## Xuexia Miao

## List of Publications by Year in descending order

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567281 610901 24 954 15 24 citations h-index g-index papers 24 24 24 1145 all docs docs citations times ranked citing authors

| #  | Article   | IF  | CITATIONS |
|----|---|-----|-----------|
| 1  | New insights into an <scp>RNAi</scp> approach for plant defence against piercingâ€sucking and stemâ€borer insect pests. Plant, Cell and Environment, 2015, 38, 2277-2285.   | 5.7 | 158       |
| 2  | Second-Generation Sequencing Supply an Effective Way to Screen RNAi Targets in Large Scale for Potential Application in Pest Insect Control. PLoS ONE, 2011, 6, e18644.   | 2.5 | 143       |
| 3  | The OsmiR396–Os <scp>GRF</scp> 8–OsF3Hâ€flavonoid pathway mediates resistance to the brown planthopper in rice ( <i>Oryza sativa</i> ). Plant Biotechnology Journal, 2019, 17, 1657-1669.                         | 8.3 | 110       |
| 4  | Modulation of plant architecture by the miR156f–OsSPL7–OsGH3.8 pathway in rice. Journal of Experimental Botany, 2018, 69, 5117-5130.  | 4.8 | 97        |
| 5  | Novel crosstalk between ethylene―and jasmonic acidâ€pathway responses to a piercing–sucking insect in rice. New Phytologist, 2020, 225, 474-487.  | 7.3 | 53        |
| 6  | OsRAMOSA2 Shapes Panicle Architecture through Regulating Pedicel Length. Frontiers in Plant Science, 2017, 8, 1538.   | 3.6 | 45        |
| 7  | OsmiR396/growth regulating factor modulate rice grain size through direct regulation of embryo-specific miR408. Plant Physiology, 2021, 186, 519-533.   | 4.8 | 36        |
| 8  | Identification of differential expression genes associated with host selection and adaptation between two sibling insect species by transcriptional profile analysis. BMC Genomics, 2013, 14, 582.                | 2.8 | 30        |
| 9  | Identification of transcription factors potential related to brown planthopper resistance in rice via microarray expression profiling. BMC Genomics, 2012, 13, 687.   | 2.8 | 29        |
| 10 | Microarray analysis of broad-spectrum resistance derived from an indica cultivar Rathu Heenati. Planta, 2012, 235, 829-840.   | 3.2 | 27        |
| 11 | Lepidopteran insect speciesâ€specific, broadâ€spectrum, and systemic <scp>RNA</scp> interference by spraying ds <scp>RNA</scp> on larvae. Entomologia Experimentalis Et Applicata, 2015, 155, 218-228.            | 1.4 | 26        |
| 12 | OsEXPA10 mediates the balance between growth and resistance to biotic stress in rice. Plant Cell Reports, 2018, 37, 993-1002.   | 5.6 | 25        |
| 13 | RNAi pest control and enhanced BT insecticidal efficiency achieved by dsRNA of chymotrypsin-like genes in Ostrinia furnacalis. Journal of Pest Science, 2017, 90, 745-757.  | 3.7 | 23        |
| 14 | OsHLH61-OsbHLH96 influences rice defense to brown planthopper through regulating the pathogen-related genes. Rice, 2019, 12, 9.   | 4.0 | 23        |
| 15 | The in vivo dsRNA Cleavage Has Sequence Preference in Insects. Frontiers in Physiology, 2018, 9, 1768.  | 2.8 | 19        |
| 16 | Advances in the Development of Microbial Double-Stranded RNA Production Systems for Application of RNA Interference in Agricultural Pest Control. Frontiers in Bioengineering and Biotechnology, 2021, 9, 753790. | 4.1 | 19        |
| 17 | OsMADS1 Represses microRNA172 in Elongation of Palea/Lemma Development in Rice. Frontiers in Plant Science, 2016, 7, 1891.  | 3.6 | 16        |
| 18 | Bab2 Functions as an Ecdysone-Responsive Transcriptional Repressor during Drosophila Development. Cell Reports, 2020, 32, 107972.   | 6.4 | 15        |

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|----|--|-----|----------|
| 19 | Knockout of the HaREase Gene Improves the Stability of dsRNA and Increases the Sensitivity of Helicoverpa armigera to Bacillus thuringiensis Toxin. Frontiers in Physiology, 2019, 10, 1368. | 2.8 | 14       |
| 20 | miR156f integrates panicle architecture through genetic modulation of branch number and pedicel length pathways. Rice, 2019, 12, 40.   | 4.0 | 14       |
| 21 | Identification of the rice genes and metabolites involved in dual resistance against brown planthopper and rice blast fungus. Plant, Cell and Environment, 2022, 45, 1914-1929.              | 5.7 | 14       |
| 22 | Proteomic Analysis of Silkworm Antennae. Journal of Chemical Ecology, 2015, 41, 1037-1042.   | 1.8 | 12       |
| 23 | Identification and phylogeny of five maleâ€specific lethal genes in the silkworm <i>Bombyx mori</i> Entomological Research, 2008, 38, S48.   | 1.1 | 3        |
| 24 | The Novel Agrotis ipsilon Nora Virus Confers Deleterious Effects to the Fitness of Spodoptera frugiperda (Lepidoptera: Noctuidae). Frontiers in Microbiology, 2021, 12, 727202.              | 3.5 | 3        |