

# Alan E Yocca

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2861585/publications.pdf>

Version: 2024-02-01

12  
papers

984  
citations

1040056

9  
h-index

1125743

13  
g-index

18  
all docs

18  
docs citations

18  
times ranked

1246  
citing authors

#	ARTICLE	IF	CITATIONS
1	Origin and evolution of the octoploid strawberry genome. <i>Nature Genetics</i> , 2019, 51, 541-547.	21.4	469
2	Haplotype-phased genome and evolution of phytonutrient pathways of tetraploid blueberry. <i>GigaScience</i> , 2019, 8, .	6.4	167
3	Exceptional subgenome stability and functional divergence in the allotetraploid Ethiopian cereal teff. <i>Nature Communications</i> , 2020, 11, 884.	12.8	101
4	Reply to: Revisiting the origin of octoploid strawberry. <i>Nature Genetics</i> , 2020, 52, 5-7.	21.4	44
5	Evolutionary history and pan-genome dynamics of strawberry ( <i>Fragaria</i> spp.). <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	43
6	A chromosome-scale assembly of the bilberry genome identifies a complex locus controlling berry anthocyanin composition. <i>Molecular Ecology Resources</i> , 2022, 22, 345-360.	4.8	28
7	There and back again; historical perspective and future directions for <i>Vaccinium</i> breeding and research studies. <i>Horticulture Research</i> , 2022, 9, .	6.3	27
8	Disease Resistance Genetics and Genomics in Octoploid Strawberry. <i>G3: Genes, Genomes, Genetics</i> , 2019, 9, 3315-3332.	1.8	26
9	Current status and future perspectives on the evolution of cis-regulatory elements in plants. <i>Current Opinion in Plant Biology</i> , 2022, 65, 102139.	7.1	19
10	Evolution of Conserved Noncoding Sequences in <i>Arabidopsis thaliana</i> . <i>Molecular Biology and Evolution</i> , 2021, 38, 2692-2703.	8.9	14
11	<i>Linnemannia elongata</i> (Mortierellaceae) stimulates <i>Arabidopsis thaliana</i> aerial growth and responses to auxin, ethylene, and reactive oxygen species. <i>PLoS ONE</i> , 2022, 17, e0261908.	2.5	10
12	Machine learning approaches to identify core and dispensable genes in pangenomes. <i>Plant Genome</i> , 2022, 15, e20135.	2.8	4