

# Longli Kang

## List of Publications by Year in descending order

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48  
papers

1,063  
citations

623734

14  
h-index

477307

29  
g-index

48  
all docs

48  
docs citations

48  
times ranked

1506  
citing authors

#	ARTICLE	IF	CITATIONS
1	Targeted Sequencing Identifies the Genetic Variants Associated with High-altitude Polycythemia in the Tibetan Population. <i>Indian Journal of Hematology and Blood Transfusion</i> , 2022, 38, 556-565.	0.6	4
2	Genomic insights into the formation of human populations in East Asia. <i>Nature</i> , 2021, 591, 413-419.	27.8	216
3	Using Composite Phenotypes to Reveal Hidden Physiological Heterogeneity in High-Altitude Acclimatization in a Chinese Han Longitudinal Cohort. <i>Phenomics</i> , 2021, 1, 3-14.	2.9	10
4	Effects of altitude on human oral microbes. <i>AMB Express</i> , 2021, 11, 41.	3.0	15
5	Effects of <i>Helicobacter pylori</i> Infection on the Oral Microbiota of Reflux Esophagitis Patients. <i>Frontiers in Cellular and Infection Microbiology</i> , 2021, 11, 732613.	3.9	8
6	Migration effects on the intestinal microbiota of Tibetans. <i>PeerJ</i> , 2021, 9, e12036.	2.0	4
7	Smaller reaction volume of triplex taqman real-time reverse transcription-PCR assays for diagnosing coronavirus disease 2019. <i>Journal of Clinical Laboratory Analysis</i> , 2021, , e24137.	2.1	2
8	The distinctive geographic patterns of common pigmentation variants at the OCA2 gene. <i>Scientific Reports</i> , 2020, 10, 15433.	3.3	8
9	Prioritizing natural-selection signals from the deep-sequencing genomic data suggests multi-variant adaptation in Tibetan highlanders. <i>National Science Review</i> , 2019, 6, 1201-1222.	9.5	30
10	Fluorescent probe for Cu <sup>2+</sup> and the secondary application of the resultant complex to detect cysteine. <i>RSC Advances</i> , 2019, 9, 16812-16818.	3.6	22
11	The massive assimilation of indigenous East Asian populations in the origin of Muslim Hui people inferred from paternal Y chromosome. <i>American Journal of Physical Anthropology</i> , 2019, 169, 341-347.	2.1	16
12	Genome-Wide Association Study Identifies a New Locus at 7q21.13 Associated with Hepatitis B Virus-Related Hepatocellular Carcinoma. <i>Clinical Cancer Research</i> , 2018, 24, 906-915.	7.0	37
13	Associations of high-altitude polycythemia with polymorphisms in PIK3CD and COL4A3 in Tibetan populations. <i>Human Genomics</i> , 2018, 12, 37.	2.9	16
14	Physiological, hematological and biochemical factors associated with high-altitude headache in young Chinese males following acute exposure at 3700m. <i>Journal of Headache and Pain</i> , 2018, 19, 59.	6.0	15
15	Genetic polymorphisms of the drug-metabolizing enzyme cytochrome P450 2A6 in a Tibetan Chinese population. <i>International Journal of Clinical and Experimental Pathology</i> , 2018, 11, 5024-5033.	0.5	0
16	Association between IL-1 gene polymorphisms and tuberculosis susceptibility in the Chinese Tibetan population. <i>International Journal of Clinical and Experimental Pathology</i> , 2018, 11, 5441-5449.	0.5	2
17	Increasing the reference populations for the 55 AISNP panel: the need and benefits. <i>International Journal of Legal Medicine</i> , 2017, 131, 913-917.	2.2	38
18	Investigation of the major cytochrome P450 1A2 genetic variant in a healthy Tibetan population in China. <i>Molecular Medicine Reports</i> , 2017, 16, 573-580.	2.4	5

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19	Differentiated demographic histories and local adaptations between Sherpas and Tibetans. <i>Genome Biology</i> , 2017, 18, 115.	8.8	67
20	Association between single nucleotide polymorphisms in <i>ADRB2</i> , <i>GNB3</i> and <i>GSTP1</i> genes and high-altitude pulmonary edema (HAPE) in the Chinese Han population. <i>Oncotarget</i> , 2017, 8, 18206-18212.	1.8	10
21	<i>TCF7L2</i> polymorphisms and the risk of schizophrenia in the Chinese Han population. <i>Oncotarget</i> , 2017, 8, 28614-28620.	1.8	10
22	Genome-wide association study of high-altitude pulmonary edema in a Han Chinese population. <i>Oncotarget</i> , 2017, 8, 31568-31580.	1.8	7
23	Associations of high altitude polycythemia with polymorphisms in <i>EPHA2</i> and <i>AGT</i> in Chinese Han and Tibetan populations. <i>Oncotarget</i> , 2017, 8, 53234-53243.	1.8	14
24	Associations between polymorphisms in the <i>IL-4</i> gene and renal cell carcinoma in Chinese Han population. <i>Oncotarget</i> , 2017, 8, 82078-82084.	1.8	3
25	Associations of high altitude polycythemia with polymorphisms in <i>EPAS1</i> , <i>ITGA6</i> and <i>ERBB4</i> in Chinese Han and Tibetan populations. <i>Oncotarget</i> , 2017, 8, 86736-86746.	1.8	13
26	Angiotensin II receptor 1 gene variants are associated with high-altitude pulmonary edema risk. <i>Oncotarget</i> , 2016, 7, 77117-77123.	1.8	8
27	Vitamin D receptor gene associations with pulmonary tuberculosis in a Tibetan Chinese population. <i>BMC Infectious Diseases</i> , 2016, 16, 469.	2.9	20
28	MtDNA analysis reveals enriched pathogenic mutations in Tibetan highlanders. <i>Scientific Reports</i> , 2016, 6, 31083.	3.3	22
29	<i>P2X7R</i> Gene Polymorphisms are Associated with Increased Risk of Pulmonary Tuberculosis in the Tibetan Chinese Population. <i>American Journal of Tropical Medicine and Hygiene</i> , 2016, 95, 1016-1020.	1.4	9
30	Ancestral Origins and Genetic History of Tibetan Highlanders. <i>American Journal of Human Genetics</i> , 2016, 99, 580-594.	6.2	208
31	Telomere length-related gene <i>ACYP2</i> polymorphism is associated with the risk of HAPE in Chinese Han population. <i>Journal of Gene Medicine</i> , 2016, 18, 244-249.	2.8	18
32	Genetic polymorphisms study of pharmacogenomic VIP variants in Han ethnic of China's Shaanxi province. <i>Environmental Toxicology and Pharmacology</i> , 2016, 46, 27-35.	4.0	8
33	Genetic polymorphisms of pharmacogenomic VIP variants in the Mongol of Northwestern China. <i>BMC Genetics</i> , 2016, 17, 70.	2.7	6
34	<i>WDR1</i> and <i>CLNK</i> gene polymorphisms correlate with serum glucose and high-density lipoprotein levels in Tibetan gout patients. <i>Rheumatology International</i> , 2016, 36, 405-412.	3.0	5
35	Genotype-phenotype analysis of <i>CYP2C19</i> in the Tibetan population and its potential clinical implications in drug therapy. <i>Molecular Medicine Reports</i> , 2016, 13, 2117-2123.	2.4	12
36	<i>CLPTM1L</i> polymorphism as a protective factor for lung cancer: a case-control study in southern Chinese population. <i>Tumor Biology</i> , 2016, 37, 10533-10538.	1.8	6

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37	The population genetics of pharmacogenomics VIP variants in the Sherpa population. <i>Drug Metabolism and Pharmacokinetics</i> , 2016, 31, 82-89.	2.2	8
38	Genetic polymorphisms analysis of drug-metabolizing enzyme CYP2C9 in the Uyghur population. <i>Xenobiotica</i> , 2016, 46, 709-714.	1.1	6
39	Genetic polymorphism analysis of the drug-metabolizing enzyme CYP1A2 in a Uyghur Chinese population: a pilot study. <i>Xenobiotica</i> , 2016, 46, 542-547.	1.1	2
40	Genetic variation in the ABCG2 gene is associated with gout risk in the Chinese Han population. <i>Clinical Rheumatology</i> , 2016, 35, 159-163.	2.2	18
41	Identification of a shared protective genetic susceptibility locus for colorectal cancer and gastric cancer. <i>Tumor Biology</i> , 2016, 37, 2443-2448.	1.8	2
42	The relationship between polymorphisms of <i>XRCC5</i> genes with astrocytoma prognosis in the Han Chinese population. <i>Oncotarget</i> , 2016, 7, 85283-85290.	1.8	8
43	<i>TERT</i> rs2853676 polymorphisms correlate with glioma prognosis in Chinese population. <i>Oncotarget</i> , 2016, 7, 73781-73791.	1.8	3
44	A 3.4-kb Copy-Number Deletion near EPAS1 Is Significantly Enriched in High-Altitude Tibetans but Absent from the Denisovan Sequence. <i>American Journal of Human Genetics</i> , 2015, 97, 54-66.	6.2	69
45	Genetic polymorphism analysis of the drug-metabolizing enzyme CYP2C9 in a Chinese Tibetan population. <i>Gene</i> , 2015, 567, 196-200.	2.2	4
46	Genetic Polymorphisms Analysis of Pharmacogenomic VIP Variants in Miao Ethnic Group of Southwest China. <i>Medical Science Monitor</i> , 2015, 21, 3769-3776.	1.1	11
47	FGFR2 gene polymorphisms are associated with breast cancer risk in the Han Chinese population. <i>American Journal of Cancer Research</i> , 2015, 5, 1854-61.	1.4	23
48	Genetic polymorphisms of pharmacogenomic VIP variants in the lhoba population of southwest China. <i>International Journal of Clinical and Experimental Pathology</i> , 2015, 8, 13293-303.	0.5	15