

# 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	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
2	Design principles of biochemical oscillators. <i>Nature Reviews Molecular Cell Biology</i> , 2008, 9, 981-991.	37.0	970
3	Integrative Analysis of Cell Cycle Control in Budding Yeast. <i>Molecular Biology of the Cell</i> , 2004, 15, 3841-3862.	2.1	584
4	Network dynamics and cell physiology. <i>Nature Reviews Molecular Cell Biology</i> , 2001, 2, 908-916.	37.0	481
5	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
6	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
7	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
8	The dynamics of cell cycle regulation. <i>BioEssays</i> , 2002, 24, 1095-1109.	2.5	277
9	A model for restriction point control of the mammalian cell cycle. <i>Journal of Theoretical Biology</i> , 2004, 230, 563-579.	1.7	272
10	Functional Motifs in Biochemical Reaction Networks. <i>Annual Review of Physical Chemistry</i> , 2010, 61, 219-240.	10.8	257
11	Downregulation of PP2A <sup>Cdc55</sup> Phosphatase by Separase Initiates Mitotic Exit in Budding Yeast. <i>Cell</i> , 2006, 125, 719-732.	28.9	230
12	Analysis of a Generic Model of Eukaryotic Cell-Cycle Regulation. <i>Biophysical Journal</i> , 2006, 90, 4361-4379.	0.5	226
13	Steady States and Oscillations in the p53/Mdm2 Network. <i>Cell Cycle</i> , 2005, 4, 488-493.	2.6	221
14	Regulation of APC/C Activity in Oocytes by a Bub1-Dependent Spindle Assembly Checkpoint. <i>Current Biology</i> , 2009, 19, 369-380.	3.9	194
15	Irreversible cell-cycle transitions are due to systems-level feedback. <i>Nature Cell Biology</i> , 2007, 9, 724-728.	10.3	178
16	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
17	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
18	Temporal Organization of the Cell Cycle. <i>Current Biology</i> , 2008, 18, R759-R768.	3.9	165

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19	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
20	Mathematical model of the cell division cycle of fission yeast. <i>Chaos</i> , 2001, 11, 277.	2.5	144
21	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
22	Phosphorylation network dynamics in the control of cell cycle transitions. <i>Journal of Cell Science</i> , 2012, 125, 4703-4711.	2.0	138
23	The BEG (PP2A-B55/ENSA/Greatwall) Pathway Ensures Cytokinesis follows Chromosome Separation. <i>Molecular Cell</i> , 2013, 52, 393-405.	9.7	136
24	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
25	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
26	A Dynamical Framework for the All-or-None G1/S Transition. <i>Cell Systems</i> , 2016, 2, 27-37.	6.2	115
27	Control of Cell Proliferation, Organ Growth, and DNA Damage Response Operate Independently of Dephosphorylation of the <i>Arabidopsis</i> Cdk1 Homolog CDKA;1. <i>Plant Cell</i> , 2009, 21, 3641-3654.	6.6	106
28	A General G1/S-Phase Cell-Cycle Control Module in the Flowering Plant <i>Arabidopsis thaliana</i> . <i>PLoS Genetics</i> , 2012, 8, e1002847.	3.5	103
29	The Influence of Catalysis on Mad2 Activation Dynamics. <i>PLoS Biology</i> , 2009, 7, e1000010.	5.6	97
30	A model of yeast cell cycle regulation based on multisite phosphorylation. <i>Molecular Systems Biology</i> , 2010, 6, 405.	7.2	97
31	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
32	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
33	Meiotic Prophase Requires Proteolysis of M Phase Regulators Mediated by the Meiosis-Specific APC/C <sup>Ama1</sup> . <i>Cell</i> , 2012, 151, 603-618.	28.9	93
34	Chemical kinetic theory: understanding cell-cycle regulation. <i>Trends in Biochemical Sciences</i> , 1996, 21, 89-96.	7.5	92
35	Irreversibility of mitotic exit is the consequence of systems-level feedback. <i>Nature</i> , 2009, 459, 592-595.	27.8	91
36	Nutritional Control of Cell Size by the Greatwall-Endosulfine-PP2A-B55 Pathway. <i>Current Biology</i> , 2016, 26, 319-330.	3.9	87

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37	Absolute quantification of cohesin, CTCF and their regulators in human cells. <i>ELife</i> , 2019, 8, .	6.0	79
38	Mathematical model of the morphogenesis checkpoint in budding yeast. <i>Journal of Cell Biology</i> , 2003, 163, 1243-1254.	5.2	78
39	Bistability by multiple phosphorylation of regulatory proteins. <i>Progress in Biophysics and Molecular Biology</i> , 2009, 100, 47-56.	2.9	74
40	Two Bistable Switches Govern M Phase Entry. <i>Current Biology</i> , 2016, 26, 3361-3367.	3.9	72
41	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
42	Antagonism and bistability in protein interaction networks. <i>Journal of Theoretical Biology</i> , 2008, 250, 209-218.	1.7	66
43	Modelling the controls of the eukaryotic cell cycle. <i>Biochemical Society Transactions</i> , 2003, 31, 1526-1529.	3.4	65
44	Dependency of the Spindle Assembly Checkpoint on Cdk1 Renders the Anaphase Transition Irreversible. <i>Current Biology</i> , 2014, 24, 630-637.	3.9	63
45	Kinetochores—microtubule error correction is driven by differentially regulated interaction modes. <i>Nature Cell Biology</i> , 2015, 17, 421-433.	10.3	63
46	Molecular mechanisms creating bistable switches at cell cycle transitions. <i>Open Biology</i> , 2013, 3, 120179.	3.6	62
47	Two Interlinked Bistable Switches Govern Mitotic Control in Mammalian Cells. <i>Current Biology</i> , 2018, 28, 3824-3832.e6.	3.9	62
48	Models in biology: lessons from modeling regulation of the eukaryotic cell cycle. <i>BMC Biology</i> , 2015, 13, 46.	3.8	61
49	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
50	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
51	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
52	PP2A/B55 and Fcp1 Regulate Greatwall and Ensa Dephosphorylation during Mitotic Exit. <i>PLoS Genetics</i> , 2014, 10, e1004004.	3.5	55
53	Regulated protein kinases and phosphatases in cell cycle decisions. <i>Current Opinion in Cell Biology</i> , 2010, 22, 801-808.	5.4	54
54	A Dynamical Paradigm for Molecular Cell Biology. <i>Trends in Cell Biology</i> , 2020, 30, 504-515.	7.9	53

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55	Cyclin A triggers Mitosis either via the Greatwall kinase pathway or Cyclin B. EMBO Journal, 2020, 39, e104419.	7.8	52
56	Finishing the Cell Cycle. Journal of Theoretical Biology, 1999, 199, 223-233.	1.7	50
57	Cell Cycle Control by a Minimal Cdk Network. PLoS Computational Biology, 2015, 11, e1004056.	3.2	49
58	Human Chromosome Segregation Involves Multi-Layered Regulation of Separase by the Peptidyl-Prolyl-Isomerase Pin1. Molecular Cell, 2015, 58, 495-506.	9.7	49
59	Protein Phosphatase 2A Controls the Order and Dynamics of Cell-Cycle Transitions. Molecular Cell, 2011, 44, 437-450.	9.7	47
60	The cell cycle. Philosophical Transactions of the Royal Society B: Biological Sciences, 2011, 366, 3494-3497.	4.0	47
61	Spatiotemporal dynamics of Spc105 regulates the assembly of the Drosophila kinetochore. Open Biology, 2012, 2, 110032.	3.6	47
62	Dilution and titration of cell-cycle regulators may control cell size in budding yeast. PLoS Computational Biology, 2018, 14, e1006548.	3.2	45
63	CDK1-CCNB1 creates a spindle checkpoint "permissive state by enabling MPS1 kinetochore localization. Journal of Cell Biology, 2019, 218, 1182-1199.	5.2	45
64	Cell cycle regulation by feed-forward loops coupling transcription and phosphorylation. Molecular Systems Biology, 2009, 5, 236.	7.2	44
65	Hypoxia-dependent sequestration of an oxygen sensor by a widespread structural motif can shape the hypoxic response - a predictive kinetic model. BMC Systems Biology, 2010, 4, 139.	3.0	44
66	Restriction point control of the mammalian cell cycle via the cyclin E/Cdk2:p27 complex. FEBS Journal, 2010, 277, 357-367.	4.7	44
67	CULLIN 4-RING FINGER-LIGASE plays a key role in the control of endoreplication cycles in Arabidopsis trichomes. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 15275-15280.	7.1	44
68	Dynamical modeling of syncytial mitotic cycles in <i>Drosophila</i> embryos. Molecular Systems Biology, 2007, 3, 131.	7.2	41
69	System-level feedbacks control cell cycle progression. FEBS Letters, 2009, 583, 3992-3998.	2.8	38
70	Modelling the fission yeast cell cycle. Briefings in Functional Genomics & Proteomics, 2004, 2, 298-307.	3.8	36
71	Time scale and dimension analysis of a budding yeast cell cycle model. BMC Bioinformatics, 2006, 7, 494.	2.6	34
72	Spatial controls for growth zone formation during the fission yeast cell cycle. Yeast, 2008, 25, 59-69.	1.7	31

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73	Control of cell growth, division and death: information processing in living cells. <i>Interface Focus</i> , 2014, 4, 20130070.	3.0	31
74	Checkpoints in the cell cycle from a modeler's perspective. , 1995, 1, 1-8.		29
75	Microtubules offset growth site from the cell centre in fission yeast. <i>Journal of Cell Science</i> , 2007, 120, 2205-2213.	2.0	27
76	Modeling the septation initiation network (SIN) in fission yeast cells. <i>Current Genetics</i> , 2007, 51, 245-255.	1.7	27
77	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
78	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
79	Interlinked bistable mechanisms generate robust mitotic transitions. <i>Cell Cycle</i> , 2017, 16, 1885-1892.	2.6	23
80	Mitotic exit in two dimensions. <i>Journal of Theoretical Biology</i> , 2007, 248, 560-573.	1.7	21
81	Model scenarios for switch-like mitotic transitions. <i>FEBS Letters</i> , 2015, 589, 667-671.	2.8	21
82	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
83	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
84	A Structural Systems Biology Approach for Quantifying the Systemic Consequences of Missense Mutations in Proteins. <i>PLoS Computational Biology</i> , 2012, 8, e1002738.	3.2	19
85	Multisite phosphoregulation of Cdc25 activity refines the mitotic entrance and exit switches. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 9899-9904.	7.1	19
86	Rewiring the Exit from Mitosis. <i>Cell Cycle</i> , 2005, 4, 4107-4112.	2.6	17
87	Analysis of a budding yeast cell cycle model using the shapes of local sensitivity functions. <i>International Journal of Chemical Kinetics</i> , 2008, 40, 710-720.	1.6	17
88	Different effects of redundant feedback loops on a bistable switch. <i>Chaos</i> , 2010, 20, 045120.	2.5	17
89	Robust mitotic entry is ensured by a latching switch. <i>Biology Open</i> , 2013, 2, 924-931.	1.2	17
90	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

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91	The role of APC/C inhibitor Emi2/XErp1 in oscillatory dynamics of early embryonic cell cycles. <i>Biophysical Chemistry</i> , 2013, 177-178, 1-6.	2.8	16
92	Cell-cycle transitions: a common role for stoichiometric inhibitors. <i>Molecular Biology of the Cell</i> , 2017, 28, 3437-3446.	2.1	16
93	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
94	Mitotic exit in mammalian cells. <i>Molecular Systems Biology</i> , 2009, 5, 324.	7.2	15
95	Dynamical Scenarios for Chromosome Bi-orientation. <i>Biophysical Journal</i> , 2013, 104, 2595-2606.	0.5	15
96	microRNA as a Potential Vector for the Propagation of Robustness in Protein Expression and Oscillatory Dynamics within a ceRNA Network. <i>PLoS ONE</i> , 2013, 8, e83372.	2.5	15
97	Reverse Engineering Models of Cell Cycle Regulation. <i>Advances in Experimental Medicine and Biology</i> , 2008, 641, 88-97.	1.6	15
98	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
99	Cell cycle commitment in budding yeast emerges from the cooperation of multiple bistable switches. <i>Open Biology</i> , 2011, 1, 110009.	3.6	14
100	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
101	Irreversible Transitions, Bistability and Checkpoint Controls in the Eukaryotic Cell Cycle. , 2013, , 265-285.		13
102	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
103	Systems-level feedback in cell-cycle control. <i>Biochemical Society Transactions</i> , 2010, 38, 1242-1246.	3.4	12
104	Time-keeping and decision-making in the cell cycle. <i>Interface Focus</i> , 2022, 12, .	3.0	10
105	Cell Cycle: Who Turns the Crank?. <i>Current Biology</i> , 2011, 21, R185-R187.	3.9	9
106	CDK-Dependent Nuclear Localization of B-Cyclin Clb1 Promotes FEAR Activation during Meiosis I in Budding Yeast. <i>PLoS ONE</i> , 2013, 8, e79001.	2.5	9
107	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
108	Rewiring the exit from mitosis. <i>Cell Cycle</i> , 2005, 4, 1107-12.	2.6	8

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109	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
110	Bistability, Oscillations, and Traveling Waves in Frog Egg Extracts. Bulletin of Mathematical Biology, 2015, 77, 796-816.	1.9	7
111	Mathematical Model for Growth Regulation of Fission Yeast <i>Schizosaccharomyces pombe</i> . PLoS ONE, 2012, 7, e49675.	2.5	6
112	Interplay of transcriptional and proteolytic regulation in driving robust cell cycle progression. Molecular BioSystems, 2012, 8, 863.	2.9	5
113	Mitotic kinase oscillation governs the latching of cell cycle switches. Current Biology, 2022, 32, 2780-2785.e2.	3.9	5
114	Systems biology of the yeast cell cycle engine. , 0, , 305-324.		4
115	Systems-level feedback regulation of cell cycle transitions in <i>Ostreococcus tauri</i> . Plant Physiology and Biochemistry, 2018, 126, 39-46.	5.8	2
116	Morphogenetic checkpoint in fission yeast? Yes!. Microbiology (United Kingdom), 2002, 148, 2270-2271.	1.8	2
117	Cell Cycle, Budding Yeast. , 2013, , 337-341.		2
118	Pom1 is not the size ruler. Cell Cycle, 2013, 12, 3463-3464.	2.6	1
119	Cell Cycle Dynamics, Bistability and Oscillations. , 2013, , 263-270.		1
120	Computational modeling of chromosome re-replication in mutant strains of fission yeast. Molecular Biology of the Cell, 2021, 32, 830-841.	2.1	0
121	Cell Cycle Transitions, Mitotic Exit. , 2013, , 333-336.		0
122	Cell Cycle Dynamics, Irreversibility. , 2013, , 270-273.		0