Helena Oakey

List of Publications by Year in descending order

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623734 610901 1,242 24 14 24 citations g-index h-index papers 27 27 27 1954 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	A Long-Term Evaluation of Facebook for Recruitment and Retention in the ENDIA Type 1 Diabetes Pregnancy-Birth Cohort Study. Journal of Diabetes Science and Technology, 2023, 17, 696-704.	2.2	2
2	Women with type 1 diabetes exhibit a progressive increase in gut Saccharomyces cerevisiae in pregnancy associated with evidence of gut inflammation. Diabetes Research and Clinical Practice, 2022, 184, 109189.	2.8	6
3	Comparison of Methods to Select Candidates for High-Density Genotyping; Practical Observations in a Cattle Breeding Program. Agriculture (Switzerland), 2022, 12, 276.	3.1	2
4	Within-breed selection is sufficient to improve terminal crossbred beef marbling: a review of reciprocal recurrent genomic selection. Animal Production Science, 2021, , .	1.3	3
5	Assessing Rice Salinity Tolerance: From Phenomics to Association Mapping. Methods in Molecular Biology, 2021, 2238, 339-375.	0.9	4
6	Genetic mapping of the early responses to salt stress in <i>Arabidopsis thaliana</i> . Plant Journal, 2021, 107, 544-563.	5.7	22
7	Evaluation of protocol amendments to the Environmental Determinants of Islet Autoimmunity (ENDIA) study during the COVIDâ€19 pandemic. Diabetic Medicine, 2021, 38, e14638.	2.3	2
8	Type 1 diabetes in pregnancy is associated with distinct changes in the composition and function of the gut microbiome. Microbiome, 2021, 9, 167.	11.1	23
9	Quinoa Phenotyping Methodologies: An International Consensus. Plants, 2021, 10, 1759.	3.5	24
10	Optimization of Blood Handling and Peripheral Blood Mononuclear Cell Cryopreservation of Low Cell Number Samples. International Journal of Molecular Sciences, 2021, 22, 9129.	4.1	9
11	Stimulation of homologous recombination in plants expressing heterologous recombinases. BMC Plant Biology, 2020, 20, 336.	3.6	11
12	Association mapping identifies quantitative trait loci (QTL) for digestibility in rice straw. Biotechnology for Biofuels, 2020, 13, 165.	6.2	7
13	Barley sodium content is regulated by natural variants of the Na+ transporter HvHKT1;5. Communications Biology, 2020, 3, 258.	4.4	21
14	Changes in pancreatic exocrine function in young atâ€risk children followed to islet autoimmunity and type 1 diabetes in the <scp>ENDIA</scp> study. Pediatric Diabetes, 2020, 21, 945-949.	2.9	9
15	<scp>RNA</scp> iâ€suppression of barley caffeic acid <i>O</i> â€methyltransferase modifies lignin despite redundancy in the gene family. Plant Biotechnology Journal, 2019, 17, 594-607.	8.3	37
16	Highâ€throughput 3D modelling to dissect the genetic control of leaf elongation in barley (<i>Hordeum vulgare</i>). Plant Journal, 2019, 98, 555-570.	5.7	20
17	Salinity tolerance loci revealed in rice using high-throughput non-invasive phenotyping. Nature Communications, 2016, 7, 13342.	12.8	218
18	Yield-related salinity tolerance traits identified in a nested association mapping (NAM) population of wild barley. Scientific Reports, 2016, 6, 32586.	3.3	118

#	Article	IF	CITATION
19	The effect of temperature on the male and female recombination landscape of barley. New Phytologist, 2015, 208, 421-429.	7.3	105
20	The Synaptonemal Complex Protein ZYP1 Is Required for Imposition of Meiotic Crossovers in Barley. Plant Cell, 2014, 26, 729-740.	6.6	88
21	Maternal and neonatal outcomes following induction of labor: a cohort study. Acta Obstetricia Et Gynecologica Scandinavica, 2012, 91, 198-203.	2.8	43
22	Effect of Treatment of Gestational Diabetes Mellitus on Obesity in the Next Generation. Diabetes Care, 2010, 33, 964-968.	8.6	224
23	Joint modeling of additive and non-additive (genetic line) effects in multi-environment trials. Theoretical and Applied Genetics, 2007, 114, 1319-1332.	3.6	70
24	Joint modeling of additive and non-additive genetic line effects in single field trials. Theoretical and Applied Genetics, 2006, 113, 809-819.	3.6	170