

# New Registration and Data System (New Registration)

# **Submitting Investigator Entry of Registration Information and Data Submission for a New Study**





# Enter Study Registration Information

+ OMB control number: 0925-0670 Expiration date: 03/31/2026

- My Studies
- Register a New Study**
- API Documentation



## Funding and Support

### NIH Funding

\* Does this study have NIH funding?

- Yes
- No

\* Look up Main NIH Grant Number

[Look up](#)

[Back](#)

[Continue](#) [Save draft](#) [Discard draft](#)

Time-saving search function

### Step 1 of 2: Look up Main NIH Grant Number

NIH grant numbers typically start with "Z0", "R0", "U", or similar. Supporting grant numbers can be added in a later step.

\* Enter full grant number

2R01NS027863-06 (assume this is the full grant number)

[Look up grant number](#) [Close](#)

### Step 2 of 2: Confirm Main NIH Grant Number

1 result found. To refine your search, select "Go Back" below.

- 2R01NS027863-06 (R01NS027863)**  
Grant name: RANDOMIZED TRIAL OF ORG 10172 IN ACUTE ISCHEMIC STROKE  
Primary ICO: NIAID  
Funding ICO: NCBI  
PI: Arnold Louie  
Organization: University of Iowa  
Fiscal year: 1997

[Confirm grant number](#) [Go back](#) [Close](#)

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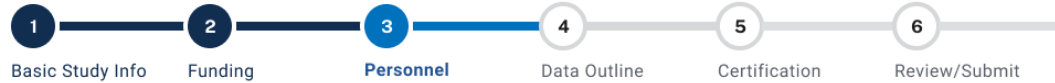
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# Enter Study Registration Information

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## Personnel

\* Select Principal Investigator (PI)

User: Anne Doe ([Change](#))

Email: [anne.doe@nih.gov](mailto:anne.doe@nih.gov)

Institute: NLM (National Library of Medicine)

[View in NED](#)

Select Main Data Submitter

[Add contact info](#)

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### Select Main Data Submitter

\* First name

\* Last name

\* Email address

[Save submitter](#)

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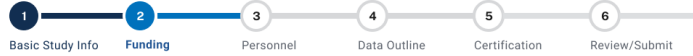
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### Funding and Support

#### Supporting Institutions

##### NIH Institutes

Select a supporting NIH institute, if applicable

NIDCR

OD

+ Add an additional supporting NIH ICO

List any additional NIH grant information below. Enter each grant on a separate line, separated by line breaks.

Z0213498734  
R0193847934  
U23213498734  
R0193847934

##### Non-NIH Institutions or Funding Organizations

List any supporting non-NIH institutions or funding organizations below. Enter each entity on a separate line, separated by line breaks

Lorem ipsum organization  
University of Colorado

List any additional grants from non-NIH institutions or funding organization below. Enter each grant on a separate line, separated by line breaks

Grant 1  
Grant 2  
Grant 3

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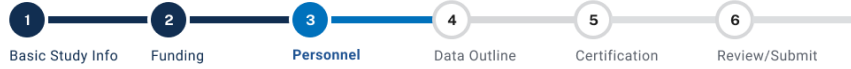
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# Enter Study Registration Information

+ OMB control number: 0925-0670 Expiration date: 03/31/2026

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## Personnel

\* Select Principal Investigator (PI)

User: Anne Doe ([Change](#))

Email: anne.doe@nih.gov

Institute: NLM (National Library of Medicine)

[View in NED](#)

Select Main Data Submitter

User: Jon Snow ([Change](#))

Email: jon.snow@nih.gov

\* Does this study have a Program Officer (PO)?

Yes

No

\* Select Program Officer (PO)

User: Charlie Cupcake ([Change](#))

Email: charlie.cupcake@nih.gov

Institute: NLM (National Library of Medicine)

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### Step 1 of 2: Look Up User

First name

Last name

Email address

[Look up](#)

[Close](#)

### Step 2 of 2: Select User

3 results found. To refine your search, select "Go Back" below.

**Charlie Cupcake** (charlie.cupcake@nih.gov)  
NED ID: 0123456789  
Institute: NLM (National Library of Medicine)  
IPD: Staff Scientist

**Charlie Cupcake II** (charlie.m.cupcake@nih.gov)  
NED ID: 0123987654  
Institute: NCI (National Cancer Institute)  
IPD: Staff Scientist

**Charlie Cupcake III** (charlie.z.cupcake@nih.gov)  
NED ID: 1123456789  
Institute: NHBLI (National Heart, Blood, and Lung Institute)  
IPD: Staff Scientist

[Select User](#)

[Go back](#)

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- My Studies
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## Study Data Outline (SDO)

\* Will data be submitted for storage by dbGaP?

- Yes
- No

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## Institutional Certification (IC)

### Upload Files

\* Upload all required Institutional Certifications

Select PDF(s)

**Selected file(s)** [Change files](#)

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- Sit amet.pdf

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**Submitting Investigator Edit Study Registration**

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**Once the study is registered, the Submitting Investigator is sent the Submission Portal to submit data to dbGaP**

### Submission Portal

Groups Accounts Templates Stats My profile

**OMB control Number: 0925-0670**  
**Expiration Date: March 31, 2026**  
 Public reporting burden for this collection of information is estimated to be about one hour per response including the time gathering and maintaining the information needed, adding information to this part of the study registration and uploading the data. An agency may not conduct or sponsor, and a person is not required to respond to a collection of information unless it displays a currently valid OMB control number. Send comments regarding this burden estimate or any other aspect of this collection of information, including suggestions for reducing this burden, to: NIH, Project Clearance Branch, 6705 Rockledge Drive, MSC 7974, Bethesda, MD 20892-7974, ATTN: PRA (0925-0670). Do not return the completed form to this address.

### dbGaP Studies

Study Id	Study Name	Investigator	Agency	Study ID	File Name	Files Updated	Last Updated	Study status
65008	MoTrPAC Multiomics Dataset – Adult Humans – Restricted Access	Shruti Marwaha	NIDDK		phs004523.v1.p1	None	2026-02-18 17:15	<a href="#">View study status</a>
59562	Human GPR15 deficiency reveals CD8+ T regulatory cell-macrophage crosstalk in intestinal inflammation	Yu Zhang	NIAID	GPR15_Deficiency_IBD	phs003950.v1.p1	2026-02-04 11:58	2026-02-18 16:32	<a href="#">View study status</a>
65470	Interrogating gene regulation during human cortical development	Yashodara Abeykoon	NIMH		phs004522.v1.p1	None	2026-02-18 16:16	<a href="#">View study status</a>
64362	Tissue Collection and Characterization: Longitudinal Senescence Analysis		None		.v1	None	2026-02-18 16:04	<a href="#">View study status</a>
65559	Liver Study Collection		None		.v1	None	2026-02-18 14:23	<a href="#">View study status</a>
57380	<a href="#">A091401 Randomized Study of Nivolumab With or Without Ipilimumab in Patients With Metastatic or Unresectable</a>	Paige Fuentes	NCI	A091401	phs003785.v1.p1	2026-02-18 13:55	2026-02-18 14:16	<a href="#">View study status</a>

Submitting Investigator can click on the registered study to enter required information for data submission

## The Genetics of Telework Stress (Example Study)

Study Data Outline [?](#)

Consent group registered for this study [?](#)

Consent Group Title	Consent Group Abbreviation
Disease-Specific (Allergy, IRB, PUB, COL, NPU, MDS, GSO, RD)	DS-ALERG-IRB-PUB-COL-NPU-MDS-GSO-RD
General Research Use	GRU

Required fields are marked with \* asterisk

Please review all questions and select all data types that will be provided by you or another submitter as part of this study

Required information provided by the Submitting Investigator for data submission

Study data type	Study data outline provided at registration	Select 'Yes' if this data type will be provided						
<p>* 1. Subject IDs, Sample IDs, and Consents <a href="#">?</a></p> <table border="1"> <thead> <tr> <th>Subject Consent <a href="#">?</a></th> <th>Subject Sample Mapping <a href="#">?</a></th> </tr> </thead> <tbody> <tr> <td><a href="#">2a_SubjectConsent_DS.txt</a></td> <td><a href="#">3a_SSM_DS.txt</a></td> </tr> <tr> <td><a href="#">2b_SubjectConsent_DD.xlsx</a></td> <td><a href="#">3b_SSM_DD.xlsx</a></td> </tr> </tbody> </table> <p>* Pedigree: Are the relationships of the subjects known? <a href="#">?</a></p>	Subject Consent <a href="#">?</a>	Subject Sample Mapping <a href="#">?</a>	<a href="#">2a_SubjectConsent_DS.txt</a>	<a href="#">3a_SSM_DS.txt</a>	<a href="#">2b_SubjectConsent_DD.xlsx</a>	<a href="#">3b_SSM_DD.xlsx</a>	Yes	<p><input checked="" type="radio"/> Yes <input type="radio"/> No</p> <p><input type="radio"/> Yes <input checked="" type="radio"/> No</p>
Subject Consent <a href="#">?</a>	Subject Sample Mapping <a href="#">?</a>							
<a href="#">2a_SubjectConsent_DS.txt</a>	<a href="#">3a_SSM_DS.txt</a>							
<a href="#">2b_SubjectConsent_DD.xlsx</a>	<a href="#">3b_SSM_DD.xlsx</a>							
<p>* 2. Phenotypes and Attributes <a href="#">?</a></p> <table border="1"> <thead> <tr> <th>Subject Phenotypes</th> <th>Sample Attributes</th> </tr> </thead> <tbody> <tr> <td><a href="#">5a_SubjectPhenotypes_DS.txt</a></td> <td><a href="#">6a_SampleAttributes_DS.txt</a></td> </tr> <tr> <td><a href="#">5b_SubjectPhenotypes_DD.xlsx</a></td> <td><a href="#">6b_SampleAttributes_DD.xlsx</a></td> </tr> </tbody> </table>	Subject Phenotypes	Sample Attributes	<a href="#">5a_SubjectPhenotypes_DS.txt</a>	<a href="#">6a_SampleAttributes_DS.txt</a>	<a href="#">5b_SubjectPhenotypes_DD.xlsx</a>	<a href="#">6b_SampleAttributes_DD.xlsx</a>	Yes	<p><input checked="" type="radio"/> Yes <input type="radio"/> No</p>
Subject Phenotypes	Sample Attributes							
<a href="#">5a_SubjectPhenotypes_DS.txt</a>	<a href="#">6a_SampleAttributes_DS.txt</a>							
<a href="#">5b_SubjectPhenotypes_DD.xlsx</a>	<a href="#">6b_SampleAttributes_DD.xlsx</a>							
<p>* 3. Molecular (DOES NOT include high throughput sequencing data such as BAM, CRAM, and FASTQ) <a href="#">?</a></p>	No	<p><input checked="" type="radio"/> Yes <input type="radio"/> No</p> <p><b>Genome Build</b></p> <p>GRCh38 <input type="text"/></p> <p><b>Array Name</b></p> <p>Illumina 1M <input type="text"/></p> <p>* Select at least one</p> <p><input type="checkbox"/> Sequence-based genotypes (multi-sample vcf)</p> <p><input type="checkbox"/> Sequence-based genotypes (single sample vcf)</p> <p><input type="checkbox"/> CNV derived from sequence (vcf)</p> <p><input checked="" type="checkbox"/> SNP array (PLINK and/or raw data)</p>						

## Required information provided by the Submitting Investigator for data submission

- SNP array (PLINK and/or raw data)
- CNV calls from array (PLINK and/or raw data)
- Imputed genotypes
- Tumor / Normal variations / Somatic SNV (maf)
- Methylation array or summaries
- Expression array or summaries
- Other

**\* Other description**

test data for sdo change email

**Describe this data**

<p><b>* 4. Links to public NCBI databases</b> <a href="#">?</a></p>	<p>No</p>	<p><input type="radio"/> Yes <input checked="" type="radio"/> No</p>
<p><b>* 5. Sequence</b> <a href="#">?</a></p>	<p>Yes</p>	<p><input checked="" type="radio"/> Yes <input type="radio"/> No</p> <p><b>Please select one option</b></p> <p><input checked="" type="radio"/> NCBI Storage (SRA)</p> <p><input type="radio"/> Cloud Storage (External Data Sources including Trusted Partners)</p> <p><b>* Select at least one</b></p> <p><input checked="" type="checkbox"/> Whole or Targeted Genome</p> <p><input type="checkbox"/> Whole or Targeted Exome</p> <p><input type="checkbox"/> RNASeq - Whole or targeted transcriptome</p> <p><input type="checkbox"/> MethylSeq / Epigenomic Marks</p> <p><input type="checkbox"/> Microbiome sequence (Metagenome, 16S rRNA, 18S rRNA, ITS)</p> <p><input type="checkbox"/> Other</p>
<p><b>* 6. Association Analysis</b> <a href="#">?</a></p>	<p>No</p>	<p><input type="radio"/> Yes <input checked="" type="radio"/> No</p>
<p><b>* 7. Documents (protocols, questionnaires, consent forms, etc)</b> <a href="#">?</a></p>	<p>No</p>	<p><input type="radio"/> Yes <input checked="" type="radio"/> No</p>
<p><b>* 8. Medical Images</b> <a href="#">?</a></p>	<p>No</p>	<p><input type="radio"/> Yes <input checked="" type="radio"/> No</p>
<p>9. Other files not covered and/or any notes about study outline</p>		<p></p>

Submit

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