# **Resource Summary Report**

Generated by <u>NIF</u> on May 25, 2025

# **forestplot**

RRID:SCR\_023633 Type: Tool

**Proper Citation** 

forestplot (RRID:SCR\_023633)

#### **Resource Information**

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

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**Description:** Software R package that allows for multiple confidence intervals per row, custom fonts for each text element, custom confidence intervals, text mixed with expressions, and more. Advanced Forest Plot using 'grid' graphics. Used to extend the use of forest plots beyond meta analyses.

Resource Type: software resource, software toolkit

**Keywords:** multiple confidence intervals per row, custom fonts for each text element, custom confidence intervals, text mixed with expressions,

Funding:

Availability: Free, Available for download, Freely available

Resource Name: forestplot

Resource ID: SCR\_023633

License: GPL v2

**Record Creation Time:** 20230601T050210+0000

Record Last Update: 20250525T032619+0000

**Ratings and Alerts** 

No rating or validation information has been found for forestplot.

No alerts have been found for forestplot.

## Data and Source Information

Source: <u>SciCrunch Registry</u>

## **Usage and Citation Metrics**

We found 8 mentions in open access literature.

Listed below are recent publications. The full list is available at <u>NIF</u>.

Egea-Rodriguez S, et al. (2025) RECQL4 affects MHC class II-mediated signalling and favours an immune-evasive signature that limits response to immune checkpoint inhibitor therapy in patients with malignant melanoma. Clinical and translational medicine, 15(1), e70094.

Bentley AR, et al. (2024) Lipidomics profiling and circulating triglyceride concentrations in sub-Saharan African individuals. Scientific reports, 14(1), 20834.

Zhen L, et al. (2024) Activated hedgehog gene pattern correlates with dismal clinical outcome and tumor microenvironment heterogeneity in hepatocellular carcinoma. Heliyon, 10(5), e26989.

Lu Z, et al. (2024) High expression of CCNB2 is an independent predictive poor prognostic biomarker and correlates with immune infiltrates in breast carcinoma. Heliyon, 10(10), e31586.

Liu W, et al. (2024) SnapHiC-G: identifying long-range enhancer-promoter interactions from single-cell Hi-C data via a global background model. Briefings in bioinformatics, 25(5).

Wang D, et al. (2024) Race as a prognostic factor of breast mucinous carcinoma. Journal of cancer research and clinical oncology, 151(1), 25.

Gallagher RI, et al. (2023) Protein signaling and drug target activation signatures to guide therapy prioritization: Therapeutic resistance and sensitivity in the I-SPY 2 Trial. Cell reports. Medicine, 4(12), 101312.

Martinez-Carrasco A, et al. (2023) Genetic meta-analysis of levodopa induced dyskinesia in Parkinson's disease. NPJ Parkinson's disease, 9(1), 128.