

Resource Summary Report

Generated by [dkNET](#) on Sep 12, 2026

FGENESH Plus

RRID:SCR_018937

Type: Tool

Proper Citation

FGENESH Plus (RRID:SCR_018937)

Resource Information

URL:

http://www.softberry.com/berry.phtml?topic=fgenes_plus&group=programs&subgroup=gfs

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Description: Web tool as HMM plus similar protein based gene prediction. Used for multiple gene prediction in genomic DNA with using information from similar protein. Used if you know protein sequence similar with protein which is encoded by gene in your sequence.

Synonyms: Fgenesh plus

Resource Type: data access protocol, simulation software, software application, software resource, web service

Keywords: HMM, gene prediction, protein, multiple gene prediction, genomic DNA, similar protein, protein sequence, Softberry

Availability: Free, Freely available

Resource Name: FGENESH Plus

Resource ID: SCR_018937

License URLs: <http://www.softberry.com/berry.phtml?topic=termsfuse>

Record Creation Time: 20220129T080342+0000

Record Last Update: 20260905T132844+0000

Ratings and Alerts

No rating or validation information has been found for FGENESH Plus.

No alerts have been found for FGENESH Plus.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 6 mentions in open access literature.

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

Overlander-Chen M, et al. (2025) Genetic analysis of the single internode dwarf 1 mutant in barley. BMC plant biology, 25(1), 797.

Gain N, et al. (2025) Molecular delineation and haplotype analysis of domain membrane protein (DMP) gene influencing in-vivo haploid induction in maize. Scientific reports, 15(1), 7697.

Kang Z, et al. (2025) Genomic and Multi-Omics Analysis of Phlebopus portentosus: Effects of Cultivation on Secondary Metabolites. Journal of fungi (Basel, Switzerland), 11(4).

Zhuang X, et al. (2025) Identification, Expression Profiling, Microbial Binding, and Agglutination Analyses of Two Cathepsin B Genes in Black Rockfish (Sebastes schlegelii). Marine drugs, 23(5).

Hewitt TC, et al. (2025) Divergent molecular pathways govern temperature-dependent wheat stem rust resistance genes. Nature communications, 16(1), 4905.

Cheng G, et al. (2025) Genomic Insights into Cobweb Disease Resistance in Agaricus bisporus: A Comparative Analysis of Resistant and Susceptible Strains. Journal of fungi (Basel, Switzerland), 11(3).