Resource Summary Report

Generated by NIF on May 20, 2025

5 prime end Serial Analysis of Gene Expression Database

RRID:SCR_001680
Type: Tool

Proper Citation

5 prime end Serial Analysis of Gene Expression Database (RRID:SCR_001680)

Resource Information

URL: http://5sage.gi.k.u-tokyo.ac.jp/

Proper Citation: 5 prime end Serial Analysis of Gene Expression Database (RRID:SCR_001680)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented on October 30, 2012. A database that displays the observed frequencies of individual 5' end SAGE tags and previously unknown transcription start sites in the promoter regions, introns and intergenic regions of known genes. 5'SAGE will be useful for analyzing promoter regions and start site variation in different tissues, and is freely available.

Abbreviations: 5'SAGE, 5SAGE

Synonyms: 5'end Serial Analysis of Gene Expression Database, 5end Serial Analysis of Gene Expression Database

Resource Type: database, data or information resource

Defining Citation: PMID:15608259

Keywords: gene expression, dna, rna, serial analysis, transcription start site, promoter region, intron, intergenic region, gene

Funding: Japanese Ministry of Education Culture Sports Science and Technology MEXT

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: 5 prime end Serial Analysis of Gene Expression Database

Resource ID: SCR_001680

Alternate IDs: nif-0000-02525

Record Creation Time: 20220129T080208+0000

Record Last Update: 20250519T204622+0000

Ratings and Alerts

No rating or validation information has been found for 5 prime end Serial Analysis of Gene Expression Database.

No alerts have been found for 5 prime end Serial Analysis of Gene Expression Database.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We have not found any literature mentions for this resource.