

## PsychENCODE Consortium Data

The PsychENCODE Consortium (<http://www.synapse.org/pec>) has a collective goal of accelerating discovery of non-coding functional genomic elements in human brain, and elucidating their role in the molecular pathophysiology of psychiatric disorders. The PEC is producing a public resource of multi-dimensional genomic data using unbiased genome-wide approaches on tissue and cell-type specific samples from approximately 1000 phenotypically well-characterized healthy and diseased human post-mortem brains. The PEC also shares data from functional characterization of disease-associated regulatory elements and variants in *in-vitro* and *in-vivo* model systems.

The PsychENCODE investigators are committed to the release of data and results with the anticipation that data shared in a rapid and transparent manner will speed the pace of research to the benefit of both the PsychENCODE teams and the greater research community.

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The PsychENCODE Consortium data consists of a set of human studies controlled through this distribution, and animal models provided as open access. All data is distributed through the Sage Bionetworks Synapse system (<http://www.synapse.org/pec>). To download the human data, applicants must obtain a Synapse data access token before submitting an application to NIMH RGR. See instructions of how to obtain the token and how to proceed with the NIMH data access application here. (<https://www.synapse.org/#!/Synapse:syn4921369/wiki/390660>).