

Sofie V Nielsen

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10971354/publications.pdf>

Version: 2024-02-01

14
papers

593
citations

933447

10
h-index

1125743

13
g-index

18
all docs

18
docs citations

18
times ranked

744
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | Molecular Basis and Regulation of OTULIN-LUBAC Interaction. <i>Molecular Cell</i> , 2014, 54, 335-348. | 9.7 | 158 |
| 2 | Predicting the impact of Lynch syndrome-causing missense mutations from structural calculations. <i>PLoS Genetics</i> , 2017, 13, e1006739. | 3.5 | 90 |
| 3 | Understanding the Origins of Loss of Protein Function by Analyzing the Effects of Thousands of Variants on Activity and Abundance. <i>Molecular Biology and Evolution</i> , 2021, 38, 3235-3246. | 8.9 | 65 |
| 4 | Toward mechanistic models for genotypeâ€“phenotype correlations in phenylketonuria using protein stability calculations. <i>Human Mutation</i> , 2019, 40, 444-457. | 2.5 | 56 |
| 5 | Computational and cellular studies reveal structural destabilization and degradation of MLH1 variants in Lynch syndrome. <i>ELife</i> , 2019, 8, . | 6.0 | 49 |
| 6 | Protein Quality Control in the Nucleus. <i>Biomolecules</i> , 2014, 4, 646-661. | 4.0 | 39 |
| 7 | Co-Chaperones in Targeting and Delivery of Misfolded Proteins to the 26S Proteasome. <i>Biomolecules</i> , 2020, 10, 1141. | 4.0 | 29 |
| 8 | Bioinformatics analysis identifies several intrinsically disordered human E3 ubiquitin-protein ligases. <i>PeerJ</i> , 2016, 4, e1725. | 2.0 | 24 |
| 9 | Blocking protein quality control to counter hereditary cancers. <i>Genes Chromosomes and Cancer</i> , 2017, 56, 823-831. | 2.8 | 23 |
| 10 | Folliculin variants linked to Birt-Hogg-DubÃ© syndrome are targeted for proteasomal degradation. <i>PLoS Genetics</i> , 2020, 16, e1009187. | 3.5 | 16 |
| 11 | Disease-linked mutations cause exposure of a protein quality control degron. <i>Structure</i> , 2022, 30, 1245-1253.e5. | 3.3 | 14 |
| 12 | Multiplexed assays reveal effects of missense variants in MSH2 and cancer predisposition. <i>PLoS Genetics</i> , 2021, 17, e1009496. | 3.5 | 13 |
| 13 | Protein destabilization and degradation as a mechanism for hereditary disease. , 2020, , 111-125. | | 5 |
| 14 | High-Throughput siRNA Screening Applied to the Ubiquitinâ€“Proteasome System. <i>Methods in Molecular Biology</i> , 2016, 1449, 421-439. | 0.9 | 2 |