

# Hai Nian

## List of Publications by Year in descending order

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11  
papers

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| #  | ARTICLE   | IF  | CITATIONS |
|----|---|-----|-----------|
| 1  | Fine mapping of an adult-plant resistance gene to powdery mildew in soybean cultivar Zhonghuang 24. <i>Crop Journal</i> , 2022, 10, 1103-1110.  | 5.2 | 7         |
| 2  | Fine mapping and genetic analysis of resistance genes, Rsc18, against soybean mosaic virus. <i>Journal of Integrative Agriculture</i> , 2022, 21, 644-653.  | 3.5 | 4         |
| 3  | QTL mapping for soybean ( <i>Glycine max L.</i> ) leaf chlorophyll-content traits in a genotyped RIL population by using RAD-seq based high-density linkage map. <i>BMC Genomics</i> , 2020, 21, 739. | 2.8 | 14        |
| 4  | Fine mapping and analyses of the RSC15ZH resistance candidate gene for the soybean mosaic virus. <i>Euphytica</i> , 2020, 216, 1.   | 1.2 | 2         |
| 5  | Fine mapping of a Phytophthora-resistance locus RpsGZ in soybean using genotyping-by-sequencing. <i>BMC Genomics</i> , 2020, 21, 280.   | 2.8 | 20        |
| 6  | Genetic mapping of powdery mildew resistance genes in soybean by high-throughput genome-wide sequencing. <i>Theoretical and Applied Genetics</i> , 2019, 132, 1833-1845.                              | 3.6 | 17        |
| 7  | QTL fine-mapping of soybean ( <i>Glycine max L.</i> ) leaf type associated traits in two RILs populations. <i>BMC Genomics</i> , 2019, 20, 260.   | 2.8 | 17        |
| 8  | Fine-mapping of QTLs for individual and total isoflavone content in soybean ( <i>Glycine max L.</i> ) using a high-density genetic map. <i>Theoretical and Applied Genetics</i> , 2018, 131, 555-568. | 3.6 | 34        |
| 9  | Fine mapping of a Phytophthora-resistance gene RpsWY in soybean ( <i>Glycine max L.</i> ) by high-throughput genome-wide sequencing. <i>Theoretical and Applied Genetics</i> , 2017, 130, 1041-1051.  | 3.6 | 80        |
| 10 | A Single Nucleotide Deletion in J Encoding GmELF3 Confers Long Juvenility and Is Associated with Adaption of Tropic Soybean. <i>Molecular Plant</i> , 2017, 10, 656-658.                              | 8.3 | 96        |
| 11 | Construction of high-density genetic map and QTL mapping of yield-related and two quality traits in soybean RILs population by RAD-sequencing. <i>BMC Genomics</i> , 2017, 18, 466.                   | 2.8 | 51        |