

Resource Summary Report

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Integrated Manually Extracted Annotation

RRID:SCR_008876

Type: Tool

Proper Citation

Integrated Manually Extracted Annotation (RRID:SCR_008876)

Resource Information

URL: https://neuinfo.org/mynif/search.php?q=*&t=indexable&list=cover&nif=nlx_154697-2

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Description: A virtual database of annotations made by 50 database providers (April 2014) - and growing (see below), that map data to publication information. All NIF Data Federation sources can be part of this virtual database as long as they indicate the publications that correspond to data records. The format that NIF accepts is the PubMed Identifier, category or type of data that is being linked to, and a data record identifier. A subset of this data is passed to NCBI, as LinkOuts (links at the bottom of PubMed abstracts), however due to NCBI policies the full data records are not currently associated with PubMed records. Database providers can use this mechanism to link to other NCBI databases including gene and protein, however these are not included in the current data set at this time. (To view databases available for linking see, http://www.ncbi.nlm.nih.gov/books/NBK3807/#files.Databases_Available_for_Linking) The categories that NIF uses have been standardized to the following types: * Resource: Registry * Resource: Software * Reagent: Plasmid * Reagent: Antibodies * Data: Clinical Trials * Data: Gene Expression * Data: Drugs * Data: Taxonomy * Data: Images * Data: Animal Model * Data: Microarray * Data: Brain connectivity * Data: Volumetric observation * Data: Value observation * Data: Activation Foci * Data: Neuronal properties * Data: Neuronal reconstruction * Data: Chemosensory receptor * Data: Electrophysiology * Data: Computational model * Data: Brain anatomy * Data: Gene annotation * Data: Disease annotation * Data: Cell Model * Data: Chemical * Data: Pathways For more information refer to Create a LinkOut file, http://neuinfo.org/nif_components/disco/interoperation.shtml Participating resources (<http://disco.neuinfo.org/webportal/discoLinkoutServiceSummary.do?id=4>): * Addgene <http://www.addgene.org/pgvec1> * Animal Imaging Database <http://aidb.crbs.ucsd.edu> * Antibody Registry <http://www.neuinfo.org/products/antibodyregistry/> * Avian Brain Circuitry Database <http://www.behav.org/abcd/abcd.php> * BAMS Connectivity <http://brancusi.usc.edu/> * Beta Cell Biology Consortium <http://www.betacell.org/> * bioDBcore <http://biodbcore.org/> * BioGRID <http://thebiogrid.org/> * BioNumbers <http://bionumbers.hms.harvard.edu/> * Brain Architecture Management System

<http://brancusi.usc.edu/bkms/> * Brede Database
<http://hendrix.imm.dtu.dk/services/jerne/brede/> * Cell Centered Database
<http://ccdb.ucsd.edu> * CellML Model Repository <http://www.cellml.org/models> * CHEBI
<http://www.ebi.ac.uk/chebi/> * Clinical Trials Network (CTN) Data Share
<http://www.ctndatashare.org/> * Comparative Toxicogenomics Database <http://ctdbase.org/>
* Coriell Cell Repositories <http://ccr.coriell.org/> * CRCNS - Collaborative Research in
Computational Neuroscience - Data sharing <http://crcns.org> * Drug Related Gene
Database <https://confluence.crbs.ucsd.edu/display/NIF/DRG> * DrugBank
<http://www.drugbank.ca/> * FLYBASE <http://flybase.org/> * Gene Expression Omnibus
<http://www.ncbi.nlm.nih.gov/geo/> * Gene Ontology Tools
<http://www.geneontology.org/GO.tools.shtml> * Gene Weaver <http://www.GeneWeaver.org>
* GeneDB <http://www.genedb.org/Homepage> * Glomerular Activity Response Archive
<http://gara.bio.uci.edu> * GO <http://www.geneontology.org/> * Internet Brain Volume
Database <http://www.cma.mgh.harvard.edu/ibvd/> * ModelDB
<http://senselab.med.yale.edu/modeldb/> * Mouse Genome Informatics Transgenes
ftp://ftp.informatics.jax.org/pub/reports/MGI_PhenotypicAllele.rpt * NCBI Taxonomy
Browser <http://www.ncbi.nlm.nih.gov/Taxonomy/taxonomyhome.html> * NeuroMorpho.Org
<http://neuromorpho.org/neuroMorpho> * NeuronDB <http://senselab.med.yale.edu/neurondb>
* SciCrunch Registry <http://neuinfo.org/nif/nifgwt.html?tab=registry> * NIF Registry
Automated Crawl Data http://lucene1.neuinfo.org/nif_resource/current/ * NITRC
<http://www.nitrc.org/> * Nuclear Receptor Signaling Atlas <http://www.nursa.org> * Olfactory
Receptor DataBase <http://senselab.med.yale.edu/ordb/> * OMIM <http://omim.org> *
OpenfMRI <http://openfmri.org> * PeptideAtlas <http://www.peptideatlas.org> * RGD
<http://rgd.mcw.edu> * SFARI Gene: AutDB <https://gene.sfari.org/autdb/Welcome.do> *
SumsDB <http://sumsdb.wustl.edu/sums/> * Temporal-Lobe: Hippocampal -
Parahippocampal Neuroanatomy of the Rat <http://www.temporal-lobe.com/> * The Cell: An
Image Library <http://www.cellimagelibrary.org/> * Visiome Platform
<http://platform.visiome.neuroinf.jp/> * WormBase <http://www.wormbase.org> * YPED
<http://medicine.yale.edu/keck/nida/yped.aspx> * ZFIN <http://zfin.org>

Abbreviations: Integrated MEA, Integrated LinkOut, Manually Extracted Annotation,

Synonyms: Integrated Manually Extracted Annotation View, NIF Integrated LinkOut, NIF
Integrated Manually Extracted Annotation, NIF LinkOut

Resource Type: database, data or information resource

Defining Citation: [PMID:20387131](#), [PMID:18975149](#)

Keywords: biomedical, bibliographic, linkout, literature

Availability: Data are licensed by their respective owners. Use and distribution is subject
to the Terms of Use by the original resource as well as the, Creative Commons Attribution
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Resource Name: Integrated Manually Extracted Annotation

Resource ID: SCR_008876

Alternate IDs: nlx_149407

Old URLs: http://neuinfo.org/nif/nifgwt.html?query=nlx_149407,
https://neuinfo.org/mynif/search.php?q=*&t=indexable&list=cover&nif=nlx_149407-1

Record Creation Time: 20220129T080249+0000

Record Last Update: 20260803T040531+0000

Ratings and Alerts

No rating or validation information has been found for Integrated Manually Extracted Annotation.

No alerts have been found for Integrated Manually Extracted Annotation.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.