

Sheng-ce Tao

List of Publications by Year in descending order

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Version: 2024-02-01

119
papers

5,286
citations

109321

35
h-index

102487

66
g-index

139
all docs

139
docs citations

139
times ranked

9481
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|------|-----------|
| 1 | Structural basis for inhibition of the RNA-dependent RNA polymerase from SARS-CoV-2 by remdesivir. <i>Science</i> , 2020, 368, 1499-1504. | 12.6 | 950 |
| 2 | SARS-CoV-2 Orf9b suppresses type I interferon responses by targeting TOM70. <i>Cellular and Molecular Immunology</i> , 2020, 17, 998-1000. | 10.5 | 280 |
| 3 | Protein Acetylation Microarray Reveals that NuA4 Controls Key Metabolic Target Regulating Gluconeogenesis. <i>Cell</i> , 2009, 136, 1073-1084. | 28.9 | 279 |
| 4 | SARS-CoV-2 proteome microarray for global profiling of COVID-19 specific IgG and IgM responses. <i>Nature Communications</i> , 2020, 11, 3581. | 12.8 | 251 |
| 5 | Lectin microarrays identify cell-specific and functionally significant cell surface glycan markers. <i>Glycobiology</i> , 2008, 18, 761-769. | 2.5 | 184 |
| 6 | Linear epitopes of SARS-CoV-2 spike protein elicit neutralizing antibodies in COVID-19 patients. <i>Cellular and Molecular Immunology</i> , 2020, 17, 1095-1097. | 10.5 | 168 |
| 7 | Linear epitope landscape of the SARS-CoV-2 Spike protein constructed from 1,051 COVID-19 patients. <i>Cell Reports</i> , 2021, 34, 108915. | 6.4 | 127 |
| 8 | Systematic identification of arsenic-binding proteins reveals that hexokinase-2 is inhibited by arsenic. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 15084-15089. | 7.1 | 126 |
| 9 | A proteome chip approach reveals new DNA damage recognition activities in <i>Escherichia coli</i> . <i>Nature Methods</i> , 2008, 5, 69-74. | 19.0 | 121 |
| 10 | Antibody dynamics to SARS-CoV-2 in asymptomatic COVID-19 infections. <i>Allergy: European Journal of Allergy and Clinical Immunology</i> , 2021, 76, 551-561. | 5.7 | 107 |
| 11 | Protein chip fabrication by capture of nascent polypeptides. <i>Nature Biotechnology</i> , 2006, 24, 1253-1254. | 17.5 | 90 |
| 12 | Advances and Utility of the Human Plasma Proteome. <i>Journal of Proteome Research</i> , 2021, 20, 5241-5263. | 3.7 | 86 |
| 13 | Identification of Serum Biomarkers for Gastric Cancer Diagnosis Using a Human Proteome Microarray. <i>Molecular and Cellular Proteomics</i> , 2016, 15, 614-623. | 3.8 | 82 |
| 14 | Global analysis of the glycoproteome in <i>Saccharomyces cerevisiae</i> reveals new roles for protein glycosylation in eukaryotes. <i>Molecular Systems Biology</i> , 2009, 5, 308. | 7.2 | 79 |
| 15 | Mycobacterium Tuberculosis Proteome Microarray for Global Studies of Protein Function and Immunogenicity. <i>Cell Reports</i> , 2014, 9, 2317-2329. | 6.4 | 77 |
| 16 | Cordycepin induces cell cycle arrest and apoptosis by inducing DNA damage and up-regulation of p53 in Leukemia cells. <i>Cell Cycle</i> , 2015, 14, 761-771. | 2.6 | 75 |
| 17 | Current applications of antibody microarrays. <i>Clinical Proteomics</i> , 2018, 15, 7. | 2.1 | 75 |
| 18 | Detection and Verification of Glycosylation Patterns of Glycoproteins from Clinical Specimens Using Lectin Microarrays and Lectin-Based Immunosorbent Assays. <i>Analytical Chemistry</i> , 2011, 83, 8509-8516. | 6.5 | 71 |

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|----|--|------|-----------|
| 19 | Functional Dissection of a HECT Ubiquitin E3 Ligase. <i>Molecular and Cellular Proteomics</i> , 2008, 7, 35-45. | 3.8 | 70 |
| 20 | Applications of Protein Microarray Technology. <i>Combinatorial Chemistry and High Throughput Screening</i> , 2007, 10, 706-718. | 1.1 | 67 |
| 21 | Microfluidic chip integrating high throughput continuous-flow PCR and DNA hybridization for bacteria analysis. <i>Talanta</i> , 2014, 122, 246-250. | 5.5 | 64 |
| 22 | Global Profiling of Protein Lysine Malonylation in <i>Escherichia coli</i> Reveals Its Role in Energy Metabolism. <i>Journal of Proteome Research</i> , 2016, 15, 2060-2071. | 3.7 | 63 |
| 23 | Bcl2-associated Athanogene 3 Interactome Analysis Reveals a New Role in Modulating Proteasome Activity. <i>Molecular and Cellular Proteomics</i> , 2013, 12, 2804-2819. | 3.8 | 62 |
| 24 | MACRO: A Combined Microchip-PCR and Microarray System for High-Throughput Monitoring of Genetically Modified Organisms. <i>Analytical Chemistry</i> , 2014, 86, 1269-1276. | 6.5 | 61 |
| 25 | Current Technologies for Complex Glycoproteomics and Their Applications to Biology/Disease-Driven Glycoproteomics. <i>Journal of Proteome Research</i> , 2018, 17, 4097-4112. | 3.7 | 60 |
| 26 | Skp1 in lung cancer: clinical significance and therapeutic efficacy of its small molecule inhibitors. <i>Oncotarget</i> , 2015, 6, 34953-34967. | 1.8 | 53 |
| 27 | YcgC represents a new protein deacetylase family in prokaryotes. <i>ELife</i> , 2015, 4, . | 6.0 | 52 |
| 28 | Characterization of Protein Lysine Propionylation in <i>Escherichia coli</i> : Global Profiling, Dynamic Change, and Enzymatic Regulation. <i>Journal of Proteome Research</i> , 2016, 15, 4696-4708. | 3.7 | 50 |
| 29 | Lectin RCA-I specifically binds to metastasis-associated cell surface glycans in triple-negative breast cancer. <i>Breast Cancer Research</i> , 2015, 17, 36. | 5.0 | 48 |
| 30 | Protein Arrays on Patterned Porous Gold Substrates Interrogated with Mass Spectrometry: Detection of Peptides in Plasma. <i>Analytical Chemistry</i> , 2008, 80, 1448-1458. | 6.5 | 47 |
| 31 | Protein microarrays for systems biology. <i>Acta Biochimica Et Biophysica Sinica</i> , 2011, 43, 161-171. | 2.0 | 47 |
| 32 | Systematic evaluation of IgG responses to SARS-CoV-2 spike protein-derived peptides for monitoring COVID-19 patients. <i>Cellular and Molecular Immunology</i> , 2021, 18, 621-631. | 10.5 | 43 |
| 33 | Identification of Novel 14-3-3 Interacting Proteins by Quantitative Immunoprecipitation Combined with Knockdown (QUICK). <i>Journal of Proteome Research</i> , 2010, 9, 5848-5858. | 3.7 | 40 |
| 34 | The Ser/Thr Protein Kinase Protein-Protein Interaction Map of <i>M. tuberculosis</i> *. <i>Molecular and Cellular Proteomics</i> , 2017, 16, 1491-1506. | 3.8 | 39 |
| 35 | Multiplex sample-to-answer detection of bacteria using a pipette-actuated capillary array comb with integrated DNA extraction, isothermal amplification, and smartphone detection. <i>Lab on A Chip</i> , 2018, 18, 2854-2864. | 6.0 | 37 |
| 36 | Profiling Lipid-protein Interactions Using Nonquenched Fluorescent Liposomal Nanovesicles and Proteome Microarrays. <i>Molecular and Cellular Proteomics</i> , 2012, 11, 1177-1190. | 3.8 | 36 |

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|----|---|-----|-----------|
| 37 | Global identification of <i>O</i> -GlcNAc transferase (OGT) interactors by a human proteome microarray and the construction of an OGT interactome. <i>Proteomics</i> , 2014, 14, 1020-1030. | 2.2 | 35 |
| 38 | Global identification of CobB interactors by an <i>Escherichia coli</i> proteome microarray. <i>Acta Biochimica Et Biophysica Sinica</i> , 2014, 46, 548-555. | 2.0 | 33 |
| 39 | Comprehensive profiling of accessible surface glycans of mammalian sperm using a lectin microarray. <i>Clinical Proteomics</i> , 2014, 11, 10. | 2.1 | 32 |
| 40 | Antibody landscape against SARS-CoV-2 reveals significant differences between non-structural/accessory and structural proteins. <i>Cell Reports</i> , 2021, 36, 109391. | 6.4 | 32 |
| 41 | Longitudinal serum autoantibody repertoire profiling identifies surgery-associated biomarkers in lung adenocarcinoma. <i>EBioMedicine</i> , 2020, 53, 102674. | 6.1 | 30 |
| 42 | Antibody Binding Epitope Mapping (AbMap) of Hundred Antibodies in a Single Run. <i>Molecular and Cellular Proteomics</i> , 2021, 20, 100059. | 3.8 | 30 |
| 43 | Systematic profiling of SARS-CoV-2-specific IgG responses elicited by an inactivated virus vaccine identifies peptides and proteins for predicting vaccination efficacy. <i>Cell Discovery</i> , 2021, 7, 67. | 6.7 | 29 |
| 44 | Quantitative Phosphoproteomics of Proteasome Inhibition in Multiple Myeloma Cells. <i>PLoS ONE</i> , 2010, 5, e13095. | 2.5 | 28 |
| 45 | Reversibly acetylated lysine residues play important roles in the enzymatic activity of <i>Escherichia coli</i> <i>N</i> -hydroxyarylamine <i>O</i> -acetyltransferase. <i>FEBS Journal</i> , 2013, 280, 1966-1979. | 4.7 | 28 |
| 46 | Interplay between the bacterial protein deacetylase CobB and the second messenger c-di-GMP . <i>EMBO Journal</i> , 2019, 38, e100948. | 7.8 | 28 |
| 47 | Lectin binding of human sperm associates with DEFB126 mutation and serves as a potential biomarker for subfertility. <i>Scientific Reports</i> , 2016, 6, 20249. | 3.3 | 25 |
| 48 | Cyclic di-GMP regulates Mycobacterium tuberculosis resistance to ethionamide. <i>Scientific Reports</i> , 2017, 7, 5860. | 3.3 | 25 |
| 49 | RIG-I regulates myeloid differentiation by promoting TRIM25-mediated ISGylation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 14395-14404. | 7.1 | 25 |
| 50 | A graphene oxide coated tapered microfiber acting as a super-sensor for rapid detection of SARS-CoV-2. <i>Lab on A Chip</i> , 2021, 21, 2398-2406. | 6.0 | 25 |
| 51 | Fluorescent Protein Nanowire-Mediated Protein Microarrays for Multiplexed and Highly Sensitive Pathogen Detection. <i>ACS Applied Materials & Interfaces</i> , 2016, 8, 17472-17477. | 8.0 | 24 |
| 52 | Multiplex and visual detection of African Swine Fever Virus (ASFV) based on Hive-Chip and direct loop-mediated isothermal amplification. <i>Analytica Chimica Acta</i> , 2020, 1140, 30-40. | 5.4 | 23 |
| 53 | Quantitative Proteomic Analysis of Tumor Reversion in Multiple Myeloma Cells. <i>Journal of Proteome Research</i> , 2011, 10, 845-855. | 3.7 | 22 |
| 54 | Discovering cancer biomarkers from clinical samples by protein microarrays. <i>Proteomics - Clinical Applications</i> , 2015, 9, 98-110. | 1.6 | 22 |

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|----|---|------|-----------|
| 55 | A Human Lectin Microarray for Sperm Surface Glycosylation Analysis. <i>Molecular and Cellular Proteomics</i> , 2016, 15, 2839-2851. | 3.8 | 22 |
| 56 | Proteomic identification of the oncoprotein STAT3 as a target of a novel Skp1 inhibitor. <i>Oncotarget</i> , 2017, 8, 2681-2693. | 1.8 | 22 |
| 57 | An array of 60,000 antibodies for proteome-scale antibody generation and target discovery. <i>Science Advances</i> , 2020, 6, eaax2271. | 10.3 | 22 |
| 58 | Protein Microarrays for Studies of Drug Mechanisms and Biomarker Discovery in the Era of Systems Biology. <i>Current Pharmaceutical Design</i> , 2014, 20, 49-55. | 1.9 | 22 |
| 59 | A universal multiplex PCR strategy for 100-plex amplification using a hydrophobically patterned microarray. <i>Lab on A Chip</i> , 2011, 11, 3609. | 6.0 | 21 |
| 60 | Visual detection of multiple genetically modified organisms in a capillary array. <i>Lab on A Chip</i> , 2017, 17, 521-529. | 6.0 | 21 |
| 61 | Comparison of Different Methods for Preparing Single Stranded DNA for Oligonucleotide Microarray. <i>Analytical Letters</i> , 2003, 36, 2849-2863. | 1.8 | 20 |
| 62 | Proteome microarray technology and application: higher, wider, and deeper. <i>Expert Review of Proteomics</i> , 2019, 16, 815-827. | 3.0 | 19 |
| 63 | Systematic Identification of Mycobacterium tuberculosis Effectors Reveals that BfrB Suppresses Innate Immunity. <i>Molecular and Cellular Proteomics</i> , 2017, 16, 2243-2253. | 3.8 | 18 |
| 64 | Lectin Microarrays: A Powerful Tool for Glycan-Based Biomarker Discovery. <i>Combinatorial Chemistry and High Throughput Screening</i> , 2011, 14, 711-719. | 1.1 | 16 |
| 65 | Construction of a metabolomics profile of arsenic trioxide effect in gastric carcinoma cell line SGC7901. <i>Acta Biochimica Et Biophysica Sinica</i> , 2016, 48, 474-481. | 2.0 | 15 |
| 66 | Systematic profiling of SARS-CoV-2-specific IgG epitopes at amino acid resolution. <i>Cellular and Molecular Immunology</i> , 2021, 18, 1067-1069. | 10.5 | 14 |
| 67 | Nsp2 has the potential to be a drug target revealed by global identification of SARS-CoV-2 Nsp2-interacting proteins. <i>Acta Biochimica Et Biophysica Sinica</i> , 2021, 53, 1134-1141. | 2.0 | 14 |
| 68 | Landscape of the RBD-specific IgG, IgM, and IgA responses triggered by the inactivated virus vaccine against the Omicron variant. <i>Cell Discovery</i> , 2022, 8, 15. | 6.7 | 14 |
| 69 | Identification of Serum Biomarkers for Systemic Lupus Erythematosus Using a Library of Phage Displayed Random Peptides and Deep Sequencing. <i>Molecular and Cellular Proteomics</i> , 2019, 18, 1851-1863. | 3.8 | 13 |
| 70 | The design and application of DNA chips for early detection of SARS-CoV from clinical samples. <i>Journal of Clinical Virology</i> , 2005, 33, 123-131. | 3.1 | 11 |
| 71 | Proteomic analysis of multiple myeloma: Current status and future perspectives. <i>Proteomics - Clinical Applications</i> , 2011, 5, 30-37. | 1.6 | 11 |
| 72 | PMD: A Resource for Archiving and Analyzing Protein Microarray data. <i>Scientific Reports</i> , 2016, 6, 19956. | 3.3 | 11 |

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|----|--|-----|-----------|
| 73 | Global Profiling of PknG Interactions Using a Human Proteome Microarray Reveals Novel Connections with CypA. <i>Proteomics</i> , 2018, 18, e1800265. | 2.2 | 11 |
| 74 | Fatty acylCoA synthetase FadD13 regulates proinflammatory cytokine secretion dependent on the NF- κ B signalling pathway by binding to eEF1A1. <i>Cellular Microbiology</i> , 2019, 21, e13090. | 2.1 | 11 |
| 75 | Blocking oligo ² a novel approach for improving chip-based DNA hybridization efficiency. <i>Molecular and Cellular Probes</i> , 2003, 17, 197-202. | 2.1 | 10 |
| 76 | An integrated micro-electro-fluidic and protein arraying system for parallel analysis of cell responses to controlled microenvironments. <i>Integrative Biology (United Kingdom)</i> , 2010, 2, 416. | 1.3 | 10 |
| 77 | Systematic identification of the protein substrates of UDP-GalNAc:polypeptide N-acetylgalactosaminyltransferase ^{1/2/3} using a human proteome microarray. <i>Proteomics</i> , 2017, 17, 1600485. | 2.2 | 10 |
| 78 | Effects of Cryopreservation on Human Sperm Glycocalyx. <i>Reproductive and Developmental Medicine</i> , 2017, 1, 233-238. | 0.5 | 10 |
| 79 | Sensitive Detection of SARS Coronavirus RNA by a Novel Asymmetric Multiplex Nested RT-PCR Amplification Coupled With Oligonucleotide Microarray Hybridization. , 2005, 114, 59-78. | | 9 |
| 80 | Global Identification of Prokaryotic Glycoproteins Based on an Escherichia coli Proteome Microarray. <i>PLoS ONE</i> , 2012, 7, e49080. | 2.5 | 9 |
| 81 | Identification of Serine 119 as an Effective Inhibitor Binding Site of M. tuberculosis Ubiquitin-like Protein Ligase PafA Using Purified Proteins and M. smegmatis. <i>EBioMedicine</i> , 2018, 30, 225-236. | 6.1 | 9 |
| 82 | Comparative analysis of human sperm glycocalyx from different freezability ejaculates by lectin microarray and identification of ABA as sperm freezability biomarker. <i>Clinical Proteomics</i> , 2018, 15, 19. | 2.1 | 9 |
| 83 | Unrestrictive identification of non-phosphorylation PTMs in yeast kinases by MS and PTMap. <i>Proteomics</i> , 2010, 10, 896-903. | 2.2 | 8 |
| 84 | Proteomics: addressing the challenges of multiple myeloma. <i>Acta Biochimica Et Biophysica Sinica</i> , 2011, 43, 89-95. | 2.0 | 8 |
| 85 | Functional protein microarray: an ideal platform for investigating protein binding property. <i>Frontiers in Biology</i> , 2012, 7, 336-349. | 0.7 | 8 |
| 86 | Rapid Production of Virus Protein Microarray Using Protein Microarray Fabrication through Gene Synthesis (PAGES). <i>Molecular and Cellular Proteomics</i> , 2017, 16, 288-299. | 3.8 | 8 |
| 87 | Mycobacterium tuberculosis Thymidyltransferase RmlA Is Negatively Regulated by Ser/Thr Protein Kinase PknB. <i>Frontiers in Microbiology</i> , 2021, 12, 643951. | 3.5 | 8 |
| 88 | COVID-ONE-hi: The One-stop Database for COVID-19-specific Humoral Immunity and Clinical Parameters. <i>Genomics, Proteomics and Bioinformatics</i> , 2021, 19, 669-678. | 6.9 | 8 |
| 89 | Room-Temperature Hybridization of Target DNA with Microarrays in Concentrated Solutions of Guanidine Thiocyanate. <i>BioTechniques</i> , 2003, 34, 1260-1262. | 1.8 | 7 |
| 90 | High-Throughput Lectin Microarray-Based Analysis of Live Cell Surface Glycosylation. <i>Current Protocols in Protein Science</i> , 2011, 63, Unit12.9. | 2.8 | 6 |

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|-----|---|-----|-----------|
| 91 | Phage display: an ideal platform for coupling protein to nucleic acid. <i>Acta Biochimica Et Biophysica Sinica</i> , 2021, 53, 389-399. | 2.0 | 6 |
| 92 | Anti-SARS-CoV-2 IgG responses are powerful predicting signatures for the outcome of COVID-19 patients. <i>Journal of Advanced Research</i> , 2022, 36, 133-145. | 9.5 | 6 |
| 93 | MULTIPLEX PCR FOR SIMULTANEOUS DETECTION OF HUMAN CARCINOMA-RELATED VIRUSES. <i>Analytical Letters</i> , 2002, 35, 1149-1162. | 1.8 | 5 |
| 94 | The binding epitope of sintilimab on PD-1 revealed by AbMap. <i>Acta Biochimica Et Biophysica Sinica</i> , 2021, 53, 628-635. | 2.0 | 5 |
| 95 | One-Tube Nested RT-PCR Enabled by Using a Plastic Film and its Application for the Rapid Detection of SARS-Virus. <i>Biotechnology Letters</i> , 2004, 26, 179-183. | 2.2 | 4 |
| 96 | Integrated Glycosylation Patterns of Glycoproteins and DNA Methylation Landscapes in Mammalian Oogenesis and Preimplantation Embryo Development. <i>Frontiers in Cell and Developmental Biology</i> , 2020, 8, 555. | 3.7 | 4 |
| 97 | SARS-CoV-2 spike linear epitope scanning via a peptide microarray through sera profiling. <i>STAR Protocols</i> , 2021, 2, 100707. | 1.2 | 4 |
| 98 | Global discovery the PstP interactions using Mtb proteome microarray and revealing novel connections with EthR. <i>Journal of Proteomics</i> , 2020, 215, 103650. | 2.4 | 3 |
| 99 | SARS-CoV-2 Antibody Signatures for Predicting the Outcome of COVID-19. <i>SSRN Electronic Journal</i> , 0, , . | 0.4 | 3 |
| 100 | Linear Epitope Landscape of SARS-CoV-2 Spike Protein Constructed from 1,051 COVID-19 Patients. <i>SSRN Electronic Journal</i> , 0, , . | 0.4 | 3 |
| 101 | Epitope Analysis of Anti-SARS-CoV-2 Neutralizing Antibodies. <i>Current Medical Science</i> , 2021, 41, 1065. | 1.8 | 3 |
| 102 | Visual Detection of Multiple Nucleic Acids in a Capillary Array. <i>Journal of Visualized Experiments</i> , 2017, , . | 0.3 | 2 |
| 103 | Assessment and comparison of recombinant proteins from different sources for the detection of SARS-CoV-2 infection by using protein microarray. <i>Acta Biochimica Et Biophysica Sinica</i> , 2021, 53, 262-264. | 2.0 | 2 |
| 104 | A visual multiplex PCR microchip with easy sample loading. <i>Yi Chuan = Hereditas / Zhongguo Yi Chuan Xue Hui Bian Ji</i> , 2017, 39, 525-534. | 0.2 | 2 |
| 105 | Protein Microarray: An Ideal Platform for Systems Biology. , 2012, , 113-134. | | 1 |
| 106 | Cell Lysate Microarray for Mapping the Network of Genetic Regulators for Histone Marks. <i>Molecular and Cellular Proteomics</i> , 2018, 17, 1720-1736. | 3.8 | 1 |
| 107 | RIBOi: a database for ribosome-interacting proteins. <i>Acta Biochimica Et Biophysica Sinica</i> , 2019, 51, 441-443. | 2.0 | 1 |
| 108 | SARS-CoV-2 proteome microarray for COVID-19 patient sera profiling. <i>STAR Protocols</i> , 2022, 3, 101238. | 1.2 | 1 |

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|-----|---|-----|-----------|
| 109 | Microarray partition using a recycled marker pen and neutral balsam. <i>Acta Biochimica Et Biophysica Sinica</i> , 2013, 45, 706-708. | 2.0 | 0 |
| 110 | Toward the development of magnetic tweezers for high-throughput measurement of protein-protein interactions. <i>Acta Biochimica Et Biophysica Sinica</i> , 2017, 49, 468-470. | 2.0 | 0 |
| 111 | Identification of serum biomarkers for systemic lupus erythematosus using a library of phage displayed random peptides and deep sequencing. , 2019, , . | | 0 |
| 112 | Sketching the Glycan Hallmark of Intact Cells Using Lectin Microarray. <i>ACS Symposium Series</i> , 2020, , 119-126. | 0.5 | 0 |
| 113 | Antibody Landscape Against SARS-CoV-2 Proteome Revealed Significant Differences between Non-Structural/ Accessory Proteins and Structural Proteins. <i>SSRN Electronic Journal</i> , 0, , . | 0.4 | 0 |
| 114 | Protein Deacetylase Cobb Interplays with C-Di-Gmp. <i>SSRN Electronic Journal</i> , 0, , . | 0.4 | 0 |
| 115 | Integrated Glycosylation Patterns of Glycoproteins and DNA Methylation Landscapes in Mammal Oogenesis and Preimplantation Embryo Development. <i>SSRN Electronic Journal</i> , 0, , . | 0.4 | 0 |
| 116 | Database Development for Antibody Arrays. <i>Methods in Molecular Biology</i> , 2021, 2237, 257-261. | 0.9 | 0 |
| 117 | EASINESS: E. coli Assisted Speedy affinity-maturation Evolution SyStem. <i>Frontiers in Immunology</i> , 2021, 12, 747267. | 4.8 | 0 |
| 118 | Quantitative plasma proteome profiling of COVID-19 patients with mild and moderate symptoms. <i>EBioMedicine</i> , 2022, 75, 103773. | 6.1 | 0 |
| 119 | Lectin Microarray: A Powerful Tool for Glycan Related Biomarker Discovery. <i>Combinatorial Chemistry and High Throughput Screening</i> , 2011, , . | 1.1 | 0 |