



CORRECTION

Correction: MobiDetails: online DNA variants interpretation

David Baux  · Charles Van Goethem · Olivier Ardouin · Thomas Guignard  · Anne Bergougnoux · Michel Koenig · Anne-Françoise Roux

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Table 2 of the original article contained some mistakes. The corrected table has been updated in the original article and is also included below, with the updated values denoted with three asterisks: ***.

In addition, the “Comparison with other online platforms” paragraph has also been slightly modified in the original article according to these changes:

“However, VEP and MARRVEL2 lack several useful features such as automated literature search. MARRVEL2 and VarSome do not provide splice site predictions” has become:

“MARRVEL2 lack several useful features such as automated literature search. MARRVEL2 and VarSome do not provide extensive splice site predictions”.

Table 2 Comparison of features between MobiDetails and similar software.

	MobiDetails	VarSome	VEP web	MARRVEL
Access and technical aspects	Free usage for academics	y	y—limited	y
	Free API usage for academics	y	n	y
	Local installation	y*	n	y
	User friendly	y	y	n—but extensive options
Specific content for interpretation	Encrypted connection by default*** (https)	y	y	n***
	Responsive website	y	y—limited	n
	Custom annotations	y—registered users	y—registered users	y—only on local instances
	Selectable wild-type and mutant*** DNA sequence surrounding variants	y	n	n
	gnomADv3: whole genome population variant frequencies	y	y***	n
	Extensive*** splice site predictions	y	n	n
	miRNA target sites predictions	y	n	n
	Literature search	y—LitVar	y—MasterMind	y***
	LOVD occurrences	y	n	n
	Pdf export	y	n	n
hg38 coordinates		y	y	n
All small variants type**		y	y	n—only substitutions

*Requires the SQL scheme available upon request.

**Substitutions and small insertions/deletions.

***Updated values.