Nino Brown

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2998241/publications.pdf

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		1937685	1474206	
17	83	4	9	
papers	citations	h-index	g-index	
17	17	17	81	
all docs	docs citations	times ranked	citing authors	

#	Article	IF	CITATIONS
1	Pyramiding novel EMS-generated mutant alleles to improve fiber quality components of elite upland cotton germplasm. Industrial Crops and Products, 2022, 178, 114594.	5.2	4
2	Improved Upland Cotton Germplasm for Multiple Fiber Traits Mediated by Transferring and Pyramiding Novel Alleles From Ethyl Methanesulfonate-Generated Mutant Lines Into Elite Genotypes. Frontiers in Plant Science, 2022, 13, 842741.	3.6	5
3	Genetic diversity assessment of Georgia peanut cultivars developed during ninety years of breeding. Plant Genome, 2021, 14, e20141.	2.8	4
4	Sound Splits as Influenced by Seed Size for Runner and Virginia Market Type Peanut Shelled on a Reciprocating Sheller. Agronomy, 2021, 11, 1869.	3.0	1
5	Registration of Spearâ€shaped Leaf peanut genetic stock. Journal of Plant Registrations, 2020, 14, 457-459.	0.5	0
6	Registration of Albinoâ€Virescent Leaf peanut genetic stock. Journal of Plant Registrations, 2020, 14, 460-463.	0.5	0
7	Registration of Revoluteâ€Leaf peanut genetic stock. Journal of Plant Registrations, 2020, 14, 464-466.	0.5	0
8	Registration of eight upland cotton (Gossypium hirsutumL.) germplasm lines withqFLâ€Chr.25, a fiberâ€length QTL introgressed fromGossypium barbadense. Journal of Plant Registrations, 2020, 14, 57-63.	0.5	1
9	Registration of six upland cotton germplasm lines with improved fiber quality through ethyl methane sulfonate treatments and selection. Journal of Plant Registrations, 2020, 14, 159-164.	0.5	1
10	Inheritance of an Albino-Virescent Leaf Mutant in the Cultivated Peanut (<i>Arachis hypogaea</i> L.). Peanut Science, 2019, 46, 203-205.	0.1	1
11	Evaluation of a Chromosome Segment from G <i>ossypium barbadense</i> Harboring the Fiber Length QTL <i>qFLâ€Chr.25</i> in Four Diverse Upland Cotton Genetic Backgrounds. Crop Science, 2019, 59, 2621-2633.	1.8	4
12	Registration of GA R01â€40â€08, a <i>Gossypium hirsutum</i> Upland Cotton Germplasm Line with <i>qFLâ€Chr.1</i> Introgressed from <i>Gossypium barbadense</i> Conferring Improved Fiber Length. Journal of Plant Registrations, 2019, 13, 406-410.	0.5	1
13	Comparative genetic variation of fiber quality traits in reciprocal advanced backcross populations. Euphytica, 2017, 213, 1.	1.2	9
14	Withinâ€Boll Yield Characteristics and Their Correlation with Fiber Quality Parameters following Mutagenesis of Upland Cotton, TAM 94Lâ€25. Crop Science, 2015, 55, 1513-1523.	1.8	7
15	Improvement of Upland Cotton Fiber Quality through Mutation of TAM 94Lâ€25. Crop Science, 2013, 53, 452-459.	1.8	5
16	Registration of TAM 94Lâ€25â€M24, TAM 94Lâ€25â€M25, and TAM 94Lâ€25â€M30 Mutant Upland Cotton Ger with Improved Fiber Length and Strength. Journal of Plant Registrations, 2012, 6, 195-199.	mplasm	7
17	Development of Extraâ€Long Staple Upland Cotton. Crop Science, 2008, 48, 1823-1831.	1.8	33