

# Jennifer S Jaqueth

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1547782/publications.pdf>

Version: 2024-02-01

9  
papers

583  
citations

1307594

7  
h-index

1474206

9  
g-index

9  
all docs

9  
docs citations

9  
times ranked

872  
citing authors

#	ARTICLE	IF	CITATIONS
1	Rate and Pattern of Mutation at Microsatellite Loci in Maize. <i>Molecular Biology and Evolution</i> , 2002, 19, 1251-1260.	8.9	248
2	An Analysis of Genetic Diversity Across the Maize Genome Using Microsatellites. <i>Genetics</i> , 2005, 169, 1617-1630.	2.9	147
3	A pair of homoeolog ClpP5 genes underlies a <i>virescent yellowâ€like</i> mutant and its modifier in maize. <i>Plant Journal</i> , 2014, 79, 192-205.	5.7	58
4	tassel-less1 Encodes a Boron Channel Protein Required for Inflorescence Development in Maize. <i>Plant and Cell Physiology</i> , 2014, 55, 1044-1054.	3.1	46
5	Fertility restoration of maize <scp>CMS</scp>â€™ altered by a single amino acid substitution within the <i>Rf4</i> <scp>bHLH</scp> transcription factor. <i>Plant Journal</i> , 2020, 101, 101-111.	5.7	35
6	Identification and fineâ€mapping of <i>RppCML496</i>, a major QTL for resistance to <i>Puccinia polysora</i> in maize. <i>Plant Genome</i> , 2021, 14, e20062.	2.8	17
7	The Identification of Two Head Smut Resistance-Related QTL in Maize by the Joint Approach of Linkage Mapping and Association Analysis. <i>PLoS ONE</i> , 2015, 10, e0145549.	2.5	12
8	Identification of genetic loci associated with rough dwarf disease resistance in maize by integrating GWAS and linkage mapping. <i>Plant Science</i> , 2022, 315, 111100.	3.6	12
9	Identification and Fine Mapping of <i>qSCR4.01</i>, a Novel Major QTL for Resistance to <i>Puccinia polysora</i> in Maize. <i>Plant Disease</i> , 2020, 104, 1944-1948.	1.4	8