Resource Summary Report

Generated by PRECISE-TBI on Sep 17, 2026

ChromImpute

RRID:SCR_023990
Type: Tool

Proper Citation

ChromImpute (RRID:SCR_023990)

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

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 PRECISE-TBI .

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.