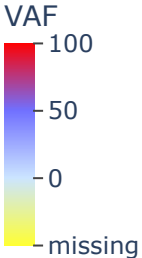
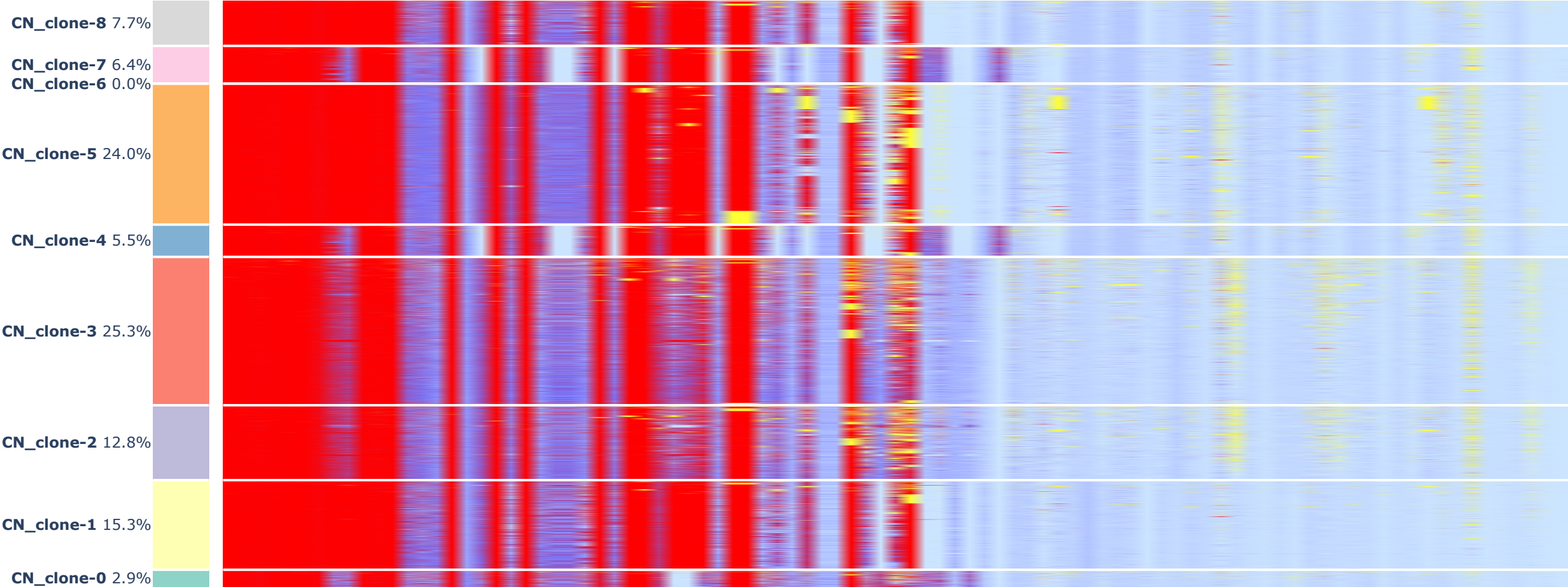


RA21_17 - RA21_17-high_conf_snvs

20123 cells, 91 features



KMT2A p.Asp1872Gly
MED12 p.Ser1697Leu
ELK1 p.Ile385Thr
KMT2D intron_variant
DNMT3A p.Ser337=
REV3L splice_site_variant
TP53 intron_variant
DHX9 p.Ile397Ser
BMPR1B p.Ile170Ser
WRN p.Glu260Asp
PER1 p.Ser655=
RNF43 intron_variant
MSH3 intron_variant
RTEL1 intron_variant
RAD17 splice_site_variant
NCOA3 p.Gln1275_Gln1276del
ERCC3 p.Ser90Phe
KMT2B p.Lys1866Asn
GATA6 5_prime_UTR_variant
BRCA1 p.Asp2Tyr
SLX4 p.Glu921Lys
FANCD2 intron_variant
NCOA3 p.Gln1255=
BRCA2 p.Leu2805Val
RNF111 p.Gln3Ter
KMT2B p.Lys1866Thr
AXIN2 3_prime_UTR_variant
RCTOR intron_variant
KMT2C intron_variant
RAD17 splice_site_variant
RAD17 5_prime_UTR_variant
ATXN3 p.Gln305del
RNF111 intron_variant
MUS81 intron_variant
WRN intron_variant
FAT3 intron_variant
IGFBR1 intron_variant
TP53 intron_variant
SMAD3 p.Ser416Phe
BMPR1B intron_variant
RNF43 p.Arg343His
KMT2C intron_variant
PAXIP1 p.Pro415=
BMPR2 p.Leu200=
TGFBRI2 p.Ser553Thr
BMPR1A p.Pro2Thr
H3C8 p.Asp124=
MTOR p.Ala1577=
BRCA2 p.Leu1521=
KMT2C intron_variant
KMT2C p.Tyr987His
WRN intron_variant
SPTA1 p.Cys1568Arg
HEM1 p.Ser115Pro
MUS81 p.Ala312=
PTPRS intron_variant
PTPRS intron_variant
CHEK2 intron_variant
RREB1 p.His338=
KRAS p.Gly12Aasp
TP53 p.Tyr91Ter
RCTOR intron_variant
HEM1 p.Ile117Val
TP53 p.Pro72Arg
MTOR p.Asn999=
RTEL1 intron_variant
AXIN1 p.Ala609=
PIK3R3 p.Ser297=
PIK3R3 p.Asn283Lys
AURKA intron_variant
FLT1 p.Tyr12T3=
FANCA intron_variant
PARP1 intron_variant
MCM3 2kb upstream_variant
ERBB2 intron_variant
PBRM1 intron_variant
PAK5 p.Leu7T2=
TP53 intron_variant
BMPR1A intron_variant
RECQL5 p.Tyr471=
FANCL p.Gly1302=
CARD11 intron_variant
KMT2C p.Pro1669Arg
BRCA1 p.Gln356Arg
KAT5 intron_variant
KRAS p.Arg161=
TP53 intron_variant
BRCA2 p.Ser2414=
FAT1 p.Gln2933Pro
DNMT3B intron_variant
CUL5 5_prime_UTR_variant

