

# Welcome to eagle-i



## Unsupported Software

The eagle-i software is still available to the open-source community, but is no longer supported or maintained.

## Making open science happen...

eagle-i is an ontology-driven, RDF-based software platform for creating, storing and searching semantically rich data about research resources of all kinds. eagle-i is built around semantic web technologies and adheres to [linked open data](#) principles.

The collage displays various features of the eagle-i platform:

- Search Interface:** A search bar with the text "Your gene's disease" and a "Try our new IPS Cell Search" button. Navigation links include ABOUT, GET INVOLVED, NEWS + EVENTS, FAQ, CONTACT US, and HELP.
- Search Results:** A page titled "Narrow Your Results" showing a list of resources with filters for "Collection Filter" and "Resource Type Filter". The results are categorized by "Core Facilities", "IPS Cells", "Micro", "Monoclonal Antibodies", "Biological Specimens", "Human Study", "Request", "Research Opportunity", and "Software".
- Resource Detail:** A detailed view of a resource titled "296-ALS (SOD1A144F)" with fields for "Request", "Description", "Location", "Source", "Status", and "Contact".
- SWEET Interface:** A "Semantic Web Entry & Editing Tool" for editing a resource, with fields for "Name", "Description", "Status", and "Contact".
- Search Filters:** A sidebar with filters for "Diagnosed Disease", "Subject Age at Diagnosis", "Genetic Alterations", "Ethnicity", "Sex", "Part of Collection", "Induction Method", and "Type of QC Performed".
- Network-wide Cell Line Stats:** A section titled "eagle-i.net network-wide cell line stats" showing a bar chart of "Total Number Subjects" and "Total Number Cell Lines".