RA16_29 - RA16_29-high_conf_snvs

10025 cells, 8 features

CN_clone-5 21.4%

CN_clone-4 27.8%

CN_clone-3 13.4%

CN_clone-2 8.6%

CN_clone-1 7.1%

CN_clone-0 21.6%

VAF
100
50
0
missing

CDKN2A 3_prime_UTR_variant
ARID2 p.Ser1229Ter
TP53 p.Gly245Asp
KRAS p.Gly12Val
KRAS p.Gly12Ser
CTNNB1 p.Thr41Ala
ARID2 p.Ser660Ter
STK11 intron_variant