

# Data Availability Statements

Why so hard?

Juliane Schneider  
Sage Bionetworks





Files » 1.IndividualStudies » Human » FreshMicro

## FreshMicro ☆

Folder Tools ▾

Synapse ID: syn25671134 ⓘ Item count: 3 Access: View Terms Report Violation Storage Location: s3://ad-knowledge-portal-main/data



## The Fresh Microglia Regulome (FreshMicro) Study

Genetics of the human microglia regulome refines Alzheimer's disease risk loci

**Sample information :** All brain specimens were obtained through informed consent and/or brain donation programs at the respective organizations. All procedures and research protocols were approved by the corresponding ethical committees of our collaborator's institutions.

**Autopsies:** The autopsy brain specimens originated from brain donation programs at Rush University Medical Center/Rush Alzheimer's Disease Center (RADC) in Chicago, IL and The Mount Sinai/JJ Peters VA Medical Center NIH Brain and Tissue Repository (NBTR) in the Bronx, NY.

**RADC:** Samples were collected at RADC in Chicago, IL as part of two prospective studies of aging: the Religious Orders Study (ROS) (1) and the Memory and Aging Project (MAP) (2). At the time of enrollment, participants had to be at least 53 (ROS) or 55 (MAP) years old and non-demented at the time of enrollment, and were required to sign an Anatomical Gift Act agreeing to donate their brain and spinal cord at the time of death. Extensive annual neuropsychologic evaluations were collected prior to death, with a structured, quantitative neuropathologic examination conducted at autopsy, described at <https://www.radc.rush.edu/docs/var/variables.htm>. After autopsy, the collected tissue was weighted, placed in ice-cold MACS tissue storage solution (Miltenyi Biotec, cat# 130-100-008) and shipped overnight at 4°C with priority shipping.

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Wiki Revision History



required to sign an Anatomical Gift Act agreeing to donate their brain and spinal cord at the time of death. Extensive annual neuropsychologic evaluations were collected prior to death, with a structured, quantitative neuropathologic examination conducted at autopsy, described at <https://www.radc.rush.edu/docs/var/variables.htm>. After autopsy, the collected tissue was weighted, placed in ice-cold MACS tissue storage solution (Miltenyi Biotec, cat# 130-100-008) and shipped overnight at 4°C with priority shipping.

### Show More














































































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Wiki Revision History

Download Options

Name		Size	Modified On	ID	Download
▶ Analysis			04/11/2022 11:14 AM	syn26213257	
▼ Data			05/04/2021 8:22 AM	syn25671135	
▶ Deprecated	✓		04/07/2022 8:51 AM	syn26708613	
▶ Epigenetics			09/30/2021 12:31 PM	syn25671141	
▶ Gene Expression			09/30/2021 12:31 PM	syn25671140	
▶ Genomic Variants			09/30/2021 12:30 PM	syn25671142	
▶ Metadata			09/30/2021 12:31 PM	syn25671143	
▶ Study Description Images			09/30/2021 12:32 PM	syn26230219	

Created by @abby.vanderlinden on May 4, 2021 8:22 AM

        	▼ <b>Gene Expression (RNA seq)</b>	 		05/04/2021 8:24 AM	syn25671146	
	▶ <b>counts</b>			01/05/2022 10:57 AM	syn25671149	
	▼ <b>raw</b>			05/04/2021 8:24 AM	syn25671148	
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▶	Genomic Variants			09/30/2021 12:30 PM	syn25671142	
▼	Metadata			09/30/2021 12:31 PM	syn25671143	
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	FreshMicro_assay_HiC_metadata.csv		3.7 KB	10/18/2021 1:00 PM	syn26254720	↓ +
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 Modified by [@abby.vanderlinden](#) on September 30, 2021 12:33 PM



Experimental Mode Off

## GeneticsoftheHumanMicroglia Regulome\_Kosoy.etal ☆

Synapse ID: syn26207321 ⓘ DOI: <https://doi.org/10.7303/syn26207321> Access: ⓘ Add Conditions for Use ⓘ Report Violation Storage Location

### Manuscript: Genetics of the human microglia regulome refines Alzheimer's disease risk loci

Authors: Roman Kosoy<sup>1,2,3,4,\*</sup>, John F. Fullard<sup>1,2,3,4,\*</sup>, Biao Zeng<sup>1,2,3,4,\*</sup>, Jaroslav Bendl<sup>1,2,3,4,\*</sup>, Pengfei Dong<sup>1,2,3,4</sup>, Samir Rahman<sup>1,2,3,4</sup>, Steven P. Kleopoulos<sup>1,2,8</sup>, Alexander W. Charney<sup>1,2,5</sup>, Towfique Raj<sup>2,4,7,8</sup>, David Bennett<sup>9,10</sup>, Christopher P. Kellner<sup>6</sup>, Vahram Haroutunian<sup>5,7,11</sup>, Gabriel E. Hoffman<sup>1,2,3,†</sup>, Panos Roussos

<sup>1</sup>Pamela Sklar Division of Psychiatric Genomics,

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<sup>3</sup>Icahn Institute for Data Science and Genomic Technology,

<sup>4</sup>Friedman Brain Institute,

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\*These first authors contributed equally to this work.

†These senior authors contributed equally to this work.

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**Panos Roussos**; Icahn School of Medicine at Mount Sinai; 1470 Madison Ave, Fl. 9, Rm 107

medRxiv preprint is available at doi: <https://doi.org/10.1101/2021.10.17.21264910>

Data described in this manuscript comes from the [AD Knowledge Portal](#)

#### Reproducible Analysis:

Data distributed through the AD Knowledge Portal is de-identified of PHI in accordance with HIPAA privacy rules as required by the Synapse system governance policy

#### Data:

Data described in this manuscript is available via the [Fresh Microglia Regulome Study](#).

#### Data access and download instructions

- [Data access](#)
- Data download
  - [individual files](#)
  - [bulk](#)

[https://docs.synapse.org/articles/downloading\\_data.html](https://docs.synapse.org/articles/downloading_data.html)



## Description of research output in manuscript

## Location and link of repo

## Description of repository/funder

"<data, analysis output, tools (describe content)> are available via the AD Knowledge Portal (<https://adknowledgeportal.org>). The AD Knowledge Portal is a platform for accessing data, analyses, and tools generated by the Accelerating Medicines Partnership (AMP-AD) Target Discovery Program and other National Institute on Aging (NIA)-supported programs to enable open-science practices and accelerate translational learning. The data, analyses and tools are shared early in the research cycle without a publication embargo on secondary use. Data is available for general research use according to the following requirements for data access and data attribution (<https://adknowledgeportal.org/DataAccess/Instructions>).

For access to content described in this manuscript see: <manuscript landing page DOI>"

## Data DOI

## Conditions for access and use with link to instructions



# AD Knowledge Portal - backend ★

Synapse ID: syn2580853



DOI: <https://doi.org/10.7303/syn2580853>

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Project Settings ▾

Wiki ⓘ

Files ⓘ

**Datasets ⓘ**

Tables ⓘ

Discussion ⓘ

Docker ⓘ

Add Dataset

## Datasets

Name	Type	File Count	Modified On	Created On	ID
AD_Portal_Dataset_Template	Dataset	44	04/14/2022 1:55 PM	04/14/2022 1:18 PM	syn29421835
Genetics of the human microglia regulome refines Alzheimer's disease risk loci	Dataset	575	04/12/2022 3:42 PM	03/30/2022 8:34 PM	syn28575168
Transcriptional landscape of human microglia implicates age, sex and APOE-related immunometabolic pathway perturbations	Dataset	147	04/13/2022 1:23 PM	03/24/2022 6:07 PM	syn28450881
Deletion of Abi3_Gngt2 influences age-progressive amyloid beta and tau pathologies in distinctive ways	Dataset	51	03/24/2022 6:43 PM	03/24/2022 12:48 PM	syn28412421
CharacterizationofMitochondrialDNAQuantityAndQuality_Klein	Dataset	3,799	04/12/2022 3:41 PM	03/01/2022 6:08 PM	syn27337082



# Genetics of the human microglia regulome refines Alzheimer's disease risk loci



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Version: Draft - [Show Version History](#)

Synapse ID: syn28575168

DOI: <https://doi.org/10.7303/syn28575168> Access: Report Violation

Dataset DOI

Manuscript: Genetics of the human microglia regulome refines Alzheimer's disease risk loci

medRxiv preprint: <https://doi.org/10.1101/2021.10.17.21264910>

Manuscript DOI

Authors: Roman Kosoy<sup>1,2,3,4,\*</sup>, John F. Fullard<sup>1,2,3,4,\*</sup>, Biao Zeng<sup>1,2,3,4,\*</sup>, Jaroslav Bendl<sup>1,2,3,4,\*</sup>, Pengfei Dong<sup>1,2,3,4</sup>, Samir Rahman<sup>1,2,3,4</sup>, Steven P. Kleopoulos<sup>1,2,3,4</sup>, Zhiping Shao<sup>1,2,3,4</sup>, Jack Humphrey<sup>2,4,7,8</sup>, Katia de Paiva Lopes<sup>2,4,7,8</sup>, Alexander W. Charney<sup>1,2,5</sup>, Towfique Raj<sup>2,4,7,8</sup>, David Bennett<sup>9,10</sup>, Christopher P. Kellner<sup>6</sup>, Vahram Haroutunian<sup>5,7,11</sup>, Gabriel E. Hoffman<sup>1,2,3,†</sup>, Panos Roussos<sup>1,2,3,4,5,11,\*</sup>

<sup>1</sup>Pamela Sklar Division of Psychiatric Genomics, <sup>2</sup>Department of Ge<sup>4</sup>Friedman Brain Institute, <sup>5</sup>Department of Psychiatry, <sup>6</sup>DepartmenMedicine at Mount Sinai, NY, USA; <sup>9</sup>Rush Alzheimer's Disease CentRush University Medical Center, Chicago, Illinois, USA; <sup>11</sup>Mental Ill

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study	
<input type="checkbox"/> FreshMicro	575
resourceType	
<input type="checkbox"/> experimentalData	515
<input type="checkbox"/> analysis	54
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## Files (575)

575 Results

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syn26249697	FM_005_ATACseq_TAGGCAT...	FreshMicro	experimentalData	epigenetics	ATACSeq	microglia
syn26249698	FM_005_ATACseq_TAGGCAT...	FreshMicro	experimentalData	epigenetics	ATACSeq	microglia
syn26249699	FM_006_ATACseq_CGTACTA...	FreshMicro	experimentalData	epigenetics	ATACSeq	microglia
syn26249700	FM_006_ATACseq_CGTACTA...	FreshMicro	experimentalData	epigenetics	ATACSeq	microglia
syn26249701	FM_007_ATACseq_TCCTGAG...	FreshMicro	experimentalData	epigenetics	ATACSeq	microglia
syn26249702	FM_007_ATACseq_TCCTGAG...	FreshMicro	experimentalData	epigenetics	ATACSeq	microglia
syn26249735	FM_008_ATACseq_GGACTCC...	FreshMicro	experimentalData	epigenetics	ATACSeq	microglia
syn26249758	FM_008_ATACseq_GGACTCC...	FreshMicro	experimentalData	epigenetics	ATACSeq	microglia
syn26249763	FM_009_ATACseq_TAGGCAT...	FreshMicro	experimentalData	epigenetics	ATACSeq	microglia
syn26249765	FM_009_ATACseq_TAGGCAT...	FreshMicro	experimentalData	epigenetics	ATACSeq	microglia

## study

☐ FreshMicro

## resourceType

☐ experimentalData☐ analysis☐ metadata

## dataType

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## assay

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## cellType

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## dataSubtype

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## Files (575)

575 Results

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## Files (575)

575 Results

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sorted cells	brain	IlluminaNovaseq6000	50	pairedEnd	female	Human	FM_012_ATACseq	prefront

Wiki Revision History

study

☐ FreshMicro 6

resourceType

☐ experimentalData 515

☐ analysis 54

☒ metadata 6

dataType

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☐ epigenetics 2

☐ geneExpression 1

☐ genomicVariants 1

assay

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☐ Hi-C 1

☐ rnaSeq 1







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cellType

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Files (575)

6 Results

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 <b>syn26254720</b>	FreshMicro_assay_HiC_metadata.csv	FreshMicro	metadata	genomicVariants	snpArray		AMP-AD	
 <b>syn26254721</b>	FreshMicro_biospecimen_metadata.csv	FreshMicro	metadata	clinical			AMP-AD	
 <b>syn26254722</b>	FreshMicro_individual_metadata.csv	FreshMicro	metadata	clinical			AMP-AD	

FreshMicro\_individual\_metadata.csv

Last updated on 04/12/2022 10:22

# AD Knowledge Portal - backend

Synapse ID: syn2580853 DOI: <https://doi.org/10.7303/syn2580853> Storage Location: s3://ad-knowledge-portal-main/data

Project Settings

Wiki

Files

Datasets

Tables

Discussion

Docker

Files » 1.IndividualStudies » Human » ADMC\_ADNI\_BakerLipidomics » Data » Metabolomics » Metabolomics (Baker Lipi... » ADMC Lipidomics Meikle L...

## ADMC Lipidomics Meikle Lab Baseline Data Matrix

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Version: 2 (Current) - [Show Version History](#)

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Synapse ID: syn22106053 Access: [View Terms](#) [Report Violation](#)

Preview

No preview is available for syn22106053.2

Provenance

Undefined

ADMC  
Lipidomic..  
v.2

Discussion about ADMC Lipidomics Meikle Lab Baseline Data Matrix

No threads found.

Created by @cblach15 on May 29, 2020 1:46 PM



## AD Knowledge Portal - backend

Synapse ID: syn2580853

DOI: <https://doi.org/10.26434/chemrxiv-2021-04-13>

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Datasets

Files » 1.IndividualStudies » Human » ADMC\_ADNI\_E

### ADMC Lipidomics Meikle Lab Baseline

Version: 2 (Current) - [Show Version History](#)

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URL: <https://ida.loni.usc.edu/data/ampad?id=59&version=2021-04-13>

Synapse ID: syn22106053 [View Terms](#) [Report Violation](#)

Preview

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Provenance

Undefined

ADMC  
Lipidomic..  
v.2

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Password

Discussion about ADMC Lipidomics Meikle Lab Baseline Data Matrix

No threads found.

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# Pros

Direct link to data

Able to choose data files from disparate studies

Immediate access to metadata

# Cons

Data has to be in the Synapse platform

There are still access issues when we link out to data from other repositories

No automatic access

System doesn't know who you are and what your access level is if you click on the DOI