

Claudia R Ball

List of Publications by Year in descending order

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Version: 2024-02-01

42
papers

4,673
citations

331670

21
h-index

289244

40
g-index

43
all docs

43
docs citations

43
times ranked

6775
citing authors

#	ARTICLE	IF	CITATIONS
1	Tumororganoide als präklinische Tumormodelle – Implikationen für Knochen- und Weichteiltumore. <i>Osteologie</i> , 2022, 31, 71-72.	0.1	0
2	Rational design of aqueous conjugated polymer nanoparticles as potential theranostic agents of breast cancer. <i>Materials Chemistry Frontiers</i> , 2021, 5, 4950-4962.	5.9	7
3	Functional States in Tumor-Initiating Cell Differentiation in Human Colorectal Cancer. <i>Cancers</i> , 2021, 13, 1097.	3.7	11
4	The balance between the intronic miR-342 and its host gene <i>Evl</i> determines hematopoietic cell fate decision. <i>Leukemia</i> , 2021, 35, 2948-2963.	7.2	9
5	Degradation of CCNK/CDK12 is a druggable vulnerability of colorectal cancer. <i>Cell Reports</i> , 2021, 36, 109394.	6.4	41
6	YAP-induced <i>Ccl2</i> expression is associated with a switch in hepatic macrophage identity and vascular remodelling in liver cancer. <i>Liver International</i> , 2021, 41, 3011-3023.	3.9	17
7	YAP Orchestrates Heterotypic Endothelial Cell Communication via HGF/c-MET Signaling in Liver Tumorigenesis. <i>Cancer Research</i> , 2020, 80, 5502-5514.	0.9	31
8	Comprehensive genomic characterization of gene therapy-induced T-cell acute lymphoblastic leukemia. <i>Leukemia</i> , 2020, 34, 2785-2789.	7.2	4
9	A perivascular niche in the bone marrow hosts quiescent and proliferating tumorigenic colorectal cancer cells. <i>International Journal of Cancer</i> , 2020, 147, 519-531.	5.1	5
10	Transcriptome Profiling of Adipose Tissue Reveals Depot-Specific Metabolic Alterations Among Patients with Colorectal Cancer. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2019, 104, 5225-5237.	3.6	21
11	Pheno-seq – linking visual features and gene expression in 3D cell culture systems. <i>Scientific Reports</i> , 2019, 9, 12367.	3.3	16
12	Salinomycin: Anti-tumor activity in a pre-clinical colorectal cancer model. <i>PLoS ONE</i> , 2019, 14, e0211916.	2.5	27
13	Systematic Generation of Patient-Derived Tumor Models in Pancreatic Cancer. <i>Cells</i> , 2019, 8, 142.	4.1	9
14	The notch target gene <i>HEYL</i> modulates metastasis forming capacity of colorectal cancer patient-derived spheroid cells in vivo. <i>BMC Cancer</i> , 2019, 19, 1181.	2.6	16
15	Targeted BiTE Expression by an Oncolytic Vector Augments Therapeutic Efficacy Against Solid Tumors. <i>Clinical Cancer Research</i> , 2018, 24, 2128-2137.	7.0	88
16	<i>NRG1</i> Fusions in <i>KRAS</i> Wild-Type Pancreatic Cancer. <i>Cancer Discovery</i> , 2018, 8, 1087-1095.	9.4	189
17	Mapping Active Gene-Regulatory Regions in Human Repopulating Long-Term HSCs. <i>Cell Stem Cell</i> , 2018, 23, 132-146.e9.	11.1	14
18	Cell-of-Origin DNA Methylation Signatures Are Maintained during Colorectal Carcinogenesis. <i>Cell Reports</i> , 2018, 23, 3407-3418.	6.4	66

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19	Succession of transiently active tumor-initiating cell clones in human pancreatic cancer xenografts. <i>EMBO Molecular Medicine</i> , 2017, 9, 918-932.	6.9	36
20	Genetic subclone architecture of tumor clone-initiating cells in colorectal cancer. <i>Journal of Experimental Medicine</i> , 2017, 214, 2073-2088.	8.5	30
21	Patient-derived xenografts of gastrointestinal cancers are susceptible to rapid and delayed B-lymphoproliferation. <i>International Journal of Cancer</i> , 2017, 140, 1356-1363.	5.1	26
22	Pan-cancer analysis of somatic copy-number alterations implicates <i>IRS4</i> and <i>IGF2</i> in enhancer hijacking. <i>Nature Genetics</i> , 2017, 49, 65-74.	21.4	326
23	Colorectal cancer-initiating cells caught in the act. <i>EMBO Molecular Medicine</i> , 2017, 9, 856-858.	6.9	12
24	Phenotypic differentiation does not affect tumorigenicity of primary human colon cancer initiating cells. <i>Cancer Letters</i> , 2016, 371, 326-333.	7.2	11
25	Wnt secretion is required to maintain high levels of Wnt activity in colon cancer cells. <i>Nature Communications</i> , 2013, 4, 2610.	12.8	213
26	Specific Elimination of CD133+ Tumor Cells with Targeted Oncolytic Measles Virus. <i>Cancer Research</i> , 2013, 73, 865-874.	0.9	115
27	Transforming Growth Factor Beta Receptor 2 (<i>TGFBR2</i>) Changes Sialylation in the Microsatellite Unstable (MSI) Colorectal Cancer Cell Line HCT116. <i>PLoS ONE</i> , 2013, 8, e57074.	2.5	28
28	Overexpression of <i>EV11</i> interferes with cytokinesis and leads to accumulation of cells with supernumerary centrosomes in G _{0/1} phase. <i>Cell Cycle</i> , 2012, 11, 3492-3503.	2.6	21
29	Targeting of <i>KRAS</i> mutant tumors by <i>HSP90</i> inhibitors involves degradation of <i>STK33</i> . <i>Journal of Experimental Medicine</i> , 2012, 209, 697-711.	8.5	63
30	Stable Long-Term Blood Formation by Stem Cells in Murine Steady-State Hematopoiesis. <i>Stem Cells</i> , 2012, 30, 1961-1970.	3.2	11
31	Extensive Methylation of Promoter Sequences Silences Lentiviral Transgene Expression During Stem Cell Differentiation In Vivo. <i>Molecular Therapy</i> , 2012, 20, 1014-1021.	8.2	87
32	Distinct Types of Tumor-Initiating Cells Form Human Colon Cancer Tumors and Metastases. <i>Cell Stem Cell</i> , 2011, 9, 357-365.	11.1	276
33	You Can Count on This: Barcoded Hematopoietic Stem Cells. <i>Cell Stem Cell</i> , 2011, 9, 390-392.	11.1	10
34	Deregulated <i>EV11</i> Expression Leads to Genomic Instability and G1 Cell Cycle Arrest. <i>Blood</i> , 2011, 118, 2431-2431.	1.4	0
35	Hematopoietic activity of human short-term repopulating cells in mobilized peripheral blood cell transplants is restricted to the first 5 months after transplantation. <i>Blood</i> , 2010, 115, 5023-5025.	1.4	3
36	Genomic instability and myelodysplasia with monosomy 7 consequent to <i>EV11</i> activation after gene therapy for chronic granulomatous disease. <i>Nature Medicine</i> , 2010, 16, 198-204.	30.7	727

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37	The Inherent Differentiation Program of Short-Term Hematopoietic Repopulating Cells Changes During Human Ontogeny. <i>Stem Cells and Development</i> , 2010, 19, 621-628.	2.1	3
38	Stem-Cell Gene Therapy for the Wiskottâ€Aldrich Syndrome. <i>New England Journal of Medicine</i> , 2010, 363, 1918-1927.	27.0	505
39	Comprehensive genomic access to vector integration in clinical gene therapy. <i>Nature Medicine</i> , 2009, 15, 1431-1436.	30.7	173
40	Stable differentiation and clonality of murine long-term hematopoiesis after extended reduced-intensity selection for MGMT P140K transgene expression. <i>Blood</i> , 2007, 110, 1779-1787.	1.4	16
41	High-resolution insertion-site analysis by linear amplificationâ€mediated PCR (LAM-PCR). <i>Nature Methods</i> , 2007, 4, 1051-1057.	19.0	281
42	Correction of X-linked chronic granulomatous disease by gene therapy, augmented by insertional activation of MDS1-EVI1, PRDM16 or SETBP1. <i>Nature Medicine</i> , 2006, 12, 401-409.	30.7	1,129