

# Resource Summary Report

Generated by [ASWG](#) on Aug 4, 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:** data analysis service, service resource, analysis service resource, production 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:** 20260804T043303+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](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.