



Research Dissemination and Acknowledgement of NIH Genomic Study Datasets

Excerpted from Data Use Certification Agreement:

https://dbgap.ncbi.nlm.nih.gov/aa/wga.cgi?view_pdf&stacc=phs000424.v1.p1

It is the intent of the NIH to promote the dissemination of research findings from NIH genomic dataset(s) as widely as possible through scientific publication or other appropriate public dissemination mechanisms. NIH intends that GTEx genomic data are released as rapidly as possible after they are produced, with no restrictions on use. Approved Users are strongly encouraged to publish their results in peer-reviewed journals and to present research findings at scientific meetings, etc. There is no formal Embargo Date associated with this dataset. However, users are asked to apply the normal standards of scientific etiquette when considering a publication based substantially on unpublished GTEx data. Additional information on GTEx Consortium publication plans and policy are available at <http://gtexportal.org> and by contacting NIH Program staff at nhgrigtex@mail.nih.gov. Approved Users agree to acknowledge GTEx and the NIH data repository, the Contributing Investigator(s) who contributed the phenotype data and DNA samples from his/her original study, and the primary funding organization that supported the contributing study in all oral and written presentations, disclosures, and publications resulting from any analyses of the data. Approved Users further agree that the acknowledgment shall include the dbGaP accession number to the specific version of the dataset(s) analyzed.

A sample statement for the acknowledgment of the Genotype-Tissue Expression (GTEx) Project dataset(s) follows:

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