

Giuseppina Logozzo

List of Publications by Year in descending order

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17
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687363

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#	ARTICLE	IF	CITATIONS
1	Root Morphology, Allometric Relations and Rhizosheath of Ancient and Modern Tetraploid Wheats (<i>Triticum durum</i> Desf.) in Response to Inoculation with <i>Trichoderma harzianum</i> T-22. <i>Plants</i> , 2022, 11, 159.	3.5	10
2	Towards the Development, Maintenance and Standardized Phenotypic Characterization of Singleâ€Seedâ€Descent Genetic Resources for Chickpea. <i>Current Protocols</i> , 2022, 2, e371.	2.9	6
3	Seed Coating with <i>Trichoderma harzianum</i> T-22 of Italian Durum Wheat Increases Protection against <i>Fusarium culmorum</i> -Induced Crown Rot. <i>Agriculture (Switzerland)</i> , 2022, 12, 714.	3.1	5
4	Understanding photothermal interactions will help expand production range and increase genetic diversity of lentil (<i>Lens culinaris</i> Medik.). <i>Plants People Planet</i> , 2021, 3, 171-181.	3.3	26
5	Intelligent Characterization of Lentil Genetic Resources: Evolutionary History, Genetic Diversity of Germplasm, and the Need for Wellâ€Represented Collections. <i>Current Protocols</i> , 2021, 1, e134.	2.9	18
6	Characterization of Nutritional Quality Traits of a Common Bean Germplasm Collection. <i>Foods</i> , 2021, 10, 1572.	4.3	20
7	Response of Two Local Common Bean Ecotypes of â€Fagioli di Sarconiâ€PGI (<i>Phaseolus vulgaris</i> L.) to Seed-Borne Pathogens and Environmental Change. <i>Agronomy</i> , 2021, 11, 1924.	3.0	2
8	The INCREASE project: Intelligent Collections of foodâ€legume genetic resources for European agrofood systems. <i>Plant Journal</i> , 2021, 108, 646-660.	5.7	29
9	Pod indehiscence in common bean is associated with the fine regulation of <i>PvMYB26</i> . <i>Journal of Experimental Botany</i> , 2021, 72, 1617-1633.	4.8	29
10	Evolution of SSR diversity from wild types to U.S. advanced cultivars in the Andean and Mesoamerican domestications of common bean (<i>Phaseolus vulgaris</i>). <i>PLoS ONE</i> , 2019, 14, e0211342.	2.5	39
11	Molecular Genotyping (SSR) and Agronomic Phenotyping for Utilization of Durum Wheat (<i>Triticum</i>) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 Varieties. <i>Genes</i> , 2018, 9, 465.	2.4	36
12	Molecular analysis of the parallel domestication of the common bean (<i>Phaseolus</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	7.3	240
13	Identification and Characterization of a Homologue to the Arabidopsis INDEHISCENT Gene in Common Bean. <i>Journal of Heredity</i> , 2013, 104, 273-286.	2.4	39
14	Evidence for Introduction Bottleneck and Extensive Inter-Gene Pool (Mesoamerica x Andes) Hybridization in the European Common Bean (<i>Phaseolus vulgaris</i> L.) Germplasm. <i>PLoS ONE</i> , 2013, 8, e75974.	2.5	50
15	Mesoamerican origin of the common bean (<i>Phaseolus vulgaris</i> L.) is revealed by sequence data. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, E788-96.	7.1	327
16	Landraces in Inland areas of the Basilicata region, Italy: monitoring and perspectives for on farm conservation. <i>Genetic Resources and Crop Evolution</i> , 2012, 59, 701-716.	1.6	26
17	Analysis of the contribution of Mesoamerican and Andean gene pools to European common bean (<i>Phaseolus vulgaris</i> L.) germplasm and strategies to establish a core collection. <i>Genetic Resources and Crop Evolution</i> , 2007, 54, 1763-1779.	1.6	63