

Bela Novak

List of Publications by Year in descending order

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122
papers

11,512
citations

38742

50
h-index

30922

102
g-index

139
all docs

139
docs citations

139
times ranked

9182
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Mitotic kinase oscillation governs the latching of cell cycle switches. <i>Current Biology</i> , 2022, 32, 2780-2785.e2. | 3.9 | 5 |
| 2 | Time-keeping and decision-making in the cell cycle. <i>Interface Focus</i> , 2022, 12, . | 3.0 | 10 |
| 3 | Mechanisms of signalling-memory governing progression through the eukaryotic cell cycle. <i>Current Opinion in Cell Biology</i> , 2021, 69, 7-16. | 5.4 | 21 |
| 4 | Computational modeling of chromosome re-replication in mutant strains of fission yeast. <i>Molecular Biology of the Cell</i> , 2021, 32, 830-841. | 2.1 | 0 |
| 5 | A Single Light-Responsive Sizer Can Control Multiple-Fission Cycles in <i>Chlamydomonas</i> . <i>Current Biology</i> , 2020, 30, 634-644.e7. | 3.9 | 16 |
| 6 | Cyclin A triggers Mitosis either via the Greatwall kinase pathway or Cyclin B. <i>EMBO Journal</i> , 2020, 39, e104419. | 7.8 | 52 |
| 7 | A Dynamical Paradigm for Molecular Cell Biology. <i>Trends in Cell Biology</i> , 2020, 30, 504-515. | 7.9 | 53 |
| 8 | CDK1-CCNB1 creates a spindle checkpoint "permissive state by enabling MPS1 kinetochore localization. <i>Journal of Cell Biology</i> , 2019, 218, 1182-1199. | 5.2 | 45 |
| 9 | Absolute quantification of cohesin, CTCF and their regulators in human cells. <i>ELife</i> , 2019, 8, . | 6.0 | 79 |
| 10 | Systems-level feedback regulation of cell cycle transitions in <i>Ostreococcus tauri</i> . <i>Plant Physiology and Biochemistry</i> , 2018, 126, 39-46. | 5.8 | 2 |
| 11 | A comprehensive model for the proliferation "quiescence decision in response to endogenous DNA damage in human cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 2532-2537. | 7.1 | 70 |
| 12 | Genome stability during cell proliferation: A systems analysis of the molecular mechanisms controlling progression through the eukaryotic cell cycle. <i>Current Opinion in Systems Biology</i> , 2018, 9, 22-31. | 2.6 | 13 |
| 13 | Two Interlinked Bistable Switches Govern Mitotic Control in Mammalian Cells. <i>Current Biology</i> , 2018, 28, 3824-3832.e6. | 3.9 | 62 |
| 14 | Dilution and titration of cell-cycle regulators may control cell size in budding yeast. <i>PLoS Computational Biology</i> , 2018, 14, e1006548. | 3.2 | 45 |
| 15 | APC/CCdh1 Enables Removal of Shugoshin-2 from the Arms of Bivalent Chromosomes by Moderating Cyclin-Dependent Kinase Activity. <i>Current Biology</i> , 2017, 27, 1462-1476.e5. | 3.9 | 8 |
| 16 | DNA damage during S-phase mediates the proliferation-quiescence decision in the subsequent G1 via p21 expression. <i>Nature Communications</i> , 2017, 8, 14728. | 12.8 | 284 |
| 17 | Cell-cycle transitions: a common role for stoichiometric inhibitors. <i>Molecular Biology of the Cell</i> , 2017, 28, 3437-3446. | 2.1 | 16 |
| 18 | Interlinked bistable mechanisms generate robust mitotic transitions. <i>Cell Cycle</i> , 2017, 16, 1885-1892. | 2.6 | 23 |

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|----|---|------|-----------|
| 19 | A Dynamical Framework for the All-or-None G1/S Transition. <i>Cell Systems</i> , 2016, 2, 27-37. | 6.2 | 115 |
| 20 | A PP2A-B55 recognition signal controls substrate dephosphorylation kinetics during mitotic exit. <i>Journal of Cell Biology</i> , 2016, 214, 539-554. | 5.2 | 164 |
| 21 | Two Bistable Switches Govern M Phase Entry. <i>Current Biology</i> , 2016, 26, 3361-3367. | 3.9 | 72 |
| 22 | Nutritional Control of Cell Size by the Greatwall-Endosulfine-PP2A-B55 Pathway. <i>Current Biology</i> , 2016, 26, 319-330. | 3.9 | 87 |
| 23 | Bistability, Oscillations, and Traveling Waves in Frog Egg Extracts. <i>Bulletin of Mathematical Biology</i> , 2015, 77, 796-816. | 1.9 | 7 |
| 24 | Cell Cycle Control by a Minimal Cdk Network. <i>PLoS Computational Biology</i> , 2015, 11, e1004056. | 3.2 | 49 |
| 25 | Model scenarios for switch-like mitotic transitions. <i>FEBS Letters</i> , 2015, 589, 667-671. | 2.8 | 21 |
| 26 | Kinetochores' microtubule error correction is driven by differentially regulated interaction modes. <i>Nature Cell Biology</i> , 2015, 17, 421-433. | 10.3 | 63 |
| 27 | Human Chromosome Segregation Involves Multi-Layered Regulation of Separase by the Peptidyl-Prolyl-Isomerase Pin1. <i>Molecular Cell</i> , 2015, 58, 495-506. | 9.7 | 49 |
| 28 | Premature Sister Chromatid Separation Is Poorly Detected by the Spindle Assembly Checkpoint as a Result of System-Level Feedback. <i>Cell Reports</i> , 2015, 13, 469-478. | 6.4 | 21 |
| 29 | Models in biology: lessons from modeling regulation of the eukaryotic cell cycle. <i>BMC Biology</i> , 2015, 13, 46. | 3.8 | 61 |
| 30 | A Model for the Epigenetic Switch Linking Inflammation to Cell Transformation: Deterministic and Stochastic Approaches. <i>PLoS Computational Biology</i> , 2014, 10, e1003455. | 3.2 | 17 |
| 31 | PP2A/B55 and Fcp1 Regulate Greatwall and Ensa Dephosphorylation during Mitotic Exit. <i>PLoS Genetics</i> , 2014, 10, e1004004. | 3.5 | 55 |
| 32 | Control of cell growth, division and death: information processing in living cells. <i>Interface Focus</i> , 2014, 4, 20130070. | 3.0 | 31 |
| 33 | Dependency of the Spindle Assembly Checkpoint on Cdk1 Renders the Anaphase Transition Irreversible. <i>Current Biology</i> , 2014, 24, 630-637. | 3.9 | 63 |
| 34 | Dynamical Scenarios for Chromosome Bi-orientation. <i>Biophysical Journal</i> , 2013, 104, 2595-2606. | 0.5 | 15 |
| 35 | Minimal Models for Cell-Cycle Control Based on Competitive Inhibition and Multisite Phosphorylations of Cdk Substrates. <i>Biophysical Journal</i> , 2013, 104, 1367-1379. | 0.5 | 13 |
| 36 | The BEG (PP2A-B55/ENSA/Greatwall) Pathway Ensures Cytokinesis follows Chromosome Separation. <i>Molecular Cell</i> , 2013, 52, 393-405. | 9.7 | 136 |

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|----|---|------|-----------|
| 37 | Role for regulated phosphatase activity in generating mitotic oscillations in <i>Xenopus</i> cell-free extracts. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 20539-20544. | 7.1 | 7 |
| 38 | The role of APC/C inhibitor Emi2/XErp1 in oscillatory dynamics of early embryonic cell cycles. Biophysical Chemistry, 2013, 177-178, 1-6. | 2.8 | 16 |
| 39 | Irreversible Transitions, Bistability and Checkpoint Controls in the Eukaryotic Cell Cycle. , 2013, , 265-285. | | 13 |
| 40 | Molecular mechanisms creating bistable switches at cell cycle transitions. Open Biology, 2013, 3, 120179. | 3.6 | 62 |
| 41 | Robust mitotic entry is ensured by a latching switch. Biology Open, 2013, 2, 924-931. | 1.2 | 17 |
| 42 | Pom1 is not the size ruler. Cell Cycle, 2013, 12, 3463-3464. | 2.6 | 1 |
| 43 | microRNA as a Potential Vector for the Propagation of Robustness in Protein Expression and Oscillatory Dynamics within a ceRNA Network. PLoS ONE, 2013, 8, e83372. | 2.5 | 15 |
| 44 | Cell Cycle Dynamics, Bistability and Oscillations. , 2013, , 263-270. | | 1 |
| 45 | CDK-Dependent Nuclear Localization of B-Cyclin Clb1 Promotes FEAR Activation during Meiosis I in Budding Yeast. PLoS ONE, 2013, 8, e79001. | 2.5 | 9 |
| 46 | Cell Cycle Transitions, Mitotic Exit. , 2013, , 333-336. | | 0 |
| 47 | Cell Cycle Dynamics, Irreversibility. , 2013, , 270-273. | | 0 |
| 48 | Cell Cycle, Budding Yeast. , 2013, , 337-341. | | 2 |
| 49 | A Structural Systems Biology Approach for Quantifying the Systemic Consequences of Missense Mutations in Proteins. PLoS Computational Biology, 2012, 8, e1002738. | 3.2 | 19 |
| 50 | A General G1/S-Phase Cell-Cycle Control Module in the Flowering Plant Arabidopsis thaliana. PLoS Genetics, 2012, 8, e1002847. | 3.5 | 103 |
| 51 | Multisite phosphoregulation of Cdc25 activity refines the mitotic entrance and exit switches. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 9899-9904. | 7.1 | 19 |
| 52 | Meiotic Prophase Requires Proteolysis of M Phase Regulators Mediated by the Meiosis-Specific APC/C ^{Am1} . Cell, 2012, 151, 603-618. | 28.9 | 93 |
| 53 | Interplay of transcriptional and proteolytic regulation in driving robust cell cycle progression. Molecular BioSystems, 2012, 8, 863. | 2.9 | 5 |
| 54 | Phosphorylation network dynamics in the control of cell cycle transitions. Journal of Cell Science, 2012, 125, 4703-4711. | 2.0 | 138 |

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|----|---|-----|-----------|
| 55 | Mathematical Model for Growth Regulation of Fission Yeast <i>Schizosaccharomyces pombe</i> . <i>PLoS ONE</i> , 2012, 7, e49675. | 2.5 | 6 |
| 56 | Spatiotemporal dynamics of Spc105 regulates the assembly of the <i>Drosophila</i> kinetochore. <i>Open Biology</i> , 2012, 2, 110032. | 3.6 | 47 |
| 57 | Protein Phosphatase 2A Controls the Order and Dynamics of Cell-Cycle Transitions. <i>Molecular Cell</i> , 2011, 44, 437-450. | 9.7 | 47 |
| 58 | Overexpression limits of fission yeast cell cycle regulators <i>in vivo</i> and <i>in silico</i> . <i>Molecular Systems Biology</i> , 2011, 7, 556. | 7.2 | 14 |
| 59 | Cell cycle commitment in budding yeast emerges from the cooperation of multiple bistable switches. <i>Open Biology</i> , 2011, 1, 110009. | 3.6 | 14 |
| 60 | Cell Cycle: Who Turns the Crank?. <i>Current Biology</i> , 2011, 21, R185-R187. | 3.9 | 9 |
| 61 | The cell cycle. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2011, 366, 3494-3497. | 4.0 | 47 |
| 62 | Computational modelling of mitotic exit in budding yeast: the role of separase and Cdc14 endocycles. <i>Journal of the Royal Society Interface</i> , 2011, 8, 1128-1141. | 3.4 | 24 |
| 63 | Switches and latches: a biochemical tug-of-war between the kinases and phosphatases that control mitosis. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2011, 366, 3584-3594. | 4.0 | 95 |
| 64 | System-level feedbacks make the anaphase switch irreversible. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 10016-10021. | 7.1 | 55 |
| 65 | Systems-level feedback in cell-cycle control. <i>Biochemical Society Transactions</i> , 2010, 38, 1242-1246. | 3.4 | 12 |
| 66 | Hypoxia-dependent sequestration of an oxygen sensor by a widespread structural motif can shape the hypoxic response - a predictive kinetic model. <i>BMC Systems Biology</i> , 2010, 4, 139. | 3.0 | 44 |
| 67 | Regulated protein kinases and phosphatases in cell cycle decisions. <i>Current Opinion in Cell Biology</i> , 2010, 22, 801-808. | 5.4 | 54 |
| 68 | Restriction point control of the mammalian cell cycle via the cyclin E/Cdk2:p27 complex. <i>FEBS Journal</i> , 2010, 277, 357-367. | 4.7 | 44 |
| 69 | The regulatory network of cell-cycle progression is fundamentally different in plants versus yeast or metazoans. <i>Plant Signaling and Behavior</i> , 2010, 5, 1613-1618. | 2.4 | 24 |
| 70 | CULLIN 4-RING FINGER-LIGASE plays a key role in the control of endoreplication cycles in <i>Arabidopsis</i> trichomes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 15275-15280. | 7.1 | 44 |
| 71 | Different effects of redundant feedback loops on a bistable switch. <i>Chaos</i> , 2010, 20, 045120. | 2.5 | 17 |
| 72 | A model of yeast cell cycle regulation based on multisite phosphorylation. <i>Molecular Systems Biology</i> , 2010, 6, 405. | 7.2 | 97 |

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|----|---|------|-----------|
| 73 | Functional Motifs in Biochemical Reaction Networks. Annual Review of Physical Chemistry, 2010, 61, 219-240. | 10.8 | 257 |
| 74 | Cell cycle regulation by feedâ€forward loops coupling transcription and phosphorylation. Molecular Systems Biology, 2009, 5, 236. | 7.2 | 44 |
| 75 | Mitotic exit in mammalian cells. Molecular Systems Biology, 2009, 5, 324. | 7.2 | 15 |
| 76 | The Influence of Catalysis on Mad2 Activation Dynamics. PLoS Biology, 2009, 7, e1000010. | 5.6 | 97 |
| 77 | Control of Cell Proliferation, Organ Growth, and DNA Damage Response Operate Independently of Dephosphorylation of the <i>Arabidopsis</i> Cdk1 Homolog CDKA;1 Å. Plant Cell, 2009, 21, 3641-3654. | 6.6 | 106 |
| 78 | Bistability by multiple phosphorylation of regulatory proteins. Progress in Biophysics and Molecular Biology, 2009, 100, 47-56. | 2.9 | 74 |
| 79 | Regulation of APC/C Activity in Oocytes by a Bub1-Dependent Spindle Assembly Checkpoint. Current Biology, 2009, 19, 369-380. | 3.9 | 194 |
| 80 | Systemâ€level feedbacks control cell cycle progression. FEBS Letters, 2009, 583, 3992-3998. | 2.8 | 38 |
| 81 | Irreversibility of mitotic exit is the consequence of systems-level feedback. Nature, 2009, 459, 592-595. | 27.8 | 91 |
| 82 | Spatial controls for growth zone formation during the fission yeast cell cycle. Yeast, 2008, 25, 59-69. | 1.7 | 31 |
| 83 | Analysis of a budding yeast cell cycle model using the shapes of local sensitivity functions. International Journal of Chemical Kinetics, 2008, 40, 710-720. | 1.6 | 17 |
| 84 | Antagonism and bistability in protein interaction networks. Journal of Theoretical Biology, 2008, 250, 209-218. | 1.7 | 66 |
| 85 | Design principles of biochemical oscillators. Nature Reviews Molecular Cell Biology, 2008, 9, 981-991. | 37.0 | 970 |
| 86 | Temporal Organization of the Cell Cycle. Current Biology, 2008, 18, R759-R768. | 3.9 | 165 |
| 87 | Reverse Engineering Models of Cell Cycle Regulation. Advances in Experimental Medicine and Biology, 2008, 641, 88-97. | 1.6 | 15 |
| 88 | Microtubules offset growth site from the cell centre in fission yeast. Journal of Cell Science, 2007, 120, 2205-2213. | 2.0 | 27 |
| 89 | Dynamical modeling of syncytial mitotic cycles in <i>Drosophila</i> embryos. Molecular Systems Biology, 2007, 3, 131. | 7.2 | 41 |
| 90 | Mitotic exit in two dimensions. Journal of Theoretical Biology, 2007, 248, 560-573. | 1.7 | 21 |

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|-----|---|------|-----------|
| 91 | Irreversible cell-cycle transitions are due to systems-level feedback. <i>Nature Cell Biology</i> , 2007, 9, 724-728. | 10.3 | 178 |
| 92 | Modeling the septation initiation network (SIN) in fission yeast cells. <i>Current Genetics</i> , 2007, 51, 245-255. | 1.7 | 27 |
| 93 | Analysis of a Generic Model of Eukaryotic Cell-Cycle Regulation. <i>Biophysical Journal</i> , 2006, 90, 4361-4379. | 0.5 | 226 |
| 94 | Downregulation of PP2ACdc55 Phosphatase by Separase Initiates Mitotic Exit in Budding Yeast. <i>Cell</i> , 2006, 125, 719-732. | 28.9 | 230 |
| 95 | Time scale and dimension analysis of a budding yeast cell cycle model. <i>BMC Bioinformatics</i> , 2006, 7, 494. | 2.6 | 34 |
| 96 | Steady States and Oscillations in the p53/Mdm2 Network. <i>Cell Cycle</i> , 2005, 4, 488-493. | 2.6 | 221 |
| 97 | Rewiring the Exit from Mitosis. <i>Cell Cycle</i> , 2005, 4, 4107-4112. | 2.6 | 17 |
| 98 | Rewiring the exit from mitosis. <i>Cell Cycle</i> , 2005, 4, 1107-12. | 2.6 | 8 |
| 99 | Modelling the fission yeast cell cycle. <i>Briefings in Functional Genomics & Proteomics</i> , 2004, 2, 298-307. | 3.8 | 36 |
| 100 | Integrative Analysis of Cell Cycle Control in Budding Yeast. <i>Molecular Biology of the Cell</i> , 2004, 15, 3841-3862. | 2.1 | 584 |
| 101 | A model for restriction point control of the mammalian cell cycle. <i>Journal of Theoretical Biology</i> , 2004, 230, 563-579. | 1.7 | 272 |
| 102 | Sniffers, buzzers, toggles and blinkers: dynamics of regulatory and signaling pathways in the cell. <i>Current Opinion in Cell Biology</i> , 2003, 15, 221-231. | 5.4 | 1,423 |
| 103 | Mathematical model of the morphogenesis checkpoint in budding yeast. <i>Journal of Cell Biology</i> , 2003, 163, 1243-1254. | 5.2 | 78 |
| 104 | Modelling the controls of the eukaryotic cell cycle. <i>Biochemical Society Transactions</i> , 2003, 31, 1526-1529. | 3.4 | 65 |
| 105 | The dynamics of cell cycle regulation. <i>BioEssays</i> , 2002, 24, 1095-1109. | 2.5 | 277 |
| 106 | Morphogenetic checkpoint in fission yeast? Yes!. <i>Microbiology (United Kingdom)</i> , 2002, 148, 2270-2271. | 1.8 | 2 |
| 107 | Mathematical model of the cell division cycle of fission yeast. <i>Chaos</i> , 2001, 11, 277. | 2.5 | 144 |
| 108 | A stochastic, molecular model of the fission yeast cell cycle: role of the nucleocytoplasmic ratio in cycle time regulation. <i>Biophysical Chemistry</i> , 2001, 92, 1-15. | 2.8 | 56 |

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|-----|--|------|-----------|
| 109 | Regulation of the Eukaryotic Cell Cycle: Molecular Antagonism, Hysteresis, and Irreversible Transitions. <i>Journal of Theoretical Biology</i> , 2001, 210, 249-263. | 1.7 | 328 |
| 110 | Network dynamics and cell physiology. <i>Nature Reviews Molecular Cell Biology</i> , 2001, 2, 908-916. | 37.0 | 481 |
| 111 | Kinetic Analysis of a Molecular Model of the Budding Yeast Cell Cycle. <i>Molecular Biology of the Cell</i> , 2000, 11, 369-391. | 2.1 | 437 |
| 112 | Modeling the fission yeast cell cycle: Quantized cycle times in <i>wee1-cdc25Delta</i> mutant cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2000, 97, 7865-7870. | 7.1 | 117 |
| 113 | Finishing the Cell Cycle. <i>Journal of Theoretical Biology</i> , 1999, 199, 223-233. | 1.7 | 50 |
| 114 | A Simple Model of Circadian Rhythms Based on Dimerization and Proteolysis of PER and TIM. <i>Biophysical Journal</i> , 1999, 77, 2411-2417. | 0.5 | 168 |
| 115 | Modeling M-phase control in <i>Xenopus</i> oocyte extracts: the surveillance mechanism for unreplicated DNA. <i>Biophysical Chemistry</i> , 1998, 72, 169-184. | 2.8 | 57 |
| 116 | Mathematical model of the fission yeast cell cycle with checkpoint controls at the G1/S, G2/M and metaphase/anaphase transitions. <i>Biophysical Chemistry</i> , 1998, 72, 185-200. | 2.8 | 121 |
| 117 | Modeling the control of DNA replication in fission yeast. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1997, 94, 9147-9152. | 7.1 | 165 |
| 118 | Chemical kinetic theory: understanding cell-cycle regulation. <i>Trends in Biochemical Sciences</i> , 1996, 21, 89-96. | 7.5 | 92 |
| 119 | Quantitative analysis of a molecular model of mitotic control in fission yeast. <i>Journal of Theoretical Biology</i> , 1995, 173, 283-305. | 1.7 | 93 |
| 120 | Checkpoints in the cell cycle from a modeler's perspective. , 1995, 1, 1-8. | | 29 |
| 121 | Modeling the Cell Division Cycle: M-phase Trigger, Oscillations, and Size Control. <i>Journal of Theoretical Biology</i> , 1993, 165, 101-134. | 1.7 | 142 |
| 122 | Systems biology of the yeast cell cycle engine. , 0, , 305-324. | | 4 |