

Resource Summary Report

Generated by SPARC SAWG on Jul 28, 2026

KegTools

RRID:SCR_006432

Type: Tool

Proper Citation

KegTools (RRID:SCR_006432)

Resource Information

URL: <http://www.kegg.jp/kegg/download/kegtools.html>

Proper Citation: KegTools (RRID:SCR_006432)

Description: Freely available desktop applications including KegHier: Java application for browsing BRITE hierarchy files, KegArray: Java application for microarray data analysis, and KegDraw: Java application for drawing compound and glycan structures. They run on the Mac OS X, Windows, and Linux platforms.

Abbreviations: KegTools

Resource Type: software application, software resource

Keywords: microarray data analysis, microarray, data analysis, compound, glycan, structure, compound structure, glycan structure

Availability: Free

Resource Name: KegTools

Resource ID: SCR_006432

Alternate IDs: nlx_29129

Old URLs: <http://www.genome.jp/download/>

Record Creation Time: 20220129T080236+0000

Record Last Update: 20260727T040414+0000

Ratings and Alerts

No rating or validation information has been found for KegTools.

No alerts have been found for KegTools.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 10 mentions in open access literature.

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

Zheng Q, et al. (2020) miR-224 targets BTRC and promotes cell migration and invasion in colorectal cancer. 3 Biotech, 10(11), 485.

Theis JL, et al. (2020) Patient-specific genomics and cross-species functional analysis implicate LRP2 in hypoplastic left heart syndrome. eLife, 9.

Lal K, et al. (2020) Computational tools for drawing, building and displaying carbohydrates: a visual guide. Beilstein journal of organic chemistry, 16, 2448.

Zhou C, et al. (2017) MiR-650 represses high-risk non-metastatic colorectal cancer progression via inhibition of AKT2/GSK3 β /E-cadherin pathway. Oncotarget, 8(30), 49534.

Bu D, et al. (2017) Transcriptome difference and potential crosstalk between liver and mammary tissue in mid-lactation primiparous dairy cows. PloS one, 12(3), e0173082.

Yabuzaki J, et al. (2017) Carotenoids Database: structures, chemical fingerprints and distribution among organisms. Database : the journal of biological databases and curation, 2017(1).

Bionaz M, et al. (2015) Transcription Adaptation during In Vitro Adipogenesis and Osteogenesis of Porcine Mesenchymal Stem Cells: Dynamics of Pathways, Biological Processes, Up-Stream Regulators, and Gene Networks. PloS one, 10(9), e0137644.

Shahzad K, et al. (2014) Integrative analyses of hepatic differentially expressed genes and blood biomarkers during the peripartal period between dairy cows overfed or restricted-fed energy prepartum. PloS one, 9(6), e99757.

Pairett AN, et al. (2013) De novo assembly and characterization of two transcriptomes reveal multiple light-mediated functions in the scallop eye (Bivalvia: Pectinidae). PloS one, 8(7), e69852.

Qin XY, et al. (2012) Identification of novel low-dose bisphenol a targets in human foreskin fibroblast cells derived from hypospadias patients. PloS one, 7(5), e36711.