

# Resource Summary Report

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## CoPreTHi

RRID:SCR\_006217

Type: Tool

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### Proper Citation

CoPreTHi (RRID:SCR\_006217)

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### Resource Information

**URL:** <http://athina.biol.uoa.gr/CoPreTHi/>

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**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

**Availability:** Free

**Resource Name:** CoPreTHi

**Resource ID:** SCR\_006217

**Alternate IDs:** nlx\_151763

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

**Record Last Update:** 20260829T182946+0000

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## Ratings and Alerts

No rating or validation information has been found for CoPreTHi.

No alerts have been found for CoPreTHi.

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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.