

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

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), THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 16,2025.

Abbreviations: HOTSPOTTER

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, c++

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: HOTSPOTTER

Resource ID: SCR_009238

Alternate IDs: nlx_154399

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260727T040446+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 3 mentions in open access literature.

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

Biffi D, et al. (2025) Identification of Individual Texas Horned Lizards (*Phrynosoma cornutum*) Using Genotypes and Ventral Spot Patterns. *Ecology and evolution*, 15(3), e71167.

Stauffer M, et al. (2023) Temporal and spatial variations in local sex ratios in a suburban population of the European green toad *Bufo 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).