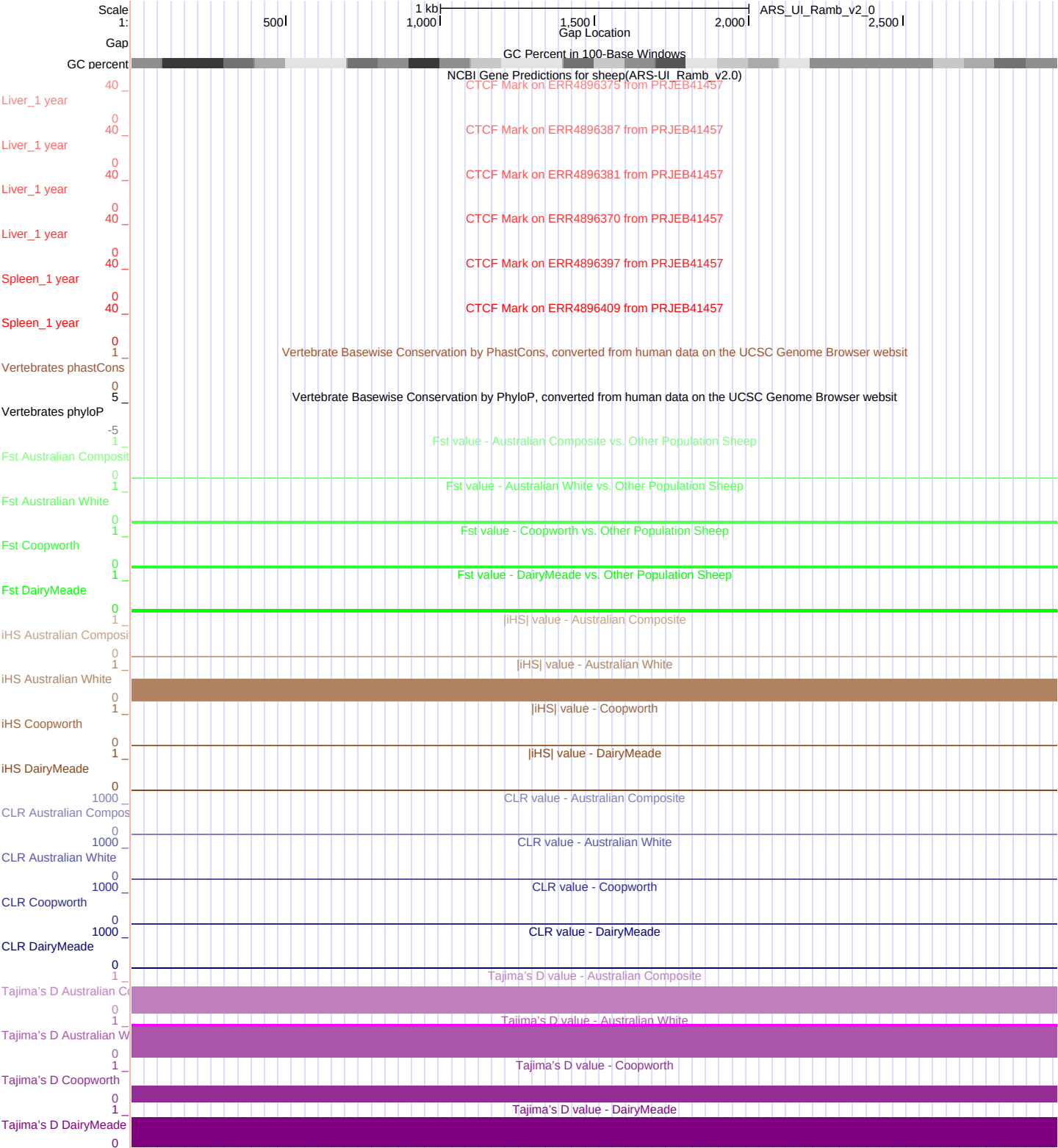
Scale
1:
1 kb
ARS_UI_Ramb_v2_0
500
1,000
1,500
2,000
2,500
Gap Location
Gap
GC percent
GC Percent in 100-Base Windows
NCBI Gene Predictions for sheep(ARS-UI_Ramb_v2.0)
CTCF Mark on ERR4896375 from PRJEB41457
Liver_1 year
CTCF Mark on ERR4896387 from PRJEB41457
Liver_1 year
CTCF Mark on ERR4896381 from PRJEB41457
Liver_1 year
CTCF Mark on ERR4896370 from PRJEB41457
Liver_1 year
CTCF Mark on ERR4896397 from PRJEB41457
Spleen_1 year
CTCF Mark on ERR4896409 from PRJEB41457
Spleen_1 year
Vertebrate Basewise Conservation by PhastCons, converted from human data on the UCSC Genome Browser websit
Vertebrates phastCons
Vertebrate Basewise Conservation by PhyloP, converted from human data on the UCSC Genome Browser websit
Vertebrates phyloP
Fst value - Australian Composite vs. Other Population Sheep
Fst Australian Composit
Fst value - Australian White vs. Other Population Sheep
Fst Australian White
Fst value - Coopworth vs. Other Population Sheep
Fst Coopworth
Fst value - DairyMeade vs. Other Population Sheep
Fst DairyMeade
|iHS| value - Australian Composite
iHS Australian Composi
|iHS| value - Australian White
iHS Australian White
|iHS| value - Coopworth
iHS Coopworth
|iHS| value - DairyMeade
iHS DairyMeade
CLR value - Australian Composite
CLR Australian Compos
CLR value - Australian White
CLR Australian White
CLR value - Coopworth
CLR Coopworth
CLR value - DairyMeade
CLR DairyMeade
Tajima's D value - Australian Composite
Tajima's D Australian C
Tajima's D value - Australian White
Tajima's D Australian W
Tajima's D value - Coopworth
Tajima's D Coopworth
Tajima's D value - DairyMeade
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