

# Xu Tang

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10703786/publications.pdf>

Version: 2024-02-01

25  
papers

2,722  
citations

430874

18  
h-index

610901

24  
g-index

27  
all docs

27  
docs citations

27  
times ranked

2430  
citing authors

#	ARTICLE	IF	CITATIONS
1	CRISPR-Cas9 mediated <i>OsMIR168a</i> knockout reveals its pleiotropy in rice. <i>Plant Biotechnology Journal</i> , 2022, 20, 310-322.	8.3	32
2	Genome-wide analyses of PAM-relaxed Cas9 genome editors reveal substantial off-target effects by ABE8e in rice. <i>Plant Biotechnology Journal</i> , 2022, 20, 1670-1682.	8.3	23
3	Optimal Power Allocation Strategy for a MIMO-Integrated Radar and Communication System Based OFDM Waveform. , 2022, , .		2
4	Single Transcript Unit CRISPR 2.0 Systems for Genome Editing in Rice. <i>Methods in Molecular Biology</i> , 2021, 2238, 193-204.	0.9	2
5	Efficient deletion of multiple circle RNA loci by CRISPR-Cas9 reveals <i>Os06circ02797</i> as a putative sponge for <i>OsMIR408</i> in rice. <i>Plant Biotechnology Journal</i> , 2021, 19, 1240-1252.	8.3	37
6	Expanding the scope of plant genome engineering with Cas12a orthologs and highly multiplexable editing systems. <i>Nature Communications</i> , 2021, 12, 1944.	12.8	79
7	Improved plant cytosine base editors with high editing activity, purity, and specificity. <i>Plant Biotechnology Journal</i> , 2021, 19, 2052-2068.	8.3	55
8	PAM-less plant genome editing using a CRISPR-SpRY toolbox. <i>Nature Plants</i> , 2021, 7, 25-33.	9.3	140
9	Rapid Vector Construction and Assessment of BE3 and Target-AID C to T Base Editing Systems in Rice Protoplasts. <i>Methods in Molecular Biology</i> , 2021, 2238, 95-113.	0.9	5
10	The Improvement of CRISPR-Cas9 System With Ubiquitin-Associated Domain Fusion for Efficient Plant Genome Editing. <i>Frontiers in Plant Science</i> , 2020, 11, 621.	3.6	12
11	Plant Prime Editors Enable Precise Gene Editing in Rice Cells. <i>Molecular Plant</i> , 2020, 13, 667-670.	8.3	148
12	Intron-Based Single Transcript Unit CRISPR Systems for Plant Genome Editing. <i>Rice</i> , 2020, 13, 8.	4.0	22
13	Bidirectional Promoter-Based CRISPR-Cas9 Systems for Plant Genome Editing. <i>Frontiers in Plant Science</i> , 2019, 10, 1173.	3.6	39
14	Multiplex QTL editing of grain-related genes improves yield in elite rice varieties. <i>Plant Cell Reports</i> , 2019, 38, 475-485.	5.6	136
15	Single transcript unit CRISPR 2.0 systems for robust Cas9 and Cas12a mediated plant genome editing. <i>Plant Biotechnology Journal</i> , 2019, 17, 1431-1445.	8.3	120
16	A Single Transcript CRISPR-Cas9 System for Multiplex Genome Editing in Plants. <i>Methods in Molecular Biology</i> , 2019, 1917, 75-82.	0.9	3
17	Application of CRISPR-Cas12a temperature sensitivity for improved genome editing in rice, maize, and Arabidopsis. <i>BMC Biology</i> , 2019, 17, 9.	3.8	172
18	Plant Genome Editing Using FxCpf1 and LbCpf1 Nucleases at Redefined and Altered PAM Sites. <i>Molecular Plant</i> , 2018, 11, 999-1002.	8.3	136

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19	A large-scale whole-genome sequencing analysis reveals highly specific genome editing by both Cas9 and Cpf1 (Cas12a) nucleases in rice. <i>Genome Biology</i> , 2018, 19, 84.	8.8	230
20	A CRISPR-Cpf1 system for efficient genome editing and transcriptional repression in plants. <i>Nature Plants</i> , 2017, 3, 17018.	9.3	425
21	CRISPR-Cas9 Based Genome Editing Reveals New Insights into MicroRNA Function and Regulation in Rice. <i>Frontiers in Plant Science</i> , 2017, 8, 1598.	3.6	150
22	Construction of a Single Transcriptional Unit for Expression of Cas9 and Single-guide RNAs for Genome Editing in Plants. <i>Bio-protocol</i> , 2017, 7, e2546.	0.4	2
23	A Single Transcript CRISPR-Cas9 System for Efficient Genome Editing in Plants. <i>Molecular Plant</i> , 2016, 9, 1088-1091.	8.3	144
24	Effective screen of CRISPR/Cas9-induced mutants in rice by single-strand conformation polymorphism. <i>Plant Cell Reports</i> , 2016, 35, 1545-1554.	5.6	74
25	A CRISPR/Cas9 Toolbox for Multiplexed Plant Genome Editing and Transcriptional Regulation. <i>Plant Physiology</i> , 2015, 169, 971-985.	4.8	532