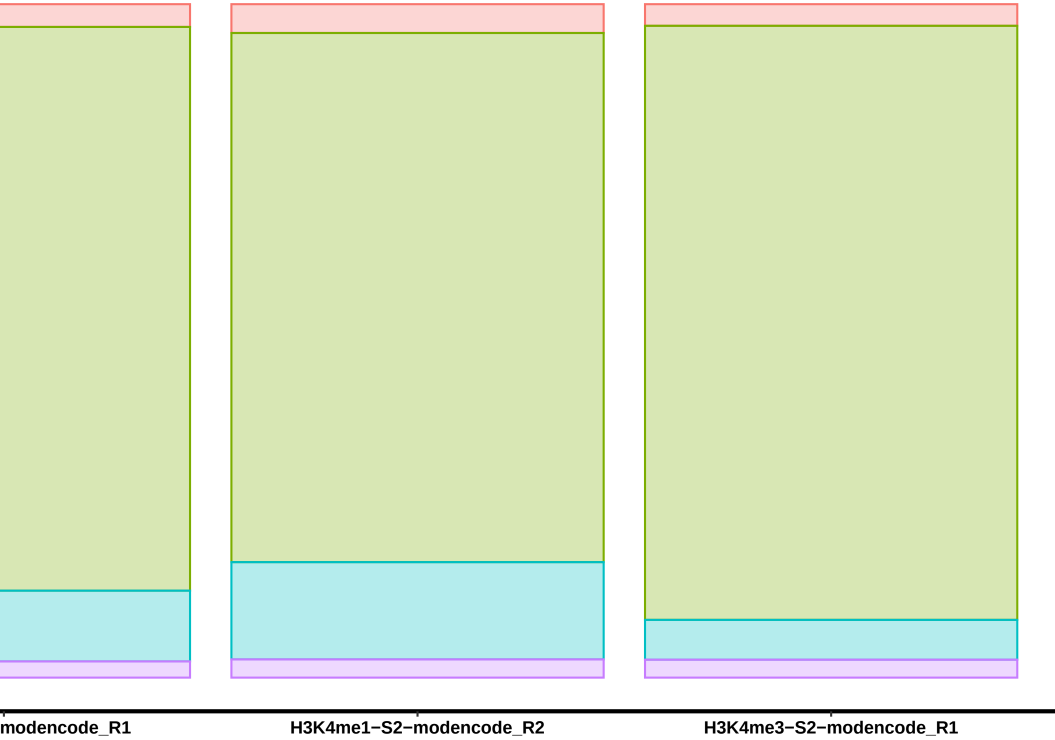
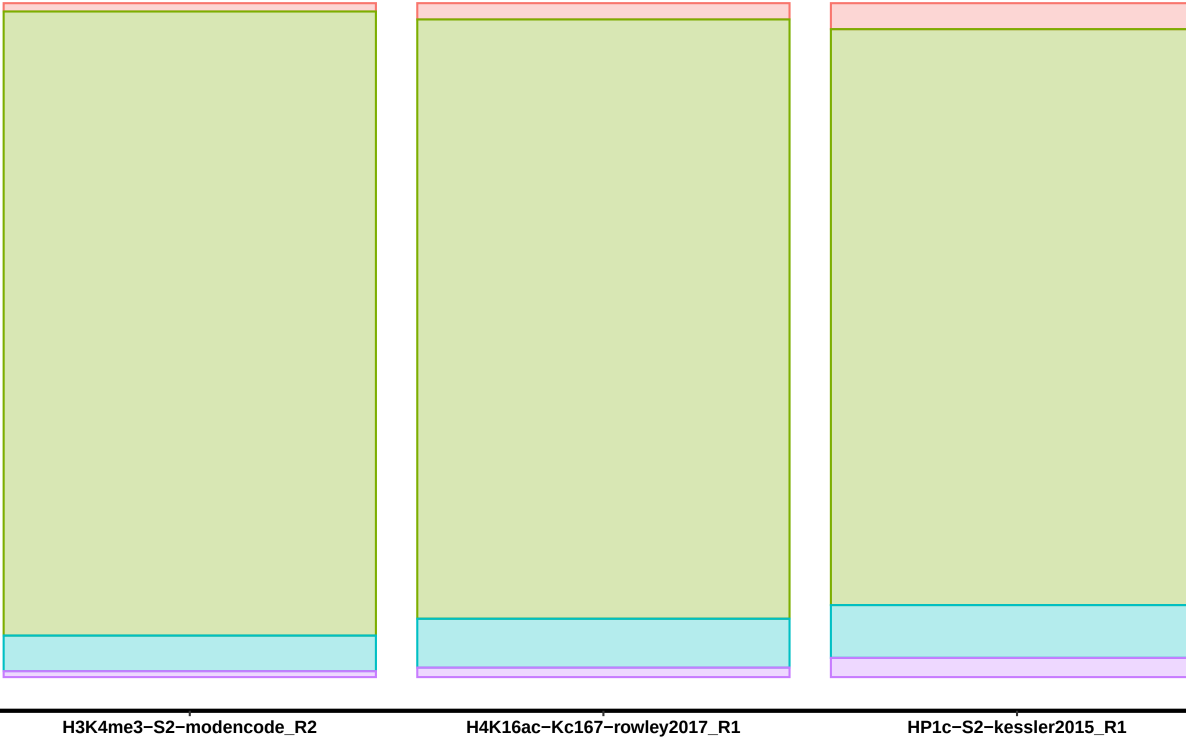
dMyc−S2−modencode_R1
dMyc−S2−modencode_R2

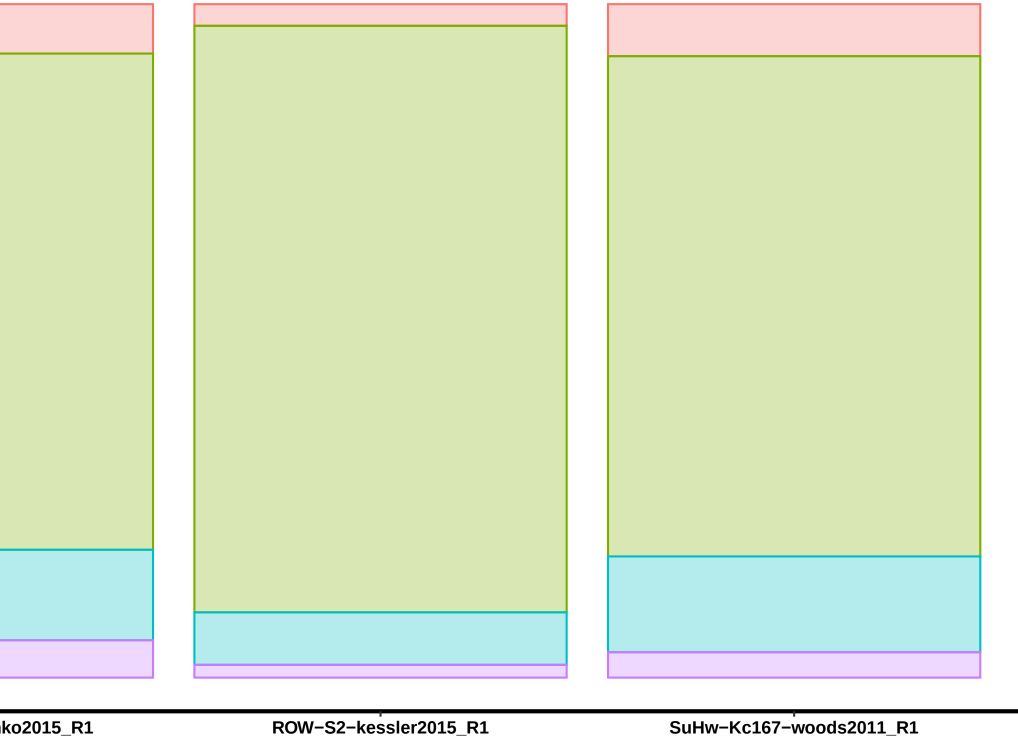

Dsk2−S2−kessler2015_R1
FLAG−S2−jox2017_R1
H3K4me1−S2−

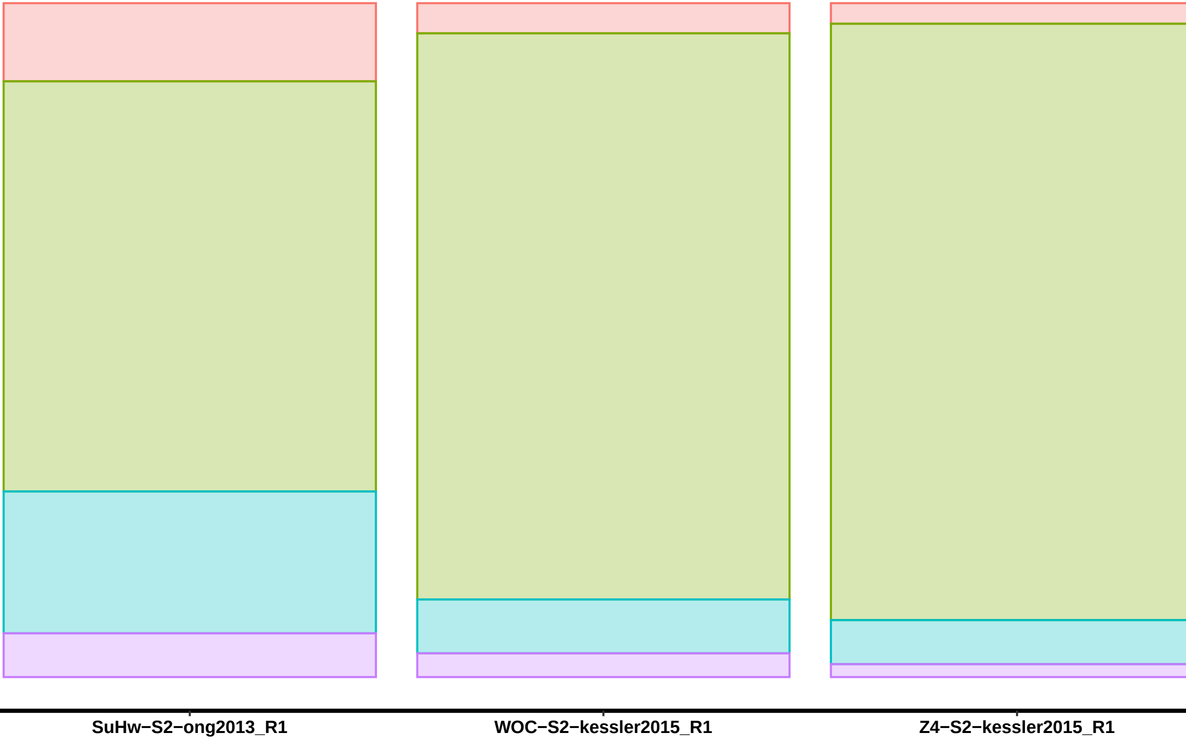


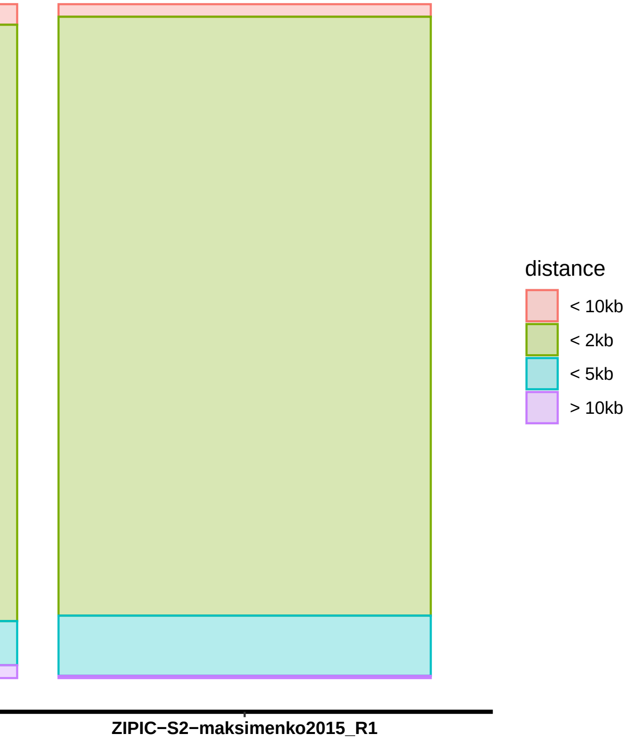
modencode_R1
H3K4me1−S2−modencode_R2
H3K4me3−S2−modencode_R1

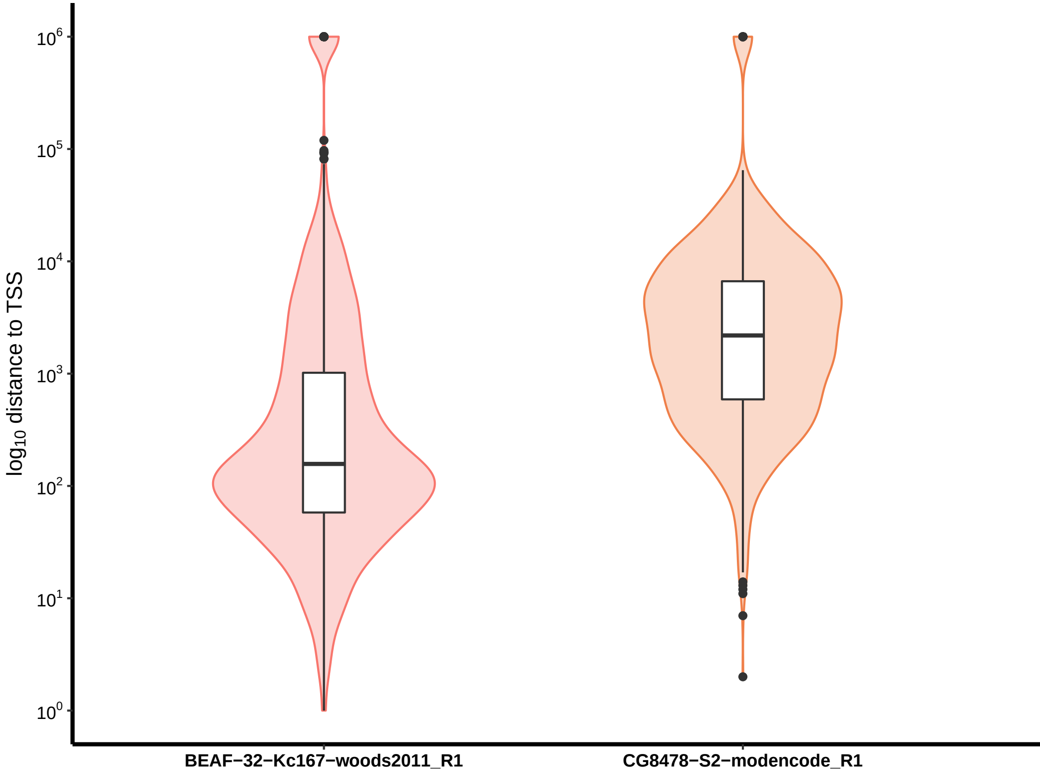
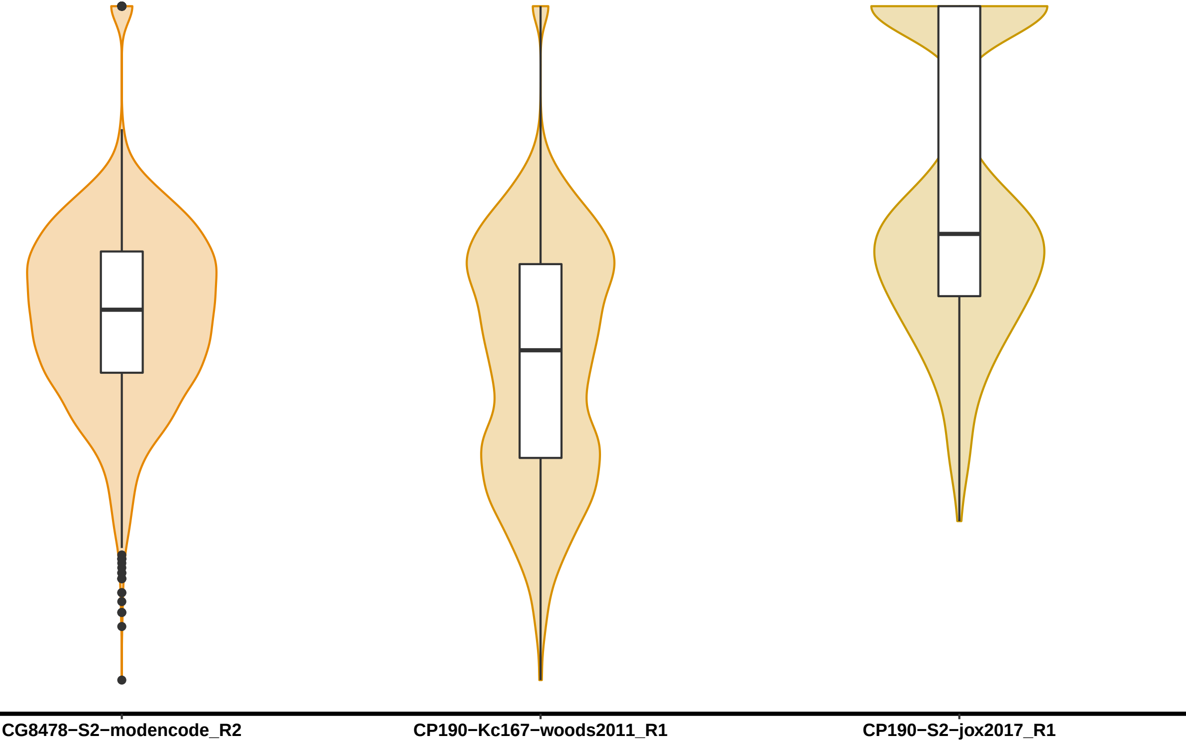
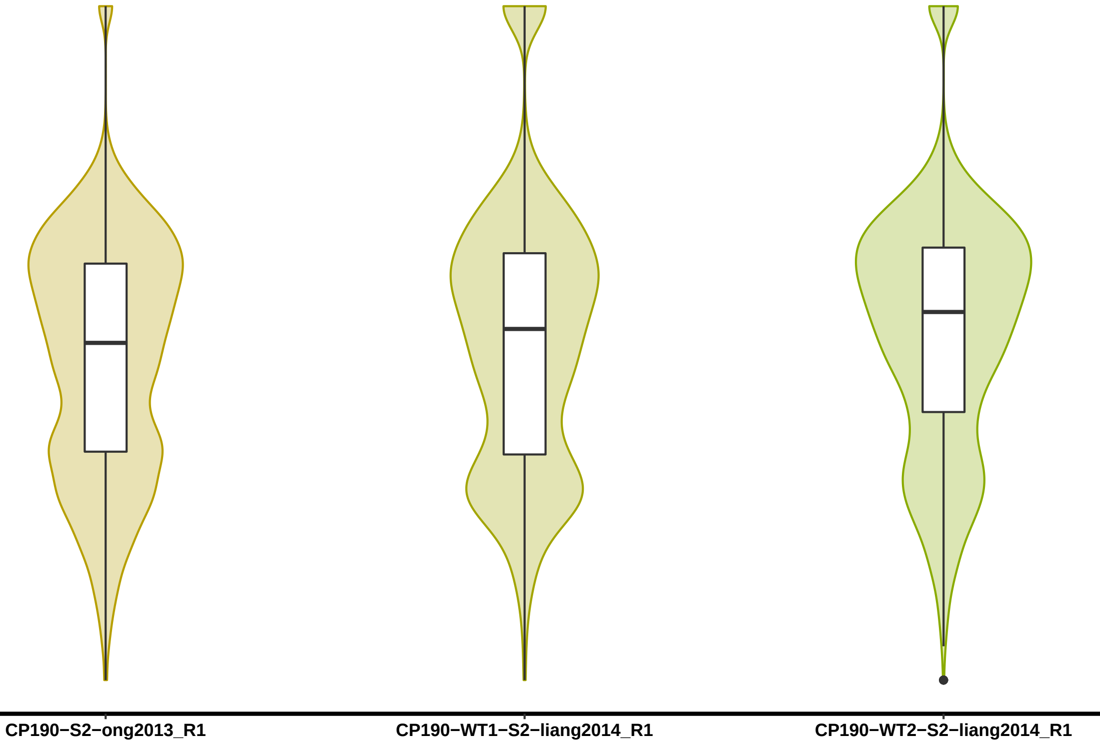
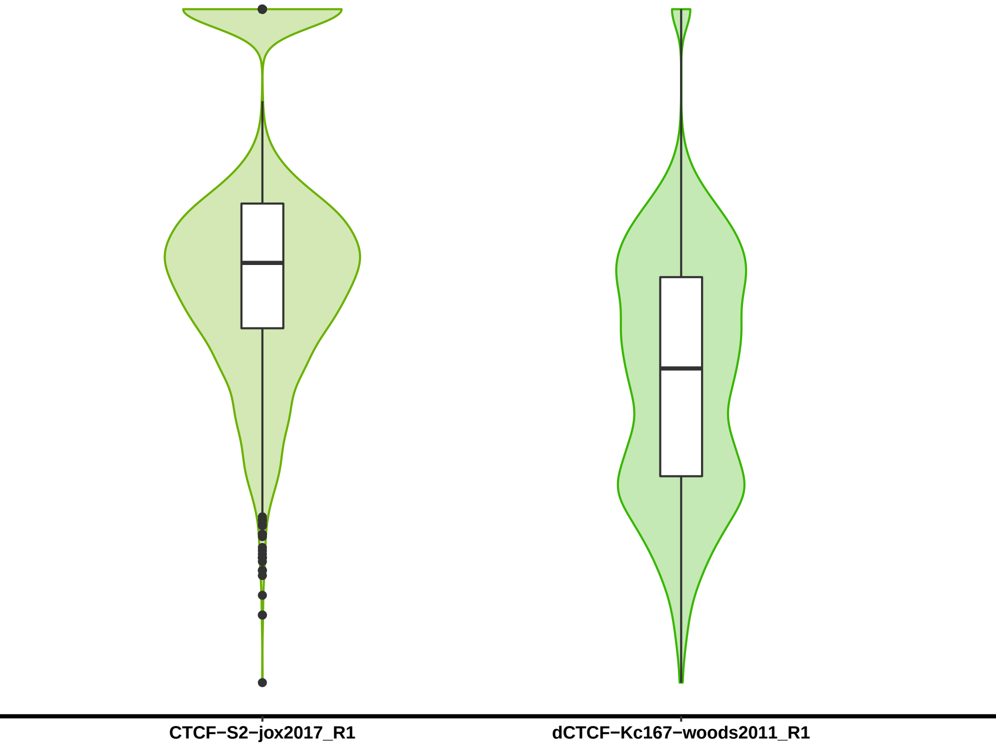
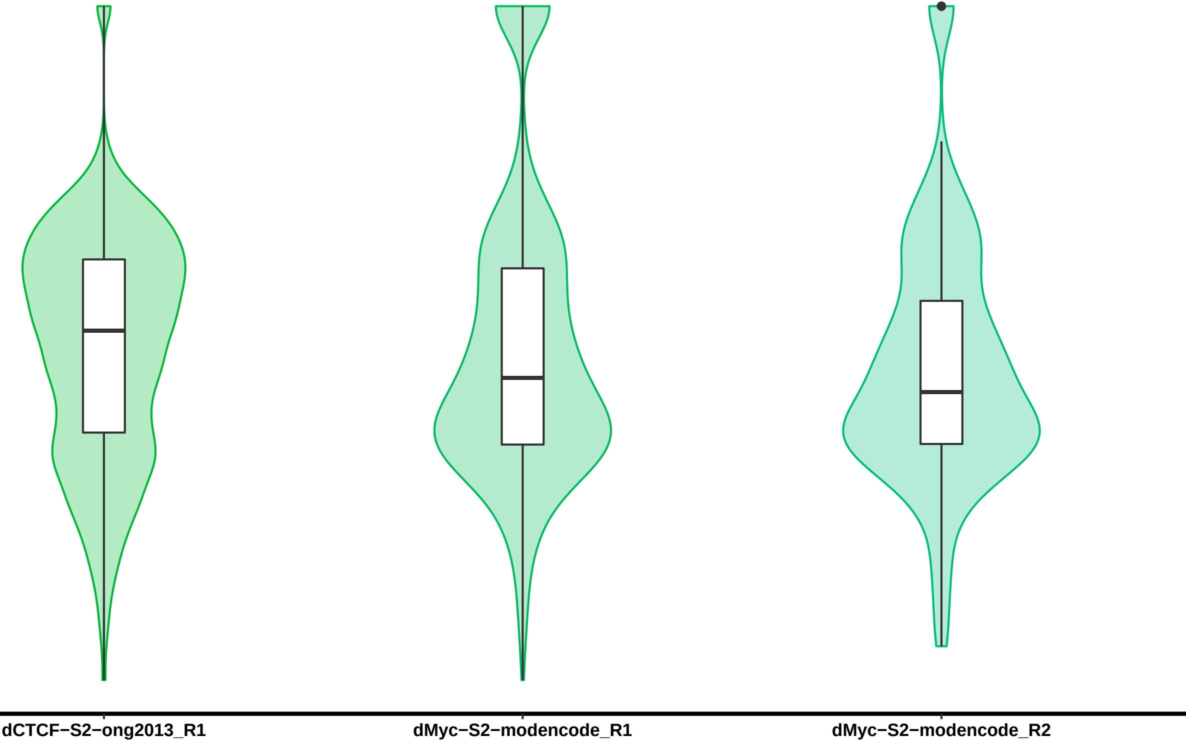
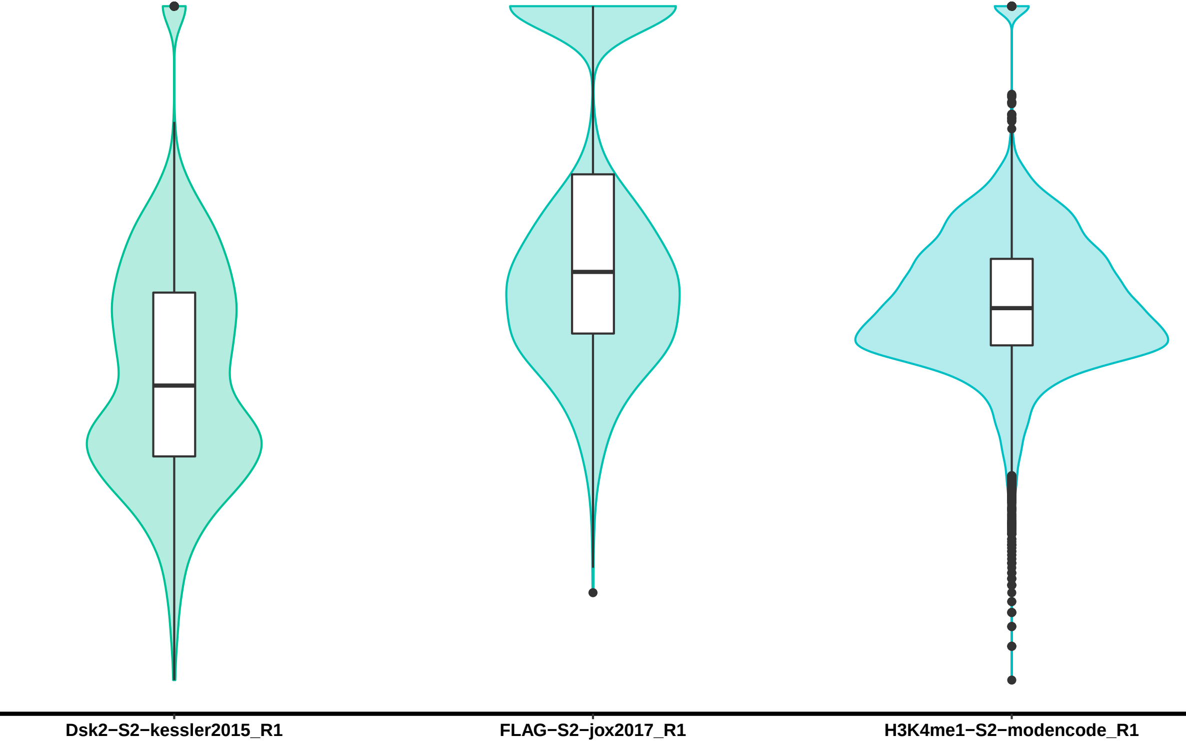
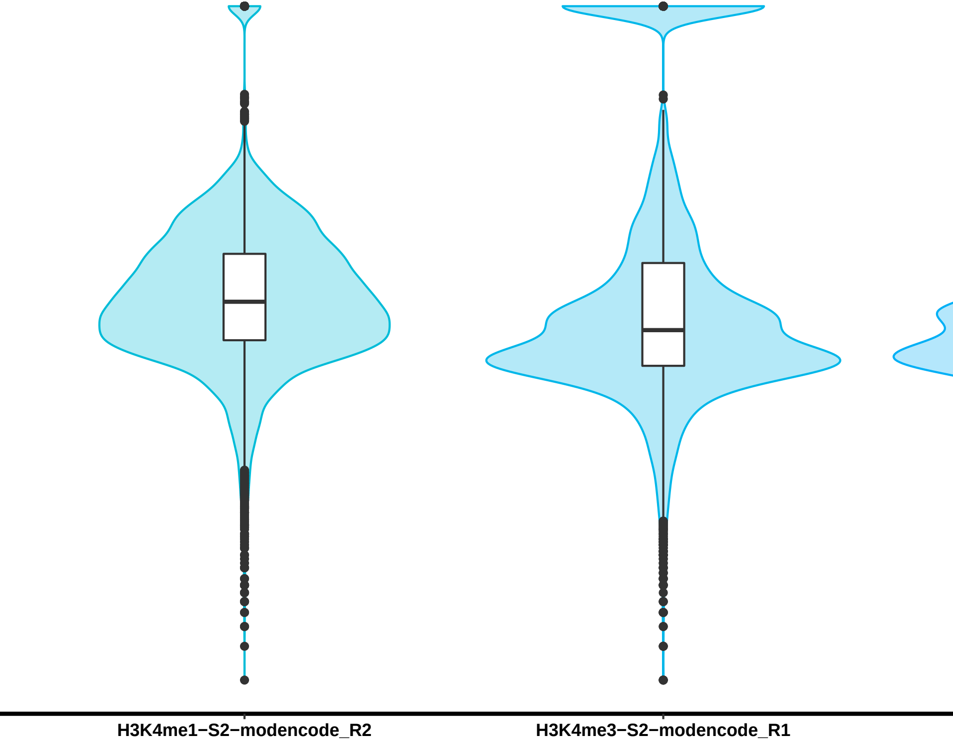
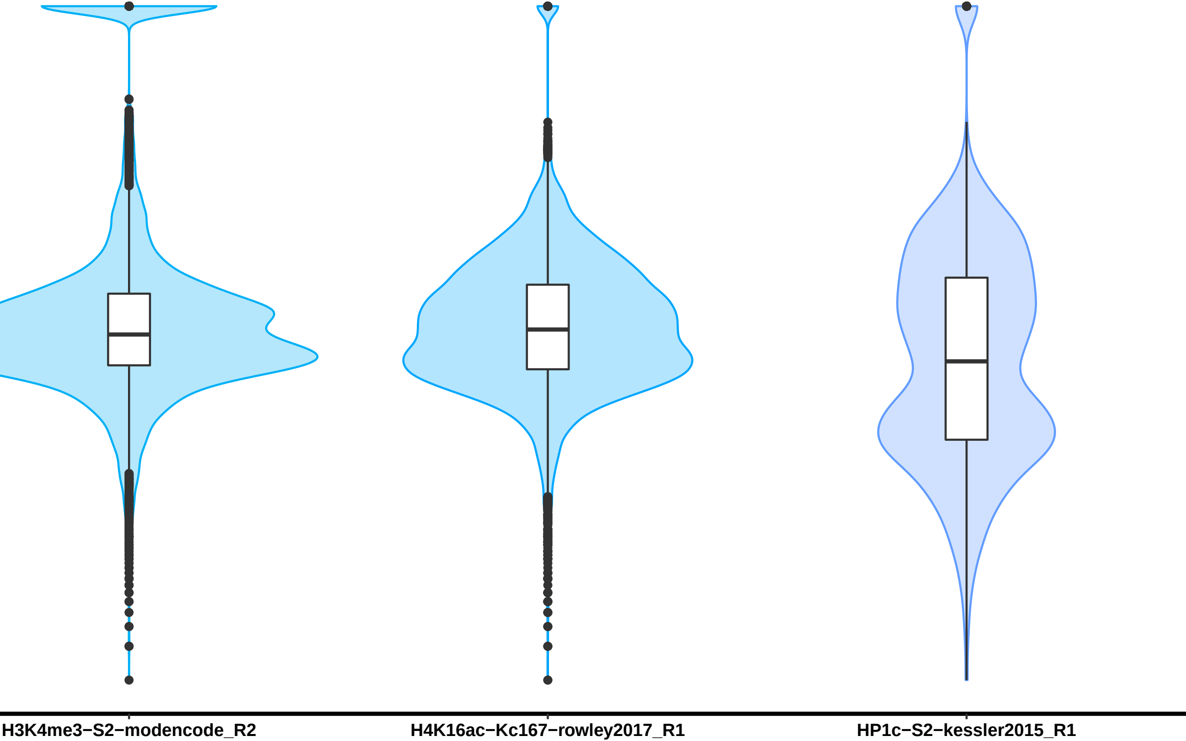
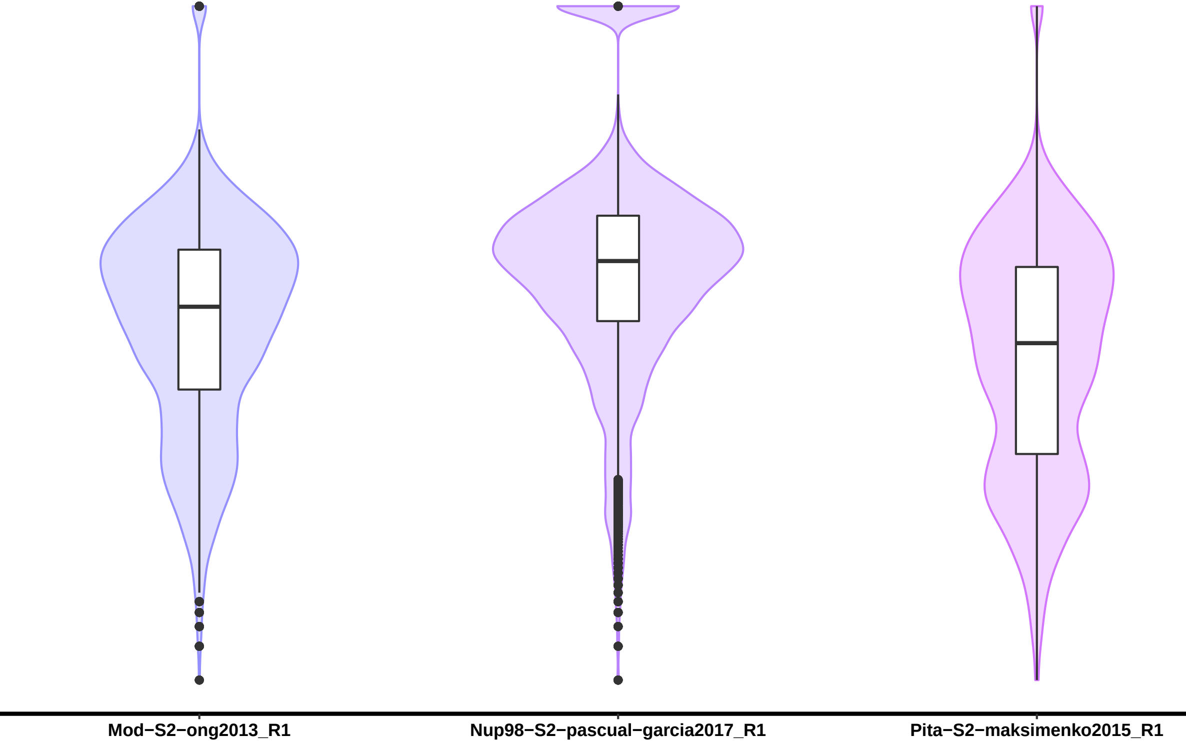
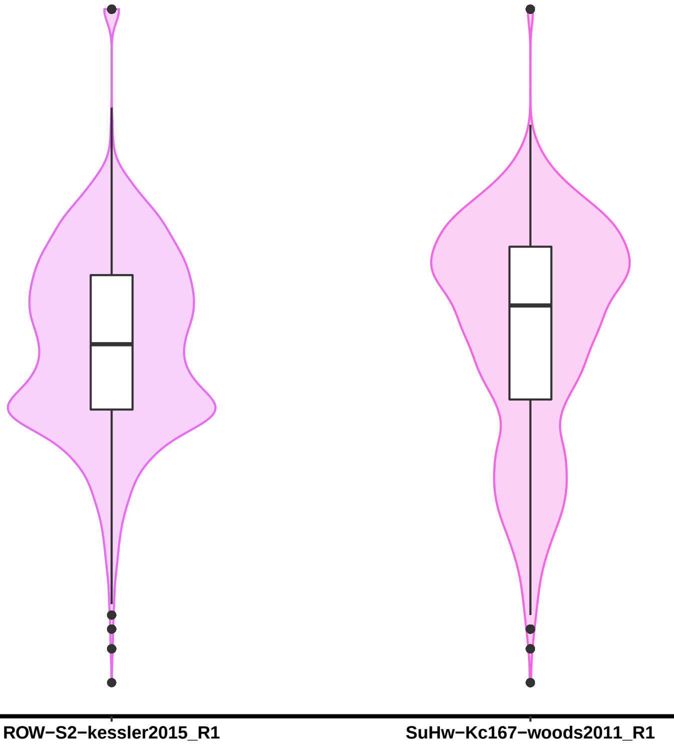
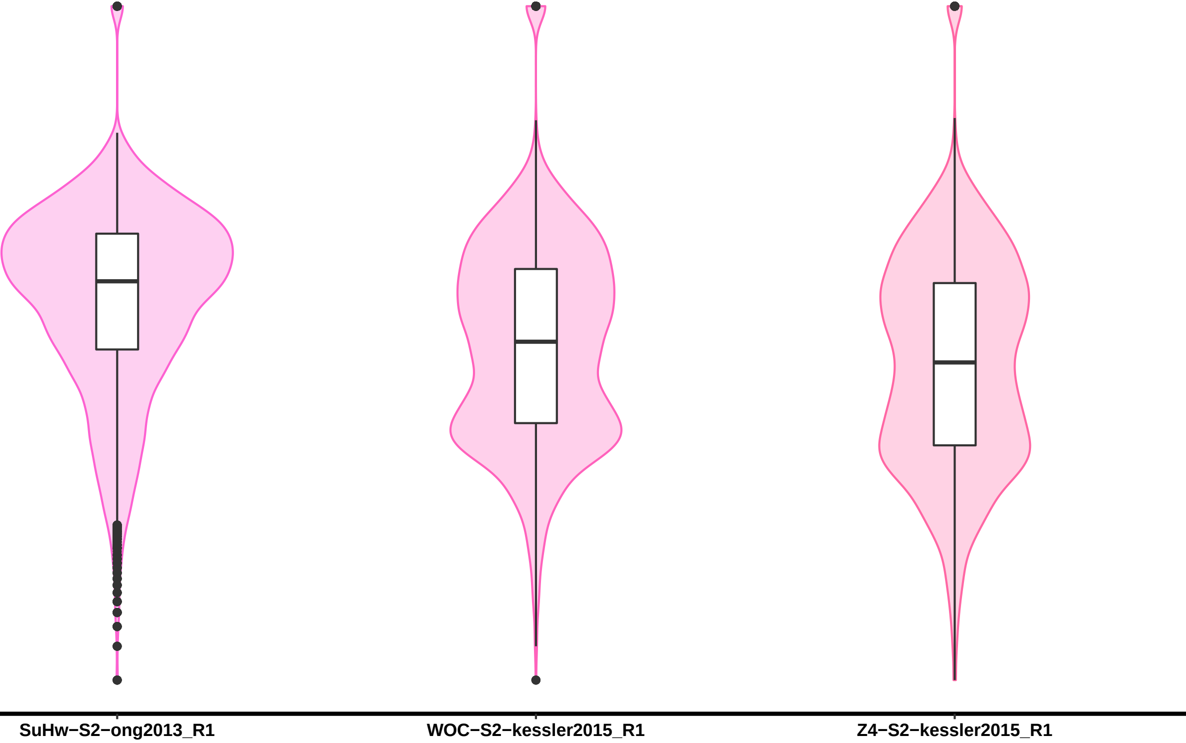
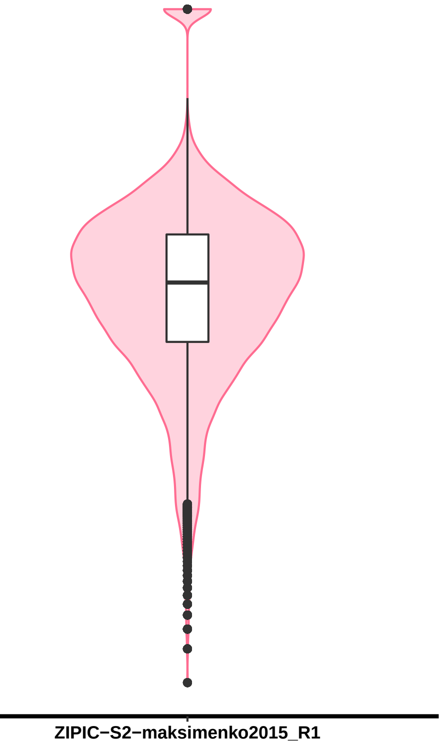

H3K4me3−S2−modencode_R2
H4K16ac−Kc167−rowley2017_R1
HP1c−S2−kessler2015_R1

Mod−S2−ong2013_R1
Nup98−S2−pascual−garcia2017_R1
Pita−S2−maksimen


ko2015_R1
ROW−S2−kessler2015_R1
SuHw−Kc167−woods2011_R1


SuHw−S2−ong2013_R1
WOC−S2−kessler2015_R1
Z4−S2−kessler2015_R1


distance
< 10kb
< 2kb
< 5kb
> 10kb
ZIPIC−S2−maksimenko2015_R1

Peak Distribution Relative to TSS