

Resource Summary Report

Generated by SPARC SAWG on Sep 14, 2026

FT

RRID:SCR_006228

Type: Tool

Proper Citation

FT (RRID:SCR_006228)

Resource Information

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

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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: analysis service resource, data analysis service, production service resource, service resource

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: 20260912T200141+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.