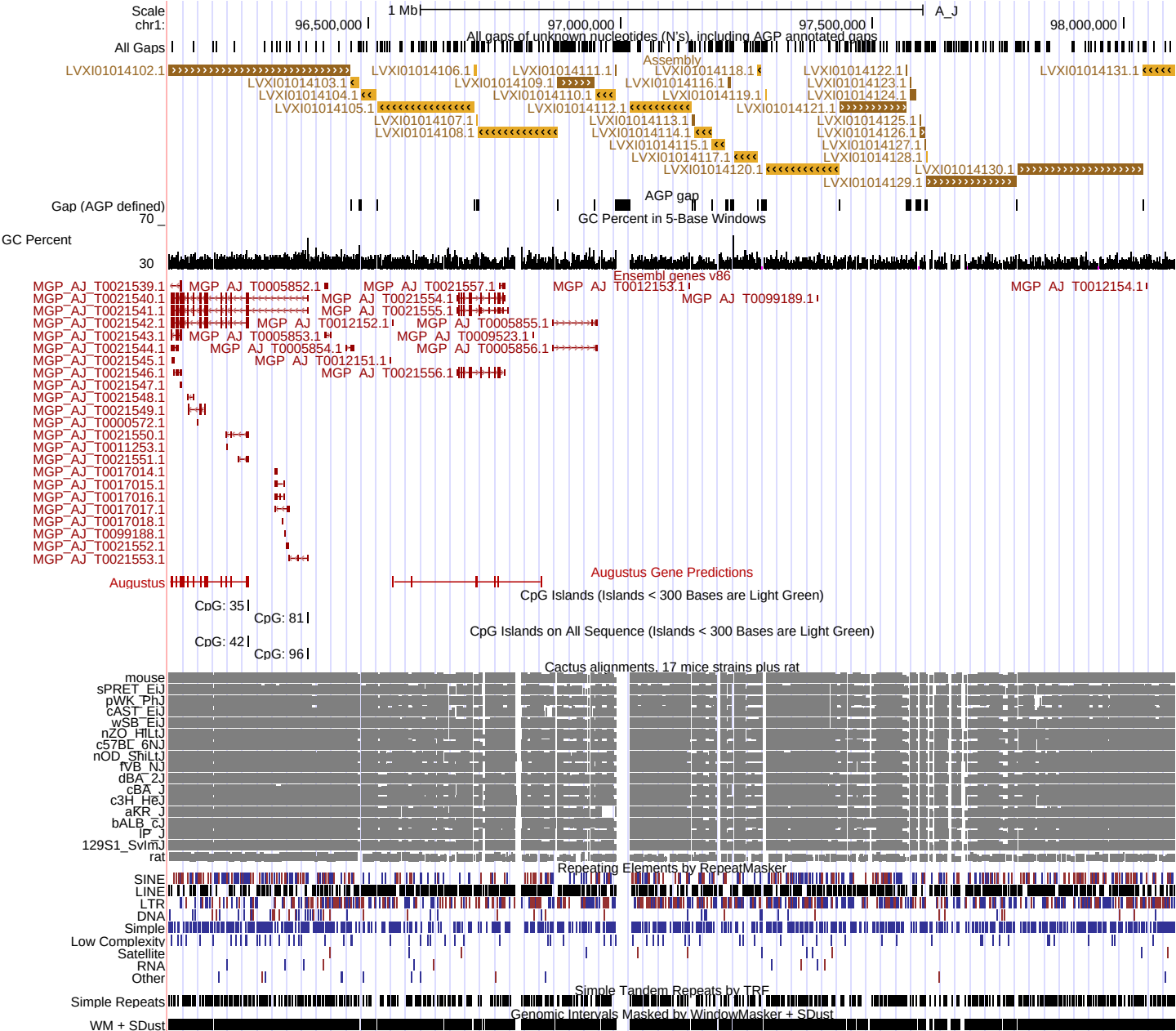
Scale
chr1:
1 Mb
A_J
96,500,000
97,000,000
97,500,000
98,000,000
All Gaps
All gaps of unknown nucleotides (N's), including AGP annotated gaps
Assembly
LVXI01014102.1
LVXI01014106.1
LVXI01014111.1
LVXI01014118.1
LVXI01014122.1
LVXI01014131.1
LVXI01014103.1
LVXI01014109.1
LVXI01014116.1
LVXI01014123.1
LVXI01014104.1
LVXI01014110.1
LVXI01014119.1
LVXI01014124.1
LVXI01014105.1
LVXI01014112.1
LVXI01014121.1
LVXI01014107.1
LVXI01014113.1
LVXI01014125.1
LVXI01014108.1
LVXI01014114.1
LVXI01014126.1
LVXI01014115.1
LVXI01014127.1
LVXI01014117.1
LVXI01014128.1
LVXI01014120.1
LVXI01014130.1
LVXI01014129.1
AGP gap
Gap (AGP defined)
GC Percent in 5-Base Windows
GC Percent
70
30
Ensembl genes v86
MGP_AJ_T0021539.1
MGP_AJ_T0005852.1
MGP_AJ_T0021557.1
MGP_AJ_T0012153.1
MGP_AJ_T0012154.1
MGP_AJ_T0021540.1
MGP_AJ_T0021554.1
MGP_AJ_T0099189.1
MGP_AJ_T0021541.1
MGP_AJ_T0021555.1
MGP_AJ_T0021542.1
MGP_AJ_T0012152.1
MGP_AJ_T0005855.1
MGP_AJ_T0021543.1
MGP_AJ_T0005853.1
MGP_AJ_T0009523.1
MGP_AJ_T0021544.1
MGP_AJ_T0005854.1
MGP_AJ_T0005856.1
MGP_AJ_T0021545.1
MGP_AJ_T0012151.1
MGP_AJ_T0021546.1
MGP_AJ_T0021556.1
MGP_AJ_T0021547.1
MGP_AJ_T0021548.1
MGP_AJ_T0021549.1
MGP_AJ_T0000572.1
MGP_AJ_T0021550.1
MGP_AJ_T0011253.1
MGP_AJ_T0021551.1
MGP_AJ_T0017014.1
MGP_AJ_T0017015.1
MGP_AJ_T0017016.1
MGP_AJ_T0017017.1
MGP_AJ_T0017018.1
MGP_AJ_T0099188.1
MGP_AJ_T0021552.1
MGP_AJ_T0021553.1
Augustus Gene Predictions
Augustus
CpG Islands (Islands < 300 Bases are Light Green)
CpG: 35
CpG: 81
CpG Islands on All Sequence (Islands < 300 Bases are Light Green)
CpG: 42
CpG: 96
Cactus alignments, 17 mice strains plus rat
mouse
sPRET_EiJ
pWK_PhJ
cAST_EiJ
wSB_EiJ
nZO_HlLtJ
c57BL_6NJ
nOD_ShiLtJ
NZB_NJ
dBA_2J
cBA_J
c3H_HeJ
aKR_J
bALB_cJ
IP_J
129S1_SvImJ
rat
Repeating Elements by RepeatMasker
SINE
LINE
LTR
DNA
Simple
Low Complexity
Satellite
RNA
Other
Simple Tandem Repeats by TRF
Simple Repeats
Genomic Intervals Masked by WindowMasker + SDust
WM + SDust