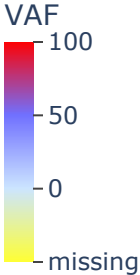
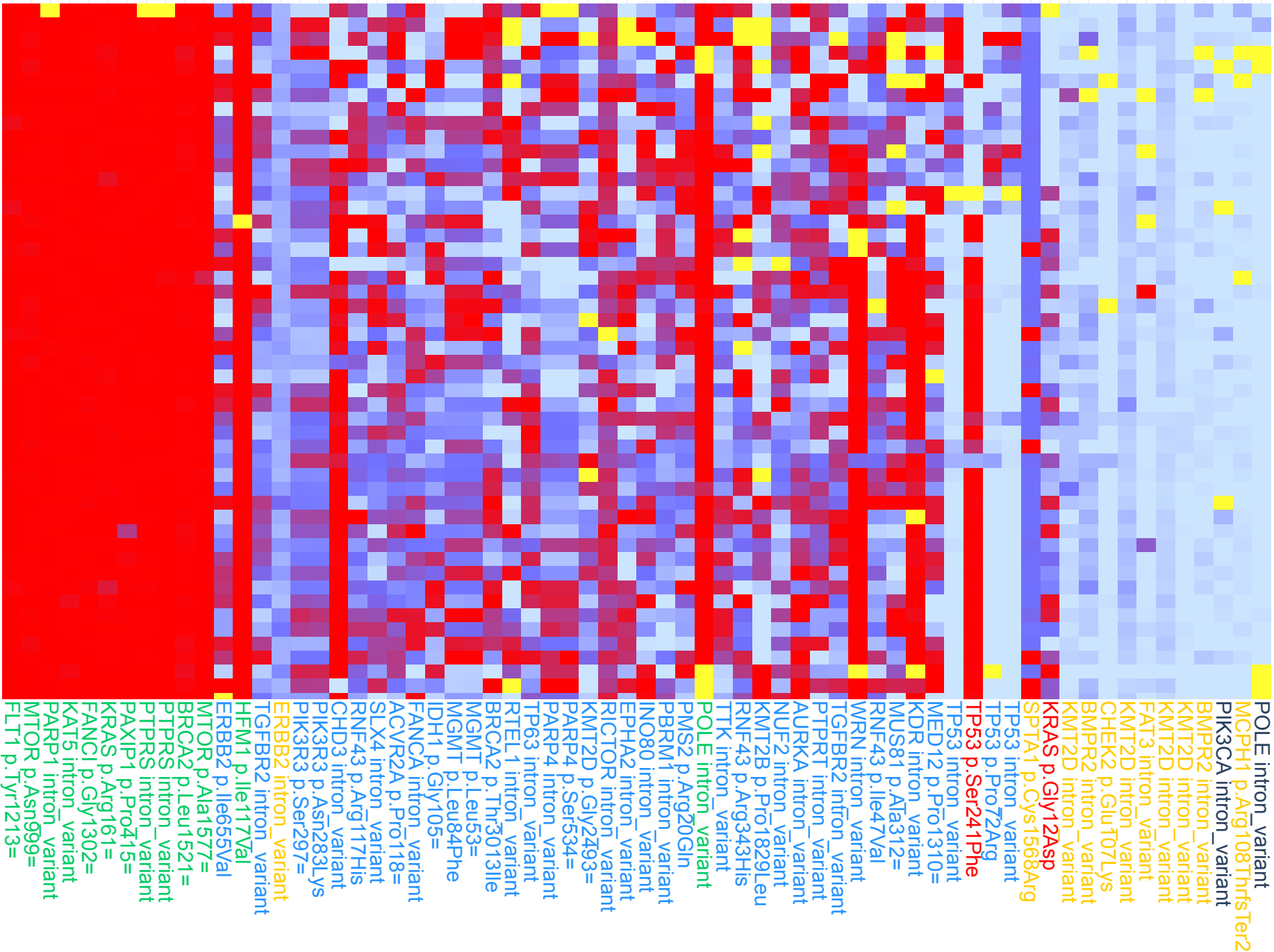


TP49-IR-0T - TP49-IR-0T-high_conf_snvs

50 cells, 66 features



POLE intron_variant
MCPH1 p.Arg1081TrfsTer2
PIK3CA intron_variant
BMPR2 intron_variant
KMT2D intron_variant
KMT2D intron_variant
FAT3 intron_variant
KMT2D intron_variant
CHEK2 p.Glu107Lys
BMPR2 intron_variant
KMT2D intron_variant
KRAS p.Gly12Asp
SPTA1 p.Cys1568Arg
TP53 intron_variant
TP53 p.Pro72Arg
TP53 p.Ser241Phe
TP53 intron_variant
MED12 p.Pro1310=
KDR intron_variant
MUS81 p.Ala312=
RNF43 p.Ile47Val
WRN intron_variant
TGFBFR2 intron_variant
PTPR1 intron_variant
AURKA intron_variant
NUF2 intron_variant
KMT2B p.Pro1829Leu
RNF43 p.Arg343His
TTK intron_variant
POLE intron_variant
PMS2 p.Arg20Gln
PBRM1 intron_variant
INO80 intron_variant
EPHA2 intron_variant
RICTOR intron_variant
KMT2D p.Gly2493=
PARP4 p.Ser534=
PARP4 intron_variant
TP63 intron_variant
RTEL1 intron_variant
BRCA2 p.Thr3013Ile
MGMT p.Leu53=
MGMT p.Leu84Phe
IDH1 p.Gly105=
FANCA intron_variant
ACVR2A p.Pro118=
SLX4 intron_variant
RNF43 p.Arg117His
CHD3 intron_variant
PIK3R3 p.Asn283Lys
PIK3R3 p.Ser297=
ERBB2 intron_variant
TGFBFR2 intron_variant
HFM1 p.Ile117Val
ERBB2 p.Ile655Val
MTOR p.Ala1577=
BRCA2 p.Leu1521=
PTPRS intron_variant
PTPRS intron_variant
PAXIP1 p.Pro415=
KRAS p.Arg161=
FANCI p.Gly1302=
KAT5 intron_variant
PARP1 intron_variant
MTOR p.Asn999=
FLT1 p.Tyr1213=

- 100.0%