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

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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

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**Ratings and Alerts**

No rating or validation information has been found for DISULFIND.

No alerts have been found for DISULFIND.

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**Data and Source Information**

**Source:** [SciCrunch Registry](https://www.sci-crunch.org)

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**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](https://www.sci-crunch.org).


and a promising tool for drug delivery in cancer cells. Scientific reports, 8(1), 14739.


