

Resource Summary Report

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TRAP

RRID:SCR_009002

Type: Tool

Proper Citation

TRAP (RRID:SCR_009002)

Resource Information

URL: <http://research.nhgri.nih.gov/software/TRAP/>

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Description: Software tool for determining a regression model of quantitative or binary trait variation when the number of possible genetic predictors is very large, considering only a moderate number of predictors at one time, using unrelated or family data. (entry from Genetic Analysis Software)

Abbreviations: TRAP

Synonyms: Tiled Regression Analysis Package

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, r, any with r installation

Resource Name: TRAP

Resource ID: SCR_009002

Alternate IDs: nlx_154010

Record Creation Time: 20220129T080250+0000

Record Last Update: 20260829T183051+0000

Ratings and Alerts

No rating or validation information has been found for TRAP.

No alerts have been found for TRAP.

Data and Source Information

Usage and Citation Metrics

We found 397 mentions in open access literature.

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

Menden B, et al. (2026) Loss-of-function variants in the CAPN1 activator CD99L2 cause X-linked spastic ataxia. *Nature communications*, 17(1), 1698.

Swanson S, et al. (2026) Seasonal Drivers of Density in a Subarctic Population of Northern Red-Backed Voles. *Ecology and evolution*, 16(2), e73142.

Motelow JE, et al. (2026) Interstitial cystitis: a phenotype and rare variant exome sequencing study. *EBioMedicine*, 125, 106150.

Faraji A, et al. (2025) Comparative resilience and precision of digitized optical counting using ImageJ during routine mosquito (Diptera: Culicidae) sample processing. *Journal of insect science (Online)*, 25(2).

Soares RRG, et al. (2025) Pooled optical screening in bacteria using chromosomally expressed barcodes. *Communications biology*, 8(1), 851.

Pal Mahadevan V, et al. (2025) Preference for and resistance to a toxic sulfur volatile opens up a unique niche in *Drosophila busckii*. *Nature communications*, 16(1), 767.

Yu JW, et al. (2025) Digenic impairments of haploinsufficient genes in patients with craniosynostosis. *JCI insight*, 10(4).

Pauwels J, et al. (2025) A 96-Well Ultrafiltration Approach for the High-Throughput Proteome Analysis of Extracellular Vesicles Isolated From Conditioned Medium. *Journal of extracellular vesicles*, 14(7), e70103.

Sermeño-Correa C, et al. (2025) Sticky traps for *Aedes aegypti* surveillance and targeted vector control in Sincelejo, Colombia. *Biomedica : revista del Instituto Nacional de Salud*, 45(1), 118.

Daly ME, et al. (2025) Proteomic Profiling of Celiac-Toxic Motifs and Allergens in Cereals Containing Gluten. *Journal of proteome research*, 24(5), 2336.

Melián-Henríquez A, et al. (2025) Geographical distribution of potential mechanical vectors implicated in Surra transmission in Spain: an entomological perspective. *Parasites & vectors*, 18(1), 305.

Wang H, et al. (2025) Astragaloside IV alleviated bone loss in mice with ovariectomy-induced osteoporosis via modulating gut microbiota and fecal metabolism. *Frontiers in pharmacology*, 16, 1548491.

Park J, et al. (2025) ATPase-dependent duplex nucleic acid unwinding by SARS-CoV-2 nSP13 relies on facile binding and translocation along single-stranded nucleic acid. *The*

Journal of biological chemistry, 301(7), 110373.

Liu C, et al. (2025) Enhanced performance of T2SLs LWIR avalanche photodiodes with a separate Al_xGa_{1-x}Sb multiplication layer. Scientific reports, 15(1), 12043.

Giuranna J, et al. (2025) Genetic and functional analyses of CTBP2 in anorexia nervosa and body weight regulation. Molecular psychiatry, 30(5), 1836.

Sánchez-González L, et al. (2025) Incompatible *Aedes aegypti* male releases as an intervention to reduce mosquito population-A field trial in Puerto Rico. PLoS neglected tropical diseases, 19(1), e0012839.

Nametov A, et al. (2025) Stray Dogs as Reservoirs and Sources of Infectious and Parasitic Diseases in the Environment of the City of Uralsk in Western Kazakhstan. Biology, 14(6).

Mehatre SH, et al. (2025) A capsular myofibroblastic niche maintains hematopoietic stem cells in the spleen. The EMBO journal, 44(14), 3983.

Aizpurua O, et al. (2025) Functional Insights Into the Effect of Feralisation on the Gut Microbiota of Cats Worldwide. Molecular ecology, 34(6), e17695.

Martínez AS, et al. (2025) Understanding key population drivers of the spotted wing *Drosophila* in cultivated and natural areas in the Andes. Scientific reports, 15(1), 6208.