## Yee Ying Lim

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4532198/publications.pdf

Version: 2024-02-01

|          |                | 1937685      | 1588992        |  |
|----------|----------------|--------------|----------------|--|
| 8        | 55             | 4            | 8              |  |
| papers   | citations      | h-index      | g-index        |  |
|          |                |              |                |  |
|          |                |              |                |  |
| 0        | 0              | 0            | 126            |  |
| 8        | 8              | 8            | 126            |  |
| all docs | docs citations | times ranked | citing authors |  |
|          |                |              |                |  |

| # | Article   | IF  | CITATIONS |
|---|---|-----|-----------|
| 1 | G-Quadruplexes as An Alternative Recognition Element in Disease-Related Target Sensing. Molecules, 2019, 24, 1079.  | 3.8 | 26        |
| 2 | Broad specificity of immune helminth scFv library to identify monoclonal antibodies targeting Strongyloides. Scientific Reports, 2021, 11, 2502.  | 3.3 | 8         |
| 3 | Computer-Aided Antibody Design: An Overview. Advances in Experimental Medicine and Biology, 2017, 1053, 221-243.  | 1.6 | 6         |
| 4 | Theoretical study of the interactions between peptide tyrosine tyrosine [PYY (1-36)], a newly identified modulator in type 2 diabetes pathophysiology, with receptors NPY1R and NPY4R. Hormones, 2021, 20, 557-569. | 1.9 | 5         |
| 5 | Human IgG1 Fc pH-dependent optimization from a constant pH molecular dynamics simulation analysis.<br>RSC Advances, 2020, 10, 13066-13075.  | 3.6 | 3         |
| 6 | Data curation to improve the pattern recognition performance of B-cell epitope prediction by support vector machine. Pure and Applied Chemistry, 2021, 93, 571-577.   | 1.9 | 3         |
| 7 | Development and structural characterisation of human scFv targeting MDM2 spliced variant MDM215kDa. Molecular Immunology, 2021, 135, 191-203.   | 2.2 | 3         |
| 8 | Structural approaches for the DNA binding motifs prediction in Bacillus thuringiensis sigma-E transcription factor ( $if$ ETF). Journal of Molecular Modeling, 2019, 25, 301.                                       | 1.8 | 1         |