

# Mark W Richards

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7711570/publications.pdf>

Version: 2024-02-01

33  
papers

1,689  
citations

236925

25  
h-index

414414

32  
g-index

37  
all docs

37  
docs citations

37  
times ranked

2643  
citing authors

#	ARTICLE	IF	CITATIONS
1	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.	1.2	178
2	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.	7.1	130
3	Ca <sup>2+</sup> channel $\beta$ -subunits: structural insights aid our understanding. <i>Trends in Pharmacological Sciences</i> , 2004, 25, 626-632.	8.7	100
4	Crystal structure of EML1 reveals the basis for Hsp90 dependence of oncogenic EML4-ALK by disruption of an atypical $\beta$ -propeller domain. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 5195-5200.	7.1	93
5	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.	3.7	86
6	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.	9.7	83
7	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.	5.4	80
8	Coordination of adjacent domains mediates TACC3-ch-TOG-clathrin assembly and mitotic spindle binding. <i>Journal of Cell Biology</i> , 2013, 202, 463-478.	5.2	76
9	Interaction via a Key Tryptophan in the I-II Linker of N-Type Calcium Channels Is Required for $\Delta$ 1 But Not for Palmitoylated $\Delta$ 2, Implicating an Additional Binding Site in the Regulation of Channel Voltage-Dependent Properties. <i>Journal of Neuroscience</i> , 2005, 25, 6984-6996.	3.6	75
10	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.	3.7	73
11	Mitotic spindle association of TACC3 requires AuroraA-dependent stabilization of a cryptic $\beta$ -helix. <i>EMBO Journal</i> , 2018, 37, .	7.8	55
12	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.	3.4	54
13	Combined inhibition of Aurora-A and ATR kinases results in regression of MYCN-amplified neuroblastoma. <i>Nature Cancer</i> , 2021, 2, 312-326.	13.2	50
14	Molecular mechanisms of human IRE1 activation through dimerization and ligand binding. <i>Oncotarget</i> , 2015, 6, 13019-13035.	1.8	49
15	Insights into the Conformational Variability and Regulation of Human Nek2 Kinase. <i>Journal of Molecular Biology</i> , 2009, 386, 476-485.	4.2	47
16	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.	5.2	44
17	Mechanistic basis of Nek7 activation through Nek9 binding and induced dimerization. <i>Nature Communications</i> , 2015, 6, 8771.	12.8	43
18	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.	3.4	40

#	ARTICLE	IF	CITATIONS
19	Orally bioavailable CDK9/2 inhibitor shows mechanism-based therapeutic potential in MYCN-driven neuroblastoma. <i>Journal of Clinical Investigation</i> , 2020, 130, 5875-5892.	8.2	40
20	Allosteric inhibition of Aurora-A kinase by a synthetic vNAR domain. <i>Open Biology</i> , 2016, 6, 160089.	3.6	39
21	The HOOK-Domain Between the SH3- and the GK-Domains of Ca <sub>v</sub> 1.2 Subunits Contains Key Determinants Controlling Calcium Channel Inactivation. <i>Channels</i> , 2007, 1, 92-101.	2.8	32
22	Phase-separated foci of EML4-ALK facilitate signalling and depend upon an active kinase conformation. <i>EMBO Reports</i> , 2021, 22, e53693.	4.5	31
23	Mitotic phosphorylation by NEK6 and NEK7 reduces the microtubule affinity of EML4 to promote chromosome congression. <i>Science Signaling</i> , 2019, 12, .	3.6	30
24	EML4-ALK V3 oncogenic fusion proteins promote microtubule stabilization and accelerated migration through NEK9 and NEK7. <i>Journal of Cell Science</i> , 2020, 133, .	2.0	30
25	The importance of occupancy rather than affinity of Ca <sub>v</sub> 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.	2.9	26
26	A moving target: structure and disorder in pursuit of Myc inhibitors. <i>Biochemical Society Transactions</i> , 2017, 45, 709-717.	3.4	26
27	Eml1 loss impairs apical progenitor spindle length and soma shape in the developing cerebral cortex. <i>Scientific Reports</i> , 2017, 7, 17308.	3.3	26
28	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.	4.2	12
29	A closed conformation of the <i>Caenorhabditis elegans</i> separase-securin complex. <i>Open Biology</i> , 2016, 6, 160032.	3.6	10
30	Mitotic phosphorylation regulates Hsp72 spindle localization by uncoupling ATP binding from substrate release. <i>Science Signaling</i> , 2018, 11, .	3.6	8
31	Discovery and Optimization of wt-RET/KDR-Selective Inhibitors of RET <sup>V804M</sup> Kinase. <i>ACS Medicinal Chemistry Letters</i> , 2020, 11, 497-505.	2.8	8
32	2-Arylamino-6-ethynylpurines are cysteine-targeting irreversible inhibitors of Nek2 kinase. <i>RSC Medicinal Chemistry</i> , 2020, 11, 707-731.	3.9	8
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