Shankha Satpathy

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

15 1,290 27 30 h-index g-index citations papers 21.5 4.03 30 2,372 avg, IF L-index ext. citations ext. papers

| # | Paper | IF | Citations |
|----|--|-------------------|--------------------|
| 27 | Integrated Proteogenomic Characterization of Clear Cell Renal Cell Carcinoma. <i>Cell</i> , 2019 , 179, 964-983 | . €8 .b | 173 |
| 26 | Time-Resolved Analysis Reveals Rapid Dynamics and Broad Scope of the CBP/p300 Acetylome. <i>Cell</i> , 2018 , 174, 231-244.e12 | 56.2 | 148 |
| 25 | Proteogenomic Characterization Reveals Therapeutic Vulnerabilities in Lung Adenocarcinoma. <i>Cell</i> , 2020 , 182, 200-225.e35 | 56.2 | 139 |
| 24 | Proteogenomic Characterization of Endometrial Carcinoma. Cell, 2020, 180, 729-748.e26 | 56.2 | 122 |
| 23 | TMT Labeling for the Masses: A Robust and Cost-efficient, In-solution Labeling Approach. <i>Molecular and Cellular Proteomics</i> , 2019 , 18, 1468-1478 | 7.6 | 106 |
| 22 | SPATA2 links CYLD to the TNF-Ireceptor signaling complex and modulates the receptor signaling outcomes. <i>EMBO Journal</i> , 2016 , 35, 1868-84 | 13 | 98 |
| 21 | Systems-wide analysis of BCR signalosomes and downstream phosphorylation and ubiquitylation. <i>Molecular Systems Biology</i> , 2015 , 11, 810 | 12.2 | 82 |
| 20 | Proteogenomic Landscape of Breast Cancer Tumorigenesis and Targeted Therapy. <i>Cell</i> , 2020 , 183, 1436 | 5- 44.5 6. | .e 3 11 |
| 19 | Proteogenomic and metabolomic characterization of human glioblastoma. <i>Cancer Cell</i> , 2021 , 39, 509-52 | 2 8. ę30 | 71 |
| 18 | Accurate Quantification of Site-specific Acetylation Stoichiometry Reveals the Impact of Sirtuin Deacetylase CobB on the Acetylome. <i>Molecular and Cellular Proteomics</i> , 2017 , 16, 759-769 | 7.6 | 54 |
| 17 | Proteogenomic insights into the biology and treatment of HPV-negative head and neck squamous cell carcinoma. <i>Cancer Cell</i> , 2021 , 39, 361-379.e16 | 24.3 | 50 |
| 16 | Rapid and deep-scale ubiquitylation profiling for biology and translational research. <i>Nature Communications</i> , 2020 , 11, 359 | 17.4 | 40 |
| 15 | Microscaled proteogenomic methods for precision oncology. <i>Nature Communications</i> , 2020 , 11, 532 | 17.4 | 31 |
| 14 | Enhancers are activated by p300/CBP activity-dependent PIC assembly, RNAPII recruitment, and pause release. <i>Molecular Cell</i> , 2021 , 81, 2166-2182.e6 | 17.6 | 19 |
| 13 | Proteome dynamics at broken replication forks reveal a distinct ATM-directed repair response suppressing DNA double-strand break ubiquitination. <i>Molecular Cell</i> , 2021 , 81, 1084-1099.e6 | 17.6 | 17 |
| 12 | A proteogenomic portrait of lung squamous cell carcinoma. <i>Cell</i> , 2021 , 184, 4348-4371.e40 | 56.2 | 15 |
| 11 | Evaluation of Advanced Precursor Determination for Tandem Mass Tag (TMT)-Based Quantitative Proteomics across Instrument Platforms. <i>Journal of Proteome Research</i> , 2019 , 18, 542-547 | 5.6 | 11 |

LIST OF PUBLICATIONS

| 1 | 10 | Neurophysiological evidence for the presence of cannabinoid CB1 receptors in the laterodorsal tegmental nucleus. <i>European Journal of Neuroscience</i> , 2014 , 40, 3635-52 | 3.5 | 10 |
|------------|----|--|------|----|
| 9 |) | SUMOylation of the ING1b tumor suppressor regulates gene transcription. <i>Carcinogenesis</i> , 2014 , 35, 2214-23 | 4.6 | 8 |
| 8 | 3 | Cancer proteogenomics: current impact and future prospects Nature Reviews Cancer, 2022, | 31.3 | 7 |
| 7 | 7 | STK3 is a therapeutic target for a subset of acute myeloid leukemias. <i>Oncotarget</i> , 2018 , 9, 25458-25473 | 3.3 | 5 |
| ϵ | 6 | SIK2 orchestrates actin-dependent host response upon infection. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021 , 118, | 11.5 | 4 |
| 5 | 5 | Genomic Profiling of Lung Adenocarcinoma in Never-Smokers. <i>Journal of Clinical Oncology</i> , 2021 , 39, 3747-3758 | 2.2 | 4 |
| 4 | 1 | A highly multiplexed quantitative phosphosite assay for biology and preclinical studies. <i>Molecular Systems Biology</i> , 2021 , 17, e10156 | 12.2 | 3 |
| 3 | } | Automating UbiFast for High-throughput and Multiplexed Ubiquitin Enrichment | | 1 |
| 2 | 2 | Automating UbiFast for High-throughput and Multiplexed Ubiquitin Enrichment. <i>Molecular and Cellular Proteomics</i> , 2021 , 20, 100154 | 7.6 | 1 |
| 1 | | Demethylating Agents as Epigenetic Anticancer Therapeutics. <i>Current Cancer Therapy Reviews</i> , 2013 , 9, 24-33 | 0.4 | |