Resource Summary Report

Generated by FDI Lab - SciCrunch.org on Apr 11, 2025

VoroMQA

RRID:SCR_021881
Type: Tool

Proper Citation

VoroMQA (RRID:SCR_021881)

Resource Information

URL: https://bioinformatics.lt/wtsam/voromqa

Proper Citation: VoroMQA (RRID:SCR_021881)

Description: Web tool for assessment of protein structure quality using interatomic contact areas. Method for estimation of single protein structure quality.

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

Defining Citation: PMID:28263393

Keywords: protein structure quality assessment, single protein structure quality estimation, interatomic contact areas

Funding: European Social Fund

Availability: Free, Freely available

Resource Name: VoroMQA

Resource ID: SCR_021881

Record Creation Time: 20220421T050137+0000

Record Last Update: 20250410T071420+0000

Ratings and Alerts

No rating or validation information has been found for VoroMQA.

No alerts have been found for VoroMQA.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 5 mentions in open access literature.

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

Araújo MC, et al. (2024) Effect of Eugenol on Detrusor Muscle: Potential for Overactive Bladder Treatment. International neurourology journal, 28(4), 253.

Wozny MR, et al. (2023) In situ architecture of the ER-mitochondria encounter structure. Nature, 618(7963), 188.

Urbelien? N, et al. (2023) Cytidine deaminases catalyze the conversion of N(S,O)4-substituted pyrimidine nucleosides. Science advances, 9(5), eade4361.

Arguelles J, et al. (2023) In Silico Analysis of a Drosophila Parasitoid Venom Peptide Reveals Prevalence of the Cation-Polar-Cation Clip Motif in Knottin Proteins. Pathogens (Basel, Switzerland), 12(1).

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