

Yiyang Wu

List of Publications by Year in descending order

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11
papers

1,104
citations

1039406

9
h-index

1281420

11
g-index

14
all docs

14
docs citations

14
times ranked

3621
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | Genotypic and phenotypic characterization of Chinese patients with osteogenesis imperfecta. <i>Human Mutation</i> , 2019, 40, 588-600. | 1.1 | 19 |
| 2 | NAA10-related syndrome. <i>Experimental and Molecular Medicine</i> , 2018, 50, 1-10. | 3.2 | 30 |
| 3 | Whole genome sequencing of one complex pedigree illustrates challenges with genomic medicine. <i>BMC Medical Genomics</i> , 2017, 10, 10. | 0.7 | 15 |
| 4 | Indel variant analysis of short-read sequencing data with Scalpel. <i>Nature Protocols</i> , 2016, 11, 2529-2548. | 5.5 | 99 |
| 5 | Genome-wide variant analysis of simplex autism families with an integrative clinical-bioinformatics pipeline. <i>Journal of Physical Education and Sports Management</i> , 2015, 1, a000422. | 0.5 | 6 |
| 6 | TAF1 Variants Are Associated with Dysmorphic Features, Intellectual Disability, and Neurological Manifestations. <i>American Journal of Human Genetics</i> , 2015, 97, 922-932. | 2.6 | 101 |
| 7 | Biochemical and cellular analysis of Ogden syndrome reveals downstream Nt-acetylation defects. <i>Human Molecular Genetics</i> , 2015, 24, 1956-1976. | 1.4 | 97 |
| 8 | Reducing INDEL calling errors in whole genome and exome sequencing data. <i>Genome Medicine</i> , 2014, 6, 89. | 3.6 | 144 |
| 9 | Accurate de novo and transmitted indel detection in exome-capture data using microassembly. <i>Nature Methods</i> , 2014, 11, 1033-1036. | 9.0 | 194 |
| 10 | Low concordance of multiple variant-calling pipelines: practical implications for exome and genome sequencing. <i>Genome Medicine</i> , 2013, 5, 28. | 3.6 | 381 |
| 11 | Congenital Syndromes Involving the Lungs: Pathogenetic Models Based on Chinese Medicine Theories. <i>Journal of Alternative and Complementary Medicine</i> , 2008, 14, 1017-1025. | 2.1 | 4 |