# **Resource Summary Report**

Generated by FDI Lab - SciCrunch.org on Apr 11, 2025

# **HOTSPOTTER**

RRID:SCR\_009238

Type: Tool

### **Proper Citation**

HOTSPOTTER (RRID:SCR\_009238)

#### **Resource Information**

**URL:** http://www.biostat.umn.edu/~nali/SoftwareListing.html

**Proper Citation:** HOTSPOTTER (RRID:SCR\_009238)

**Description:** Software application that implements the PAC (Products of Approximate Conditional) model for estimating recombination rate (entry from Genetic Analysis Software)

**Abbreviations: HOTSPOTTER** 

Resource Type: software resource, software application

Keywords: gene, genetic, genomic, c++

Funding:

**Resource Name: HOTSPOTTER** 

Resource ID: SCR\_009238

Alternate IDs: nlx\_154399

**Record Creation Time:** 20220129T080251+0000

**Record Last Update:** 20250411T055306+0000

### **Ratings and Alerts**

No rating or validation information has been found for HOTSPOTTER.

No alerts have been found for HOTSPOTTER.

### **Data and Source Information**

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 2 mentions in open access literature.

**Listed below are recent publications.** The full list is available at FDI Lab - SciCrunch.org.

Staufer M, et al. (2023) Temporal and spatial variations in local sex ratios in a suburban population of the European green toad Bufotes viridis. BMC ecology and evolution, 23(1), 6.

Nepovinnykh E, et al. (2022) SealID: Saimaa Ringed Seal Re-Identification Dataset. Sensors (Basel, Switzerland), 22(19).