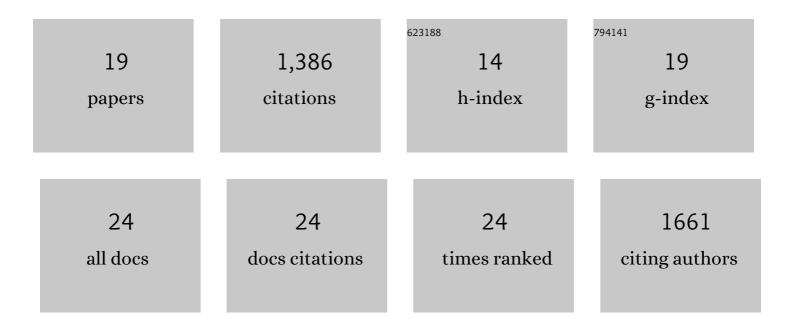
Ismail Y Rabbi

List of Publications by Year in descending order

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ISMAIL Y PARRI

#	Article	IF	CITATIONS
1	Selection for resistance to cassava mosaic disease in African cassava germplasm using single nucleotide polymorphism markers. South African Journal of Science, 2022, 118, .	0.3	3
2	Largeâ€scale genomeâ€wide association study, using historical data, identifies conserved genetic architecture of cyanogenic glucoside content in cassava (<i>Manihot esculenta</i> Crantz) root. Plant Journal, 2021, 105, 754-770.	2.8	26
3	Conversion and Validation of Uniplex SNP Markers for Selection of Resistance to Cassava Mosaic Disease in Cassava Breeding Programs. Agronomy, 2021, 11, 420.	1.3	10
4	ldentifying New Resistance to Cassava Mosaic Disease and Validating Markers for the CMD2 Locus. Agriculture (Switzerland), 2021, 11, 829.	1.4	8
5	The Cassava Source–Sink project: opportunities and challenges for crop improvement by metabolic engineering. Plant Journal, 2020, 103, 1655-1665.	2.8	33
6	Genomic prediction and quantitative trait locus discovery in a cassava training population constructed from multiple breeding stages. Crop Science, 2020, 60, 896-913.	0.8	9
7	Poverty Reduction Effects of Agricultural Technology Adoption: The Case of Improved Cassava Varieties in Nigeria. Journal of Agricultural Economics, 2019, 70, 392-407.	1.6	51
8	Estimating the Productivity Impacts of Technology Adoption in the Presence of Misclassification. American Journal of Agricultural Economics, 2019, 101, 1-16.	2.4	47
9	Cassava haplotype map highlights fixation of deleterious mutations during clonal propagation. Nature Genetics, 2017, 49, 959-963.	9.4	208
10	Genomeâ€Wide Association Mapping of Correlated Traits in Cassava: Dry Matter and Total Carotenoid Content. Plant Genome, 2017, 10, plantgenome2016.09.0094.	1.6	63
11	Prospects for Genomic Selection in Cassava Breeding. Plant Genome, 2017, 10, plantgenome2017.03.0015.	1.6	101
12	Marker-Based Estimates Reveal Significant Nonadditive Effects in Clonally Propagated Cassava (<i>Manihot esculenta</i>): Implications for the Prediction of Total Genetic Value and the Selection of Varieties. G3: Genes, Genomes, Genetics, 2016, 6, 3497-3506.	0.8	34
13	Genomeâ€Wide Association and Prediction Reveals Genetic Architecture of Cassava Mosaic Disease Resistance and Prospects for Rapid Genetic Improvement. Plant Genome, 2016, 9, plantgenome2015.11.0118.	1.6	120
14	Sequencing wild and cultivated cassava and related species reveals extensive interspecific hybridization and genetic diversity. Nature Biotechnology, 2016, 34, 562-570.	9.4	340
15	Tracking crop varieties using genotyping-by-sequencing markers: a case study using cassava (Manihot) Tj ETQq1	1 9.78431	l4rgBT /Ove
16	The Effects of Restrictionâ€Enzyme Choice on Properties of Genotypingâ€byâ€5equencing Libraries: A Study in Cassava (<i>Manihot esculenta</i>). Crop Science, 2014, 54, 2603-2608.	0.8	51
17	solGS: a web-based tool for genomic selection. BMC Bioinformatics, 2014, 15, 398.	1.2	18
18	High-resolution mapping of resistance to cassava mosaic geminiviruses in cassava using genotyping-by-sequencing and its implications for breeding. Virus Research, 2014, 186, 87-96.	1.1	143

#	Article	IF	CITATIONS
19	Impact of farmers' practices and seed systems on the genetic structure of common sorghum varieties in Kenya and Sudan. Plant Genetic Resources: Characterisation and Utilisation, 2010, 8, 116-126.	0.4	12