

# Shiaw-Yih Lin

## List of Publications by Year in descending order

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Version: 2024-02-01

63  
papers

3,649  
citations

147801

31  
h-index

138484

58  
g-index

65  
all docs

65  
docs citations

65  
times ranked

6314  
citing authors

#	ARTICLE	IF	CITATIONS
1	Proteogenomic Analysis of Salivary Adenoid Cystic Carcinomas Defines Molecular Subtypes and Identifies Therapeutic Targets. <i>Clinical Cancer Research</i> , 2023, 27, 852-864.	7.0	61
2	Combined IL-2, agonistic CD3 and 4-1BB stimulation preserve clonotype hierarchy in propagated non-small cell lung cancer tumor-infiltrating lymphocytes. , 2022, 10, e003082.		11
3	CNGPLD: caseâ€“control copy-number analysis using Gaussian process latent difference. <i>Bioinformatics</i> , 2022, , .	4.1	0
4	Spontaneous tumor regression following COVID-19 vaccination. , 2022, 10, e004371.		26
5	Exploiting induced vulnerability to overcome PARPi resistance and clonal heterogeneity in BRCA mutant triple-negative inflammatory breast cancer.. <i>American Journal of Cancer Research</i> , 2022, 12, 337-354.	1.4	0
6	Mechanisms of immunogenic cell death and immune checkpoint blockade therapy. <i>Kaohsiung Journal of Medical Sciences</i> , 2021, 37, 448-458.	1.9	15
7	A Gene Expression Signature to Predict Nucleotide Excision Repair Defects and Novel Therapeutic Approaches. <i>International Journal of Molecular Sciences</i> , 2021, 22, 5008.	4.1	3
8	Combined Inhibition of Rad51 and Wee1 Enhances Cell Killing in HNSCC Through Induction of Apoptosis Associated With Excessive DNA Damage and Replication Stress. <i>Molecular Cancer Therapeutics</i> , 2021, 20, 1257-1269.	4.1	15
9	Replication stress response defects are associated with response to immune checkpoint blockade in nonhypermuted cancers. <i>Science Translational Medicine</i> , 2021, 13, eabe6201.	12.4	19
10	Integrated Genomic Characterization of the Human Immunome in Cancer. <i>Cancer Research</i> , 2020, 80, 4854-4867.	0.9	11
11	PBRM1 loss defines a nonimmunogenic tumor phenotype associated with checkpoint inhibitor resistance in renal carcinoma. <i>Nature Communications</i> , 2020, 11, 2135.	12.8	114
12	Nucleostemin Modulates Outcomes of Hepatocellular Carcinoma via a Tumor Adaptive Mechanism to Genomic Stress. <i>Molecular Cancer Research</i> , 2020, 18, molcanres.0777.2019.	3.4	8
13	Role of DNA repair defects in predicting immunotherapy response. <i>Biomarker Research</i> , 2020, 8, 23.	6.8	47
14	Proteome Instability Is a Therapeutic Vulnerability in Mismatch Repair-Deficient Cancer. <i>Cancer Cell</i> , 2020, 37, 371-386.e12.	16.8	68
15	BRIT1 dysfunction confers synergistic inhibition of hepatocellular carcinoma by targeting poly (ADP-ribose) polymerases and PI3K. <i>American Journal of Cancer Research</i> , 2020, 10, 1900-1918.	1.4	2
16	Genetic alterations and expression characteristics of ARID1A impact tumor immune contexture and survival in early-onset gastric cancer. <i>American Journal of Cancer Research</i> , 2020, 10, 3947-3972.	1.4	3
17	Nucleostemin reveals a dichotomous nature of genome maintenance in mammary tumor progression. <i>Oncogene</i> , 2019, 38, 3919-3931.	5.9	11
18	Sequential Therapy with PARP and WEE1 Inhibitors Minimizes Toxicity while Maintaining Efficacy. <i>Cancer Cell</i> , 2019, 35, 851-867.e7.	16.8	156

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19	PARP inhibitors synergize with gemcitabine by potentiating DNA damage in non-small cell lung cancer. <i>International Journal of Cancer</i> , 2019, 144, 1092-1103.	5.1	38
20	The Tale of CHD4 in DNA Damage Response and Chemotherapeutic Response. <i>Cancer Research and Cellular Therapeutics</i> , 2019, 3, .	0.0	3
21	Multi-omics analysis reveals neoantigen-independent immune cell infiltration in copy-number driven cancers. <i>Nature Communications</i> , 2018, 9, 1317.	12.8	94
22	BRD4 Inhibition Is Synthetic Lethal with PARP Inhibitors through the Induction of Homologous Recombination Deficiency. <i>Cancer Cell</i> , 2018, 33, 401-416.e8.	16.8	215
23	Defective Replication Stress Response Is Inherently Linked to the Cancer Stem Cell Phenotype. <i>Cell Reports</i> , 2018, 23, 2095-2106.	6.4	37
24	CHD4 mutations promote endometrial cancer stemness by activating TGF-beta signaling. <i>American Journal of Cancer Research</i> , 2018, 8, 903-914.	1.4	6
25	A murine preclinical syngeneic transplantation model for breast cancer precision medicine. <i>Science Advances</i> , 2017, 3, e1600957.	10.3	10
26	Rational combination therapy with PARP and MEK inhibitors capitalizes on therapeutic liabilities in <i>RAS</i> mutant cancers. <i>Science Translational Medicine</i> , 2017, 9, .	12.4	174
27	Improved prediction of PARP inhibitor response and identification of synergizing agents through use of a novel gene expression signature generation algorithm. <i>Npj Systems Biology and Applications</i> , 2017, 3, 8.	3.0	55
28	MicroPET/CT Imaging of AXL Downregulation by HSP90 Inhibition in Triple-Negative Breast Cancer. <i>Contrast Media and Molecular Imaging</i> , 2017, 2017, 1-11.	0.8	9
29	The role of Rak in the regulation of stability and function of BRCA1. <i>Oncotarget</i> , 2017, 8, 86799-86815.	1.8	9
30	BRIT1 Gene. , 2017, , 699-702.		0
31	Connecting the Dots: From DNA Damage and Repair to Aging. <i>International Journal of Molecular Sciences</i> , 2016, 17, 685.	4.1	53
32	mTOR Inhibitors Suppress Homologous Recombination Repair and Synergize with PARP Inhibitors via Regulating SUV39H1 in BRCA-Proficient Triple-Negative Breast Cancer. <i>Clinical Cancer Research</i> , 2016, 22, 1699-1712.	7.0	95
33	Genomic-Glycosylation Aberrations in Tumor Initiation, Progression and Management. <i>AIMS Medical Science</i> , 2016, 3, 386-416.	0.4	3
34	Local generation of fumarate promotes DNA repair through inhibition of histone H3 demethylation. <i>Nature Cell Biology</i> , 2015, 17, 1158-1168.	10.3	154
35	Nuclear PTEN tumor-suppressor functions through maintaining heterochromatin structure. <i>Cell Cycle</i> , 2015, 14, 2323-2332.	2.6	38
36	TUSC4 Functions as a Tumor Suppressor by Regulating BRCA1 Stability. <i>Cancer Research</i> , 2015, 75, 378-386.	0.9	24

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37	New insights into tumor dormancy: Targeting DNA repair pathways. <i>World Journal of Clinical Oncology</i> , 2015, 6, 80.	2.3	15
38	BRIT1 Gene. , 2015, , 1-4.		0
39	Genome-wide transcriptome profiling of homologous recombination DNA repair. <i>Nature Communications</i> , 2014, 5, 3361.	12.8	182
40	Tumor dormancy: potential therapeutic target in tumor recurrence and metastasis prevention. <i>Experimental Hematology and Oncology</i> , 2013, 2, 29.	5.0	40
41	Zinc finger protein 668 interacts with Tip60 to promote H2AX acetylation after DNA damage. <i>Cell Cycle</i> , 2013, 12, 2033-2041.	2.6	19
42	BRIT1 regulates p53 stability and functions as a tumor suppressor in breast cancer. <i>Carcinogenesis</i> , 2013, 34, 2271-2280.	2.8	13
43	Chromodomain Helicase DNA-binding Protein 4 (CHD4) Regulates Homologous Recombination DNA Repair, and Its Deficiency Sensitizes Cells to Poly(ADP-ribose) Polymerase (PARP) Inhibitor Treatment. <i>Journal of Biological Chemistry</i> , 2012, 287, 6764-6772.	3.4	85
44	DNA damage and breast cancer. <i>World Journal of Clinical Oncology</i> , 2011, 2, 329.	2.3	69
45	DNA Damage Response Is Suppressed by the High Cyclin-dependent Kinase 1 Activity in Mitotic Mammalian Cells. <i>Journal of Biological Chemistry</i> , 2011, 286, 35899-35905.	3.4	31
46	<i>ZNF668</i> Functions as a Tumor Suppressor by Regulating p53 Stability and Function in Breast Cancer. <i>Cancer Research</i> , 2011, 71, 6524-6534.	0.9	26
47	Exploiting the homologous recombination DNA repair network for targeted cancer therapy. <i>World Journal of Clinical Oncology</i> , 2011, 2, 73.	2.3	38
48	BRIT1 Gene. , 2011, , 567-570.		0
49	Multiple Roles of BRIT1/MCPH1 in DNA Damage Response, DNA Repair, and Cancer Suppression. <i>Yonsei Medical Journal</i> , 2010, 51, 295.	2.2	32
50	BRIT1/MCPH1 Is Essential for Mitotic and Meiotic Recombination DNA Repair and Maintaining Genomic Stability in Mice. <i>PLoS Genetics</i> , 2010, 6, e1000826.	3.5	86
51	The DNA damage response: Balancing the scale between cancer and ageing. <i>Aging</i> , 2010, 2, 900-907.	3.1	52
52	Exploring Rak tyrosine kinase function in breast cancer. <i>Cell Cycle</i> , 2009, 8, 2360-2364.	2.6	20
53	BRIT1/MCPH1 is a multifunctional DNA damage responsive protein mediating DNA repair-associated chromatin remodeling. <i>Cell Cycle</i> , 2009, 8, 3071-3072.	2.6	12
54	The linkage of chromatin remodeling to genome maintenance: Contribution from a human disease gene BRIT1/MCPH1. <i>Epigenetics</i> , 2009, 4, 457-461.	2.7	7

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55	Rak Functions as a Tumor Suppressor by Regulating PTEN Protein Stability and Function. <i>Cancer Cell</i> , 2009, 15, 304-314.	16.8	175
56	DNA Damage Response Pathways in Tumor Suppression and Cancer Treatment. <i>World Journal of Surgery</i> , 2009, 33, 661-666.	1.6	77
57	BRIT1/MCPH1 links chromatin remodelling to DNA damage response. <i>Nature Cell Biology</i> , 2009, 11, 865-872.	10.3	175
58	TRF2 functions as a protein hub and regulates telomere maintenance by recognizing specific peptide motifs. <i>Nature Structural and Molecular Biology</i> , 2009, 16, 372-379.	8.2	118
59	Differential regulation of centrosome integrity by DNA damage response proteins. <i>Cell Cycle</i> , 2008, 7, 2225-2233.	2.6	52
60	BRIT1 regulates early DNA damage response, chromosomal integrity, and cancer. <i>Cancer Cell</i> , 2006, 10, 145-157.	16.8	137
61	BRIT1/MCPH1: A Guardian of Genome and an Enemy of Tumors. <i>Cell Cycle</i> , 2006, 5, 2579-2583.	2.6	35
62	BRIT1/MCPH1 is a DNA damage responsive protein that regulates the Brca1-Chk1 pathway, implicating checkpoint dysfunction in microcephaly. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 15105-15109.	7.1	160
63	Multiple Tumor Suppressor Pathways Negatively Regulate Telomerase. <i>Cell</i> , 2003, 113, 881-889.	28.9	400