

Consequences of variation in coding regions are also reported such as synonymous, missense, frameshift, stop gained. PolyPhen2 and SIFT scores indicate likely effects of coding variation along with dbSNP clinical significance. Links to other databases such as Ensemble, UniPort and dbSNP are given, as well as PubMed IDs. Variant frequencies from populations in other projects such as 1000G and ExAC are also reported.

Final VEP output is searchable HTML table and tab delimited text