

Resource Summary Report

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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 [RRID](#) .

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.