## Yuriko Osakabe

## List of Publications by Citations

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76
papers

7,945
citations

41
p-index

81
g-index

81
ext. papers

6.8
avg, IF

L-index

| #              | Paper  | IF   | Citations |
|----------------|--|------|-----------|
| 76             | Functional analysis of an Arabidopsis transcription factor, DREB2A, involved in drought-responsive gene expression. <i>Plant Cell</i> , <b>2006</b> , 18, 1292-309   | 11.6 | 780       |
| 75             | Response of plants to water stress. Frontiers in Plant Science, 2014, 5, 86  | 6.2  | 740       |
| 74             | Dual function of an Arabidopsis transcription factor DREB2A in water-stress-responsive and heat-stress-responsive gene expression. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2006</b> , 103, 18822-7 | 11.5 | 561       |
| 73             | Positive regulatory role of strigolactone in plant responses to drought and salt stress. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2014</b> , 111, 851-6   | 11.5 | 370       |
| 7 <sup>2</sup> | Regulation and functional analysis of ZmDREB2A in response to drought and heat stresses in Zea mays L. <i>Plant Journal</i> , <b>2007</b> , 50, 54-69  | 6.9  | 353       |
| 71             | Sensing the environment: key roles of membrane-localized kinases in plant perception and response to abiotic stress. <i>Journal of Experimental Botany</i> , <b>2013</b> , 64, 445-58  | 7    | 274       |
| 70             | Arabidopsis HsfA1 transcription factors function as the main positive regulators in heat shock-responsive gene expression. <i>Molecular Genetics and Genomics</i> , <b>2011</b> , 286, 321-32  | 3.1  | 253       |
| 69             | RPK2 is an essential receptor-like kinase that transmits the CLV3 signal in Arabidopsis. <i>Development</i> (Cambridge), <b>2010</b> , 137, 3911-20  | 6.6  | 249       |
| 68             | Leucine-rich repeat receptor-like kinase1 is a key membrane-bound regulator of abscisic acid early signaling in Arabidopsis. <i>Plant Cell</i> , <b>2005</b> , 17, 1105-19   | 11.6 | 239       |
| 67             | Osmotic stress responses and plant growth controlled by potassium transporters in Arabidopsis. <i>Plant Cell</i> , <b>2013</b> , 25, 609-24  | 11.6 | 237       |
| 66             | Site-directed mutagenesis in Arabidopsis using custom-designed zinc finger nucleases. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2010</b> , 107, 12034-9  | 11.5 | 231       |
| 65             | ABA control of plant macroelement membrane transport systems in response to water deficit and high salinity. <i>New Phytologist</i> , <b>2014</b> , 202, 35-49   | 9.8  | 217       |
| 64             | A small peptide modulates stomatal control via abscisic acid in long-distance signalling. <i>Nature</i> , <b>2018</b> , 556, 235-238   | 50.4 | 214       |
| 63             | Co-expression of the stress-inducible zinc finger homeodomain ZFHD1 and NAC transcription factors enhances expression of the ERD1 gene in Arabidopsis. <i>Plant Journal</i> , <b>2007</b> , 49, 46-63  | 6.9  | 204       |
| 62             | Optimization of CRISPR/Cas9 genome editing to modify abiotic stress responses in plants. <i>Scientific Reports</i> , <b>2016</b> , 6, 26685  | 4.9  | 192       |
| 61             | Efficient Genome Editing in Apple Using a CRISPR/Cas9 system. Scientific Reports, 2016, 6, 31481   | 4.9  | 179       |
| 60             | Genome editing with engineered nucleases in plants. Plant and Cell Physiology, 2015, 56, 389-400   | 4.9  | 154       |

## (2020-2009)

| 59 | The phytochrome-interacting factor PIF7 negatively regulates DREB1 expression under circadian control in Arabidopsis. <i>Plant Physiology</i> , <b>2009</b> , 151, 2046-57  | 6.6  | 154 |
|----|---|------|-----|
| 58 | Rapid breeding of parthenocarpic tomato plants using CRISPR/Cas9. <i>Scientific Reports</i> , <b>2017</b> , 7, 507  | 4.9  | 147 |
| 57 | Receptor-like protein kinase 2 (RPK 2) is a novel factor controlling anther development in Arabidopsis thaliana. <i>Plant Journal</i> , <b>2007</b> , 50, 751-66  | 6.9  | 147 |
| 56 | Abiotic stress-inducible receptor-like kinases negatively control ABA signaling in Arabidopsis. <i>Plant Journal</i> , <b>2012</b> , 70, 599-613  | 6.9  | 130 |
| 55 | Functional analysis of an Arabidopsis thaliana abiotic stress-inducible facilitated diffusion transporter for monosaccharides. <i>Journal of Biological Chemistry</i> , <b>2010</b> , 285, 1138-46  | 5.4  | 120 |
| 54 | GmDREB2A;2, a canonical DEHYDRATION-RESPONSIVE ELEMENT-BINDING PROTEIN2-type transcription factor in soybean, is posttranslationally regulated and mediates dehydration-responsive element-dependent gene expression. <i>Plant Physiology</i> , <b>2013</b> , 161, 346-61                 | 6.6  | 113 |
| 53 | Overproduction of the membrane-bound receptor-like protein kinase 1, RPK1, enhances abiotic stress tolerance in Arabidopsis. <i>Journal of Biological Chemistry</i> , <b>2010</b> , 285, 9190-201   | 5.4  | 107 |
| 52 | Arabidopsis DPB3-1, a DREB2A interactor, specifically enhances heat stress-induced gene expression by forming a heat stress-specific transcriptional complex with NF-Y subunits. <i>Plant Cell</i> , <b>2014</b> , 26, 4954-73  | 11.6 | 95  |
| 51 | The karrikin receptor KAI2 promotes drought resistance in Arabidopsis thaliana. <i>PLoS Genetics</i> , <b>2017</b> , 13, e1007076   | 6    | 87  |
| 50 | Generation of Bolanine-free hairy roots of potato by CRISPR/Cas9 mediated genome editing of the St16DOX gene. <i>Plant Physiology and Biochemistry</i> , <b>2018</b> , 131, 70-77   | 5.4  | 86  |
| 49 | CRISPR-Cas9-mediated genome editing in apple and grapevine. <i>Nature Protocols</i> , <b>2018</b> , 13, 2844-2863   | 18.8 | 86  |
| 48 | Rice phytochrome-interacting factor-like protein OsPIL1 functions as a key regulator of internode elongation and induces a morphological response to drought stress. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2012</b> , 109, 15947-52 | 11.5 | 84  |
| 47 | Isolation and characterization of the RAD54 gene from Arabidopsis thaliana. <i>Plant Journal</i> , <b>2006</b> , 48, 827-42   | 6.9  | 75  |
| 46 | An efficient DNA- and selectable-marker-free genome-editing system using zygotes in rice. <i>Nature Plants</i> , <b>2019</b> , 5, 363-368   | 11.5 | 72  |
| 45 | Overexpression of Arabidopsis response regulators, ARR4/ATRR1/IBC7 and ARR8/ATRR3, alters cytokinin responses differentially in the shoot and in callus formation. <i>Biochemical and Biophysical Research Communications</i> , <b>2002</b> , 293, 806-15                                 | 3.4  | 72  |
| 44 | Characterization of the promoter region of an Arabidopsis gene for 9-cis-epoxycarotenoid dioxygenase involved in dehydration-inducible transcription. <i>DNA Research</i> , <b>2013</b> , 20, 315-24  | 4.5  | 67  |
| 43 | Secondary xylem-specific expression of caffeoyl-coenzyme A 3-O-methyltransferase plays an important role in the methylation pathway associated with lignin biosynthesis in loblolly pine. <i>Plant Molecular Biology</i> , <b>1999</b> , 40, 555-65                                       | 4.6  | 64  |
| 42 | Precision genome editing in plants: state-of-the-art in CRISPR/Cas9-based genome engineering. <i>BMC Plant Biology</i> , <b>2020</b> , 20, 234  | 5.3  | 63  |

| 41 | Monosaccharide absorption activity of Arabidopsis roots depends on expression profiles of transporter genes under high salinity conditions. <i>Journal of Biological Chemistry</i> , <b>2011</b> , 286, 43577-86 | 5.4  | 57 |
|----|--|------|----|
| 40 | Direct conversion of carlactonoic acid to orobanchol by cytochrome P450 CYP722C in strigolactone biosynthesis. <i>Science Advances</i> , <b>2019</b> , 5, eaax9067   | 14.3 | 52 |
| 39 | Efficient Multiplex Genome Editing Induces Precise, and Self-Ligated Type Mutations in Tomato Plants. <i>Frontiers in Plant Science</i> , <b>2018</b> , 9, 916   | 6.2  | 48 |
| 38 | Genome editing in the mushroom-forming basidiomycete Coprinopsis cinerea, optimized by a high-throughput transformation system. <i>Scientific Reports</i> , <b>2017</b> , 7, 1260                                | 4.9  | 47 |
| 37 | MYB transcription factor gene involved in sex determination in Asparagus officinalis. <i>Genes To Cells</i> , <b>2017</b> , 22, 115-123  | 2.3  | 42 |
| 36 | Genetic engineering of woody plants: current and future targets in a stressful environment. <i>Physiologia Plantarum</i> , <b>2011</b> , 142, 105-17   | 4.6  | 42 |
| 35 | Stabilization of Arabidopsis DREB2A is required but not sufficient for the induction of target genes under conditions of stress. <i>PLoS ONE</i> , <b>2013</b> , 8, e80457                                       | 3.7  | 41 |
| 34 | Responses to environmental stresses in woody plants: key to survive and longevity. <i>Journal of Plant Research</i> , <b>2012</b> , 125, 1-10  | 2.6  | 27 |
| 33 | Immunocytochemical localization of phenylalanine ammonia-lyase in tissues of Populus kitakamiensis. <i>Planta</i> , <b>1996</b> , 200, 13-9  | 4.7  | 26 |
| 32 | Efficient and Heritable Targeted Mutagenesis in Mosses Using the CRISPR/Cas9 System. <i>Plant and Cell Physiology</i> , <b>2016</b> , 57, 2600-2610  | 4.9  | 24 |
| 31 | Structure and tissue-specific expression of genes for phenylalanine ammonia-lyase from a hybrid aspen, Populus kitakamiensis. <i>Plant Science</i> , <b>1995</b> , 105, 217-226                                  | 5.3  | 24 |
| 30 | Characterization of the structure and determination of mRNA levels of the phenylalanine ammonia-lyase gene family from Populus kitakamiensis. <i>Plant Molecular Biology</i> , <b>1995</b> , 28, 1133-41         | 4.6  | 23 |
| 29 | Lignin characterization of rice CONIFERALDEHYDE 5-HYDROXYLASE loss-of-function mutants generated with the CRISPR/Cas9 system. <i>Plant Journal</i> , <b>2019</b> , 97, 543-554                                   | 6.9  | 22 |
| 28 | Genome Editing to Improve Abiotic Stress Responses in Plants. <i>Progress in Molecular Biology and Translational Science</i> , <b>2017</b> , 149, 99-109   | 4    | 21 |
| 27 | OsMYB108 loss-of-function enriches p-coumaroylated and tricin lignin units in rice cell walls. <i>Plant Journal</i> , <b>2019</b> , 98, 975-987  | 6.9  | 21 |
| 26 | Genome editing in plants using CRISPR type I-D nuclease. <i>Communications Biology</i> , <b>2020</b> , 3, 648  | 6.7  | 20 |
| 25 | A mutated cytosine deaminase gene, codA (D314A), as an efficient negative selection marker for gene targeting in rice. <i>Plant and Cell Physiology</i> , <b>2014</b> , 55, 658-65                               | 4.9  | 20 |
| 24 | Comparative functional analyses of DWARF14 and KARRIKIN INSENSITIVE in drought adaptation of Arabidopsis thaliana. <i>Plant Journal</i> , <b>2020</b> , 103, 111-127   | 6.9  | 19 |

|                | Characterization of the tissue-specific expression of phenylalanine ammonia-lyase gene promoter from loblolly pine (Pinus taeda) in Nicotiana tabacum. <i>Plant Cell Reports</i> , <b>2009</b> , 28, 1309-17   | 5.1             | 18                                      |
|----------------|--|-----------------|---|
| 22             | Sugar compartmentation as an environmental stress adaptation strategy in plants. <i>Seminars in Cell and Developmental Biology</i> , <b>2018</b> , 83, 106-114   | 7.5             | 15                                      |
| 21             | Isolation of 4-coumarate Co-A ligase gene promoter from loblolly pine (Pinus taeda) and characterization of tissue-specific activity in transgenic tobacco. <i>Plant Physiology and Biochemistry</i> , <b>2009</b> , 47, 1031-6  | 5.4             | 14                                      |
| 20             | Genome engineering of woody plants: past, present and future. Journal of Wood Science, 2016, 62, 217   | 7-2 <u>32.5</u> | 13                                      |
| 19             | Lotus japonicus Triterpenoid Profile and Characterization of the CYP716A51 and LjCYP93E1 Genes Involved in Their Biosynthesis In Planta. <i>Plant and Cell Physiology</i> , <b>2019</b> , 60, 2496-2509  | 4.9             | 12                                      |
| 18             | A C-terminal motif contributes to the plasma membrane localization of Arabidopsis STP transporters. <i>PLoS ONE</i> , <b>2017</b> , 12, e0186326   | 3.7             | 10                                      |
| 17             | Double knockout of OsWRKY36 and OsWRKY102 boosts lignification with altering culm morphology of rice. <i>Plant Science</i> , <b>2020</b> , 296, 110466   | 5.3             | 9                                       |
| 16             | RPK2 is an essential receptor-like kinase that transmits the CLV3 signal in Arabidopsis. <i>Development (Cambridge)</i> , <b>2010</b> , 137, 4327-4327   | 6.6             | 9                                       |
| 15             | Characterization of steroid 5E eductase involved in Etomatine biosynthesis in tomatoes. <i>Plant Biotechnology</i> , <b>2019</b> , 36, 253-263   | 1.3             | 9                                       |
| 14             | Genome editing in mammalian cells using the CRISPR type I-D nuclease. <i>Nucleic Acids Research</i> , <b>2021</b> , 49, 6347-6363  | 20.1            | 7                                       |
|                |  |                 |   |
| 13             | Plant Environmental Stress Responses for Survival and Biomass Enhancement 2013, 79-108   |                 | 5                                       |
| 13             | Plant Environmental Stress Responses for Survival and Biomass Enhancement 2013, 79-108  Plant Light Stress 2012,   |                 | 5                                       |
|                |  | 1.9             |   |
| 12             | Plant Light Stress 2012,  Targeted mutagenesis of CENTRORADIALIS using CRISPR/Cas9 system through the improvement of genetic transformation efficiency of tetraploid highbush blueberry. <i>Journal of Horticultural</i>   | 1.9             | 5                                       |
| 12             | Plant Light Stress 2012,  Targeted mutagenesis of CENTRORADIALIS using CRISPR/Cas9 system through the improvement of genetic transformation efficiency of tetraploid highbush blueberry. <i>Journal of Horticultural Science and Biotechnology</i> , 2021, 96, 153-161  Overexpression of a fungal laccase gene induces nondehiscent anthers and morphological changes   |                 | 5                                       |
| 12<br>11<br>10 | Plant Light Stress 2012,  Targeted mutagenesis of CENTRORADIALIS using CRISPR/Cas9 system through the improvement of genetic transformation efficiency of tetraploid highbush blueberry. <i>Journal of Horticultural Science and Biotechnology</i> , 2021, 96, 153-161  Overexpression of a fungal laccase gene induces nondehiscent anthers and morphological changes in flowers of transgenic tobacco. <i>Journal of Wood Science</i> , 2010, 56, 460-469  Expanding the plant genome editing toolbox with recently developed CRISPR-Cas systems <i>Plant</i>                    | 2.4             | <ul><li>5</li><li>5</li><li>4</li></ul> |
| 12<br>11<br>10 | Plant Light Stress 2012,  Targeted mutagenesis of CENTRORADIALIS using CRISPR/Cas9 system through the improvement of genetic transformation efficiency of tetraploid highbush blueberry. <i>Journal of Horticultural Science and Biotechnology</i> , 2021, 96, 153-161  Overexpression of a fungal laccase gene induces nondehiscent anthers and morphological changes in flowers of transgenic tobacco. <i>Journal of Wood Science</i> , 2010, 56, 460-469  Expanding the plant genome editing toolbox with recently developed CRISPR-Cas systems <i>Plant Physiology</i> , 2022, | 2.4             | <ul><li>5</li><li>5</li><li>4</li></ul> |

| 5 | Genome editing in mammals using CRISPR type I-D nuclease                                  |     | 2 |
|---|---|-----|---|
| 4 | Genome Editing in Higher Plants <b>2015</b> , 197-205                                     |     | 1 |
| 3 | Crop Breeding Using CRISPR/Cas9 <b>2018</b> , 451-464                                     |     | 1 |
| 2 | Transcription Factors: Improving Abiotic Stress Tolerance in Plants <b>2012</b> , 591-621 |     | 1 |
| 1 | Genome Editing in Apple. Compendium of Plant Genomes 2021, 213-225                        | 0.8 | 0 |