

# Resource Summary Report

Generated by [NIF](#) on May 16, 2025

## TAGIMPUTE

RRID:SCR\_013338

Type: Tool

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### Proper Citation

TAGIMPUTE (RRID:SCR\_013338)

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### Resource Information

**URL:** <http://www.bios.unc.edu/~lin/software/tagIMPUTE/>

**Proper Citation:** TAGIMPUTE (RRID:SCR\_013338)

**Description:** A command-line program for the imputation of untyped SNPs. tagIMPUTE is based on a few flanking SNPs that can optimally predict the SNP under imputation. (entry from Genetic Analysis Software)

**Synonyms:** TAGging-snp based IMPUTATE

**Resource Type:** software resource, software application

**Keywords:** gene, genetic, genomic

**Funding:**

**Resource Name:** TAGIMPUTE

**Resource ID:** SCR\_013338

**Alternate IDs:** nlx\_154670

**Record Creation Time:** 20220129T080315+0000

**Record Last Update:** 20250513T061438+0000

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### Ratings and Alerts

No rating or validation information has been found for TAGIMPUTE.

No alerts have been found for TAGIMPUTE.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.