# **Resource Summary Report**

Generated by FDI Lab - SciCrunch.org on Apr 12, 2025

# **HARSH**

RRID:SCR\_010792

Type: Tool

### **Proper Citation**

HARSH (RRID:SCR\_010792)

#### **Resource Information**

URL: http://genetics.cs.ucla.edu/harsh/

**Proper Citation:** HARSH (RRID:SCR\_010792)

Description: Software that provides a method to infer the haplotype using haplotype

reference panel and high throughput sequencing data.

**Abbreviations:** HARSH

**Synonyms:** HAplotype inference using Reference and Sequencing tecHnology

**Resource Type:** software resource

Keywords: bio.tools

**Funding:** 

Resource Name: HARSH

Resource ID: SCR\_010792

Alternate IDs: OMICS\_00199, biotools:harsh

Alternate URLs: https://bio.tools/harsh

**Record Creation Time:** 20220129T080300+0000

**Record Last Update:** 20250410T070021+0000

### Ratings and Alerts

No rating or validation information has been found for HARSH.

No alerts have been found for HARSH.

#### Data and Source Information

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 14 mentions in open access literature.

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

Pope-Caldwell S, et al. (2024) Variability and harshness shape flexible strategy-use in support of the constrained flexibility framework. Scientific reports, 14(1), 7236.

Guo J, et al. (2024) Response of the gut microbiota to changes in the nutritional status of red deer during winter. Scientific reports, 14(1), 24961.

Lee YB, et al. (2023) Sub-10 fJ/bit radiation-hard nanoelectromechanical non-volatile memory. Nature communications, 14(1), 460.

Li M, et al. (2023) The relationship between harsh parenting and adolescent depression. Scientific reports, 13(1), 20647.

Chen L, et al. (2021) Analysis of RNA conformation in endogenously assembled RNPs by icSHAPE. STAR protocols, 2(2), 100477.

Moss JB, et al. (2021) Constrained flexibility of parental cooperation limits adaptive responses to harsh conditions. Evolution; international journal of organic evolution, 75(7), 1835.

Carr JJ, et al. (2021) Staying Strong Toolbox: Co-design of a physical activity and lifestyle program for Aboriginal families with Machado-Joseph disease in the Top End of Australia. PloS one, 16(2), e0244311.

Gupta AK, et al. (2021) An efficient method to generate kidney organoids at the air-liquid interface. Journal of biological methods, 8(2), e150.

Al-Maqdi KA, et al. (2021) Challenges and Recent Advances in Enzyme-Mediated Wastewater Remediation-A Review. Nanomaterials (Basel, Switzerland), 11(11).

Chen J, et al. (2021) Diversity increases yield but reduces harvest index in crop mixtures. Nature plants, 7(7), 893.

Ullah R, et al. (2021) Understanding Variations in the Tracking and Erosion Performance of

HTV-SR-Based Composites due to AC-Stressed Aging. Polymers, 13(21).

He D, et al. (2018) Efficient algorithms for polyploid haplotype phasing. BMC genomics, 19(Suppl 2), 110.

Jiang Z, et al. (2018) An improved advertising CTR prediction approach based on the fuzzy deep neural network. PloS one, 13(5), e0190831.

Mokhtari S, et al. (2012) Mechanisms of cyst formation in metastatic lymph nodes of head and neck squamous cell carcinoma. Diagnostic pathology, 7, 6.