

# E Mace

## List of Publications by Year in descending order

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Version: 2024-02-01

14  
papers

893  
citations

932766

10  
h-index

1058022

14  
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14  
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docs citations

14  
times ranked

1345  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic control of leaf angle in sorghum and its effect on light interception. <i>Journal of Experimental Botany</i> , 2022, 73, 801-816.	2.4	10
2	Genetic diversity of Ethiopian sorghum reveals signatures of climatic adaptation. <i>Theoretical and Applied Genetics</i> , 2021, 134, 731-742.	1.8	23
3	Extensive variation within the pan-genome of cultivated and wild sorghum. <i>Nature Plants</i> , 2021, 7, 766-773.	4.7	94
4	Large-scale GWAS in sorghum reveals common genetic control of grain size among cereals. <i>Plant Biotechnology Journal</i> , 2020, 18, 1093-1105.	4.1	72
5	A Graph-Based Pan-Genome Guides Biological Discovery. <i>Molecular Plant</i> , 2020, 13, 1247-1249.	3.9	10
6	Large-scale genome-wide association study reveals that drought-induced lodging in grain sorghum is associated with plant height and traits linked to carbon remobilisation. <i>Theoretical and Applied Genetics</i> , 2020, 133, 3201-3215.	1.8	14
7	Differences in temperature response of phenological development among diverse Ethiopian sorghum genotypes are linked to racial grouping and agroecological adaptation. <i>Crop Science</i> , 2020, 60, 977-990.	0.8	12
8	Crop Genomics Goes Beyond a Single Reference Genome. <i>Trends in Plant Science</i> , 2019, 24, 1072-1074.	4.3	9
9	The Sorghum QTL Atlas: a powerful tool for trait dissection, comparative genomics and crop improvement. <i>Theoretical and Applied Genetics</i> , 2019, 132, 751-766.	1.8	114
10	Integrating modelling and phenotyping approaches to identify and screen complex traits: transpiration efficiency in cereals. <i>Journal of Experimental Botany</i> , 2018, 69, 3181-3194.	2.4	76
11	Fine Mapping of qDor7, a Major QTL Affecting Seed Dormancy in Sorghum ( <i>Sorghum bicolor</i> (L.) Tj). <i>ETQq1</i> 1.0.784314 rgBT / Overlooked	1.0	4
12	Whole-genome sequencing reveals untapped genetic potential in Africa's indigenous cereal crop sorghum. <i>Nature Communications</i> , 2013, 4, 2320.	5.8	405
13	Mapping spot blotch resistance genes in four barley populations. <i>Molecular Breeding</i> , 2010, 26, 653-666.	1.0	24
14	Mapping of adult plant resistance to net form of net blotch in three Australian barley populations. <i>Australian Journal of Agricultural Research</i> , 2007, 58, 1191.	1.5	26