

Resource Summary Report

Generated by SPARC SAWG on Jul 31, 2026

FT

RRID:SCR_006228

Type: Tool

Proper Citation

FT (RRID:SCR_006228)

Resource Information

URL: <http://athina.biol.uoa.gr/FT/>

Proper Citation: FT (RRID:SCR_006228)

Description: A web program that can locate residue periodicities in either amino acid or DNA sequences. It is based on an algorithm of Dr. A.D. McLachlan (1977). NOTE: You must use a Java compatible browser to run the application.

Abbreviations: FT

Synonyms: FT - Study of Residue Periodicities in Sequences

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

Keywords: residue, periodicity, sequence, amino acid, dna, algorithm

Resource Name: FT

Resource ID: SCR_006228

Alternate IDs: nlx_151783

Record Creation Time: 20220129T080235+0000

Record Last Update: 20260731T042720+0000

Ratings and Alerts

No rating or validation information has been found for FT.

No alerts have been found for FT.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.