

# Congcong Jiang

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6798432/publications.pdf>

Version: 2024-02-01

7  
papers

114  
citations

1684188  
5  
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1720034  
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7  
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docs citations

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times ranked

79  
citing authors

#	ARTICLE	IF	CITATIONS
1	Bymovirus-induced yellow mosaic diseases in barley and wheat: viruses, genetic resistances and functional aspects. <i>Theoretical and Applied Genetics</i> , 2020, 133, 1623-1640.	3.6	37
2	Bulked segregant RNA-sequencing (BSR-seq) identified a novel rare allele of eIF4E effective against multiple isolates of BaYMV/BaMMV. <i>Theoretical and Applied Genetics</i> , 2019, 132, 1777-1788.	3.6	30
3	Simultaneous editing of host factor gene <i>TaPDIL5</i> homoeoalleles confers wheat yellow mosaic virus resistance in hexaploid wheat. <i>New Phytologist</i> , 2022, 234, 340-344.	7.3	20
4	A reference-guided TILLING by amplicon-sequencing platform supports forward and reverse genetics in barley. <i>Plant Communications</i> , 2022, 3, 100317.	7.7	15
5	Marker-assisted selection in a global barley ( <i>Hordeum vulgare</i> subsp. <i>vulgare</i> ) collection revealed a unique genetic determinant of the naked barley controlled by the nud locus. <i>Genetic Resources and Crop Evolution</i> , 2020, 67, 273-280.	1.6	6
6	Genomic and pathogenic diversity of barley yellow mosaic virus and barley mild mosaic virus isolates in fields of China and their compatibility with resistance genes of cultivated barley. <i>Plant Disease</i> , 2022, , .	1.4	4
7	Evaluating the genetic effects of seed dormancy regulatory genes Qsd1 and Qsd2 in a global collection of cultivated barley ( <i>Hordeum vulgare</i> ssp. <i>vulgare</i> ) with functional kompetitive allele-specific PCR markers. <i>Plant Breeding</i> , 2021, 140, 827.	1.9	2