

# Timothy J Tranbarger

## List of Publications by Year in descending order

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39  
papers

2,288  
citations

304743

22  
h-index

302126

39  
g-index

40  
all docs

40  
docs citations

40  
times ranked

2827  
citing authors

#	ARTICLE	IF	CITATIONS
1	SSR markers in transcripts of genes linked to post-transcriptional and transcriptional regulatory functions during vegetative and reproductive development of <i>Elaeis guineensis</i> . <i>BMC Plant Biology</i> , 2012, 12, 1.	3.6	275
2	Regulation of the nitrate transporter gene <i>AtNRT2.1</i> in <i>Arabidopsis thaliana</i> : responses to nitrate, amino acids and developmental stage. <i>Plant Molecular Biology</i> , 2003, 52, 689-703.	3.9	213
3	Pluripotent versus totipotent plant stem cells: dependence versus autonomy?. <i>Trends in Plant Science</i> , 2007, 12, 245-252.	8.8	211
4	Comparative Transcriptome Analysis of Three Oil Palm Fruit and Seed Tissues That Differ in Oil Content and Fatty Acid Composition. <i>Plant Physiology</i> , 2013, 162, 1337-1358.	4.8	200
5	Regulatory Mechanisms Underlying Oil Palm Fruit Mesocarp Maturation, Ripening, and Functional Specialization in Lipid and Carotenoid Metabolism. <i>Plant Physiology</i> , 2011, 156, 564-584.	4.8	190
6	The soybean 94-kilodalton vegetative storage protein is a lipoxygenase that is localized in paraveinal mesophyll cell vacuoles. <i>Plant Cell</i> , 1991, 3, 973-987.	6.6	184
7	Temporal responses of <i>Arabidopsis</i> root architecture to phosphate starvation: evidence for the involvement of auxin signalling. <i>Plant, Cell and Environment</i> , 2003, 26, 1053-1066.	5.7	135
8	Nitrate-dependent control of root architecture and N nutrition are altered by a plant growth-promoting <i>Phyllobacterium</i> sp. <i>Planta</i> , 2006, 223, 591-603.	3.2	110
9	A new family of lipolytic plant enzymes with members in rice, <i>Arabidopsis</i> and maize. <i>FEBS Letters</i> , 1995, 377, 475-480.	2.8	97
10	The isolation of a novel metallothionein-related cDNA expressed in somatic and zygotic embryos of Douglas-fir: regulation by ABA, osmoticum, and metal ions. <i>Plant Molecular Biology</i> , 1997, 34, 243-254.	3.9	65
11	Improving palm oil quality through identification and mapping of the lipase gene causing oil deterioration. <i>Nature Communications</i> , 2013, 4, 2160.	12.8	62
12	Transcriptome analysis during somatic embryogenesis of the tropical monocot <i>Elaeis guineensis</i> : evidence for conserved gene functions in early development. <i>Plant Molecular Biology</i> , 2009, 70, 173-192.	3.9	59
13	Temporal and spatial expression of polygalacturonase gene family members reveals divergent regulation during fleshy fruit ripening and abscission in the monocot species oil palm. <i>BMC Plant Biology</i> , 2012, 12, 150.	3.6	55
14	Conservation of the abscission signaling peptide IDA during Angiosperm evolution: withstanding genome duplications and gain and loss of the receptors HAE/HSL2. <i>Frontiers in Plant Science</i> , 2015, 6, 931.	3.6	50
15	Post-germination-induced and hormonally dependent expression of low-molecular-weight heat shock protein genes in Douglas fir. <i>Plant Molecular Biology</i> , 1996, 30, 1115-1128.	3.9	35
16	Transcription factor genes with expression correlated to nitrate-related root plasticity of <i>Arabidopsis thaliana</i> . <i>Plant, Cell and Environment</i> , 2003, 26, 459-469.	5.7	34
17	The Soybean 94-Kilodalton Vegetative Storage Protein Is a Lipoxygenase That Is Localized in Paraveinal Mesophyll Cell Vacuoles. <i>Plant Cell</i> , 1991, 3, 973.	6.6	33
18	Acquisition of callogenic capacity in date palm leaf tissues in response to 2,4-D treatment. <i>Plant Cell, Tissue and Organ Culture</i> , 2009, 99, 35-45.	2.3	32

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19	Cellular and Pectin Dynamics during Abscission Zone Development and Ripe Fruit Abscission of the Monocot Oil Palm. <i>Frontiers in Plant Science</i> , 2016, 7, 540.	3.6	32
20	Transcriptome Analysis of Cell Wall and NAC Domain Transcription Factor Genes during <i>Elaeis guineensis</i> Fruit Ripening: Evidence for Widespread Conservation within Monocot and Eudicot Lineages. <i>Frontiers in Plant Science</i> , 2017, 8, 603.	3.6	31
21	Structure and expression of a developmentally regulated cDNA encoding a cysteine protease (pseudotzain) from Douglas fir. <i>Gene</i> , 1996, 172, 221-226.	2.2	28
22	Expression and Accumulation Patterns of Nitrogen-Responsive Lipoxygenase in Soybeans. <i>Plant Physiology</i> , 1993, 103, 457-466.	4.8	24
23	The PIP Peptide of INFLORESCENCE DEFICIENT IN ABSCISSION Enhances <i>Populus</i> Leaf and <i>Elaeis guineensis</i> Fruit Abscission. <i>Plants</i> , 2019, 8, 143.	3.5	22
24	The molecular characterization of a set of cDNAs differentially expressed during Douglas-fir germination and early seedling development. <i>Physiologia Plantarum</i> , 1995, 95, 456-464.	5.2	16
25	Editorial: Plant Organ Abscission: From Models to Crops. <i>Frontiers in Plant Science</i> , 2017, 8, 196.	3.6	15
26	Improvement of the content in bioaccessible lipophilic micronutrients in raw and processed drumstick leaves ( <i>Moringa oleifera</i> Lam.). <i>LWT - Food Science and Technology</i> , 2017, 75, 279-285.	5.2	12
27	Characterization of proteinase activity in stratified Douglas-fir seeds. <i>Tree Physiology</i> , 2001, 21, 625-629.	3.1	10
28	EPIP-Evoked Modifications of Redox, Lipid, and Pectin Homeostasis in the Abscission Zone of Lupine Flowers. <i>International Journal of Molecular Sciences</i> , 2021, 22, 3001.	4.1	8
29	A phenotypic test for delay of abscission and non-abscission oil palm fruit and validation by abscission marker gene expression analysis. <i>Acta Horticulturae</i> , 2016, , 97-104.	0.2	7
30	The acceleration of yellow lupine flower abscission by jasmonates is accompanied by lipid-related events in abscission zone cells. <i>Plant Science</i> , 2022, 316, 111173.	3.6	7
31	Regulation of NADPH-cytochrome P450 reductase expressed during Douglas-fir germination and seedling development. <i>Plant Molecular Biology</i> , 2000, 44, 141-153.	3.9	5
32	Multi-scale comparative transcriptome analysis reveals key genes and metabolic reprogramming processes associated with oil palm fruit abscission. <i>BMC Plant Biology</i> , 2021, 21, 92.	3.6	5
33	A macro-array-based screening approach to identify transcriptional factors involved in the nitrogen-related root plasticity response of <i>Arabidopsis thaliana</i> . <i>Agronomy for Sustainable Development</i> , 2003, 23, 519-528.	0.8	5
34	Environmental and trophic determinism of fruit abscission and outlook with climate change in tropical regions. <i>Plant-Environment Interactions</i> , 2020, 1, 17-28.	1.5	4
35	EPIP as an abscission promoting agent in the phytohormonal pathway. <i>Plant Physiology and Biochemistry</i> , 2022, 178, 137-145.	5.8	4
36	The molecular characterization of a set of cDNAs differentially expressed during Douglas-fir germination and early seedling development. <i>Physiologia Plantarum</i> , 1995, 95, 456-464.	5.2	3

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37	Elaeis oleifera (Kunth) Cortáez: A neglected palm from the Ecuadorian Amazon. Revista Ecuatoriana De Medicina Y Ciencias Biológicas, 2018, 39, .	0.1	3
38	Differentially Regulated Gene Sets in Douglas Fir Seeds and Somatic Embryos.. Forestry Sciences, 1996, , 197-204.	0.4	1
39	A Non-Shedding Fruit Elaeis oleifera Palm Reveals Perturbations to Hormone Signaling, ROS Homeostasis, and Hemicellulose Metabolism. Genes, 2021, 12, 1724.	2.4	1