

# Resource Summary Report

Generated by ASWG on Aug 4, 2026

## WHICHRUN

RRID:SCR\_009064

Type: Tool

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### Proper Citation

WHICHRUN (RRID:SCR\_009064)

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### Resource Information

**URL:** <https://github.com/gaow/genetic-analysis-software/blob/master/pages/WHICHRUN.md>

**Proper Citation:** WHICHRUN (RRID:SCR\_009064)

**Description:** THIS RESOURCE IS NO LONGER IN SERVICE, documented September 22, 2016. Software application that uses multilocus genotypic data to allocate individuals to their most likely source population.

**Abbreviations:** WHICHRUN

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

**Keywords:** gene, genetic, genomic, c++, ms-windows, (95/98/nt)

**Availability:** THIS RESOURCE IS NO LONGER IN SERVICE

**Resource Name:** WHICHRUN

**Resource ID:** SCR\_009064

**Alternate IDs:** nlx\_154051

**Old URLs:** <http://www.bml.ucdavis.edu/whichrun.htm>

**Record Creation Time:** 20220129T080250+0000

**Record Last Update:** 20260801T191043+0000

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### Ratings and Alerts

No rating or validation information has been found for WHICHRUN.

No alerts have been found for WHICHRUN.

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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.