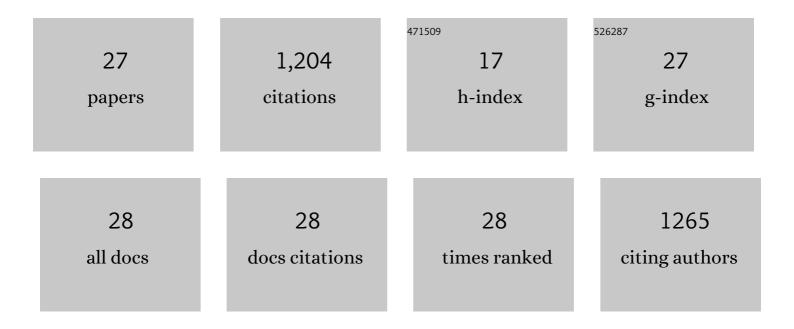
Weiwei Qi

List of Publications by Year in descending order

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| # | Article | IF | CITATIONS |
|----|---|-----|-----------|
| 1 | High-efficiency CRISPR/Cas9 multiplex gene editing using the glycine tRNA-processing system-based strategy in maize. BMC Biotechnology, 2016, 16, 58. | 3.3 | 162 |
| 2 | Genome-Wide Characterization of <i>cis</i> -Acting DNA Targets Reveals the Transcriptional Regulatory Framework of <i>Opaque2</i> in Maize. Plant Cell, 2015, 27, 532-545. | 6.6 | 130 |
| 3 | Dek35 Encodes a PPR Protein that Affects cis -Splicing of Mitochondrial nad4 Intron 1 andÂSeed Development in Maize. Molecular Plant, 2017, 10, 427-441. | 8.3 | 106 |
| 4 | OPAQUE11 Is a Central Hub of the Regulatory Network for Maize Endosperm Development and Nutrient Metabolism. Plant Cell, 2018, 30, 375-396. | 6.6 | 103 |
| 5 | Mitochondrial Function and Maize Kernel Development Requires Dek2, a Pentatricopeptide Repeat Protein Involved in nad1 mRNA Splicing. Genetics, 2017, 205, 239-249. | 2.9 | 82 |
| 6 | The ZmbZIP22 Transcription Factor Regulates 27-kD γ-Zein Gene Transcription during Maize Endosperm Development. Plant Cell, 2018, 30, 2402-2424. | 6.6 | 65 |
| 7 | ZmMADS47 Regulates Zein Gene Transcription through Interaction with Opaque2. PLoS Genetics, 2016, 12, e1005991. | 3.5 | 62 |
| 8 | Editing of Mitochondrial Transcripts <i>nad3</i> and <i>cox2</i> by Dek10 Is Essential for Mitochondrial Function and Maize Plant Development. Genetics, 2017, 205, 1489-1501. | 2.9 | 56 |
| 9 | Maize <i>Dek37</i> Encodes a P-type PPR Protein That Affects <i>cis</i> -Splicing of Mitochondrial <i>nad2</i> Intron 1 and Seed Development. Genetics, 2018, 208, 1069-1082. | 2.9 | 55 |
| 10 | Identification and Characterization of Maize floury4 as a Novel Semidominant Opaque Mutant That Disrupts Protein Body Assembly Â. Plant Physiology, 2014, 165, 582-594. | 4.8 | 52 |
| 11 | Maize opaque10 Encodes a Cereal-Specific Protein That Is Essential for the Proper Distribution of Zeins in Endosperm Protein Bodies. PLoS Genetics, 2016, 12, e1006270. | 3.5 | 43 |
| 12 | Maize <i>reas1</i> Mutant Stimulates Ribosome Use Efficiency and Triggers Distinct Transcriptional and Translational Responses. Plant Physiology, 2016, 170, 971-988. | 4.8 | 41 |
| 13 | <i>>Dek42</i> encodes an RNAâ€binding protein that affects alternative preâ€mRNA splicing and maize kernel development. Journal of Integrative Plant Biology, 2019, 61, 728-748. | 8.5 | 38 |
| 14 | Maize <i>Dek15</i> Encodes the Cohesin-Loading Complex Subunit SCC4 and Is Essential for Chromosome Segregation and Kernel Development. Plant Cell, 2019, 31, 465-485. | 6.6 | 35 |
| 15 | Maize pentatricopeptide repeat protein DEK41 affects cis-splicing of mitochondrial nad4 intron 3 and is required for normal seed development. Journal of Experimental Botany, 2019, 70, 3795-3808. | 4.8 | 35 |
| 16 | Maize <i>Dek44</i> Encodes Mitochondrial Ribosomal Protein L9 and Is Required for Seed Development. Plant Physiology, 2019, 180, 2106-2119. | 4.8 | 28 |
| 17 | A SnRK1- <i>Zm</i> RFWD3-Opaque2 Signaling Axis Regulates Diurnal Nitrogen Accumulation in Maize Seeds. Plant Cell, 2020, 32, 2823-2841. | 6.6 | 22 |
| 18 | Maize Dek33 encodes a pyrimidine reductase in riboflavin biosynthesis that is essential for oil-body formation and ABA biosynthesis during seed development. Journal of Experimental Botany, 2019, 70, 5173-5187. | 4.8 | 16 |

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| # | Article | IF | CITATIONS |
|----|---|-----|-----------|
| 19 | Maize pentatricopeptide repeat protein DEK53 is required for mitochondrial RNA editing at multiple sites and seed development. Journal of Experimental Botany, 2020, 71, 6246-6261. | 4.8 | 16 |
| 20 | <i>ENB1</i> encodes a cellulose synthase 5 that directs synthesis of cell wall ingrowths in maize basal endosperm transfer cells. Plant Cell, 2022, 34, 1054-1074. | 6.6 | 13 |
| 21 | <i>shrunken4</i> is a mutant allele of <i>ZmYSL2</i> that affects aleurone development and starch synthesis in maize. Genetics, 2021, 218, . | 2.9 | 12 |
| 22 | Comparative Study between the CRISPR/Cpf1 (Cas12a) and CRISPR/Cas9 Systems for Multiplex Gene Editing in Maize. Agriculture (Switzerland), 2021, 11, 429. | 3.1 | 11 |
| 23 | Maize ZmVPP5 is a truncated Vacuole H ⁺ â€PPase that confers hypersensitivity to salt stress. Journal of Integrative Plant Biology, 2016, 58, 518-528. | 8.5 | 7 |
| 24 | Accumulation of 22 kDa αâ€zeinâ€mediated nonzein protein in protein body of maize endosperm. New Phytologist, 2022, 233, 265-281. | 7.3 | 5 |
| 25 | Lactobacillus paracasei BD5115-Derived 2-Hydroxy-3-Methylbutyric Acid Promotes Intestinal Epithelial Cells Proliferation by Upregulating the MYC Signaling Pathway. Frontiers in Nutrition, 2022, 9, 799053. | 3.7 | 4 |
| 26 | Pollen-Specific CRISPR/Cas9 System to Increase Heritable Gene Mutations in Maize. Agriculture (Switzerland), 2021, 11, 751. | 3.1 | 3 |
| 27 | Establishment of a Bivector Genetic Transformation System in Recalcitrant Maize Inbred Lines. Agriculture (Switzerland), 2021, 11, 663. | 3.1 | 1 |