Release Notes for v1.10:

1. GenomOncology annotations, Cancer Gene Census, ClinVar and Cosmic databases are updated
2. Genotypes and supporting reads for GL pertinent findings are added to CSV file on Interpretation portal
3. Link to IGV viewer for genomic coordinates for structural variants is added to WGA results HTML file on Interpretation portal
4. ClinVar star rating and pathogenicity for GL pertinent findings are added to WGA results HTML file on Interpretation portal
5. Bug with asterisk (\*) that is highlighting genes in the panel(s) for the patient's disease type for tier 3 GL pertinent findings is fixed in WGA results HTML file on Interpretation portal
6. Germline variant chr3:10046720 CTTAGTAAG>TTTAT in FANCD2 gene is blacklisted from the WGA results HTML file on Interpretation portal (arise as alignment artefact with high recurrence)