Preclinical Tool Registration Guide

Step-by-Step Instructions for Registering Preclinical Research Tools

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Preclinical Research Tools

This document provides a step-by-step guide to registering preclinical research tools.

The ASAP Open Science Policy requires researchers to (1) cite all newly-generated lab resources (antibodies, cell lines, plasmids/clones, transgenic models, and other reagents) and (2) to deposit those resources in a publicly accessible repository.
For additional help and questions, email **openscience@parkinsonsroadmap.org**.

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What is an RRID?

Research Resource Identifiers (RRIDs) are persistent and unique IDs that allow researchers to reference the tools and resources that were utilized to support the research efforts described in preprints and final publications.

Registering lab resources for an RRID is a free process that provides several benefits:

- **Centralized system:** RRIDs are a way to centralize the resource database for materials being used in biomedical research

- **Impact assessment:** RRIDs are machine-readable, so the resource can be easily identified in any text

- **Consistent:** RRIDs are consistent across both publishers and vendors

- **Resolve URLs and catalog numbers:** Resource catalog numbers may change. The RRID is constant.

- **Identification of problematic tools:** Whenever there is a problematic tool (e.g., contaminated cell line), it is flagged in the RRID database

Antibodies

1. To register an antibody, visit https://scicrunch.org/resources/about/resource

2. Select Antibody

3. Login to the Antibody Registry with your ORCID or email and password. If you are a new user, create an account.

4. Select 'Personal Antibody'

5. Add Antibody details and select 'Submit'

2. Antibody Details

Identifier (Mandatory)

An catalogNumber unique to your antibody (e.g. Labname_001 or myab_1023)

Note: Submit unregistered antibodies only

Principal Investigator - Institution or Vendor (Mandatory)

(e.g. J. Doe - Harvard)

Principal Investigator's/Institution's Website or Vendor Website (Mandatory)

http:// or https://

Antibody name (Mandatory)

Anti-phospho-Glo1(Y136) Antibody

Host Species (Mandatory)

Rabbit

Target/Reactive Species (Mandatory)

Mouse

Antibody Target (Mandatory)

phospho-Glo1(Y136)

Clonality (Mandatory)

Unknown ▼

Clone ID

C200234

Isotype

IgG

Step 2/2:Antibody Details

‹ Previous   ✓ Submit

Example of successful antibody registration

🔍 RRID Portal

ABOUT ˅

Resource Summary Report   🔍 New Search   ↺ Previous Search Results

Home / Resource Reports / Antibodies / Resource Summary Report

🧪 Antibody Name ❓   *NOTICE: Multiple vendors found, please select your record: Novus - NB120-15534 ˅

alpha Synuclein Antibody 🔗 ☐

RRID:AB_792157 📋

PDF REPORT   HOW TO CITE

ℹ️ Antibody Information ❓

URL: http://antibodyregistry.org/AB_792157

Proper Citation: (Novus Cat# NB120-15534, RRID:AB_792157)

Target Antigen: alpha Synuclein

Host Organism: rabbit

Clonality: polyclonal

Comments: Immunohistochemistry-Paraffin

Expand All

📈 Usage and Citation Metrics ❓⚠️

We have not found any literature mentions for this resource.

Check Google Scholar for all resource mentions.

👥 Collaborator Network ❓

A list of researchers who have used the resource and an author search tool. This is available for resources that have literature mentions.

💬 Ratings and Alerts ❓   Report Information

No rating or validation information has been found for alpha Synuclein Antibody.

No alerts have been found for alpha Synuclein Antibody.

🗄️ Data and Source Information ❓

Source: Antibody Registry

Cell Lines

1. Confirm novelty by searching for similar registered cell lines at scicrunch.org/resources and www.cellosaurus.org

2. Register cell lines through the Cellosaurus Home Page by selecting "Contact" on the top panel menu.
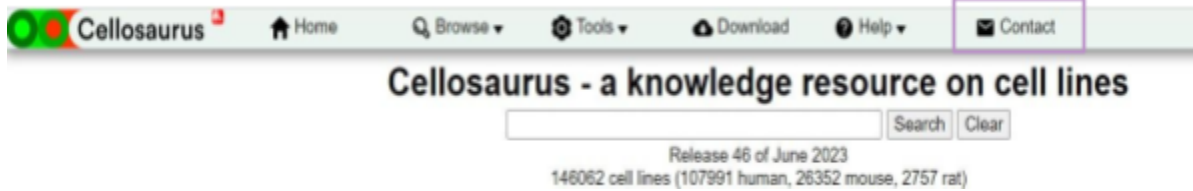
3. Fill in the information and click Submit. The information that should be included in the request varies depending on if the cell lines are described in a publication.

Cell Lines Described in a Publication
    Subject: Indicate that this cell line is described in a publication

    Message: Provide a link to the publication where your cell line was first described

Cell Lines are NOT Described in a Publication
    Subject: Indicate that this is a request to create a new Cellosaurus entry

    Message: Provide the following information
- Name - Species of origin (if not human, include strain/breed)
- Gender and age of donor
- Category of cell line (e.g., cancer, hybridoma, iPSC, ESC, etc.)
- If the donor is suffering from a disease, include the disease name

Plasmids

1. Visit Addgene.com and select "Deposit a Plasmid" on the home page

2. Scroll down to "Submit Plasmids Using a Spreadsheet"
3. Select "Download Deposit Spreadsheet"

4. Save the file as "Lab/PI Name_Addgene_BatchUpload"
   Example: SmithLab_Addgene_BatchUpload

Plasmid Name	Plasmid Type	Purpose	PubMed ID	Gene or insert Name	Alternative Gene/Insert Name 1	Alternative Gene/Insert Name 2	Entrez Gene ID	gRNA/shRNA Sequence	Insert Size	Species of gene or insert	Additional Species of gene or insert	Species Other	Relevant Mutations

If you are depositing more than 75 plasmids, you must email deposit@addgene.org BEFORE completing this spreadsheet.

5. Fill in the spreadsheet with as much information as possible.
   Please review the <u>REQUIRED</u> information below:

- Plasmid Name
- Plasmid Type
- Purpose
- Gene or Insert Name
- Species of gene or insert
- Relevant Mutations
- Backbone Name

- Primary vector type
- Cloning Method
- Bacterial Resistance
- High or Low Copy
- Growth Temperature
- Growth SStrain
- Hazardous
- Patents or licenses

6. Upon completion, email the following to deposit@addgene.org
   a. Completed spreadsheet
   b. Any plasmid sequence files, maps, and/or GenBank files; each file name containing the provided plasmid name
   c. Shipping address for Addgene to send a Deposit Kit
   d. Name of Principal Investigator and Organization where the constructs were first created
   e. Addgene account username (optional: include this information if you want the plasmids to appear in your account under "My Plasmids"

Mice

1. Register mouse strains with Mouse Genome Informatics (MGI) by visiting http://www.informatics.jax.org/mgihome/submissions/amsp_submission.cgi

2. Fill in contact details

Mutant Alleles, Strains, and Phenotypes Submission Form

Contact Details: * = required field

Last name: *		Street Address:	
First name (& middle initial): *		City:	
E-mail address: *		State/Province:	
E-mail address (repeat): *		Postal Code:	
Principal Investigator:		Country:	
Institute/Organization:		Telephone:	
		Fax:	

3. Select publication preference

Citing your data:

Are your data published?   ○ yes ○ no

If no, would you prefer that your data: ○ be public at the MGI website now ○ be held private until publication

Provide reference(s) or PubMed IDs for published data or authors & descriptive title for unpublished data:

If data are available from a website, please list URL:

4. Provide any additional information by selecting the appropriate **Purple** heading

Choose information type(s) to submit.
Click a heading to open or close its respective section.
   Allele Name and describe a new allele, mutation, or transgene
   Strain Register a new mouse strain
   Phenotypes Submit phenotype data for given genotypes
   File Submissions Submit data files (e.g. images, text files, Excel, or bulk data)

5. Provide any additional comments. Click 'verify'

6. For Alleles, provide the following:

Enter Allele Data: * = required field

Suggest **symbol** and/or **name** for this mutation: * [   ]

If this mutation is an allele of a **known gene** enter the **gene symbol** or **MGI ID**: [   ] (**Check by** searching MGI.)

Common nicknames for this mutant allele [   ]

Class of Allele (check all that apply): *

☐ spontaneous ☐ ENU induced ☐ chemical (non-ENU) induced ☐ irradiation induced ☐ transgenic ☐ gene trapped ☐ targeted
☐ conditional/targeted ☐ endonuclease-mediated ☐ recombinase (cre or other) containing ☐ transposon induced ☐ other (specify) [   ]

For transgenes, specify transgene promoter: [   ]
For targeted mutations or gene traps, specify ES cell line used: [   ] (**Example**: E14.1, JM8A3)
For gene traps, specify the resulting mutant ES cell line: [   ] (**Example**: AD0888)

Inheritance: ○ dominant ○ codominant ○ semidominant ○ recessive ○ X-linked ○ other (specify) [   ] ○ unknown/not applicable

Strain background in which the mutation occurred [   ] (**Examples**: C57BL/6J, 129P2/OlaHsd)

Genome Location (Chromosome, genome coordinates, cM) and molecular detail about this allele, such as "exon 1 deletion, etc.":
[   ]

- For hints on mutant allele nomenclature, see Quick Guide to Nomenclature for Alleles and Mutations.
- Lab codes are available from ILAR (Institute of Laboratory Animal Resources).
- **If you would like assistance with allele nomenclature:**
  ☐ Check the box here and continue with your submission. We will contact you about nomenclature for this mutation.

7. For Strains, provide the following:

Register a New Mouse Strain: * = required field

Enter a suggested **strain name**.
When mutant alleles are part of the strain name, use < > to indicate the superscripted alleles.
Example: **C57BL/6J-Kit**$^{W-39J}$ should be entered as **C57BL/6J-Kit<W-39J>**
* [   ]

Enter the **gene symbols** corresponding to alleles carried on this strain.
Example: for the strain **NOD/LtSz-Prkdc<scid> B2m<tm1Unc>**, the gene symbols entered into this box would be **Prkdc** and **B2m**.
[   ]
(one gene symbol per line)

If this strain is in a repository, please list the repository [   ] (**View**: list of repositories)
Enter its repository ID or MGI ID for this strain, if known [   ]

Strain categories: Choose one or more. *

☐ inbred strain   ☐ wild-derived ☐ recombinant inbred (RI)
☐ segregating inbred ☐ outbred   ☐ recombinant congenic (RC)
☐ mutant strain   ☐ coisogenic   ☐ major histocompatibility congenic
☐ mutant stock   ☐ congenic   ☐ minor histocompatibility congenic
                       ☐ consomic   ☐ other, specify [   ]

- For hints on strain nomenclature, see Guidelines for Nomenclature of Mouse and Rat Strains.
- Lab codes are available from ILAR (Institute of Laboratory Animal Resources).
- **If you would like assistance with strain nomenclature:**
  ☐ Check the box and continue with your submission. We will contact you about nomenclature for this strain.

8. For Phenotypes, provide the following:

Submit Phenotype Data: * = required field

Mutant allele(s) analyzed.

List one or more allele pairs analyzed in the animal (one allele pair per line, with the alleles comma separated).
When entering mutant alleles, use < > to indicate the superscripted portion of an allele.

Example:
If you phenotyped animals that were heterozygous for **Kit^{W-39J}** and homozygous for **Tec^{tm1Welm}**, they should be entered as

```
Kit<W-39J>, Kit<+>
Tec<tm1Welm>, Tec<tm1Welm>
```

Enter allele pairs of your phenotyped animals: *

(Find the correct allele symbol by searching MGI.)

Additional allele information not currently in MGI (allele synonyms, ES cell line, strain of origin, mutation type, molecular description, etc.):

Genetic Background: Genetic background can have a significant effect on phenotype.

Enter the Strain/Genetic Background on which phenotypes were analyzed: *

Other Strain/Background Information (e.g. specify crosses): Click here for an example.

- **If you would like assistance with the Genetic Background Section:**
  ☐ Check the box here and continue with your submission. We will contact you about determining the correct genetic background.

Phenotype:

Phenotypic Description (enter text, describing details of phenotypes observed, etc.): *
Click here for an example. You may browse the Mammalian Phenotype Ontology and use these terms to describe the phenotype.

If this genotype + genetic background is a model for a human disease based on phenotypic similarity, please name the disease and include any associated information:

Other known information (gene function/pathway, available clones, GenBank numbers, etc.) that will enhance these data:

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9. For File Submission, provide the following:

File submissions:

You may submit a limited number of files using this form. Please limit file size to <5 MB.
If you have larger files, or many files to submit, please contact us at: mgi-submissions@jax.org.

Are the file(s) that you submitted copyrighted? ○ yes ○ no
If you have entered copyrighted information we will contact you.

Upload your data files (images, text descriptions, Excel, or text data):
File 1: [Choose File] No file chosen   [Submit more files]

See examples and templates for file submissions.

10. Select "Verify" to review and complete your submission

Completing your submission:

Are there any additional commments or information you would like to convey about your data?

[   ]

Use the buttons below to verify your data before submission or to reset the entire form. Thank you!

[Verify] [Reset]

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Rats

1. Register rat strains with Rat Genome Database (RGB) by visiting https://rgd.mcw.edu/ rgdweb/models/strainSubmissionForm. html?new=true

2. Provide Strain information

Strain Information

Strain Symbol *

[Enter strain symbol]

Please enter a symbol for the strain being submitted, for example: BN/Crl, or SHR.BN-(D13Arb5-Ren)/Ipcv. For more information, please refer to the rules for strain nomenclature and the rules for naming endonuclease-mediated mutants for strains produced using CRISPR/Cas, TALEN or ZFN mutagenesis. For help naming your strain please contact RGD.Data@mcw.edu.

Type
[mutant ⌄]

Genetic Status
[Homozygous ⌄]

Method
[CRISPR/Cas9 ⌄]

Other Method

[Enter modification method]

If you selected "other" as the method above, please specify the method you used here.

Background Strain

[Enter background strain symbol]

Provide a description of strain's origin

[   ]

Example:"The CRISPR/Cas9 system was used to introduce a 5-bp deletion in exon 3 of the Spp1 gene of SHRSP/A3NCrl rat embryos.

Reference/Pubmed ID

[Enter reference/pubmed id]

If this strain has been mentioned in any published paper please give that citation here. (author, journal, vol, page number, or PMID).

Research Use

[Eg: Cancer, Cardiovascular, etc.]

Optional. If this strain has been or could be used for research in a particular area, please indicate this. For example, this could be a disease area such as cardiovascular or cancer, or a phenotype category such as reproduction.

ILAR Code

[Enter ILAR code]

This is the laboratory code assigned by The Institute of Laboratory and Animal Research to each lab or group. Registering a group at ILAR identifies the group as a place where this strain was originated and maintained.

Upload image file if available

[Choose File] No file chosen

Upload image of size less than 5MB. Acceptable file formats: .PNG, .JPEG, .GIF

3. Provide Gene/Allele information

Gene/Allele Information

Gene Symbol

Enter gene symbol

Gene RGD ID (if known)

Enter gene RGD ID

Allele Symbol

Enter allele symbol

Allele RGD ID (if known)

Enter allele RGD ID

4. Indicate Availability

Availability

Current Status: ☐ Live Animals ☐ Cryopreserved Embryo ☐ Cryopreserved Sperm

Where could this strain be obtained? *

Availability Contact Email

Availability Contact URL

5. Provide Contact information

Submitter Contact Details

Last Name/Surname *

Enter submitter last name/surname

First Name *

Enter submitter first name/given name

Email Address *

Enter submitter email address

Laboratory PI

Enter PI First Name & Last Name

PI Email Address

Enter PI email address

Institution/Orgainzation

Enter Institution or Organization name

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6. Provide any additional information and indicate preference for public sharing. At the time of publication, the strain will need to be public per ASAP Open Science Policies.

Additional Information

Additional Information about the STRAIN or ALLELE or GENE or any information you want to provide.

Please let us know if you want this strain to be displayed on the RGD website. If not, check Non Public (we can hold a strain until instructed by you to release it).

○ **Public** ◉ **Non Public**

7. Confirm you are not a robot and submit.

☐ I'm not a robot

reCAPTCHA
Privacy - Terms

Submit