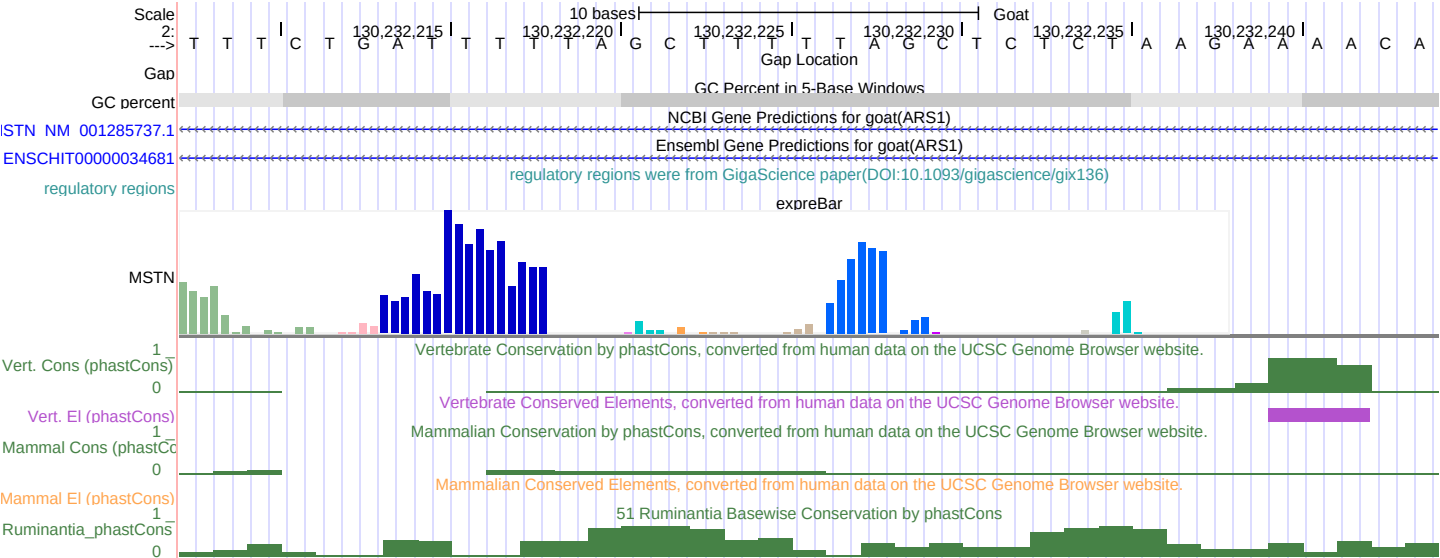
Scale
2:
--->
10 bases
Goat
130,232,215
130,232,220
130,232,225
130,232,230
130,232,235
130,232,240
T T T C T G A T T T T T A G C T T T T T A G C T C T C T A A G A A A A C A
Gap Location
Gap
GC percent
GC Percent in 5-Base Windows
ISTN NM 001285737.1
NCBI Gene Predictions for goat(ARS1)
ENSCHIT00000034681
Ensembl Gene Predictions for goat(ARS1)
regulatory regions
regulatory regions were from GigaScience paper(DOI:10.1093/gigascience/gix136)
expreBar
MSTN
1
Vert. Cons (phastCons)
0
Vertebrate Conservation by phastCons, converted from human data on the UCSC Genome Browser website.
Vert. El (phastCons)
Vertebrate Conserved Elements, converted from human data on the UCSC Genome Browser website.
1
Mammal Cons (phastCo
Mammalian Conservation by phastCons, converted from human data on the UCSC Genome Browser website.
0
Mammal El (phastCons)
Mammalian Conserved Elements, converted from human data on the UCSC Genome Browser website.
1
Ruminantia_phastCons
51 Ruminantia Basewise Conservation by phastCons
0