

Li-Li Tang

List of Publications by Year in descending order

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Version: 2024-02-01

23
papers

424
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1307594

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#	ARTICLE	IF	CITATIONS
1	Chromatin accessibility and transcriptome integrative analysis revealed AP-1-mediated genes potentially modulate histopathology features in psoriasis. <i>Clinical Epigenetics</i> , 2022, 14, 38.	4.1	6
2	Case Report: Chanarin-Dorfman Syndrome: A Novel Homozygous Mutation in ABHD5 Gene in a Chinese Case and Genotype-Phenotype Correlation Analysis. <i>Frontiers in Genetics</i> , 2022, 13, 847321.	2.3	2
3	<i>CYP2S1</i> might regulate proliferation and immune response of keratinocyte in psoriasis. <i>Epigenetics</i> , 2021, 16, 618-628.	2.7	7
4	Assay for Transposase-Accessible Chromatin Using Sequencing Analysis Reveals a Widespread Increase in Chromatin Accessibility in Psoriasis. <i>Journal of Investigative Dermatology</i> , 2021, 141, 1745-1753.	0.7	8
5	A novel mutation of COL7A1 in a Chinese DEB family and review of the literature. <i>Journal of Cosmetic Dermatology</i> , 2020, 19, 1508-1512.	1.6	3
6	Identification of a Novel Mutation in SASH1 Gene in a Chinese Family With Dyschromatosis Universalis Hereditaria and Genotype-Phenotype Correlation Analysis. <i>Frontiers in Genetics</i> , 2020, 11, 841.	2.3	9
7	Annular epidermolytic ichthyosis with palmoplantar keratosis: a unique phenotype associated with interfamilial phenotypic heterogeneity. <i>European Journal of Dermatology</i> , 2020, 30, 294-299.	0.6	1
8	Inflammasomes in Common Immune-Related Skin Diseases. <i>Frontiers in Immunology</i> , 2020, 11, 882.	4.8	50
9	Genomic DNA methylation in HLA-Cw*0602 carriers and non-carriers of psoriasis. <i>Journal of Dermatological Science</i> , 2020, 99, 23-29.	1.9	1
10	Novel Mutation of the <i>NCSTN</i> Gene Identified in a Chinese Acne Inversa Family. <i>Annals of Dermatology</i> , 2020, 32, 237.	0.9	7
11	DNA methylation profile of psoriatic skins from different body locations. <i>Epigenomics</i> , 2019, 11, 1613-1625.	2.1	3
12	A KRT16 mutation in the first Chinese pedigree with Pachyonychia congenita and review of the literatures. <i>Journal of Cosmetic Dermatology</i> , 2019, 18, 1930-1934.	1.6	4
13	Novel compound heterozygous variants in the <i>XPC</i> gene identified in a Chinese xeroderma pigmentosum group C patient with ovarian teratoma. <i>Journal of Dermatology</i> , 2018, 45, e300-e301.	1.2	4
14	Integrative methylome and transcriptome analysis to dissect key biological pathways for psoriasis in Chinese Han population. <i>Journal of Dermatological Science</i> , 2018, 91, 285-291.	1.9	20
15	Novel mutations in Chinese Han patients with tuberous sclerosis complex: Case series and review of the published work. <i>Journal of Dermatology</i> , 2018, 45, 867-870.	1.2	5
16	Genetic Analysis of KRT9 Gene Revealed Previously Known Mutations and Genotype-Phenotype Correlations in Epidermolytic Palmoplantar Keratoderma. <i>Frontiers in Genetics</i> , 2018, 9, 645.	2.3	5
17	Genotype combination contributes to psoriasis: An exhaustive algorithm perspective. <i>PLoS ONE</i> , 2017, 12, e0186067.	2.5	5
18	Deep sequencing of the MHC region in the Chinese population contributes to studies of complex disease. <i>Nature Genetics</i> , 2016, 48, 740-746.	21.4	188

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19	Î±-Solanine inhibits vascular endothelial growth factor expression by down-regulating the ERK1/2-HIF-1Î± and STAT3 signaling pathways. <i>European Journal of Pharmacology</i> , 2016, 771, 93-98.	3.5	26
20	Changes of Regulatory T Cells in the Early Stage of Obesity Mice and Their Modulation on Macrophage Subtypes in Visceral Adipose Tissue. <i>Zhongguo Yi Xue Ke Xue Yuan Xue Bao Acta Academiae Medicinae Sinicae</i> , 2016, 38, 399-403.	0.2	0
21	Universal Stem-Loop Primer Method for Screening and Quantification of MicroRNA. <i>PLoS ONE</i> , 2014, 9, e115293.	2.5	63
22	The progression of the tubulointerstitial fibrosis driven by stress-induced “proliferation” death vicious circle. <i>Medical Hypotheses</i> , 2014, 82, 643-647.	1.5	2
23	Rare mutations in NLRP3 and NLRP12 associated with familial cold autoinflammatory syndrome: two Chinese pedigrees. <i>Clinical Rheumatology</i> , 0, , .	2.2	5