

Development of a 63K SNP Array for Cotton and High-D Interspecific Populations of *Gossypium* spp.

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Citation Report

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| 207 | Identification and genomic characterization of major effect bacterial blight resistance locus (BB-13) in Upland cotton (<i>Gossypium hirsutum</i> L.). <i>Theoretical and Applied Genetics</i> , 2022, 135, 4421-4436. | 1.8 | 2 |
| 208 | Recent progression and future perspectives in cotton genomic breeding. <i>Journal of Integrative Plant Biology</i> , 2023, 65, 548-569. | 4.1 | 20 |
| 209 | Quantitative Trait Locus Mapping for Fusarium Wilt Race 4 Resistance in a Recombinant Inbred Line Population of Pima Cotton (<i>Gossypium Barbadense</i>). <i>Pathogens</i> , 2022, 11, 1143. | 1.2 | 5 |
| 210 | Population Genomics of Cotton. <i>Population Genomics</i> , 2022, , . | 0.2 | 0 |
| 211 | A method for low-coverage single-gamete sequence analysis demonstrates adherence to Mendel's first law across a large sample of human sperm. <i>ELife</i> , 0, 11, . | 2.8 | 2 |
| 212 | Inter-Specific Hybridization in Cotton (<i>Gossypium hirsutum</i>) for Crop Improvement. <i>Agronomy</i> , 2022, 12, 3158. | 1.3 | 2 |

| # | ARTICLE | IF | CITATIONS |
|-----|--|-----|-----------|
| 213 | Identification of quantitative trait loci for fiber quality, yield, and plant height traits in Upland cotton. <i>Crop Science</i> , 2023, 63, 1421-1441. | 0.8 | 1 |
| 214 | Detecting Cotton Leaf Curl Virus Resistance Quantitative Trait Loci in <i>Gossypium hirsutum</i> and iCottonQTL a New R/Shiny App to Streamline Genetic Mapping. <i>Plants</i> , 2023, 12, 1153. | 1.6 | 1 |
| 215 | First insight into the phylogeny of fine-leaved <i>Festuca</i> in the Altai Mountain Country based on genome-wide genotyping. <i>Ecology and Evolution</i> , 2023, 13, . | 0.8 | 1 |
| 216 | Haplotyping interspecific hybrids by dual alignment to both parental genomes. <i>Plant Genome</i> , 0, , . | 1.6 | 1 |