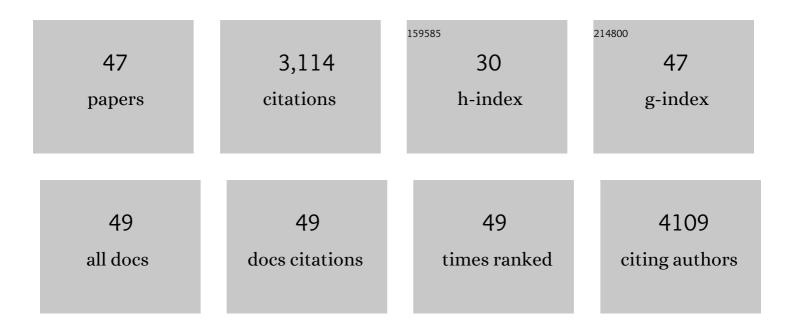
## Adnan Halim

List of Publications by Year in descending order

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Δηνιάν Ηλιίμα

| #  | Article  | IF   | CITATIONS |
|----|--|------|-----------|
| 1  | Enrichment of glycopeptides for glycan structure and attachment site identification. Nature<br>Methods, 2009, 6, 809-811.  | 19.0 | 309       |
| 2  | Engineered CHO cells for production of diverse, homogeneous glycoproteins. Nature Biotechnology, 2015, 33, 842-844.  | 17.5 | 213       |
| 3  | Site-specific characterization of threonine, serine, and tyrosine glycosylations of amyloid precursor protein/amyloid Î2-peptides in human cerebrospinal fluid. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 11848-11853. | 7.1  | 205       |
| 4  | Mining the O-mannose glycoproteome reveals cadherins as major O-mannosylated glycoproteins.<br>Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 21018-21023.  | 7.1  | 143       |
| 5  | Molecular Logic of Neuronal Self-Recognition through Protocadherin Domain Interactions. Cell, 2015, 163, 629-642.  | 28.9 | 141       |
| 6  | Human Urinary Glycoproteomics; Attachment Site Specific Analysis of N- and O-Linked Glycosylations by CID and ECD. Molecular and Cellular Proteomics, 2012, 11, M111.013649.   | 3.8  | 137       |
| 7  | Assignment of Saccharide Identities through Analysis of Oxonium Ion Fragmentation Profiles in LC–MS/MS of Glycopeptides. Journal of Proteome Research, 2014, 13, 6024-6032.  | 3.7  | 129       |
| 8  | Characterizing the O-glycosylation landscape of human plasma, platelets, and endothelial cells.<br>Blood Advances, 2017, 1, 429-442.   | 5.2  | 121       |
| 9  | Advances in mass spectrometry driven O-glycoproteomics. Biochimica Et Biophysica Acta - General<br>Subjects, 2015, 1850, 33-42.  | 2.4  | 104       |
| 10 | Interlaboratory Study on Differential Analysis of Protein Glycosylation by Mass Spectrometry: The<br>ABRF Glycoprotein Research Multi-Institutional Study 2012. Molecular and Cellular Proteomics, 2013,<br>12, 2935-2951.   | 3.8  | 103       |
| 11 | A systematic study of modulation of ADAM-mediated ectodomain shedding by site-specific<br>O-glycosylation. Proceedings of the National Academy of Sciences of the United States of America,<br>2015, 112, 14623-14628.   | 7.1  | 98        |
| 12 | LC–MS/MS Characterization of O-Glycosylation Sites and Glycan Structures of Human Cerebrospinal Fluid Glycoproteins. Journal of Proteome Research, 2013, 12, 573-584.  | 3.7  | 97        |
| 13 | Discovery of an O-mannosylation pathway selectively serving cadherins and protocadherins.<br>Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 11163-11168.  | 7.1  | 83        |
| 14 | An online nanoâ€LCâ€ESIâ€FTICRâ€MS method for comprehensive characterization of endogenous fragments<br>from amyloid β and amyloid precursor protein in human and cat cerebrospinal fluid. Journal of Mass<br>Spectrometry, 2012, 47, 591-603.                           | 1.6  | 78        |
| 15 | The GalNAc-type O-Glycoproteome of CHO Cells Characterized by the SimpleCell Strategy. Molecular and Cellular Proteomics, 2014, 13, 3224-3235.   | 3.8  | 72        |
| 16 | SnapShot: O-Glycosylation Pathways across Kingdoms. Cell, 2018, 172, 632-632.e2.   | 28.9 | 72        |
| 17 | Complex Regulation of Prolyl-4-Hydroxylases Impacts Root Hair Expansion. Molecular Plant, 2015, 8, 734-746.  | 8.3  | 70        |
| 18 | Discovery of a nucleocytoplasmic O-mannose glycoproteome in yeast. Proceedings of the National<br>Academy of Sciences of the United States of America, 2015, 112, 15648-15653.   | 7.1  | 67        |

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|----|--|------|-----------|
| 19 | Display of the human mucinome with defined O-glycans by gene engineered cells. Nature<br>Communications, 2021, 12, 4070.   | 12.8 | 67        |
| 20 | Mapping the O-Mannose Glycoproteome in Saccharomyces cerevisiae. Molecular and Cellular Proteomics, 2016, 15, 1323-1337.   | 3.8  | 61        |
| 21 | Glycoproteomics. Nature Reviews Methods Primers, 2022, 2, .  | 21.2 | 61        |
| 22 | Different affinity of galectins for human serum glycoproteins: Galectin-3 binds many protease inhibitors and acute phase proteins. Glycobiology, 2008, 18, 384-394.  | 2.5  | 59        |
| 23 | NleB/SseK effectors from Citrobacter rodentium, Escherichia coli, and Salmonella enterica display distinct differences in host substrate specificity. Journal of Biological Chemistry, 2017, 292, 11423-11430.             | 3.4  | 56        |
| 24 | Low Density Lipoprotein Receptor Class A Repeats Are O-Glycosylated in Linker Regions. Journal of<br>Biological Chemistry, 2014, 289, 17312-17324.   | 3.4  | 46        |
| 25 | Differentiation of glycosphingolipid-derived glycan structural isomers by liquid chromatography/mass spectrometry. Clycobiology, 2010, 20, 1103-1116.  | 2.5  | 45        |
| 26 | Exploring Regulation of Protein O-Glycosylation in Isogenic Human HEK293 Cells by Differential O-Glycoproteomics. Molecular and Cellular Proteomics, 2019, 18, 1396-1409.  | 3.8  | 44        |
| 27 | Characterization of Binding Epitopes of CA125 Monoclonal Antibodies. Journal of Proteome Research, 2014, 13, 3349-3359.  | 3.7  | 42        |
| 28 | Mammalian O-mannosylation of cadherins and plexins is independent of protein<br>O-mannosyltransferases 1 and 2. Journal of Biological Chemistry, 2017, 292, 11586-11598.   | 3.4  | 39        |
| 29 | Multiple distinct O-Mannosylation pathways in eukaryotes. Current Opinion in Structural Biology, 2019, 56, 171-178.  | 5.7  | 37        |
| 30 | Targeting the glycoproteome. Glycoconjugate Journal, 2013, 30, 119-136.  | 2.7  | 35        |
| 31 | O-Mannose and O–N-acetyl galactosamine glycosylation of mammalian α-dystroglycan is conserved in a<br>region-specific manner. Glycobiology, 2012, 22, 1413-1423.   | 2.5  | 32        |
| 32 | Glycoproteomic Analysis of Seven Major Allergenic Proteins Reveals Novel Post-translational<br>Modifications. Molecular and Cellular Proteomics, 2015, 14, 191-204.  | 3.8  | 32        |
| 33 | GlycoDomainViewer: a bioinformatics tool for contextual exploration of glycoproteomes.<br>Glycobiology, 2018, 28, 131-136.   | 2.5  | 25        |
| 34 | Site-specific O -glycosylation of N-terminal serine residues by polypeptide GalNAc-transferase 2<br>modulates human δ-opioid receptor turnover at the plasma membrane. Cellular Signalling, 2018, 42,<br>184-193.          | 3.6  | 24        |
| 35 | O-Linked Glycosylation of the Mucin Domain of the Herpes Simplex Virus Type 1-specific Glycoprotein<br>gC-1 Is Temporally Regulated in a Seed-and-spread Manner. Journal of Biological Chemistry, 2015, 290,<br>5078-5091. | 3.4  | 21        |
| 36 | Endoplasmic reticulum transmembrane protein TMTC3 contributes to O-mannosylation of E-cadherin, cellular adherence, and embryonic gastrulation. Molecular Biology of the Cell, 2020, 31, 167-183.                          | 2.1  | 21        |

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|----|--|-----|-----------|
| 37 | O-linked mucin-type glycosylation regulates the transcriptional programme downstream of EGFR.<br>Glycobiology, 2021, 31, 200-210.  | 2.5 | 18        |
| 38 | Genetically engineered cell factories produce glycoengineered vaccines that target<br>antigen-presenting cells and reduce antigen-specific T-cell reactivity. Journal of Allergy and Clinical<br>Immunology, 2018, 142, 1983-1987. | 2.9 | 17        |
| 39 | A strategy for generating cancer-specific monoclonal antibodies to aberrantO-glycoproteins:<br>identification of a novel dysadherin-Tn antibody. Glycobiology, 2019, 29, 307-319.  | 2.5 | 17        |
| 40 | Molecular pathogenesis of a new glycogenosis caused by a glycogenin-1 mutation. Biochimica Et<br>Biophysica Acta - Molecular Basis of Disease, 2012, 1822, 493-499.  | 3.8 | 13        |
| 41 | Novel Leb-like Helicobacter pylori-binding glycosphingolipid created by the expression of human<br>Â-1,3/4-fucosyltransferase in FVB/N mouse stomach. Glycobiology, 2008, 19, 182-191.   | 2.5 | 11        |
| 42 | The N-terminal domain of α-dystroglycan, released as a 38kDa protein, is increased in cerebrospinal<br>fluid in patients with Lyme neuroborreliosis. Biochemical and Biophysical Research Communications,<br>2011, 412, 494-499.   | 2.1 | 11        |
| 43 | LC–MS/MS characterization of combined glycogenin-1 and glycogenin-2 enzymatic activities reveals their self-glucosylation preferences. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2014, 1844, 398-405.               | 2.3 | 11        |
| 44 | A Bacterial Mannose Binding Lectin as a Tool for the Enrichment of C- and O-Mannosylated Peptides.<br>Analytical Chemistry, 2022, 94, 7329-7338.   | 6.5 | 8         |
| 45 | Involvement of viral glycoprotein <scp>gC</scp> â€1 in expression of the selectin ligand sialylâ€Lewis X induced after infection with herpes simplex virus type 1. Apmis, 2013, 121, 280-289.                                      | 2.0 | 7         |
| 46 | Microbial glycoproteomics. Current Opinion in Structural Biology, 2017, 44, 143-150.   | 5.7 | 6         |
| 47 | Drosophila O-GlcNAcase Mutants Reveal an Expanded Glycoproteome and Novel Growth and Longevity<br>Phenotypes. Cells, 2021, 10, 1026.   | 4.1 | 6         |