DISULFIND
RRID:SCR_016072
Type: Tool

Proper Citation

DISULFIND (RRID:SCR_016072)

Resource Information

URL: http://disulfind.dsi.unifi.it/

Description: Software for predicting the disulfide bonding state of cysteines and their disulfide connectivity, starting from a protein sequence alone and may be useful in other genomic annotation tasks.

Resource Name: DISULFIND

Proper Citation: DISULFIND (RRID:SCR_016072)

Resource Type: Resource, software resource, software application, data processing software, data analysis software, sequence analysis software

Keywords: predict, disulfide, bonding, state, cysteine, protein, sequence, genomic, annotation

Resource ID: SCR_016072

Parent Organization: University of Florence; Florence; Italy

Funding Agency: Embark Fellowship from the Irish Research Council for Science, Engineering and Technology, EU NoE BIOPATTERN, EU STREP APrIL II

References: PMID:16844986

Availability: Free, Freely available

Website Status: Last checked up
Abbreviations: Disulfinder

Mentions Count: 44

Ratings and Alerts

No rating or validation information has been found for DISULFIND.

No alerts have been found for DISULFIND.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 44 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch Infrastructure.


