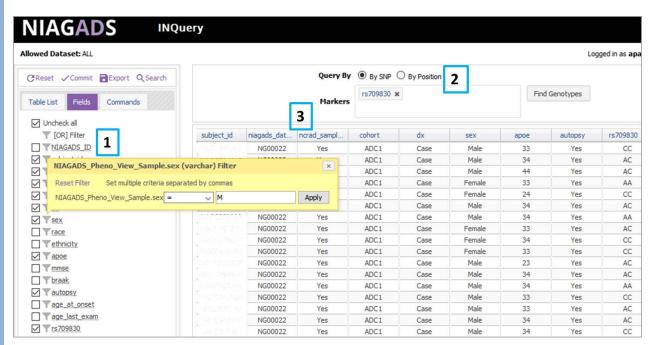
Coming Soon: INQuery

INQuery is a web-based user interface enabling access of data including phenotypes and genotypes associated with GWAS studies to qualified investigators. Harmonized phenotypes and genotypes (genome build hg37) will be available for all GWAS datasets housed in NIAGADS. In addition, information on the availability of a sample stored at NCRAD for request is provided. The database enables users to query a set of samples based on a particular phenotype/genotype of interest to determine if there is a sample available at NCRAD for further analysis. We will release INQuery by the fall of 2017.

Query for a set of samples based on genotype and phenotype:

- 1. Filter for subjects based on phenotype
- 2. Search for genotypes
- 3. See if NCRAD has a sample available for the subjects of interest



New Datasets available at https://www.niagads.org/datasets

NG00053: IGAP Summary Statistics, ADGC Subset— Lambert et al. (2013)

PI: Dr. Gerard Schellenberg, U of Pennsylvania These data are summary statistics generated for the International Genomics of Alzheimer's Project (IGAP) meta-analysis of Genome-Wise Association data in Alzheimer's Disease. This dataset contains the ADGC subset of the data analyzed in Stage 1 of the metaanalysis. NG00056: Transethnic GWAS for AD Summary Statistics—Jun et al. (2017)

PI: Dr. Lindsay Farrer, Boston University
These data are summary statistics generated from a
transethnic GWAS analysis obtained from cohorts
including European, African American, Japanese, and
Israeli Arab Ancestries.