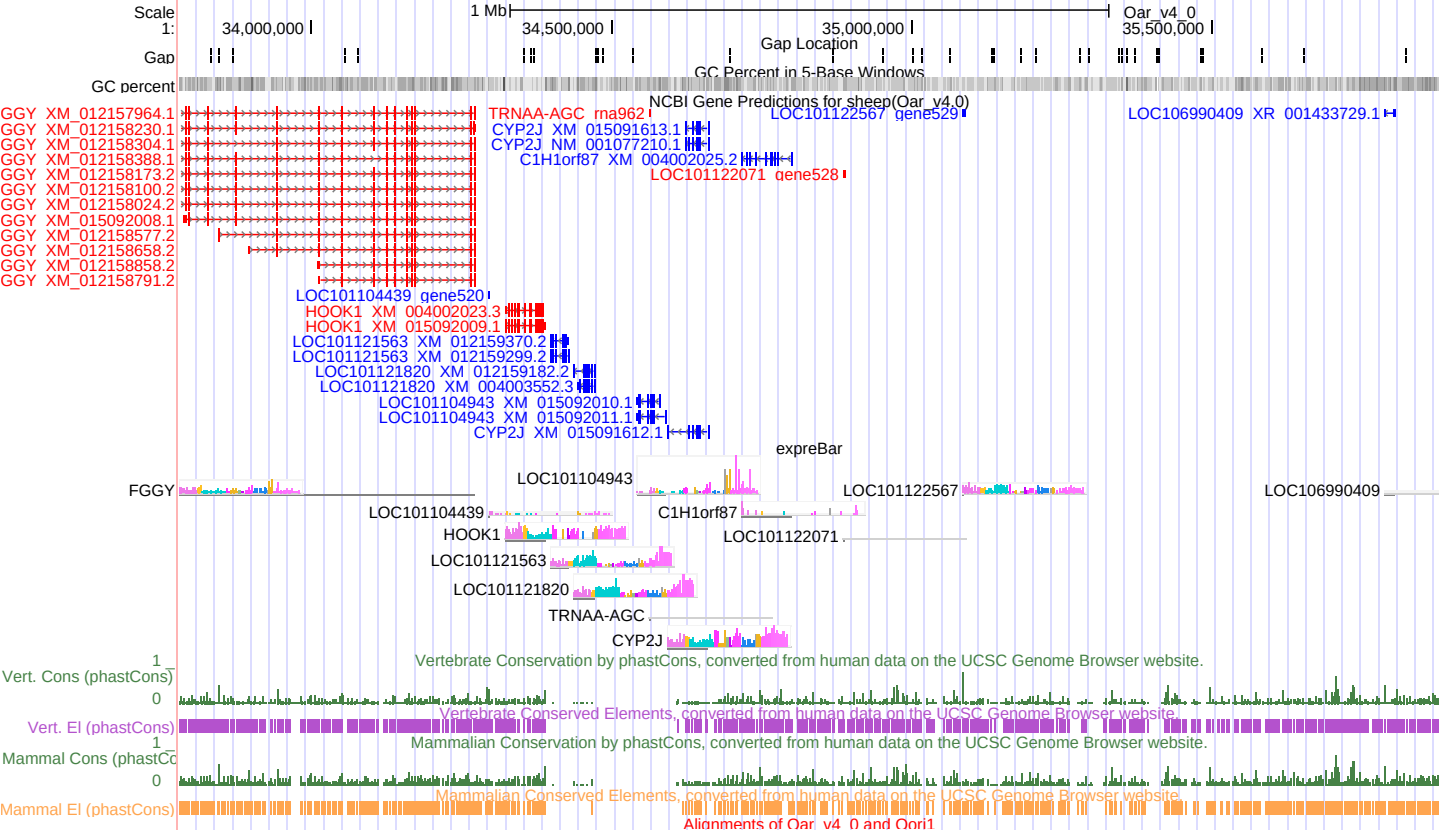
Scale
1:
1 Mb
Oar_v4_0
34,000,000
34,500,000
35,000,000
35,500,000
Gap
Gap Location
GC percent
GC Percent in 5-Base Windows
NCBI Gene Predictions for sheep(Oar_v4.0)
GGY XM_012157964.1
GGY XM_012158230.1
GGY XM_012158304.1
GGY XM_012158388.1
GGY XM_012158173.2
GGY XM_012158100.2
GGY XM_012158024.2
GGY XM_015092008.1
GGY XM_012158577.2
GGY XM_012158658.2
GGY XM_012158858.2
GGY XM_012158791.2
TRNAA-AGC rna962
LOC101122567 gene529
LOC106990409 XR 001433729.1
CYP2J XM 015091613.1
CYP2J NM 001077210.1
C1H1orf87 XM 004002025.2
LOC101122071 gene528
LOC101104439 gene520
HOOK1 XM 004002023.3
HOOK1 XM 015092009.1
LOC101121563 XM 012159370.2
LOC101121563 XM 012159299.2
LOC101121820 XM 012159182.2
LOC101121820 XM 004003552.3
LOC101104943 XM 015092010.1
LOC101104943 XM 015092011.1
CYP2J XM 015091612.1
expreBar
FGGY
LOC101104943
LOC101122567
LOC106990409
LOC101104439
C1H1orf87
HOOK1
LOC101122071
LOC101121563
LOC101121820
TRNAA-AGC
CYP2J
Vertebrate Conservation by phastCons, converted from human data on the UCSC Genome Browser website.
Vert. Cons (phastCons)
1
0
Vertebrate Conserved Elements, converted from human data on the UCSC Genome Browser website.
Vert. El (phastCons)
Mammalian Conservation by phastCons, converted from human data on the UCSC Genome Browser website.
Mammal Cons (phastCo
1
0
Mammalian Conserved Elements, converted from human data on the UCSC Genome Browser website.
Mammal El (phastCons)
Alignments of Oar_v4_0 and Oori1