

Resource Summary Report

Generated by PRECISE-TBI on Sep 8, 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: 20260905T133237+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 [PRECISE-TBI](#) .

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.