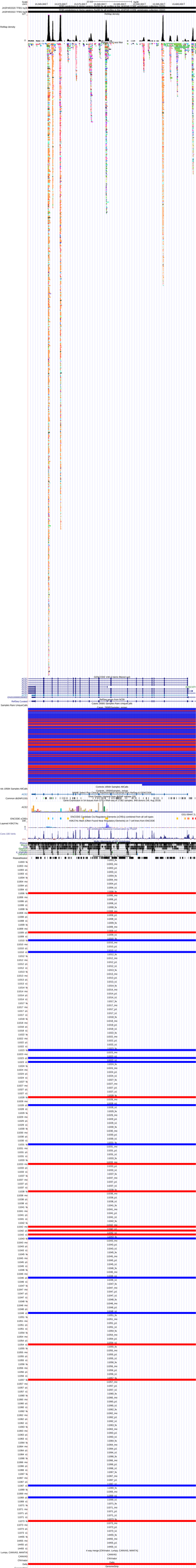
Scale
chr1
15,585,000 f
15,590,000 f
15,595,000 f
15,600,000 f
15,605,000 f
15,610,000 f
15,615,000 f
15,620,000 f
15,625,000 f
15,630,000 f
15,635,000 f
15,640,000 f
15,645,000 f
15,650,000 f
15,655,000 f
15,660,000 f
JASPAR2020 TFBS
JASPAR2022 TFBS
hg38
Reflog density
Reflog density
Gene Map
Reflog density
ENCODE cell 2 items filtered out
ENCODE DNase Clusters
Reflog Curated
Reflog Curated from NCBI
Samples Rank UniqueCalls
Cases 1000K Samples Rank UniqueCalls
Cases 1000K Samples Rank
Cores 1000K Samples AlCalls
Cores 1000K Samples Rank
Cores 1000K Samples Rank Reflog & ENCODE
Gene Exon/Intron on 54 tissues from GTEx v7 RNA-seq of 17382 samples, 948 donors (UE, Aug 2019)
Common dbSNP(151)
ENCODE cCREs
ENCODE Candidate Cis-Regulatory Elements (cCREs) combined from all cell types
HiCUTs Well-Ziffen Fourie Neve Regulatory Elements on 7 cell lines from ENCODE
Layered H3K27ac
GSLSMAT 2
Cores 100 Vols
Phastcons100way
Phastcons100way
Phastcons100way