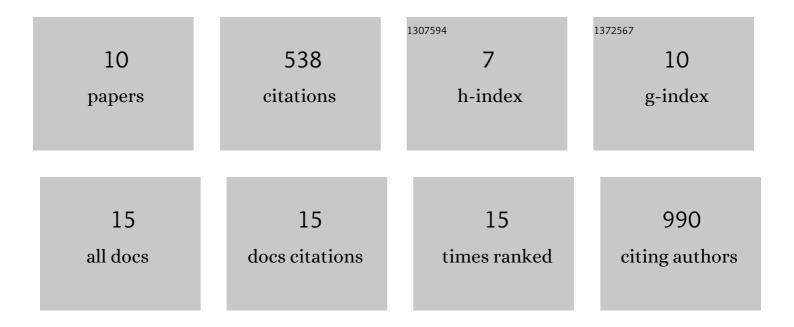
Hyeonuk Woo

List of Publications by Year in descending order

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| # | Article | IF | CITATIONS |
|----|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------|-----------|
| 1 | Structural basis of neuropeptide Y signaling through Y1 receptor. Nature Communications, 2022, 13, 853. | 12.8 | 20 |
| 2 | Cryo-EM structure of the human somatostatin receptor 2 complex with its agonist somatostatin delineates the ligand-binding specificity. ELife, 2022, 11, . | 6.0 | 10 |
| 3 | Structure, Dynamics, Receptor Binding, and Antibody Binding of the Fully Glycosylated Full-Length SARS-CoV-2 Spike Protein in a Viral Membrane. Journal of Chemical Theory and Computation, 2021, 17, 2479-2487. | 5.3 | 62 |
| 4 | Protein oligomer structure prediction using GALAXY in CASP14. Proteins: Structure, Function and Bioinformatics, 2021, 89, 1844-1851. | 2.6 | 4 |
| 5 | Prediction of protein assemblies, the next frontier: The <scp>CASP14â€CAPRI</scp> experiment. Proteins: Structure, Function and Bioinformatics, 2021, 89, 1800-1823. | 2.6 | 73 |
| 6 | Dynamic Interactions of Fully Glycosylated SARS-CoV-2 Spike Protein with Various Antibodies. Journal of Chemical Theory and Computation, 2021, 17, 6559-6569. | 5.3 | 13 |
| 7 | Structure prediction of biological assemblies using GALAXY in CAPRI rounds 38â€45. Proteins: Structure, Function and Bioinformatics, 2020, 88, 1009-1017. | 2.6 | 5 |
| 8 | Developing a Fully Glycosylated Full-Length SARS-CoV-2 Spike Protein Model in a Viral Membrane. Journal of Physical Chemistry B, 2020, 124, 7128-7137. | 2.6 | 240 |
| 9 | Prediction of protein oligomer structures using GALAXY in CASP13. Proteins: Structure, Function and Bioinformatics, 2019, 87, 1233-1240. | 2.6 | 6 |
| 10 | Blind prediction of homo―and heteroâ€protein complexes: The CASP13â€CAPRI experiment. Proteins: Structure, Function and Bioinformatics, 2019, 87, 1200-1221. | 2.6 | 99 |