

Resource Summary Report

Generated by ASWG on Aug 23, 2026

Pompep

RRID:SCR_010536

Type: Tool

Proper Citation

Pompep (RRID:SCR_010536)

Resource Information

URL: http://ftp://ftp.ebi.ac.uk/pub/databases/pombase/pombe/Protein_data/

Proper Citation: Pompep (RRID:SCR_010536)

Description: FTP site to access Schizosaccharomyces pombe protein data.

Abbreviations: Pompep

Synonyms: Schizosaccharomyces pombe protein data

Resource Type: data or information resource, data set

Keywords: schizosaccharomyces pombe, yeast, s. pombe

Resource Name: Pompep

Resource ID: SCR_010536

Alternate IDs: nlx_24725

Old URLs: ftp://ftp.sanger.ac.uk/pub/yeast/pombe/Protein_data/

Record Creation Time: 20220129T080259+0000

Record Last Update: 20260821T193924+0000

Ratings and Alerts

No rating or validation information has been found for Pompep.

No alerts have been found for Pompep.

Data and Source Information

Usage and Citation Metrics

We found 942 mentions in open access literature.

Listed below are recent publications. The full list is available at [ASWG](#) .

Barton RE, et al. (2022) Eco1-dependent cohesin acetylation anchors chromatin loops and cohesion to define functional meiotic chromosome domains. *eLife*, 11.

Snell CE, et al. (2014) Proline-hydroxylated hypoxia-inducible factor 1? (HIF-1?) upregulation in human tumours. *PloS one*, 9(2), e88955.

Miyagishima SY, et al. (2014) DipM is required for peptidoglycan hydrolysis during chloroplast division. *BMC plant biology*, 14, 57.

Boubaker G, et al. (2014) Echinococcus P29 antigen: molecular characterization and implication on post-surgery follow-up of CE patients infected with different species of the Echinococcus granulosus complex. *PloS one*, 9(5), e98357.

Rouhani F, et al. (2014) Genetic background drives transcriptional variation in human induced pluripotent stem cells. *PLoS genetics*, 10(6), e1004432.

Garnett MJ, et al. (2014) The evolving role of cancer cell line-based screens to define the impact of cancer genomes on drug response. *Current opinion in genetics & development*, 24(100), 114.

Köhler S, et al. (2014) The Human Phenotype Ontology project: linking molecular biology and disease through phenotype data. *Nucleic acids research*, 42(Database issue), D966.

Tenedini E, et al. (2014) Targeted cancer exome sequencing reveals recurrent mutations in myeloproliferative neoplasms. *Leukemia*, 28(5), 1052.

Garcia HA, et al. (2014) Microsatellite analysis supports clonal propagation and reduced divergence of *Trypanosoma vivax* from asymptomatic to fatally infected livestock in South America compared to West Africa. *Parasites & vectors*, 7, 210.

Lu YB, et al. (2014) Identification of boron-deficiency-responsive microRNAs in *Citrus sinensis* roots by Illumina sequencing. *BMC plant biology*, 14, 123.

Hahnel S, et al. (2014) Gonad RNA-specific qRT-PCR analyses identify genes with potential functions in schistosome reproduction such as SmFz1 and SmFGFRs. *Frontiers in genetics*, 5, 170.

Buro C, et al. (2014) Imatinib treatment causes substantial transcriptional changes in adult *Schistosoma mansoni* in vitro exhibiting pleiotropic effects. *PLoS neglected tropical diseases*, 8(6), e2923.

Kwon YJ, et al. (2014) Alternative splicing and nonsense-mediated decay of circadian clock genes under environmental stress conditions in *Arabidopsis*. *BMC plant biology*, 14,

Hiley L, et al. (2014) Distribution of Gifsy-3 and of variants of ST64B and Gifsy-1 prophages amongst *Salmonella enterica* Serovar Typhimurium isolates: evidence that combinations of prophages promote clonality. *PloS one*, 9(1), e86203.

Saad H, et al. (2014) DNA dynamics during early double-strand break processing revealed by non-intrusive imaging of living cells. *PLoS genetics*, 10(3), e1004187.

Collins MO, et al. (2014) Confident and sensitive phosphoproteomics using combinations of collision induced dissociation and electron transfer dissociation. *Journal of proteomics*, 103(100), 1.

Nguyen C, et al. (2014) Comparison of sequence variants in transcriptomic control regions across 17 mouse genomes. *Database : the journal of biological databases and curation*, 2014, bau020.

Guo S, et al. (2014) Synergistic effects of targeted PI3K signaling inhibition and chemotherapy in liposarcoma. *PloS one*, 9(4), e93996.

Goonesekere NC, et al. (2014) A meta analysis of pancreatic microarray datasets yields new targets as cancer genes and biomarkers. *PloS one*, 9(4), e93046.

Zhu Y, et al. (2014) Soybean (*Glycine max*) expansin gene superfamily origins: segmental and tandem duplication events followed by divergent selection among subfamilies. *BMC plant biology*, 14, 93.