Alexander Yermishin

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3883003/publications.pdf

Version: 2024-02-01

1684188 1720034 9 49 5 7 citations g-index h-index papers 9 9 9 29 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Development of chromosome-specific markers for a study on introgressive hybridization of potato with the wild Mexican allotetraploid species Solanum stoloniferum Schltdl. Plant Biotechnology and Breeding, 2020, 2, 24-35.	2.0	2
2	Marker assisted selection of potato breeding lines with combination of PVY resistance genes from different wild species. Plant Biotechnology and Breeding, 2020, 2, 6-14.	2.0	2
3	Overcoming unilateral incompatibility in crosses with wild allotetraploid potato species Solanum stoloniferum Schldtl. & Duchet. Euphytica, 2017, 213, 1.	1.2	7
4	SvSv-lines is an effective tool for involvement of the valuable genepool of 1 EBN diploid potato species into breeding. Vavilovskii Zhurnal Genetiki I Selektsii, 2017, 21, 42-50.	1.1	3
5	Determination of the composition and the allelic state of disease and pest resistance genes in potato parental lines using DNA markers. Russian Journal of Genetics, 2016, 52, 498-506.	0.6	7
6	Production of Hybrids Between the 2EBN Bridge Species Solanum verrucosum and 1EBN diploid Potato Species. American Journal of Potato Research, 2014, 91, 610-617.	0.9	13
7	Production of potato breeding material using somatic hybrids between Solanum tuberosum L. dihaploids and the wild diploid species Solanum bulbocastanum Dunal. from Mexico. Russian Journal of Genetics, 2008, 44, 559-566.	0.6	5
8	Diploid hybrids between allotetraploid wild potato species Solanum acaule Bitt., S. stoloniferum Schltdl. and dihaploids of S. tuberosum L Russian Journal of Genetics, 2007, 43, 882-889.	0.6	4
9	Application of somatic hybrids between dihaploids of potato Solanum tuberosum L. and wild diploid species from Mexico in breeding: Generation and backcrossing of dihaploids of somatic hybrids. Russian Journal of Genetics, 2006, 42, 1414-1421.	0.6	6