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##fileformat=VCFv4.1
##source=COSMICv74
##reference=GRCh38
##fileDate=20150908
##comment="Missing nucleotide details indicate ambiguity during curation process"
##comment="URL stub for COSM ID field (use numeric portion of ID)='http://cancer.sanger.ac.uk/cosmic/mutati
##comment="REF and ALT sequences are both forward strand
##INFO=<ID=GENE,Number=1,Type=String,Description="Gene name">
##INFO=<ID=STRAND,Number=1,Type=String,Description="Gene strand">
##INFO=<ID=CDS,Number=1,Type=String,Description="CDS annotation">
##INFO=<ID=AA,Number=1,Type=String,Description="Peptide annotation">
##INFO=<ID=CNT,Number=1,Type=Integer,Description="How many samples have this mutation">
##INFO=<ID=SNP,Number=0,Type=Flag,Description="classified as SNP">

```

#CHROM	POS	ID	link	REF	ALT	number of times observed
19	4363794	328588	Cosmic	G	A	15
22	29668418	1415635	Cosmic	A	G	4
22	29668418	1415636	Cosmic	A	G	4
5	1294976	1717369	Cosmic	G	A	6
5	1294985	1717370	Cosmic	G	T	4
5	1294988	1717366	Cosmic	A	G	22
7	73477669	3381976	Cosmic	T	C	3

smic/mutation/overview?id=""

Origin of tumor	INFO
Kidney	GENE=SH3GL1;STRAND=-;GENE=SH3GL1;STRAND=-;CDS=c.550C>T;AA=p.R184C;CN
Large Intestine	GENE=NF2;STRAND=+;GENE=NF2;STRAND=+;CDS=c.971A>G;AA=p.Q324R;CNT=4
Large Intestine	GENE=NF2_ENST00000403999;STRAND=+;GENE=NF2_ENST00000403999;STRAND:
Urinary tract	GENE=TERT;STRAND=-;GENE=TERT;STRAND=-;CDS=c.1-45G>T;AA=p.?;CNT=6
Urinary tract	GENE=TERT;STRAND=-;GENE=TERT;STRAND=-;CDS=c.1-54C>A;AA=p.?;CNT=4
Urinary tract	GENE=TERT;STRAND=-;GENE=TERT;STRAND=-;CDS=c.1-57A>C;AA=p.?;CNT=27
Pancreas	GENE=BAZ1B;STRAND=-;GENE=BAZ1B;STRAND=-;CDS=c.1792A>G;AA=p.T598A;CNT

p.R184C;CNT=15

99;STRAND=+;CDS=c.971A>G;AA=p.Q324R;CNT=4

.T598A;CNT=3