

HGVA 1.1.0 released!

- Improved web interface appearance
- Web interface includes new *beta* features:
 - Beta Genome Browser available to visualise variant context
 - Summary (*beta*) tab to get a quick, visual description of filtering result
 - Facets to get graphical descriptions of the data
- Upgraded backend. HGVA backend is now powered by OpenCGA 1.1.1

Try now!

<http://hgva.opencb.org>