

Resource Summary Report

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

ARNIE

RRID:SCR_000514

Type: Tool

Proper Citation

ARNIE (RRID:SCR_000514)

Resource Information

URL: <http://www.sanger.ac.uk/cgi-bin/teams/team30/arnie>

Proper Citation: ARNIE (RRID:SCR_000514)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 1, 2023. Database that integrates the extracellular protein interaction network generated in our lab using AVEXIS technology with spatiotemporal expression patterns for all genes in the network. The tool allows users to browse the network by clicking on individual proteins, or by specifying the spatiotemporal parameters. Clicking on connector lines will allow users to compare stage-matched expression patterns for genes encoding interacting proteins. Additionally, users can rapidly search for their genes in the network using the BLAST server provided.

Abbreviations: ARNIE

Synonyms: AVEXIS Receptor Network with Integrated Expression

Resource Type: database, data or information resource

Defining Citation: [PMID:20802085](#)

Keywords: network, orthologue, paralogue, gene, orthologue, protein interaction, protein, blast, extracellular, expression profiling, interaction network, ligand, interaction, signaling

Funding: MRC ;
Wellcome Trust

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: ARNIE

Resource ID: SCR_000514

Alternate IDs: OMICS_01529

Record Creation Time: 20220129T080202+0000

Record Last Update: 20260725T191115+0000

Ratings and Alerts

No rating or validation information has been found for ARNIE.

No alerts have been found for ARNIE.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.