Genotype Imputation (Minimac4) 1.7.1

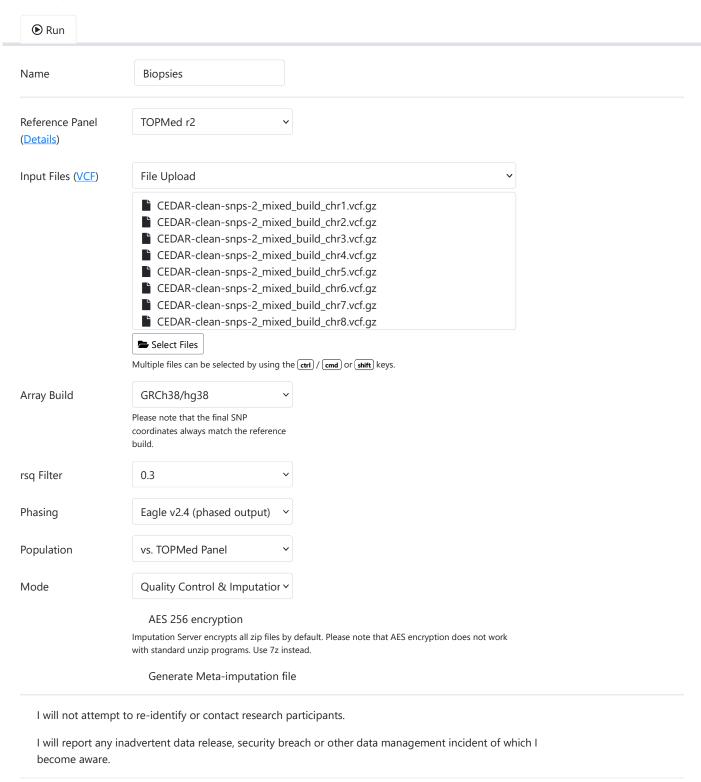
Submit Job

This is the new Michigan Imputation Server Pipeline using Minimac4. Documentation can be found here.

If your input data is **GRCh37/hg19** please ensure chromosomes are encoded without prefix (e.g. **20**).

If your input data is **GRCh38hg38** please ensure chromosomes are encoded with prefix 'chr' (e.g. **chr20**).

Mttps://imputationserver.readthedocs.io



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Back to top