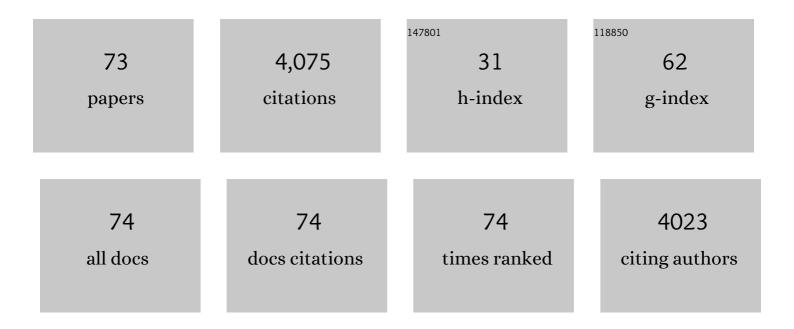
Marco Muzi-Falconi

List of Publications by Year in descending order

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MARCO MUZI-EALCONI

#	Article	IF	CITATIONS
1	The DNA replication checkpoint response stabilizes stalled replication forks. Nature, 2001, 412, 557-561.	27.8	693
2	Mutations in the mitochondrial protease gene AFG3L2 cause dominant hereditary ataxia SCA28. Nature Genetics, 2010, 42, 313-321.	21.4	291
3	The DNA Damage Checkpoint Response Requires Histone H2B Ubiquitination by Rad6-Bre1 and H3 Methylation by Dot1. Journal of Biological Chemistry, 2005, 280, 9879-9886.	3.4	249
4	Regulation of the replication initiator protein p65 ^{<i>cdc18</i>} by CDK phosphorylation. Genes and Development, 1997, 11, 2767-2779.	5.9	161
5	Histone methyltransferase Dot1 and Rad9 inhibit single-stranded DNA accumulation at DSBs and uncapped telomeres. EMBO Journal, 2008, 27, 1502-12.	7.8	159
6	RNase H and Postreplication Repair Protect Cells from Ribonucleotides Incorporated in DNA. Molecular Cell, 2012, 45, 99-110.	9.7	153
7	DNA damage checkpoint in budding yeast. EMBO Journal, 1998, 17, 5525-5528.	7.8	145
8	Ribonucleotides Misincorporated into DNA Act as Strand-Discrimination Signals in Eukaryotic Mismatch Repair. Molecular Cell, 2013, 50, 323-332.	9.7	139
9	cdc18+ regulates initiation of DNA replication in Schizosaccharomyces pombe Proceedings of the National Academy of Sciences of the United States of America, 1996, 93, 1566-1570.	7.1	124
10	Physical and functional interactions between nucleotide excision repair and DNA damage checkpoint. EMBO Journal, 2004, 23, 429-438.	7.8	112
11	DNA damage checkpoints and DNA replication controls in Saccharomyces cerevisiae. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2000, 451, 187-196.	1.0	110
12	Phosphorylation of the Budding Yeast 9-1-1 Complex Is Required for Dpb11 Function in the Full Activation of the UV-Induced DNA Damage Checkpoint. Molecular and Cellular Biology, 2008, 28, 4782-4793.	2.3	109
13	DNA nucleotide excision repair-dependent signaling to checkpoint activation. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 17325-17330.	7.1	107
14	The 9-1-1 Checkpoint Clamp Physically Interacts with Polζ and Is Partially Required for Spontaneous Polζ-dependent Mutagenesis in Saccharomyces cerevisiae. Journal of Biological Chemistry, 2005, 280, 38657-38665.	3.4	104
15	Exo1 Competes with Repair Synthesis, Converts NER Intermediates to Long ssDNA Gaps, and Promotes Checkpoint Activation. Molecular Cell, 2010, 40, 50-62.	9.7	99
16	Orp1, a member of the Cdc18/Cdc6 family of S-phase regulators, is homologous to a component of the origin recognition complex Proceedings of the National Academy of Sciences of the United States of America, 1995, 92, 12475-12479.	7.1	98
17	Checkpoint mechanisms at the intersection between DNA damage and repair. DNA Repair, 2009, 8, 1055-1067.	2.8	82
18	Reduction of hRNase H2 activity in Aicardi–GoutiÔres syndrome cells leads to replication stress and genome instability. Human Molecular Genetics, 2015, 24, 649-658.	2.9	67

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19	Human exonuclease 1 connects nucleotide excision repair (NER) processing with checkpoint activation in response to UV irradiation. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 13647-13652.	7.1	64
20	Dynamics of Rad9 Chromatin Binding and Checkpoint Function Are Mediated by Its Dimerization and Are Cell Cycle–Regulated by CDK1 Activity. PLoS Genetics, 2010, 6, e1001047.	3.5	59
21	The DNA PolymerasePrimase Complex: Multiple Functions and Interactions. Scientific World Journal, The, 2003, 3, 21-33.	2.1	56
22	Nucleotide sequence and characterization of temperature-sensitive pol1 mutants of Saccharomyces cerevisiae. Gene, 1990, 90, 99-104.	2.2	54
23	Yeast Rev1 is cell cycle regulated, phosphorylated in response to DNA damage and its binding to chromosomes is dependent upon MEC1. DNA Repair, 2007, 6, 121-127.	2.8	53
24	Elevated Levels of the Polo Kinase Cdc5 Override the Mec1/ATR Checkpoint in Budding Yeast by Acting at Different Steps of the Signaling Pathway. PLoS Genetics, 2010, 6, e1000763.	3.5	49
25	One-step high-throughput assay for quantitative detection of β-galactosidase activity in intact Gram-negative bacteria, yeast, and mammalian cells. BioTechniques, 2006, 40, 433-440.	1.8	48
26	14-3-3 Proteins Regulate Exonuclease 1–Dependent Processing of Stalled Replication Forks. PLoS Genetics, 2011, 7, e1001367.	3.5	45
27	Sensing of Replication Stress and Mec1 Activation Act through Two Independent Pathways Involving the 9-1-1 Complex and DNA Polymerase ε. PLoS Genetics, 2011, 7, e1002022.	3.5	40
28	Exploring the pH Sensitivity of Poly(allylamine) Phosphate Supramolecular Nanocarriers for Intracellular siRNA Delivery. ACS Applied Materials & Interfaces, 2017, 9, 38242-38254.	8.0	38
29	DNA replication: Controlling initiation during the cell cycle. Current Biology, 1996, 6, 229-233.	3.9	33
30	Mind the gap: Keeping UV lesions in check. DNA Repair, 2011, 10, 751-759.	2.8	33
31	NER and DDR: Classical music with new instruments. Cell Cycle, 2012, 11, 668-674.	2.6	32
32	The ribonuclease DIS3 promotes let-7 miRNA maturation by degrading the pluripotency factor LIN28B mRNA. Nucleic Acids Research, 2015, 43, 5182-5193.	14.5	31
33	Alk1 and Alk2 are Two New Cell Cycle-Regulated Haspin-Like Proteins in Budding Yeast. Cell Cycle, 2006, 5, 1464-1471.	2.6	28
34	Exploring Quantitative Yeast Phenomics with Single-Cell Analysis of DNA Damage Foci. Cell Systems, 2016, 3, 264-277.e10.	6.2	26
35	Coordinated Activity of Y Family TLS Polymerases and EXO1 Protects Non-S Phase Cells from UV-Induced Cytotoxic Lesions. Molecular Cell, 2018, 70, 34-47.e4.	9.7	26
36	EXO1: A tightly regulated nuclease. DNA Repair, 2020, 93, 102929.	2.8	23

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37	Yeast Haspin Kinase Regulates Polarity Cues Necessary for Mitotic Spindle Positioning and Is Required to Tolerate Mitotic Arrest. Developmental Cell, 2013, 26, 483-495.	7.0	22
38	Novel <i>DYT11</i> gene mutation in patients without dopaminergic deficit (SWEDD) screened for dystonia. Neurology, 2014, 83, 1155-1162.	1.1	22
39	Tau and DNA Damage in Neurodegeneration. Brain Sciences, 2020, 10, 946.	2.3	22
40	RNase H activities counteract a toxic effect of Polymerase \hat{I} in cells replicating with depleted dNTP pools. Nucleic Acids Research, 2019, 47, 4612-4623.	14.5	21
41	One, No One, and One Hundred Thousand: The Many Forms of Ribonucleotides in DNA. International Journal of Molecular Sciences, 2020, 21, 1706.	4.1	17
42	The yeast DNA polymerase-primase complex: Genes and proteins. Biochimica Et Biophysica Acta Gene Regulatory Mechanisms, 1988, 951, 268-273.	2.4	16
43	Saccharomyces CDK1 Phosphorylates Rad53 Kinase in Metaphase, Influencing Cellular Morphogenesis. Journal of Biological Chemistry, 2009, 284, 32627-32634.	3.4	16
44	Physical and functional crosstalk between Fanconi anemia core components and the GINS replication complex. DNA Repair, 2011, 10, 149-158.	2.8	15
45	DNA decay and limited Rad53 activation after liquid holding of UV-treated nucleotide excision repair deficient S. cerevisiae cells. DNA Repair, 2004, 3, 1591-1599.	2.8	14
46	A dominant-negative MEC3 mutant uncovers new functions for the Rad17 complex and Tel1. Proceedings of the National Academy of Sciences of the United States of America, 2002, 99, 12997-13002.	7.1	13
47	Mechanisms Controlling the Integrity of Replicating Chromosomes in Budding Yeast. Cell Cycle, 2003, 2, 563-566.	2.6	12
48	To trim or not to trim: Progression and control of DSB end resection. Cell Cycle, 2013, 12, 1848-1860.	2.6	12
49	Haspin regulates Ras localization to promote Cdc24-driven mitotic depolarization. Cell Discovery, 2020, 6, 42.	6.7	12
50	The Incorporation of Ribonucleotides Induces Structural and Conformational Changes in DNA. Biophysical Journal, 2017, 113, 1373-1382.	0.5	11
51	Overproduction and functional analysis of DNA primase subunits from yeast and mouse. Gene, 1992, 113, 199-205.	2.2	9
52	Non-Canonical CRL4A/4BCDT2 Interacts with RAD18 to Modulate Post Replication Repair and Cell Survival. PLoS ONE, 2013, 8, e60000.	2.5	8
53	Generation of three iPSC lines from fibroblasts of a patient with Aicardi Goutières Syndrome mutated in TREX1. Stem Cell Research, 2019, 41, 101580.	0.7	8
54	Roles and regulation of Haspin kinase and its impact on carcinogenesis. Cellular Signalling, 2022, 93, 110303.	3.6	8

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55	In vivo and in silico analysis of PCNA ubiquitylation in the activation of the Post Replication Repair pathway in S. cerevisiae. BMC Systems Biology, 2013, 7, 24.	3.0	7
56	A blooming resolvase at chromosomal fragile sites. Nature Cell Biology, 2013, 15, 883-885.	10.3	7
57	Phosphorylation of H3-Thr3 by Haspin Is Required for Primary Cilia Regulation. International Journal of Molecular Sciences, 2021, 22, 7753.	4.1	7
58	Mechanisms controlling the integrity of replicating chromosomes in budding yeast. Cell Cycle, 2003, 2, 564-7.	2.6	7
59	Establishment of three iPSC lines from fibroblasts of a patient with Aicardi Goutières syndrome mutated in RNaseH2B. Stem Cell Research, 2019, 41, 101620.	0.7	6
60	Correlation between Checkpoint Activation and in Vivo Assembly of the Yeast Checkpoint Complex Rad17-Mec3-Ddc1. Journal of Biological Chemistry, 2003, 278, 22303-22308.	3.4	5
61	Haspin Modulates the G2/M Transition Delay in Response to Polarization Failures in Budding Yeast. Frontiers in Cell and Developmental Biology, 2020, 8, 625717.	3.7	5
62	Measuring the Levels of Ribonucleotides Embedded in Genomic DNA. Methods in Molecular Biology, 2018, 1672, 319-327.	0.9	5
63	<i>VID22</i> counteracts G-quadruplex-induced genome instability. Nucleic Acids Research, 2021, 49, 12785-12804.	14.5	5
64	Study of UV-induced DNA Repair Factor Recruitment: Kinetics and Dynamics. Methods in Molecular Biology, 2018, 1672, 101-105.	0.9	4
65	Generation of three isogenic induced Pluripotent Stem Cell lines (iPSCs) from fibroblasts of a patient with Aicardi GoutiA¨res Syndrome carrying a c.2471G>A dominant mutation in IFIH1 gene. Stem Cell Research, 2019, 41, 101623.	0.7	4
66	gRASping Depolarization: Contribution of RAS GTPases to Mitotic Polarity Clusters Resolution. Frontiers in Cell and Developmental Biology, 2020, 8, 589993.	3.7	4
67	Checkpoint response to DNA damage. DNA Repair, 2009, 8, 973-973.	2.8	3
68	Characterization of Structural and Configurational Properties of DNA by Atomic Force Microscopy. Methods in Molecular Biology, 2018, 1672, 557-573.	0.9	1
69	Co-immunoprecipitation of human mitochondrial proteases AFG3L2 and paraplegin heterologously expressed in yeast cells. Protocol Exchange, 0, , .	0.3	1
70	Budding Yeast DNA Damage Checkpoint: A Signal Transduction-Mediated Surveillance System. , 2003, , 197-202.		1
71	Sometimes size does matter. European Journal of Cancer, 2003, 39, 1337-1338.	2.8	0
72	Replication DNA Polymerase α – Primase Complex: A Fundamental Element in the Maintenance of Genome Integrity. , 2021, , 71-79.		0

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CITATIONS

73 Cell Cycle Checkpoints. , 2013, , 254-259.