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| Submitter Contact Information |
| Name | Institution | Email |
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| Study NameExample: 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 DescriptionDescribe 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 ReturnIf 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 InformationSupply 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 |
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| 2. |  |
| 3. |  |

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| Acknowledgment |
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| Related PublicationsProvide 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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