

Resource Summary Report

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Pathway Articulator

RRID:SCR_002101

Type: Tool

Proper Citation

Pathway Articulator (RRID:SCR_002101)

Resource Information

URL: <http://www.jubilantbiosys.com/pathart.html>

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Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented on July 15, 2013. A comprehensive collection of manually curated information from literature as well as public domain databases on signaling and metabolic pathways. PathArt includes a dynamic pathway articulator component, which builds molecular interaction networks from curated databases. PathArt provides a tool for analysis, biological interpretation and visualization of microarray data results in these curated pathways. In addition, PathArt provides a collection of high priority disease and physiology pathways with emphasis on pathway responsive genes and knockouts. The coverage is for pathways of Human, Rat and Mouse for cell specific, tissue specific and organism specific data. The present version of PathArt covers the following: -Includes 3527 regulatory and signaling pathways across diseases and physiologies. -Provides information on 39 high priority diseases, and pathway and disease responsive genes. -Provides pathway information on 23 diverse physiologies. -Covers information on ~8783 Knockouts and ~18000 mutation data points. - Coverage of pathways for Human, Mouse and Rat for cell specificity, tissue specificity and organism specific data.

Abbreviations: PathArt

Synonyms: Pathway Articulator

Resource Type: data or information resource, database

Keywords: cell, human, metabolic, molecular, mouse, pathway, physiology, rat, regulatory, signaling

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Pathway Articulator

Resource ID: SCR_002101

Alternate IDs: nif-0000-20883

Record Creation Time: 20220129T080211+0000

Record Last Update: 20260905T133116+0000

Ratings and Alerts

No rating or validation information has been found for Pathway Articulator.

No alerts have been found for Pathway Articulator.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Chen YY, et al. (2010) Microarray analysis reveals the inhibition of nuclear factor-kappa B signaling by aristolochic acid in normal human kidney (HK-2) cells. Acta pharmacologica Sinica, 31(2), 227.