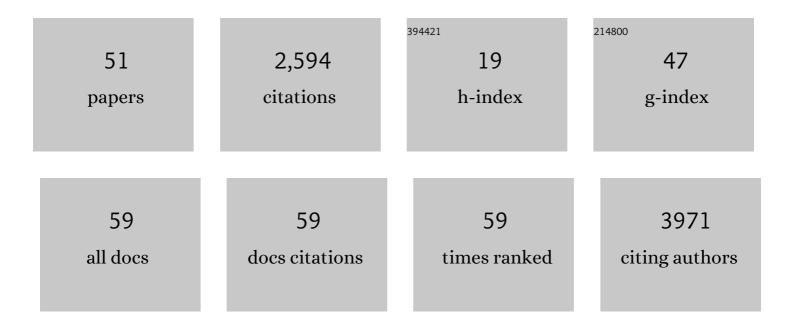
## **Dapeng Wang**

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7145539/publications.pdf Version: 2024-02-01



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| #  | Article  | IF   | CITATIONS |
|----|--|------|-----------|
| 1  | KaKs_Calculator 2.0: A Toolkit Incorporating Gamma-Series Methods and Sliding Window Strategies.<br>Genomics, Proteomics and Bioinformatics, 2010, 8, 77-80.   | 6.9  | 1,301     |
| 2  | A blood atlas of COVID-19 defines hallmarks of disease severity and specificity. Cell, 2022, 185, 916-938.e58.   | 28.9 | 164       |
| 3  | An immunodominant NP105–113-B*07:02 cytotoxic T cell response controls viral replication and is associated with less severe COVID-19 disease. Nature Immunology, 2022, 23, 50-61.  | 14.5 | 110       |
| 4  | Identification of LZTFL1 as a candidate effector gene at a COVID-19 risk locus. Nature Genetics, 2021, 53, 1606-1615.  | 21.4 | 93        |
| 5  | Single-Cell Network Analysis Identifies DDIT3 as a Nodal Lineage Regulator in Hematopoiesis. Cell<br>Reports, 2015, 11, 1503-1510.   | 6.4  | 70        |
| 6  | A Human IPS Model Implicates Embryonic B-Myeloid Fate Restriction as Developmental Susceptibility to<br>BÂAcute Lymphoblastic Leukemia-Associated ETV6-RUNX1. Developmental Cell, 2018, 44, 362-377.e7.                            | 7.0  | 65        |
| 7  | Prevalence and antimicrobial susceptibility of Vibrio parahaemolyticus isolated from retail shellfish<br>in Shanghai. Food Control, 2016, 60, 263-268.   | 5.5  | 60        |
| 8  | Distribution of norovirus in oyster tissues. Journal of Applied Microbiology, 2008, 105, 1966-1972.  | 3.1  | 51        |
| 9  | Inactivation conditions for human norovirus measured by an in situ capture-qRT-PCR method.<br>International Journal of Food Microbiology, 2014, 172, 76-82.  | 4.7  | 42        |
| 10 | Nonsynonymous substitution rate (Ka) is a relatively consistent parameter for defining fast-evolving and slow-evolving protein-coding genes. Biology Direct, 2011, 6, 13.  | 4.6  | 37        |
| 11 | Phosphoproteomic analysis reveals plant <scp>DNA</scp> damage signalling pathways with a functional role for histone H2 <scp>AX</scp> phosphorylation in plant growth under genotoxic stress. Plant Journal, 2019, 100, 1007-1021. | 5.7  | 37        |
| 12 | Retention of Vibrio parahaemolyticus in oyster tissues after chlorine dioxide treatment. International<br>Journal of Food Microbiology, 2010, 137, 76-80.  | 4.7  | 36        |
| 13 | Bacteriophage potential against Vibrio parahaemolyticus biofilms. Food Control, 2019, 98, 156-163.   | 5.5  | 34        |
| 14 | hppRNA—a Snakemake-based handy parameter-free pipeline for RNA-Seq analysis of numerous samples.<br>Briefings in Bioinformatics, 2018, 19, bbw143.   | 6.5  | 32        |
| 15 | Prevalence and characterization of Salmonella serovars isolated from farm products in Shanghai.<br>Food Control, 2018, 85, 269-275.  | 5.5  | 32        |
| 16 | A Novel Role for Minimal Introns: Routing mRNAs to the Cytosol. PLoS ONE, 2010, 5, e10144.   | 2.5  | 27        |
| 17 | New <i>In Situ</i> Capture Quantitative (Real-Time) Reverse Transcription-PCR Method as an<br>Alternative Approach for Determining Inactivation of Tulane Virus. Applied and Environmental<br>Microbiology, 2014, 80, 2120-2124.   | 3.1  | 26        |
| 18 | The Rice Genome Knowledgebase (RGKbase): an annotation database for rice comparative genomics and evolutionary biology. Nucleic Acids Research, 2012, 41, D1199-D1205.   | 14.5 | 25        |

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|----|---|-----|-----------|
| 19 | Hypoxic adaptation of leukemic cells infiltrating the CNS affords a therapeutic strategy targeting VEGFA. Blood, 2017, 129, 3126-3129.  | 1.4 | 23        |
| 20 | Development and evaluation of a novel in situ target-capture approach for aptamer selection of human noroviruses. Talanta, 2019, 193, 199-205.  | 5.5 | 20        |
| 21 | Endometrium On-a-Chip Reveals Insulin- and Glucose-induced Alterations in the Transcriptome and Proteomic Secretome. Endocrinology, 2021, 162, .  | 2.8 | 18        |
| 22 | Trehalose-Induced Remodelling of the Human Microbiota Affects Clostridioides difficile Infection<br>Outcome in an In Vitro Colonic Model: A Pilot Study. Frontiers in Cellular and Infection<br>Microbiology, 2021, 11, 670935. | 3.9 | 18        |
| 23 | Both Size and GC-Content of Minimal Introns Are Selected in Human Populations. PLoS ONE, 2011, 6, e17945.   | 2.5 | 18        |
| 24 | Transposon-Derived and Satellite-Derived Repetitive Sequences Play Distinct Functional Roles in Mammalian Intron Size Expansion. Evolutionary Bioinformatics, 2012, 8, EBO.S9758.   | 1.2 | 17        |
| 25 | Seasonal dynamics and diversity of bacteria in retail oyster tissues. International Journal of Food<br>Microbiology, 2014, 173, 14-20.  | 4.7 | 17        |
| 26 | Cytoplasmic long noncoding RNAs are differentially regulated and translated during human neuronal differentiation. Rna, 2021, 27, 1082-1101.  | 3.5 | 17        |
| 27 | The role of CAPG in molecular communication between the embryo and the uterine endometrium: Is its function conserved in species with different implantation strategies?. FASEB Journal, 2020, 34, 11015-11029.                 | 0.5 | 15        |
| 28 | Redesigned Duplex RT-qPCR for the Detection of GI and GII Human Noroviruses. Engineering, 2020, 6, 442-448.   | 6.7 | 15        |
| 29 | Surveillance of human norovirus in oysters collected from production area in Shandong Province,<br>China during 2017–2018. Food Control, 2021, 121, 107649.   | 5.5 | 15        |
| 30 | Seed DNA damage responses promote germination and growth in <i>Arabidopsis thaliana</i> .<br>Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .                                      | 7.1 | 15        |
| 31 | Engineering Bacterial Surface Displayed Human Norovirus Capsid Proteins: A Novel System to Explore<br>Interaction Between Norovirus and Ligands. Frontiers in Microbiology, 2015, 6, 1448.                                      | 3.5 | 12        |
| 32 | In Situ Capture RT-qPCR: A New Simple and Sensitive Method to Detect Human Norovirus in Oysters.<br>Frontiers in Microbiology, 2017, 8, 554.  | 3.5 | 12        |
| 33 | Characterization of a Histo-Blood Group Antigen-like Substance in Romaine Lettuce That Contributes to Human Norovirus Attachment. Journal of Agricultural and Food Chemistry, 2020, 68, 1207-1212.                              | 5.2 | 12        |
| 34 | Culturable bacteria resident on lettuce might contribute to accumulation of human noroviruses.<br>International Journal of Food Microbiology, 2020, 317, 108492.  | 4.7 | 11        |
| 35 | Fingerprinting of human noroviruses co-infections in a possible foodborne outbreak by metagenomics. International Journal of Food Microbiology, 2020, 333, 108787.  | 4.7 | 11        |
| 36 | Bacterial Surface-Displayed GII.4 Human Norovirus Capsid Proteins Bound to HBGA-Like Molecules in<br>Romaine Lettuce. Frontiers in Microbiology, 2017, 8, 251.  | 3.5 | 10        |

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|----|---|------|-----------|
| 37 | A Bacterial Surface Display System Expressing Cleavable Capsid Proteins of Human Norovirus: A Novel<br>System to Discover Candidate Receptors. Frontiers in Microbiology, 2017, 8, 2405.            | 3.5  | 10        |
| 38 | Broad-range and effective detection of human noroviruses by colloidal gold<br>immunochromatographic assay based on the shell domain of the major capsid protein. BMC<br>Microbiology, 2021, 21, 22. | 3.3  | 9         |
| 39 | LCGbase: A Comprehensive Database for Lineage-Based Co-regulated Genes. Evolutionary<br>Bioinformatics, 2012, 8, EBO.S8540.   | 1.2  | 7         |
| 40 | GCevobase: an evolution-based database for GC content in eukaryotic genomes. Bioinformatics, 2018, 34, 2129-2131.   | 4.1  | 6         |
| 41 | IntronDB: a database for eukaryotic intron features. Bioinformatics, 2019, 35, 4400-4401.   | 4.1  | 6         |
| 42 | Oyster Heat Shock Protein 70 Plays a Role in Binding of Human Noroviruses. Applied and Environmental Microbiology, 2021, 87, e0079021.  | 3.1  | 6         |
| 43 | KGCAK: a K-mer based database for genome-wide phylogeny and complexity evaluation. Biology Direct, 2015, 10, 53.  | 4.6  | 5         |
| 44 | Plastid-LCGbase: a collection of evolutionarily conserved plastid-associated gene pairs. Nucleic Acids Research, 2015, 43, D990-D995.   | 14.5 | 4         |
| 45 | Functional Networking of Human Divergently Paired Genes (DPGs). PLoS ONE, 2013, 8, e78896.  | 2.5  | 3         |
| 46 | An Effective Platform for Exploring Rotavirus Receptors by Bacterial Surface Display System.<br>Virologica Sinica, 2020, 35, 103-109.   | 3.0  | 3         |
| 47 | Library Preparation Based on Transposase Assisted RNA/DNA Hybrid Co-Tagmentation for Next-Generation Sequencing of Human Noroviruses. Viruses, 2021, 13, 65.  | 3.3  | 3         |
| 48 | Detection of group A rotavirus in oyster tissues by in situ capture RT-qPCR. Food Control, 2021, 127, 108161.   | 5.5  | 2         |
| 49 | LCCserver: A Webserver for Exploring Evolutionary Trajectory of Gene Orders in a Large Number of Genomes. OMICS A Journal of Integrative Biology, 2015, 19, 574-577.                                | 2.0  | 0         |
| 50 | DLCP: A database for lineage-conserved and lineage-specific gene pairs in animal and plant genomes.<br>Biochemical and Biophysical Research Communications, 2016, 469, 542-545.                     | 2.1  | 0         |
| 51 | RNA-Seq analysis of a Pax3-expressing myoblast clone in-vitro and effect of culture surface stiffness on differentiation. Scientific Reports, 2022, 12, 2841.                                       | 3.3  | 0         |