Shuwei Li

List of Publications by Year in descending order

19 papers	1,135 citations	840776 11 h-index	940533 16 g-index
19 all docs	19 docs citations	19 times ranked	3047 citing authors

#	Article	IF	Citations
1	Validation of a prostate cancer polygenic risk score. Prostate, 2020, 80, 1314-1321.	2.3	23
2	Rare Germline Pathogenic Mutations of DNA Repair Genes Are Most Strongly Associated with Grade Group 5 Prostate Cancer. European Urology Oncology, 2020, 3, 224-230.	5.4	41
3	Tumour characteristics provide evidence for germline mismatch repair missense variant pathogenicity. Journal of Medical Genetics, 2020, 57, 62-69.	3.2	11
4	Prevalence of Germline Pathogenic and Likely Pathogenic Variants in Patients With Second Breast Cancers. JNCI Cancer Spectrum, 2020, 4, pkaa094.	2.9	10
5	Association of Breast and Ovarian Cancers With Predisposition Genes Identified by Large-Scale Sequencing. JAMA Oncology, 2019, 5, 51.	7.1	145
6	Clinical features and cancer risk in families with pathogenic CDH1 variants irrespective of clinical criteria. Journal of Medical Genetics, 2019, 56, 838-843.	3.2	84
7	Clinical germline diagnostic exome sequencing for hereditary cancer: Findings within novel candidate genes are prevalent. Cancer Genetics, 2018, 224-225, 12-20.	0.4	7
8	A Bayesian framework for efficient and accurate variant prediction. PLoS ONE, 2018, 13, e0203553.	2.5	12
9	Male breast cancer in a multi-gene panel testing cohort: insights and unexpected results. Breast Cancer Research and Treatment, 2017, 161, 575-586.	2.5	116
10	Candidate-gene criteria for clinical reporting: diagnostic exome sequencing identifies altered candidate genes among 8% of patients with undiagnosed diseases. Genetics in Medicine, 2017, 19, 224-235.	2.4	47
11	Multigene Panel Testing Provides a New Perspective on Lynch Syndrome. Journal of Clinical Oncology, 2017, 35, 2568-2575.	1.6	122
12	Diagnosing Hereditary Cancer Susceptibility Through Multigene Panel Testing. , 2017, , 123-153.		0
13	Sanger Confirmation Is Required to Achieve Optimal Sensitivity and Specificity in Next-Generation Sequencing Panel Testing. Journal of Molecular Diagnostics, 2016, 18, 923-932.	2.8	143
14	Breast cancer risk is similar for CHEK2 founder and non-founder mutation carriers. Cancer Genetics, 2016, 209, 403-407.	0.4	41
15	Diagnostic exome sequencing provides a molecular diagnosis for a significant proportion of patients with epilepsy. Genetics in Medicine, 2016, 18, 898-905.	2.4	299
16	Abstract P4-12-16: Teasing out the PALB2 phenotype. , 2015, , .		0
17	Women with breast and uterine cancer in relation to genetic mutation risk: A case-control analysis.. Journal of Clinical Oncology, 2015, 33, 1549-1549.	1.6	0
18	Functional and evolutionary correlates of gene constellations in the Drosophila melanogaster genome that deviate from the stereotypical gene architecture. BMC Genomics, 2010, 11, 322.	2.8	1

#	Article	IF	Citations
19	A hyperactive quantitative trait locus allele of Arabidopsis BRX contributes to natural variation in root growth vigor. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 8475-8480.	7.1	33