

Bela Novak

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

119
papers

9,294
citations

47
h-index

95
g-index

139
ext. papers

10,772
ext. citations

8.4
avg, IF

6.32
L-index

#	Paper	IF	Citations
119	Mechanisms of signalling-memory governing progression through the eukaryotic cell cycle. <i>Current Opinion in Cell Biology</i> , 2021 , 69, 7-16	9	7
118	Computational modeling of chromosome re-replication in mutant strains of fission yeast. <i>Molecular Biology of the Cell</i> , 2021 , 32, 830-841	3.5	
117	Cyclin A triggers Mitosis either via the Greatwall kinase pathway or Cyclin B. <i>EMBO Journal</i> , 2020 , 39, e104419	13	25
116	A Dynamical Paradigm for Molecular Cell Biology. <i>Trends in Cell Biology</i> , 2020 , 30, 504-515	18.3	27
115	A Single Light-Responsive Sizer Can Control Multiple-Fission Cycles in <i>Chlamydomonas</i> . <i>Current Biology</i> , 2020 , 30, 634-644.e7	6.3	6
114	CDK1-CCNB1 creates a spindle checkpoint-permissive state by enabling MPS1 kinetochore localization. <i>Journal of Cell Biology</i> , 2019 , 218, 1182-1199	7.3	27
113	Absolute quantification of cohesin, CTCF and their regulators in human cells. <i>ELife</i> , 2019 , 8,	8.9	44
112	Systems-level feedback regulation of cell cycle transitions in <i>Ostreococcus tauri</i> . <i>Plant Physiology and Biochemistry</i> , 2018 , 126, 39-46	5.4	0
111	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	11.5	44
110	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	3.2	9
109	Two Interlinked Bistable Switches Govern Mitotic Control in Mammalian Cells. <i>Current Biology</i> , 2018 , 28, 3824-3832.e6	6.3	32
108	Dilution and titration of cell-cycle regulators may control cell size in budding yeast. <i>PLoS Computational Biology</i> , 2018 , 14, e1006548	5	28
107	APC/C 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	6.3	5
106	DNA damage during S-phase mediates the proliferation-quiescence decision in the subsequent G1 via p21 expression. <i>Nature Communications</i> , 2017 , 8, 14728	17.4	176
105	Cell-cycle transitions: a common role for stoichiometric inhibitors. <i>Molecular Biology of the Cell</i> , 2017 , 28, 3437-3446	3.5	12
104	Interlinked bistable mechanisms generate robust mitotic transitions. <i>Cell Cycle</i> , 2017 , 16, 1885-1892	4.7	12
103	A Dynamical Framework for the All-or-None G1/S Transition. <i>Cell Systems</i> , 2016 , 2, 27-37	10.6	77

102	A PP2A-B55 recognition signal controls substrate dephosphorylation kinetics during mitotic exit. <i>Journal of Cell Biology</i> , 2016 , 214, 539-54	7.3	105
101	Two Bistable Switches Govern M Phase Entry. <i>Current Biology</i> , 2016 , 26, 3361-3367	6.3	48
100	Nutritional Control of Cell Size by the Greatwall-Endosulfine-PP2A/B55 Pathway. <i>Current Biology</i> , 2016 , 26, 319-30	6.3	63
99	Bistability, oscillations, and traveling waves in frog egg extracts. <i>Bulletin of Mathematical Biology</i> , 2015 , 77, 796-816	2.1	6
98	Cell cycle control by a minimal Cdk network. <i>PLoS Computational Biology</i> , 2015 , 11, e1004056	5	41
97	Model scenarios for switch-like mitotic transitions. <i>FEBS Letters</i> , 2015 , 589, 667-71	3.8	15
96	Kinetochore-microtubule error correction is driven by differentially regulated interaction modes. <i>Nature Cell Biology</i> , 2015 , 17, 421-33	23.4	47
95	Human chromosome segregation involves multi-layered regulation of separase by the peptidyl-prolyl-isomerase Pin1. <i>Molecular Cell</i> , 2015 , 58, 495-506	17.6	37
94	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	10.6	13
93	Models in biology: lessons from modeling regulation of the eukaryotic cell cycle. <i>BMC Biology</i> , 2015 , 13, 46	7.3	43
92	Control of cell growth, division and death: information processing in living cells. <i>Interface Focus</i> , 2014 , 4, 20130070	3.9	24
91	Dependency of the spindle assembly checkpoint on Cdk1 renders the anaphase transition irreversible. <i>Current Biology</i> , 2014 , 24, 630-7	6.3	53
90	A model for the epigenetic switch linking inflammation to cell transformation: deterministic and stochastic approaches. <i>PLoS Computational Biology</i> , 2014 , 10, e1003455	5	13
89	PP2A/B55 and Fcp1 regulate Greatwall and Ensa dephosphorylation during mitotic exit. <i>PLoS Genetics</i> , 2014 , 10, e1004004	6	49
88	Dynamical scenarios for chromosome bi-orientation. <i>Biophysical Journal</i> , 2013 , 104, 2595-606	2.9	11
87	Minimal models for cell-cycle control based on competitive inhibition and multisite phosphorylations of Cdk substrates. <i>Biophysical Journal</i> , 2013 , 104, 1367-79	2.9	13
86	The BEG (PP2A-B55/ENSA/Greatwall) pathway ensures cytokinesis follows chromosome separation. <i>Molecular Cell</i> , 2013 , 52, 393-405	17.6	107
85	Role for regulated phosphatase activity in generating mitotic oscillations in <i>Xenopus</i> cell-free extracts. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013 , 110, 20539-44	11.5	3

84	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	3.5	15
83	Irreversible Transitions, Bistability and Checkpoint Controls in the Eukaryotic Cell Cycle: A Systems-Level Understanding 2013 , 265-285		9
82	Molecular mechanisms creating bistable switches at cell cycle transitions. <i>Open Biology</i> , 2013 , 3, 1201797		48
81	Robust mitotic entry is ensured by a latching switch. <i>Biology Open</i> , 2013 , 2, 924-31	2.2	12
80	Pom1 is not the size ruler. <i>Cell Cycle</i> , 2013 , 12, 3463-4	4.7	1
79	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	3.7	12
78	CDK-dependent nuclear localization of B-cyclin Clb1 promotes FEAR activation during meiosis I in budding yeast. <i>PLoS ONE</i> , 2013 , 8, e79001	3.7	6
77	Meiotic prophase requires proteolysis of M phase regulators mediated by the meiosis-specific APC/C _{Am1} . <i>Cell</i> , 2012 , 151, 603-18	56.2	62
76	Interplay of transcriptional and proteolytic regulation in driving robust cell cycle progression. <i>Molecular BioSystems</i> , 2012 , 8, 863-70		4
75	Phosphorylation network dynamics in the control of cell cycle transitions. <i>Journal of Cell Science</i> , 2012 , 125, 4703-11	5.3	115
74	Mathematical model for growth regulation of fission yeast <i>Schizosaccharomyces pombe</i> . <i>PLoS ONE</i> , 2012 , 7, e49675	3.7	6
73	Spatiotemporal dynamics of Spc105 regulates the assembly of the <i>Drosophila</i> kinetochore. <i>Open Biology</i> , 2012 , 2, 110032	7	28
72	A structural systems biology approach for quantifying the systemic consequences of missense mutations in proteins. <i>PLoS Computational Biology</i> , 2012 , 8, e1002738	5	17
71	A general G1/S-phase cell-cycle control module in the flowering plant <i>Arabidopsis thaliana</i> . <i>PLoS Genetics</i> , 2012 , 8, e1002847	6	86
70	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, 9899-904	11.5	13
69	Protein phosphatase 2A controls the order and dynamics of cell-cycle transitions. <i>Molecular Cell</i> , 2011 , 44, 437-50	17.6	31
68	Overexpression limits of fission yeast cell-cycle regulators in vivo and in silico. <i>Molecular Systems Biology</i> , 2011 , 7, 556	12.2	12
67	Cell cycle commitment in budding yeast emerges from the cooperation of multiple bistable switches. <i>Open Biology</i> , 2011 , 1, 110009	7	13

66	Cell cycle: who turns the crank?. <i>Current Biology</i> , 2011 , 21, R185-7	6.3	8
65	The cell cycle. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2011 , 366, 3494-7	5.8	33
64	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, 1128-41	4.1	21
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, 3584-94	5.8	89
62	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, 10016-21	11.5	43
61	Restriction point control of the mammalian cell cycle via the cyclin E/Cdk2:p27 complex. <i>FEBS Journal</i> , 2010 , 277, 357-67	5.7	37
60	The regulatory network of cell-cycle progression is fundamentally different in plants versus yeast or metazoans. <i>Plant Signaling and Behavior</i> , 2010 , 5, 1613-8	2.5	21
59	Cullin 4-ring finger-ligase plays a key role in the control of endoreplication cycles in Arabidopsis trichomes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010 , 107, 15275-80	11.5	38
58	Different effects of redundant feedback loops on a bistable switch. <i>Chaos</i> , 2010 , 20, 045120	3.3	15
57	A model of yeast cell-cycle regulation based on multisite phosphorylation. <i>Molecular Systems Biology</i> , 2010 , 6, 405	12.2	88
56	Functional motifs in biochemical reaction networks. <i>Annual Review of Physical Chemistry</i> , 2010 , 61, 219-40	5.7	212
55	Systems-level feedback in cell-cycle control. <i>Biochemical Society Transactions</i> , 2010 , 38, 1242-6	5.1	10
54	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	3.5	35
53	Regulated protein kinases and phosphatases in cell cycle decisions. <i>Current Opinion in Cell Biology</i> , 2010 , 22, 801-8	9	41
52	Cell cycle regulation by feed-forward loops coupling transcription and phosphorylation. <i>Molecular Systems Biology</i> , 2009 , 5, 236	12.2	35
51	Mitotic exit in mammalian cells. <i>Molecular Systems Biology</i> , 2009 , 5, 324	12.2	12
50	The influence of catalysis on mad2 activation dynamics. <i>PLoS Biology</i> , 2009 , 7, e10	9.7	79
49	Control of cell proliferation, organ growth, and DNA damage response operate independently of dephosphorylation of the Arabidopsis Cdk1 homolog CDKA;1. <i>Plant Cell</i> , 2009 , 21, 3641-54	11.6	92

48	Bistability by multiple phosphorylation of regulatory proteins. <i>Progress in Biophysics and Molecular Biology</i> , 2009 , 100, 47-56	4.7	62
47	Regulation of APC/C activity in oocytes by a Bub1-dependent spindle assembly checkpoint. <i>Current Biology</i> , 2009 , 19, 369-80	6.3	166
46	System-level feedbacks control cell cycle progression. <i>FEBS Letters</i> , 2009 , 583, 3992-8	3.8	30
45	Irreversibility of mitotic exit is the consequence of systems-level feedback. <i>Nature</i> , 2009 , 459, 592-5	50.4	73
44	Design principles of biochemical oscillators. <i>Nature Reviews Molecular Cell Biology</i> , 2008 , 9, 981-91	48.7	757
43	Temporal organization of the cell cycle. <i>Current Biology</i> , 2008 , 18, R759-R768	6.3	131
42	Spatial controls for growth zone formation during the fission yeast cell cycle. <i>Yeast</i> , 2008 , 25, 59-69	3.4	29
41	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	1.4	13
40	Antagonism and bistability in protein interaction networks. <i>Journal of Theoretical Biology</i> , 2008 , 250, 209-18	2.3	56
39	Reverse engineering models of cell cycle regulation. <i>Advances in Experimental Medicine and Biology</i> , 2008 , 641, 88-97	3.6	11
38	Mitotic exit in two dimensions. <i>Journal of Theoretical Biology</i> , 2007 , 248, 560-73	2.3	19
37	Irreversible cell-cycle transitions are due to systems-level feedback. <i>Nature Cell Biology</i> , 2007 , 9, 724-8	23.4	146
36	Modeling the septation initiation network (SIN) in fission yeast cells. <i>Current Genetics</i> , 2007 , 51, 245-55	2.9	23
35	Microtubules offset growth site from the cell centre in fission yeast. <i>Journal of Cell Science</i> , 2007 , 120, 2205-13	5.3	26
34	Dynamical modeling of syncytial mitotic cycles in <i>Drosophila</i> embryos. <i>Molecular Systems Biology</i> , 2007 , 3, 131	12.2	37
33	Time scale and dimension analysis of a budding yeast cell cycle model. <i>BMC Bioinformatics</i> , 2006 , 7, 494	3.6	26
32	Analysis of a generic model of eukaryotic cell-cycle regulation. <i>Biophysical Journal</i> , 2006 , 90, 4361-79	2.9	189
31	Downregulation of PP2A(Cdc55) phosphatase by separase initiates mitotic exit in budding yeast. <i>Cell</i> , 2006 , 125, 719-32	56.2	201

30	Rewiring the Exit from Mitosis. <i>Cell Cycle</i> , 2005 , 4, 4107-4112	4.7	15
29	Steady states and oscillations in the p53/Mdm2 network. <i>Cell Cycle</i> , 2005 , 4, 488-93	4.7	182
28	Rewiring the exit from mitosis. <i>Cell Cycle</i> , 2005 , 4, 1107-12	4.7	8
27	Modelling the fission yeast cell cycle. <i>Briefings in Functional Genomics & Proteomics</i> , 2004 , 2, 298-307		34
26	Integrative analysis of cell cycle control in budding yeast. <i>Molecular Biology of the Cell</i> , 2004 , 15, 3841-62,5	3.5	478
25	A model for restriction point control of the mammalian cell cycle. <i>Journal of Theoretical Biology</i> , 2004 , 230, 563-79	2.3	227
24	Checkpoints in the Cell Cycle 2003 ,		5
23	Modelling the controls of the eukaryotic cell cycle. <i>Biochemical Society Transactions</i> , 2003 , 31, 1526-9	5.1	54
22	Sniffers, buzzers, toggles and blinkers: dynamics of regulatory and signaling pathways in the cell. <i>Current Opinion in Cell Biology</i> , 2003 , 15, 221-31	9	1165
21	Mathematical model of the morphogenesis checkpoint in budding yeast. <i>Journal of Cell Biology</i> , 2003 , 163, 1243-54	7.3	67
20	The dynamics of cell cycle regulation. <i>BioEssays</i> , 2002 , 24, 1095-109	4.1	236
19	Morphogenetic checkpoint in fission yeast? Yes!. <i>Microbiology (United Kingdom)</i> , 2002 , 148, 2270-2271	2.9	1
18	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	3.5	47
17	Regulation of the eukaryotic cell cycle: molecular antagonism, hysteresis, and irreversible transitions. <i>Journal of Theoretical Biology</i> , 2001 , 210, 249-63	2.3	279
16	Network dynamics and cell physiology. <i>Nature Reviews Molecular Cell Biology</i> , 2001 , 2, 908-16	48.7	407
15	Mathematical model of the cell division cycle of fission yeast. <i>Chaos</i> , 2001 , 11, 277-286	3.3	124
14	Kinetic analysis of a molecular model of the budding yeast cell cycle. <i>Molecular Biology of the Cell</i> , 2000 , 11, 369-91	3.5	369
13	Modeling the fission yeast cell cycle: quantized cycle times in wee1- cdc25Delta mutant cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2000 , 97, 7865-70	11.5	101

12	Finishing the cell cycle. <i>Journal of Theoretical Biology</i> , 1999 , 199, 223-33	2.3	47
11	A simple model of circadian rhythms based on dimerization and proteolysis of PER and TIM. <i>Biophysical Journal</i> , 1999 , 77, 2411-7	2.9	147
10	Modeling M-phase control in <i>Xenopus</i> oocyte extracts: the surveillance mechanism for unreplicated DNA. <i>Biophysical Chemistry</i> , 1998 , 72, 169-84	3.5	51
9	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	3.5	106
8	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, 9147-52	11.5	144
7	Chemical kinetic theory: understanding cell-cycle regulation. <i>Trends in Biochemical Sciences</i> , 1996 , 21, 89-96	10.3	79
6	Quantitative analysis of a molecular model of mitotic control in fission yeast. <i>Journal of Theoretical Biology</i> , 1995 , 173, 283-305	2.3	77
5	Checkpoints in the cell cycle from a modeler's perspective. <i>Progress in Cell Cycle Research</i> , 1995 , 1, 1-8		20
4	Modeling the Cell Division Cycle: M-phase Trigger, Oscillations, and Size Control. <i>Journal of Theoretical Biology</i> , 1993 , 165, 101-134	2.3	116
3	Systems biology of the yeast cell cycle engine		3
2	Computational modelling of chromosome re-replication in mutant strains of fission yeast		1
1	Mechanisms of signalling-memory governing progression through the eukaryotic cell cycle		1