

# Resource Summary Report

Generated by [Kravitz](#) on Sep 14, 2026

## ChromImpute

RRID:SCR\_023990

Type: Tool

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### Proper Citation

ChromImpute (RRID:SCR\_023990)

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### Resource Information

**URL:** <http://www.biolchem.ucla.edu/labs/ernst/ChromImpute/>

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**Description:** Software tool for large scale systematic epigenome imputation. ChromImpute takes existing compendium of epigenomic data and uses it to predict signal tracks for mark-sample combinations not experimentally mapped or to generate a potentially more robust version of data sets that have been mapped experimentally.

**Synonyms:** chromimpute

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

**Keywords:** systematic epigenome imputation, epigenomic data, predict signal tracks, mark-sample combinations, not experimentally mapped data,

**Availability:** Free, Available for download, Freely available

**Resource Name:** ChromImpute

**Resource ID:** SCR\_023990

**Alternate IDs:** OMICS\_11185

**Alternate URLs:** <https://sources.debian.org/src/chromimpute/>

**Record Creation Time:** 20230824T050211+0000

**Record Last Update:** 20260912T200044+0000

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### Ratings and Alerts

No rating or validation information has been found for ChromImpute.

No alerts have been found for ChromImpute.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [Kravitz](#) .

Welzenbach J, et al. (2021) Integrative approaches generate insights into the architecture of non-syndromic cleft lip with or without cleft palate. HGG advances, 2(3), 100038.