

Kimberly Stegmaier

List of Publications by Year in descending order

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Version: 2024-02-01

108
papers

13,302
citations

61984

43
h-index

33894

99
g-index

116
all docs

116
docs citations

116
times ranked

26068
citing authors

#	ARTICLE	IF	CITATIONS
1	Targeting serine hydroxymethyltransferases 1 and 2 for T-cell acute lymphoblastic leukemia therapy. <i>Leukemia</i> , 2022, 36, 348-360.	7.2	23
2	An <i>In Vivo</i> CRISPR Screening Platform for Prioritizing Therapeutic Targets in AML. <i>Cancer Discovery</i> , 2022, 12, 432-449.	9.4	32
3	SHMT2 inhibition disrupts the TCF3 transcriptional survival program in Burkitt lymphoma. <i>Blood</i> , 2022, 139, 538-553.	1.4	27
4	EP300 Selectively Controls the Enhancer Landscape of MYCN-Amplified Neuroblastoma. <i>Cancer Discovery</i> , 2022, 12, 730-751.	9.4	64
5	The proteogenomic subtypes of acute myeloid leukemia. <i>Cancer Cell</i> , 2022, 40, 301-317.e12.	16.8	43
6	A distinct core regulatory module enforces oncogene expression in KMT2A-rearranged leukemia. <i>Genes and Development</i> , 2022, 36, 368-389.	5.9	14
7	Cystine uptake inhibition potentiates front-line therapies in acute myeloid leukemia. <i>Leukemia</i> , 2022, 36, 1585-1595.	7.2	24
8	IKAROS and MENIN coordinate therapeutically actionable leukemogenic gene expression in MLL-r acute myeloid leukemia. <i>Nature Cancer</i> , 2022, 3, 595-613.	13.2	16
9	Identification of an Epi-metabolic dependency on EHMT2/G9a in T-cell acute lymphoblastic leukemia. <i>Cell Death and Disease</i> , 2022, 13, .	6.3	6
10	Abstract 3889: Identification of ADRN-specific, MES-specific, and pan-subtype therapeutic targets in neuroblastoma. <i>Cancer Research</i> , 2022, 82, 3889-3889.	0.9	0
11	Transcriptional Plasticity Drives Leukemia Immune Escape. <i>Blood Cancer Discovery</i> , 2022, 3, 394-409.	5.0	8
12	Transition to a mesenchymal state in neuroblastoma confers resistance to anti-GD2 antibody via reduced expression of ST8SIA1. <i>Nature Cancer</i> , 2022, 3, 976-993.	13.2	23
13	Network-based systems pharmacology reveals heterogeneity in LCK and BCL2 signaling and therapeutic sensitivity of T-cell acute lymphoblastic leukemia. <i>Nature Cancer</i> , 2021, 2, 284-299.	13.2	70
14	Matched Targeted Therapy for Pediatric Patients with Relapsed, Refractory, or High-Risk Leukemias: A Report from the LEAP Consortium. <i>Cancer Discovery</i> , 2021, 11, 1424-1439.	9.4	16
15	Targeting acute myeloid leukemia dependency on VCP-mediated DNA repair through a selective second-generation small-molecule inhibitor. <i>Science Translational Medicine</i> , 2021, 13, .	12.4	29
16	RAD21 is a driver of chromosome 8 gain in Ewing sarcoma to mitigate replication stress. <i>Genes and Development</i> , 2021, 35, 556-572.	5.9	28
17	A first-generation pediatric cancer dependency map. <i>Nature Genetics</i> , 2021, 53, 529-538.	21.4	76
18	Selective Modulation of a Pan-Essential Protein as a Therapeutic Strategy in Cancer. <i>Cancer Discovery</i> , 2021, 11, 2282-2299.	9.4	21

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19	Gene Fusions Create Partner and Collateral Dependencies Essential to Cancer Cell Survival. <i>Cancer Research</i> , 2021, 81, 3971-3984.	0.9	11
20	The synergy of BET inhibitors with aurora A kinase inhibitors in MYCN-amplified neuroblastoma is heightened with functional TP53. <i>Neoplasia</i> , 2021, 23, 624-633.	5.3	8
21	TRIM8 modulates the EWS/FLI oncoprotein to promote survival in Ewing sarcoma. <i>Cancer Cell</i> , 2021, 39, 1262-1278.e7.	16.8	49
22	Structure-activity relationship study of THZ531 derivatives enables the discovery of BSJ-01-175 as a dual CDK12/13 covalent inhibitor with efficacy in Ewing sarcoma. <i>European Journal of Medicinal Chemistry</i> , 2021, 221, 113481.	5.5	27
23	Transcriptional Immunoediting of AML Cells after Allogeneic Hematopoietic Stem Cell Transplantation. <i>Blood</i> , 2021, 138, 647-647.	1.4	0
24	Alisertib Synergistically Strengthens the Anti-Leukemia Activity of Venetoclax in <i>TCF3-Hlf</i> B-ALL. <i>Blood</i> , 2021, 138, 705-705.	1.4	0
25	Resistance Mechanisms to SYK Inhibition in Acute Myeloid Leukemia. <i>Cancer Discovery</i> , 2020, 10, 214-231.	9.4	27
26	Rapid and direct control of target protein levels with VHL-recruiting dTAG molecules. <i>Nature Communications</i> , 2020, 11, 4687.	12.8	129
27	The Folate Cycle Enzyme MTHFR Is a Critical Regulator of Cell Response to MYC-Targeting Therapies. <i>Cancer Discovery</i> , 2020, 10, 1894-1911.	9.4	13
28	Synthetic Lethal Interaction between the ESCRT Paralog Enzymes VPS4A and VPS4B in Cancers Harboring Loss of Chromosome 18q or 16q. <i>Cell Reports</i> , 2020, 33, 108493.	6.4	28
29	Selective USP7 inhibition elicits cancer cell killing through a p53-dependent mechanism. <i>Scientific Reports</i> , 2020, 10, 5324.	3.3	69
30	Blockade of Oncogenic NOTCH1 with the SERCA Inhibitor CAD204520 in T Cell Acute Lymphoblastic Leukemia. <i>Cell Chemical Biology</i> , 2020, 27, 678-697.e13.	5.2	27
31	Targeting DUBs to degrade oncogenic proteins. <i>British Journal of Cancer</i> , 2020, 122, 1121-1123.	6.4	6
32	Preclinical efficacy for a novel tyrosine kinase inhibitor, ArQule 531 against acute myeloid leukemia. <i>Journal of Hematology and Oncology</i> , 2020, 13, 8.	17.0	16
33	Salt-inducible kinase inhibition suppresses acute myeloid leukemia progression in vivo. <i>Blood</i> , 2020, 135, 56-70.	1.4	49
34	Targeting EZH2 for the treatment of hepatosplenic T-cell lymphoma. <i>Blood Advances</i> , 2020, 4, 1265-1269.	5.2	5
35	EWS-FLI1 modulated alternative splicing of ARID1A reveals novel oncogenic function through the BAF complex. <i>Nucleic Acids Research</i> , 2019, 47, 9619-9636.	14.5	35
36	Small-Molecule and CRISPR Screening Converge to Reveal Receptor Tyrosine Kinase Dependencies in Pediatric Rhabdoid Tumors. <i>Cell Reports</i> , 2019, 28, 2331-2344.e8.	6.4	24

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37	Single-cell cloning of human T-cell lines reveals clonal variation in cell death responses to chemotherapeutics. <i>Cancer Genetics</i> , 2019, 237, 69-77.	0.4	6
38	Synthetic Lethality of Wnt Pathway Activation and Asparaginase in Drug-Resistant Acute Leukemias. <i>Cancer Cell</i> , 2019, 35, 664-676.e7.	16.8	70
39	Ushering in the next generation of precision trials for pediatric cancer. <i>Science</i> , 2019, 363, 1175-1181.	12.6	41
40	Targeting chromatin complexes in fusion protein-driven malignancies. <i>Nature Reviews Cancer</i> , 2019, 19, 255-269.	28.4	55
41	MDM2 and MDM4 Are Therapeutic Vulnerabilities in Malignant Rhabdoid Tumors. <i>Cancer Research</i> , 2019, 79, 2404-2414.	0.9	43
42	Virtual Screening Identifies Irreversible FMS-like Tyrosine Kinase 3 Inhibitors with Activity toward Resistance-Confering Mutations. <i>Journal of Medicinal Chemistry</i> , 2019, 62, 2428-2446.	6.4	32
43	Creatine kinase pathway inhibition alters GSK3 and WNT signaling in EVI1-positive AML. <i>Leukemia</i> , 2019, 33, 800-804.	7.2	10
44	A Combination CDK4/6 and IGF1R Inhibitor Strategy for Ewing Sarcoma. <i>Clinical Cancer Research</i> , 2019, 25, 1343-1357.	7.0	69
45	Biology and Therapy of Dominant Fusion Oncoproteins Involving Transcription Factor and Chromatin Regulators in Sarcomas. <i>Annual Review of Cancer Biology</i> , 2019, 3, 299-321.	4.5	25
46	Therapeutic discovery for marrow failure with MDS predisposition using pluripotent stem cells. <i>JCI Insight</i> , 2019, 4, .	5.0	10
47	The Folate Cycle Enzyme MTHFR Is a Critical Regulator of Cell Response to MYC-Targeting Therapies. <i>Blood</i> , 2019, 134, 877-877.	1.4	1
48	Bepridil exhibits anti-leukemic activity associated with NOTCH1 pathway inhibition in chronic lymphocytic leukemia. <i>International Journal of Cancer</i> , 2018, 143, 958-970.	5.1	32
49	Exploiting an Asp-Glu "switch" in glycogen synthase kinase 3 to design paralog-selective inhibitors for use in acute myeloid leukemia. <i>Science Translational Medicine</i> , 2018, 10, .	12.4	69
50	EWS/FLI1 increases transcription to cause R-loops and block BRCA1 repair in Ewing sarcoma. <i>Nature</i> , 2018, 555, 387-391.	27.8	222
51	Phase I trial of the mTOR inhibitor everolimus in combination with multi-agent chemotherapy in relapsed childhood acute lymphoblastic leukemia. <i>Pediatric Blood and Cancer</i> , 2018, 65, e27062.	1.5	48
52	EWS/FLI Confers Tumor Cell Synthetic Lethality to CDK12 Inhibition in Ewing Sarcoma. <i>Cancer Cell</i> , 2018, 33, 202-216.e6.	16.8	116
53	Leukemia-specific delivery of mutant NOTCH1 targeted therapy. <i>Journal of Experimental Medicine</i> , 2018, 215, 197-216.	8.5	30
54	Resistance to Epigenetic-Targeted Therapy Engenders Tumor Cell Vulnerabilities Associated with Enhancer Remodeling. <i>Cancer Cell</i> , 2018, 34, 922-938.e7.	16.8	63

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55	Precision Targeting of BFL-1/A1 and an ATM Co-dependency in Human Cancer. <i>Cell Reports</i> , 2018, 24, 3393-3403.e5.	6.4	15
56	TRPS1 Is a Lineage-Specific Transcriptional Dependency in Breast Cancer. <i>Cell Reports</i> , 2018, 25, 1255-1267.e5.	6.4	46
57	Genome-scale CRISPR-Cas9 screen identifies druggable dependencies in TP53 wild-type Ewing sarcoma. <i>Journal of Experimental Medicine</i> , 2018, 215, 2137-2155.	8.5	55
58	Targeted therapy for fusion-driven high-risk acute leukemia. <i>Blood</i> , 2018, 132, 1241-1247.	1.4	22
59	Selective gene dependencies in MYCN-amplified neuroblastoma include the core transcriptional regulatory circuitry. <i>Nature Genetics</i> , 2018, 50, 1240-1246.	21.4	199
60	Comparative proteomics reveals a diagnostic signature for pulmonary head&neck cancer metastasis. <i>EMBO Molecular Medicine</i> , 2018, 10, .	6.9	41
61	Detection of circulating tumour DNA is associated with inferior outcomes in Ewing sarcoma and osteosarcoma: a report from the Children's Oncology Group. <i>British Journal of Cancer</i> , 2018, 119, 615-621.	6.4	83
62	Matched Targeted Therapy for Pediatric Patients with Relapsed, Refractory or High-Risk Leukemias: A Report from the LEAP Consortium. <i>Blood</i> , 2018, 132, 261-261.	1.4	3
63	KLF5 controls glutathione metabolism to suppress p190-BCR-ABL+ B-cell lymphoblastic leukemia. <i>Oncotarget</i> , 2018, 9, 29665-29679.	1.8	6
64	The creatine kinase pathway is a metabolic vulnerability in EVI1-positive acute myeloid leukemia. <i>Nature Medicine</i> , 2017, 23, 301-313.	30.7	79
65	Synergistic Drug Combinations with a CDK4/6 Inhibitor in T-cell Acute Lymphoblastic Leukemia. <i>Clinical Cancer Research</i> , 2017, 23, 1012-1024.	7.0	88
66	Hoxa9 and Meis1 Cooperatively Induce Addiction to Syk Signaling by Suppressing miR-146a in Acute Myeloid Leukemia. <i>Cancer Cell</i> , 2017, 31, 549-562.e11.	16.8	89
67	Computational correction of copy number effect improves specificity of CRISPR-Cas9 essentiality screens in cancer cells. <i>Nature Genetics</i> , 2017, 49, 1779-1784.	21.4	1,436
68	TOX Regulates Growth, DNA Repair, and Genomic Instability in T-cell Acute Lymphoblastic Leukemia. <i>Cancer Discovery</i> , 2017, 7, 1336-1353.	9.4	48
69	Identification of an allosteric benzothiazolopyrimidone inhibitor of the oncogenic protein tyrosine phosphatase SHP2. <i>Bioorganic and Medicinal Chemistry</i> , 2017, 25, 6479-6485.	3.0	43
70	Scratching the Surface of Immunotherapeutic Targets in Neuroblastoma. <i>Cancer Cell</i> , 2017, 32, 271-273.	16.8	3
71	CRISPR-Cas9 screen reveals a MYCN-amplified neuroblastoma dependency on EZH2. <i>Journal of Clinical Investigation</i> , 2017, 128, 446-462.	8.2	117
72	Characterization of midostaurin as a dual inhibitor of FLT3 and SYK and potentiation of FLT3 inhibition against FLT3-ITD-driven leukemia harboring activated SYK kinase. <i>Oncotarget</i> , 2017, 8, 52026-52044.	1.8	19

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73	The second European interdisciplinary Ewing sarcoma research summit - A joint effort to deconstructing the multiple layers of a complex disease. <i>Oncotarget</i> , 2016, 7, 8613-8624.	1.8	55
74	Integrated genetic and pharmacologic interrogation of rare cancers. <i>Nature Communications</i> , 2016, 7, 11987.	12.8	45
75	The Public Repository of Xenografts Enables Discovery and Randomized Phase II-like Trials in Mice. <i>Cancer Cell</i> , 2016, 29, 574-586.	16.8	227
76	Inhibitors of Glycogen Synthase Kinase 3 with Exquisite Kinome-Wide Selectivity and Their Functional Effects. <i>ACS Chemical Biology</i> , 2016, 11, 1952-1963.	3.4	55
77	Targeting MTHFD2 in acute myeloid leukemia. <i>Journal of Experimental Medicine</i> , 2016, 213, 1285-1306.	8.5	118
78	Genomic Copy Number Dictates a Gene-Independent Cell Response to CRISPR/Cas9 Targeting. <i>Cancer Discovery</i> , 2016, 6, 914-929.	9.4	485
79	Multicenter Feasibility Study of Tumor Molecular Profiling to Inform Therapeutic Decisions in Advanced Pediatric Solid Tumors. <i>JAMA Oncology</i> , 2016, 2, 608.	7.1	172
80	Leukemia-Specific Delivery of Mutant NOTCH1 Targeted Therapy. <i>Blood</i> , 2016, 128, 889-889.	1.4	0
81	Evaluation of Improved Glycogen Synthase Kinase-3 β Inhibitors in Models of Acute Myeloid Leukemia. <i>Journal of Medicinal Chemistry</i> , 2015, 58, 8907-8919.	6.4	25
82	Increased SYK activity is associated with unfavorable outcome among patients with acute myeloid leukemia. <i>Oncotarget</i> , 2015, 6, 25575-25587.	1.8	20
83	Functional, chemical genomic, and super-enhancer screening identify sensitivity to cyclin D1/CDK4 pathway inhibition in Ewing sarcoma. <i>Oncotarget</i> , 2015, 6, 30178-30193.	1.8	68
84	Aberrant Activation of the PI3K/mTOR Pathway Promotes Resistance to Sorafenib in AML. <i>Blood</i> , 2015, 126, 2472-2472.	1.4	0
85	New Approaches to Target T-ALL. <i>Frontiers in Oncology</i> , 2014, 4, 170.	2.8	48
86	The Genomic Landscape of Pediatric Ewing Sarcoma. <i>Cancer Discovery</i> , 2014, 4, 1326-1341.	9.4	415
87	Targeting Csnk1a1 in leukemia. <i>Journal of Experimental Medicine</i> , 2014, 211, 594-594.	8.5	0
88	An epigenetic mechanism of resistance to targeted therapy in T cell acute lymphoblastic leukemia. <i>Nature Genetics</i> , 2014, 46, 364-370.	21.4	333
89	Triplication of a 21q22 region contributes to B cell transformation through HMGN1 overexpression and loss of histone H3 Lys27 trimethylation. <i>Nature Genetics</i> , 2014, 46, 618-623.	21.4	117
90	SYK Is a Critical Regulator of FLT3 in Acute Myeloid Leukemia. <i>Cancer Cell</i> , 2014, 25, 226-242.	16.8	126

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91	A phase II study of the EGFR inhibitor gefitinib in patients with acute myeloid leukemia. <i>Leukemia Research</i> , 2014, 38, 430-434.	0.8	23
92	Parallel genome-scale loss of function screens in 216 cancer cell lines for the identification of context-specific genetic dependencies. <i>Scientific Data</i> , 2014, 1, 140035.	5.3	328
93	Molecular rationale for the use of PI3K/AKT/mTOR pathway inhibitors in combination with crizotinib in <i><i>ALK</i>-mutated neuroblastoma. <i>Oncotarget</i>, 2014, 5, 8737-8749.</i>	1.8	72
94	Selective HDAC1/HDAC2 Inhibitors Induce Neuroblastoma Differentiation. <i>Chemistry and Biology</i> , 2013, 20, 713-725.	6.0	89
95	Complementary Genomic Screens Identify SERCA as a Therapeutic Target in NOTCH1 Mutated Cancer. <i>Cancer Cell</i> , 2013, 23, 390-405.	16.8	130
96	Mutational heterogeneity in cancer and the search for new cancer-associated genes. <i>Nature</i> , 2013, 499, 214-218.	27.8	4,761
97	Targeting MYCN in Neuroblastoma by BET Bromodomain Inhibition. <i>Cancer Discovery</i> , 2013, 3, 308-323.	9.4	549
98	Targeting Folate Metabolism In Acute Myelogenous Leukemia. <i>Blood</i> , 2013, 122, 3798-3798.	1.4	1
99	The intersection of genetic and chemical genomic screens identifies GSK-3 β as a target in human acute myeloid leukemia. <i>Journal of Clinical Investigation</i> , 2012, 122, 935-947.	8.2	96
100	Intersecting High-Throughput Screens Identifies SERCA As a Target for Modulating NOTCH1 In Hematopoietic Malignancies. <i>Blood</i> , 2011, 118, 555-555.	1.4	0
101	In Vivo RNA Interference Screening Identifies a Leukemia-Specific Dependence on Integrin Beta 3 Signaling. <i>Blood</i> , 2011, 118, 758-758.	1.4	0
102	Intersecting Chemical Genomic and Genetic Screens Identifies Glycogen Synthase Kinase-3 β (GSK-3 β) as a Modulator of Differentiation In Acute Myeloid Leukemia. <i>Blood</i> , 2010, 116, 1000-1000.	1.4	0
103	Proteomic and Genetic Approaches Identify Syk as an AML Target. <i>Cancer Cell</i> , 2009, 16, 281-294.	16.8	140
104	Chemical Genomic Screen Identifies Ionophores as Modulators of Notch1 in T-ALL. <i>Blood</i> , 2008, 112, 200-200.	1.4	0
105	Signature-Based Small Molecule Screening Identifies Cytosine Arabinoside as an EWS/FLI Modulator in Ewing Sarcoma. <i>PLoS Medicine</i> , 2007, 4, e122.	8.4	129
106	Genomic approaches in acute leukemia. <i>Best Practice and Research in Clinical Haematology</i> , 2006, 19, 263-268.	1.7	8
107	Modulating AML1-ETO with Signature-Based Small Molecule Library Screening.. <i>Blood</i> , 2006, 108, 729-729.	1.4	1
108	Gene expression \hat{c} -based high-throughput screening(GE-HTS) and application to leukemia differentiation. <i>Nature Genetics</i> , 2004, 36, 257-263.	21.4	276