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| Submitter Contact Information | | |
| Name | Institution | Email |
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| Study Name  Example: Large scale pQTL and mQTL atlas in brain and CSF  <https://dss.niagads.org/studies/sa000031/> |
| We recommend using the title of the associated publication or a descriptive sentence about the overall goal of the experiments being conducted.  If you would like to add a dataset to an existing study in the DSS, please provide the accession number. |
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| Study Description  Describe your study and the overall goal of the experiments being conducted for the data being submitted.  Example: Large scale pQTL and mQTL atlas in brain and CSF  <https://dss.niagads.org/studies/sa000031/> |
| We recommend including (1) Background/History (2) Study Inclusion/Exclusion Criteria (3) Population (4) Disease type (5) Study hypothesis (6) Autopsy availability (if applicable) |
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| Derived/Secondary Data Return  If the data you are returning is derived from another dataset in NIAGADS, please provide the dataset accession number and release version.  Example: ADSP Umbrella – NG00067v11 |
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| Dataset Name (if different than the study)  Example: Large scale genetics study on CSF and brain metabolite levels<https://dss.niagads.org/datasets/ng00131/> |
| We recommend a title that describes what is being analyzed in the dataset. |
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| Dataset Description  Example: Large scale genetics study on CSF and brain metabolite levels<https://dss.niagads.org/datasets/ng00131/> |
| We recommend providing as much detail about your dataset as possible. I.e., what would be your elevator pitch to another researcher if they were interested in using your data in their research. |
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| Cohort Information  Supply a list of the participating cohorts and descriptions for each. Not required for submission of analysis result. Please add a new set of rows to enter information for each cohort. | |
| Cohort Name |  |
| Cohort Abbreviation |  |
| Cohort Description |  |
| Recruiting Countries |  |

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| Data Type  Check any that apply: |  |  |
| General  Individual Phenotype  Individual Genotype  Individual Sequencing  Supporting Documents  Proteomic/Metabolomic  Images | **Array Data**  SNP Array  Expression Array  Methylation Array | **Genotypes**  Array derived Genotypes  CNV calls from microarray  CNV calls derived from Sequencing  Genotype calls derived from Sequence |
| Sequencing  Whole Genome  Whole Exome  Targeted Genome  Targeted Exome  Whole Transcriptome  Targeted Transcriptome  Sanger | **Analyses**  Association/Linkage Results  Array derived Expression  RNA Seq derived Expression  Array derived Methylation | |

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| Genome Build |
| GRCh37  GRCh38 |

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| Grant Funding |
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| --- |
| NIH Sponsoring Institute |
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| --- | --- |
| Principal Investigators and Institutions (to be displayed on website) | |
| Name | Affiliation or Institute |
| 1. |  |
| 2. |  |
| 3. |  |

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| Acknowledgment |
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| Related Publications  Provide either Pubmed ID or title, first author and publication date |
| 1. |
| 2. |
| 3. |

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| Website if applicable |
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| Logo if applicable, or URL to link the logo |
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