20150903_data_release_notes

Overview of changes in the release of September 3, 2015

Note: During Sept 3-17, 2015, an incorrect version of the VCF file was posted to the ftp site. The corrected file was added on Sept 17, 2015. If you downloaded ClinVar's July VCF file before Sept 17, please download the new corrected file, which has the correct gene symbols.

Please also see our RSS feed for information about accessing the data, and our release notes for recent changes to ClinVar's web display.

Overview of submissions: 2015

Date	Total Submissions
Jan 01, 2015	149013
Feb 01, 2015	156999
Mar 1, 2015	162455
Apr 01, 2015	171408
May 1, 2015	172044
Jun 1, 2015	173236
July 1, 2015	184506
Aug 1, 2015	154686
Sep 1, 2015	158580

Content

Brief	Explanation
Coming soon - change in variant types for duplications	Coming soon - Clinvar will use the more specific variant type "tandem duplication" (SO:1000173), rather than "duplication", for duplications defined at the level of sequence. This includes short variants that are typically described as "dup" in HGVS as well as large structural variants that are known to be tandem duplications, not copy number gains.
change to variant_summary.txt	A column was added to represent the accession and version of the RefSeq sequence on which position data are based. This is especially important when the location is on an alternate locus or sub-chromosomal sequence; otherwise chromosome and location data may be misleading.
New reporting strategy for clinical significance	Apologies for the late announcement. In the 20150806 release, the reporting of clinical significance was changed. Conflicts are reported only relative to the terms describing pathogenicity. If other clinical significance values are submitted, clinical significance is represented as a list of values. For example, if a variant has been reported as pathogenic and as a risk factor, the clinical significance is reported as "Pathogenic, risk factor" rather than as a conflict.
GeneDx	GeneDx submitted interpretations for 4454 variants identified on an epilepsy panel and for variants in AHDC1, ARID2, DDX3X, POGZ, and SPATA5.

Overview of submissions: 2014

Date	Total Submissions
Jan 01, 2014	68204
Feb 01, 2014	73492
Mar 01, 2014	83343
Apr 01, 2014	111501
May 01, 2014	112349
Jun 01, 2014	117209
Jul 01, 2014	127132
Aug 01, 2014	127557
Sep 1, 2014	143114
Oct 1, 2014	143601
Nov 1, 2014	144117
Dec 1, 2014	148008

Overview of Submissions: 2013

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