### **Resource Summary Report**

Generated by FDI Lab - SciCrunch.org on Apr 14, 2025

## **WHICHRUN**

RRID:SCR\_009064

Type: Tool

### **Proper Citation**

WHICHRUN (RRID:SCR\_009064)

#### **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 SERVCE, 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 application, software resource

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

**Funding:** 

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: 20250412T055338+0000

### Ratings and Alerts

No rating or validation information has been found for WHICHRUN.

No alerts have been found for WHICHRUN.

### Data and Source Information

Source: SciCrunch Registry

# Usage and Citation Metrics

We have not found any literature mentions for this resource.