Gunter Backes

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Association Mapping for Common Bunt Resistance in Wheat Landraces and Cultivars. Agronomy, 2022, 12, 642.	3.0	2
2	Pathogenic variability of a Uruguayan population of Bipolaris sorokiniana in barley suggests a mix of qualitative interactions. Journal of Plant Diseases and Protection, 2020, 127, 25-33.	2.9	4
3	Identification of Ideal Allele Combinations for the Adaptation of Spring Barley to Northern Latitudes. Frontiers in Plant Science, 2019, 10, 542.	3.6	10
4	Exposure to Ultraviolet (UV-C) Radiation Increases Germination Rate of Maize (Zea maize L.) and Sugar Beet (Beta vulgaris) Seeds. Plants, 2019, 8, 49.	3.5	17
5	Mapping of common bunt resistance gene Bt9 in wheat. Theoretical and Applied Genetics, 2017, 130, 1031-1040.	3.6	20
6	Genetic diversity, population structure and linkage disequilibrium in Nordic spring barley (Hordeum) Tj ETQq0 0 C) rgBT /Ove	erlgck 10 Tf S
7	Chromosomal regions associated with the <i>in vitro</i> culture response of wheat (<i><scp>T</scp>riticum aestivum </i> <scp>L</scp> .) microspores. Plant Breeding, 2015, 134, 255-263.	1.9	13
8	Significant decrease in yield under future climate conditions: Stability and production of 138 spring barley accessions. European Journal of Agronomy, 2015, 63, 105-113.	4.1	43
9	Genetic Diversity and Population Structure Analysis of European Hexaploid Bread Wheat (Triticum) Tj ETQq $1\ 1\ 0$.	784314 rg 2.5	gBT_/Overloci
10	Genetic diversity and structure found in samples of Eritrean bread wheat. Plant Genetic Resources: Characterisation and Utilisation, 2014, 12, 151-155.	0.8	1
11	Changes in allelic frequency over time in European bread wheat (Triticum aestivum L.) varieties revealed using DArT and SSR markers. Euphytica, 2014, 197, 447-462.	1.2	32
12	TILLING and EcoTILLING. , 2013, , 145-165.		2

13	Genome-wide Association Study of Resistant Starch (RS) Phenotypes in a Barley Variety Collection. Journal of Agricultural and Food Chemistry, 2012, 60, 10302-10311.	5.2	27
14	Integration of the barley genetic and seed proteome maps for chromosome 1H, 2H, 3H, 5H and 7H. Functional and Integrative Genomics, 2009, 9, 135-143.	3.5	18
15	High genetic diversity revealed in barley (Hordeum vulgare) collected from small-scale farmer's fields in Eritrea. Genetic Resources and Crop Evolution, 2009, 56, 85-97.	1.6	19
16	Genetic diversity and population structure of wild and cultivated barley from West Asia and North Africa. Plant Breeding, 2009, 128, 606-614.	1.9	17
17	Identification of barley mutants in the cultivar â€~Lux' at the <i>Dhn</i> loci through TILLING. Plant Breeding, 2009, 128, 332-336.	1.9	42
18	Molecular markers to exploit genotype–environment interactions of relevance in organic growing	1.2	17

systems. Euphytica, 2008, 163, 523-531.

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19	The Horn of Africa as a centre of barley diversification and a potential domestication site. Theoretical and Applied Genetics, 2007, 114, 1117-1127.	3.6	56
20	Barley. , 2006, , 155-210.		2
21	EcoTILLING for the identification of allelic variation in the powdery mildew resistance genes mlo and Mla of barley. Plant Breeding, 2006, 125, 461-467.	1.9	74
22	QTLs for straw quality characteristics identified in recombinant inbred lines of a Hordeum vulgare � H. spontaneum cross in a Mediterranean environment. Theoretical and Applied Genetics, 2005, 110, 688-695.	3.6	31
23	New molecular markers linked to qualitative and quantitative powdery mildew and scald resistance genes in barley for dry areas. Euphytica, 2004, 135, 225-228.	1.2	18
24	QTLs and Genes for Disease Resistance in Barley and Wheat. , 2004, , 199-251.		5
25	Localisation of genes for resistance against Blumeria graminis f.sp. hordei and Puccinia graminis in a cross between a barley cultivar and a wild barley (Hordeum vulgare ssp. spontaneum) line. Theoretical and Applied Genetics, 2003, 106, 353-362.	3.6	64
26	QTLs for agronomic traits in the Mediterranean environment identified in recombinant inbred lines of the cross 'Arta' × H. spontaneum 41-1. Theoretical and Applied Genetics, 2003, 107, 1215-1225.	3.6	196
27	RFLP diversity within and between major groups of barley in Europe. Plant Breeding, 2003, 122, 291-299.	1.9	33
28	Localising QTLs for leaf rust resistance and agronomic traits in barley (Hordeum vulgare L.). Theoretical and Applied Genetics, 2000, 100, 881-888.	3.6	56
29	Short Communication Comparison between QTL analysis of powdery mildew resistance in barley based on detached primary leaves and on field data. Plant Breeding, 1996, 115, 419-421.	1.9	31
30	Localization of quantitative trait loci (QTL) for agronomic important characters by the use of a RFLP map in barley (Hordeum vulgare L.). Theoretical and Applied Genetics, 1995, 90, 294-302.	3.6	165
31	RFLP markers to identify the alleles on the Mla locus conferring powdery mildew resistance in barley. Theoretical and Applied Genetics, 1992, 84-84, 330-338.	3.6	55
32	Development of RFLP Markers for Barley. Plant Breeding, 1991, 107, 73-76.	1.9	22