Hsap_KVLQT1

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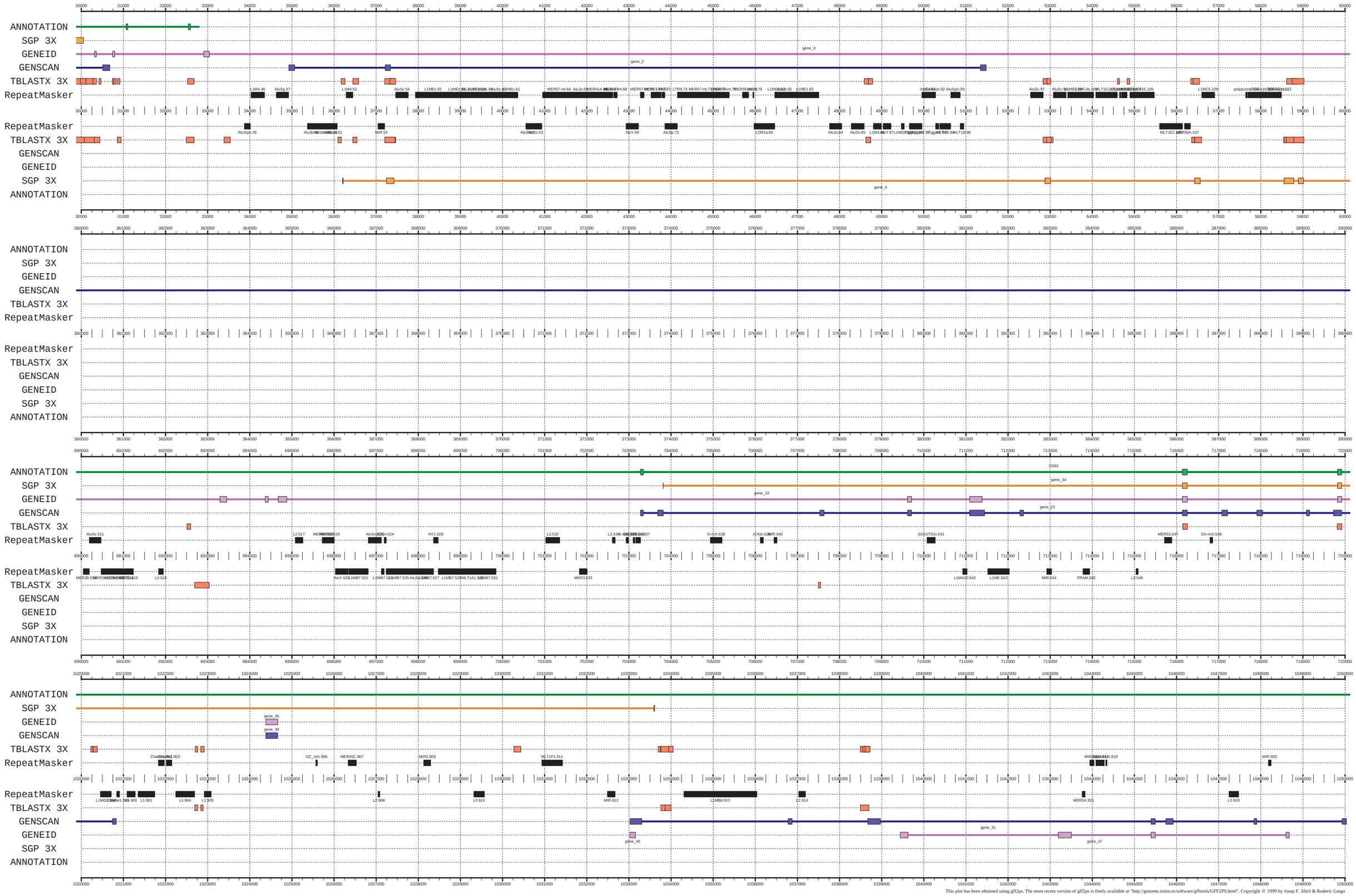
Hsap_KVLQT1



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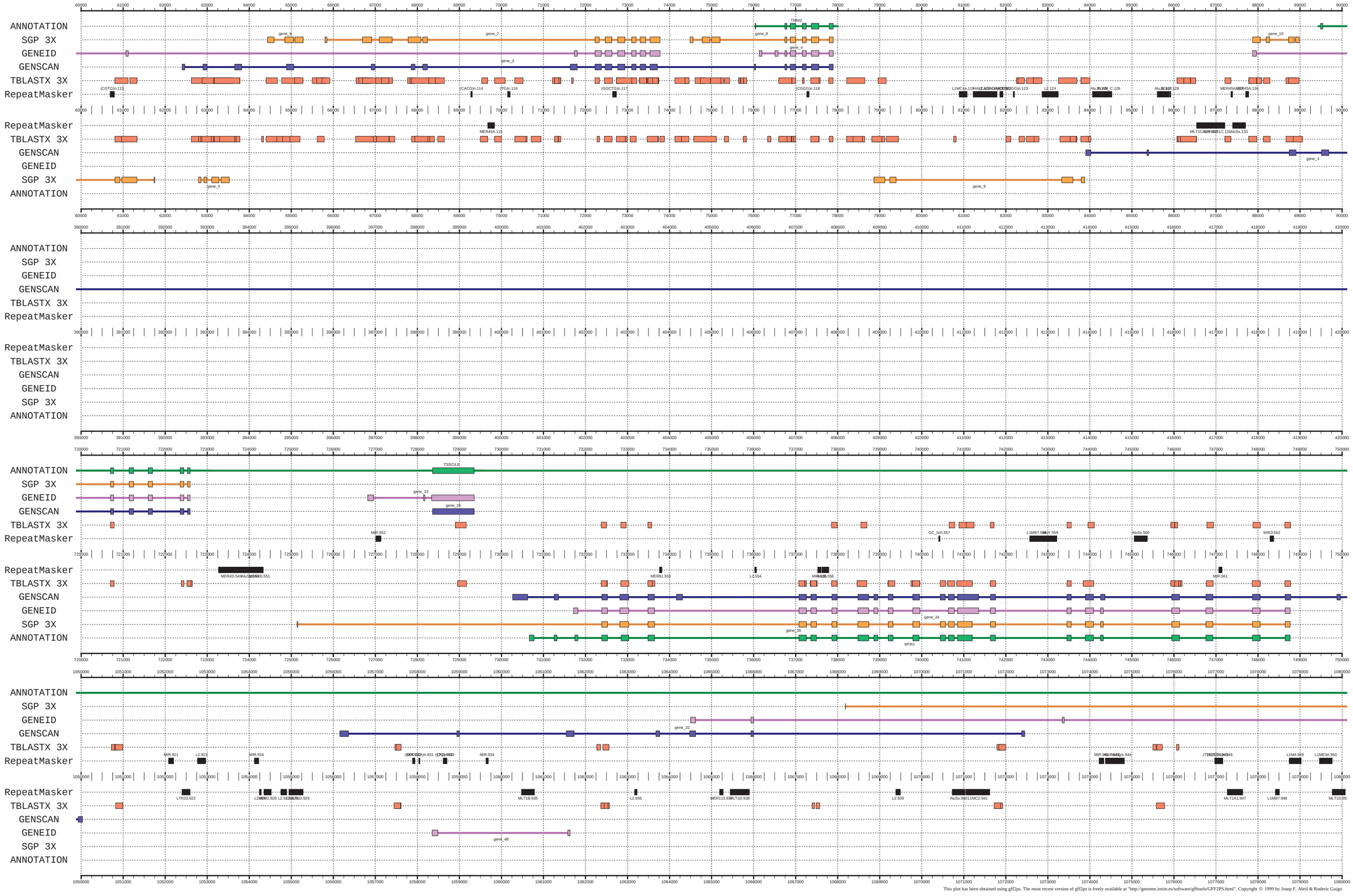
Hsap_KVLQT1



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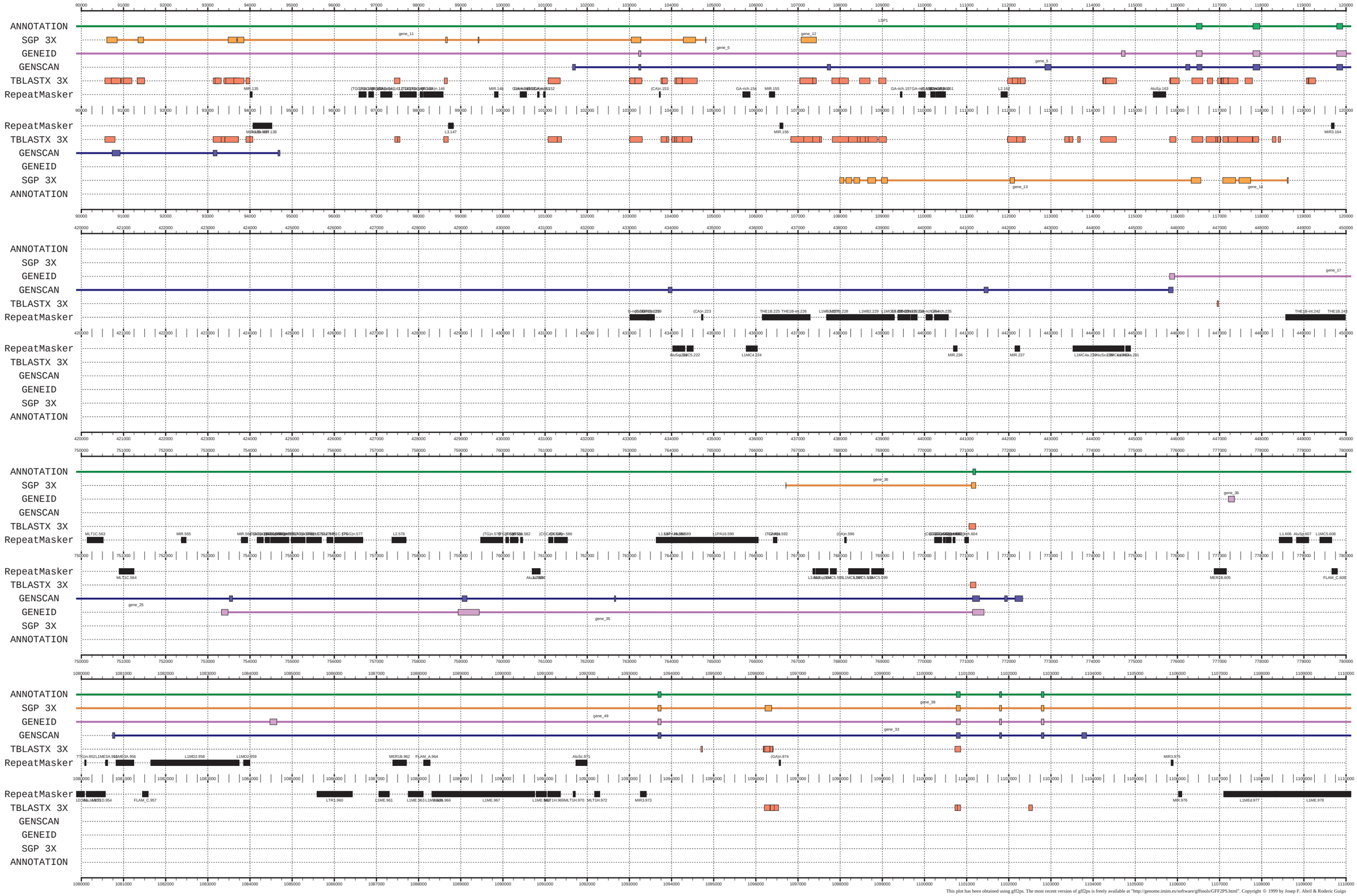

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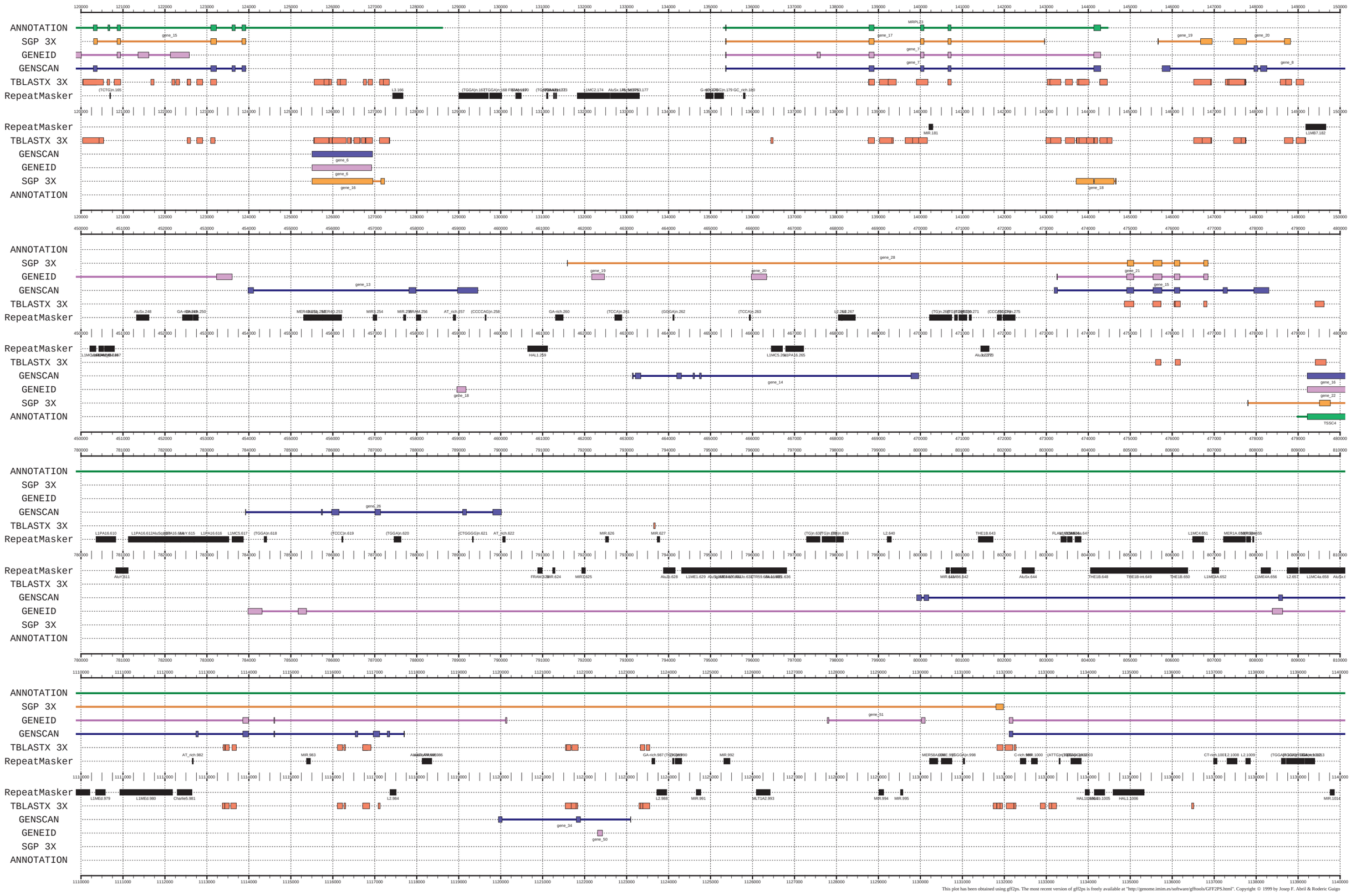
ANNOTATION
SGP 3X
GENEID
GENSCAN
TBLASTX 3X
RepeatMasker
gene_15
gene_17
gene_19
gene_20
gene_7
gene_8
MRPL23
(TCTG)n.165
L3.166
(TGGA)n.167
(TGGA)n.168
(TGGA)n.169
(TGGA)n.170
L1MC2.174
AluSx.175
L1MC3.177
GC_rich.180
MIR.181
L1MB7.182
gene_6
gene_16
gene_18
gene_28
gene_21
gene_13
AluSx.248
GA-rich.249
GA-rich.250
AT_rich.257
(CCCCAG)n.258
GA-rich.260
(TCCA)n.261
(GGGA)n.262
(TCCA)n.263
(TG)n.266
(TG)n.267
(TG)n.270
(TG)n.271
(CCCAG)n.275
HAL1.259
L1MC5.264
L1PA16.265
gene_14
gene_16
gene_22
TSSC4
(CTCTG)n.618
L1PA16.610
L1PA16.612
L1MC5.617
(TGGA)n.618
(TCCC)n.619
(TGGA)n.620
(CTGGGG)n.621
AT_rich.622
MIR.626
MIR.627
(TG)n.637
(TG)n.638
(TG)n.639
L2.640
THE1B.643
L1MC4.651
MER1A.653
MER1B.655
AluY.611
FRAM.623
MIR.624
MIR.625
AluJb.628
L1ME1.629
THE1B.636
MIR.641
MIR.642
AluSx.644
THE1B.648
TBE1B-int.649
THE1B.650
L1ME4A.652
L1ME4A.656
L2.657
L1MC4a.658
AT_rich.982
MIR.983
GA-rich.987
MIR.992
C-rich.999
MIR.1000
CT-rich.1007
L2.1008
L2.1009
L1MEd.979
L1MEd.980
Charlie5.981
L2.984
L2.988
MIR.991
MLT1A2.993
MIR.994
MIR.995
HAL1.1004
HAL1.1005
HAL1.1006
MIR.1014
gene_34
gene_50
gene_51
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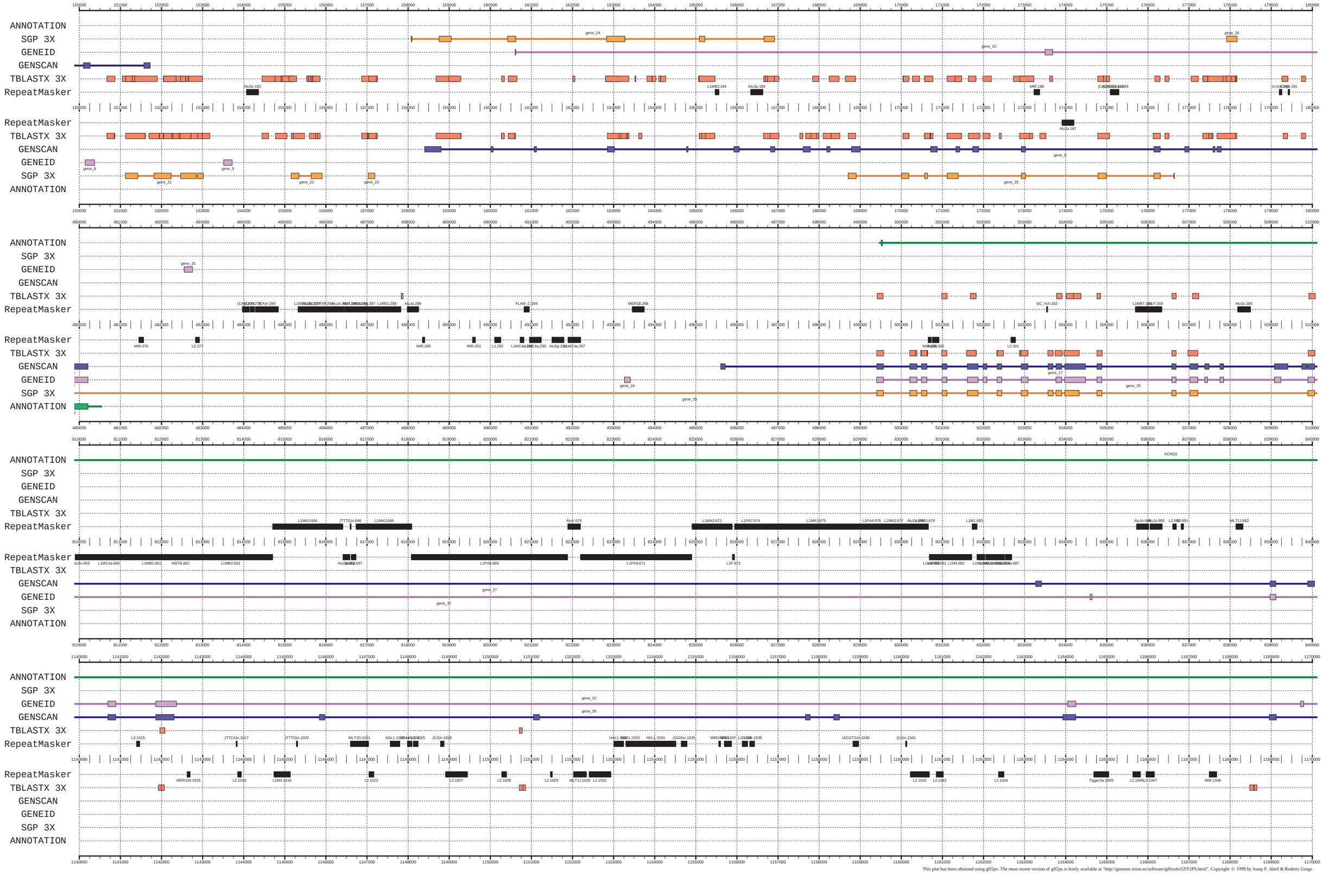
Hsap_KVLQT1



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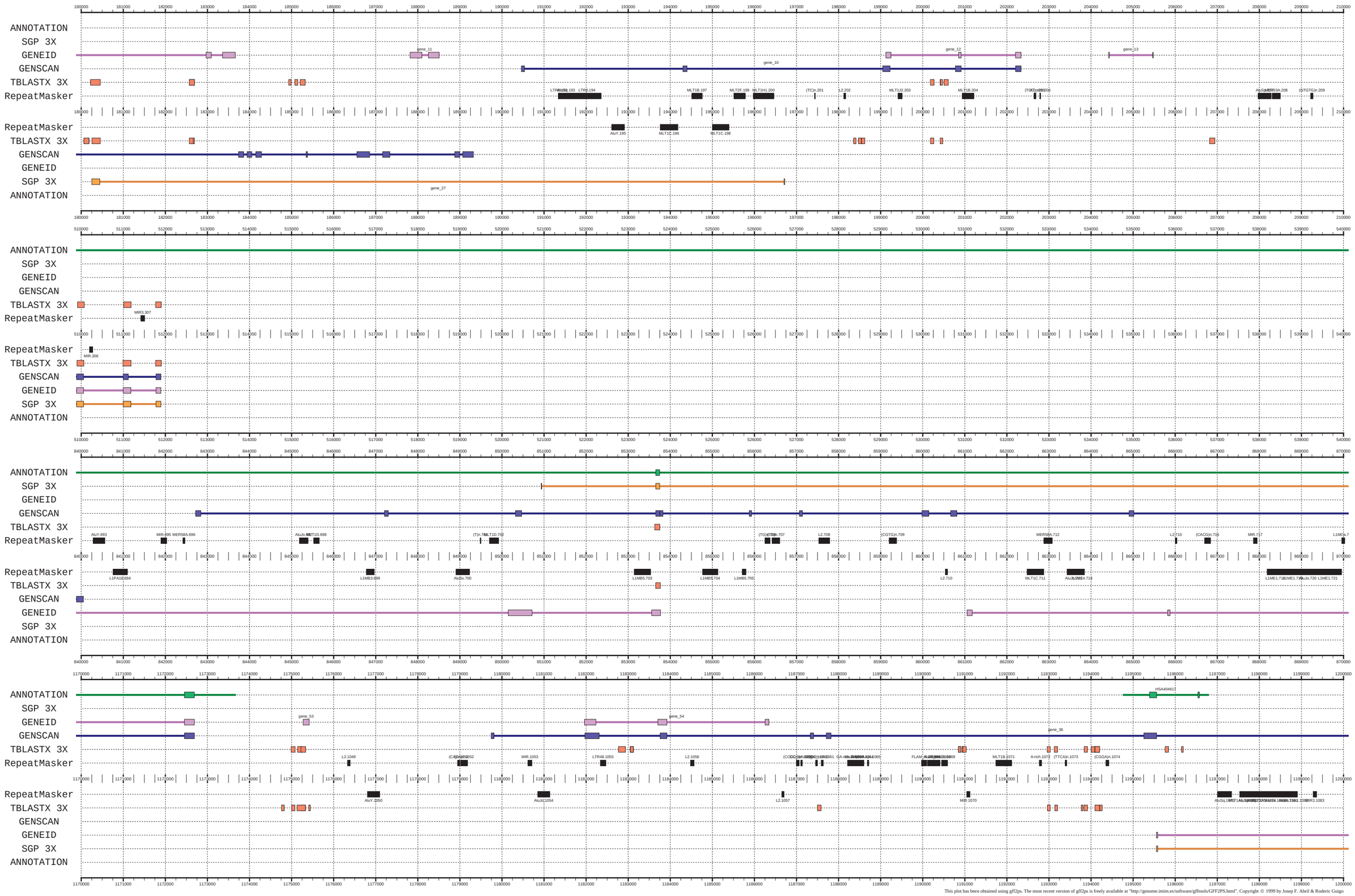

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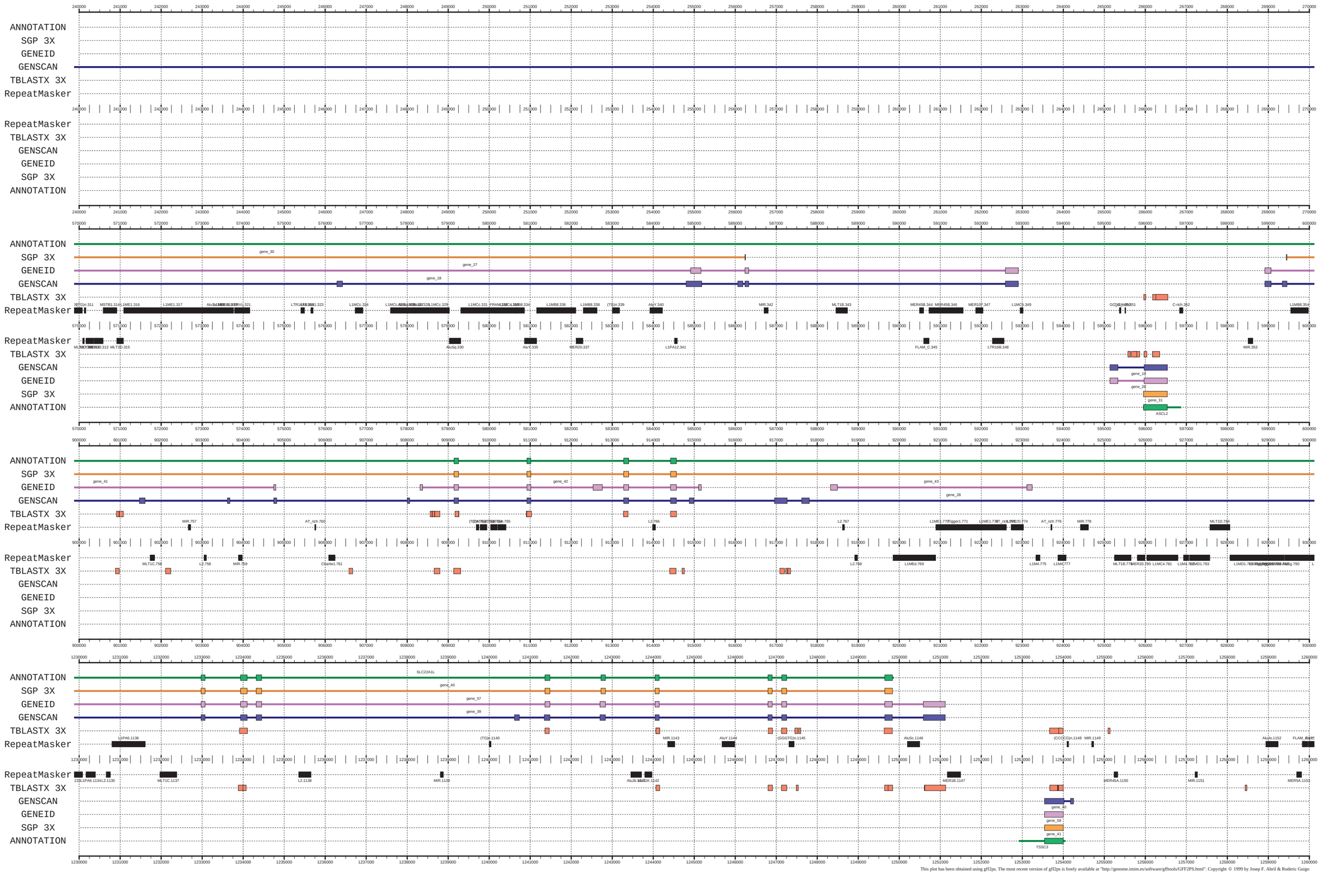

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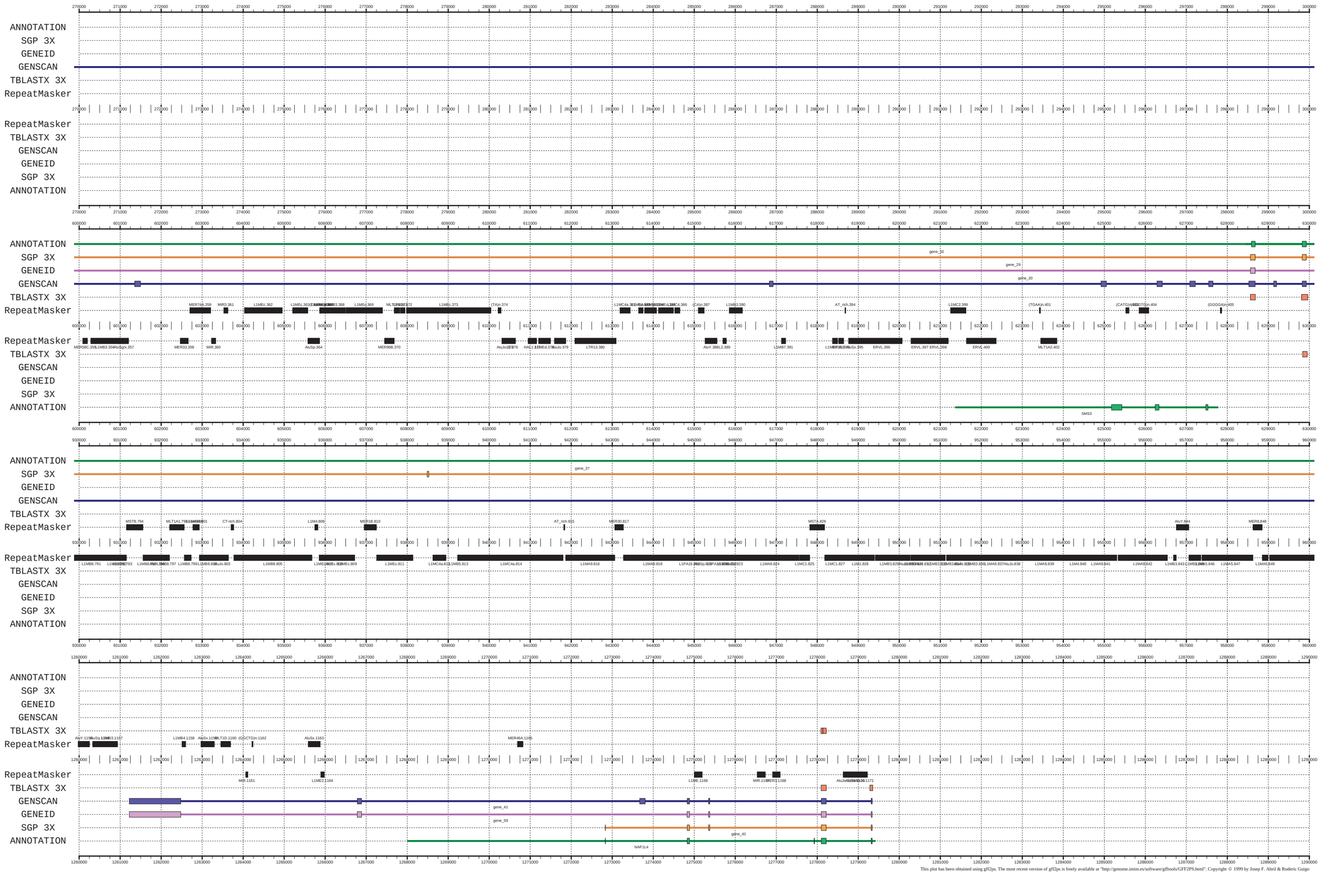
Hsap_KVLQT1



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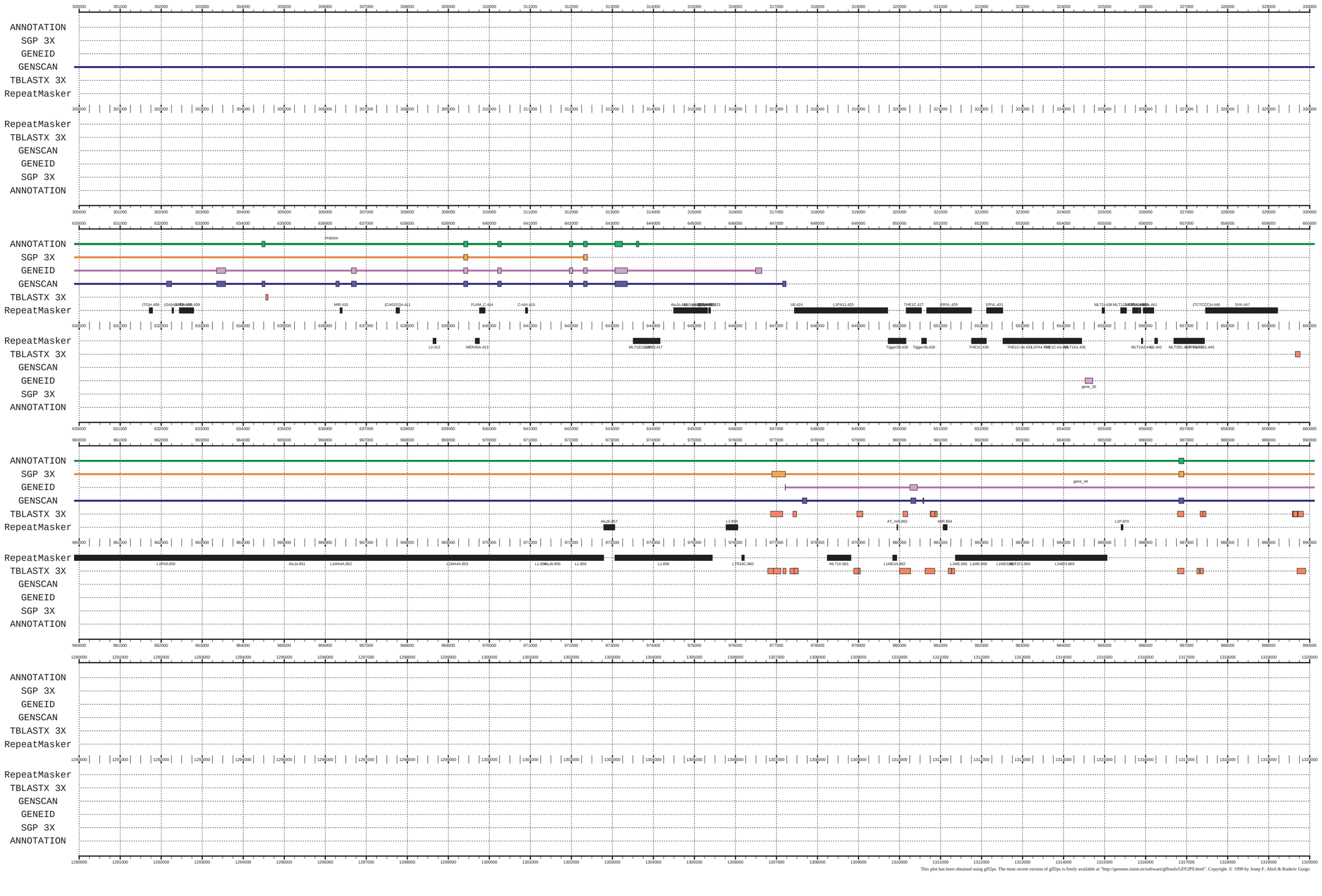
Hsap_KVLQT1



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