

Mihkel Ærd

List of Publications by Year in descending order

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Version: 2024-02-01

17
papers

678
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932766

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887659

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23
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23
docs citations

23
times ranked

690
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 1 | The Eukaryotic Linear Motif resource: 2022 release. <i>Nucleic Acids Research</i> , 2022, 50, D497-D508. | 6.5 | 144 |
| 2 | Multisite phosphorylation by Cdk1 initiates delayed negative feedback to control mitotic transcription. <i>Current Biology</i> , 2022, 32, 256-263.e4. | 1.8 | 7 |
| 3 | Cdc6 is sequentially regulated by PP2A-Cdc55, Cdc14, and Sic1 for origin licensing in <i>S. cerevisiae</i> . <i>ELife</i> , 2022, 11, . | 2.8 | 6 |
| 4 | Cdc4 phospho-degrons allow differential regulation of Ame1CENP-U protein stability across the cell cycle. <i>ELife</i> , 2021, 10, . | 2.8 | 6 |
| 5 | Docking to a Basic Helix Promotes Specific Phosphorylation by G1-Cdk1. <i>International Journal of Molecular Sciences</i> , 2021, 22, 9514. | 1.8 | 2 |
| 6 | Regulation of trehalase activity by multi-site phosphorylation and 14-3-3 interaction. <i>Scientific Reports</i> , 2021, 11, 962. | 1.6 | 11 |
| 7 | A new linear cyclin docking motif that mediates exclusively Sâ€phase CDKâ€specific signaling. <i>EMBO Journal</i> , 2021, 40, e105839. | 3.5 | 23 |
| 8 | The sequence at Spike S1/S2 site enables cleavage by furin and phospho-regulation in SARS-CoV2 but not in SARS-CoV1 or MERS-CoV. <i>Scientific Reports</i> , 2020, 10, 16944. | 1.6 | 125 |
| 9 | Comprehensive Analysis of G1 Cyclin Docking Motif Sequences that Control CDK Regulatory Potency InÂVivo. <i>Current Biology</i> , 2020, 30, 4454-4466.e5. | 1.8 | 21 |
| 10 | Proline-Rich Motifs Control G2-CDK Target Phosphorylation and Priming an Anchoring Protein for Polo Kinase Localization. <i>Cell Reports</i> , 2020, 31, 107757. | 2.9 | 16 |
| 11 | A processive phosphorylation circuit with multiple kinase inputs and mutually diversional routes controls G1/S decision. <i>Nature Communications</i> , 2020, 11, 1836. | 5.8 | 12 |
| 12 | Detection of Multisite Phosphorylation of Intrinsically Disordered Proteins Using Phos-tag SDS-PAGE. <i>Methods in Molecular Biology</i> , 2020, 2141, 779-792. | 0.4 | 8 |
| 13 | Multisite phosphorylation code of CDK. <i>Nature Structural and Molecular Biology</i> , 2019, 26, 649-658. | 3.6 | 64 |
| 14 | Cyclin-Specific Docking Mechanisms Reveal the Complexity of M-CDK Function in the Cell Cycle. <i>Molecular Cell</i> , 2019, 75, 76-89.e3. | 4.5 | 50 |
| 15 | How the cell cycle clock ticks. <i>Molecular Biology of the Cell</i> , 2019, 30, 169-172. | 0.9 | 29 |
| 16 | Multistep phosphorylation systems: tunable components of biological signaling circuits. <i>Molecular Biology of the Cell</i> , 2014, 25, 3456-3460. | 0.9 | 38 |
| 17 | Multisite phosphorylation networks as signal processors for Cdk1. <i>Nature Structural and Molecular Biology</i> , 2013, 20, 1415-1424. | 3.6 | 112 |