## Graham R Teakle

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7855831/publications.pdf

Version: 2024-02-01

22 papers 1,411 citations

623734 14 h-index 677142 22 g-index

24 all docs

24 docs citations

times ranked

24

1988 citing authors

#	Article	IF	Citations
1	Identification of microbial signatures linked to oilseed rape yield decline at the landscape scale. Microbiome, 2021, 9, 19.	11.1	31
2	The Evolutionary History of Wild, Domesticated, and Feral <i>Brassica oleracea</i> (Brassicaceae). Molecular Biology and Evolution, 2021, 38, 4419-4434.	8.9	49
3	Quantitative Trait Locus Mapping of Resistance to Turnip Yellows Virus in Brassica rapa and Brassica oleracea and Introgression of These Resistances by Resynthesis Into Allotetraploid Plants for Deployment in Brassica napus. Frontiers in Plant Science, 2021, 12, 781385.	3.6	9
4	Identification and QTL mapping of resistance to Turnip yellows virus (TuYV) in oilseed rape, Brassica napus. Theoretical and Applied Genetics, 2020, 133, 383-393.	3.6	19
5	Cordycepin, a metabolite of Cordyceps militaris, reduces immune-related gene expression in insects. Journal of Invertebrate Pathology, 2020, 177, 107480.	3.2	13
6	Transcriptome and organellar sequencing highlights the complex origin and diversification of allotetraploid Brassica napus. Nature Communications, $2019$ , $10$ , $2878$ .	12.8	78
7	Assembly and characterisation of a unique onion diversity set identifies resistance to Fusarium basal rot and improved seedling vigour. Theoretical and Applied Genetics, 2019, 132, 3245-3264.	3.6	20
8	Addressing the threat of climate change to agriculture requires improving crop resilience to short-term abiotic stress. Outlook on Agriculture, 2018, 47, 270-276.	3.4	14
9	Towards new sources of resistance to the currant-lettuce aphid (Nasonovia ribisnigri). Molecular Breeding, 2017, 37, 4.	2.1	17
10	Development of a Statistical Crop Model to Explain the Relationship between Seed Yield and Phenotypic Diversity within the Brassica napus Genepool. Agronomy, 2017, 7, 31.	3.0	13
11	The pangenome of an agronomically important crop plant Brassica oleracea. Nature Communications, 2016, 7, 13390.	12.8	375
12	An introduction to the contemporary breeding of oil seed rape. Lipid Technology, 2013, 25, 251-254.	0.3	3
13	Functional alleles of the flowering time regulator FRIGIDA in the Brassica oleraceagenome. BMC Plant Biology, 2012, 12, 21.	3.6	51
14	Developing genetic resources for pre-breeding in Brassica oleracea L.: an overview of the UK perspective. Journal of Plant Biotechnology, 2012, 39, 62-68.	0.4	13
15	Double haploids, markers and QTL analysis in vegetable brassicas. Euphytica, 2008, 164, 509-514.	1.2	46
16	Shoot Calcium and Magnesium Concentrations Differ between Subtaxa, Are Highly Heritable, and Associate with Potentially Pleiotropic Loci in <i>Brassica oleracea</i> Â Â Â. Plant Physiology, 2008, 146, 1707-1720.	4.8	107
17	The reference genetic linkage map for the multinational Brassica rapa genome sequencing project. Theoretical and Applied Genetics, 2007, 115, 777-792.	3.6	160
18	Genetic analysis of the bracting trait in cauliflower and broccoli. Plant Science, 2003, 164, 803-808.	3.6	15

#	Article	IF	CITATIONS
19	Arabidopsis thaliana GATA factors: organisation, expression and DNA-binding characteristics. Plant Molecular Biology, 2002, 50, 43-56.	3.9	101
20	Integration of the Cytogenetic and Genetic Linkage Maps of <i>Brassica oleracea</i> . Genetics, 2002, 161, 1225-1234.	2.9	108
21	The GATA-binding protein CGF-1 is closely related to GT-1. Plant Molecular Biology, 1995, 29, 1253-1266.	3.9	31
22	Circadian clock- and phytochrome-regulated transcription is conferred by a 78 bp cis-acting domain of the Arabidopsis CAB2 promoter. Plant Journal, 1994, 6, 457-470.	5.7	136