

# Qiaojun Jia

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4098341/publications.pdf>

Version: 2024-02-01

10  
papers

221  
citations

1307594

7  
h-index

1372567

10  
g-index

10  
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10  
docs citations

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times ranked

333  
citing authors

#	ARTICLE	IF	CITATIONS
1	Transcriptome Analysis of Different Sections of Rhizome in <i>Polygonatum sibiricum</i> Red. and Mining Putative Genes Participate in Polysaccharide Biosynthesis. <i>Biochemical Genetics</i> , 2022, 60, 1547-1566.	1.7	8
2	SSR Loci Analysis in Transcriptome and Molecular Marker Development in <i>Polygonatum sibiricum</i> . <i>BioMed Research International</i> , 2022, 2022, 1-9.	1.9	9
3	Structure characterization and bioactivity of neutral polysaccharides from different sources of <i>Polygonatum</i> Mill. <i>Biopolymers</i> , 2022, 113, e23490.	2.4	12
4	Advances in Pathophysiology of Triple-Negative Breast Cancer: The Potential of lncRNAs for Clinical Diagnosis, Treatment, and Prognostic Monitoring. <i>Molecular Biotechnology</i> , 2021, 63, 1093-1102.	2.4	4
5	Evaluation of genetic diversity and construction of DNA fingerprinting in <i>Polygonatum</i> Mill. based on EST-SSR and SRAP molecular markers. <i>3 Biotech</i> , 2020, 10, 322.	2.2	15
6	Long non-coding RNA MALAT1 regulates BLCAP mRNA expression through binding to miR-339-5p and promotes poor prognosis in breast cancer. <i>Bioscience Reports</i> , 2019, 39, .	2.4	48
7	Characterization of the <i>sdw1</i> semi-dwarf gene in barley. <i>BMC Plant Biology</i> , 2017, 17, 11.	3.6	78
8	Gene Deletion in Barley Mediated by LTR-retrotransposon BARE. <i>Scientific Reports</i> , 2017, 7, 43766.	3.3	13
9	Marker development using SLAF-seq and whole-genome shotgun strategy to fine-map the semi-dwarf gene <i>ari-e</i> in barley. <i>BMC Genomics</i> , 2016, 17, 911.	2.8	27
10	Characterization and mapping of a <i>Prbs</i> gene controlling spike development in <i>Hordeum vulgare</i> L.. <i>Genes and Genomics</i> , 2014, 36, 275-282.	1.4	7