

# Takashi Akagi

## List of Publications by Year in descending order

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Version: 2024-02-01

49  
papers

2,592  
citations

257357

24  
h-index

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48  
g-index

53  
all docs

53  
docs citations

53  
times ranked

2065  
citing authors

| #  | ARTICLE   | IF   | CITATIONS |
|----|---|------|-----------|
| 1  | A Y-chromosome-encoded small RNA acts as a sex determinant in persimmons. <i>Science</i> , 2014, 346, 646-650.  | 6.0  | 330       |
| 2  | Functional and Expressional Analyses of <i>PmDAM</i> Genes Associated with Endodormancy in Japanese Apricot. <i>Plant Physiology</i> , 2011, 157, 485-497.  | 2.3  | 219       |
| 3  | <i>DkMyb4</i> Is a Myb Transcription Factor Involved in Proanthocyanidin Biosynthesis in Persimmon Fruit. <i>Plant Physiology</i> , 2009, 151, 2028-2045.   | 2.3  | 193       |
| 4  | A Y-Encoded Suppressor of Feminization Arose via Lineage-Specific Duplication of a Cytokinin Response Regulator in Kiwifruit. <i>Plant Cell</i> , 2018, 30, 780-795.  | 3.1  | 151       |
| 5  | Two Y-chromosome-encoded genes determine sex in kiwifruit. <i>Nature Plants</i> , 2019, 5, 801-809.   | 4.7  | 148       |
| 6  | Expression balances of structural genes in shikimate and flavonoid biosynthesis cause a difference in proanthocyanidin accumulation in persimmon ( <i>Diospyros kaki</i> Thunb.) fruit. <i>Planta</i> , 2009, 230, 899-915. | 1.6  | 108       |
| 7  | Antagonistic regulation of the gibberellic acid response during stem growth in rice. <i>Nature</i> , 2020, 584, 109-114.  | 13.7 | 98        |
| 8  | Epigenetic Regulation of the Sex Determination Gene <i>MeGI</i> in Polyploid Persimmon. <i>Plant Cell</i> , 2016, 28, 2905-2915.  | 3.1  | 97        |
| 9  | Proanthocyanidin biosynthesis of persimmon ( <i>Diospyros kaki</i> Thunb.) fruit. <i>Scientia Horticulturae</i> , 2011, 130, 373-380.   | 1.7  | 95        |
| 10 | <i>DkMyb2</i> wound-induced transcription factor of persimmon ( <i>Diospyros kaki</i> Thunb.), contributes to proanthocyanidin regulation. <i>Planta</i> , 2010, 232, 1045-1059.  | 1.6  | 81        |
| 11 | One Hundred Ways to Invent the Sexes: Theoretical and Observed Paths to Dioecy in Plants. <i>Annual Review of Plant Biology</i> , 2018, 69, 553-575.  | 8.6  | 78        |
| 12 | Seasonal Abscisic Acid Signal and a Basic Leucine Zipper Transcription Factor, <i>DkbZIP5</i> , Regulate Proanthocyanidin Biosynthesis in Persimmon Fruit. <i>Plant Physiology</i> , 2012, 158, 1089-1102.                  | 2.3  | 66        |
| 13 | Genome-wide view of genetic diversity reveals paths of selection and cultivar differentiation in peach domestication. <i>DNA Research</i> , 2016, 23, 271-282.  | 1.5  | 64        |
| 14 | The persimmon genome reveals clues to the evolution of a lineage-specific sex determination system in plants. <i>PLoS Genetics</i> , 2020, 16, e1008566.  | 1.5  | 54        |
| 15 | Quantitative characterization of fruit shape and its differentiation pattern in diverse persimmon ( <i>Diospyros kaki</i> ) cultivars. <i>Scientia Horticulturae</i> , 2018, 228, 41-48.                                    | 1.7  | 53        |
| 16 | Evolution and diversification of the plant gibberellin receptor <i>GID1</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E7844-E7853.                              | 3.3  | 51        |
| 17 | Molecular identification of 1-Cys peroxiredoxin and anthocyanidin/flavonol 3-O-galactosyltransferase from proanthocyanidin-rich young fruits of persimmon ( <i>Diospyros kaki</i> ) Tj ETQq1 1 0.7843d 4 rgBT40verloc       |      |           |
| 18 | Gene networks orchestrated by <i>MeG</i> : a single-factor mechanism underlying sex determination in persimmon. <i>Plant Journal</i> , 2019, 98, 97-111.  | 2.8  | 47        |

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|----|---|-----|-----------|
| 19 | Genome Re-Sequencing of Diverse Sweet Cherry ( <i>Prunus avium</i> ) Individuals Reveals a Modifier Gene Mutation Conferring Pollen-Part Self-Compatibility. <i>Plant and Cell Physiology</i> , 2018, 59, 1265-1275.                          | 1.5 | 37        |
| 20 | SCAR Markers for Practical Application of Marker-assisted Selection in Persimmon ( <i>Diospyros kaki</i> ) Tj ETQq0 0 0 rgBT/Overlogk 10 Tf 50  | 0.8 | 35        |
| 21 | Insights into the <i>Prunus</i> -Specific S-RNase-Based Self-Incompatibility System from a Genome-Wide Analysis of the Evolutionary Radiation of <i>S</i> -Locus-Related F-box Genes. <i>Plant and Cell Physiology</i> , 2016, 57, 1281-1294. | 1.5 | 32        |
| 22 | Condensed Tannin Composition Analysis in Persimmon ( <i>Diospyros kaki</i> Thunb.) Fruit by Acid Catalysis in the Presence of Excess Phloroglucinol. <i>Japanese Society for Horticultural Science</i> , 2010, 79, 275-281.                   | 0.8 | 30        |
| 23 | Development of Molecular Markers Associated with Sexuality in <i>Diospyros lotus</i> L. and Their Application in <i>D. kaki</i> Thunb.. <i>Japanese Society for Horticultural Science</i> , 2014, 83, 214-221.                                | 0.8 | 29        |
| 24 | <i>Shy Girl</i> , a kiwifruit suppressor of feminization, restricts gynoceium development via regulation of cytokinin metabolism and signalling. <i>New Phytologist</i> , 2021, 230, 1461-1475.   | 3.5 | 29        |
| 25 | Pleiotropic effects of sex-determining genes in the evolution of dioecy in two plant species. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2019, 286, 20191805.   | 1.2 | 28        |
| 26 | Low temperature modulates natural peel degreening in lemon fruit independently of endogenous ethylene. <i>Journal of Experimental Botany</i> , 2020, 71, 4778-4796.   | 2.4 | 26        |
| 27 | Fine genotyping of a highly polymorphic ASTRINGENCY-linked locus reveals variable hexasomic inheritance in persimmon ( <i>Diospyros kaki</i> Thunb.) cultivars. <i>Tree Genetics and Genomes</i> , 2012, 8, 195-204.                          | 0.6 | 25        |
| 28 | Quantitative Genotyping for the Astringency Locus in Hexaploid Persimmon Cultivars using Quantitative Real-time PCR. <i>Journal of the American Society for Horticultural Science</i> , 2010, 135, 59-66.                                     | 0.5 | 25        |
| 29 | Effects of seasonal temperature changes on <i>DkMyb4</i> expression involved in proanthocyanidin regulation in two genotypes of persimmon ( <i>Diospyros kaki</i> Thunb.) fruit. <i>Planta</i> , 2011, 233, 883-894.                          | 1.6 | 24        |
| 30 | Evolutionary Analysis of Genes for S-RNase-based Self-incompatibility Reveals <i>S</i> -Locus Duplications in the Ancestral Rosaceae. <i>Horticulture Journal</i> , 2015, 84, 233-242.  | 0.3 | 24        |
| 31 | Quantitative real-time PCR to determine allele number for the astringency locus by analysis of a linked marker in <i>Diospyros kaki</i> Thunb. <i>Tree Genetics and Genomes</i> , 2009, 5, 483-492.   | 0.6 | 21        |
| 32 | A male determinant gene in diploid dioecious <i>Diospyros</i> , <i>OGL</i> , is required for male flower production in monoecious individuals of Oriental persimmon ( <i>D. kaki</i> ). <i>Scientia Horticulturae</i> , 2016, 213, 243-251.   | 1.7 | 21        |
| 33 | The effect of layer-by-layer edible coating on the shelf life and transcriptome of "Kosui" Japanese pear fruit. <i>Postharvest Biology and Technology</i> , 2022, 185, 111787.  | 2.9 | 21        |
| 34 | Reinvention of hermaphroditism via activation of a <i>RADIALIS</i> -like gene in hexaploid persimmon. <i>Nature Plants</i> , 2022, 8, 217-224.  | 4.7 | 21        |
| 35 | Development of Molecular Markers Linked to the Allele Associated with the Non-astringent Trait of the Chinese Persimmon ( <i>Diospyros kaki</i> Thunb.). <i>Japanese Society for Horticultural Science</i> , 2011, 80, 150-155.               | 0.8 | 19        |
| 36 | "Passe Crassane" pear fruit ( <i>Pyrus communis</i> L.) ripening: Revisiting the role of low temperature via integrated physiological and transcriptome analysis. <i>Postharvest Biology and Technology</i> , 2019, 158, 110949.              | 2.9 | 18        |

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|----|---|-----|-----------|
| 37 | Characterization of post-mating interspecific cross-compatibility in <i>Prunus</i> (Rosaceae). <i>Scientia Horticulturae</i> , 2019, 246, 693-699.  | 1.7 | 17        |
| 38 | Evolution of Lineage-Specific Gene Networks Underlying the Considerable Fruit Shape Diversity in Persimmon. <i>Plant and Cell Physiology</i> , 2019, 60, 2464-2477.   | 1.5 | 16        |
| 39 | Interspecific introgression and natural selection in the evolution of Japanese apricot ( <i>Prunus</i> ) Tj ETQq1 1 0.784314 rgBT /Overlock 16  | 2.8 | 16        |
| 40 | Explainable Deep Learning Reproduces a "Professional Eye" on the Diagnosis of Internal Disorders in Persimmon Fruit. <i>Plant and Cell Physiology</i> , 2020, 61, 1967-1973.                                | 1.5 | 14        |
| 41 | Epigenetic Flexibility Underlies Somaclonal Sex Conversions in Hexaploid Persimmon. <i>Plant and Cell Physiology</i> , 2020, 61, 393-402.   | 1.5 | 12        |
| 42 | Noninvasive Diagnosis of Seedless Fruit Using Deep Learning in Persimmon. <i>Horticulture Journal</i> , 2021, 90, 172-180.  | 0.3 | 10        |
| 43 | Genome-wide cis-decoding for expression design in tomato using cistrome data and explainable deep learning. <i>Plant Cell</i> , 2022, 34, 2174-2187.  | 3.1 | 10        |
| 44 | Detection of a novel locus involved in non-seed-shattering behaviour of Japonica rice cultivar, <i>Oryzasativa</i> "Nipponbare". <i>Theoretical and Applied Genetics</i> , 2019, 132, 2615-2623.            | 1.8 | 8         |
| 45 | Molecular Mechanism Underlying Derepressed Male Production in Hexaploid Persimmon. <i>Frontiers in Plant Science</i> , 2020, 11, 567249.  | 1.7 | 8         |
| 46 | Genome-wide study on the polysomic genetic factors conferring plasticity of flower sexuality in hexaploid persimmon. <i>DNA Research</i> , 2020, 27, .  | 1.5 | 8         |
| 47 | The Relationship Between a Maleness-associated Region in <i>Diospyros lotus</i> L. and Maleness of Persimmon ( <i>D. kaki</i> Thunb.) Cultivars. <i>Horticultural Research (Japan)</i> , 2015, 14, 121-126. | 0.1 | 7         |
| 48 | Examining the Role of Low Temperature in Satsuma Mandarin Fruit Peel Degreening via Comparative Physiological and Transcriptomic Analysis. <i>Frontiers in Plant Science</i> , 0, 13, .                     | 1.7 | 6         |
| 49 | Evolutionary alterations in gene expression and enzymatic activities of gibberellin 3-oxidase 1 in <i>Oryza</i> . <i>Communications Biology</i> , 2022, 5, 67.  | 2.0 | 4         |