

Table S2 Input variant format of CVE. The input file needs to be provided as a comma-separated (csv) file containing the location information chromosome, start, end, as well as the information about the reference and the observed allele in the displayed order.

chr	start	end	reference allele	observed allele
1	120437782	120437783	GG	TT
1	145273405	145273406	CG	AT
1	153963239	153963239	C	T