WacÅ,aw Orczyk

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Silencing of the HvCKX1 gene decreases the cytokinin oxidase/dehydrogenase level in barley and leads to higher plant productivity. Journal of Experimental Botany, 2010, 61, 1839-1851.	4.8	183
2	Major genes determining yield-related traits in wheat and barley. Theoretical and Applied Genetics, 2017, 130, 1081-1098.	3.6	175
3	Somatic hybrids of Solanum tuberosum – application to genetics and breeding. Plant Cell, Tissue and Organ Culture, 2003, 74, 1-13.	2.3	59
4	HvCKX2 gene silencing by biolistic or Agrobacterium-mediated transformation in barley leads to different phenotypes. BMC Plant Biology, 2012, 12, 206.	3.6	49
5	Expression Patterns of HvCKX Genes Indicate Their Role in Growth and Reproductive Development of Barley. PLoS ONE, 2014, 9, e115729.	2.5	45
6	The RNA-mediated silencing of one of the Pin genes in allohexaploid wheat simultaneously decreases the expression of the other, and increases grain hardness. Journal of Experimental Botany, 2011, 62, 4025-4036.	4.8	38
7	TaWAK6 encoding wall-associated kinase is involved in wheat resistance to leaf rust similar to adult plant resistance. PLoS ONE, 2020, 15, e0227713.	2.5	36
8	The Agrobacterium-mediated transformation of common wheat (Triticum aestivum L.) and triticale (x) Tj ETQq Genetics, 2012, 53, 1-8.	0 0 0 rgBT / 1.9	Overlock 10 ⁻ 29
9	Structural characteristics of ScBx genes controlling the biosynthesis of hydroxamic acids in rye (Secale cereale L.). Journal of Applied Genetics, 2015, 56, 287-298.	1.9	29
10	Agrobacterium-mediated transformation of polyploid cereals. The efficiency of selection and transgene expression in wheat. Cellular and Molecular Biology Letters, 2004, 9, 903-17.	7.0	28
11	Pathogen-regulated genes in wheat isogenic lines differing in resistance to brown rust Puccinia triticina. BMC Genomics, 2015, 16, 742.	2.8	27
12	Identification and VIGS-based characterization of Bx1 ortholog in rye (Secale cereale L.). PLoS ONE, 2017, 12, e0171506.	2.5	23
13	Specificity of expression of TaCKX family genes in developing plants of wheat and their co-operation within and among organs. PLoS ONE, 2019, 14, e0214239.	2.5	22
14	Silencing of TaCKX1 Mediates Expression of Other TaCKX Genes to Increase Yield Parameters in Wheat. International Journal of Molecular Sciences, 2020, 21, 4809.	4.1	22
15	Different sets of TaCKX genes affect yield-related traits in wheat plants grown in a controlled environment and in field conditions. BMC Plant Biology, 2020, 20, 496.	3.6	13
16	Changes in benzoxazinoid contents and the expression of the associated genes in rye (Secale cereale) Tj ETQc	10 0 0 rgBT /	Overlock 10
17	Annotation and profiling of barley GLYCOGEN SYNTHASE3/Shaggy-like genes indicated shift in organ-preferential expression, PLoS ONF, 2018, 13, e0199364.	2.5	11

18Silencing of HvGSK1.1â€"A GSK3/SHAGGY-Like Kinaseâ€"Enhances Barley (Hordeum vulgare L.) Growth in
Normal and in Salt Stress Conditions. International Journal of Molecular Sciences, 2020, 21, 6616.4.111

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19	TaCKX2.2 Genes Coordinate Expression of Other TaCKX Family Members, Regulate Phytohormone Content and Yield-Related Traits of Wheat. International Journal of Molecular Sciences, 2021, 22, 4142.	4.1	10
20	Genotype-Dependent Effect of Silencing of TaCKX1 and TaCKX2 on Phytohormone Crosstalk and Yield-Related Traits in Wheat. International Journal of Molecular Sciences, 2021, 22, 11494.	4.1	8
21	A new BSMV-based vector with modified \hat{l}^2 molecule allows simultaneous and stable silencing of two genes. Cellular and Molecular Biology Letters, 2012, 17, 107-23.	7.0	7
22	Polyamine Oxidation Is Indispensable for Wheat (Triticum aestivum L.) Oxidative Response and Necrotic Reactions during Leaf Rust (Puccinia triticina Eriks.) Infection. Plants, 2021, 10, 2787.	3.5	1