

Resource Summary Report

Generated by T1D on Aug 1, 2026

HCLUST

RRID:SCR_009154

Type: Tool

Proper Citation

HCLUST (RRID:SCR_009154)

Resource Information

URL: <http://wpicr.wpic.pitt.edu/WPICCompGen/hclust/hclust.htm>

Proper Citation: HCLUST (RRID:SCR_009154)

Description: Software application that is a simple clustering method that can be used to rapidly identify a set of tag SNP's based upon genotype data (entry from Genetic Analysis Software), THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 16,2025.

Abbreviations: HCLUST

Synonyms: R/HCLUST

Resource Type: software resource, software application

Keywords: gene, genetic, genomic, r, bio.tools

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: HCLUST

Resource ID: SCR_009154

Alternate IDs: biotools:h-clust, SCR_009102, nlx_154195, nlx_154331

Alternate URLs: <https://bio.tools/h-clust>

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260801T042646+0000

Ratings and Alerts

No rating or validation information has been found for HCLUST.

No alerts have been found for HCLUST.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1459 mentions in open access literature.

Listed below are recent publications. The full list is available at [T1D](#).

Bleve A, et al. (2025) ROR?? Bridges Cancer-Driven Lipid Dysmetabolism and Myeloid Immunosuppression. *Cancer discovery*, 15(7), 1505.

Azizi M, et al. (2025) Assessment of the phenotypic and physicochemical traits of nine Iranian endemic fenugreek (*Trigonella foenum-graecum* L.). *Scientific reports*, 15(1), 3303.

Garcia-Cabau C, et al. (2025) Mis-splicing of a neuronal microexon promotes CPEB4 aggregation in ASD. *Nature*, 637(8045), 496.

Ausems ANMA, et al. (2025) Sex-specific variation in breeding-ground departure decisions and detection rates during southbound migration in West Atlantic Flyway Whimbrels (*Numenius phaeopus*). *Scientific reports*, 15(1), 6735.

McCarthy OS, et al. (2025) Using Community Composition and Successional Theory to Guide Site-Specific Coral Reef Management. *Global change biology*, 31(1), e70050.

Wu J, et al. (2025) Flavones enrich rhizosphere *Pseudomonas* to enhance nitrogen utilization and secondary root growth in *Populus*. *Nature communications*, 16(1), 1461.

Uffenorde J, et al. (2025) Enhancing glioblastoma therapy: unveiling synergistic anticancer effects of Onalespib - radiotherapy combination therapy. *Frontiers in oncology*, 15, 1451156.

Ma Y, et al. (2025) Longitudinal Viral Load Clustering for People With HIV Using Functional Principal Component Analysis. *AIDS research and treatment*, 2025, 5890464.

Mahmoudian-Hamedani S, et al. (2025) Investigating combined hypoxia and stemness indices for prognostic transcripts in gastric cancer: Machine learning and network analysis approaches. *Biochemistry and biophysics reports*, 41, 101897.

Liu Y, et al. (2025) Differential transcriptomic profiling of lipid metabolism and collagen remodeling in fast- and slow-twitch skeletal muscles in aging. *FASEB journal : official publication of the Federation of American Societies for Experimental Biology*, 39(2), e70335.

Miebach J, et al. (2025) Importance, structure, cultivability, and resilience of the bacterial microbiota during infection of laboratory-grown *Haematococcus* spp. by the blastocladialean pathogen *Paraphysoderma sedebokerense*: evidence for a domesticated microbiota and its potential for biocontrol. *FEMS microbiology ecology*, 101(2).

Geng D, et al. (2025) Identification of key genes associated with residual feed intake in small-sized meat ducks through integrated analysis of mRNA and miRNA transcriptomes. *Poultry science*, 104(5), 105058.

Liu R, et al. (2025) Biochemical properties and substrate specificity of GOB-38 in *Elizabethkingia anophelis*. *Scientific reports*, 15(1), 351.

Sun M, et al. (2025) Exploring the potential mechanisms of sorafenib resistance in hepatocellular carcinoma cell lines based on RNA sequencing. *Cancer cell international*, 25(1), 91.

Zhang N, et al. (2025) Integration of multi-omics data accelerates molecular analysis of common wheat traits. *Nature communications*, 16(1), 2200.

Kolachala VL, et al. (2025) Altered inflammatory mucosal signatures within their spatial and cellular context during active ileal Crohn's disease. *JCI insight*, 10(5).

Torrado H, et al. (2025) Evolutionary Genomics of Two Co-occurring Congeneric Fore Reef Coral Species on Guam (Mariana Islands). *Genome biology and evolution*, 17(1).

Zeng PYF, et al. (2025) Cellular blueprint of healthy and diseased human epiglottis and subglottis-a study of the Canadian Airways Research (CARE) group. *EBioMedicine*, 114, 105631.

Ni C, et al. (2025) Single-cell transcriptomic analysis reveals prognosis-related stromal signatures that potentiate stratification of patients with extrahepatic cholangiocarcinoma. *BMC gastroenterology*, 25(1), 235.

Bao Q, et al. (2025) Identification of immune-related cervical cancer prognostic biomarkers and construction of prognostic model based on tumor microenvironment. *European journal of medical research*, 30(1), 261.