

```

##fileformat=VCFv4.1
##fileDate=2016-05-17 22:06:44.296109
##source=GnuBio0.1
##reference=.
##run=None
##INFO=<ID=DP,Number=1,Type=Integer,Description="Total Depth">
##INFO=<ID=AF,Number=A,Type=Float,Description="Allele Frequency">
##INFO=<ID=PC,Number=1,Type=Integer,Description="Positive coverage">
##INFO=<ID=CO,Number=0,Type=String,Description="COSMIC ID">
##INFO=<ID=GN,Number=0,Type=String,Description="Gene identifier">
##INFO=<ID=EH,Number=0,Type=Integer,Description="high error band">
##INFO=<ID=EL,Number=0,Type=Integer,Description="low error band">
##FILTER=<ID=q10,Description="Quality below 10">
##FORMAT=<ID=DP,Number=1,Type=Integer,Description="Read Depth">
#CHROM POS ID REF ALT QUAL FILTER INFO FORMAT
None -135 . C T . . DP=1968.0;AF=0.395177664975;PC=777.5;CO=.;GN=ABL16;EH=[
0.41685913];EL=[ 0.37369779]

```

Research Use Only