

# Resource Summary Report

Generated by [NIF](#) on Aug 20, 2026

## Eagle I

RRID:SCR\_013153

Type: Tool

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### Proper Citation

Eagle I (RRID:SCR\_013153)

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### Resource Information

**URL:** <https://www.eagle-i.net/>

**Proper Citation:** Eagle I (RRID:SCR\_013153)

**Description:** Web application to discover resources available at participating networked universities. This distributed platform for creating and sharing semantically rich data is built around semantic web technologies and follows linked open data principles.

**Abbreviations:** eagle-i, eagle i, eaglei

**Resource Type:** data or information resource, database

**Defining Citation:** [PMID:22434835](#)

**Keywords:** ontology, semantic web, rdf, sparql endpoint, linked open data, distributed platform, protocol

**Funding:** NCRR U24 RR029825;  
ARRA

**Availability:** Available to external user, The community can contribute to this resource

**Resource Name:** Eagle I

**Resource ID:** SCR\_013153

**Alternate IDs:** r3d100011564, nlx\_143592

**Alternate URLs:** <https://www.eagle-i.org/>, <https://www.force11.org/node/4661>

**License:** BSD License v3

**Record Creation Time:** 20220129T080314+0000

**Record Last Update:** 20260820T054736+0000

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## Ratings and Alerts

No rating or validation information has been found for Eagle I.

No alerts have been found for Eagle I.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 10 mentions in open access literature.

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

Chen Y, et al. (2021) Integrated Collection of Stem Cell Bank Data, a Data Portal for Standardized Stem Cell Information. Stem cell reports, 16(4), 997.

He Y, et al. (2019) OSCI: standardized stem cell ontology representation and use cases for stem cell investigation. BMC bioinformatics, 20(Suppl 5), 180.

Fujimori K, et al. (2016) Modeling neurological diseases with induced pluripotent cells reprogrammed from immortalized lymphoblastoid cell lines. Molecular brain, 9(1), 88.

Thessen AE, et al. (2015) Emerging semantics to link phenotype and environment. PeerJ, 3, e1470.

Walls RL, et al. (2014) Semantics in support of biodiversity knowledge discovery: an introduction to the biological collections ontology and related ontologies. PloS one, 9(3), e89606.

Tsiliki G, et al. (2014) Collaborative mining and interpretation of large-scale data for biomedical research insights. PloS one, 9(9), e108600.

Soares FA, et al. (2014) International coordination of large-scale human induced pluripotent stem cell initiatives: Wellcome Trust and ISSCR workshops white paper. Stem cell reports, 3(6), 931.

Mabile L, et al. (2013) Quantifying the use of bioresources for promoting their sharing in scientific research. GigaScience, 2(1), 7.

Vasilevsky N, et al. (2012) Research resources: curating the new eagle-i discovery system. Database : the journal of biological databases and curation, 2012, bar067.

Bandrowski AE, et al. (2012) A hybrid human and machine resource curation pipeline for the Neuroscience Information Framework. Database : the journal of biological databases and curation, 2012, bas005.