

Marcel A T M Van Vugt

List of Publications by Year in descending order

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Version: 2024-02-01

78
papers

5,397
citations

101496

36
h-index

88593

70
g-index

82
all docs

82
docs citations

82
times ranked

10765
citing authors

#	ARTICLE	IF	CITATIONS
1	Systematic Discovery of In Vivo Phosphorylation Networks. <i>Cell</i> , 2007, 129, 1415-1426.	13.5	702
2	Polo-like Kinase-1 Controls Recovery from a G2 DNA Damage-Induced Arrest in Mammalian Cells. <i>Molecular Cell</i> , 2004, 15, 799-811.	4.5	336
3	Gene expression analysis identifies global gene dosage sensitivity in cancer. <i>Nature Genetics</i> , 2015, 47, 115-125.	9.4	313
4	Relevance of Tumor-Infiltrating Immune Cell Composition and Functionality for Disease Outcome in Breast Cancer. <i>Journal of the National Cancer Institute</i> , 2017, 109, djw192.	3.0	296
5	Getting in and out of mitosis with Polo-like kinase-1. <i>Oncogene</i> , 2005, 24, 2844-2859.	2.6	258
6	Selective Loss of PARC Restores PARylation and Counteracts PARP Inhibitor-Mediated Synthetic Lethality. <i>Cancer Cell</i> , 2018, 33, 1078-1093.e12.	7.7	238
7	14-3-3 β controls mitotic translation to facilitate cytokinesis. <i>Nature</i> , 2007, 446, 329-332.	13.7	217
8	DNA Damage Activates a Spatially Distinct Late Cytoplasmic Cell-Cycle Checkpoint Network Controlled by MK2-Mediated RNA Stabilization. <i>Molecular Cell</i> , 2010, 40, 34-49.	4.5	210
9	A Mitotic Phosphorylation Feedback Network Connects Cdk1, Plk1, 53BP1, and Chk2 to Inactivate the G2/M DNA Damage Checkpoint. <i>PLoS Biology</i> , 2010, 8, e1000287.	2.6	201
10	Polo-like Kinase-1 Is Required for Bipolar Spindle Formation but Is Dispensable for Anaphase Promoting Complex/Cdc20 Activation and Initiation of Cytokinesis. <i>Journal of Biological Chemistry</i> , 2004, 279, 36841-36854.	1.6	173
11	cGAS \rightarrow STING drives the IL-6-dependent survival of chromosomally instable cancers. <i>Nature</i> , 2022, 607, 366-373.	13.7	132
12	Loss of Rb proteins causes genomic instability in the absence of mitogenic signaling. <i>Genes and Development</i> , 2010, 24, 1377-1388.	2.7	107
13	RIF1 promotes replication fork protection and efficient restart to maintain genome stability. <i>Nature Communications</i> , 2019, 10, 3287.	5.8	91
14	BRCA2 deficiency instigates cGAS-mediated inflammatory signaling and confers sensitivity to tumor necrosis factor-alpha-mediated cytotoxicity. <i>Nature Communications</i> , 2019, 10, 100.	5.8	91
15	The DNA damage response during mitosis. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2013, 750, 45-55.	0.4	85
16	Progression through mitosis promotes PARP inhibitor-induced cytotoxicity in homologous recombination-deficient cancer cells. <i>Nature Communications</i> , 2017, 8, 15981.	5.8	83
17	Functional Dynamics of Polo-Like Kinase 1 at the Centrosome. <i>Molecular and Cellular Biology</i> , 2009, 29, 3134-3150.	1.1	82
18	Testicular cancer: Determinants of cisplatin sensitivity and novel therapeutic opportunities. <i>Cancer Treatment Reviews</i> , 2020, 88, 102054.	3.4	76

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19	Rif1 Is Required for Resolution of Ultrafine DNA Bridges in Anaphase to Ensure Genomic Stability. <i>Developmental Cell</i> , 2015, 34, 466-474.	3.1	74
20	CLIP-170 facilitates the formation of kinetochore–microtubule attachments. <i>EMBO Journal</i> , 2006, 25, 45-57.	3.5	72
21	Chk2 controls CtIP stability during the cell cycle and in response to DNA damage. <i>EMBO Journal</i> , 2014, 33, 2860-2879.	3.5	65
22	PLK1 (polo like kinase 1) inhibits MTOR complex 1 and promotes autophagy. <i>Autophagy</i> , 2017, 13, 486-505.	4.3	63
23	Uncoupling Anaphase-Promoting Complex/Cyclosome Activity from Spindle Assembly Checkpoint Control by Deregulating Polo-Like Kinase 1. <i>Molecular and Cellular Biology</i> , 2005, 25, 2031-2044.	1.1	62
24	Restarting the Cell Cycle When the Checkpoint Comes to a Halt: Figure 1.. <i>Cancer Research</i> , 2005, 65, 7037-7040.	0.4	62
25	Premature mitotic entry induced by ATR inhibition potentiates olaparib inhibition–mediated genomic instability, inflammatory signaling, and cytotoxicity in BRCA2–deficient cancer cells. <i>Molecular Oncology</i> , 2019, 13, 2422-2440.	2.1	62
26	Regulators of homologous recombination repair as novel targets for cancer treatment. <i>Frontiers in Genetics</i> , 2015, 6, 96.	1.1	58
27	Checkpoint Adaptation and Recovery: Back with Polo after the Break. <i>Cell Cycle</i> , 2004, 3, 1383-1386.	1.3	56
28	Genomic instability, inflammatory signaling and response to cancer immunotherapy. <i>Biochimica Et Biophysica Acta: Reviews on Cancer</i> , 2022, 1877, 188661.	3.3	56
29	Breaking the DNA damage response to improve cervical cancer treatment. <i>Cancer Treatment Reviews</i> , 2016, 42, 30-40.	3.4	54
30	Transcriptional effects of copy number alterations in a large set of human cancers. <i>Nature Communications</i> , 2020, 11, 715.	5.8	53
31	Studying platinum sensitivity and resistance in high-grade serous ovarian cancer: Different models for different questions. <i>Drug Resistance Updates</i> , 2016, 24, 55-69.	6.5	52
32	A haploid genetic screen identifies the G ₁ /S regulatory machinery as a determinant of Wee1 inhibitor sensitivity. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 15160-15165.	3.3	50
33	TPX2/Aurora kinase A signaling as a potential therapeutic target in genomically unstable cancer cells. <i>Oncogene</i> , 2019, 38, 852-867.	2.6	43
34	Shaping the BRCAness mutational landscape by alternative double-strand break repair, replication stress and mitotic aberrancies. <i>Nucleic Acids Research</i> , 2021, 49, 4239-4257.	6.5	42
35	Elongation factor ELOF1 drives transcription-coupled repair and prevents genome instability. <i>Nature Cell Biology</i> , 2021, 23, 608-619.	4.6	41
36	Claspin: Timing the Cell Cycle Arrest When the Genome is Damaged. <i>Cell Cycle</i> , 2006, 5, 2831-2834.	1.3	39

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37	Functional validation of putative tumor suppressor gene <i>C13ORF18</i> in cervical cancer by Artificial Transcription Factors. <i>Molecular Oncology</i> , 2013, 7, 669-679.	2.1	39
38	miR-371a-3p, miR-373-3p and miR-367-3p as Serum Biomarkers in Metastatic Testicular Germ Cell Cancers Before, During and After Chemotherapy. <i>Cells</i> , 2019, 8, 1221.	1.8	39
39	Overexpression of Cyclin E1 or Cdc25A leads to replication stress, mitotic aberrancies, and increased sensitivity to replication checkpoint inhibitors. <i>Oncogenesis</i> , 2020, 9, 88.	2.1	37
40	The role of ATM and 53BP1 as predictive markers in cervical cancer. <i>International Journal of Cancer</i> , 2012, 131, 2056-2066.	2.3	35
41	CXCR4 inhibition enhances radiosensitivity, while inducing cancer cell mobilization in a prostate cancer mouse model. <i>Clinical and Experimental Metastasis</i> , 2014, 31, 829-839.	1.7	35
42	Therapeutic targeting and patient selection for cancers with homologous recombination defects. <i>Expert Opinion on Drug Discovery</i> , 2017, 12, 565-581.	2.5	32
43	Polo-Like Kinase-1 Controls Aurora A Destruction by Activating APC/C-Cdh1. <i>PLoS ONE</i> , 2009, 4, e5282.	1.1	30
44	Checkpoint kinase 2 (Chk2) supports sensitivity to platinum-based treatment in high grade serous ovarian cancer. <i>Gynecologic Oncology</i> , 2014, 133, 591-598.	0.6	28
45	Cyclin E expression is associated with high levels of replication stress in triple-negative breast cancer. <i>Npj Breast Cancer</i> , 2020, 6, 40.	2.3	27
46	When breaks get hot: inflammatory signaling in BRCA1/2-mutant cancers. <i>Trends in Cancer</i> , 2022, 8, 174-189.	3.8	27
47	Modeling of Cisplatin-Induced Signaling Dynamics in Triple-Negative Breast Cancer Cells Reveals Mediators of Sensitivity. <i>Cell Reports</i> , 2019, 28, 2345-2357.e5.	2.9	25
48	Cell cycle re-entry mechanisms after DNA damage checkpoints: Giving it some gas to shut off the breaks!. <i>Cell Cycle</i> , 2010, 9, 2097-2101.	1.3	21
49	Toward Molecular Imaging-Driven Drug Development in Oncology. <i>Cancer Discovery</i> , 2011, 1, 25-28.	7.7	21
50	Inflammatory signaling in genomically instable cancers. <i>Cell Cycle</i> , 2019, 18, 1830-1848.	1.3	21
51	Improving gene function predictions using independent transcriptional components. <i>Nature Communications</i> , 2021, 12, 1464.	5.8	20
52	Identification of relevant drugable targets in diffuse large B-cell lymphoma using a genome-wide unbiased CD20 guilt-by association approach. <i>PLoS ONE</i> , 2018, 13, e0193098.	1.1	20
53	Controlling the response to DNA damage by the APC/C-Cdh1. <i>Cellular and Molecular Life Sciences</i> , 2016, 73, 949-960.	2.4	19
54	Identification of Two Protein-Signaling States Delineating Transcriptionally Heterogeneous Human Medulloblastoma. <i>Cell Reports</i> , 2018, 22, 3206-3216.	2.9	19

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55	An mRNA expression-based signature for oncogene-induced replication-stress. <i>Oncogene</i> , 2022, 41, 1216-1224.	2.6	17
56	Replication stress: Driver and therapeutic target in genomically unstable cancers. <i>Advances in Protein Chemistry and Structural Biology</i> , 2019, 115, 157-201.	1.0	15
57	CXCR4 and CXCL12 Expression in Rectal Tumors of Stage IV Patients Before and After Local Radiotherapy and Systemic Neoadjuvant Treatment. <i>Current Pharmaceutical Design</i> , 2015, 21, 2276-2283.	0.9	15
58	Comparison of Carboplatin With 5-Fluorouracil vs. Cisplatin as Concomitant Chemoradiotherapy for Locally Advanced Head and Neck Squamous Cell Carcinoma. <i>Frontiers in Oncology</i> , 2020, 10, 761.	1.3	14
59	The H3.3K27M oncohistone affects replication stress outcome and provokes genomic instability in pediatric glioma. <i>PLoS Genetics</i> , 2021, 17, e1009868.	1.5	14
60	Nuclear COMMD1 Is Associated with Cisplatin Sensitivity in Ovarian Cancer. <i>PLoS ONE</i> , 2016, 11, e0165385.	1.1	13
61	Harnessing Integrative Omics to Facilitate Molecular Imaging of the Human Epidermal Growth Factor Receptor Family for Precision Medicine. <i>Theranostics</i> , 2017, 7, 2111-2133.	4.6	12
62	Perspectives for tailored chemoprevention and treatment of colorectal cancer in Lynch syndrome. <i>Critical Reviews in Oncology/Hematology</i> , 2011, 80, 264-277.	2.0	11
63	Quantitative proteomics analysis identifies MUC1 as an effect sensor of EGFR inhibition. <i>Oncogene</i> , 2019, 38, 1477-1488.	2.6	11
64	Dual mTORC1/2 Inhibition Sensitizes Testicular Cancer Models to Cisplatin Treatment. <i>Molecular Cancer Therapeutics</i> , 2020, 19, 590-601.	1.9	8
65	Shutting down the power supply for DNA repair in cancer cells. <i>Journal of Cell Biology</i> , 2017, 216, 295-297.	2.3	7
66	Polo-Like Kinase-1: Activity Measurement and RNAi-Mediated Knockdown. , 2005, 296, 355-370.		6
67	Never tear us a-PARP: Dealing with DNA lesions during mitosis. <i>Molecular and Cellular Oncology</i> , 2018, 5, e1382670.	0.3	6
68	Deposition Bias of Chromatin Proteins Inverts under DNA Replication Stress Conditions. <i>ACS Chemical Biology</i> , 2021, 16, 2193-2201.	1.6	6
69	Data-Driven prioritisation of antibody-drug conjugate targets in head and neck squamous cell carcinoma. <i>Oral Oncology</i> , 2018, 80, 33-39.	0.8	5
70	Establishment and characterisation of testicular cancer patient-derived xenograft models for preclinical evaluation of novel therapeutic strategies. <i>Scientific Reports</i> , 2020, 10, 18938.	1.6	4
71	The RECQL helicase prevents replication fork collapse during replication stress. <i>Life Science Alliance</i> , 2020, 3, e202000668.	1.3	4
72	Identification and Validation of Esophageal Squamous Cell Carcinoma Targets for Fluorescence Molecular Endoscopy. <i>International Journal of Molecular Sciences</i> , 2021, 22, 9270.	1.8	3

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73	Building a great wall around mitosis: Evolutionary conserved roles for the Greatwall/MASTL kinases in securing chromosome stability. <i>Cell Cycle</i> , 2010, 9, 3842-3847.	1.3	1
74	Editorial: Cancer-Associated Defects in the DNA Damage Response: Drivers for Malignant Transformation and Potential Therapeutic Targets. <i>Frontiers in Genetics</i> , 2015, 6, 355.	1.1	1
75	Plk1 Manages DNA break repair during mitosis. <i>Cell Cycle</i> , 2015, 14, 1356-1357.	1.3	1
76	A synthetic lethal screen identifies HDAC4 as a potential target in MELK overexpressing cancers. <i>G3: Genes, Genomes, Genetics</i> , 2021, 11, .	0.8	1
77	MBRS-36. IDENTIFICATION OF TWO PROTEIN-SIGNALING STATES DELINEATING TRANSCRIPTIONALLY HETEROGENEOUS HUMAN MEDULLOBLASTOMA. <i>Neuro-Oncology</i> , 2018, 20, i136-i136.	0.6	0
78	A large pooled analysis refines gene expression-based molecular subclasses in cutaneous melanoma. <i>Oncolmmunology</i> , 2019, 8, 1558664.	2.1	0