

Mark W Richards

List of Publications by Year in descending order

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Version: 2024-02-01

33
papers

1,689
citations

270111

25
h-index

466096

32
g-index

37
all docs

37
docs citations

37
times ranked

2867
citing authors

#	ARTICLE	IF	CITATIONS
1	Combined inhibition of Aurora-A and ATR kinases results in regression of MYCN-amplified neuroblastoma. <i>Nature Cancer</i> , 2021, 2, 312-326.	5.7	50
2	Phase-separated foci of EML4-ALK facilitate signalling and depend upon an active kinase conformation. <i>EMBO Reports</i> , 2021, 22, e53693.	2.0	31
3	Discovery and Optimization of wt-RET/KDR-Selective Inhibitors of RET ^{V804M} Kinase. <i>ACS Medicinal Chemistry Letters</i> , 2020, 11, 497-505.	1.3	8
4	EML4-ALK V3 oncogenic fusion proteins promote microtubule stabilization and accelerated migration through NEK9 and NEK7. <i>Journal of Cell Science</i> , 2020, 133, .	1.2	30
5	2-Arylamino-6-ethynylpurines are cysteine-targeting irreversible inhibitors of Nek2 kinase. <i>RSC Medicinal Chemistry</i> , 2020, 11, 707-731.	1.7	8
6	Orally bioavailable CDK9/2 inhibitor shows mechanism-based therapeutic potential in MYCN-driven neuroblastoma. <i>Journal of Clinical Investigation</i> , 2020, 130, 5875-5892.	3.9	40
7	Mitotic phosphorylation by NEK6 and NEK7 reduces the microtubule affinity of EML4 to promote chromosome congression. <i>Science Signaling</i> , 2019, 12, .	1.6	30
8	Mitotic spindle association of TACC3 requires AuroraA-dependent stabilization of a cryptic α -helix. <i>EMBO Journal</i> , 2018, 37, .	3.5	55
9	Mitotic phosphorylation regulates Hsp72 spindle localization by uncoupling ATP binding from substrate release. <i>Science Signaling</i> , 2018, 11, .	1.6	8
10	A moving target: structure and disorder in pursuit of Myc inhibitors. <i>Biochemical Society Transactions</i> , 2017, 45, 709-717.	1.6	26
11	Differential protein stability and clinical responses of EML4-ALK fusion variants to various ALK inhibitors in advanced ALK-rearranged non-small cell lung cancer. <i>Annals of Oncology</i> , 2017, 28, 791-797.	0.6	178
12	Characterization of Three Druggable Hot-Spots in the Aurora-A/TPX2 Interaction Using Biochemical, Biophysical, and Fragment-Based Approaches. <i>ACS Chemical Biology</i> , 2017, 12, 2906-2914.	1.6	40
13	Eml1 loss impairs apical progenitor spindle length and soma shape in the developing cerebral cortex. <i>Scientific Reports</i> , 2017, 7, 17308.	1.6	26
14	A closed conformation of the <i>Caenorhabditis elegans</i> separase-securin complex. <i>Open Biology</i> , 2016, 6, 160032.	1.5	10
15	Allosteric inhibition of Aurora-A kinase by a synthetic vNAR domain. <i>Open Biology</i> , 2016, 6, 160089.	1.5	39
16	Structural basis of N-Myc binding by Aurora-A and its destabilization by kinase inhibitors. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 13726-13731.	3.3	130
17	Molecular mechanisms that underpin EML4-ALK driven cancers and their response to targeted drugs. <i>Cellular and Molecular Life Sciences</i> , 2016, 73, 1209-1224.	2.4	80
18	Microtubule association of EML proteins and the EML4-ALK variant 3 oncoprotein require an N-terminal trimerization domain. <i>Biochemical Journal</i> , 2015, 467, 529-536.	1.7	73

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19	Hsp72 is targeted to the mitotic spindle by Nek6 to promote K-fiber assembly and mitotic progression. <i>Journal of Cell Biology</i> , 2015, 209, 349-358.	2.3	44
20	Mechanistic basis of Nek7 activation through Nek9 binding and induced dimerization. <i>Nature Communications</i> , 2015, 6, 8771.	5.8	43
21	Molecular mechanisms of human IRE1 activation through dimerization and ligand binding. <i>Oncotarget</i> , 2015, 6, 13019-13035.	0.8	49
22	Crystal structure of EML1 reveals the basis for Hsp90 dependence of oncogenic EML4-ALK by disruption of an atypical $\hat{1}^2$ -propeller domain. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 5195-5200.	3.3	93
23	Coordination of adjacent domains mediates TACC3- $\hat{1}$ -TOG- $\hat{1}$ -clathrin assembly and mitotic spindle binding. <i>Journal of Cell Biology</i> , 2013, 202, 463-478.	2.3	76
24	Structural Basis of Poly(ADP-ribose) Recognition by the Multizinc Binding Domain of Checkpoint with Forkhead-associated and RING Domains (CHFR). <i>Journal of Biological Chemistry</i> , 2010, 285, 39348-39358.	1.6	54
25	Crystal structure of an Aurora-A mutant that mimics Aurora-B bound to MLN8054: insights into selectivity and drug design. <i>Biochemical Journal</i> , 2010, 427, 19-28.	1.7	86
26	A Pocket on the Surface of the N-Terminal BRCT Domain of Mcph1 Is Required to Prevent Abnormal Chromosome Condensation. <i>Journal of Molecular Biology</i> , 2010, 395, 908-915.	2.0	12
27	Insights into the Conformational Variability and Regulation of Human Nek2 Kinase. <i>Journal of Molecular Biology</i> , 2009, 386, 476-485.	2.0	47
28	An Autoinhibitory Tyrosine Motif in the Cell-Cycle-Regulated Nek7 Kinase Is Released through Binding of Nek9. <i>Molecular Cell</i> , 2009, 36, 560-570.	4.5	83
29	The HOOK-Domain Between the SH3- and the GK-Domains of Ca _v 1.2 Subunits Contains Key Determinants Controlling Calcium Channel Inactivation. <i>Channels</i> , 2007, 1, 92-101.	1.5	32
30	The importance of occupancy rather than affinity of Ca _v 1.2 subunits for the calcium channel I-II linker in relation to calcium channel function. <i>Journal of Physiology</i> , 2006, 574, 387-398.	1.3	26
31	Interaction via a Key Tryptophan in the I-II Linker of N-Type Calcium Channels Is Required for $\hat{1}$ But Not for Palmitoylated $\hat{2}$, Implicating an Additional Binding Site in the Regulation of Channel Voltage-Dependent Properties. <i>Journal of Neuroscience</i> , 2005, 25, 6984-6996.	1.7	75
32	Ca ²⁺ channel $\hat{1}^2$ -subunits: structural insights AID our understanding. <i>Trends in Pharmacological Sciences</i> , 2004, 25, 626-632.	4.0	100
33	EML4-ALK V3 Drives Cell Migration Through NEK9 and NEK7 Kinases in Non-Small-Cell Lung Cancer. <i>SSRN Electronic Journal</i> , 0, , .	0.4	0