

Resource Summary Report

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F-Seq

RRID:SCR_010880

Type: Tool

Proper Citation

F-Seq (RRID:SCR_010880)

Resource Information

URL: <http://fureylab.web.unc.edu/software/fseq/>

Proper Citation: F-Seq (RRID:SCR_010880)

Description: A software package that generates a continuous tag sequence density estimation allowing identification of biologically meaningful sites whose output can be displayed directly in the UCSC Genome Browser.

Abbreviations: F-Seq

Resource Type: software resource

Resource Name: F-Seq

Resource ID: SCR_010880

Alternate IDs: OMICS_00482

Record Creation Time: 20220129T080301+0000

Record Last Update: 20261017T202737+0000

Ratings and Alerts

No rating or validation information has been found for F-Seq.

No alerts have been found for F-Seq.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.