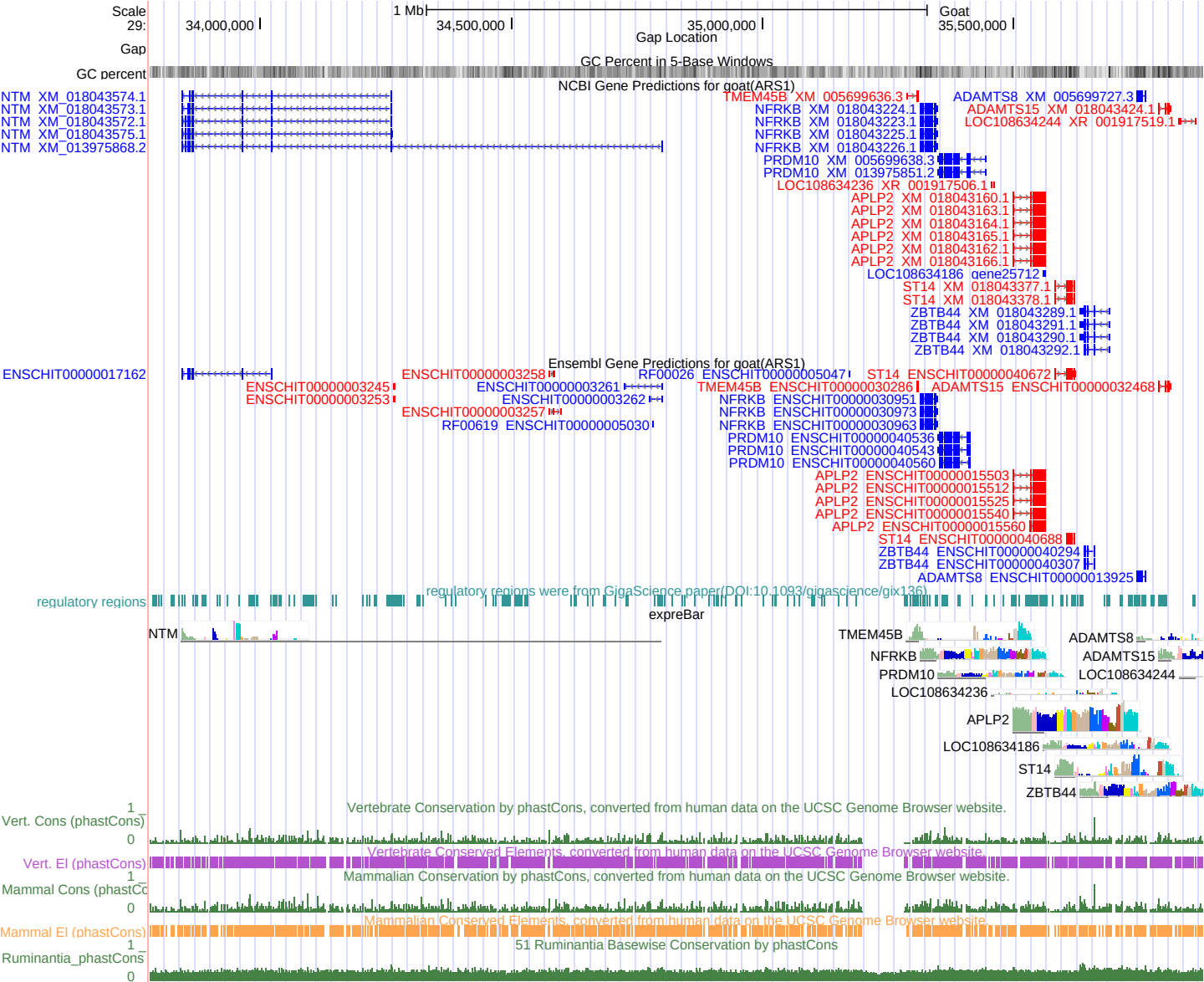
Scale
29:
1 Mb
34,000,000
34,500,000
35,000,000
35,500,000
Goat
Gap
Gap Location
GC percent
GC Percent in 5-Base Windows
NCBI Gene Predictions for goat(ARS1)
NTM XM_018043574.1
NTM XM_018043573.1
NTM XM_018043572.1
NTM XM_018043575.1
NTM XM_013975868.2
TMEM45B XM 005699636.3
NFRKB XM 018043224.1
NFRKB XM 018043223.1
NFRKB XM 018043225.1
NFRKB XM 018043226.1
PRDM10 XM 005699638.3
PRDM10 XM 013975851.2
LOC108634236 XR 001917506.1
APLP2 XM 018043160.1
APLP2 XM 018043163.1
APLP2 XM 018043164.1
APLP2 XM 018043165.1
APLP2 XM 018043162.1
APLP2 XM 018043166.1
LOC108634186 gene25712
ST14 XM 018043377.1
ST14 XM 018043378.1
ZBTB44 XM 018043289.1
ZBTB44 XM 018043291.1
ZBTB44 XM 018043290.1
ZBTB44 XM 018043292.1
ADAMTS8 XM 005699727.3
ADAMTS15 XM 018043424.1
LOC108634244 XR 001917519.1
Ensembl Gene Predictions for goat(ARS1)
ENSCHIT00000017162
ENSCHIT00000003258
ENSCHIT00000003245
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ENSCHIT00000003262
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RF00619 ENSCHIT00000005030
RF00026 ENSCHIT00000005047
TMEM45B ENSCHIT00000030286
NFRKB ENSCHIT00000030951
NFRKB ENSCHIT00000030973
NFRKB ENSCHIT00000030963
PRDM10 ENSCHIT00000040536
PRDM10 ENSCHIT00000040543
PRDM10 ENSCHIT00000040560
ST14 ENSCHIT00000040672
ADAMTS15 ENSCHIT00000032468
APLP2 ENSCHIT00000015503
APLP2 ENSCHIT00000015512
APLP2 ENSCHIT00000015525
APLP2 ENSCHIT00000015540
APLP2 ENSCHIT00000015560
ST14 ENSCHIT00000040688
ZBTB44 ENSCHIT00000040294
ZBTB44 ENSCHIT00000040307
ADAMTS8 ENSCHIT00000013925
regulatory regions
regulatory regions were from GigaScience paper(DOI:10.1093/gigascience/gix136)
expreBar
NTM
TMEM45B
NFRKB
PRDM10
LOC108634236
APLP2
LOC108634186
ST14
ZBTB44
ADAMTS8
ADAMTS15
LOC108634244
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
0
Mammalian Conservation by phastCons, converted from human data on the UCSC Genome Browser website.
Mammal El (phastCons)
Mammalian Conserved Elements, converted from human data on the UCSC Genome Browser website.
1
Ruminantia_phastCons
0
51 Ruminantia Basewise Conservation by phastCons