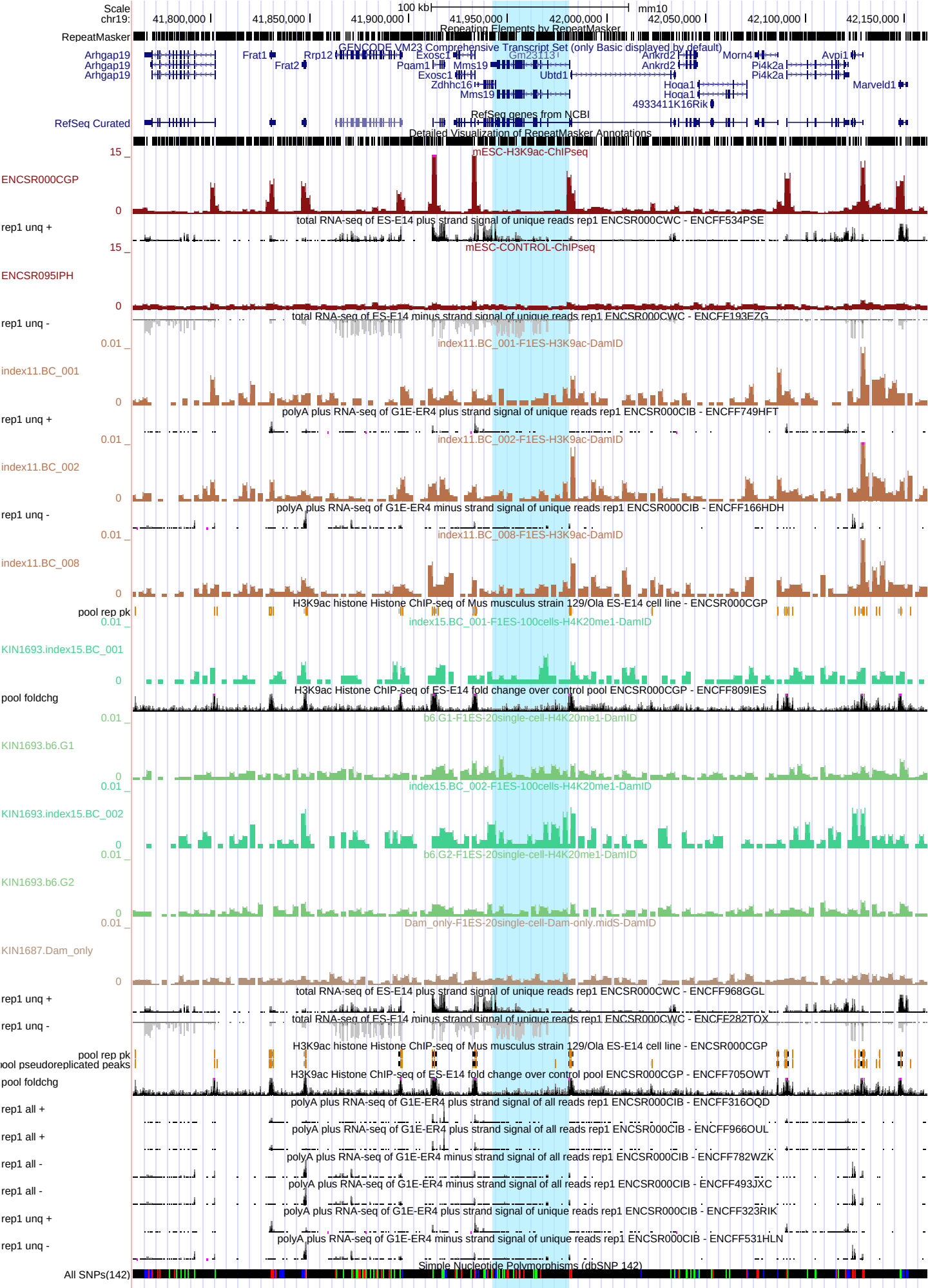
Scale
chr19:
100 kb
mm10
41,800,000
41,850,000
41,900,000
41,950,000
42,000,000
42,050,000
42,100,000
42,150,000
RepeatMasker
Repeating Elements by RepeatMasker
GENCODE VM23 Comprehensive Transcript Set (only Basic displayed by default)
Arhgap19
Arhgap19
Arhgap19
Frat1
Frat2
Rrp12
Exosc1
Pqam1
Mms19
Gm23113
Exosc1
Zdhhc16
Mms19
Ubtd1
Ankrd2
Ankrd2
Morn4
Hoga1
Hoga1
4933411K16Rik
Pi4k2a
Pi4k2a
Avpi1
Marveld1
RefSeq Curated
RefSeq genes from NCBI
Detailed Visualization of RepeatMasker Annotations
mESC-H3K9ac-ChIPseq
ENCSR000CGP
15
0
rep1 unq +
total RNA-seq of ES-E14 plus strand signal of unique reads rep1 ENCSR000CWC - ENCFF534PSE
mESC-CONTROL-ChIPseq
ENCSR095IPH
rep1 unq -
total RNA-seq of ES-E14 minus strand signal of unique reads rep1 ENCSR000CWC - ENCFF193EZG
0.01
index11.BC_001-F1ES-H3K9ac-DamID
index11.BC_001
polyA plus RNA-seq of G1E-ER4 plus strand signal of unique reads rep1 ENCSR000CIB - ENCFF749HFT
index11.BC_002-F1ES-H3K9ac-DamID
index11.BC_002
polyA plus RNA-seq of G1E-ER4 minus strand signal of unique reads rep1 ENCSR000CIB - ENCFF166HDH
index11.BC_008-F1ES-H3K9ac-DamID
index11.BC_008
H3K9ac histone Histone ChIP-seq of Mus musculus strain 129/Ola ES-E14 cell line - ENCSR000CGP
pool rep pk
index15.BC_001-F1ES-100cells-H4K20me1-DamID
KIN1693.index15.BC_001
H3K9ac Histone ChIP-seq of ES-E14 fold change over control pool ENCSR000CGP - ENCFF809IES
pool foldchg
b6.G1-F1ES-20single-cell-H4K20me1-DamID
KIN1693.b6.G1
index15.BC_002-F1ES-100cells-H4K20me1-DamID
KIN1693.index15.BC_002
b6.G2-F1ES-20single-cell-H4K20me1-DamID
KIN1693.b6.G2
Dam_only-F1ES-20single-cell-Dam-only.midS-DamID
KIN1687.Dam_only
total RNA-seq of ES-E14 plus strand signal of unique reads rep1 ENCSR000CWC - ENCFF968GGL
total RNA-seq of ES-E14 minus strand signal of unique reads rep1 ENCSR000CWC - ENCFF282TOX
H3K9ac histone Histone ChIP-seq of Mus musculus strain 129/Ola ES-E14 cell line - ENCSR000CGP
pool pseudoreplicated peaks
H3K9ac Histone ChIP-seq of ES-E14 fold change over control pool ENCSR000CGP - ENCFF705OWT
rep1 all +
polyA plus RNA-seq of G1E-ER4 plus strand signal of all reads rep1 ENCSR000CIB - ENCFF316OQD
polyA plus RNA-seq of G1E-ER4 plus strand signal of all reads rep1 ENCSR000CIB - ENCFF966OUL
rep1 all -
polyA plus RNA-seq of G1E-ER4 minus strand signal of all reads rep1 ENCSR000CIB - ENCFF782WZK
polyA plus RNA-seq of G1E-ER4 minus strand signal of all reads rep1 ENCSR000CIB - ENCFF493JXC
polyA plus RNA-seq of G1E-ER4 plus strand signal of unique reads rep1 ENCSR000CIB - ENCFF323RIK
polyA plus RNA-seq of G1E-ER4 minus strand signal of unique reads rep1 ENCSR000CIB - ENCFF531HLN
Simple Nucleotide Polymorphisms (dbSNP 142)
All SNPs(142)