## **Resource Summary Report**

Generated by <u>dkNET</u> on Apr 30, 2025

## **CoPreTHi**

RRID:SCR\_006217 Type: Tool

**Proper Citation** 

CoPreTHi (RRID:SCR\_006217)

## **Resource Information**

URL: http://athina.biol.uoa.gr/CoPreTHi/

Proper Citation: CoPreTHi (RRID:SCR\_006217)

**Description:** A Java based web application, which combines the results of methods that predict the location of transmembrane segments in protein sequences into a joint prediction histogram. Clearly, the joint prediction algorithm, produces superior quality results than individual prediction schemes.

Abbreviations: CoPreTHi

**Synonyms:** CoPreTHi - A Java-program which Combines the results of several methods (available throught the Internet ) that Predict Transmembrane regions in proteins in a joint prediction Histogram

**Resource Type:** analysis service resource, data analysis service, production service resource, service resource

Defining Citation: PMID:11471236

**Keywords:** java, predict, transmembrane, region, protein, histogram, algorithm, joint prediction

Funding:

Availability: Free

Resource Name: CoPreTHi

Resource ID: SCR\_006217

Alternate IDs: nlx\_151763

**Record Creation Time:** 20220129T080234+0000

Record Last Update: 20250430T055425+0000

## **Ratings and Alerts**

No rating or validation information has been found for CoPreTHi.

No alerts have been found for CoPreTHi.

Data and Source Information

Source: <u>SciCrunch Registry</u>

**Usage and Citation Metrics** 

We have not found any literature mentions for this resource.