## Steve van Nocker

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Fineâ€ŧuning of SUMOylation modulates drought tolerance of apple. Plant Biotechnology Journal, 2022, 20, 903-919.	8.3	16
2	Metaâ€analysis of apple ( <i>Malus</i> × <i>domestica</i> Borkh.) fruit and juice quality traits for potential use in hard cider production. Plants People Planet, 2022, 4, 463-475.	3.3	4
3	Genetic mechanisms associated with floral initiation and the repressive effect of fruit on flowering in apple (Malus x domestica Borkh). PLoS ONE, 2021, 16, e0245487.	2.5	9
4	Overexpression of the Apple (Malus× domestica) MdERF100 in Arabidopsis Increases Resistance to Powdery Mildew. International Journal of Molecular Sciences, 2021, 22, 5713.	4.1	13
5	Physiological and transcriptome analyses of the effects of exogenous dopamine on drought tolerance in apple. Plant Physiology and Biochemistry, 2020, 148, 260-272.	5.8	65
6	Genetic mechanisms in the repression of flowering by gibberellins in apple (Malus x domestica Borkh.). BMC Genomics, 2019, 20, 747.	2.8	56
7	The jasmonate-ZIM domain gene VqJAZ4 from the Chinese wild grape Vitis quinquangularis improves resistance to powdery mildew in Arabidopsis thaliana. Plant Physiology and Biochemistry, 2019, 143, 329-339.	5.8	21
8	Genome-Wide Analysis of the YABBY Gene Family in Grapevine and Functional Characterization of VvYABBY4. Frontiers in Plant Science, 2019, 10, 1207.	3.6	37
9	Contribution of methylation regulation of MpDREB2A promoter to drought resistance of Mauls prunifolia. Plant and Soil, 2019, 441, 15-32.	3.7	16
10	Apple whole genome sequences: recent advances and new prospects. Horticulture Research, 2019, 6, 59.	6.3	77
11	The grapevine homeobox gene VvHB58 influences seed and fruit development through multiple hormonal signaling pathways. BMC Plant Biology, 2019, 19, 523.	3.6	18
12	Overexpression of a protein kinase gene MpSnRK2.10 from Malus prunifolia confers tolerance to drought stress in transgenic Arabidopsis thaliana and apple. Gene, 2019, 692, 26-34.	2.2	28
13	Overexpression of <i>MdIAA9</i> confers high tolerance to osmotic stress in transgenic tobacco. PeerJ, 2019, 7, e7935.	2.0	11
14	Singleâ€base methylome analysis reveals dynamic epigenomic differences associated with water deficit in apple. Plant Biotechnology Journal, 2018, 16, 672-687.	8.3	130
15	Comprehensive genomic analysis of the TYROSINE AMINOTRANSFERASE (TAT) genes in apple (Malus) Tj ETQq1 1 stresses in plants. Plant Physiology and Biochemistry, 2018, 133, 81-91.	0.784314 5.8	rgBT /Over 16
16	Genome-wide identification and expression analyses of the homeobox transcription factor family during ovule development in seedless and seeded grapes. Scientific Reports, 2017, 7, 12638.	3.3	27
17	Identification and expression analysis of the apple (Malus × domestica) basic helix-loop-helix transcription factor family. Scientific Reports, 2017, 7, 28.	3.3	43
18	Expression of the Grape VaSTS19 Gene in Arabidopsis Improves Resistance to Powdery Mildew and Botrytis cinerea but Increases Susceptibility to Pseudomonas syringe pv Tomato DC3000. International Journal of Molecular Sciences, 2017, 18, 2000.	4.1	16

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19	Improved hybrid de novo genome assembly of domesticated apple (Malus x domestica). GigaScience, 2016, 5, 35.	6.4	56
20	Breeding better cultivars, faster: applications of new technologies for the rapid deployment of superior horticultural tree crops. Horticulture Research, 2014, 1, 14022.	6.3	169
21	Genomic and Gene-Level Distribution of Histone H3 Dimethyl Lysine-27 (H3K27me2) in Arabidopsis. PLoS ONE, 2012, 7, e52855.	2.5	11
22	Gene Expression Associated with Apple Aroma Biosynthesis. Hortscience: A Publication of the American Society for Hortcultural Science, 2006, 41, 977C-977.	1.0	0