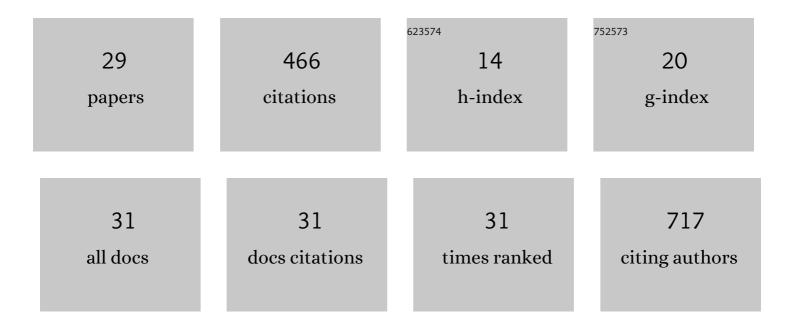
Nerea Osinalde

List of Publications by Year in descending order

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NEDEA OSINALDE

| # | Article | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | Interleukin-2 signaling pathway analysis by quantitative phosphoproteomics. Journal of Proteomics, 2011, 75, 177-191. | 1.2 | 42 |
| 2 | Fundamental constraints in synchronous muscle limit superfast motor control in vertebrates. ELife, 2017, 6, . | 2.8 | 41 |
| 3 | Nuclear Phosphoproteomic Screen Uncovers ACLY as Mediator of IL-2-induced Proliferation of CD4+ T lymphocytes. Molecular and Cellular Proteomics, 2016, 15, 2076-2092. | 2.5 | 40 |
| 4 | Differential proteomic analysis of endometrial fluid suggests increased inflammation and impaired glucose metabolism in non-implantative IVF cycles and pinpoints PYGB as a putative implantation marker. Human Reproduction, 2018, 33, 1898-1906. | 0.4 | 38 |
| 5 | Phosphoproteomic and Functional Analyses Reveal Sperm-specific Protein Changes Downstream of Kappa Opioid Receptor in Human Spermatozoa. Molecular and Cellular Proteomics, 2019, 18, S118-S131. | 2.5 | 31 |
| 6 | How to Inactivate Human Ubiquitin E3 Ligases by Mutation. Frontiers in Cell and Developmental Biology, 2020, 8, 39. | 1.8 | 31 |
| 7 | Quantitative proteomics reveals neuronal ubiquitination of Rngo/Ddi1 and several proteasomal subunits by Ube3a, accounting for the complexity of Angelman syndrome. Human Molecular Genetics, 2018, 27, 1955-1971. | 1.4 | 30 |
| 8 | Simultaneous dissection and comparison of <scp>IL</scp> â€2 and <scp>IL</scp> â€15 signaling pathways by global quantitative phosphoproteomics. Proteomics, 2015, 15, 520-531. | 1.3 | 22 |
| 9 | Targeted mass spectrometry: An emerging powerful approach to unblock the bottleneck in phosphoproteomics. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2017, 1055-1056, 29-38. | 1.2 | 22 |
| 10 | Guanine Nucleotide Exchange Factor αPIX Leads to Activation of the Rac 1 GTPase/Glycogen Phosphorylase Pathway in Interleukin (IL)-2-stimulated T Cells. Journal of Biological Chemistry, 2015, 290, 9171-9182. | 1.6 | 19 |
| 11 | Detailed Dissection of UBE3A-Mediated DDI1 Ubiquitination. Frontiers in Physiology, 2019, 10, 534. | 1.3 | 17 |
| 12 | The Nuclear Protein ALY Binds to and Modulates the Activity of Transcription Factor E2F2. Molecular and Cellular Proteomics, 2013, 12, 1087-1098. | 2.5 | 16 |
| 13 | SILAC-based quantification of changes in protein tyrosine phosphorylation induced by Interleukin-2 (IL-2) and IL-15 in T-lymphocytes. Data in Brief, 2015, 5, 53-58. | 0.5 | 16 |
| 14 | Cylindromatosis Tumor Suppressor Protein (CYLD) Deubiquitinase is Necessary for Proper Ubiquitination and Degradation of the Epidermal Growth Factor Receptor. Molecular and Cellular Proteomics, 2017, 16, 1433-1446. | 2.5 | 15 |
| 15 | Impaired proteostasis in rare neurological diseases. Seminars in Cell and Developmental Biology, 2019, 93, 164-177. | 2.3 | 14 |
| 16 | Neuronal Proteomic Analysis of the Ubiquitinated Substrates of the Disease-Linked E3 Ligases Parkin and Ube3a. BioMed Research International, 2018, 2018, 1-14. | 0.9 | 12 |
| 17 | SPANX-A/D protein subfamily plays a key role in nuclear organisation, metabolism and flagellar motility of human spermatozoa. Scientific Reports, 2020, 10, 5625. | 1.6 | 10 |
| 18 | Changes in Gab2 phosphorylation and interaction partners in response to interleukin (IL)-2 stimulation in T-lymphocytes. Scientific Reports, 2016, 6, 23530. | 1.6 | 9 |

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|----|---|-----|-----------|
| 19 | Detection of E2F-Induced Transcriptional Activity Using a Dual Luciferase Reporter Assay. Methods in Molecular Biology, 2018, 1726, 153-166. | 0.4 | 7 |
| 20 | NADH dehydrogenase complex21 is overexpressed in incipient metastatic murine colon cancer cells. Oncology Reports, 2019, 41, 742-752. | 1.2 | 7 |
| 21 | A Proteomic Approach for Systematic Mapping of Substrates of Human Deubiquitinating Enzymes. International Journal of Molecular Sciences, 2021, 22, 4851. | 1.8 | 6 |
| 22 | The ubiquitin ligase Ariadne-1 regulates neurotransmitter release via ubiquitination of NSF. Journal of Biological Chemistry, 2021, 296, 100408. | 1.6 | 6 |
| 23 | Identification of substrates for human deubiquitinating enzymes (DUBs): An up-to-date review and a case study for neurodevelopmental disorders. Seminars in Cell and Developmental Biology, 2022, 132, 120-131. | 2.3 | 4 |
| 24 | Characterization of Receptor-Associated Protein Complex Assembly in Interleukin (IL)-2- and IL-15-Activated T-Cell Lines. Journal of Proteome Research, 2017, 16, 106-121. | 1.8 | 3 |
| 25 | Mass Spectrometry-Based Characterization of Ub- and UbL-Modified Proteins. Methods in Molecular Biology, 2020, 2051, 265-276. | 0.4 | 3 |
| 26 | Kappa- opioid receptor regulates human sperm functions via SPANX-A/D protein family. Reproductive Biology, 2020, 20, 300-306. | 0.9 | 2 |
| 27 | The multifunctional role of SPANX-A/D protein subfamily in the promotion of pro-tumoural processes in human melanoma. Scientific Reports, 2021, 11, 3583. | 1.6 | 2 |
| 28 | Data on interleukin (IL)-2- and IL-15-dependent changes in IL-2R Î ² and IL-2RÎ ³ complexes. Data in Brief, 2017, 11, 499-506. | 0.5 | 0 |
| 29 | Data on mass spectrometry-based proteomics for studying the involvement of CYLD in the ubiquitination events downstream of EGFR activation. Data in Brief, 2018, 18, 1856-1863. | 0.5 | 0 |