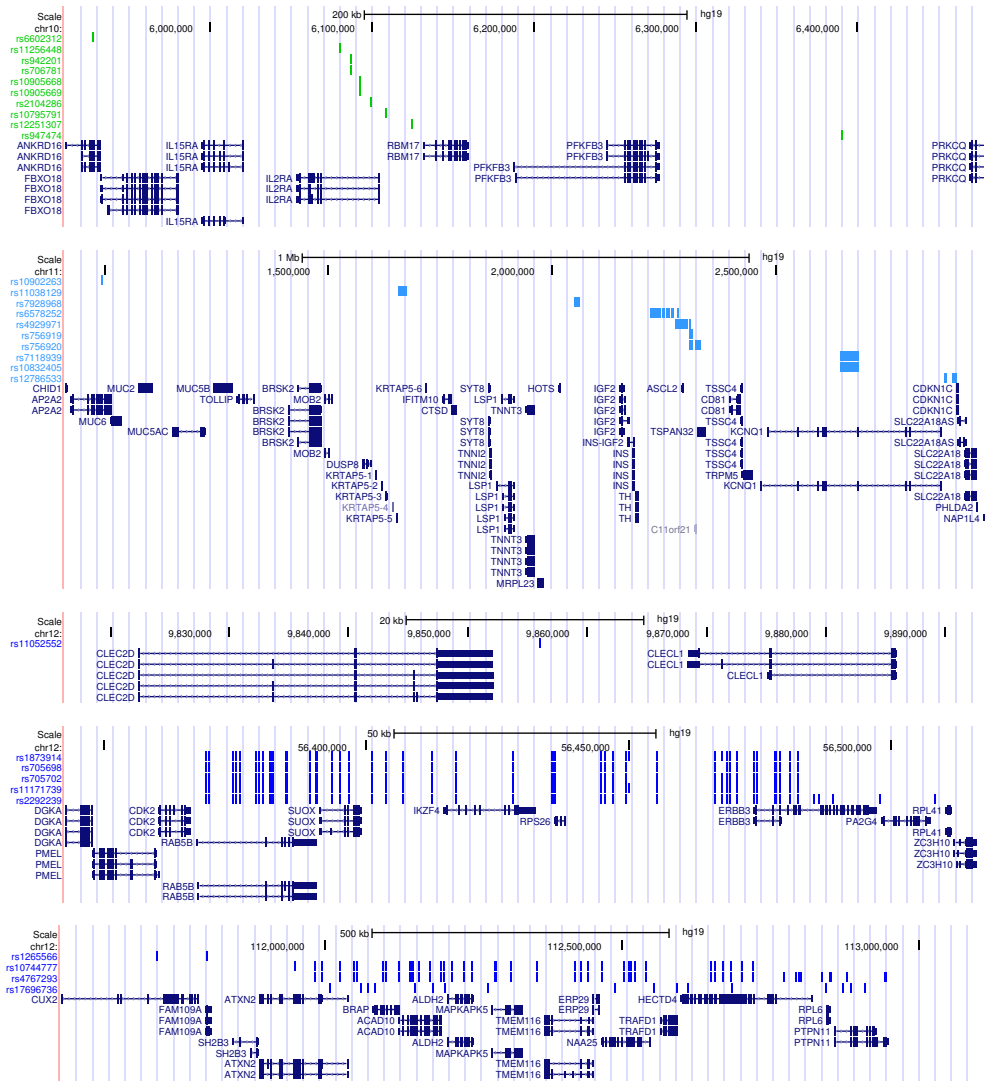
Scale
chr1:
100 kb
hg19
114,150,000
114,200,000
114,250,000
114,300,000
114,350,000
114,400,000
rs4456089
rs17013326
rs1217227
rs1217200
rs1230666
rs1230658
rs1230649
rs6679677
rs1217396
rs2488457
MAGI3
MAGI3
PHTF1
RSBN1
PTPN22
PTPN22
PTPN22
PTPN22
BCL2L15
Scale
chr2:
100 kb
hg19
204,750,000
204,800,000
204,850,000
204,900,000
rs231790
rs231797
rs11571304
rs11571293
rs11571292
rs3087243
rs231726
rs1427676
rs6748358
rs2556345
CTLA4
CTLA4
ICOS
Scale
chr6:
500 kb
hg19
31,500,000
32,000,000
32,500,000
rs3132460
rs2248459
rs2248617
rs1046089
rs760293
rs805301
rs805294
rs3130484
rs3131379
rs2763979
rs1150753
rs2071278
rs3131294
rs377763
rs9268230
rs3132959
rs910049
rs3129932
rs9268403
rs9268418
rs9268429
rs2894254
rs9268480
rs3763307
rs3135377
rs9268557
rs9268645
rs9268853
rs9268877
rs7749092
rs9272219
rs9273363
rs9275134
rs2856688
rs9469220
rs9275418
rs9275523
rs3957146
rs6936863
rs3916765
rs9276429
rs9276431
rs9276432
rs2051549
rs1015575
rs2857212
rs241427
rs4148882
rs241403
rs3101942
HLA-C
HLA-C
MICA
MICA
HLA-B
MICA
MICA
MICA
MICB
MICB
MCCD1
DDX39B
DDX39B
ATP6V1G2
ATP6V1G2
ATP6V1G2
NFKBIL1
NFKBIL1
NFKBIL1
NFKBIL1
LTB
LST1
LST1
LST1
LST1
LST1
NCR3
NCR3
NCR3
PRRC2A
PRRC2A
BAG6
BAG6
BAG6
BAG6
C6orf47
GPANK1
GPANK1
GPANK1
GPANK1
GPANK1
CSNK2B
CSNK2B
LY6G5B
ABHD16A
ABHD16A
C6orf25
C6orf25
LTA
TNF
LTB
AIF1
AIF1
APOM
APOM
LY6G5C
C6orf25
C6orf25
C6orf25
C6orf25
DDAH2
DDAH2
MSH5
MSH5
MSH5
MSH5
SAPCD1
VWA7
LY6G6F
LY6G6D
LY6G6C
DDAH2
CLIC1
CLIC1
CLIC1
VARS
LSM2
HSPA1L
HSPA1A
HSPA1B
C6orf48
C6orf48
C6orf48
C6orf48
C6orf48
C6orf48
C6orf48
C6orf48
C6orf48
C6orf48
SLC44A4
SLC44A4
C2
NEU1
SLC44A4
EHMT2
EHMT2
EHMT2
ZBTB12
C2
C2
C2
C2
C2
NELFE
SKIV2L
STK19
STK19
C4B_2
CYP21A2
CYP21A2
STK19
STK19
C4B_2
C4B
CYP21A2
CYP21A2
TNXB
CFB
C4A
DXO
C4A
C4B
TNXB
TNXB
ATF6B
ATF6B
FKBPL
PRRT1
PPT2
PPT2
PPT2
EGFL8
AGPAT1
RNF5
AGPAT1
AGER
AGER
AGER
AGER
AGER
AGER
AGER
AGER
AGER
PBX2
GPSM3
GPSM3
NOTCH4
C6orf10
C6orf10
C6orf10
BTNL2
HLA-DRA
HLA-DRB5
HLA-DRB1
HLA-DRB1
HLA-DQA1
HLA-DQB1
HLA-DQB1
HLA-DQB2
HLA-DQB2
HLA-DQA2
HLA-DOB
TAP2
TAP2
TAP1
TAP2
PSMB8
PSMB8
TAP1
PSMB9
HLA-DMB
HLA-DMA
BRD2
BRD2
BRD2
BRD2
BRD2
HLA-DOA


Scale
chr10:
200 kb
hg19
6,000,000
6,100,000
6,200,000
6,300,000
6,400,000
rs6602312
rs11256448
rs942201
rs706781
rs10905668
rs10905669
rs2104286
rs10795791
rs12251307
rs947474
ANKRD16
FBXO18
IL15RA
IL2RA
RBM17
PFKFB3
PRKCQ
Scale
chr11:
1 Mb
hg19
1,500,000
2,000,000
2,500,000
rs10902263
rs11038129
rs7928968
rs6578252
rs4929971
rs756919
rs756920
rs7118939
rs10832405
rs12786533
CHID1
AP2A2
MUC2
MUC6
MUC5AC
MUC5B
TOLLIP
BRSK2
MOB2
DUSP8
KRTAP5-1
KRTAP5-2
KRTAP5-3
KRTAP5-4
KRTAP5-5
KRTAP5-6
IFITM10
CTSD
SYT8
LSP1
TNNI2
TNNT3
MRPL23
HOTS
IGF2
INS-IGF2
INS
TH
C11orf21
ASCL2
TSSC4
CD81
TSPAN32
KCNQ1
TRPM5
CDKN1C
SLC22A18AS
SLC22A18
PHLDA2
NAP1L4
Scale
chr12:
20 kb
hg19
9,830,000
9,840,000
9,850,000
9,860,000
9,870,000
9,880,000
9,890,000
rs11052552
CLEC2D
CLECL1
Scale
chr12:
50 kb
hg19
56,400,000
56,450,000
56,500,000
rs1873914
rs705698
rs705702
rs11171739
rs2292239
DGKA
PMEL
CDK2
RAB5B
SUOX
IKZF4
RPS26
ERBB3
PA2G4
RPL41
ZC3H10
Scale
chr12:
500 kb
hg19
112,000,000
112,500,000
113,000,000
rs1265566
rs10744777
rs4767293
rs17696736
CUX2
FAM109A
SH2B3
ATXN2
BRAP
ALDH2
ACAD10
MAPKAPK5
TMEM116
ERP29
NAA25
TRAFD1
HECTD4
RPL6
PTPN11