

Zdenka Galova

List of Publications by Year in descending order

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Version: 2024-02-01

60
papers

192
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1478505

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times ranked

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#	ARTICLE	IF	CITATIONS
1	Molecular Characterization of Novel x-Type HMW Glutenin Subunit 1B Å— 6.5 in Wheat. <i>Plants</i> , 2021, 10, 2108.	3.5	0
2	MicroRNA-Based and Proteomics Fingerprinting of <i>Avena sativa</i> L. Genotypes. , 2021, 11, .		0
3	Detection of Celiac Active Polypeptides in Wheat, Oat and Buckwheat Using Immunochemical Methods. , 2021, 11, .		0
4	Genetic Diversity of Oat Genotypes Using SCoT Markers. , 2021, 11, .		1
5	Proteomic and Genetic Approach for Lunasin Peptide and Gene Presence Detection in Various Plants. , 2021, 11, .		0
6	Genetic diversity and relationship of Tunisian castor (<i>Ricinus communis</i> L.) genotypes revealed by SSR markers. <i>Genetika</i> , 2020, 52, 765-776.	0.4	0
7	CHARACTERISTICS OF CEREALS, PSEUDOCEREALS AND LEGUMES FOR THEIR COELIAC ACTIVE POLYPEPTIDES. <i>Journal of Microbiology, Biotechnology and Food Sciences</i> , 2019, 9, 390-395.	0.8	1
8	Genetic diversity in Tunisian castor genotypes (<i>Ricinus communis</i> L.) detected using RAPD markers. <i>Potravinarstvo</i> , 2019, 13, 294-300.	0.6	4
9	Determination of HMW â€“ GS in wheat using SDS â€“ PAGE and Lab-on-chip methods. <i>Potravinarstvo</i> , 2019, 13, 477-481.	0.6	1
10	Electrophoretic profiles of storage proteins in selected maize (<i>Zea mays</i> L.) genotypes. <i>Journal of Central European Agriculture</i> , 2019, 20, 911-918.	0.6	1
11	Comparison of selected wheat, oat and buckwheat genotypes on proteomic level. <i>Journal of Central European Agriculture</i> , 2019, 20, 891-899.	0.6	1
12	Start codon targeted polymorphism for evaluation of functional genetic variation and relationships in cultivated castor (<i>Ricinus communis</i> L.) genotypes. <i>Genetika</i> , 2019, 51, 137-146.	0.4	5
13	PROTEIN PROFILES OF CEREALS AND PSEUDOCEREALS DETERMINED BY TWO-DIMENSIONAL GEL ELECTROPHORESIS. <i>Journal of Microbiology, Biotechnology and Food Sciences</i> , 2019, 9, 359-365.	0.8	1
14	Characterization of Tunisian castor bean genotypes using SDS-PAGE of total seed storage proteins. <i>Potravinarstvo</i> , 2018, 12, 701-706.	0.6	1
15	Comparison of American and European maize (<i>Zea mays</i> L.) protein profiles. <i>Journal of Central European Agriculture</i> , 2018, 19, 453-465.	0.6	3
16	Comparison of nutritional and technological quality of winter wheat (<i>Triticum aestivum</i> L.) and hybrid wheat (<i>Triticum aestivum</i> L. x <i>Triticum spelta</i> L.). <i>Journal of Central European Agriculture</i> , 2018, 19, 437-452.	0.6	3
17	Genetic diversity and population structure in tunisian castor genotypes (<i>Ricinus communis</i> L.) Detected using scot markers. <i>Potravinarstvo</i> , 2018, 12, .	0.6	4
18	Study of polymorphism of maize using dna and protein markers. <i>Potravinarstvo</i> , 2018, 12, .	0.6	1

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19	Molecular analysis of buckwheat using gene specific markers. <i>Potravinarstvo</i> , 2018, 12, 546-552.	0.6	2
20	Chitinase Activities in Wheat and Its Relative Species. <i>Agriculture</i> , 2017, 63, 14-22.	0.4	2
21	POTENTIAL OF SELECTED SSR MARKERS FOR IDENTIFICATION OF MALTING BARLEY GENOTYPES. <i>Journal of Microbiology, Biotechnology and Food Sciences</i> , 2017, 6, 1276-1279.	0.8	3
22	Detection genetic variability of <i>secale cereale</i> L. by scot markers. <i>Potravinarstvo</i> , 2017, 11, .	0.6	7
23	GENETIC VARIATION OF MAIZE GENOTYPES (<i>ZEAMAYS</i> L.) DETECTED USING SDS-PAGE. <i>Journal of Microbiology, Biotechnology and Food Sciences</i> , 2017, 6, 1086-1089.	0.8	0
24	Genetic variation of european maize genotypes (<i>Zeamays</i> L.) Detected using ssr markers. <i>Potravinarstvo</i> , 2017, 11, 126-131.	0.6	4
25	GENETIC DIVERSITY ANALYSIS OF MAIZE (<i>ZEAMAYS</i> L.) USING SCoT MARKERS. <i>Journal of Microbiology, Biotechnology and Food Sciences</i> , 2017, 6, 1170-1173.	0.8	10
26	Molecular variability of oat based on gene specific markers. <i>Potravinarstvo</i> , 2017, 11, .	0.6	2
27	Perception of biotech trees by Slovak university students – a comparative survey. <i>Nova Biotechnologica Et Chimica</i> , 2017, 16, 12-19.	0.1	0
28	THE FAMILY OF CHITINASES IN COTTON <i>G. raimondii</i> . <i>Journal of Microbiology, Biotechnology and Food Sciences</i> , 2017, 6, 1284-1289.	0.8	2
29	GENETIC DIVERSITY OF WINTER WHEAT (<i>Triticum aestivum</i> L.). <i>Journal of Microbiology, Biotechnology and Food Sciences</i> , 2017, 6, 1233-1236.	0.8	0
30	Beta-1,3-Glucanase Activities in Wheat and Relative Species. <i>Nova Biotechnologica Et Chimica</i> , 2016, 15, 122-132.	0.1	5
31	Agrochemicals affect the antioxidative defense potential of cotton plants. <i>Journal of Microbiology, Biotechnology and Food Sciences</i> , 2016, 05, 505-508.	0.8	2
32	Genetic variation and relationships of old maize genotypes (<i>Zeamays</i> l.) detected using SDS-page. <i>Potravinarstvo</i> , 2016, 10, .	0.6	5
33	Genetic diversity of European cultivars of common wheat (<i>Triticum aestivum</i> L.) based on RAPD and protein markers. <i>Journal of Central European Agriculture</i> , 2016, 17, 957-969.	0.6	5
34	Polymorphism of proteins in selected slovak winter wheat genotypes using SDS-PAGE. <i>Journal of Central European Agriculture</i> , 2016, 17, 970-985.	0.6	7
35	RAPD analysis of the genetic polymorphism in european wheat genotypes. <i>Potravinarstvo</i> , 2016, 10, .	0.6	4
36	Evaluation of molecular diversity of central European maize cultivars. <i>Emirates Journal of Food and Agriculture</i> , 2016, 28, 93.	1.0	4

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37	Lunasin detection in coloured wheat genotype. <i>Potravinarstvo</i> , 2016, 10, 152-156.	0.6	0
38	Molecular characterization of rye cultivars. <i>Potravinarstvo</i> , 2016, 10, .	0.6	4
39	Assessment of rapd polymorphism in ricin genotypes. <i>Journal of Microbiology, Biotechnology and Food Sciences</i> , 2016, 05, 386-388.	0.8	1
40	Application of rye SSR markers for detection of genetic diversity in triticale. <i>Journal of Microbiology, Biotechnology and Food Sciences</i> , 2016, 05, 623-626.	0.8	2
41	Start codon targeted (scot) polymorphism reveals genetic diversity in european old maize (<i>zea mays</i> l.) Genotypes. <i>Potravinarstvo</i> , 2016, 10, .	0.6	18
42	Comparison of 2-de proteome maps of wheat, rye and amaranth. <i>Journal of Microbiology, Biotechnology and Food Sciences</i> , 2015, 04, 7-10.	0.8	1
43	Application of wheat SSR markers for detection of genetic diversity in triticale (<i>x Triticosecale witt.</i>). <i>Genetika</i> , 2015, 47, 983-992.	0.4	1
44	Genetic diversity of <i>glu-1</i> in European wheat genetic resources and varieties. <i>Journal of Microbiology, Biotechnology and Food Sciences</i> , 2015, 04, 23-25.	0.8	5
45	Protein extraction of maize (<i>Zea mays</i>) for proteomic 2-de analysis. <i>Journal of Microbiology, Biotechnology and Food Sciences</i> , 2015, 04, 263-265.	0.8	0
46	STUDY OF DNA POLYMORPHISM OF THE CASTOR NEW LINES BASED ON RAPD MARKERS. <i>Journal of Microbiology, Biotechnology and Food Sciences</i> , 2015, 4, 125-127.	0.8	3
47	Identification of technologically important genes and their products in the collection of bread wheat genotypes. <i>Journal of Microbiology, Biotechnology and Food Sciences</i> , 2015, 04, 26-29.	0.8	4
48	The differences between the old and new barley varieties on the basis of hordein polymorphism with respect to qualitative parameters. <i>Journal of Microbiology, Biotechnology and Food Sciences</i> , 2015, 04, 108-110.	0.8	1
49	Assessment of rapd polymorphism in rye (<i>Secale cereale</i> L.) genotypes. <i>Journal of Microbiology, Biotechnology and Food Sciences</i> , 2015, 04, 94-97.	0.8	4
50	Bioinformatic approach in the identification of arabidopsis gene homologous in amaranthus. <i>Potravinarstvo</i> , 2015, 9, .	0.6	1
51	Identification and differentiation of <i>Ricinus communis</i> L. using SSR markers. <i>Potravinarstvo</i> , 2015, 9, 556-561.	0.6	10
52	Molecular marker-based characterization of a set of wheat genotypes adapted to Central Europe. <i>Cereal Research Communications</i> , 2014, 42, 189-198.	1.6	4
53	Detection and characterisation of Plum pox virus (PPV) isolates from Eastern Slovakia revealed the presence of three main viral strains.. <i>Potravinarstvo</i> , 2014, 8, 1-7.	0.6	0
54	Protein maps of buckwheat and amaranth. <i>Current Opinion in Biotechnology</i> , 2013, 24, S133.	6.6	0

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55	Detection of genetic relationships among spring and winter triticale (<i>Triticosecale</i> Witt.) and rye cultivars (<i>Secale cereale</i> L.) by using retrotransposon-based markers. <i>Czech Journal of Genetics and Plant Breeding</i> , 2013, 49, 171-174.	0.8	8
56	Identification of <i>Triticum aestivum</i> L., <i>Triticum spelta</i> L. and <i>Triticum durum</i> DESF. genotypes on the HMW-GS base. <i>Plant, Soil and Environment</i> , 2010, 56, 82-86.	2.2	4
57	Stress-induced expression of cucumber chitinase and <i>Nicotiana plumbaginifolia</i> β -1,3-glucanase genes in transgenic potato plants. <i>Acta Physiologiae Plantarum</i> , 2007, 29, 133-141.	2.1	20
58	Variation in HMW glutenin subunits of different species of wheat. <i>Plant, Soil and Environment</i> , 2002, 48, 15-19.	2.2	4
59	Effect of Structural Variability of the Ceramide Part on the Saccharide-Ceramide Linkage in Model Glycolipids Studied by Molecular Mechanics and Molecular Dynamics Methods. <i>Collection of Czechoslovak Chemical Communications</i> , 1996, 61, 1405-1431.	1.0	0
60	Genetic divergence in Tunisian castor bean genotypes based on trap markers. <i>Potravinarstvo</i> , 0, 14, 510-518.	0.6	2