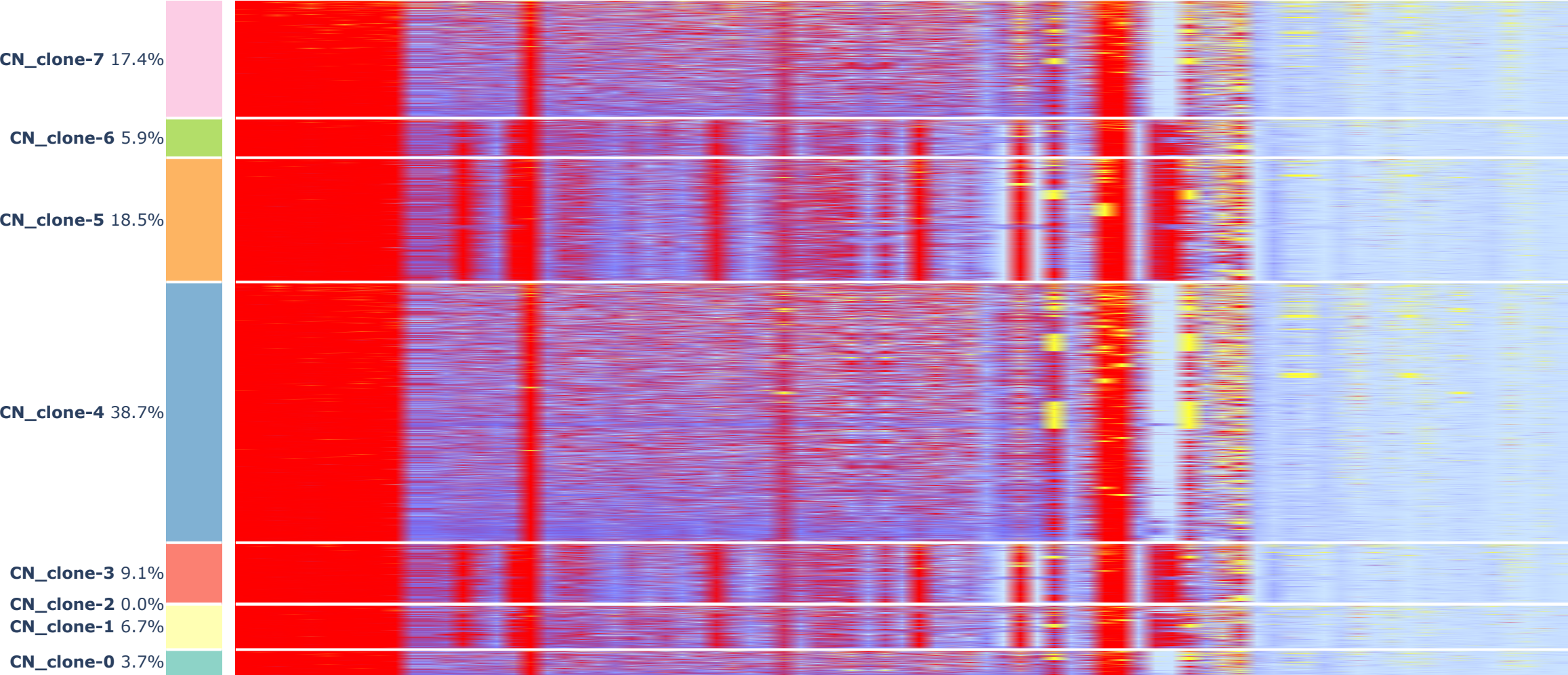


RA19\_10 - RA19\_10-high\_conf\_snvs

6796 cells, 79 features



MLH3 intron variant  
NCOA3 p.Gln1274Arg  
PER1 p.Ser656del  
ARID1A intron variant  
SMAD4 intron variant  
BMPR1B intron variant  
RICTOR intron variant  
PAXIP1 p.Gln413del  
BMPR2 intron variant  
HUWE1 intron variant  
ATXN3 p.Gln305del  
SMAD4 p.Met294Ile  
FAT3 intron variant  
WRN intron variant  
SMAD4 intron variant  
HUWE1 intron variant  
HUWE1 intron variant  
SMAD4 intron variant  
KMT2C intron variant  
BAP1 p.Ala47T=  
FGFR3 intron variant  
NRAS p.Ala59Gly  
MEN1 p.Asp418=  
TP53 p.Cys242Phe  
KRAS p.Gly12Val  
KRAS intron variant  
BRCAC2 p.Leu1521=  
AKT1 intron variant  
MTOR p.Ala1577=  
FANCD2 intron variant  
MEN1 p.His433=  
TP53 intron variant  
INO80 intron variant  
TP53 intron variant  
SPTA1 p.Cys1568Arg  
DNMT3B intron variant  
BAP1 p.Trp613Met  
RECQL5 p.Tyr471=  
TP53 p.Pro72AArg  
EPHA2 intron variant  
PTPRS intron variant  
BAP1 p.Gln261=  
PTPRS intron variant  
TGFB2 intron variant  
PARP2 intron variant  
CHEK2 intron variant  
RICTOR intron variant  
PARP4 p.Ser534=  
MTOR p.Asn999=  
PAK5 intron variant  
POLE intron variant  
BMPR1A intron variant  
BRCA2 p.Ser2474=  
ATM p.Cys219=  
POLN p.Gln121His  
GNAS p.Ile828=  
FLT1 p.Tyr1213=  
RTEL1 intron variant  
BAP1 p.Ser482=  
RPL22 intron variant  
KAT5 intron variant  
HFM1 p.Ile117Val  
KMT2D p.Gly2493=  
DNMT3B p.Asp627=  
TGFB2 p.Leu333=  
HERC2 intron variant  
TGFB2 intron variant  
PIK3R3 p.Ser297=  
PIK3R3 p.Asn283Lys  
TP63 intron variant  
PARP1 intron variant  
KRAS p.Arg161=  
PBRM1 intron variant  
AXIN1 p.Ala609=  
FAT1 p.Gln2933Pro  
RPL22 intron variant  
DNMT3B intron variant  
TP53 intron variant  
CUL5 5\_prime\_UTR variant

