

# Resource Summary Report

Generated by RRID on Aug 10, 2026

## PRINTS

RRID:SCR\_003412

Type: Tool

### Proper Citation

PRINTS (RRID:SCR\_003412)

### Resource Information

**URL:** <http://www.bioinf.man.ac.uk/dbbrowser/PRINTS/>

**Proper Citation:** PRINTS (RRID:SCR\_003412)

**Description:** Compendium of protein fingerprints. Diagnostic fingerprint database.

**Abbreviations:** PRINTS

**Resource Type:** database, data or information resource

**Keywords:** compedium, protein, fingerprint, diagnostic, database, FASEB list

**Availability:** THIS RESOURCE IS NO LONGER IN SERVICE

**Resource Name:** PRINTS

**Resource ID:** SCR\_003412

**Alternate IDs:** nif-0000-03338

**Alternate URLs:** <http://130.88.97.239/PRINTS/index.php>

**Record Creation Time:** 20220129T080218+0000

**Record Last Update:** 20260808T190343+0000

### Ratings and Alerts

No rating or validation information has been found for PRINTS.

No alerts have been found for PRINTS.

### Data and Source Information

## Usage and Citation Metrics

We found 427 mentions in open access literature.

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

Zhang L, et al. (2026) Quercetin upregulates steroid hormone biosynthesis to enhance reproductive performance in roosters. *Poultry science*, 105(4), 106590.

Du Z, et al. (2026) High-resolution phage-host assignment through key proteins using large language models. *Nature communications*, 17(1).

Ouyang Y, et al. (2026) Chromosome-scale Genome Assembly of the Critically Endangered Blue-crowned Laughingthrush (*Pterorhinus courtoisi*, Leiothrichidae). *Scientific data*, 13(1).

Chen J, et al. (2026) Phylogenomics-driven metalloantibiotic engineering: Overcoming ciprofloxacin resistance in *Pseudomonasaeruginosa* with sideromycin-bismuth synergy. *Cell reports. Medicine*, 7(3), 102637.

Jiang K, et al. (2026) Telomere-to-telomere-level genome assembly and annotation of the Zi goose *Anser cygnoides*. *Scientific data*, 13(1).

Liu T, et al. (2026) Chromosome-level genome assembly and annotation of the endangered plant *Primula kwangtungensis* (Primulaceae). *Scientific data*, 13(1), 212.

Pan Y, et al. (2026) Proteomic Analysis of Lotus-Derived NnAP2 Regulation of Soluble Sugar and Starch Content in Potato (*Solanum tuberosum*). *Plants (Basel, Switzerland)*, 15(4).

K??stlbacher S, et al. (2026) Prediction of eukaryotic cellular complexity in Asgard archaea using structural modelling. *Nature microbiology*, 11(3), 747.

Soman SP, et al. (2026) IsoProDB: an integrated map of human protein isoforms for accelerated research. *Database : the journal of biological databases and curation*, 2026.

Lu D, et al. (2026) An annotated genome of freshwater amphipod (*Gammarus nekkensis*). *Scientific data*, 13(1).

Yu ZX, et al. (2026) Chromosome-level genome of *Mosla chinensis* from alpine ecotype provides insights into terpenoid biosynthesis and germplasm exploration. *Frontiers in plant science*, 17, 1819008.

Zou R, et al. (2026) A chromosome-level genome assembly of *Cistanche deserticola* provides insights into its evolution and molecular mechanisms of parasitism. *Plant communications*, 7(1), 101581.

Huang Z, et al. (2026) Integrated Proteomics and Metabolomics Analysis Reveals Potential Pathways Underlying Onion-Mediated Regulation of Spleen Immune Function in

Liangshan Black Sheep. *Veterinary sciences*, 13(5).

Liber JA, et al. (2026) *Aimea* gen. nov. defines a novel plant-associated yeast genus in Microbotryomycetes with three novel species. *bioRxiv : the preprint server for biology*.

Yin S, et al. (2026) A telomere-to-telomere genome assembly of *Castanopsis orthacantha* (Fagaceae). *Scientific data*, 13(1).

Nneji LM, et al. (2026) First Chromosome-level Genome Assembly and Annotation of an Endangered Freshwater Stingray (*Fontitrygon garouaensis*) from Africa. *Scientific data*, 13(1).

De Panis D, et al. (2026) The reference genome of the Asian Elephant (*Elephas maximus*): a foundation for conservation and genomic research. *BMC genomics*, 27(1).

Stilwell PA, et al. (2026) The genome of *Istocheta aldrichi* (Diptera: Tachinidae), a parasitoid of the Japanese beetle, *Popillia japonica* (Coleoptera: Scarabaeidae). *G3* (Bethesda, Md.), 16(4).

Sedaghatkish A, et al. (2026) A contamination-reduced genome of *Plasmodiophora brassicae* with comprehensive functional annotation. *BMC microbiology*, 26(1), 118.

Dos Santos Menezes F, et al. (2026) Comparative analysis of PRR receptors and R genes reveals structural and functional divergence between wild, ancestral, and commercial citrus species. *BMC genomics*, 27(1), 173.