beta version

Under development

CHR	POS	ID	REF	ALT	QUAL	FILT	INFO	FORMAT	I1	I2
20	14373,14374	SNP1, SNP2	A,C	C,G	29,3	PASS,q-1000	SSID=-14370^20;MOD=ALT;VAR1=SNP1;VAR2=SNP2;SEQ= TTGTACGTG,ttgtaGgtg,ttgtCcgtg;SCORE=-10001,1.5277311,3.4223458	GT	0|1	0|0
...										

Description of variations of and extensions to the alpha version

Attribute	Description
POS	comma-separated list with the position(s) of each variant impacting the splice site
ID	comma-separated list with the ID(s) of each variant impacting the splice site (same ordering as in POS)
REF	comma-separated list with the reference string of each variant impacting the splice site (same ordering as in POS)
ALT	comma-separated list with the variant string of each variant in single (same ordering as in POS) A comma-separated list of the variants (and all info deferred from them) can lead to ambiguous results if one of the variants already describes multiple alternatives, e.g. ... rs6040355 A G,T ... ... microsat1 GTCT G,GTACT ... as provided as examples on the VCF definition page.
QUAL	comma-separated list of the quality for the corresponding assertions in ALT Possibly ambiguous in the case of variants with multiple alternatives, as above.
FILT	comma-separated list whether the variant position has passed the filtering As long as there is only one value per variant/SNP, and not per alternative/ALT, then there should be no problem.
INFO	MOD: either alternative (ALT) or constitutive (CON) splice site VARx: (combinations of) variants that form each alternative variant (same ordering x as in other columns POS, ALT, ...) SEQ: splice site sequence(s) for the reference and all variants applied described by the VARx attributes SCORE: comma-separated list of first the score of the reference site, and then of all variants in the usual ordering