

Resource Summary Report

Generated by [RRID](#) on Aug 10, 2026

Rice Pipeline

RRID:SCR_013344

Type: Tool

Proper Citation

Rice Pipeline (RRID:SCR_013344)

Resource Information

URL: <http://cdna01.dna.affrc.go.jp/PIPE>

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Description: A unification tool which dynamically collects and compiles data from scientific databases in National Institute of Agrobiological Sciences (NIAS), and thereby attempts to encapsulate the genetics and molecular biology of genes from the genomes of *Oryza sativa* into easy to navigate. The mission of Rice PIPELINE is to provide a unique scientific resource of rice that pools publicly available data commonly sought after for any clone sequence, clone name, GenBank accession number, or keyword.

Synonyms: Rice Pipeline

Resource Type: database, data or information resource

Resource Name: Rice Pipeline

Resource ID: SCR_013344

Alternate IDs: nif-0000-03408

Record Creation Time: 20220129T080315+0000

Record Last Update: 20260808T190433+0000

Ratings and Alerts

No rating or validation information has been found for Rice Pipeline.

No alerts have been found for Rice Pipeline.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Sumiyoshi M, et al. (2013) Increase in cellulose accumulation and improvement of saccharification by overexpression of arabinofuranosidase in rice. PloS one, 8(11), e78269.

Ricachenevsky FK, et al. (2011) ZINC-INDUCED FACILITATOR-LIKE family in plants: lineage-specific expansion in monocotyledons and conserved genomic and expression features among rice (*Oryza sativa*) paralogs. BMC plant biology, 11, 20.

Jo Y, et al. (2011) Plasmodesmal receptor-like kinases identified through analysis of rice cell wall extracted proteins. Protoplasma, 248(1), 191.

Galperin MY, et al. (2005) The Molecular Biology Database Collection: 2005 update. Nucleic acids research, 33(Database issue), D5.