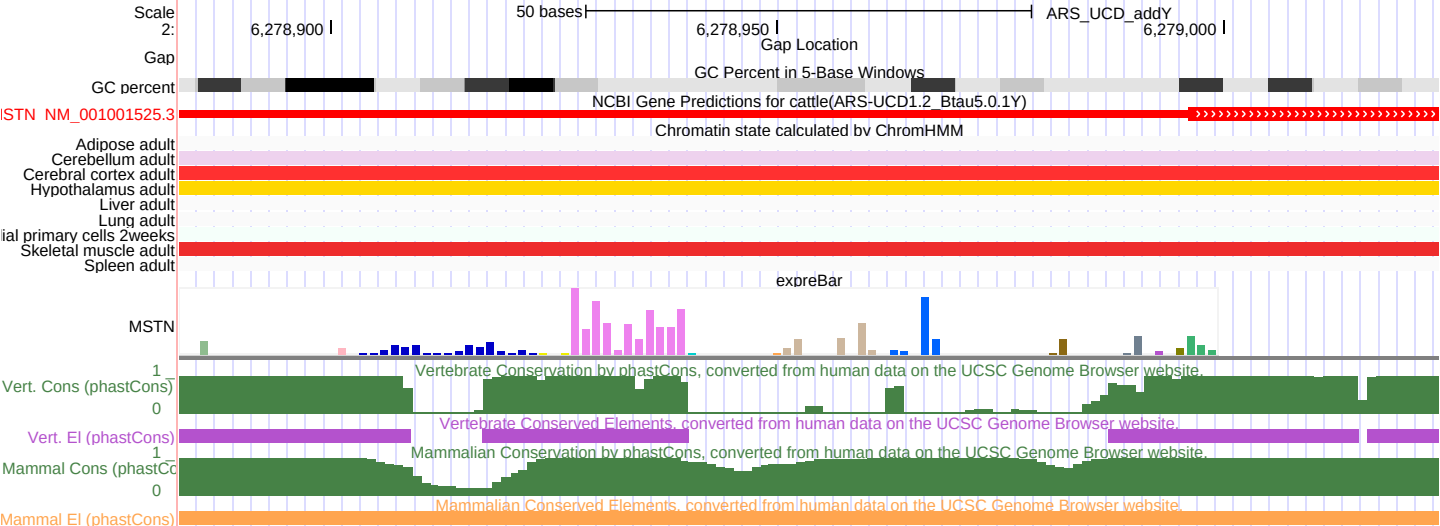
Scale
2:
50 bases
ARS_UCD_addY
6,278,900
6,278,950
6,279,000
Gap Location
Gap
GC Percent in 5-Base Windows
GC percent
ISTN NM_001001525.3
NCBI Gene Predictions for cattle(ARS-UCD1.2 Btau5.0.1Y)
Chromatin state calculated by ChromHMM
Adipose adult
Cerebellum adult
Cerebral cortex adult
Hypothalamus adult
Liver adult
Lung adult
ial primary cells 2weeks
Skeletal muscle adult
Spleen adult
expreBar
MSTN
1
Vertebrate Conservation by phastCons, converted from human data on the UCSC Genome Browser website.
Vert. Cons (phastCons)
0
Vertebrate Conserved Elements, converted from human data on the UCSC Genome Browser website.
Vert. El (phastCons)
1
Mammalian Conservation by phastCons, converted from human data on the UCSC Genome Browser website.
Mammal Cons (phastCo
0
Mammalian Conserved Elements, converted from human data on the UCSC Genome Browser website.
Mammal El (phastCons)