

Resource Summary Report

Generated by [NIF](#) on Aug 22, 2026

ChromImpute

RRID:SCR_023990

Type: Tool

Proper Citation

ChromImpute (RRID:SCR_023990)

Resource Information

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

Proper Citation: ChromImpute (RRID:SCR_023990)

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: 20260821T194253+0000

Ratings and Alerts

No rating or validation information has been found for ChromImpute.

No alerts have been found for ChromImpute.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

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.