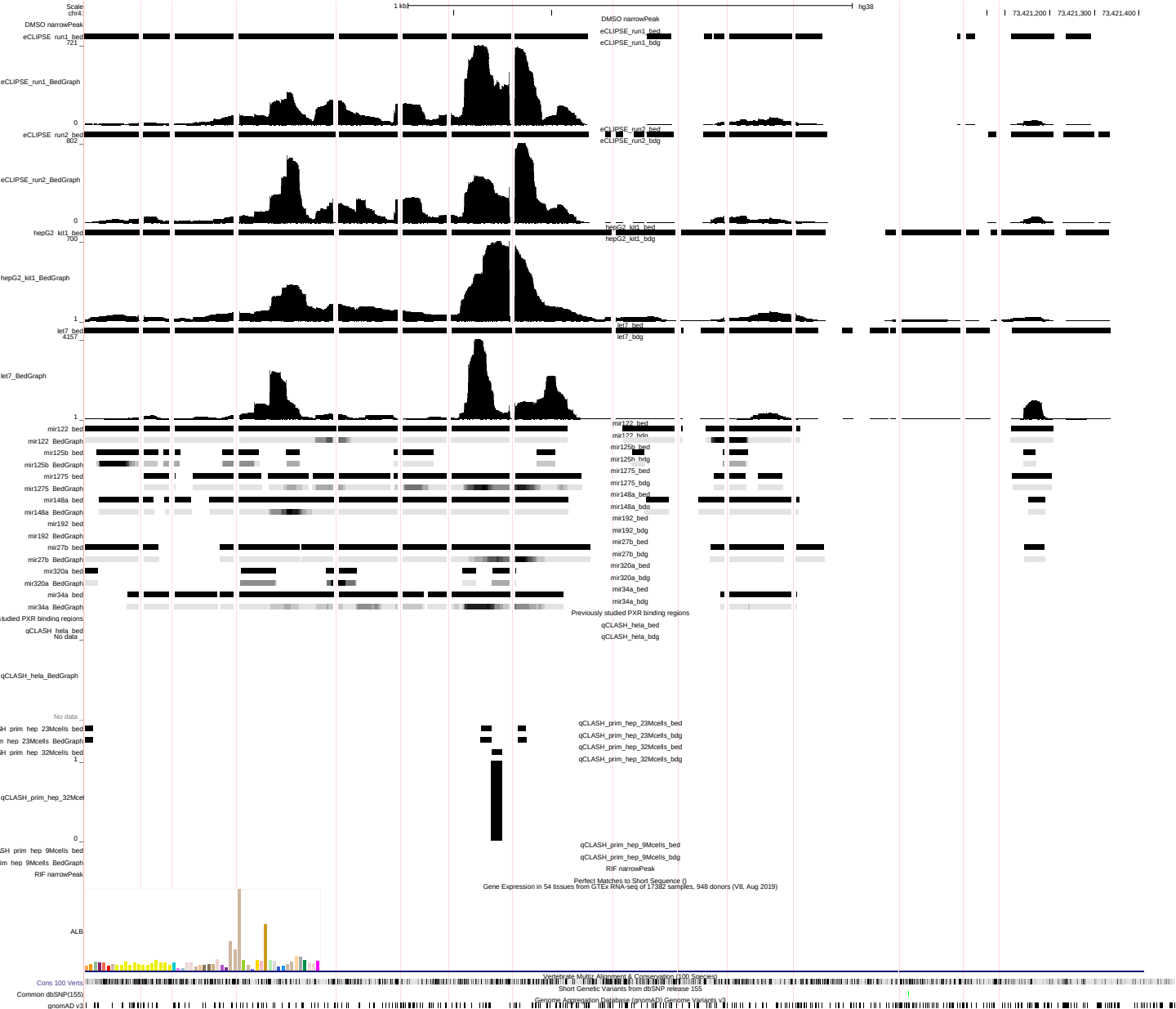
Scale
chr4:
1 kb
hg38
73,421,200
73,421,300
73,421,400
DMSO narrowPeak
eCLIPSE_run1_bed
eCLIPSE_run1_bdg
eCLIPSE_run1_BedGraph
eCLIPSE_run2_bed
eCLIPSE_run2_bdg
eCLIPSE_run2_BedGraph
hepG2_klf1_bed
hepG2_klf1_bdg
hepG2_klf1_BedGraph
let7_bed
let7_bdg
let7_BedGraph
mir122_bed
mir122_BedGraph
mir125b_bed
mir125b_BedGraph
mir1275_bed
mir1275_BedGraph
mir148a_bed
mir148a_BedGraph
mir192_bed
mir192_BedGraph
mir27b_bed
mir27b_BedGraph
mir320a_bed
mir320a_BedGraph
mir34a_bed
mir34a_BedGraph
Previously studied PXR binding regions
qCLASH_hela_bed
qCLASH_hela_bdg
qCLASH_hela_BedGraph
qCLASH_prim_hep_23Mcells_bed
qCLASH_prim_hep_23Mcells_bdg
qCLASH_prim_hep_32Mcells_bed
qCLASH_prim_hep_32Mcells_bdg
qCLASH_prim_hep_9Mcells_bed
qCLASH_prim_hep_9Mcells_bdg
RIF narrowPeak
Perfect Matches to Short Sequence ()
Gene Expression in 54 tissues from GTEx RNA-seq of 17382 samples, 948 donors (V8, Aug 2019)
ALB
Vertebrate Multiz Alignment & Conservation (100 Species)
Cons 100 Verts
Common dbSNP(155)
Short Genetic Variants from dbSNP release 155
gnomAD v3
Genome Aggregation Database (gnomAD) Genome Variants v3