

Resource Summary Report

Generated by PRECISE-TBI on Jul 31, 2026

FALCON

RRID:SCR_018804

Type: Tool

Proper Citation

FALCON (RRID:SCR_018804)

Resource Information

URL: <http://protein.ict.ac.cn/FALCON/>

Proper Citation: FALCON (RRID:SCR_018804)

Description: Web tool as high throughput protein structure prediction service. High throughput server for protein structure prediction.

Synonyms: FALCON@home

Resource Type: service resource, data access protocol, software resource, web service, production service resource, analysis service resource

Keywords: Protein structure, protein structure prediction, simulation, 3D structure, protein, high throughput protein, structure prediction

Availability: Restricted

Resource Name: FALCON

Resource ID: SCR_018804

Record Creation Time: 20220129T080342+0000

Record Last Update: 20260731T043039+0000

Ratings and Alerts

No rating or validation information has been found for FALCON.

No alerts have been found for FALCON.

Data and Source Information

Usage and Citation Metrics

We found 428 mentions in open access literature.

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

Bernstein HR, et al. (2026) Periventricular gradients in axonal and myelin microstructure are present in people with multiple sclerosis having low and high disability. Multiple sclerosis (Houndmills, Basingstoke, England), 32(1), 80.

Weissensteiner MH, et al. (2025) Combining Individual-Based Radio-Tracking With Whole-Genome Sequencing Data Reveals Candidate for Genetic Basis of Partial Migration in a Songbird. Ecology and evolution, 15(1), e70800.

Morisaki S, et al. (2025) Dendritic cell-derived exosomes induce monocyte antigen-presentation and immune amplification in neoantigen vaccine therapy. Frontiers in immunology, 16, 1565696.

Akter Y, et al. (2025) Combining nanobody labeling with STED microscopy reveals input-specific and layer-specific organization of neocortical synapses. PLoS biology, 23(4), e3002649.

Ye Y, et al. (2025) Intermediate filament IFFO1 negatively regulates the migration of lung cancer cells by inhibiting the IQGAP3-Cdc42 interaction. Cell death & disease, 16(1), 509.

Seigneuret F, et al. (2025) The HBV large envelope protein initiates virion assembly by recruiting capsids at membrane rich domains related to late endosome. Cellular and molecular life sciences : CMLS, 82(1), 128.

Tu M, et al. (2025) Integrative omics reveals mechanisms of biosynthesis and regulation of floral scent in Cymbidium tracyanum. Plant biotechnology journal, 23(6), 2162.

Pinheiro-Machado E, et al. (2025) Secretome Analysis of Human and Rat Pancreatic Islets Co-Cultured with Adipose-Derived Stromal Cells Reveals a Signature with Enhanced Regenerative Capacities. Cells, 14(4).

Lu D, et al. (2025) Multimeric protein interaction and complex prediction: Structure, dynamics and function. Computational and structural biotechnology journal, 27, 1975.

Baumeister T, et al. (2025) Loss of FXR or Bile Acid-dependent Inhibition Accelerate Carcinogenesis of Gastroesophageal Adenocarcinoma. Cellular and molecular gastroenterology and hepatology, 19(8), 101505.

Safronov O, et al. (2025) Independent evolution of betulin biosynthesis in Inonotus obliquus. Scientific reports, 15(1), 21319.

Zhang L, et al. (2025) Chronic Stress-Induced and Tumor Derived SP1+ Exosomes Polarizing IL-1?+ Neutrophils to Increase Lung Metastasis of Breast Cancer. Advanced science (Weinheim, Baden-Wurttemberg, Germany), 12(4), e2310266.

Lee J, et al. (2025) A chromosome-level genome assembly of wild silkworm, *Bombyx mandarina*. *Scientific data*, 12(1), 27.

Tu XD, et al. (2025) Genomic insights into *Castanopsis carlesii* and *Castanea henryi*: flower and fruit development and evolution of NLR genes in the beech-oak family. *Molecular horticulture*, 5(1), 33.

Svobodová B, et al. (2025) Revised Diffusion Law Permits Quantitative Nanoscale Characterization of Membrane Organization. *Analytical chemistry*, 97(22), 11478.

Tsujimoto M, et al. (2025) Differentiation and correlation of regional uptake heterogeneity with cardiac dysfunction in biopsy-proven transthyretin amyloid cardiomyopathy using quantitative single-photon emission computed tomography/computed tomography: a single-center, cross-sectional study. *EJNMMI research*, 15(1), 97.

Woo SJ, et al. (2025) Isolation, Identification, and Characteristics of *Aeromonas salmonicida* subsp. *masoucida* from Diseased Starry Flounder (*Platichthys stellatus*). *Pathogens (Basel, Switzerland)*, 14(3).

Gu C, et al. (2025) Long-read genome sequencing reveals the sequence characteristics of pear self-incompatibility locus. *Molecular horticulture*, 5(1), 13.

Xu QZ, et al. (2025) Chromosome-level genome assembly and single-cell analysis unveil molecular mechanisms of arm regeneration in the ophiuroid *Ophiura sarsii vadicola*. *Genome biology*, 26(1), 82.

Zhu M, et al. (2025) Transcriptomic and spatial GABAergic neuron subtypes in zona incerta mediate distinct innate behaviors. *Nature communications*, 16(1), 3107.