Andrea D Wolfe

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3687398/publications.pdf

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23 papers

2,081 citations

706676 14 h-index 759306 22 g-index

26 all docs

 $\begin{array}{c} 26 \\ \text{docs citations} \end{array}$

times ranked

26

2334 citing authors

#	Article	IF	CITATIONS
1	Phylogeographic analysis of shrubby beardtongues reveals range expansions during the Last Glacial Maximum and implicates the Klamath Mountains as a hotspot for hybridization. Molecular Ecology, 2021, 30, 3826-3839.	2.0	8
2	Back to the future of a rare plant species of the Chihuahuan desert: tracing distribution patterns across time and genetic diversity as a basis for conservation actions. Biodiversity and Conservation, 2020, 29, 1821-1840.	1.2	6
3	Genetic diversity and population structure in Cary's Beardtongue Penstemon caryi (Plantaginaceae), a rare plant endemic to the eastern Rocky Mountains of Wyoming and Montana. Conservation Genetics, 2019, 20, 1149-1161.	0.8	12
4	Penstemon reidmoranii (Plantaginaceae), a new species from Baja California, Mexico. Phytotaxa, 2019, 387, 63.	0.1	1
5	HyDe: A Python Package for Genome-Scale Hybridization Detection. Systematic Biology, 2018, 67, 821-829.	2.7	177
6	Investigating the genetic diversity and differentiation patterns in the Penstemon scariosus species complex under different sample sizes using AFLPs and SSRs. Conservation Genetics, 2018, 19, 1335-1348.	0.8	18
7	Fluidigm2 <scp>PURC</scp> : Automated processing and haplotype inference for doubleâ€barcoded <scp>PCR</scp> amplications in Plant Sciences, 2018, 6, e01156.	0.8	12
8	Population genetics of Penstemon albomarginatus (Plantaginaceae), a rare Mojave Desert species of conservation concern. Conservation Genetics, 2016, 17, 1245-1255.	0.8	11
9	Accounting for genotype uncertainty in the estimation of allele frequencies in autopolyploids. Molecular Ecology Resources, 2016, 16, 742-754.	2.2	40
10	Gene Prediction and Annotation inPenstemon(Plantaginaceae): A Workflow for Marker Development from Extremely Low-Coverage Genome Sequencing. Applications in Plant Sciences, 2014, 2, 1400044.	0.8	15
11	Sequenceâ€related amplified polymorphism (SRAP) markers: A potential resource for studies in plant molecular biology ¹ . Applications in Plant Sciences, 2014, 2, 1400017.	0.8	110
12	Conservation genetics and breeding system of Penstemon debilis (Plantaginaceae), a rare beardtongue endemic to oil shale talus in western Colorado, USA. Journal of Systematics and Evolution, 2014, 52, 598-611.	1.6	14
13	Constrained lability in floral evolution: counting convergent origins of hummingbird pollination in <i>Penstemon</i> and <i>Keckiella</i> . New Phytologist, 2007, 176, 883-890.	3.5	116
14	Phylogeny, taxonomic affinities, and biogeography of <i>Penstemon</i> (Plantaginaceae) based on ITS and cpDNA sequence data. American Journal of Botany, 2006, 93, 1699-1713.	0.8	81
15	Phylogeny and biogeography of Orobanchaceae. Folia Geobotanica, 2005, 40, 115-134.	0.4	95
16	ISSR Techniques for Evolutionary Biology. Methods in Enzymology, 2005, 395, 134-144.	0.4	61
17	Relationships among the Macaronesian members of Tolpis (Asteraceae: Lactuceae) based upon analyses of inter simple sequence repeat (ISSR) markers. Taxon, 2003, 52, 511-518.	0.4	39
18	Intersimple sequence repeat (ISSR) variation in Lactoris fernandeziana (Lactoridaceae), a rare endemic of the Juan Fernandez Archipelago, Chile. Plant Species Biology, 2001, 16, 185-192.	0.6	19

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#	Article	IF	CITATIONS
19	Population genetic structure of the cleistogamous plant species Viola pubescens Aiton (Violaceae), as indicated by allozyme and ISSR molecular markers. Heredity, 2001, 86, 545-556.	1.2	128
20	Disintegration of the Scrophulariaceae. American Journal of Botany, 2001, 88, 348-361.	0.8	523
21	Maize R2R3 Myb Genes: Sequence Analysis Reveals Amplification in the Higher Plants. Genetics, 1999, 153, 427-444.	1.2	150
22	Assessing hybridization in natural populations of Penstemon (Scrophulariaceae) using hypervariable intersimple sequence repeat (ISSR) bands. Molecular Ecology, 1998, 7, 1107-1125.	2.0	329
23	ORIGIN AND BIOGEOGRAPHY OF <i>AESCULUS</i> L. (HIPPOCASTANACEAE): A MOLECULAR PHYLOGENETIC PERSPECTIVE. Evolution; International Journal of Organic Evolution, 1998, 52, 988-997.	1.1	97