

NIAGADS Update: Spring 2024

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COI: Nothing to Disclose

NIAGADS at a Glance



Data Sharing

113 datasets
>183,000 samples
497 Unique Requests
238 Institutional Certifications

Userbase



>330 articles cite U24

>4,900 verified users

ADSP

73 cohorts23 current grant subawards

Exomes 20,503

Genomes

~350% increase from 2021 to 2024

(16,905 in 2021 to ~60,000 genomes in 2024)

ADSP Support

14 workgroups /
 5 consortia
 >500 members
 52 institutions

NIAGADS 2023 Growth





+17% datasets

+24%

articles that cite U24

+28% verified users

ADSP WGS Releases



Released in 2023

December

R4 WGS Annotation PHC 2023 Release

Coming Soon in 2024

| Q2 | SV joint genotype called pVCF |
|----|-------------------------------|
| 4- | QC'd chrX/multi-allelic pVCF |

R5 WGS- ~62k genomes CRAMs, gVCFs, preview pVCF

ADSP 2023 Deliverables – NG00067v11 Phenotype Harmonization Consortium (PHC)



| Cognition | Fluid Biomarker | Neuropathology | Vascular Risk Factors |
|--|---|---|---|
| •Harmonized composite scores (PHC_EF, PHC_MEM, PHC_LAN, PHC_VSP) | Amyloid ß,Tau, pTau Z-scores A/T classification Raw data Platform Information | Mapped AD Phenotypes (NIA-AA ABC score, Thal phasing, Braak staging, CERAD score) Cerebral amyloid angiopathy, lewy bodies, vascular brain injury, TDP- 43 proteinopathy and hippocampal sclerosis | •Vascular Composite (Hypertension, diabetes, heart disease [myocardial infarct, congestive heart failure, etc.], BMI, height, weight, stroke, smoking, medication, blood pressure, CVR score) |

| Imaging – DTI | Imaging – FLAIR | Imaging – MRI | Imaging – PET |
|---|-------------------|---|---|
| PreQual Processed Data (Image-Based BIDS) Free Water Corrected Metrics Scanner Harmonization Multi-Site Calibration FA and MD Scalars | •Total WMH Volume | Preprocessed Imaging Collection MUSE Regions of Interest FreeSurfer Regions of Interest SPARE-AD Scores AD Signature (Schwartz) | Preprocessed Imaging Collection Global SUVRs Centilloid Values GMM Probabilities Amyloid Status |



Courtesy Tim Hohman / Vanderbilt: timothy.j.hohman@vumc.org

ADSP 2023 Deliverables – NG00067v11 Phenotype Harmonization Consortium (PHC)

Included in ng00067.v11 Harmonization in Progress; to be included in future release Domain data not available for harmonization

| Cohorts | Autopsy | Cardiovascular Risk Factors | Cognition | Fluid Biomarkers | Imaging – DTI | Imaging – FLAIR | Imaging – PET | Imaging – T1 |
|---------------------|--------------|--------------------------------|--------------|---------------------|---------------|-----------------|---------------|--------------|
| A4 | N/A | | \sim | N/A | N/A | \checkmark | \sim | |
| ACT | \checkmark | | \checkmark | N/A | | | N/A | |
| ADNI | N/A | \sim | \checkmark | \checkmark | \sim | | \sim | \sim |
| ADFBS (NIA-LOAD) | \checkmark | \checkmark | \checkmark | N/A | N/A | N/A | N/A | N/A |
| EFIGA | | \sim | \sim | N/A | N/A | N/A | N/A | N/A |
| Knight ADRC | | | \checkmark | \checkmark | | | | |
| NACC | \checkmark | \checkmark | \checkmark | \checkmark | \checkmark | \checkmark | \checkmark | \checkmark |
| ROS/MAP/MARS | \checkmark | \checkmark | \checkmark | N/A | \checkmark | | N/A | |
| WHICAP | | | \sim | N/A | | \sim | | \sim |
| UPenn | | | | \checkmark | | | | |
| WHICAP | | \sim | \sim | | | \sim | N/A | |
| WRAP | N/A | \checkmark | \checkmark | | \sim | | \checkmark | \sim |

ADSP 2024 Deliverables R5 WGS Release



| ADSP Release | Primary Ancestry/Ethnicity | Cases | Controls | Other | Totals |
|-------------------|-------------------------------|--------|----------|-------|--------|
| GCAD Release 5 | Black/African American | 2,646 | 5,608 | 777 | 9,031 |
| | Asian | 1,273 | 4,083 | 5 | 5,361 |
| | Latino/Hispanic | 4,488 | 10,248 | 2,777 | 17,513 |
| | Non-Hispanic White | 9,124 | 14,459 | 6,067 | 29,650 |
| | Total | 17,531 | 34,398 | 9,626 | 61,555 |

ADRC samples in R5

4,590

EXOMES 22% of total ADSP WES

12,629

genomes 20% of total ADSP WGS

ADRCs can get WGS/WES data from their own samples from us! Send an email to <u>niagads@pennmedicine.upenn.edu</u> to initiate.

ADRC GWAS Data Release



ADC 1–15 GWAS available via DSS

- QC'd PLINK files on GRCh38
- Updated ID schemas for seamless mapping of phenotypes to genotypes
- TOPMedR2 imputed files in .bgen format
- Curated phenotypes provided by the ADSP-PHC
- ADSP to NACC ID mapping file for any ADRC participants that also have WGS, WES, or ADSP-PHC harmonized phenotypes available through the ADSP dataset (ng00067)

ADC 16 & 17 GWAS

available to the ADRCs

- Raw array data available
- Total of 2,292 participants from 38 ADRCs

GWAS SNP Array Data for

29,676

ADRC Samples Passing QC and available for request.

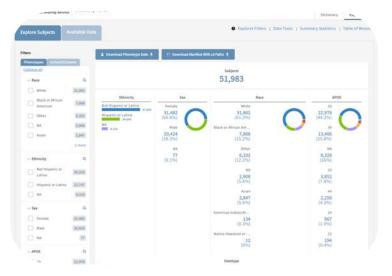
Supporting FAIR Principles



| Initiatives | Findable | Accessible | Interoperable | Reusable |
|---------------------------|----------|------------|---------------|----------|
| Gen3 Platform | Q | | \leftarrow | |
| RAS Authentication | | | \leftarrow | |
| Data Submission Portal | Q | | | |
| DOI Implementation | Q | | | |
| ADGC GWAS Standardization | | | | |
| Outreach Initiatives | Q | | | |

Supporting FAIR Principles







're-approval et preliminary details from submitter

ssign data intake manager for assessmer

stake data manger communicates approva

tatus over email and updates submission

tatus using the portal API

status

Experiment Details Upload & Validation user-dashboard to track submission templates provided for metadata files (e.g., subject phenotypes, file manifests)

documented, workflow-driven forms metadata files and forms validate guiding users from study -> dataset -> automatically at submission xperiment details



ADSP Data Discovery Platform Integration powered by Gen3 Technology

Streamline data submission and validation using **DSS Data Submission Portal**

Digital Object Identifiers (DOI) Enabled through NIH DataCite **Consortium** membership

NIAGADS Open Access Data and Annotation Resources

unrestricted data and annotations (variant, gene, sequence) available for immediate access



Open Access Portal



& advp²

GenomicsDB³

direct data downloads for GWAS summary statistics (p-values) and ADSP Variant Annotations

https://dss.niagads.org

75,207

standardized, searchable functional genomics tracks

https://tf.lisanwanglab.org/FILER/

curated AD-GWAS catalog, documenting

>18k

published associations

https://advp.niagads.org

AD-genetics knowledgebase and Genome Browser, containing

>260M

annotated variants

https://www.niagads.org/genomics



1. Kuksa et al. 2022, NAR Genomics and Bioinformatics, 4(1): lqab123. 2. Kuksa et al. 2021, Journal of Alzheimer's Disease, 86(1): 461-477. 3. Greenfest-Allen et al. 20204, Alzheimer's & Dementia, 20(2):1123-1136

NIAGADS Open Access NIAGADS API*

Lead: Emily Greenfest-Allen



filer

Alzheimer's Disease Variant Portal





Open Access Portal

GenomicsDB

Alzheimer's Genomics Database





- Programmatic access
- Bulk, ranged, and genome wide queries
- Integration into analysis pipelines
- Better support ADSP activities and AI/ML pipelines

conforms to OpenAPI standards (& is FAIR)

* under development; beta-testing version available summer 2024

Supporting Data Use

User Support

NIAGADS offers several support resources launched over the last year:



Held at least once a quarter

https://www.youtube.com/@niagads1814



All NIAGADS documentation in one place, new informational pages to address user questions, dedicated FAQ page



Users can now submit a ticket to work on resolving technical issues with the NIAGADS team.

Awareness

Expanding NIAGADS reach through increased investigator awareness will be accomplished through informational outreach sessions at AAIC and ASHG in 2024



Exhibit Hall Booth

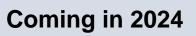
Visit us at booth 1329D at AAIC!



Poster Presentations



ADRC workshop at AAIC in partnership with NACC, Sage, and NCRAD



- Monthly newsletter
- More video tutorials, user support pages, blog posts



AAIC 2024 Activities



AAIC Exhibit Hall Booth Visit us at booth 1329D at AAIC



See you this July in Philadelphia!

