Resource Summary Report

Generated by FDI Lab - SciCrunch.org on May 15, 2025

POINT: Prediction Of INTeractome

RRID:SCR_007866
Type: Tool

Proper Citation

POINT: Prediction Of INTeractome (RRID:SCR_007866)

Resource Information

URL: http://point.bioinformatics.tw/Welcome.do

Proper Citation: POINT: Prediction Of INTeractome (RRID:SCR_007866)

Description: POINT is a protein-protein interaction database. It includes annotation of interologs and protein phsophorylation. This work analyzes the applicability of orthologs-based PPI prediction and provide the theoretical upper-bound of this approach.

Synonyms: POINT

Resource Type: data or information resource, database

Funding:

Resource Name: POINT: Prediction Of INTeractome

Resource ID: SCR_007866

Alternate URLs: http://point.bioinformatics.tw

Record Creation Time: 20220129T080244+0000

Record Last Update: 20250507T060529+0000

Ratings and Alerts

No rating or validation information has been found for POINT: Prediction Of INTeractome.

No alerts have been found for POINT: Prediction Of INTeractome.

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 FDI Lab - SciCrunch.org.

Liu X, et al. (2021) SARS-CoV-2-host proteome interactions for antiviral drug discovery. Molecular systems biology, 17(11), e10396.