

20140501_data_release_notes

Overview of changes in the release of May 1, 2014

Number of submissions

Date	Total Submissions
Apr 05, 2013	30333
May 01, 2013	30386
Jun 01, 2013	39047
Jul 01, 2013	39170
Aug 01, 2013	45901
Sep 01, 2013	50263
Oct 01, 2013	52047
Nov 01, 2013	64750
Dec 01, 2013	64881
Jan 01, 2014	68204
Feb 01, 2014	73492
Mar 01, 2014	83343
April 01, 2014	111501
May 01, 2014	112349

Content

Brief	Explanation
Corrections made	<p>In the last release, some of the values reported in the start attribute of the SequenceLocation element of the XML export were off by 1. The number should have been incremented by 1. These data have been corrected in this release; for single nucleotide changes the value of start now equals the value of stop.</p> <p>There were also more than 100 submissions from BIC that were reported as inserts, but which have now been converted to duplications to conform to the HGVS standard. The preferred HGVS expressions were updated accordingly.</p>
Support for previous genome assemblies	ClinVar now supports variants reported on NCBI36 as well as GRCh37 and GRCh38.
Submission from the PALB2 database.	Data submitted last fall are now public in ClinVar

modified ftp directory

ClinVar made the following changes to the ftp site in April:

- new subdirectories for vcf files based on [GRCh37](#) and [GRCh38](#)
- moved *.xsd files to subdirectories specific to submission and data export
- moved the community folder to document_archives