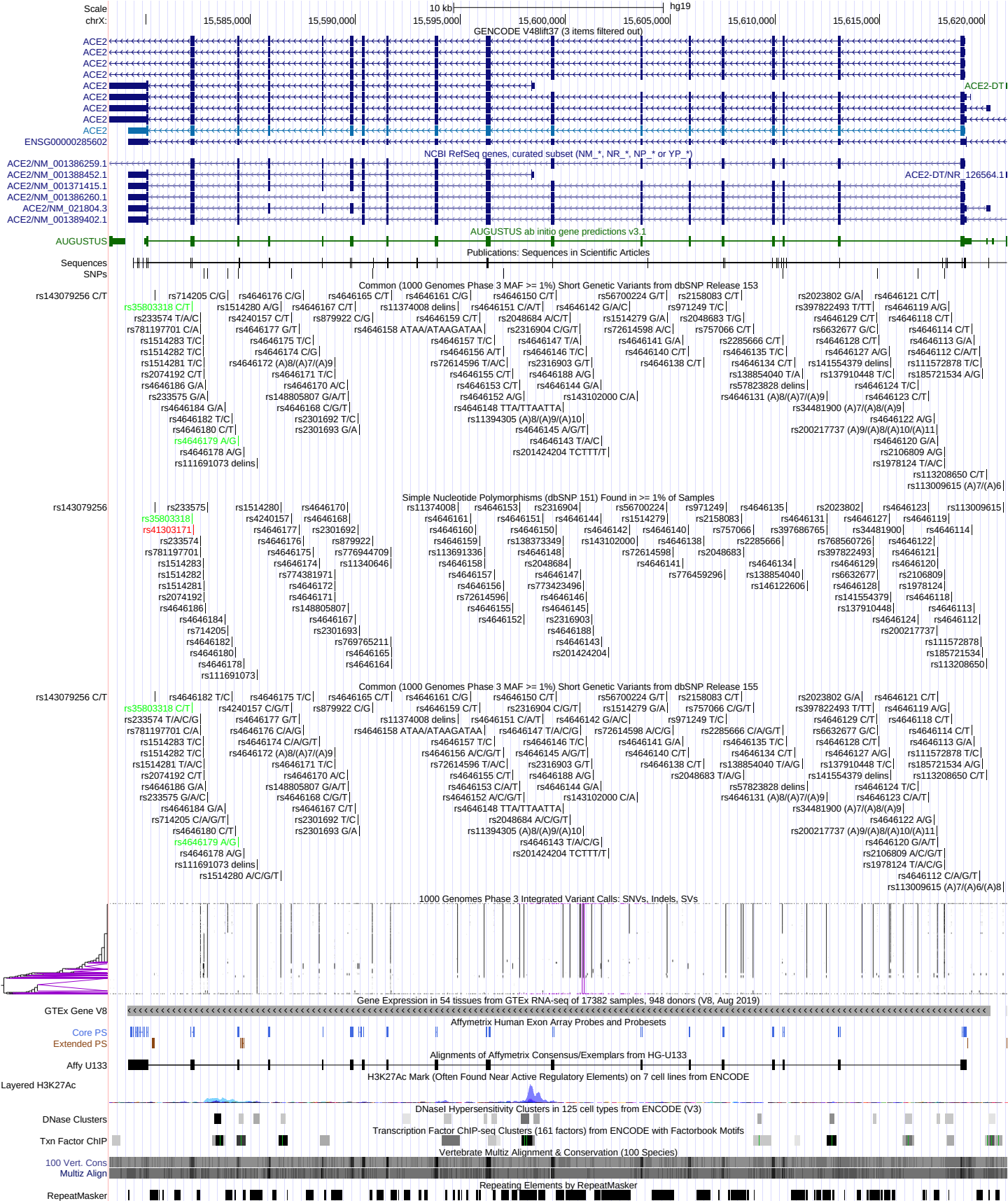
Scale
chrX:
10 kb
hg19
15,585,000
15,590,000
15,595,000
15,600,000
15,605,000
15,610,000
15,615,000
15,620,000
GENCODE V48lift37 (3 items filtered out)
ACE2
ACE2-DT
ENSG00000285602
NCBI RefSeq genes, curated subset (NM_*, NR_*, NP_* or YP_*)
ACE2/NM_001386259.1
ACE2/NM_001388452.1
ACE2/NM_001371415.1
ACE2/NM_001386260.1
ACE2/NM_021804.3
ACE2/NM_001389402.1
ACE2-DT/NR_126564.1
AUGUSTUS
AUGUSTUS ab initio gene predictions v3.1
Sequences
SNPs
Publications: Sequences in Scientific Articles
Common (1000 Genomes Phase 3 MAF >= 1%) Short Genetic Variants from dbSNP Release 153
rs143079256 C/T
Simple Nucleotide Polymorphisms (dbSNP 151) Found in >= 1% of Samples
Common (1000 Genomes Phase 3 MAF >= 1%) Short Genetic Variants from dbSNP Release 155
1000 Genomes Phase 3 Integrated Variant Calls: SNVs, Indels, SVs
GTEx Gene V8
Gene Expression in 54 tissues from GTEx RNA-seq of 17382 samples, 948 donors (V8, Aug 2019)
Core PS
Extended PS
Affymetrix Human Exon Array Probes and Probesets
Affy U133
Alignments of Affymetrix Consensus/Exemplars from HG-U133
Layered H3K27Ac
H3K27Ac Mark (Often Found Near Active Regulatory Elements) on 7 cell lines from ENCODE
DNase Clusters
DNaseI Hypersensitivity Clusters in 125 cell types from ENCODE (V3)
Txn Factor ChIP
Transcription Factor ChIP-seq Clusters (161 factors) from ENCODE with Factorbook Motifs
100 Vert. Cons
Multiz Align
Vertebrate Multiz Alignment & Conservation (100 Species)
RepeatMasker
Repeating Elements by RepeatMasker