








September 2021

## geneXplain® platform 6.4 release


### Database updates:

-  HumanPSD™ is updated to version 2021.2 (September 2021).
-  TRANSFAC® is updated to version 2021.2 (September 2021).
-  TRANSPATH® is updated to version 2021.2 (September 2021).

### Enhancements:

-  Detection of Genome variants and indels from RNA-seq data can now be processed for both [hg19](#) and [hg38](#) builds.
-  Transfac 'Matrix to Ensembl' matching was extended by links from matrices to orthologous and paralogous Ensembl identifiers. These links are used in hub, which converts Matrices to Ensembl genes in [Convert table](#) analysis.
-  New proteomics example with Gene Expression Omnibus data (GSE66789): Study of Myc induced regulation of protein translation. Find it at Data → Examples → Proteome profiling upon Myc activation, GSE66789.
-  Extended descriptions were added for the following examples:
  - COVID-19 suppress innate immune responses GSE156063, Illumina high throughput sequencing
  - Case study for RNA-seq data analysis
  - Chronic Myeloid Leukemia Patient Genotyping
  - Cytokine-triggered gene expression in cell cycle stages, GSE52465, Agilent-014850 microarray
  - E2F1 binding regions in HeLa cells, ChIP-Seq
  - HCV infection in liver GSE31193, Affymetrix U133 Plus 2.0 microarray
  - Proteome profiling upon Myc activation, GSE66789
  - TNF-stimulation of HUVECs GSE2639, Affymetrix HG-U133A microarray

- miRNA regulation by Myocardin and ERalpha, GSE44139, Affymetrix Multispecies miRNA-2

Check them out by clicking on the example name (in Data → Examples) and finding the respective extended descriptions at the bottom panel (opens with a click on *Toggle UI* mode button ).