

Resource Summary Report

Generated by RRID on Aug 12, 2026

yEd

RRID:SCR_025325

Type: Tool

Proper Citation

yEd (RRID:SCR_025325)

Resource Information

URL: <https://www.yworks.com/products/yed>

Proper Citation: yEd (RRID:SCR_025325)

Description: Software desktop application as general purpose diagramming program with multi-document interface. Cross-platform Java application to generate high-quality diagrams. Runs on Windows, Linux, Mac OS, and other platforms that support Java Virtual Machine.

Synonyms: yWorks yEd Graph Editor

Resource Type: software application, software resource

Keywords: yWorks, Java, general purpose diagramming program, diagrams,

Availability: Restricted

Resource Name: yEd

Resource ID: SCR_025325

Record Creation Time: 20240514T053253+0000

Record Last Update: 20260810T040842+0000

Ratings and Alerts

No rating or validation information has been found for yEd.

No alerts have been found for yEd.

Data and Source Information

Usage and Citation Metrics

We found 7 mentions in open access literature.

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

Eulopa-Hernandez JF, et al. (2025) Dimensional validation of CubeSat structures using 3D optical scanning technology. Scientific reports, 15(1), 45536.

Mazein A, et al. (2025) An explorable model of an adverse outcome pathway of cytokine release syndrome related to the administration of immunomodulatory biotherapeutics and cellular therapies. Frontiers in immunology, 16, 1601670.

Ralf A, et al. (2024) Male Pedigree Toolbox: A Versatile Software for Y-STR Data Analyses. Genes, 15(2).

B??nfalvi Z, et al. (2024) Transcriptome, hormonal, and secondary metabolite changes in leaves of DEFENSE NO DEATH 1 (DND1) silenced potato plants. Scientific reports, 14(1), 20601.

Mazein A, et al. (2023) A guide for developing comprehensive systems biology maps of disease mechanisms: planning, construction and maintenance. Frontiers in bioinformatics, 3, 1197310.

Bahl A, et al. (2023) PROTEOMAS: a workflow enabling harmonized proteomic meta-analysis and proteomic signature mapping. Journal of cheminformatics, 15(1), 34.

Degeneve C, et al. (2022) Analysing the digital transformation of the market for fake documents using a computational linguistic approach. Forensic science international. Synergy, 5, 100287.