Resource Summary Report

Generated by dkNET on Sep 3, 2026

CSpritz

RRID:SCR_021884
Type: Tool

Proper Citation

CSpritz (RRID:SCR_021884)

Resource Information

URL: http://old.protein.bio.unipd.it/cspritz/

Proper Citation: CSpritz (RRID:SCR_021884)

Description: Web tool for prediction of intrinsic protein disorder segments with annotation for homology, secondary structure and linear motifs.

Resource Type: data access protocol, software resource, web service

Defining Citation: DOI:10.1093/nar/gkr411

Keywords: Intrinsic protein disorder segments prediction, homology, secondary structure, linear motifs, annotation

Funding: Italian Ministry of Education ;
University of Padova

Availability: Free, Freely available

Resource Name: CSpritz

Resource ID: SCR_021884

Record Creation Time: 20220421T050137+0000

Record Last Update: 20260903T235829+0000

Ratings and Alerts

No rating or validation information has been found for CSpritz.

No alerts have been found for CSpritz.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at dkNET .

Flynn AJ, et al. (2023) Mechanosensitive ion channels MSL8, MSL9, and MSL10 have environmentally sensitive intrinsically disordered regions with distinct biophysical characteristics in vitro. Plant direct, 7(8), e515.

Philip J, et al. (2022) Cdc6 is sequentially regulated by PP2A-Cdc55, Cdc14, and Sic1 for origin licensing in S. cerevisiae. eLife, 11.