

Roland Rad

List of Publications by Year in descending order

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Version: 2024-02-01

94
papers

7,938
citations

76326

40
h-index

54911

84
g-index

99
all docs

99
docs citations

99
times ranked

14480
citing authors

#	ARTICLE	IF	CITATIONS
1	MondoA drives malignancy in B-ALL through enhanced adaptation to metabolic stress. <i>Blood</i> , 2022, 139, 1184-1197.	1.4	7
2	Disease Modeling on Tumor Organoids Implicates AURKA as a Therapeutic Target in Liver Metastatic Colorectal Cancer. <i>Cellular and Molecular Gastroenterology and Hepatology</i> , 2022, 13, 517-540.	4.5	11
3	A novel Cereblon E3 ligase modulator with antitumor activity in gastrointestinal cancer. <i>Bioorganic Chemistry</i> , 2022, 119, 105505.	4.1	13
4	Selective multi-kinase inhibition sensitizes mesenchymal pancreatic cancer to immune checkpoint blockade by remodeling the tumor microenvironment. <i>Nature Cancer</i> , 2022, 3, 318-336.	13.2	42
5	Genetic alterations of the SUMO isopeptidase SENP6 drive lymphomagenesis and genetic instability in diffuse large B-cell lymphoma. <i>Nature Communications</i> , 2022, 13, 281.	12.8	18
6	Identification of treatment-induced vulnerabilities in pancreatic cancer patients using functional model systems. <i>EMBO Molecular Medicine</i> , 2022, 14, e14876.	6.9	20
7	Indirect targeting of MYC sensitizes pancreatic cancer cells to mechanistic target of rapamycin (mTOR) inhibition. <i>Cancer Communications</i> , 2022, , .	9.2	1
8	CRISPR somatic genome engineering and cancer modeling in the mouse pancreas and liver. <i>Nature Protocols</i> , 2022, 17, 1142-1188.	12.0	13
9	Spontaneous activity of the mitochondrial apoptosis pathway drives chromosomal defects, the appearance of micronuclei and cancer metastasis through the Caspase-Activated DNase. <i>Cell Death and Disease</i> , 2022, 13, 315.	6.3	14
10	Comparative Study of the Role of Interepithelial Mucosal Mast Cells in the Context of Intestinal Adenoma-Carcinoma Progression. <i>Cancers</i> , 2022, 14, 2248.	3.7	3
11	Abstract 2514: Pancreatic cancer subtype-specific secreted factors determine the immunosuppressive tumor microenvironment. <i>Cancer Research</i> , 2022, 82, 2514-2514.	0.9	0
12	Tutorial: design and execution of CRISPR in vivo screens. <i>Nature Protocols</i> , 2022, 17, 1903-1925.	12.0	12
13	IL-24 intrinsically regulates Th17 cell pathogenicity in mice. <i>Journal of Experimental Medicine</i> , 2022, 219, .	8.5	4
14	Synergistic targeting and resistance to PARP inhibition in DNA damage repair-deficient pancreatic cancer. <i>Gut</i> , 2021, 70, 743-760.	12.1	49
15	Notch2-mediated plasticity between marginal zone and follicular B cells. <i>Nature Communications</i> , 2021, 12, 1111.	12.8	26
16	The NFIB-ERO1A axis promotes breast cancer metastatic colonization of disseminated tumour cells. <i>EMBO Molecular Medicine</i> , 2021, 13, e13162.	6.9	27
17	Auto-aggressive CXCR6+ CD8 T cells cause liver immune pathology in NASH. <i>Nature</i> , 2021, 592, 444-449.	27.8	233
18	NASH limits anti-tumour surveillance in immunotherapy-treated HCC. <i>Nature</i> , 2021, 592, 450-456.	27.8	649

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19	Anti-inflammatory chemoprevention attenuates the phenotype in a mouse model of esophageal adenocarcinoma. <i>Carcinogenesis</i> , 2021, 42, 1068-1078.	2.8	4
20	Multilevel proteomics reveals host perturbations by SARS-CoV-2 and SARS-CoV. <i>Nature</i> , 2021, 594, 246-252.	27.8	475
21	SETBP1 overexpression acts in the place of class-defining mutations to drive FLT3-ITD mutant AML. <i>Blood Advances</i> , 2021, 5, 2412-2425.	5.2	10
22	Skin and gut imprinted helper T cell subsets exhibit distinct functional phenotypes in central nervous system autoimmunity. <i>Nature Immunology</i> , 2021, 22, 880-892.	14.5	34
23	Deep learning boosts sensitivity of mass spectrometry-based immunopeptidomics. <i>Nature Communications</i> , 2021, 12, 3346.	12.8	90
24	Modeling plasticity and dysplasia of pancreatic ductal organoids derived from human pluripotent stem cells. <i>Cell Stem Cell</i> , 2021, 28, 1105-1124.e19.	11.1	53
25	Whole Exome Sequencing of Biliary Tubulopapillary Neoplasms Reveals Common Mutations in Chromatin Remodeling Genes. <i>Cancers</i> , 2021, 13, 2742.	3.7	10
26	Targeted PI3K/AKT-hyperactivation induces cell death in chronic lymphocytic leukemia. <i>Nature Communications</i> , 2021, 12, 3526.	12.8	34
27	Genetic Screens Identify a Context-Specific PI3K/p27Kip1 Node Driving Extrahepatic Biliary Cancer. <i>Cancer Discovery</i> , 2021, 11, 3158-3177.	9.4	12
28	AGR2-Dependent Nuclear Import of RNA Polymerase II Constitutes a Specific Target of Pancreatic Ductal Adenocarcinoma in the Context of Wild-Type p53. <i>Gastroenterology</i> , 2021, 161, 1601-1614.e23.	1.3	10
29	Important role of Nfkb2 in the KrasG12D-driven carcinogenesis in the pancreas. <i>Pancreatology</i> , 2021, 21, 912-919.	1.1	3
30	Brief homogeneous TCR signals instruct common iNKT progenitors whose effector diversification is characterized by subsequent cytokine signaling. <i>Immunity</i> , 2021, 54, 2497-2513.e9.	14.3	19
31	Functional analysis of peripheral and intratumoral neoantigen-specific TCRs identified in a patient with melanoma. , 2021, 9, e002754.		7
32	XIAP restrains TNF-driven intestinal inflammation and dysbiosis by promoting innate immune responses of Paneth and dendritic cells. <i>Science Immunology</i> , 2021, 6, eabf7235.	11.9	17
33	High-Fructose Diet Alters Intestinal Microbial Profile and Correlates with Early Tumorigenesis in a Mouse Model of Barrett's Esophagus. <i>Microorganisms</i> , 2021, 9, 2432.	3.6	7
34	IFN-Gamma Producing Regulatory T Cells Counterbalance T Cell-Mediated Injury to the Intestinal Stem Cell Compartment in Mice and Humans. <i>Blood</i> , 2021, 138, 89-89.	1.4	1
35	Analysis pipelines for cancer genome sequencing in mice. <i>Nature Protocols</i> , 2020, 15, 266-315.	12.0	25
36	Angiocrine Hepatocyte Growth Factor Signaling Controls Physiological Organ and Body Size and Dynamic Hepatocyte Proliferation to Prevent Liver Damage during Regeneration. <i>American Journal of Pathology</i> , 2020, 190, 358-371.	3.8	24

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37	PiggyBac mutagenesis and exome sequencing identify genetic driver landscapes and potential therapeutic targets of EGFR-mutant gliomas. <i>Genome Biology</i> , 2020, 21, 181.	8.8	18
38	Targeting the ubiquitin-proteasome system in a pancreatic cancer subtype with hyperactive MYC. <i>Molecular Oncology</i> , 2020, 14, 3048-3064.	4.6	13
39	Linkage of genetic drivers and strain-specific germline variants confound mouse cancer genome analyses. <i>Nature Communications</i> , 2020, 11, 4474.	12.8	1
40	Genetically Engineered Mouse Models of Liver Tumorigenesis Reveal a Wide Histological Spectrum of Neoplastic and Non-Neoplastic Liver Lesions. <i>Cancers</i> , 2020, 12, 2265.	3.7	5
41	Mir34a constrains pancreatic carcinogenesis. <i>Scientific Reports</i> , 2020, 10, 9654.	3.3	10
42	Novel role for CRK adaptor proteins as essential components of SRC/FAK signaling for epithelial-mesenchymal transition and colorectal cancer aggressiveness. <i>International Journal of Cancer</i> , 2020, 147, 1715-1731.	5.1	14
43	In vivo functional screening for systems-level integrative cancer genomics. <i>Nature Reviews Cancer</i> , 2020, 20, 573-593.	28.4	44
44	Generation of An Endogenous FGFR2-BICC1 Gene Fusion/58 Megabase Inversion Using Single-Plasmid CRISPR/Cas9 Editing in Biliary Cells. <i>International Journal of Molecular Sciences</i> , 2020, 21, 2460.	4.1	3
45	c-Rel gain in B cells drives germinal center reactions and autoantibody production. <i>Journal of Clinical Investigation</i> , 2020, 130, 3270-3286.	8.2	11
46	Setbp1 Overexpression Acts in the Place of Class-Defining Somatic Mutations to Drive Mouse and Human FLT3-ITD-Mutant AMLs. <i>Blood</i> , 2020, 136, 31-32.	1.4	0
47	Editorial overview: Functionalizing cancer genomes in the era of big data. <i>Current Opinion in Genetics and Development</i> , 2019, 54, iii-vi.	3.3	1
48	Targeting c-MYC through Interference with NAMPT and SIRT1 and Their Association to Oncogenic Drivers in Murine Serrated Intestinal Tumorigenesis. <i>Neoplasia</i> , 2019, 21, 974-988.	5.3	9
49	RIG-I activation is critical for responsiveness to checkpoint blockade. <i>Science Immunology</i> , 2019, 4, .	11.9	80
50	Engineering CRISPR mouse models of cancer. <i>Current Opinion in Genetics and Development</i> , 2019, 54, 88-96.	3.3	25
51	PiggyBac transposon tools for recessive screening identify B-cell lymphoma drivers in mice. <i>Nature Communications</i> , 2019, 10, 1415.	12.8	37
52	Platelet GPIIb/IIIa is a mediator and potential interventional target for NASH and subsequent liver cancer. <i>Nature Medicine</i> , 2019, 25, 641-655.	30.7	259
53	Blimp1 Prevents Methylation of Foxp3 and Loss of Regulatory T Cell Identity at Sites of Inflammation. <i>Cell Reports</i> , 2019, 26, 1854-1868.e5.	6.4	91
54	DNA methylation instability by BRAF-mediated TET silencing and lifestyle-exposure divides colon cancer pathways. <i>Clinical Epigenetics</i> , 2019, 11, 196.	4.1	22

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55	PiggyBac Transposon-Based Insertional Mutagenesis in Mice. <i>Methods in Molecular Biology</i> , 2019, 1907, 171-183.	0.9	6
56	Evolutionary routes and KRAS dosage define pancreatic cancer phenotypes. <i>Nature</i> , 2018, 554, 62-68.	27.8	328
57	MTOR inhibitor-based combination therapies for pancreatic cancer. <i>British Journal of Cancer</i> , 2018, 118, 366-377.	6.4	35
58	Oncogenic Amplification of Zygotic Dux Factors in Regenerating p53-Deficient Muscle Stem Cells Defines a Molecular Cancer Subtype. <i>Cell Stem Cell</i> , 2018, 23, 794-805.e4.	11.1	21
59	SRPK1 maintains acute myeloid leukemia through effects on isoform usage of epigenetic regulators including BRD4. <i>Nature Communications</i> , 2018, 9, 5378.	12.8	60
60	Myeloid-derived suppressor cells control B cell accumulation in the central nervous system during autoimmunity. <i>Nature Immunology</i> , 2018, 19, 1341-1351.	14.5	82
61	Preclinical Evaluation of the Hsp70 Peptide Tracer TPP-PEG24-DFO[89Zr] for Tumor-Specific PET/CT Imaging. <i>Cancer Research</i> , 2018, 78, 6268-6281.	0.9	32
62	Genome-wide transposon screening and quantitative insertion site sequencing for cancer gene discovery in mice. <i>Nature Protocols</i> , 2017, 12, 289-309.	12.0	41
63	Resistance mechanisms to TP53-MDM2 inhibition identified by in vivo piggyBac transposon mutagenesis screen in an Arf ^{fl/fl} mouse model. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 3151-3156.	7.1	48
64	Tissue-specific tumorigenesis: context matters. <i>Nature Reviews Cancer</i> , 2017, 17, 239-253.	28.4	234
65	Kupffer Cell-Derived Tnf Triggers Cholangiocellular Tumorigenesis through JNK due to Chronic Mitochondrial Dysfunction and ROS. <i>Cancer Cell</i> , 2017, 31, 771-789.e6.	16.8	140
66	A single-copy Sleeping Beauty transposon mutagenesis screen identifies new PTEN-cooperating tumor suppressor genes. <i>Nature Genetics</i> , 2017, 49, 730-741.	21.4	53
67	Molecular synergy underlies the co-occurrence patterns and phenotype of NPM1-mutant acute myeloid leukemia. <i>Blood</i> , 2017, 130, 1911-1922.	1.4	63
68	PD-1 is a haploinsufficient suppressor of T cell lymphomagenesis. <i>Nature</i> , 2017, 552, 121-125.	27.8	199
69	RIPK3 Restricts Myeloid Leukemogenesis by Promoting Cell Death and Differentiation of Leukemia Initiating Cells. <i>Cancer Cell</i> , 2016, 30, 75-91.	16.8	144
70	siRNA-coupled nanoparticles for improved therapeutic targeting of pancreatic cancer. <i>Gut</i> , 2016, 65, 1780-1781.	12.1	1
71	Multiplexed pancreatic genome engineering and cancer induction by transfection-based CRISPR/Cas9 delivery in mice. <i>Nature Communications</i> , 2016, 7, 10770.	12.8	145
72	Characterisation of worldwide <i>Helicobacter pylori</i> strains reveals genetic conservation and essentiality of serine protease HtrA. <i>Molecular Microbiology</i> , 2016, 99, 925-944.	2.5	70

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73	Development and validation of a comprehensive genomic diagnostic tool for myeloid malignancies. <i>Blood</i> , 2016, 128, e1-e9.	1.4	49
74	Direct identification of clinically relevant neoepitopes presented on native human melanoma tissue by mass spectrometry. <i>Nature Communications</i> , 2016, 7, 13404.	12.8	613
75	Leukemia-Associated Somatic Mutations Drive Distinct Patterns of Age-Related Clonal Hemopoiesis. <i>Cell Reports</i> , 2015, 10, 1239-1245.	6.4	443
76	A Synergistic Interaction between Chk1- and MK2 Inhibitors in KRAS-Mutant Cancer. <i>Cell</i> , 2015, 162, 146-159.	28.9	100
77	CRISPR/Cas9 somatic multiplex-mutagenesis for high-throughput functional cancer genomics in mice. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 13982-13987.	7.1	172
78	The E3 ligase RNF43 inhibits Wnt signaling downstream of mutated β -catenin by sequestering TCF4 to the nuclear membrane. <i>Science Signaling</i> , 2015, 8, ra90.	3.6	67
79	A conditional piggyBac transposition system for genetic screening in mice identifies oncogenic networks in pancreatic cancer. <i>Nature Genetics</i> , 2015, 47, 47-56.	21.4	77
80	Tumor Imaging and Targeting Potential of an Hsp70-Derived 14-Mer Peptide. <i>PLoS ONE</i> , 2014, 9, e105344.	2.5	29
81	Chromatin Landscapes of Retroviral and Transposon Integration Profiles. <i>PLoS Genetics</i> , 2014, 10, e1004250.	3.5	80
82	Disruption of the PRKCD-FBXO25-HAX-1 axis attenuates the apoptotic response and drives lymphomagenesis. <i>Nature Medicine</i> , 2014, 20, 1401-1409.	30.7	50
83	A next-generation dual-recombinase system for time- and host-specific targeting of pancreatic cancer. <i>Nature Medicine</i> , 2014, 20, 1340-1347.	30.7	188
84	A Genetic Progression Model of BrafV600E-Induced Intestinal Tumorigenesis Reveals Targets for Therapeutic Intervention. <i>Cancer Cell</i> , 2013, 24, 15-29.	16.8	183
85	Selective Requirement of PI3K/PDK1 Signaling for Kras Oncogene-Driven Pancreatic Cell Plasticity and Cancer. <i>Cancer Cell</i> , 2013, 23, 406-420.	16.8	291
86	Clonal Expansion Analysis of Transposon Insertions by High-Throughput Sequencing Identifies Candidate Cancer Genes in a PiggyBac Mutagenesis Screen. <i>PLoS ONE</i> , 2013, 8, e72338.	2.5	12
87	Mutant nucleophosmin and cooperating pathways drive leukemia initiation and progression in mice. <i>Nature Genetics</i> , 2011, 43, 470-475.	21.4	194
88	<i>PiggyBac</i> Transposon Mutagenesis: A Tool for Cancer Gene Discovery in Mice. <i>Science</i> , 2010, 330, 1104-1107.	12.6	217
89	Extracellular and Intracellular Pattern Recognition Receptors Cooperate in the Recognition of <i>Helicobacter pylori</i> . <i>Gastroenterology</i> , 2009, 136, 2247-2257.	1.3	162
90	Toll-Like Receptor-Dependent Activation of Antigen-Presenting Cells Affects Adaptive Immunity to <i>Helicobacter pylori</i> . <i>Gastroenterology</i> , 2007, 133, 150-163.e3.	1.3	80

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91	CD25+/Foxp3+ T Cells Regulate Gastric Inflammation and <i>Helicobacter pylori</i> Colonization In Vivo. <i>Gastroenterology</i> , 2006, 131, 525-537.	1.3	251
92	Synergistic Effect of <i>Helicobacter pylori</i> Virulence Factors and Interleukin-1 Polymorphisms for the Development of Severe Histological Changes in the Gastric Mucosa. <i>Journal of Infectious Diseases</i> , 2003, 188, 272-281.	4.0	175
93	The <i>Helicobacter pylori</i> Blood Group Antigen-Binding Adhesin Facilitates Bacterial Colonization and Augments a Nonspecific Immune Response. <i>Journal of Immunology</i> , 2002, 168, 3033-3041.	0.8	166
94	Analyse von Krebsgenen: Schnelle Suche nach der "Nadel im Heuhaufen". , 0, , .		0