

20151203_data_release_notes

Overview of changes in the release of December 3, 2015

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 01, 2015	162455
Apr 01, 2015	171408
May 01, 2015	172044
Jun 01, 2015	173236
Jul 01, 2015	184506
Aug 01, 2015	154686
Sep 01, 2015	158580
Oct 01, 2015	160538
Nov 01, 2015	170931
Dec 01, 2015	172006

Content

Brief	Explanation
Mayo Clinic submission	Mayo Clinic Genetic Testing Laboratories submitted 403 variant interpretations.
GeneDx submission	GeneDx provided updates for 12 PTEN variants, as well as interpretations for variants in CHAMP1, PPP2R5D, and several nuclear genes for mitochondrial proteins.
CHOP submission	Children's Hospital of Philadelphia provided interpretations for variants in the ABCC8 and ALK genes.
format of reporting dates	The format used to report dates in tab-delimited files is now standardized as MMM DD, YYYY. The affected files are ftp://ftp.ncbi.nlm.nih.gov/pub/clinvar/tab_delimited/summary_of_conflicting_data.txt and ftp://ftp.ncbi.nlm.nih.gov/pub/clinvar/tab_delimited/variant_summary.txt.gz
reporting of GTR test accessions as data sources for variant information	ClinVar will be phasing out reporting of GTR test accessions as sources of variant-specific information. Looking for variants in ClinVar referenced explicitly by tests in GTR can still be identified by looking for linked data, e.g. http://www.ncbi.nlm.nih.gov/clinvar/?term=clinvar_gtr[filter] , or following the link at the right side of the page from a ClinVar record to GTR.
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.

Coming soon - multiple identifiers for citations in XML	<p>Coming soon - the ClinVar XML file will allow multiple identifiers from different sources for a single citation. e.g.:</p> <pre> <Citation Type="review" Abbrev="GeneReviews"> <ID Source="BookShelf">NBK1287</ID> <ID Source="PubMed">20301464</ID> </Citation> </pre>
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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