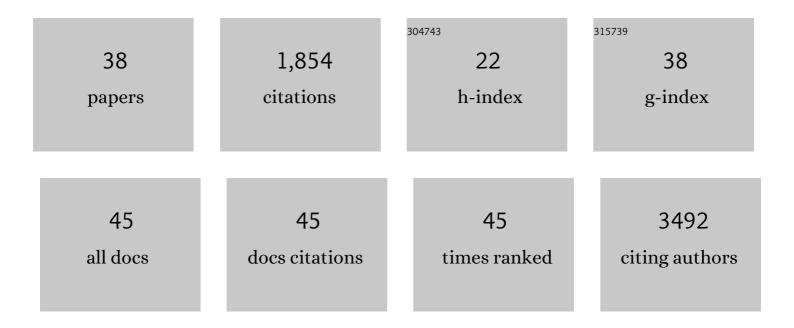
Dennis Kappei

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	The H. pylori CagA Oncoprotein Induces DNA Double Strand Breaks through Fanconi Anemia Pathway Downregulation and Replication Fork Collapse. International Journal of Molecular Sciences, 2022, 23, 1661.	4.1	6
2	PLK1 inhibition selectively induces apoptosis in ARID1A deficient cells through uncoupling of oxygen consumption from ATP production. Oncogene, 2022, 41, 1986-2002.	5.9	5
3	Multilayered control of splicing regulatory networks by DAP3 leads to widespread alternative splicing changes in cancer. Nature Communications, 2022, 13, 1793.	12.8	9
4	A degradative to secretory autophagy switch mediates mitochondria clearance in the absence of the mATG8-conjugation machinery. Nature Communications, 2022, 13, .	12.8	40
5	RNA editing mediates the functional switch of COPA in a novel mechanism of hepatocarcinogenesis. Journal of Hepatology, 2021, 74, 135-147.	3.7	41
6	Novel carfilzomib-based combinations as potential therapeutic strategies for liposarcomas. Cellular and Molecular Life Sciences, 2021, 78, 1837-1851.	5.4	6
7	Long-read transcriptome sequencing reveals abundant promoter diversity in distinct molecular subtypes of gastric cancer. Genome Biology, 2021, 22, 44.	8.8	46
8	Covalent conjugation of extracellular vesicles with peptides and nanobodies for targeted therapeutic delivery. Journal of Extracellular Vesicles, 2021, 10, e12057.	12.2	103
9	The double-stranded DNA-binding proteins TEBP-1 and TEBP-2 form a telomeric complex with POT-1. Nature Communications, 2021, 12, 2668.	12.8	12
10	Systematic Analysis of Intronic miRNAs Reveals Cooperativity within the Multicomponent <i>FTX</i> Locus to Promote Colon Cancer Development. Cancer Research, 2021, 81, 1308-1320.	0.9	14
11	HIV-1 Packaging Visualised by In-Gel SHAPE. Viruses, 2021, 13, 2389.	3.3	4
12	A common MET polymorphism harnesses HER2 signaling to drive aggressive squamous cell carcinoma. Nature Communications, 2020, 11, 1556.	12.8	12
13	Characterization of the mechanism by which the RB/E2F pathway controls expression of the cancer genomic DNA deaminase APOBEC3B. ELife, 2020, 9, .	6.0	25
14	The role of GRHL2 and epigenetic remodeling in epithelial–mesenchymal plasticity in ovarian cancer cells. Communications Biology, 2019, 2, 272.	4.4	58
15	MELK mediates the stability of EZH2 through site-specific phosphorylation in extranodal natural killer/T-cell lymphoma. Blood, 2019, 134, 2046-2058.	1.4	25
16	c-Met activation leads to the establishment of a TGFβ-receptor regulatory network in bladder cancer progression. Nature Communications, 2019, 10, 4349.	12.8	44
17	Programmed DNA elimination of germline development genes in songbirds. Nature Communications, 2019, 10, 5468.	12.8	66
18	ZBTB10 binds the telomeric variant repeat TTGGGG and interacts with TRF2. Nucleic Acids Research, 2019, 47, 1896-1907.	14.5	28

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19	Targetable BET proteins- and E2F1-dependent transcriptional program maintains the malignancy of glioblastoma. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E5086-E5095.	7.1	87
20	RUNX Poly(ADP-Ribosyl)ation and BLM Interaction Facilitate the Fanconi Anemia Pathway of DNA Repair. Cell Reports, 2018, 24, 1747-1755.	6.4	27
21	Overexpressed Melk Promotes the Stability of EZH2 through Phosphorylation in Natural Killer/T Cell Lymphoma (NKTL). Blood, 2018, 132, 2858-2858.	1.4	0
22	Phylointeractomics reconstructs functional evolution of protein binding. Nature Communications, 2017, 8, 14334.	12.8	26
23	<scp>ZBTB</scp> 48 is both a vertebrate telomereâ€binding protein and a transcriptional activator. EMBO Reports, 2017, 18, 929-946.	4.5	50
24	Epigenomic Promoter Alterations Amplify Gene Isoform and Immunogenic Diversity in Gastric Adenocarcinoma. Cancer Discovery, 2017, 7, 630-651.	9.4	48
25	P1.02-041 Characterization of MET-N375S as an Activating Mutation in Squamous Cell Carcinoma of the Lung. Journal of Thoracic Oncology, 2017, 12, S512.	1.1	2
26	The developmental proteome of <i>Drosophila melanogaster</i> . Genome Research, 2017, 27, 1273-1285.	5.5	135
27	TIP60 represses telomerase expression by inhibiting Sp1 binding to the TERT promoter. PLoS Pathogens, 2017, 13, e1006681.	4.7	24
28	Ageing and the telomere connection: An intimate relationship with inflammation. Ageing Research Reviews, 2016, 25, 55-69.	10.9	280
29	Identification of TTAGGG-binding proteins in Neurospora crassa, a fungus with vertebrate-like telomere repeats. BMC Genomics, 2015, 16, 965.	2.8	16
30	The Coilin Interactome Identifies Hundreds of Small Noncoding RNAs that Traffic through Cajal Bodies. Molecular Cell, 2014, 56, 389-399.	9.7	88
31	Quantitative interaction screen of telomeric repeat-containing RNA reveals novel TERRA regulators. Genome Research, 2013, 23, 2149-2157.	5.5	69
32	HOT1 is a mammalian direct telomere repeat-binding protein contributing to telomerase recruitment. EMBO Journal, 2013, 32, 1681-1701.	7.8	74
33	Combined RNAi and localization for functionally dissecting long noncoding RNAs. Nature Methods, 2012, 9, 360-362.	19.0	62
34	A domesticated transposon mediates the effects of a singleâ€nucleotide polymorphism responsible for enhanced muscle growth. EMBO Reports, 2010, 11, 305-311.	4.5	53
35	Monitoring Glucagon and Glucagon Antagonist-Mediated Internalization: A Useful Approach to Study Glucagon Receptor Pharmacology. Advances in Experimental Medicine and Biology, 2009, 611, 325-326.	1.6	2
36	Telomere length inheritance and aging. Mechanisms of Ageing and Development, 2008, 129, 17-26.	4.6	56

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37	Microsatellite Length Differences Between Humans and Chimpanzees at Autosomal Loci Are Not Found at Equivalent Haploid Y Chromosomal Loci. Genetics, 2006, 173, 2179-2186.	2.9	12
38	Four Different Subunits Are Essential for Expressing the Synaptic Glutamate Receptor at Neuromuscular Junctions of Drosophila. Journal of Neuroscience, 2005, 25, 3209-3218.	3.6	193