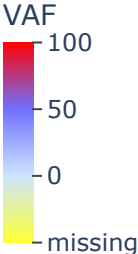
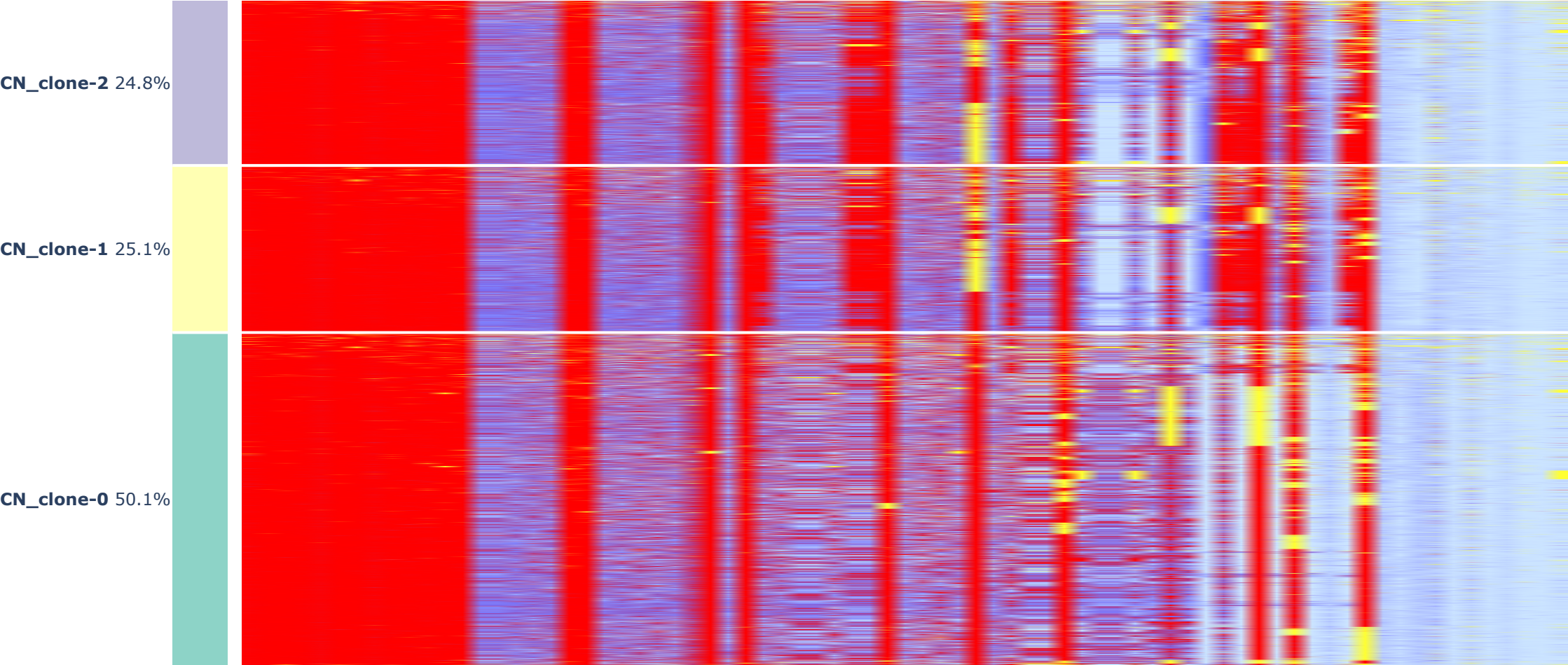


M11 - high\_conf\_snvs\_pre\_condor\_clone\_assignment

6294 cells, 75 features



NCOA3 p.Gln1274Arg  
REV3L splice site variant  
MED12 p.Ser1697Leu  
MLH3 intron variant  
FANCD2 intron variant  
HERC2 p.His313Gln  
SMAD4 intron variant  
TP53 intron variant  
SMAD4 intron variant  
SMAD4 p.Met294Ile  
BMPR1B intron variant  
BRCA2 p.Leu1521=  
SMAD4 p.Gly168AspTer34  
SMAD4 intron variant  
KRAS p.Gly12Val  
INO80 intron variant  
SMAD3 p.Arg268Cys  
MEN1 p.Asp418=  
TP53 p.Gln165Ter  
RREB1 p.His338=  
MAP2K4 p.Pro232Leu  
SMAD2 intron variant  
MEN1 p.His433=  
SMAD2 intron variant  
NCOA3 p.Gln1264=  
TP53 intron variant  
TP53 intron variant  
NCOA3 p.Gln1251=  
BMPR1A p.Pro2Thr  
PTPRS intron variant  
PTPRS intron variant  
TP53 p.Pro72Arg  
PAXIP1 p.Pro415=  
CDKN2A 3 prime UTR variant  
PARP1 intron variant  
WRN intron variant  
TP63 intron variant  
RNF43 p.Ile47Val  
MUS81 p.Ala312=  
PAK5 intron variant  
PAK5 intron variant  
AXIN1 intron variant  
PARP4 intron variant  
PARP4 p.Ser534=  
PBRM1 intron variant  
TTK intron variant  
RTEL1 intron variant  
FANCI p.Gly1302=  
PARP1 intron variant  
RICTOR intron variant  
TENT4A p.Lys373Glu  
FANCA intron variant  
OGG1 intron variant  
CREBBP intron variant  
ATM p.Glu2990Lys  
TGFB2 intron variant  
HFM1 p.Ile117Val  
AKT2 intron variant  
DNMT3B intron variant  
BRCA1 p.Ser694=  
PIK3R3 p.Asn283Lys  
PIK3R3 p.Ser297=  
RECQL5 p.Tyr471=  
AXIN1 p.Ala609=  
CARD11 intron variant  
FLT1 p.Tyr1213=  
KRAS p.Arg161=  
KDR intron variant  
TP53 intron variant  
BMPR1A intron variant  
KAT5 intron variant  
TGFB2 intron variant  
CUL5 5 prime UTR variant  
RNF111 intron variant  
FAT1 p.Gln2933Pro

