

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

FuncExpression

RRID:SCR_005773

Type: Tool

Proper Citation

FuncExpression (RRID:SCR_005773)

Resource Information

URL: <http://www.plexdb.org/plex.php?database=Barley/funcexpression.php>

Proper Citation: FuncExpression (RRID:SCR_005773)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented on July 11, 2012. FuncExpression is a web-based resource for functional interpretation of large scale genomics data. FuncExpression can be used for the functional comparison of plant, animal, and fungal gene name lists generated from genomics and proteomics experiments. Multiple gene lists can be classified, compared and visualized. FuncExpression supports two way-integration of plant gene functional information and the gene expression data, which allows for further cross-validation with plant microarray data from related experiments at BarleyBase. Platform: Online tool

Abbreviations: FuncExpression

Resource Type: analysis service resource, service resource, data analysis service, production service resource

Keywords: statistical analysis, genomics, compare, function, plant, animal, fungus, gene, proteomics, gene expression, microarray

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: FuncExpression

Resource ID: SCR_005773

Alternate IDs: nlx_149237

Record Creation Time: 20220129T080232+0000

Record Last Update: 20260728T043222+0000

Ratings and Alerts

No rating or validation information has been found for FuncExpression.

No alerts have been found for FuncExpression.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.