Resource Summary Report

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ggpubr

RRID:SCR 021139

Type: Tool

Proper Citation

ggpubr (RRID:SCR_021139)

Resource Information

URL: https://CRAN.R-project.org/package=ggpubr

Proper Citation: ggpubr (RRID:SCR_021139)

Description: Software R package provides functions for creating and customizing ggplot2

based publication ready plots. Used for creation of ggplot2 based graphs.

Resource Type: software toolkit, software resource

Keywords: ggplot2, plots, data visualization, ggplot2 based graphs creation

Funding:

Availability: Free, Available for download, Freely available

Resource Name: ggpubr

Resource ID: SCR_021139

License: GPL-2

Record Creation Time: 20220129T080354+0000

Record Last Update: 20250412T060316+0000

Ratings and Alerts

No rating or validation information has been found for ggpubr.

No alerts have been found for ggpubr.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 112 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch.org.

Jiang J, et al. (2025) Identification and validation of glucose metabolism-related gene signature in endometrial cancer. BMC cancer, 25(1), 30.

Xu W, et al. (2025) Progressive natural killer cell dysfunction in advanced-stage clear-cell renal cell carcinoma and association with clinical outcomes. ESMO open, 10(2), 104105.

Wyatt NJ, et al. (2025) Evaluation of intestinal biopsy tissue preservation methods to facilitate large-scale mucosal microbiota research. EBioMedicine, 112, 105550.

Salo W, et al. (2024) Influence of mixed and single infection of grapevine leafroll-associated viruses and viral load on berry quality. Tree physiology, 44(5).

Waizumi R, et al. (2024) A major endogenous glycoside hydrolase mediating quercetin uptake in Bombyx mori. PLoS genetics, 20(1), e1011118.

Leng D, et al. (2024) Single nucleus/cell RNA-seq of the chicken hypothalamic-pituitary-ovarian axis offers new insights into the molecular regulatory mechanisms of ovarian development. Zoological research, 45(5), 1088.

Najafi A, et al. (2024) Protocol for inferring epithelial-to-mesenchymal transition trajectories from single-cell RNA sequencing data using R. STAR protocols, 5(1), 102819.

Liu D, et al. (2024) Protocol for analyzing plasma cell-free DNA fragment end motifs from ultra-low-pass whole-genome sequencing. STAR protocols, 5(4), 103357.

González-Iglesias A, et al. (2024) Intron detention tightly regulates the stemness/differentiation switch in the adult neurogenic niche. Nature communications, 15(1), 2837.

Huang X, et al. (2024) Protocol to identify fungal profile from fecal metagenomes in cancer patients prior to immunotherapy. STAR protocols, 5(1), 102847.

Dong J, et al. (2024) Age-related changes of dopamine D1 and D2 receptors expression in parvalbumin-positive cells of the orbitofrontal and prelimbic cortices of mice. Frontiers in neuroscience, 18, 1364067.

Borgmästars E, et al. (2024) Multi-omics profiling to identify early plasma biomarkers in prediagnostic pancreatic ductal adenocarcinoma: a nested case-control study. Translational oncology, 48, 102059.

Poort VM, et al. (2024) Transient Differentiation-State Plasticity Occurs during Acute Lymphoblastic Leukemia Initiation. Cancer research, 84(16), 2720.

Marques-Pamies M, et al. (2024) Personalized Medicine in Acromegaly: The ACROFAST Study. The Journal of clinical endocrinology and metabolism, 110(1), 30.

Hai L, et al. (2024) A clinically applicable connectivity signature for glioblastoma includes the tumor network driver CHI3L1. Nature communications, 15(1), 968.

Burdett NL, et al. (2024) Timing of whole genome duplication is associated with tumor-specific MHC-II depletion in serous ovarian cancer. Nature communications, 15(1), 6069.

Bebelman MP, et al. (2024) Hepatocyte differentiation requires anisotropic expansion of bile canaliculi. Development (Cambridge, England), 151(22).

Zhou LZ, et al. (2024) The RALF signaling pathway regulates cell wall integrity during pollen tube growth in maize. The Plant cell, 36(5), 1673.

Zhao L, et al. (2024) Identification of significant m6A regulators and immune microenvironment characterization in ischemic stroke. Scientific reports, 14(1), 3456.

Hoekstra ME, et al. (2024) Distinct spatiotemporal dynamics of CD8+ T cell-derived cytokines in the tumor microenvironment. Cancer cell, 42(1), 157.