

Resource Summary Report

Generated by [dkNET](#) on Apr 25, 2025

GGally

RRID:SCR_026114

Type: Tool

Proper Citation

GGally (RRID:SCR_026114)

Resource Information

URL: <https://github.com/ggobi/ggally>

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Description: Software R package that extends ggplot2 by adding several functions to reduce complexity of combining geoms with transformed data. Some of these functions include pairwise plot matrix, scatterplot plot matrix, parallel coordinates plot, survival plot, and several functions to plot networks.

Resource Type: software toolkit, source code, software resource

Keywords: extends ggplot2, reduce complexity of combining geoms with transformed data, reduce complexity, combining geoms, transformed data,

Funding:

Availability: Free, Available for download, Freely available

Resource Name: GGally

Resource ID: SCR_026114

Record Creation Time: 20241203T053255+0000

Record Last Update: 20250425T060719+0000

Ratings and Alerts

No rating or validation information has been found for GGally.

No alerts have been found for GGally.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

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.

Akohoue F, et al. (2022) Separation of the effects of two reduced height (Rht) genes and genomic background to select for less Fusarium head blight of short-strawed winter wheat (*Triticum aestivum* L.) varieties. *TAG. Theoretical and applied genetics. Theoretische und angewandte Genetik*, 135(12), 4303.

García-Ortiz H, et al. (2021) The genomic landscape of Mexican Indigenous populations brings insights into the peopling of the Americas. *Nature communications*, 12(1), 5942.