

Elisa Bellucci

List of Publications by Year in descending order

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Version: 2024-02-01

29
papers

1,845
citations

394421

19
h-index

526287

27
g-index

31
all docs

31
docs citations

31
times ranked

1617
citing authors

#	ARTICLE	IF	CITATIONS
1	Mesoamerican origin of the common bean (<i>Phaseolus vulgaris</i> L.) is revealed by sequence data. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, E788-96.	7.1	327
2	Molecular analysis of the parallel domestication of the common bean (<i>Phaseolus vulgaris</i>). <i>Overlook</i> , 2010, 10, 240.	7.3	240
3	Beans (<i>Phaseolus</i> spp.) as a Model for Understanding Crop Evolution. <i>Frontiers in Plant Science</i> , 2017, 8, 722.	3.6	177
4	Linkage disequilibrium and population structure in wild and domesticated populations of <i>Phaseolus vulgaris</i> L.. <i>Evolutionary Applications</i> , 2009, 2, 504-522.	3.1	139
5	Cytogenetic map of common bean (<i>Phaseolus vulgaris</i> L.). <i>Chromosome Research</i> , 2010, 18, 487-502.	2.2	108
6	Decreased Nucleotide and Expression Diversity and Modified Coexpression Patterns Characterize Domestication in the Common Bean. <i>Plant Cell</i> , 2014, 26, 1901-1912.	6.6	103
7	Landscape genetics, adaptive diversity and population structure in <i>Phaseolus vulgaris</i> . <i>New Phytologist</i> , 2016, 209, 1781-1794.	7.3	86
8	Tagging the Signatures of Domestication in Common Bean (<i>Phaseolus vulgaris</i>) by Means of Pooled DNA Samples. <i>Annals of Botany</i> , 2007, 100, 1039-1051.	2.9	84
9	Genomics of Origin, Domestication and Evolution of <i>Phaseolus vulgaris</i> . , 2014, , 483-507.		60
10	Genomic dissection of pod shattering in common bean: mutations at non-orthologous loci at the basis of convergent phenotypic evolution under domestication of leguminous species. <i>Plant Journal</i> , 2019, 97, 693-714.	5.7	54
11	Evidence for Introduction Bottleneck and Extensive Inter-Gene Pool (Mesoamerica x Andes) Hybridization in the European Common Bean (<i>Phaseolus vulgaris</i> L.) <i>Germplasm. PLoS ONE</i> , 2013, 8, e75974.	2.5	50
12	A Comprehensive Phenotypic Investigation of the "Pod-Shattering Syndrome" in Common Bean. <i>Frontiers in Plant Science</i> , 2017, 8, 251.	3.6	47
13	Population Structure of Barley Landrace Populations and Gene-Flow with Modern Varieties. <i>PLoS ONE</i> , 2013, 8, e83891.	2.5	42
14	Convergent Evolution of the Seed Shattering Trait. <i>Genes</i> , 2019, 10, 68.	2.4	41
15	European <i>Phaseolus coccineus</i> L. landraces: Population Structure and Adaptation, as Revealed by cpSSRs and Phenotypic Analyses. <i>PLoS ONE</i> , 2013, 8, e57337.	2.5	31
16	Ancient genomes reveal early Andean farmers selected common beans while preserving diversity. <i>Nature Plants</i> , 2021, 7, 123-128.	9.3	29
17	The INCREASE project: Intelligent Collections of food-legume genetic resources for European agrofood systems. <i>Plant Journal</i> , 2021, 108, 646-660.	5.7	29
18	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

#	ARTICLE	IF	CITATIONS
19	Biodiversity studies in <i>Phaseolus</i> species by DNA barcoding. <i>Genome</i> , 2011, 54, 529-545.	2.0	27
20	Co-evolution in a landrace meta-population: two closely related pathogens interacting with the same host can lead to different adaptive outcomes. <i>Scientific Reports</i> , 2015, 5, 12834.	3.3	27
21	Adaptation to novel environments during crop diversification. <i>Current Opinion in Plant Biology</i> , 2020, 56, 203-217.	7.1	22
22	Characterization of Nutritional Quality Traits of a Common Bean Germplasm Collection. <i>Foods</i> , 2021, 10, 1572.	4.3	20
23	High Level of Nonsynonymous Changes in Common Bean Suggests That Selection under Domestication Increased Functional Diversity at Target Traits. <i>Frontiers in Plant Science</i> , 2016, 7, 2005.	3.6	19
24	Towards the Development, Maintenance, and Standardized Phenotypic Characterization of Single-Seed-Descent Genetic Resources for Common Bean. <i>Current Protocols</i> , 2021, 1, e133.	2.9	13
25	European Flint Landraces Grown In Situ Reveal Adaptive Introgression from Modern Maize. <i>PLoS ONE</i> , 2015, 10, e0121381.	2.5	11
26	Towards Development, Maintenance, and Standardized Phenotypic Characterization of Single-Seed-Descent Genetic Resources for Lupins. <i>Current Protocols</i> , 2021, 1, e191.	2.9	9
27	The Development of a European and Mediterranean Chickpea Association Panel (EMCAP). <i>Agronomy</i> , 2020, 10, 1417.	3.0	7
28	Domestication and Crop History. <i>Compendium of Plant Genomes</i> , 2017, , 21-55.	0.5	5
29	Sustainable Crop Production. , 2020, , 583-600.		2