

Paul H Huang

List of Publications by Year in descending order

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Version: 2024-02-01

96
papers

4,216
citations

159585

30
h-index

123424

61
g-index

103
all docs

103
docs citations

103
times ranked

6393
citing authors

#	ARTICLE	IF	CITATIONS
1	EGFR Exon 20 Insertion Mutations in Sinonasal Squamous Cell Carcinoma. <i>Cancers</i> , 2022, 14, 394.	3.7	7
2	KIT Exon 9-Mutated Gastrointestinal Stromal Tumours: Biology and Treatment. <i>Chemotherapy</i> , 2022, 67, 81-90.	1.6	10
3	Proteomic and Metabolomic Profiling in Soft Tissue Sarcomas. <i>Current Treatment Options in Oncology</i> , 2022, 23, 78-88.	3.0	10
4	Current Status and Future Directions of Immunotherapies in Soft Tissue Sarcomas. <i>Biomedicines</i> , 2022, 10, 573.	3.2	8
5	Amivantamab for the treatment of EGFR exon 20 insertion mutant non-small cell lung cancer. <i>Expert Review of Anticancer Therapy</i> , 2022, 22, 3-16.	2.4	9
6	Machine learning for rhabdomyosarcoma histopathology. <i>Modern Pathology</i> , 2022, 35, 1193-1203.	5.5	9
7	Proteomic Profiling Identifies Co-Regulated Expression of Splicing Factors as a Characteristic Feature of Intravenous Leiomyomatosis. <i>Cancers</i> , 2022, 14, 2907.	3.7	2
8	Systemic treatment of advanced clear cell sarcoma: results from a retrospective international series from the World Sarcoma Network. <i>ESMO Open</i> , 2022, 7, 100522.	4.5	11
9	Data-independent acquisition mass spectrometry (DIA-MS) for proteomic applications in oncology. <i>Molecular Omics</i> , 2021, 17, 29-42.	2.8	93
10	Sirolimus for patients with progressive epithelioid hemangioendothelioma. <i>Cancer</i> , 2021, 127, 504-506.	4.1	3
11	3D Functional Genomics Screens Identify CREBBP as a Targetable Driver in Aggressive Triple-Negative Breast Cancer. <i>Cancer Research</i> , 2021, 81, 847-859.	0.9	7
12	Unmet Medical Needs and Future Perspectives for Leiomyosarcoma Patients—A Position Paper from the National Leiomyosarcoma Foundation (NLMSF) and Sarcoma Patients EuroNet (SPAEN). <i>Cancers</i> , 2021, 13, 886.	3.7	17
13	Predictive and prognostic transcriptomic biomarkers in soft tissue sarcomas. <i>Npj Precision Oncology</i> , 2021, 5, 17.	5.4	23
14	Tackling Drug Resistance in EGFR Exon 20 Insertion Mutant Lung Cancer. <i>Pharmacogenomics and Personalized Medicine</i> , 2021, Volume 14, 301-317.	0.7	11
15	Next-generation sequencing for the management of sarcomas with no known driver mutations. <i>Current Opinion in Oncology</i> , 2021, 33, 315-322.	2.4	13
16	Targeting the Fibroblast Growth Factor Receptor (FGFR) Family in Lung Cancer. <i>Cells</i> , 2021, 10, 1154.	4.1	21
17	Clinical management and outcomes of primary ovarian leiomyosarcoma—Experience from a sarcoma specialist unit. <i>Gynecologic Oncology Reports</i> , 2021, 36, 100737.	0.6	9
18	Proteomic profiling of soft tissue sarcomas with SWATH mass spectrometry. <i>Journal of Proteomics</i> , 2021, 241, 104236.	2.4	12

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19	Epithelioid hemangioendothelioma, an ultra-rare cancer: a consensus paper from the community of experts. <i>ESMO Open</i> , 2021, 6, 100170.	4.5	65
20	Fibroblast Growth Factor Receptor (FGFR) Signaling in GIST and Soft Tissue Sarcomas. <i>Cells</i> , 2021, 10, 1533.	4.1	14
21	Pharmacotherapy for liposarcoma: current and emerging synthetic treatments. <i>Future Oncology</i> , 2021, 17, 2659-2670.	2.4	6
22	Solitary fibrous tumor: molecular hallmarks and treatment for a rare sarcoma. <i>Future Oncology</i> , 2021, 17, 3627-3636.	2.4	15
23	Gastrointestinal leiomyosarcoma demonstrate a predilection for distant recurrence and poor response to systemic treatments. <i>European Journal of Surgical Oncology</i> , 2021, 47, 2595-2601.	1.0	3
24	Future Directions in the Treatment of Osteosarcoma. <i>Cells</i> , 2021, 10, 172.	4.1	102
25	Advances in the proteomic profiling of the matrisome and adhesome. <i>Expert Review of Proteomics</i> , 2021, 18, 781-794.	3.0	16
26	The perplexing role of immuno-oncology drugs in osteosarcoma. <i>Journal of Bone Oncology</i> , 2021, 31, 100400.	2.4	4
27	The Extracellular Matrix in Soft Tissue Sarcomas: Pathobiology and Cellular Signalling. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 763640.	3.7	7
28	Rare epidermal growth factor receptor (EGFR) mutations in non-small cell lung cancer. <i>Seminars in Cancer Biology</i> , 2020, 61, 167-179.	9.6	302
29	Proteomic research in sarcomas – current status and future opportunities. <i>Seminars in Cancer Biology</i> , 2020, 61, 56-70.	9.6	50
30	Optimal Clinical Management and the Molecular Biology of Angiosarcomas. <i>Cancers</i> , 2020, 12, 3321.	3.7	15
31	Efficacy of Gemcitabine-based Chemotherapy in Clear Cell Sarcoma of Soft Tissue. <i>Anticancer Research</i> , 2020, 40, 7003-7007.	1.1	3
32	Targeting the Src Pathway Enhances the Efficacy of Selective FGFR Inhibitors in Urothelial Cancers with FGFR3 Alterations. <i>International Journal of Molecular Sciences</i> , 2020, 21, 3214.	4.1	11
33	A mouse SWATH-MS reference spectral library enables deconvolution of species-specific proteomic alterations in human tumour xenografts. <i>DMM Disease Models and Mechanisms</i> , 2020, 13, .	2.4	16
34	Tropomyosin receptor kinase inhibitors in the management of sarcomas. <i>Current Opinion in Oncology</i> , 2020, 32, 307-313.	2.4	9
35	Translational genomics for rare cancers: Challenges and opportunity. <i>Seminars in Cancer Biology</i> , 2020, 61, iii-iv.	9.6	0
36	Is the IDH Mutation a Good Target for Chondrosarcoma Treatment?. <i>Current Molecular Biology Reports</i> , 2020, 6, 1-9.	1.6	20

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37	Avapritinib in the treatment of PDGFRA exon 18 mutated gastrointestinal stromal tumors. <i>Future Oncology</i> , 2020, 16, 1641-1648.	2.4	7
38	The adequacy of tissue microarrays in the assessment of inter- and intra-tumoural heterogeneity of infiltrating lymphocyte burden in leiomyosarcoma. <i>Scientific Reports</i> , 2019, 9, 14602.	3.3	22
39	The landscape of tyrosine kinase inhibitors in sarcomas: looking beyond pazopanib. <i>Expert Review of Anticancer Therapy</i> , 2019, 19, 971-991.	2.4	31
40	Pazopanib in advanced soft tissue sarcomas. <i>Signal Transduction and Targeted Therapy</i> , 2019, 4, 16.	17.1	57
41	Pazopanib in patients with advanced intermediate-grade or high-grade liposarcoma. <i>Expert Opinion on Investigational Drugs</i> , 2019, 28, 505-511.	4.1	13
42	Targeting EGFR exon 20 insertion mutations in non-small cell lung cancer. <i>Signal Transduction and Targeted Therapy</i> , 2019, 4, 5.	17.1	231
43	Negative phase III trials announce the need for biomarkers in sarcoma. <i>European Journal of Cancer</i> , 2019, 123, 81-82.	2.8	1
44	Fibroblastic Reticular Cells Control Conduit Matrix Deposition during Lymph Node Expansion. <i>Cell Reports</i> , 2019, 29, 2810-2822.e5.	6.4	58
45	Ewing-like sarcomas: New molecular diagnoses in need of optimized treatment approaches. <i>Indian Journal of Medical Research</i> , 2019, 150, 521.	1.0	1
46	Olaratumab in soft tissue sarcoma – Current status and future perspectives. <i>European Journal of Cancer</i> , 2018, 92, 33-39.	2.8	16
47	SWATH mass spectrometry as a tool for quantitative profiling of the matrisome. <i>Journal of Proteomics</i> , 2018, 189, 11-22.	2.4	75
48	Quantitative phosphoproteomic analysis of acquired cancer drug resistance to pazopanib and dasatinib. <i>Journal of Proteomics</i> , 2018, 170, 130-140.	2.4	27
49	Clinical and Molecular Spectrum of Liposarcoma. <i>Journal of Clinical Oncology</i> , 2018, 36, 151-159.	1.6	183
50	Spatial localisation of Discoidin Domain Receptor 2 (DDR2) signalling is dependent on its collagen binding and kinase activity. <i>Biochemical and Biophysical Research Communications</i> , 2018, 501, 124-130.	2.1	2
51	Exploiting vulnerabilities in cancer signalling networks to combat targeted therapy resistance. <i>Essays in Biochemistry</i> , 2018, 62, 583-593.	4.7	25
52	Primary Cilia Mediate Diverse Kinase Inhibitor Resistance Mechanisms in Cancer. <i>Cell Reports</i> , 2018, 23, 3042-3055.	6.4	77
53	Novel therapeutic approaches in chondrosarcoma. <i>Future Oncology</i> , 2017, 13, 637-648.	2.4	96
54	Advances in mass spectrometry based strategies to study receptor tyrosine kinases. <i>IUCr</i> , 2017, 4, 119-130.	2.2	13

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55	Exploiting Synthetic Lethality and Network Biology to Overcome EGFR Inhibitor Resistance in Lung Cancer. <i>Journal of Molecular Biology</i> , 2017, 429, 1767-1786.	4.2	14
56	Phase III Soft Tissue Sarcoma Trials: Success or Failure?. <i>Current Treatment Options in Oncology</i> , 2017, 18, 19.	3.0	19
57	Exploiting receptor tyrosine kinase co-activation for cancer therapy. <i>Drug Discovery Today</i> , 2017, 22, 72-84.	6.4	30
58	Targeting SWI/SNF mutant cancers with tyrosine kinase inhibitor therapy. <i>Expert Review of Anticancer Therapy</i> , 2017, 17, 1-3.	2.4	11
59	Analysis of Phosphotyrosine Signaling Networks in Lung Cancer Cell Lines. <i>Methods in Molecular Biology</i> , 2017, 1636, 253-262.	0.9	1
60	Targeted Analysis of Phosphotyrosine Signaling by Multiple Reaction Monitoring Mass Spectrometry. <i>Methods in Molecular Biology</i> , 2017, 1636, 263-281.	0.9	3
61	Progress and impact of clinical phosphoproteomics on precision oncology. <i>Translational Cancer Research</i> , 2017, 6, S1108-S1114.	1.0	1
62	Systematic analysis of tumour cell-extracellular matrix adhesion identifies independent prognostic factors in breast cancer. <i>Oncotarget</i> , 2016, 7, 62939-62953.	1.8	26
63	Expanding the computational toolbox for interrogating cancer kinomes. <i>Pharmacogenomics</i> , 2016, 17, 95-97.	1.3	3
64	Retinoblastoma family proteins: New players in DNA repair by non-homologous end-joining. <i>Molecular and Cellular Oncology</i> , 2016, 3, e1053596.	0.7	5
65	Drug repositioning in sarcomas and other rare tumors. <i>EBioMedicine</i> , 2016, 6, 4-5.	6.1	4
66	Discoidin Domain Receptor Signalling Networks. , 2016, , 201-216.		0
67	Three-dimensional modelling identifies novel genetic dependencies associated with breast cancer progression in the isogenic MCF10 model. <i>Journal of Pathology</i> , 2016, 240, 315-328.	4.5	35
68	Dual Targeting of PDGFR α and FGFR1 Displays Synergistic Efficacy in Malignant Rhabdoid Tumors. <i>Cell Reports</i> , 2016, 17, 1265-1275.	6.4	44
69	Phosphoproteomics in translational research: a sarcoma perspective. <i>Annals of Oncology</i> , 2016, 27, 787-794.	1.2	34
70	Alterations in the phosphoproteomic profile of cells expressing a non-functional form of the SHP2 phosphatase. <i>New Biotechnology</i> , 2016, 33, 524-536.	4.4	7
71	Comparative proteomic assessment of matrisome enrichment methodologies. <i>Biochemical Journal</i> , 2016, 473, 3979-3995.	3.7	41
72	Direct Involvement of Retinoblastoma Family Proteins in DNA Repair by Non-homologous End-Joining. <i>Cell Reports</i> , 2015, 10, 2006-2018.	6.4	62

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73	RB in DNA repair. <i>Oncotarget</i> , 2015, 6, 20746-20747.	1.8	12
74	Glycosylation at Asn211 Regulates the Activation State of the Discoidin Domain Receptor 1 (DDR1). <i>Journal of Biological Chemistry</i> , 2014, 289, 9275-9287.	3.4	33
75	Discoidin domain receptors: a proteomic portrait. <i>Cellular and Molecular Life Sciences</i> , 2014, 71, 3269-3279.	5.4	28
76	Discoidin Domain Receptor 2 Signaling Networks and Therapy in Lung Cancer. <i>Journal of Thoracic Oncology</i> , 2014, 9, 900-904.	1.1	40
77	Discoidin Domain Receptors: Unique Receptor Tyrosine Kinases in Collagen-mediated Signaling. <i>Journal of Biological Chemistry</i> , 2013, 288, 7430-7437.	3.4	182
78	Phosphoproteomics of collagen receptor networks reveals SHP-2 phosphorylation downstream of wild-type DDR2 and its lung cancer mutants. <i>Biochemical Journal</i> , 2013, 454, 501-513.	3.7	68
79	The Pathobiology of Collagens in Glioma. <i>Molecular Cancer Research</i> , 2013, 11, 1129-1140.	3.4	121
80	Phosphoproteomic analysis identifies insulin enhancement of discoidin domain receptor 2 phosphorylation. <i>Cell Adhesion and Migration</i> , 2013, 7, 161-164.	2.7	26
81	Phosphoproteomic studies of receptor tyrosine kinases: future perspectives. <i>Molecular BioSystems</i> , 2012, 8, 1100-1107.	2.9	15
82	Dacomitinib. <i>Drugs of the Future</i> , 2012, 37, 393.	0.1	3
83	Discoidin Domain Receptors Promote $\alpha 1 \beta 1$ - and $\alpha 2 \beta 1$ -Integrin Mediated Cell Adhesion to Collagen by Enhancing Integrin Activation. <i>PLoS ONE</i> , 2012, 7, e52209.	2.5	122
84	A HIF-Regulated VHL-PTP1B-Src Signaling Axis Identifies a Therapeutic Target in Renal Cell Carcinoma. <i>Science Translational Medicine</i> , 2011, 3, 85ra47.	12.4	54
85	EGFRvIV: a previously uncharacterized oncogenic mutant reveals a kinase autoinhibitory mechanism. <i>Oncogene</i> , 2010, 29, 5850-5860.	5.9	58
86	Receptor Tyrosine Kinase Coactivation Networks in Cancer. <i>Cancer Research</i> , 2010, 70, 3857-3860.	0.9	161
87	Phosphotyrosine signaling analysis of site-specific mutations on EGFRvIII identifies determinants governing glioblastoma cell growth. <i>Molecular BioSystems</i> , 2010, 6, 1227.	2.9	40
88	Oncogenic EGFR Signaling Networks in Glioma. <i>Science Signaling</i> , 2009, 2, re6.	3.6	299
89	Melanoma troops massed. <i>Nature</i> , 2009, 459, 336-337.	27.8	11
90	An integrated comparative phosphoproteomic and bioinformatic approach reveals a novel class of MPM-2 motifs upregulated in EGFRvIII-expressing glioblastoma cells. <i>Molecular BioSystems</i> , 2009, 5, 59-67.	2.9	27

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91	Combinatorial Therapeutic Strategies for Blocking Kinase Pathways in Brain Tumors. , 2009, , 953-975.		1
92	Phosphoproteomics: Unraveling the Signaling Web. Molecular Cell, 2008, 31, 777-781.	9.7	50
93	Uncovering Therapeutic Targets FOR Glioblastoma: A Systems Biology Approach. Cell Cycle, 2007, 6, 2750-2754.	2.6	63
94	Quantitative analysis of EGFRvIII cellular signaling networks reveals a combinatorial therapeutic strategy for glioblastoma. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 12867-12872.	7.1	365
95	14-3-3 β controls mitotic translation to facilitate cytokinesis. Nature, 2007, 446, 329-332.	27.8	217
96	Virtual Biopsy in Soft Tissue Sarcoma. How Close Are We?. Frontiers in Oncology, 0, 12, .	2.8	6