



PROTEOME™ (HumanPSD™+TRANSPATH®) release 2020.3

The Human Proteome Survey Database (HumanPSD[™]) with focus on human proteins as disease biomarkers and drug targets contains these new features:

Quick access box on start page

Browse all human gene, disease, and drug entries in HumanPSD and access the detailed reports directly.

Keyword Search

UniProtKB accessions, EntrezGene IDs, and BKL primary and secondary accessions can now be used as input in a keyword search.

Disease entry for COVID-19

A dedicated entry for COVID-19 based on data from the <u>MeSH</u> 2021 preview release was included. It contains more than 1,900 clinical trials related to the disease provided by the <u>AACT</u> database and <u>clinicaltrials.gov</u>.

• More Clinical Trial and Biomarker data

New data from <u>clinicaltrials.gov</u> and manual disease biomarker curation by experts have increased the number of CT-Disease-Drug assignments to 695,136 and the number of disease annotations to 344,832.

Improved processing of clinical trials data using the <u>AACT</u> database has contributed to these elevated numbers.

The TRANSPATH® database on mammalian signal transduction and metabolic pathways contains these new data features:

• Increase in number of reactions

15,202 new binding reactions between proteins in human have been added, among them e.g. from the IFN-beta stimulated and unstimulated interactome or the beta-actin interactome.

• Interactions between SARS-CoV-2 and human proteins

For 26 SARS-CoV-2 proteins, 330 interactions with protein targets in human cells were imported (Gordon, D. E. et al., Nature 583, 459-468 (2020), Pubmed <u>32353859</u>).

• Update of links to pathway databases

Links from genes/proteins to the pathway databases <u>Reactome</u> (version 73) and <u>Wikipathways</u> (20200710) have been updated. Wikipathways links now also include zebra fish (Danio rerio) and SARS-CoV-2 related pathways.