

# Resource Summary Report

Generated by [kravitz2](#) on Jul 31, 2026

## BeWith

RRID:SCR\_016573

Type: Tool

### Proper Citation

BeWith (RRID:SCR\_016573)

### Resource Information

**URL:** <https://www.ncbi.nlm.nih.gov/CBBresearch/Przytycka/index.cgi#bewith>

**Proper Citation:** BeWith (RRID:SCR\_016573)

**Description:** Software tool for discovering relationships between cancer modules via integrated analysis of mutual exclusivity, co-occurrence and functional interactions.

**Abbreviations:** BeWith

**Synonyms:** BEtweenWITHin, Between-Within, BeWith

**Resource Type:** data processing software, software application, data analysis software, software resource

**Defining Citation:** [DOI:10.1371/journal.pcbi.1005695](https://doi.org/10.1371/journal.pcbi.1005695)

**Keywords:** clustering, framework, identify, module, mutation, interaction, pattern, gene, cancer

**Funding:** Naomi Kadar award ;  
NIH

**Availability:** Free, Available for download, Freely available

**Resource Name:** BeWith

**Resource ID:** SCR\_016573

**Record Creation Time:** 20220129T080331+0000

**Record Last Update:** 20260731T043004+0000

### Ratings and Alerts

No rating or validation information has been found for BeWith.

No alerts have been found for BeWith.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.