23andMe is committed to supporting genetic research by enabling access to data we have included in our publications. We provide access to full summary statistics through a Data Transfer Agreement that protects the privacy of our participants’ data and a Statement of Work that describes how the data will be used.

**DATASET REQUEST PROCESS**

1. **Prepare and submit your online request**
2. **Review and approve your request**
3. **Receive the SOW/DTA with your project details and dataset request**
4. **Have your institution’s contracts office review and sign the SOW/DTA**
5. **Sign the SOW and share both the SOW/DTA with 23andMe**
6. **Fully execute the SOW/DTA**
7. **Provide information on how to access your requested GWAS**
8. **Share any publication using data with 23andMe for review**
9. **Publish (if applicable)**

**ITEMS NEEDED TO COMPLETE THE REQUEST FORM**

- Please compile the following in advance of submitting the dataset request form:
  - The GWAS you would like to request
  - A project proposal including:
    1. Title
    2. Hypotheses
    3. Non-Technical Summary
    4. Proposed Analyses
  - If the GWAS data will be used for:
    1. Polygenic Risk Scores
    2. Meta-Analysis
    3. To identify drug targets
  - Collaborator information
  - Publication plans and timeline

**TIPS FOR STREAMLINING THE REQUEST PROCESS**

- Read all of the information provided on the dataset access webpage and within the FAQs
- Communicate with your institution's contracts office early
- Inform collaborators at other institutions that they will also need to submit a request to access GWAS data

**23ANDME CONTACT INFO**

23andMe Research Team
data-request@23andMe.com

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**23ANDME COLLABORATIONS**

*Research Innovations Collaborations*

Academic researchers can work with 23andMe to study de-identified, aggregated data from the 23andMe Research Cohort of millions of participants who have answered survey questions on wide-ranging topics.

*Populations Collaborations*

This program aims to expand and improve our Reference Data Panel through capturing the genetic diversity of populations around the world who are underrepresented in our database and in genetics research globally.