

# Jennifer A Tate

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/66137/publications.pdf>

Version: 2024-02-01

44  
papers

2,601  
citations

361413

20  
h-index

265206

42  
g-index

44  
all docs

44  
docs citations

44  
times ranked

2562  
citing authors

#	ARTICLE	IF	CITATIONS
1	Advances in the study of polyploidy since <i>Plant speciation</i> . <i>New Phytologist</i> , 2004, 161, 173-191.	7.3	640
2	Recent and recurrent polyploidy in <i>Tragopogon</i> (Asteraceae): cytogenetic, genomic and genetic comparisons. <i>Biological Journal of the Linnean Society</i> , 2004, 82, 485-501.	1.6	328
3	Transcriptomic Shock Generates Evolutionary Novelty in a Newly Formed, Natural Allopolyploid Plant. <i>Current Biology</i> , 2011, 21, 551-556.	3.9	192
4	Rapid Chromosome Evolution in Recently Formed Polyploids in <i>Tragopogon</i> (Asteraceae). <i>PLoS ONE</i> , 2008, 3, e3353.	2.5	173
5	Evolution and Expression of Homeologous Loci in <i>Tragopogon miscellus</i> (Asteraceae), a Recent and Reciprocally Formed Allopolyploid. <i>Genetics</i> , 2006, 173, 1599-1611.	2.9	166
6	Rapid, Repeated, and Clustered Loss of Duplicate Genes in Allopolyploid Plant Populations of Independent Origin. <i>Current Biology</i> , 2012, 22, 248-252.	3.9	159
7	On the road to diploidization? Homoeolog loss in independently formed populations of the allopolyploid <i>Tragopogon miscellus</i> (Asteraceae). <i>BMC Plant Biology</i> , 2009, 9, 80.	3.6	98
8	Polyploidy in Plants. , 2005, , 371-426.		83
9	Synthetic polyploids of <i>Tragopogon miscellus</i> and <i>T. mirus</i> (Asteraceae): 60 Years after Ownbey's discovery. <i>American Journal of Botany</i> , 2009, 96, 979-988.	1.7	70
10	Phylogenetic relationships within the tribe Malveae (Malvaceae, subfamily Malvoideae) as inferred from ITS sequence data. <i>American Journal of Botany</i> , 2005, 92, 584-602.	1.7	69
11	Similar patterns of rDNA evolution in synthetic and recently formed natural populations of <i>Tragopogon</i> (Asteraceae) allotetraploids. <i>BMC Evolutionary Biology</i> , 2010, 10, 291.	3.2	62
12	Cytonuclear responses to genome doubling. <i>American Journal of Botany</i> , 2017, 104, 1277-1280.	1.7	62
13	The biogeography of <i>Hoffmannseggia</i> (Leguminosae, Caesalpinioideae, Caesalpinieae): a tale of many travels. <i>Journal of Biogeography</i> , 2004, 32, 15-27.	3.0	61
14	<i>Andeimalva</i> , a New Genus of Malvaceae from Andean South America. <i>Lundellia</i> , 2003, 6, 10-18.	0.1	58
15	Concerted Evolution of rDNA in Recently Formed <i>Tragopogon</i> Allotetraploids Is Typically Associated With an Inverse Correlation Between Gene Copy Number and Expression. <i>Genetics</i> , 2007, 176, 2509-2519.	2.9	58
16	New targets acquired: Improving locus recovery from the Angiosperms353 probe set. <i>Applications in Plant Sciences</i> , 2021, 9, .	2.1	36
17	Exploring Angiosperms353: An open, community toolkit for collaborative phylogenomic research on flowering plants. <i>American Journal of Botany</i> , 2021, 108, 1059-1065.	1.7	36
18	Breeding system evolution in <i>Tarasa</i> (Malvaceae) and selection for reduced pollen grain size in the polyploid species. <i>American Journal of Botany</i> , 2004, 91, 207-213.	1.7	35

#	ARTICLE	IF	CITATIONS
19	Cytonuclear Coordination Is Not Immediate upon Allopolyploid Formation in <i>Tragopogon miscellus</i> (Asteraceae) Allopolyploids. <i>PLoS ONE</i> , 2015, 10, e0144339.	2.5	31
20	Gene silencing via DNA methylation in naturally occurring <i>Tragopogon miscellus</i> (Asteraceae) allopolyploids. <i>BMC Genomics</i> , 2014, 15, 701.	2.8	25
21	Phylogeny and Character Evolution in <i>Nierembergia</i> (Solanaceae): Molecular, Morphological, and Cytogenetic Evidence. <i>Systematic Botany</i> , 2009, 34, 198-206.	0.5	21
22	Polyploidy on Islands: Its Emergence and Importance for Diversification. <i>Frontiers in Plant Science</i> , 2021, 12, 637214.	3.6	19
23	The compounding effects of high pollen limitation, selfing rates and inbreeding depression leave a New Zealand tree with few viable offspring. <i>Annals of Botany</i> , 2015, 116, 833-843.	2.9	17
24	Phylogeny and Character Evolution of <i>Hoffmannseggia</i> (Caesalpinieae: Caesalpinioideae: Leguminosae). <i>Systematic Botany</i> , 2004, 29, 933-946.	0.5	14
25	Exploring Angiosperms353: Developing and applying a universal toolkit for flowering plant phylogenomics. <i>Applications in Plant Sciences</i> , 2021, 9, .	2.1	13
26	Ribosomal RNA genes evolution in <i>Tragopogon</i> : A story of New and Old World allotetraploids and the synthetic lines. <i>Taxon</i> , 2011, 60, 348-354.	0.7	11
27	Evolutionary significance of long-distance dispersal and hybridisation in the New Zealand endemic genus <i>Hoheria</i> (Malvaceae). <i>Australian Systematic Botany</i> , 2010, 23, 112.	0.9	8
28	Microsatellite Markers for the New Zealand Endemic <i>Myosotis pygmaea</i> Species Group (Boraginaceae) Amplify Across Species. <i>Applications in Plant Sciences</i> , 2015, 3, 1500027.	2.1	8
29	Bolstering Species Delimitation in Difficult Species Complexes by Analyzing Herbarium and Common Garden Morphological Data: A Case Study Using the New Zealand Native <i>Myosotis pygmaea</i> Species Group (Boraginaceae). <i>Systematic Botany</i> , 2018, 43, 266-289.	0.5	8
30	Comparing and analysing microsatellite and morphological data for species delimitation in the New Zealand native <i>Myosotis pygmaea</i> species group (Boraginaceae). <i>Taxon</i> , 2019, 68, 731-750.	0.7	6
31	Host range, host specificity, regional host preferences and genetic variability of <i>Korthalsella</i> Tiegh. (Viscaceae) mistletoes in New Zealand. <i>New Zealand Journal of Botany</i> , 2018, 56, 127-162.	1.1	5
32	The status of <i>Urocarpidium</i> (Malvaceae): Insight from nuclear and plastid-based phylogenies. <i>Taxon</i> , 2011, 60, 1330-1338.	0.7	4
33	Phylogeny and Character Evolution in the New Zealand Endemic Genus <i>Plagianthus</i> (Malveae). <i>Tj ETQq1 1 0,784314 rgBT /Ove</i>	0.5	4
34	<i>Sophora microphylla</i> (Fabaceae) Microsatellite Markers and their Utility Across the Genus. <i>Applications in Plant Sciences</i> , 2014, 2, 1300081.	2.1	4
35	A new record of <i>Abutilon grandifolium</i> (subfamily Malvoideae, Malvaceae) from Qena Governorate, Egypt. <i>Phytotaxa</i> , 2018, 382, 255.	0.3	3
36	Polyploidy on islands – concerted evolution and gene loss amid chromosomal stasis. <i>Annals of Botany</i> , 2023, 131, 33-44.	2.9	3

#	ARTICLE	IF	CITATIONS
37	Microsatellite markers for the endangered root holoparasite <i>Dactylanthus taylorii</i> (Balanophoraceae) from 454 pyrosequencing. <i>American Journal of Botany</i> , 2012, 99, e323-5.	1.7	2
38	Microsatellite Markers for the New Zealand Endemic Tree <i>Fuchsia excorticata</i> (Onagraceae). <i>Applications in Plant Sciences</i> , 2013, 1, 1300045.	2.1	2
39	Pollinator service affects quantity but not quality of offspring in a widespread New Zealand endemic tree species. <i>Conservation Genetics</i> , 2018, 19, 815-826.	1.5	2
40	Inferring the biogeography of New Zealand's only endemic holoparasitic plant, the threatened <i>Dactylanthus taylorii</i> (Mystropetalaceae). <i>New Zealand Journal of Botany</i> , 2022, 60, 331-353.	1.1	2
41	Microsatellite markers for <i>Corybas</i> (Orchidaceae) species in New Zealand. <i>Applications in Plant Sciences</i> , 2018, 6, e01192.	2.1	1
42	Widespread morphological parallelism in <i>Korthalsella</i> (Santalaceae, tribe Visceae): A molecular phylogenetic perspective. <i>Taxon</i> , 2019, 68, 1204-1218.	0.7	1
43	Taxonomic revision of the southern hemisphere pygmy forget-me-not group ( <i>Myosotis</i> ; Boraginaceae) based on morphological, population genetic and climate-edaphic niche modelling data. <i>Australian Systematic Botany</i> , 2022, 35, 63.	0.9	1
44	Can morphology and chromosome number contribute to species delimitation? A case in the <i>Hibiscus trionum</i> complex (Tribe Hibisceae.) <i>Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50.457 Td (Malvaceae)</i>	0.45	1