# Resource Summary Report

Generated by NIF on May 14, 2025

## Monster

RRID:SCR_006463
Type: Tool

### Proper Citation

Monster (RRID:SCR_006463)

### Resource Information

**URL:** http://www.monster.com/

**Proper Citation:** Monster (RRID:SCR_006463)

**Description:** Global online employment solution for people seeking jobs and the employers who need great people. They"ve been doing this for over ten years, and have expanded from their roots as a job board to a global provider of a full array of job seeking, career management, recruitment and talent management products and services.

**Abbreviations:** Monster

**Synonyms:** Monster.com

**Resource Type:** job resource

**Keywords:** career, advice, resume, community building portal

**Funding:**

**Resource Name:** Monster

**Resource ID:** SCR_006463

**Alternate IDs:** nlx_144164

**Record Creation Time:** 20220129T080236+0000

**Record Last Update:** 20250420T014329+0000

### Ratings and Alerts

No rating or validation information has been found for Monster.

No alerts have been found for Monster.

## Data and Source Information

**Source:** SciCrunch Registry

## Usage and Citation Metrics

We found 76 mentions in open access literature.

**Listed below are recent publications.** The full list is available at NIF.


Rossi N, et al. (2024) Rare variants at KCNJ2 are associated with LDL-cholesterol levels in a cross-population study. NPJ genomic medicine, 9(1), 36.

Calia G, et al. (2024) Identification and characterization of specific motifs in effector proteins of plant parasites using MOnSTER. Communications biology, 7(1), 850.

Calia G, et al. (2024) Definition of the effector landscape across 13 phytoplasma proteomes with LEAPH and EffectorComb. NAR genomics and bioinformatics, 6(3), lqae087.

Mueller SH, et al. (2023) Aggregation tests identify new gene associations with breast cancer in populations with diverse ancestry. Genome medicine, 15(1), 7.

Ben Guebila M, et al. (2023) The Network Zoo: a multilingual package for the inference and analysis of gene regulatory networks. Genome biology, 24(1), 45.

Volkmar N, et al. (2022) Regulation of membrane fluidity by RNF145-triggered degradation of the lipid hydrolase ADIPOR2. The EMBO journal, 41(19), e110777.

Seid M, et al. (2021) A science of collaborative learning health systems. Learning health systems, 5(3), e10278.

Miyabayashi H, et al. (2021) DNA Polymerase B1 Binding Protein 1 Is Important for DNA Repair by Holoenzyme PolB1 in the Extremely Thermophilic Crenarchaeon Sulfolobus acidocaldarius. Microorganisms, 9(2).

Miyabayashi H, et al. (2020) PolB1 Is Sufficient for DNA Replication and Repair Under Normal Growth Conditions in the Extremely Thermophilic Crenarchaeon Sulfolobus acidocaldarius. Frontiers in microbiology, 11, 613375.

Junghans BM, et al. (2019) Populations Norms for "SLURP"-An iPad App for Quantification of Visuomotor Coordination Testing. Frontiers in neuroscience, 13, 711.

Kember RL, et al. (2018) Genetic pleiotropy between mood disorders, metabolic, and

endocrine traits in a multigenerational pedigree. Translational psychiatry, 8(1), 218.

Gilly A, et al. (2018) Cohort-wide deep whole genome sequencing and the allelic architecture of complex traits. Nature communications, 9(1), 4674.

Lee S, et al. (2018) WISARD: workbench for integrated superfast association studies for related datasets. BMC medical genomics, 11(Suppl 2), 39.

Temmink RJM, et al. (2018) Azolla along a phosphorus gradient: biphasic growth response linked to diazotroph traits and phosphorus-induced iron chlorosis. Scientific reports, 8(1), 4451.

Zhang Q, et al. (2018) Impact of rare and low-frequency sequence variants on reliability of genomic prediction in dairy cattle. Genetics, selection, evolution : GSE, 50(1), 62.

Hanschen FS, et al. (2017) Optimizing isothiocyanate formation during enzymatic glucosinolate breakdown by adjusting pH value, temperature and dilution in Brassica vegetables and Arabidopsis thaliana. Scientific reports, 7, 40807.

Forero DA, et al. (2017) Ten simple rules for international short-term research stays. PLoS computational biology, 13(12), e1005832.

Jeschke V, et al. (2017) How Glucosinolates Affect Generalist Lepidopteran Larvae: Growth, Development and Glucosinolate Metabolism. Frontiers in plant science, 8, 1995.

Suzuki S, et al. (2017) Development of the Multiple Gene Knockout System with One-Step PCR in Thermoacidophilic Crenarchaeon Sulfolobus acidocaldarius. Archaea (Vancouver, B.C.), 2017, 7459310.

Aben RCH, et al. (2017) Cross continental increase in methane ebullition under climate change. Nature communications, 8(1), 1682.