## Miao Gui

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1970960/publications.pdf

Version: 2024-02-01

687363 1058476 1,980 14 13 14 citations h-index g-index papers 15 15 15 4016 citing authors all docs docs citations times ranked

| #  | Article  | IF   | CITATIONS |
|----|--|------|-----------|
| 1  | Cryo-EM structure of the SARS coronavirus spike glycoprotein in complex with its host cell receptor ACE2. PLoS Pathogens, 2018, 14, e1007236.  | 4.7  | 716       |
| 2  | Cryo-electron microscopy structures of the SARS-CoV spike glycoprotein reveal a prerequisite conformational state for receptor binding. Cell Research, 2017, 27, 119-129.                              | 12.0 | 547       |
| 3  | Structural and molecular basis for Ebola virus neutralization by protective human antibodies.<br>Science, 2016, 351, 1343-1346.  | 12.6 | 176       |
| 4  | Structures of radial spokes and associated complexes important for ciliary motility. Nature Structural and Molecular Biology, 2021, 28, 29-37.   | 8.2  | 81        |
| 5  | De novo identification of mammalian ciliary motility proteins using cryo-EM. Cell, 2021, 184, 5791-5806.e19.   | 28.9 | 73        |
| 6  | Cryo-EM structure of an activated GPCR–G protein complex in lipid nanodiscs. Nature Structural and Molecular Biology, 2021, 28, 258-267.   | 8.2  | 71        |
| 7  | Electron microscopy studies of the coronavirus ribonucleoprotein complex. Protein and Cell, 2017, 8, 219-224.  | 11.0 | 62        |
| 8  | Structure and activation mechanism of the BBSome membrane protein trafficking complex. ELife, 2020, 9, .   | 6.0  | 62        |
| 9  | Potent neutralizing monoclonal antibodies against Ebola virus infection. Scientific Reports, 2016, 6, 25856.   | 3.3  | 46        |
| 10 | Structural assembly of the tailed bacteriophage ï•29. Nature Communications, 2019, 10, 2366.   | 12.8 | 44        |
| 11 | The bacteriophage ϕ29 tail possesses a pore-forming loop for cell membrane penetration. Nature, 2016, 534, 544-547.  | 27.8 | 33        |
| 12 | Ciliary central apparatus structure reveals mechanisms of microtubule patterning. Nature Structural and Molecular Biology, 2022, 29, 483-492.  | 8.2  | 33        |
| 13 | The interactions between mitochondria and sarcoplasmic reticulum and the proteome characterization of mitochondrionâ€associated membrane from rabbit skeletal muscle. Proteomics, 2015, 15, 2701-2704. | 2.2  | 21        |
| 14 | Structural intermediates in the low pH-induced transition of influenza hemagglutinin. PLoS Pathogens, 2020, 16, e1009062.  | 4.7  | 15        |