Welcome to HGVA!

We are proud to announce HGVA project. HGVA aims to index most relevant human datasets to provide a unified high-performance NoSQL database. Variant dataset have been normalised and annotated using OpenCB CellBase. HGVA provides a complete RESTful API and client libraris and command line to access all data.

HGVA is developed and maintained by researchers at University of Cambridge and Genomics England.
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