

**Example Plan addressing Key Elements for a Data Sharing Plan under NIH Extramural Support**  
(For questions, contact the NIH Office of Extramural Research (OER), Email [Sharing@nih.gov](mailto:Sharing@nih.gov))

The NIH [Example Plan \(link is external\)doc](#) for a template you can modify to fit the data you plan to share.

**Example Data Sharing Plan for FOA-XX-XXXX**

**What data that will be shared:**

I will share phenotypic data associated with the collected samples by depositing these data at \_\_\_\_\_ which is an NIH-funded repository. Genotype data will be shared by depositing these data at \_\_\_\_\_. Additional data documentation and de-identified data will be deposited for sharing along with phenotypic data, which includes demographics, family history of XXXXXX disease, and diagnosis, consistent with applicable laws and regulations. I will comply with the NIH GWAS Policy and the funding IC's existing policies on sharing data on XXXXXX disease genetics to include secondary analysis of data resulting from a genome wide association study through the repository. Meta-analysis data and associated phenotypic data, along with data content, format, and organization, will be available at \_\_\_\_\_. Submitted data will confirm with relevant data and terminology standards.

**Who will have access to the data:**

I agree that data will be deposited and made available through \_\_\_\_\_ which is an NIH-funded repository, and that these data will be shared with investigators working under an institution with a Federal Wide Assurance (FWA) and could be used for secondary study purposes such as finding genes that contribute to process of XXXXXX. I agree that the names and Institutions of persons either given or denied access to the data, and the bases for such decisions, will be summarized in the annual progress report. Meta-analysis data and associated phenotypic data, along with data content, format, and organization, will be made available to investigators through \_\_\_\_\_.

**Where will the data be available:**

I agree to deposit and maintain the phenotypic data, and secondary analysis of data (if any) at \_\_\_\_\_, which is an NIH-funded repository and that the repository has data access policies and procedures consistent with NIH data sharing policies.

**When will the data be shared:**

I agree to deposit genetic outcome data into \_\_\_\_\_ repository as soon as possible but no later than within one year of the completion of the funded project period for the parent award or upon acceptance of the data for publication, or public disclosure of a submitted patent application, whichever is earlier.

**How will researchers locate and access the data:**

I agree that I will identify where the data will be available and how to access the data in any publications and presentations that I author or co-author about these data, as well as acknowledge the repository and funding source in any publications and presentations. As I will be using \_\_\_\_\_, which is an NIH-funded repository, this repository has policies and procedures in place that will provide data access to qualified researchers, fully consistent with NIH data sharing policies and applicable laws and regulations.