

Resource Summary Report

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BOTTLENECK

RRID:SCR_009082

Type: Tool

Proper Citation

BOTTLENECK (RRID:SCR_009082)

Resource Information

URL: <http://www.montpellier.inra.fr/URLB/bottleneck/bottleneck.html>

Proper Citation: BOTTLENECK (RRID:SCR_009082)

Description: Software application for detecting recent effective population size reductions from allele data frequencies (entry from Genetic Analysis Software)

Abbreviations: BOTTLENECK

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, ms-windows, (95)

Resource Name: BOTTLENECK

Resource ID: SCR_009082

Alternate IDs: nlx_154093

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260905T133243+0000

Ratings and Alerts

No rating or validation information has been found for BOTTLENECK.

No alerts have been found for BOTTLENECK.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 493 mentions in open access literature.

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

Marín JC, et al. (2026) Genetic Status of the Northernmost Population of the Endangered and Elusive Huemul Deer, *Hippocamelus bisulcus*. *Animals : an open access journal from MDPI*, 16(11).

Keshavmurthy S, et al. (2026) Microgeographic differentiation in the genetic structure of catch bowl coral, *Isopora cf. palifera* (Scleractinia; Acroporidae), in Kenting National Park, Taiwan. *Scientific reports*, 16(1).

Jeong D, et al. (2026) Population genetic structure of the gold-spotted pond frog (*Pelophylax chosenicus*) and its phylogenetic context within the genus *Pelophylax*. *Scientific reports*, 16(1), 4470.

Islam KMT, et al. (2026) SIRT7 as a context-dependent biomarker and therapeutic target: Insights from a pan-cancer study. *PloS one*, 21(2), e0342269.

Meléndez-Cal-Y-Mayor JF, et al. (2026) Translocated populations are genetically similar to natural populations and populations resulting from natural colonizations. *PloS one*, 21(1), e0340580.

Lin Y, et al. (2025) Potential common mechanisms between primary Sjögren's syndrome and Hashimoto's thyroiditis: a public databases-based study. *Frontiers in genetics*, 16, 1520332.

Liu F, et al. (2025) Pod-pose : an efficient top-down keypoint detection model for fine-grained pod phenotyping in mature soybean. *Plant methods*, 21(1), 82.

Lado TF, et al. (2025) Microsatellite Variation of African Elephants *Loxodonta africana* Blumenbach 1797 in a Savannah Protected Area of South Sudan. *Ecology and evolution*, 15(6), e71383.

Parreira BR, et al. (2025) Effects of Social Structure on Effective Population Size Change Estimates. *Evolutionary applications*, 18(1), e70063.

Liang H, et al. (2025) The multi-target mechanism of action of *Selaginella doederleinii* Hieron in the treatment of nasopharyngeal carcinoma: a network pharmacology and multi-omics analysis. *Scientific reports*, 15(1), 159.

Wang L, et al. (2025) Single-cell sequencing reveals the response mechanisms of vascular endothelial cells to glucocorticoids in diabetic retinopathy. *PloS one*, 20(11), e0334230.

Song Z, et al. (2025) High-throughput end-to-end aphid honeydew excretion behavior recognition method based on rapid adaptive motion-feature fusion. *Frontiers in plant science*, 16, 1609222.

Zhao J, et al. (2025) TERT links telomere length to cancer risk by integrating genomic instability and immune modulation. *Discover oncology*, 16(1), 1788.

Kamdem NG, et al. (2025) Evidence of high genetic differentiation driven by limited gene flow in a lower canopy African tropical rainforest tree species, *Coula edulis* Baill. (Coulaceae). *BMC ecology and evolution*, 25(1), 140.

Li ZF, et al. (2025) Cellular and molecular features of acute myeloid leukemia investigation based on single-cell RNA sequencing. *Cancer cell international*, 25(1), 326.

Fan Y, et al. (2025) Identification and validation of shared key genes between Parkinson's disease and erectile dysfunction: a bioinformatics approach. *Hereditas*, 163(1), 3.

Li H, et al. (2025) Integrated Transcriptome and Metabolomics Analyses Show MYC as a Potential Therapeutic Target for Behçet's Uveitis. *Advanced science (Weinheim, Baden-Wurttemberg, Germany)*, 12(36), e17843.

Luo B, et al. (2025) Hippo pathway and NLRP3-driven NETosis in macrophages: Mechanisms of viral pneumonia aggravation. *Cell death discovery*, 11(1), 323.

Mao L, et al. (2025) Identification and Validation of Key Genes Involved in the Coupling of Mitochondria-Associated Endoplasmic Reticulum Membrane in Hemorrhoidal Disease. *International journal of general medicine*, 18, 2781.

Lehoczky I, et al. (2025) Genetic survey of crucian carp *Carassius carassius* populations in Hungary for a conservation project to establish live gene bank. *Scientific reports*, 15(1), 8835.