

Martijn van Zanten

List of Publications by Year in descending order

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Version: 2024-02-01

45
papers

2,554
citations

257450

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docs citations

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times ranked

3314
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | 2D morphometric analysis of <i>Arabidopsis thaliana</i> nuclei reveals characteristic profiles of different cell types and accessions. <i>Chromosome Research</i> , 2022, 30, 5-24. | 2.2 | 7 |
| 2 | Epigenetic regulation of thermomorphogenesis and heat stress tolerance. <i>New Phytologist</i> , 2022, 234, 1144-1160. | 7.3 | 54 |
| 3 | Genetic diversity reveals synergistic interaction between yield components could improve the sink size and yield in rice. <i>Food and Energy Security</i> , 2022, 11, . | 4.3 | 6 |
| 4 | Plant thermotropism: an underexplored thermal engagement and avoidance strategy. <i>Journal of Experimental Botany</i> , 2021, , . | 4.8 | 4 |
| 5 | The membrane-localized protein kinase MAP4K4/TOT3 regulates thermomorphogenesis. <i>Nature Communications</i> , 2021, 12, 2842. | 12.8 | 30 |
| 6 | The chemical compound Heatin™ stimulates hypocotyl elongation and interferes with the <i>Arabidopsis</i> NIT1 subfamily of nitrilases. <i>Plant Journal</i> , 2021, 106, 1523-1540. | 5.7 | 7 |
| 7 | Protein kinase and phosphatase control of plant temperature responses. <i>Journal of Experimental Botany</i> , 2021, , . | 4.8 | 6 |
| 8 | Examiners™ use of rubric criteria for grading bachelor theses. <i>Assessment and Evaluation in Higher Education</i> , 2021, 46, 1269-1284. | 5.6 | 5 |
| 9 | The diverse and unanticipated roles of histone deacetylase 9 in coordinating plant development and environmental acclimation. <i>Journal of Experimental Botany</i> , 2020, 71, 6211-6225. | 4.8 | 18 |
| 10 | A high throughput method for quantifying number and size distribution of <i>Arabidopsis</i> seeds using large particle flow cytometry. <i>Plant Methods</i> , 2020, 16, 27. | 4.3 | 7 |
| 11 | HISTONE DEACETYLASE 9 stimulates auxin-dependent thermomorphogenesis in <i>Arabidopsis thaliana</i> by mediating H2A.Z depletion. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 25343-25354. | 7.1 | 91 |
| 12 | Root Tropisms: Investigations on Earth and in Space to Unravel Plant Growth Direction. <i>Frontiers in Plant Science</i> , 2019, 10, 1807. | 3.6 | 66 |
| 13 | POWERDRESS-mediated histone deacetylation is essential for thermomorphogenesis in <i>Arabidopsis thaliana</i> . <i>PLoS Genetics</i> , 2018, 14, e1007280. | 3.5 | 99 |
| 14 | Thermosensing Enlightened. <i>Trends in Plant Science</i> , 2017, 22, 185-187. | 8.8 | 32 |
| 15 | Genetic Dissection of Morphometric Traits Reveals That Phytochrome B Affects Nucleus Size and Heterochromatin Organization in <i>Arabidopsis thaliana</i> . <i>G3: Genes, Genomes, Genetics</i> , 2017, 7, 2519-2531. | 1.8 | 14 |
| 16 | Molecular and genetic control of plant thermomorphogenesis. <i>Nature Plants</i> , 2016, 2, 15190. | 9.3 | 432 |
| 17 | Genome-Wide Association Analysis of Adaptation Using Environmentally Predicted Traits. <i>PLoS Genetics</i> , 2015, 11, e1005594. | 3.5 | 7 |
| 18 | Ethylene-Mediated Regulation of A2-Type CYCLINs Modulates Hyponastic Growth in <i>Arabidopsis</i> . <i>Plant Physiology</i> , 2015, 169, 194-208. | 4.8 | 22 |

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|----|--|-----|-----------|
| 19 | HISTONE DEACETYLASE 9 represses seedling traits in <i>Arabidopsis thaliana</i> dry seeds. <i>Plant Journal</i> , 2014, 80, 475-488. | 5.7 | 107 |
| 20 | Environment-Induced Chromatin Reorganisation and Plant Acclimation. <i>Signaling and Communication in Plants</i> , 2013, , 21-40. | 0.7 | 4 |
| 21 | High temperature acclimation through PIF4 signaling. <i>Trends in Plant Science</i> , 2013, 18, 59-64. | 8.8 | 94 |
| 22 | Ethylene promotes hyponastic growth through interaction with ROTUNDIFOLIA3/CYP90C1 in <i>Arabidopsis</i> . <i>Journal of Experimental Botany</i> , 2013, 64, 613-624. | 4.8 | 40 |
| 23 | Antiphase Light and Temperature Cycles Affect PHYTOCHROME B-Controlled Ethylene Sensitivity and Biosynthesis, Limiting Leaf Movement and Growth of <i>Arabidopsis</i> . <i>Plant Physiology</i> , 2013, 163, 882-895. | 4.8 | 28 |
| 24 | Epigenetic Signalling During the Life of Seeds. <i>Signaling and Communication in Plants</i> , 2013, , 127-153. | 0.7 | 13 |
| 25 | Shedding Light on Large-Scale Chromatin Reorganization in <i>Arabidopsis thaliana</i> . <i>Molecular Plant</i> , 2012, 5, 583-590. | 8.3 | 42 |
| 26 | Control and consequences of chromatin compaction during seed maturation in <i>Arabidopsis thaliana</i> . <i>Plant Signaling and Behavior</i> , 2012, 7, 338-341. | 2.4 | 23 |
| 27 | Haemoglobin modulates NO emission and hyponasty under hypoxia-related stress in <i>Arabidopsis thaliana</i> . <i>Journal of Experimental Botany</i> , 2012, 63, 5581-5591. | 4.8 | 108 |
| 28 | Illumina Sequencing Technology as a Method of Identifying T-DNA Insertion Loci in Activation-Tagged <i>Arabidopsis thaliana</i> Plants. <i>Molecular Plant</i> , 2012, 5, 948-950. | 8.3 | 22 |
| 29 | Ethylene-induced differential petiole growth in <i>Arabidopsis thaliana</i> involves local microtubule reorientation and cell expansion. <i>New Phytologist</i> , 2012, 193, 339-348. | 7.3 | 74 |
| 30 | Modulation of ethylene- and heat-controlled hyponastic leaf movement in <i>Arabidopsis thaliana</i> by the plant defence hormones jasmonate and salicylate. <i>Planta</i> , 2012, 235, 677-685. | 3.2 | 15 |
| 31 | Identification of the <i>Arabidopsis</i> REDUCED DORMANCY 2 Gene Uncovers a Role for the Polymerase Associated Factor 1 Complex in Seed Dormancy. <i>PLoS ONE</i> , 2011, 6, e22241. | 2.5 | 77 |
| 32 | Expression of rice SUB1A and SUB1C transcription factors in <i>Arabidopsis</i> uncovers flowering inhibition as a submergence tolerance mechanism. <i>Plant Journal</i> , 2011, 67, 434-446. | 5.7 | 62 |
| 33 | Seed maturation in <i>Arabidopsis thaliana</i> is characterized by nuclear size reduction and increased chromatin condensation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 20219-20224. | 7.1 | 141 |
| 34 | Ethylene-induced hyponastic growth in <i>Arabidopsis thaliana</i> is controlled by ERECTA. <i>Plant Journal</i> , 2010, 61, 83-95. | 5.7 | 39 |
| 35 | Photoreceptors CRYTOCHROME2 and Phytochrome B Control Chromatin Compaction in <i>Arabidopsis</i> . <i>Plant Physiology</i> , 2010, 154, 1686-1696. | 4.8 | 44 |
| 36 | Large-scale chromatin de-compaction induced by low light is not accompanied by nucleosomal displacement. <i>Plant Signaling and Behavior</i> , 2010, 5, 1677-1678. | 2.4 | 11 |

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|----|---|-----|-----------|
| 37 | ERECTA controls low light intensity-induced differential petiole growth independent of Phytochrome B and Cryptochrome 2 action in <i>Arabidopsis thaliana</i> . <i>Plant Signaling and Behavior</i> , 2010, 5, 284-286. | 2.4 | 14 |
| 38 | Kinome Profiling Reveals an Interaction Between Jasmonate, Salicylate and Light Control of Hyponastic Petiole Growth in <i>Arabidopsis thaliana</i> . <i>PLoS ONE</i> , 2010, 5, e14255. | 2.5 | 21 |
| 39 | Auxin perception and polar auxin transport are not always a prerequisite for differential growth. <i>Plant Signaling and Behavior</i> , 2009, 4, 899-901. | 2.4 | 11 |
| 40 | Hormone- and Light-Mediated Regulation of Heat-Induced Differential Petiole Growth in <i>Arabidopsis</i> . <i>Plant Physiology</i> , 2009, 151, 1446-1458. | 4.8 | 78 |
| 41 | PHYTOCHROME B and HISTONE DEACETYLASE 6 Control Light-Induced Chromatin Compaction in <i>Arabidopsis thaliana</i> . <i>PLoS Genetics</i> , 2009, 5, e1000638. | 3.5 | 123 |
| 42 | Differential petiole growth in <i>Arabidopsis thaliana</i> : photocontrol and hormonal regulation. <i>New Phytologist</i> , 2009, 184, 141-152. | 7.3 | 77 |
| 43 | The many functions of ERECTA. <i>Trends in Plant Science</i> , 2009, 14, 214-218. | 8.8 | 187 |
| 44 | Abscisic Acid Antagonizes Ethylene-Induced Hyponastic Growth in <i>Arabidopsis</i> . <i>Plant Physiology</i> , 2007, 143, 1013-1023. | 4.8 | 59 |
| 45 | How to decide? Different methods of calculating gene expression from short oligonucleotide array data will give different results. <i>BMC Bioinformatics</i> , 2006, 7, 137. | 2.6 | 124 |