

Resource Summary Report

Generated by [RRID](#) on Aug 12, 2026

[gridExtra](#)

RRID:SCR_025249

Type: Tool

Proper Citation

gridExtra (RRID:SCR_025249)

Resource Information

URL: <https://CRAN.R-project.org/package=gridExtra>

Proper Citation: gridExtra (RRID:SCR_025249)

Description: Software R package to work with "grid" graphics, notably to arrange multiple grid-based plots on page, and draw tables. Miscellaneous functions for Grid graphics.

Resource Type: software toolkit, software resource

Keywords: Miscellaneous functions for Grid graphics, Grid, functions for graphics, arrange multiple grid based plots, draw tables,

Availability: Free, Available for download, Freely available

Resource Name: gridExtra

Resource ID: SCR_025249

License: GNU GPL v3

Record Creation Time: 20240411T053243+0000

Record Last Update: 20260810T040911+0000

Ratings and Alerts

No rating or validation information has been found for gridExtra.

No alerts have been found for gridExtra.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 20 mentions in open access literature.

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

Cs??sz S, et al. (2026) A Taxonomic Revision of the East Mediterranean Species of the *Crematogaster scutellaris* Complex (Hymenoptera: Formicidae). *Insects*, 17(6).

Fang K, et al. (2026) Pharmaco-genomic characterization of pancreatic and biliary tract cancer tumoroids for drug response. *iScience*, 29(3), 115062.

Chen A, et al. (2025) Symbiotic T6SS affects horizontal transmission of *Paraburkholderia bonniea* among *Dictyostelium discoideum* amoeba hosts. *ISME communications*, 5(1), ycaf005.

Ravi Kumar V, et al. (2025) Founders predict trait evolution and population performance after evolutionary rescue in the red flour beetle. *Proceedings of the National Academy of Sciences of the United States of America*, 122(36), e2506244122.

Popp C, et al. (2025) *Hymenoplectella* and *Diaporthe* dominate the fungal community of dieback affected sea Buckthorn from Northern Germany. *Environmental microbiome*, 20(1), 141.

Ellegaard KM, et al. (2025) SARS-CoV-2 sequencing artifacts associated with targeted PCR enrichment and read mapping. *PloS one*, 20(10), e0334009.

Jayasinghe RG, et al. (2025) Single-cell transcriptomic profiling reveals diversity in human iNKT cells across hematologic tissues. *Cell reports*, 44(5), 115587.

Okoye GD, et al. (2025) Single-cell map of innate-like lymphocyte response to *Francisella tularensis* infection reveals interleukin-17-dependent protection by MAIT cells. *iScience*, 28(3), 111810.

Rocafort M, et al. (2024) HIV-associated gut microbial alterations are dependent on host and geographic context. *Nature communications*, 15(1), 1055.

Golov AK, et al. (2024) A genome-wide nucleosome-resolution map of promoter-centered interactions in human cells corroborates the enhancer-promoter looping model. *eLife*, 12.

Conte AL, et al. (2023) Atopic dermatitis-derived *Staphylococcus aureus* strains: what makes them special in the interplay with the host. *Frontiers in cellular and infection microbiology*, 13, 1194254.

Peris D, et al. (2023) Macroevolutionary diversity of traits and genomes in the model yeast genus *Saccharomyces*. *Nature communications*, 14(1), 690.

Heitger M, et al. (2023) Respiration, Production, and Growth Efficiency of Marine Pelagic Fungal Isolates. *Journal of fungi (Basel, Switzerland)*, 9(4).

Ilgisonis EV, et al. (2022) Evolution of Protein Functional Annotation: Text Mining Study. *Journal of personalized medicine*, 12(3).

von Huth S, et al. (2021) Intestinal protozoan infections shape fecal bacterial microbiota in children from Guinea-Bissau. *PLoS neglected tropical diseases*, 15(3), e0009232.

Christiansen H, et al. (2021) Facilitating population genomics of non-model organisms through optimized experimental design for reduced representation sequencing. *BMC genomics*, 22(1), 625.

Osimo EF, et al. (2020) Longitudinal population subgroups of CRP and risk of depression in the ALSPAC birth cohort. *Comprehensive psychiatry*, 96, 152143.

Khatri M, et al. (2020) The association between serum chloride levels and chronic kidney disease progression: a cohort study. *BMC nephrology*, 21(1), 165.

Kern F, et al. (2019) Systematic Assessment of Blood-Borne MicroRNAs Highlights Molecular Profiles of Endurance Sport and Carbohydrate Uptake. *Cells*, 8(9).

Lappan R, et al. (2018) A microbiome case-control study of recurrent acute otitis media identified potentially protective bacterial genera. *BMC microbiology*, 18(1), 13.