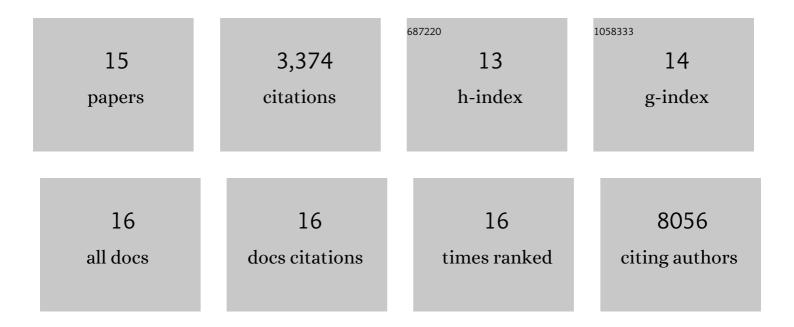
Yuen-Yi Tseng

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Dependency of a therapy-resistant state of cancer cells on a lipid peroxidase pathway. Nature, 2017, 547, 453-457.	13.7	1,194
2	PVT1 dependence in cancer with MYC copy-number increase. Nature, 2014, 512, 82-86.	13.7	617
3	Patient-derived xenografts undergo mouse-specific tumor evolution. Nature Genetics, 2017, 49, 1567-1575.	9.4	546
4	Genome-scale activation screen identifies a IncRNA locus regulating a gene neighbourhood. Nature, 2017, 548, 343-346.	13.7	336
5	WRN helicase is a synthetic lethal target in microsatellite unstable cancers. Nature, 2019, 568, 551-556.	13.7	253
6	Whole Exome Sequencing Identifies TSC1/TSC2 Biallelic Loss as the Primary and Sufficient Driver Event for Renal Angiomyolipoma Development. PLoS Genetics, 2016, 12, e1006242.	1.5	93
7	Assessment of heavy metal bioavailability in contaminated sediments and soils using green fluorescent protein-based bacterial biosensors. Environmental Pollution, 2006, 142, 17-23.	3.7	87
8	A Combination CDK4/6 and IGF1R Inhibitor Strategy for Ewing Sarcoma. Clinical Cancer Research, 2019, 25, 1343-1357.	3.2	69
9	Activation of endothelial cells to pathological status by down-regulation of connexin43. Cardiovascular Research, 2008, 79, 509-518.	1.8	54
10	The PVT1-MYC duet in cancer. Molecular and Cellular Oncology, 2015, 2, e974467.	0.3	44
11	Caenorhabditis elegans expresses a functional ArsA. FEBS Journal, 2007, 274, 2566-2572.	2.2	27
12	<i>MYC</i> and <i>PVT1</i> synergize to regulate RSPO1 levels in breast cancer. Cell Cycle, 2016, 15, 881-885.	1.3	27
13	From cell lines to living biosensors: new opportunities to prioritize cancer dependencies using ex vivo tumor cultures. Current Opinion in Genetics and Development, 2019, 54, 33-40.	1.5	20
14	C-reactive protein, sodium azide, and endothelial connexin43 gap junctions. Cell Biology and Toxicology, 2010, 26, 153-163.	2.4	2
15	Engineering Large Genomic Rearrangement in Mouse Embryonic Stem Cell for Cancer Gene Discovery. Methods in Molecular Biology, 2019, 1907, 197-212.	0.4	0