

Sanger Imputation Service

This is a free genotype imputation service provided by the [Wellcome Trust Sanger Institute](#). You can upload GWAS data in VCF or 23andMe format and receive imputed genomes back. Click [here](#) to learn more.

Before you start

Be sure to [read through the instructions](#).

You will need to set up a free account with [Globus](#) and have [Globus Connect](#) running at your institute or on your computer to transfer files to and from the service.

Ready to start?

If you are ready to upload your data, please fill in the details below to **register an imputation job**. If you need more information, see the [about](#) page.

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News

24/04/2015

The HRC.r1 panel is now available to HRC consortium members.

21/04/2015

Added page to find out the [status](#) of your imputation job.

08/04/2015

Soft launch. Please help us improve the documentation on the site by sending us [feedback](#).

[See older news...](#)

University of Michigan Imputation Server

This server provides a free genotype imputation service. You can upload GWAS genotypes and receive imputed genomes in return. Our server offers imputation from HapMap, 1000 Genomes (Phase 1 and Phase 3), and [HRC](#).



1. Upload your genotypes

Upload your data to our server. All interactions with the server are secured.

News

24. April 2015

- [HRC](#) release 1 (64,976 haplotypes at 39,235,157 SNPs) is now ready for use for HRC consortium members

23. March 2015

- New Fair Use Policy: please contact us if you want to impute >6,000 samples

Statistics

Chromosomes 11,794,743

Genomes 536,125

Server Status

145,244 Chromosomes are currently in progress.

0 Chromosomes are in the queue.