Yi Han

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1247518/publications.pdf

Version: 2024-02-01

257450 434195 3,644 29 24 31 citations h-index g-index papers 34 34 34 6041 all docs docs citations times ranked citing authors

| # | Article | IF | CITATIONS |
|----|--|--------------|-----------|
| 1 | Natural variation in genome architecture among 205 <i>Drosophila melanogaster</i> Genetic Reference Panel lines. Genome Research, 2014, 24, 1193-1208. | 5.5 | 565 |
| 2 | The house spider genome reveals an ancient whole-genome duplication during arachnid evolution. BMC Biology, 2017, 15, 62. | 3.8 | 286 |
| 3 | Genome of the Asian longhorned beetle (Anoplophora glabripennis), a globally significant invasive species, reveals key functional and evolutionary innovations at the beetle–plant interface. Genome Biology, 2016, 17, 227. | 8.8 | 244 |
| 4 | Hemimetabolous genomes reveal molecular basis of termite eusociality. Nature Ecology and Evolution, 2018, 2, 557-566. | 7.8 | 223 |
| 5 | The First Myriapod Genome Sequence Reveals Conservative Arthropod Gene Content and Genome Organisation in the Centipede Strigamia maritima. PLoS Biology, 2014, 12, e1002005. | 5.6 | 221 |
| 6 | Unique features of a global human ectoparasite identified through sequencing of the bed bug genome. Nature Communications, 2016, 7, 10165. | 12.8 | 184 |
| 7 | A Massive Expansion of Effector Genes Underlies Gall-Formation in the Wheat Pest Mayetiola destructor. Current Biology, 2015, 25, 613-620. | 3.9 | 171 |
| 8 | Evolutionary History of Chemosensory-Related Gene Families across the Arthropoda. Molecular Biology and Evolution, 2017, 34, 1838-1862. | 8.9 | 157 |
| 9 | Multifaceted biological insights from a draft genome sequence of the tobacco hornworm moth, Manduca sexta. Insect Biochemistry and Molecular Biology, 2016, 76, 118-147. | 2.7 | 154 |
| 10 | Assessing structural variation in a personal genomeâ€"towards a human reference diploid genome. BMC Genomics, 2015, 16, 286. | 2.8 | 153 |
| 11 | Gene content evolution in the arthropods. Genome Biology, 2020, 21, 15. | 8.8 | 150 |
| 12 | Comparative validation of the <i>D. melanogaster</i> modENCODE transcriptome annotation. Genome Research, 2014, 24, 1209-1223. | 5 . 5 | 147 |
| 13 | The whole genome sequence of the Mediterranean fruit fly, Ceratitis capitata (Wiedemann), reveals insights into the biology and adaptive evolution of a highly invasive pest species. Genome Biology, 2016, 17, 192. | 8.8 | 130 |
| 14 | Molecular evolutionary trends and feeding ecology diversification in the Hemiptera, anchored by the milkweed bug genome. Genome Biology, 2019, 20, 64. | 8.8 | 114 |
| 15 | Ampullary Cancers Harbor ELF3 Tumor Suppressor Gene Mutations and Exhibit Frequent WNT Dysregulation. Cell Reports, 2016, 14, 907-919. | 6.4 | 107 |
| 16 | Sooty mangabey genome sequence provides insight into AIDS resistance in a natural SIV host. Nature, 2018, 553, 77-81. | 27.8 | 81 |
| 17 | The Toxicogenome of <i>Hyalella azteca</i> : A Model for Sediment Ecotoxicology and Evolutionary Toxicology. Environmental Science & Ecotoxicology. 2018, 52, 6009-6022. | 10.0 | 79 |
| 18 | Lucilia cuprina genome unlocks parasitic fly biology to underpin future interventions. Nature Communications, 2015, 6, 7344. | 12.8 | 67 |

| # | ARTICLE | lF | CITATION |
|----|---|-----|----------|
| 19 | PacBio-LITS: a large-insert targeted sequencing method for characterization of human disease-associated chromosomal structural variations. BMC Genomics, 2015, 16, 214. | 2.8 | 63 |
| 20 | Brown marmorated stink bug, Halyomorpha halys (StåI), genome: putative underpinnings of polyphagy, insecticide resistance potential and biology of a top worldwide pest. BMC Genomics, 2020, 21, 227. | 2.8 | 60 |
| 21 | Comparative genomics of the miniature wasp and pest control agent Trichogramma pretiosum. BMC Biology, 2018, 16, 54. | 3.8 | 57 |
| 22 | Genome-enabled insights into the biology of thrips as crop pests. BMC Biology, 2020, 18, 142. | 3.8 | 54 |
| 23 | Genomic Signatures of Cooperation and Conflict in the Social Amoeba. Current Biology, 2015, 25, 1661-1665. | 3.9 | 51 |
| 24 | The genome of the water strider Gerris buenoi reveals expansions of gene repertoires associated with adaptations to life on the water. BMC Genomics, 2018, 19, 832. | 2.8 | 47 |
| 25 | Evidence for Stabilizing Selection on Codon Usage in Chromosomal Rearrangements of (i>Drosophila pseudoobscura (i>. G3: Genes, Genomes, Genetics, 2014, 4, 2433-2449. | 1.8 | 17 |
| 26 | Sawfly Genomes Reveal Evolutionary Acquisitions That Fostered the Mega-Radiation of Parasitoid and Eusocial Hymenoptera. Genome Biology and Evolution, 2020, 12, 1099-1188. | 2.5 | 17 |
| 27 | NMDA-evoked [Ca2+]i increase in salamander retinal ganglion cells: Modulation by PKA and adrenergic receptors. Visual Neuroscience, 2002, 19, 249-256. | 1.0 | 12 |
| 28 | Fully resolved assembly of <i>Cryptosporidium parvum</i> . GigaScience, 2022, 11, . | 6.4 | 8 |
| 29 | SVachra: a tool to identify genomic structural variation in mate pair sequencing data containing inward and outward facing reads. BMC Genomics, 2017, 18, 691. | 2.8 | 7 |