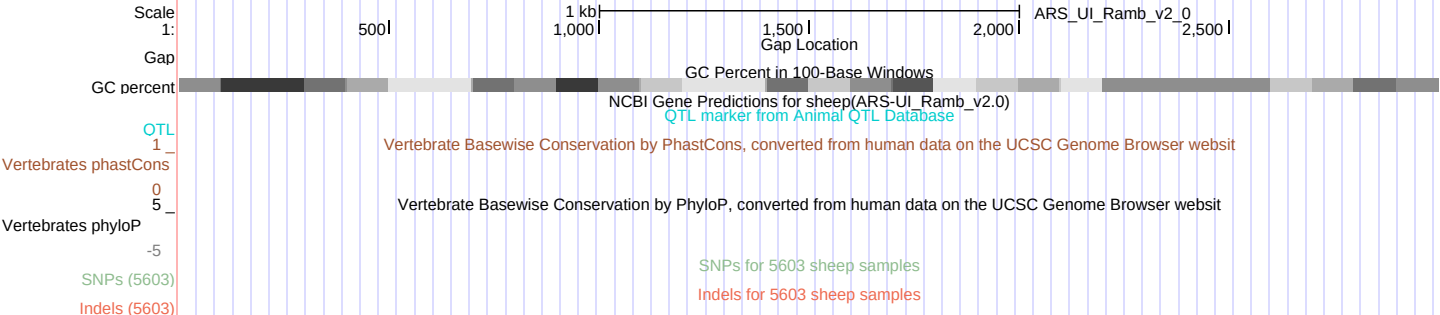
Scale
1:
500
1 kb
1,000
1,500
2,000
ARS_UI_Ramb_v2_0
2,500
Gap Location
Gap
GC percent
GC Percent in 100-Base Windows
NCBI Gene Predictions for sheep(ARS-UI_Ramb_v2.0)
QTL
QTL marker from Animal QTL Database
1
Vertebrate Basewise Conservation by PhastCons, converted from human data on the UCSC Genome Browser websit
Vertebrates phastCons
0
5
Vertebrate Basewise Conservation by PhyloP, converted from human data on the UCSC Genome Browser websit
Vertebrates phyloP
-5
SNPs (5603)
SNPs for 5603 sheep samples
Indels (5603)
Indels for 5603 sheep samples