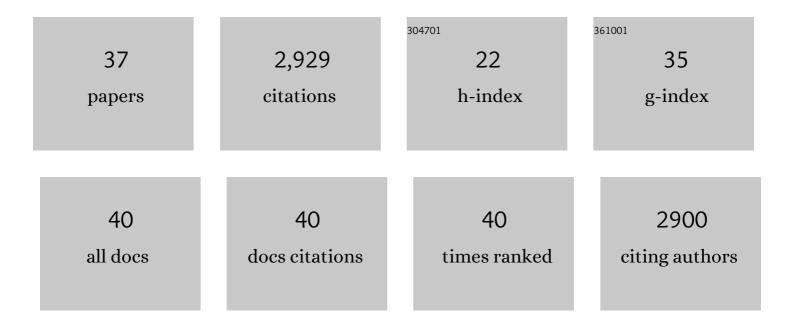
## Nick W Albert

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6501141/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	A Conserved Network of Transcriptional Activators and Repressors Regulates Anthocyanin Pigmentation in Eudicots. Plant Cell, 2014, 26, 962-980.	6.6	610
2	Members of an R2R3â€MYB transcription factor family in <i>Petunia</i> are developmentally and environmentally regulated to control complex floral and vegetative pigmentation patterning. Plant Journal, 2011, 65, 771-784.	5.7	401
3	From landing lights to mimicry: the molecular regulation of flower colouration and mechanisms for pigmentation patterning. Functional Plant Biology, 2012, 39, 619.	2.1	263
4	Light-induced vegetative anthocyanin pigmentation in Petunia. Journal of Experimental Botany, 2009, 60, 2191-2202.	4.8	256
5	The Evolution of Flavonoid Biosynthesis: A Bryophyte Perspective. Frontiers in Plant Science, 2020, 11, 7.	3.6	126
6	Genetic analysis of the liverwort <i>Marchantia polymorpha</i> reveals that R2R3 <scp>MYB</scp> activation of flavonoid production in response to abiotic stress is an ancient character in land plants. New Phytologist, 2018, 218, 554-566.	7.3	98
7	UVR8â€mediated induction of flavonoid biosynthesis for UVB tolerance is conserved between the liverwort <i>Marchantia polymorpha</i> and flowering plants. Plant Journal, 2018, 96, 503-517.	5.7	93
8	The Onion (Allium cepa L.) R2R3-MYB Gene MYB1 Regulates Anthocyanin Biosynthesis. Frontiers in Plant Science, 2016, 7, 1865.	3.6	91
9	Subspecialization of R2R3-MYB Repressors for Anthocyanin and Proanthocyanidin Regulation in Forage Legumes. Frontiers in Plant Science, 2015, 6, 1165.	3.6	70
10	Whole genome sequencing of enriched chloroplast DNA using the Illumina GAII platform. Plant Methods, 2010, 6, 22.	4.3	67
11	Auronidins are a previously unreported class of flavonoid pigments that challenges when anthocyanin biosynthesis evolved in plants. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 20232-20239.	7.1	63
12	Anthocyanin leaf markings are regulated by a family of <i>R2R3â€MYB</i> genes in the genus <i><scp>T</scp>rifolium</i> . New Phytologist, 2015, 205, 882-893.	7.3	62
13	Epigenetics in plants—vernalisation and hybrid vigour. Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms, 2011, 1809, 427-437.	1.9	61
14	Gene regulation networks generate diverse pigmentation patterns in plants. Plant Signaling and Behavior, 2014, 9, e29526.	2.4	58
15	MYBA and MYBPA transcription factors coâ€regulate anthocyanin biosynthesis in blueâ€coloured berries. New Phytologist, 2021, 232, 1350-1367.	7.3	56
16	MYBA From Blueberry (Vaccinium Section Cyanococcus) Is a Subgroup 6 Type R2R3MYB Transcription Factor That Activates Anthocyanin Production. Frontiers in Plant Science, 2018, 9, 1300.	3.6	55
17	Aromatic Decoration Determines the Formation of Anthocyanic Vacuolar Inclusions. Current Biology, 2017, 27, 945-957.	3.9	49
18	Spatiotemporal Modulation of Flavonoid Metabolism in Blueberries. Frontiers in Plant Science, 2020, 11, 545.	3.6	42

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19	Temporal and spatial regulation of anthocyanin biosynthesis provide diverse flower colour intensities and patterning in Cymbidium orchid. Planta, 2014, 240, 983-1002.	3.2	39
20	Activation of anthocyanin synthesis in Cymbidium orchids: variability between known regulators. Plant Cell, Tissue and Organ Culture, 2010, 100, 355-360.	2.3	36
21	CRISPR-Cas9 enrichment and long read sequencing for fine mapping in plants. Plant Methods, 2020, 16, 121.	4.3	31
22	Discrete bHLH transcription factors play functionally overlapping roles in pigmentation patterning in flowers of <i>Antirrhinum majus</i> . New Phytologist, 2021, 231, 849-863.	7.3	28
23	A chromosomeâ€scale assembly of the bilberry genome identifies a complex locus controlling berry anthocyanin composition. Molecular Ecology Resources, 2022, 22, 345-360.	4.8	28
24	Infiltration-RNAseq: transcriptome profiling of Agrobacterium-mediated infiltration of transcription factors to discover gene function and expression networks in plants. Plant Methods, 2016, 12, 41.	4.3	26
25	Hierarchical regulation of <i>MYBPA1</i> by anthocyanin- and proanthocyanidin-related MYB proteins is conserved in <i>Vaccinium</i> species. Journal of Experimental Botany, 2022, 73, 1344-1356.	4.8	20
26	Transformation and Regeneration of Petunia. , 2009, , 395-409.		16
27	Changes in 1-aminocyclopropane-1-carboxlate (ACC) oxidase expression and enzyme activity in response to excess manganese in white clover (Trifolium repens L.). Plant Physiology and Biochemistry, 2011, 49, 1013-1019.	5.8	15
28	Failure to launch: the self-regulating Md-MYB10 R6 gene from apple is active in flowers but not leaves of Petunia. Plant Cell Reports, 2015, 34, 1817-1823.	5.6	11
29	Stress, senescence and specialised metabolites in bryophytes. Journal of Experimental Botany, 2022, , .	4.8	11
30	Genotypic variation in sulfur assimilation and metabolism of onion (Allium cepa L.) III. Characterization of sulfite reductase. Phytochemistry, 2012, 83, 34-42.	2.9	10
31	REPRESSION - THE DARK SIDE OF ANTHOCYANIN REGULATION?. Acta Horticulturae, 2014, , 129-136.	0.2	9
32	Flavonoids – flowers, fruit, forage and the future. Journal of the Royal Society of New Zealand, 2023, 53, 304-331.	1.9	9
33	Genotypic variation in sulphur assimilation and metabolism of onion (Allium cepa L.). II: Characterisation of ATP sulphurylase activity. Phytochemistry, 2011, 72, 888-896.	2.9	8
34	The Coordinated Action of MYB Activators and Repressors Controls Proanthocyanidin and Anthocyanin Biosynthesis in Vaccinium. Frontiers in Plant Science, 0, 13, .	3.6	8
35	Identification of a Strong Anthocyanin Activator, VbMYBA, From Berries of Vaccinium bracteatum Thunb Frontiers in Plant Science, 2021, 12, 697212.	3.6	7
36	Control of anthocyanin pigmentation during flower development inCymbidiumorchid. Acta Horticulturae, 2015, , 333-340.	0.2	4

#	Article	IF	CITATIONS
37	LONG-TERM STABLE EXPRESSION OF MULTIPLE TRANSGENES UNDER CONTROL OF THE SAME PROMOTER IN CYMBIDIUM ORCHID. Acta Horticulturae, 2012, , 597-604.	0.2	1