

Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

Fungi Sequencing Projects

RRID:SCR_008524

Type: Tool

Proper Citation

Fungi Sequencing Projects (RRID:SCR_008524)

Resource Information

URL: <http://www.sanger.ac.uk/Projects/Fungi/>

Proper Citation: Fungi Sequencing Projects (RRID:SCR_008524)

Description: Fungal genomes available from the Sanger Institute. Data are accessible in a number of ways; for each organism there is a BLAST server, allowing search of the sequences. Sequences can also be down-loaded directly by FTP. In addition, for those organisms being sequenced using a cosmid approach, finished and annotated cosmids are submitted to EMBL and other public databases.

Synonyms: Fungi Sequencing

Resource Type: data or information resource, data set

Keywords: genome, genomics, sequence, fungus, blast, data analysis service

Resource Name: Fungi Sequencing Projects

Resource ID: SCR_008524

Alternate IDs: nif-0000-30593

Record Creation Time: 20220129T080247+0000

Record Last Update: 20260905T133329+0000

Ratings and Alerts

No rating or validation information has been found for Fungi Sequencing Projects.

No alerts have been found for Fungi Sequencing Projects.

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 [Kravitz](#) .

Martin R, et al. (2011) The Candida albicans-specific gene EED1 encodes a key regulator of hyphal extension. PloS one, 6(4), e18394.

Lavoie H, et al. (2010) Evolutionary tinkering with conserved components of a transcriptional regulatory network. PLoS biology, 8(3), e1000329.

González-Fernández R, et al. (2010) Proteomics of plant pathogenic fungi. Journal of biomedicine & biotechnology, 2010, 932527.

Machida M, et al. (2008) Genomics of Aspergillus oryzae: learning from the history of Koji mold and exploration of its future. DNA research : an international journal for rapid publication of reports on genes and genomes, 15(4), 173.

Hogues H, et al. (2008) Transcription factor substitution during the evolution of fungal ribosome regulation. Molecular cell, 29(5), 552.

Gupta R, et al. (2007) Ubiquitination screen using protein microarrays for comprehensive identification of Rsp5 substrates in yeast. Molecular systems biology, 3, 116.