## 2015-05-07\_Web\_Release

## Release notes, May 7, 2015

## New this month

There were no major changes this month in the display of ClinVar records. We want to announce, however, that we have started to compute connections between PubMed and ClinVar based on data mining. Every week, citations connected to the expected gene and variant are linked to ClinVar. The PubMed-gene connection is based on NCBI Gene's processing (go to http://www.ncbi.nlm.nih.gov/entrez/query/static/entrezlinks.ht ml#gene and review the gene\_pubmed section). Text is then evaluated for matches to variant descriptions in ClinVar (HGVS expression, abbreviations for alterations in protein sequence). From PubMed, the link will read **ClinVar (calculated)** (e.g. http://www.ncbi.nlm.nih.gov/pubme d?LinkName=clinvar\_pubmed\_calculated&from\_uid=182909). From ClinVar, the link will read **PubMed (calculated)**.

## **Coming soon**

In the next month, ClinVar's display will be modified to split the current Evidence tab into Summary evidence and Supporting observations. This change is being introduced to make it easier to scan an overview (Summary evidence) and look at detailed observations (Supporting observations).