## Zongli Zheng

List of Publications by Year in descending order

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52 12,256 29 51 papers citations h-index g-index

54 54 54 18996
all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	Deep RNA Sequencing Revealed Fusion Junctional Heterogeneity May Predict Crizotinib Treatment Efficacy in ALK-Rearranged NSCLC. Journal of Thoracic Oncology, 2022, 17, 264-276.	1.1	15
2	High-fidelity KKH variant of <i>Staphylococcus aureus</i> Cas9 nucleases with improved base mismatch discrimination. Nucleic Acids Research, 2022, 50, 1650-1660.	14.5	11
3	Machine learning-coupled combinatorial mutagenesis enables resource-efficient engineering of CRISPR-Cas9 genome editor activities. Nature Communications, 2022, 13, 2219.	12.8	8
4	Noncanonical Gene Fusions Detected at the DNA Level Necessitate Orthogonal Diagnosis Methods Before Targeted Therapy. Journal of Thoracic Oncology, 2021, 16, 344-348.	1.1	6
5	Evaluation of endocrine resistance using ESR1 genotyping of circulating tumor cells and plasma DNA. Breast Cancer Research and Treatment, 2021, 188, 43-52.	2.5	8
6	Defining genome-wide CRISPR–Cas genome-editing nuclease activity with GUIDE-seq. Nature Protocols, 2021, 16, 5592-5615.	12.0	27
7	A Three-Way Combinatorial CRISPR Screen for Analyzing Interactions among Druggable Targets. Cell Reports, 2020, 32, 108020.	6.4	27
8	Combinatorial mutagenesis en masse optimizes the genome editing activities of SpCas9. Nature Methods, 2019, 16, 722-730.	19.0	44
9	Rationally engineered <i>Staphylococcus aureus</i> Cas9 nucleases with high genome-wide specificity. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 20969-20976.	7.1	81
10	Bone Sarcoma With <i>EWSR1-NFATC2</i> Fusion: Sarcoma With Varied Morphology and Amplification of Fusion Gene Distinct From Ewing Sarcoma. International Journal of Surgical Pathology, 2019, 27, 561-567.	0.8	17
11	Highly Multiplexed Fluorescence in Situ Hybridization for in Situ Genomics. Journal of Molecular Diagnostics, 2019, 21, 390-407.	2.8	15
12	Expressed Gene Fusions as Frequent Drivers of Poor Outcomes in Hormone Receptor–Positive Breast Cancer. Cancer Discovery, 2018, 8, 336-353.	9.4	32
13	Artificial Intelligence Approach for Variant Reporting. JCO Clinical Cancer Informatics, 2018, 2, 1-13.	2.1	13
14	Clinical and radiographic response following targeting of BCAN-NTRK1 fusion in glioneuronal tumor. Npj Precision Oncology, 2017, 1, 5.	5.4	49
15	Recurrent and functional regulatory mutations in breast cancer. Nature, 2017, 547, 55-60.	27.8	269
16	<i>MET</i> Exon 14 Skipping in Non-Small Cell Lung Cancer. Oncologist, 2016, 21, 481-486.	3.7	94
17	Next-Generation Sequencing and Fluorescence in Situ Hybridization Have Comparable Performance Characteristics in the Analysis of Pancreaticobiliary Brushings for Malignancy. Journal of Molecular Diagnostics, 2016, 18, 124-130.	2.8	79
18	Severity of Acute Cholecystitis and Risk of latrogenic Bile Duct Injury During Cholecystectomy, a Populationâ€Based Case–Control Study. World Journal of Surgery, 2016, 40, 1060-1067.	1.6	81

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19	High-fidelity CRISPR–Cas9 nucleases with no detectable genome-wide off-target effects. Nature, 2016, 529, 490-495.	27.8	2,126
20	Impact of next-generation sequencing on the clinical diagnosis of pancreatic cysts. Gastrointestinal Endoscopy, 2016, 83, 140-148.	1.0	119
21	Unique Genetic and Survival Characteristics of Invasive Mucinous Adenocarcinoma of the Lung. Journal of Thoracic Oncology, 2015, 10, 1156-1162.	1.1	137
22	Durable Clinical Response to Entrectinib in NTRK1-Rearranged Non-Small Cell Lung Cancer. Journal of Thoracic Oncology, 2015, 10, 1670-1674.	1.1	197
23	Engineered CRISPR-Cas9 nucleases with altered PAM specificities. Nature, 2015, 523, 481-485.	27.8	1,388
24	Broadening the targeting range of Staphylococcus aureus CRISPR-Cas9 by modifying PAM recognition. Nature Biotechnology, 2015, 33, 1293-1298.	17.5	511
25	Identification of insertions in PTEN and TP53 in anaplastic thyroid carcinoma with angiogenic brain metastasis. Endocrine-Related Cancer, 2015, 22, L23-L28.	3.1	5
26	Incidence of gastric cancer among patients with gastric precancerous lesions: observational cohort study in a low risk Western population. BMJ, The, 2015, 351, h3867.	6.0	198
27	Detection of Dual IDH1 and IDH2 Mutations by Targeted Next-Generation Sequencing in Acute Myeloid Leukemia and Myelodysplastic Syndromes. Journal of Molecular Diagnostics, 2015, 17, 661-668.	2.8	31
28	Variant Profiling of Candidate Genes in Pancreatic Ductal Adenocarcinoma. Clinical Chemistry, 2015, 61, 1408-1416.	3.2	21
29	GUIDE-seq enables genome-wide profiling of off-target cleavage by CRISPR-Cas nucleases. Nature Biotechnology, 2015, 33, 187-197.	17.5	1,757
30	High p53 protein expression in therapy-related myeloid neoplasms is associated with adverse karyotype and poor outcome. Modern Pathology, 2015, 28, 552-563.	5.5	42
31	Clinical implementation of anchored multiplex PCR with targeted next-generation sequencing for detection of ALK, ROS1, RET and NTRK1 fusions in non-small cell lung carcinoma Journal of Clinical Oncology, 2015, 33, 8095-8095.	1.6	1
32	LIN28 Is Involved in Glioma Carcinogenesis and Predicts Outcomes of Glioblastoma Multiforme Patients. PLoS ONE, 2014, 9, e86446.	2.5	31
33	Identification of Oncogenic Mutations and Gene Fusions in the Follicular Variant of Papillary Thyroid Carcinoma. Journal of Clinical Endocrinology and Metabolism, 2014, 99, E2457-E2462.	3.6	55
34	Anchored multiplex PCR for targeted next-generation sequencing. Nature Medicine, 2014, 20, 1479-1484.	30.7	705
35	Crizotinib in <i>ROS1</i> -Rearranged Non–Small-Cell Lung Cancer. New England Journal of Medicine, 2014, 371, 1963-1971.	27.0	1,656
36	Ex vivo culture of circulating breast tumor cells for individualized testing of drug susceptibility. Science, 2014, 345, 216-220.	12.6	808

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37	A Novel Fusion of TPR and ALK in Lung Adenocarcinoma. Journal of Thoracic Oncology, 2014, 9, 563-566.	1.1	83
38	Metagenomic <i>De Novo</i> Assembly of an Aquatic Representative of the Verrucomicrobial Class <i>Spartobacteria</i> MBio, 2013, 4, e00569-12.	4.1	107
39	Titration-free 454 sequencing using Y adapters. Nature Protocols, 2011, 6, 1367-1376.	12.0	24
40	Rapid Screening of Complex DNA Samples by Single-Molecule Amplification and Sequencing. PLoS ONE, 2011, 6, e19723.	2.5	2
41	A Method for Metagenomics of Helicobacter pylori from Archived Formalin-Fixed Gastric Biopsies Permitting Longitudinal Studies of Carcinogenic Risk. PLoS ONE, 2011, 6, e26442.	2.5	14
42	A comprehensive analysis of common genetic variation in MUC1, MUC5AC, MUC6 genes and risk of stomach cancer. Cancer Causes and Control, 2010, 21, 313-321.	1.8	76
43	A Pyrosequencing Study in Twins Shows That Gastrointestinal Microbial Profiles Vary With Inflammatory Bowel Disease Phenotypes. Gastroenterology, 2010, 139, 1844-1854.e1.	1.3	916
44	Titration-free massively parallel pyrosequencing using trace amounts of starting material. Nucleic Acids Research, 2010, 38, e137-e137.	14.5	28
45	Metagenomic study of Helicobacter pylori microdissected from archived formalin-fixed paraffin-embedded biopsy sections. Genome Biology, 2010, 11, P42.	9.6	0
46	Is There a Link between the Lipopolysaccharide of Helicobacter pylori Gastric MALT Lymphoma Associated Strains and Lymphoma Pathogenesis?. PLoS ONE, 2009, 4, e7297.	2.5	12
47	Genetic Variation in <i>a4GnT</i> in Relation to <i>HelicobacterÂpylori</i> Serology and Gastric Cancer Risk. Helicobacter, 2009, 14, 472-477.	3 <b>.</b> 5	8
48	Long-Term Effects of latrogenic Bile Duct Injury During Cholecystectomy. Clinical Gastroenterology and Hepatology, 2009, 7, 1013-1018.	4.4	54
49	Postmenopausal Hormone Therapy as a Risk Factor for Gastroesophageal Reflux Symptoms Among Female Twins. Gastroenterology, 2008, 134, 921-928.	1.3	47
50	Effects of Estrogen With and Without Progestin and Obesity on Symptomatic Gastroesophageal Reflux. Gastroenterology, 2008, 135, 72-81.	1.3	24
51	Lifestyle Factors and Risk for Symptomatic Gastroesophageal Reflux in Monozygotic Twins. Gastroenterology, 2007, 132, 87-95.	1.3	139
52	Risk factors for the gastric cardia cancer: a case-control study in Fujian Province. World Journal of Gastroenterology, 2003, 9, 214.	3.3	58