

# Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

## RACE

RRID:SCR\_010950

Type: Tool

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### Proper Citation

RACE (RRID:SCR\_010950)

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### Resource Information

**URL:** <http://race.unil.ch/>

**Proper Citation:** RACE (RRID:SCR\_010950)

**Description:** A collection of web tools designed to assist with the analysis of DNA microarray data and results. RACE performs probe level data preprocessing, quality checks, normalization, and visualization for Affymetrix GeneChips. In addition, it performs clustering and differential analysis of normalized expression levels or ratios for arbitrary platforms, and estimates the false discovery rates in lists of potentially regulated genes. A Gene Ontology (GO)-term analysis assists in the biological interpretation of gene lists. The user can customize each analysis request; upon submission the analysis is executed in a fully automated way., THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 16,2025.

**Abbreviations:** RACE

**Synonyms:** Remote Analysis Computation for gene Expression data

**Resource Type:** analysis service resource, data analysis service, production service resource, service resource

**Defining Citation:** [PMID:15980552](#)

**Keywords:** dna microarray, bio.tools

**Availability:** THIS RESOURCE IS NO LONGER IN SERVICE

**Resource Name:** RACE

**Resource ID:** SCR\_010950

**Alternate IDs:** OMICS\_00777, biotools:race

**Alternate URLs:** <https://bio.tools/race>

**Record Creation Time:** 20220129T080301+0000

**Record Last Update:** 20260905T133201+0000

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## Ratings and Alerts

No rating or validation information has been found for RACE.

No alerts have been found for RACE.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 357 mentions in open access literature.

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

Wei C, et al. (2025) Genetic characterization and pathogenicity of two recombinant PRRSV-2 strains from lineages 1, 3, 5, and 8 emerged in China. BMC veterinary research, 21(1), 341.

Gadad SS, et al. (2025) X-Linked Cancer-Associated Polypeptide (XCP) from lncRNA1456 Cooperates with PHF8 to Regulate Gene Expression and Cellular Pathways in Breast Cancer. bioRxiv : the preprint server for biology.

Cheng SY, et al. (2025) An EGFR Co-Amplified and De Novo Long Noncoding RNA HELDR Promotes Glioblastoma Malignancy through KAT7-Driven Gene Programs. Research square.

Schneider C, et al. (2025) Functional and sequence-based comparison of Ctenocephalides felis and Rhipicephalus sanguineus sensu lato isolates from different geographic regions. Parasites & vectors, 18(1), 170.

Tang X, et al. (2025) SLC27A5 inhibits cancer stem cells by inducing alternative polyadenylation of METTL14 in hepatocellular carcinoma. Genes & diseases, 12(4), 101488.

Pollak C, et al. (2025) Virtual Life Story Club Intervention to Improve Loneliness and Apathy in Community-Dwelling Older Adults: Protocol for a Mixed Methods Feasibility Study. JMIR research protocols, 14, e70518.

Matsuda N, et al. (2025) Analysis of gene expression of Babesia gibsoni cultured with diminazene aceturate using RNA sequencing. The Journal of veterinary medical science, 87(2), 181.

Bach SV, et al. (2025) BDNF-DT and BDNF-AS-DT: Novel Genes in the BDNF locus. medRxiv : the preprint server for health sciences.

B??meke P, et al. (2025) Phenolic metabolism in *Sarcandra glabra* is mediated by distinct BAHD hydroxycinnamoyltransferases. *The Plant journal : for cell and molecular biology*, 121(5), e70035.

Wang B, et al. (2025) ebv-sisRNA-3 contributes to the formation of G4-associated R-loop upstream of EBV lytic replication origin in latently infected cells. *Cell & bioscience*, 15(1), 91.

Zhang Q, et al. (2025) LINC01235 Promotes Clonal Evolution through DNA Replication Licensing-Induced Chromosomal Instability in Breast Cancer. *Advanced science* (Weinheim, Baden-Wurttemberg, Germany), 12(14), e2413527.

Varano G, et al. (2025) B-cell Receptor Silencing Reveals the Origin and Dependencies of High-Grade B-cell Lymphomas with MYC and BCL2 Rearrangements. *Blood cancer discovery*, 6(4), 364.

Yi X, et al. (2025) Immunoglobulin gene loci structure and diversity of raccoon dog (*Nyctereutes procyonoides*). *BMC genomics*, 26(1), 424.

Feng KH, et al. (2025) Diversity and evolution analysis of RNA viruses in three wheat aphid species. *BMC genomics*, 26(1), 353.

Olmos M, et al. (2025) Evaluation of a 3D printed training model with realistic spatial-anatomical conditions for head and neck microsurgery. *Clinical oral investigations*, 29(5), 228.

Dekker L, et al. (2025) Prehospital Large-Vessel Occlusion Stroke Detection Scales: A Pooled Individual Patient Data Analysis of 2 Prospective Cohorts. *Neurology*, 104(9), e213570.

Zhao S-S, et al. (2025) *Neospora caninum* infection specifically suppresses the expression of a host lncRNA XR\_001919077.1 to facilitate parasite propagation by modulating host cell mitochondrial function and autophagy. *Microbiology spectrum*, 13(2), e0158024.

Ge S, et al. (2025) A cis-natural antisense RNA regulates alternative polyadenylation of *SISPX5* under Pi starvation in tomato. *Nature communications*, 16(1), 7981.

Yu SP, et al. (2025) A Japanese encephalitis virus biological clone with an E gene point mutation exhibits in vitro and in vivo attenuation of neurovirulence. *The Journal of general virology*, 106(9).

Luo X, et al. (2025) IGF2BP2 binding to CPSF6 facilitates m6A-mediated alternative polyadenylation of PUM2 and promotes malignant progression in ovarian cancer. *Clinical and translational medicine*, 15(7), e70388.