New Look of NIAGADS

The NIAGADS (NIA Genetics of Alzheimer's Disease Data Storage Site) website has been overhauled to enhance the user experience. NIAGADS is a national repository that facilitates access to genetic, genomic, and related data to qualified investigators for the study of Alzheimer's disease (AD).

https://www.niagads.org

Third Batch of ADSP Data Coming Soon!

The third ADSP data release is anticipated for October and will contain whole exome sequence data from ~11,000 additional subjects. Approved investigators can browse and download available data through the <u>ADSP Data Portal</u> or through the <u>dbGaP Study page</u>.

https://www.niagads.org/adsp

Most Recent Datasets Added to NIAGADS

NG00035

- •"GWAS of cerebrospinal fluid tau levels identifies risk variants for Alzheimer's disease"
- •Largest GWAS for CSF tau/ptau levels published to date (n = 1,269).
- •Imputed data consists of 5,815,690 SNPS using HapMap release 22 CEU (build 36) as a reference panel.
- www.niagads.org/dataset/ng00035

NG00034

- ACT AND GENETIC DIFFERENCES GWAS
- •The Adult Changes in Thought (ACT) study is a longitudinal prospective cohort study and the Genetic Differences study was an epidemiologic case control study.
- www.niagads.org/dataset/ng00034

NG00033

- IDENTIFYING RARE VARIANTS THAT INCREASE RISK FOR ALZHEIMER'S DISEASE
- •Exome sequencing data for families with Late-Onset AD.
- www.niagads.org/dataset/ng00033

Request Access to NIAGADS Data

To access NIAGADS data, please submit a new account request through the NIAGADS website. If you would like further information about data available at NIAGADS or how to submit your own data to NIAGADS, contact support@niagads.org.



The National Institute on Aging Genetics of Alzheimer's Disease Data Storage Site

NIAGADS Genomics Database

The NIAGADS Genomics Database provides a simple, but powerful, workspace for searching and identifying genes, SNPs, and genomics locations of interest or with special relevance to Alzheimer's Disease. New AD risk variants identified in a meta-analysis of 74,046 individuals can now be viewed on the genome browser via www.niagads.org/genomics/track_highlights.jsp. See www.niagads.org/genomics/track_listing.jsp for newly added functional genomic tracks, e.g. FANTOM5, GWAS catalog. Users can now find genes based on KEGG pathway. Register with NIAGADS to bookmark favorite search results or save and share search results and workflows (strategies).

https://www.niagads.org/genomics/



