

Resource Summary Report

Generated by RRID on Sep 10, 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 application, software resource

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: 20260905T133045+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 1460 mentions in open access literature.

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

Zhou Z, et al. (2026) Using an Unbiased Coexpression Network to Reveal Cross-Talking Pathways of Phosphoinositide-3-Kinase Regulatory Subunit 1 in Skin Aging and Rejuvenation. *FASEB journal : official publication of the Federation of American Societies for Experimental Biology*, 40(2), e71466.

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

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.

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

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.

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

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).

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.

Tsementzi D, et al. (2025) Associations among HPV persistence, the vaginal microbiome, and cervical cancer recurrence. *Journal of translational medicine*, 23(1), 858.

Castaneda E, et al. (2025) Spectral divergence prioritizes key classes, genes, and pathways shared between substance use disorders and cardiovascular disease. *Frontiers in neuroscience*, 19, 1572243.

Meek MH, et al. (2025) Range-wide climate risk and adaptive potential in a cold-water fish species. *Nature communications*, 16(1), 7514.

D'Onofrio V, et al. (2025) AS04 drives superior cross-protective antibody response by increased NOTCH signaling of dendritic cells and proliferation of memory B cells. *Frontiers in immunology*, 16, 1623405.

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

Wu J, et al. (2025) Genetic and phenotypic insights into *Cyberlindnera jadinii* as a promising yeast for industrial biotechnology. *G3 (Bethesda, Md.)*, 15(9).

Wang K, et al. (2025) Circadian rhythm genes and immune cell infiltration in myasthenia gravis: A comprehensive analysis. *PloS one*, 20(7), e0327829.

An Y, et al. (2025) Identification of a LncRNA based CeRNA network signature to establish a prognostic model and explore potential therapeutic targets in gastric cancer. *Scientific reports*, 15(1), 20891.

Hamazaki Y, et al. (2025) Generation and characterization of induced pluripotent stem cells of small apes. *Frontiers in cell and developmental biology*, 13, 1536947.

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

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.