## Yuanqiang Zhang

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	PhenoApt leverages clinical expertise to prioritize candidate genes via machine learning. American Journal of Human Genetics, 2022, 109, 270-281.	6.2	5
2	Expanding the mutation and phenotype spectrum of MYH3-associated skeletal disorders. Npj Genomic Medicine, 2022, 7, 11.	3.8	7
3	Diagnostic yield and clinical impact of exome sequencing in early-onset scoliosis (EOS). Journal of Medical Genetics, 2021, 58, 41-47.	3.2	40
4	Factors and predictive model associated with perioperative complications after long fusion in the treatment of adult non-degenerative scoliosis. BMC Musculoskeletal Disorders, 2021, 22, 483.	1.9	3
5	Deciphering the mutational signature of congenital limb malformations. Molecular Therapy - Nucleic Acids, 2021, 24, 961-970.	5.1	8
6	Fexofenadine Protects Against Intervertebral Disc Degeneration Through TNF Signaling. Frontiers in Cell and Developmental Biology, 2021, 9, 687024.	3.7	10
7	Whole Exome Sequencing Uncovered the Genetic Architecture of Growth Hormone Deficiency Patients. Frontiers in Endocrinology, 2021, 12, 711991.	3.5	6
8	The GDF11 Promotes Nerve Regeneration After Sciatic Nerve Injury in Adult Rats by Promoting Axon Growth and Inhibiting Neuronal Apoptosis. Frontiers in Bioengineering and Biotechnology, 2021, 9, 803052.	4.1	4
9	<i>TBX6</i> missense variants expand the mutational spectrum in a nonâ€Mendelian inheritance disease. Human Mutation, 2020, 41, 182-195.	2.5	27
10	Front Cover, Volume 41, Issue 1. Human Mutation, 2020, 41, i.	2.5	0
11	Identification of novel FBN1 variations implicated in congenital scoliosis. Journal of Human Genetics, 2020, 65, 221-230.	2.3	20
12	Genetic and molecular mechanism for distinct clinical phenotypes conveyed by allelic truncating mutations implicated in <i>FBN1</i> . Molecular Genetics & Genomic Medicine, 2020, 8, e1023.	1.2	19
13	Phenotypic and genetic spectrum of isolated macrodactyly: somatic mosaicism of PIK3CA and AKT1 oncogenic variants. Orphanet Journal of Rare Diseases, 2020, 15, 288.	2.7	15
14	Mutational burden and potential oligogenic model of <i>TBX6</i> â€mediated genes in congenital scoliosis. Molecular Genetics & Genomic Medicine, 2020, 8, e1453.	1.2	6
15	Cost-effectiveness analysis of using the TBX6-associated congenital scoliosis risk score (TACScore) in genetic diagnosis of congenital scoliosis. Orphanet Journal of Rare Diseases, 2020, 15, 250.	2.7	2
16	A novel COMP mutation in a Chinese family with multiple epiphyseal dysplasia. BMC Medical Genetics, 2020, 21, 115.	2.1	2
17	The mutational burden and oligogenic inheritance in Klippel-Feil syndrome. BMC Musculoskeletal Disorders, 2020, 21, 220.	1.9	15
18	Estrogen Receptors (ESRs) Mutations in Adolescent Idiopathic Scoliosis: A Cross-Sectional Study. Medical Science Monitor, 2020, 26, e921611.	1.1	2

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19	A Recurrent Rare SOX9 Variant (M469V) is Associated with Congenital Vertebral Malformations. Current Gene Therapy, 2019, 19, 242-247.	2.0	11
20	IL-21 Is Positively Associated with Intervertebral Disc Degeneration by Interaction with TNF-α Through the JAK-STAT Signaling Pathway. Inflammation, 2017, 40, 612-622.	3.8	22
21	Follistatin-like protein 1 promotes inflammatory reactions in nucleus pulposus cells by interacting with the MAPK and NFI®B signaling pathways. Oncotarget, 2017, 8, 43023-43034.	1.8	31
22	Interleukin-9 Promotes TNF-α and PGE2 Release in Human Degenerated Intervertebral Disc Tissues. Spine, 2016, 41, 1631-1640.	2.0	23
23	Salubrinal Suppresses IL-17-Induced Upregulation of MMP-13 and Extracellular Matrix Degradation Through the NF-kB Pathway in Human Nucleus Pulposus Cells. Inflammation, 2016, 39, 1997-2007.	3.8	41
24	Production of CCL20 on nucleus pulposus cells recruits IL-17-producing cells to degenerated IVD tissues in rat models. Journal of Molecular Histology, 2016, 47, 81-89.	2.2	22