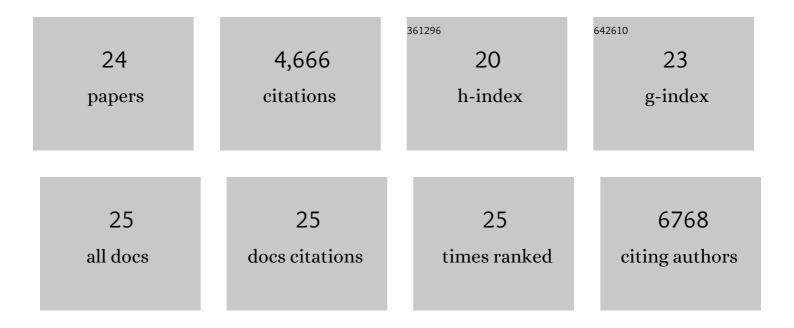
## Henry D Priest

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2873306/publications.pdf Version: 2024-02-01



HENDY D DDIEST

#	Article	IF	CITATIONS
1	The genome of woodland strawberry (Fragaria vesca). Nature Genetics, 2011, 43, 109-116.	9.4	1,091
2	Genome-wide mapping of alternative splicing in <i>Arabidopsis thaliana</i> . Genome Research, 2010, 20, 45-58.	2.4	825
3	Network Discovery Pipeline Elucidates Conserved Time-of-Day–Specific cis-Regulatory Modules. PLoS Genetics, 2008, 4, e14.	1.5	474
4	The pineapple genome and the evolution of CAM photosynthesis. Nature Genetics, 2015, 47, 1435-1442.	9.4	472
5	Alternative splicing in plants: directing traffic at the crossroads of adaptation and environmental stress. Current Opinion in Plant Biology, 2015, 24, 125-135.	3.5	215
6	A Morning-Specific Phytohormone Gene Expression Program underlying Rhythmic Plant Growth. PLoS Biology, 2008, 6, e225.	2.6	197
7	Global Profiling of Rice and Poplar Transcriptomes Highlights Key Conserved Circadian-Controlled Pathways and cis-Regulatory Modules. PLoS ONE, 2011, 6, e16907.	1.1	188
8	Transcription Factors in Light and Circadian Clock Signaling Networks Revealed by Genomewide Mapping of Direct Targets for Neurospora White Collar Complex. Eukaryotic Cell, 2010, 9, 1549-1556.	3.4	187
9	Dynamic DNA cytosine methylation in the Populus trichocarpa genome: tissue-level variation and relationship to gene expression. BMC Genomics, 2012, 13, 27.	1.2	136
10	The genome of black raspberry ( <i>Rubus occidentalis</i> ). Plant Journal, 2016, 87, 535-547.	2.8	111
11	Grasses suppress shoot-borne roots to conserve water during drought. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 8861-8866.	3.3	111
12	cis-Regulatory elements in plant cell signaling. Current Opinion in Plant Biology, 2009, 12, 643-649.	3.5	105
13	Extensive Transcriptome Changes During Natural Onset and Release of Vegetative Bud Dormancy in Populus. Frontiers in Plant Science, 2015, 6, 989.	1.7	91
14	Functional characterization of cinnamyl alcohol dehydrogenase and caffeic acid O-methyltransferase in Brachypodium distachyon. BMC Biotechnology, 2013, 13, 61.	1.7	84
15	Analysis of Global Gene Expression in Brachypodium distachyon Reveals Extensive Network Plasticity in Response to Abiotic Stress. PLoS ONE, 2014, 9, e87499.	1.1	80
16	Temporal and spatial transcriptomic and micro <scp>RNA</scp> dynamics of <scp>CAM</scp> photosynthesis in pineapple. Plant Journal, 2017, 92, 19-30.	2.8	78
17	Supersplat—spliced RNA-seq alignment. Bioinformatics, 2010, 26, 1500-1505.	1.8	41
18	Conserved Daily Transcriptional Programs in Carica papaya. Tropical Plant Biology, 2008, 1, 236-245.	1.0	37

HENRY D PRIEST

#	Article	IF	CITATIONS
19	Assembly and Characterization of the European Hazelnut †Jefferson' Transcriptome. Crop Science, 2012, 52, 2679-2686.	0.8	35
20	Comparative Analysis of Vertebrate Diurnal/Circadian Transcriptomes. PLoS ONE, 2017, 12, e0169923.	1.1	29
21	Detection and Quantification of Alternative Splicing Variants Using RNA-seq. Methods in Molecular Biology, 2012, 883, 97-110.	0.4	22
22	Sequencing and characterization of the anadromous steelhead (Oncorhynchus mykiss) transcriptome. Marine Genomics, 2014, 15, 13-15.	0.4	18
23	Transgenic insertion of the cyanobacterial membrane protein ictB increases grain yield in Zea mays through increased photosynthesis and carbohydrate production. PLoS ONE, 2021, 16, e0246359.	1.1	10
24	Developmental variation in DNA methylation in poplar (Populus trichocarpa). BMC Proceedings, 2011, 5, P177.	1.8	0