

# Biobank Sample Request Form



## Instructions for Submitting Sample Requests

1. Fill out this form with all fields completed.
2. Email the completed form to [gtex-biobank@broadinstitute.zendesk.com](mailto:gtex-biobank@broadinstitute.zendesk.com).  
If you have a list of samples that you are interested in, please include that file as well.  
\*Please use the following subject header: **GTEx Biobank Request - [Date] [Last\_Name]**
3. Note that if your request is approved, you will need to fill out a Material Transfer Agreement (MTA).

Contact us at [gtex-biobank@broadinstitute.zendesk.com](mailto:gtex-biobank@broadinstitute.zendesk.com) for assistance with sample selection.

## Request Information

GTEx sample request # (LEAVE BLANK - will be assigned)

Request date

## Requestor Information

Name

Email

Affiliation

## Shipping Contact Information

Name

Shipping address

City

State

Zip code

Email

Phone number

*Please continue to the next page and fill out details about your project.*

## Project Information

Project title

### Research use statement

Describe the proposed use of the biospecimens, including goals of the study, justification for using GTEx specimens over other sources, molecular assay details, statistical analysis plans, etc. Please describe your research in enough detail that it can be evaluated by our scientific committee.

*Please continue to the next page and fill out further details about your project.*

## Project Information (cont.)

### Sample selection

Describe how GTEx samples were, or will be, selected for the study (e.g. tissues of interest, those with associated RNA-seq or other molecular data, etc).

### Sample amount

Specify the amount of material you would like for each material type. Indicate the desired amount of tissue mass (mg), and DNA/RNA amount (ng) and preferred concentration (ng/ $\mu$ L). Please specify if there is a minimum amount or concentration cut-off.

### Funding source(s)

Please indicate the source of funds to be used to carry out the work.

### Request for NIH grant application

- |   |     |    |
|---|-----|----|
| <input type="radio"/> Is your request for an NIH grant application you plan to submit?  | Yes | No |
| <input type="radio"/> Are you requesting a preliminary review/approval of your request to submit with your grant application? | Yes | No |

### Data sharing plans

Describe how you plan to share the data and resources that will be generated.

PLEASE NOTE: As stated in the MTA, ALL protected access data (e.g. all raw genotype, RNA-seq, or other sequence data) MUST be submitted to a protected access database, such as dbGaP, using the sample IDs that will be provided to you. Non-protected, derived data may be shared in a non-protected access database. In some circumstances, the GTEx portal may be able to host your data – please indicate if you plan to reach out to the GTEx portal about hosting such data, and/or describe where else you will host/share such data. Please include an estimated timeline for all data sharing.

*Please continue to the next page and fill out further details about your project.*

## Project Information (cont.)

### Project summary

Please provide a one paragraph summary of your work to be shared on the GTEx portal website. Your request will be made public only when approved and the samples have been sent.

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