

# Frank Koopmans

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4776658/publications.pdf>

Version: 2024-02-01

24  
papers

2,356  
citations

567281

15  
h-index

642732

23  
g-index

25  
all docs

25  
docs citations

25  
times ranked

3081  
citing authors

| #  | ARTICLE   | IF   | CITATIONS |
|----|---|------|-----------|
| 1  | Mapping genomic loci implicates genes and synaptic biology in schizophrenia. <i>Nature</i> , 2022, 604, 502-508.  | 27.8 | 929       |
| 2  | SynGO: An Evidence-Based, Expert-Curated Knowledge Base for the Synapse. <i>Neuron</i> , 2019, 103, 217-234.e4.   | 8.1  | 518       |
| 3  | Single-cell isoform RNA sequencing characterizes isoforms in thousands of cerebellar cells. <i>Nature Biotechnology</i> , 2018, 36, 1197-1202.  | 17.5 | 253       |
| 4  | Stitching the synapse: Cross-linking mass spectrometry into resolving synaptic protein interactions. <i>Science Advances</i> , 2020, 6, eaax5783.   | 10.3 | 74        |
| 5  | Comparative Analyses of Data Independent Acquisition Mass Spectrometric Approaches: DIA, WiSIM-DIA, and Untargeted DIA. <i>Proteomics</i> , 2018, 18, 1700304.  | 2.2  | 71        |
| 6  | Interaction Proteomics Reveals Brain Region-Specific AMPA Receptor Complexes. <i>Journal of Proteome Research</i> , 2014, 13, 5695-5706.  | 3.7  | 63        |
| 7  | MIR137 schizophrenia-associated locus controls synaptic function by regulating synaptogenesis, synapse maturation and synaptic transmission. <i>Human Molecular Genetics</i> , 2018, 27, 1879-1891.                   | 2.9  | 58        |
| 8  | Correlation profiling of brain sub-cellular proteomes reveals co-assembly of synaptic proteins and subcellular distribution. <i>Scientific Reports</i> , 2017, 7, 12107.  | 3.3  | 55        |
| 9  | Single-nuclei isoform RNA sequencing unlocks barcoded exon connectivity in frozen brain tissue. <i>Nature Biotechnology</i> , 2022, 40, 1082-1092.  | 17.5 | 52        |
| 10 | Recent Developments in Data Independent Acquisition (DIA) Mass Spectrometry: Application of Quantitative Analysis of the Brain Proteome. <i>Frontiers in Molecular Neuroscience</i> , 2020, 13, 564446.               | 2.9  | 47        |
| 11 | Comparative Hippocampal Synaptic Proteomes of Rodents and Primates: Differences in Neuroplasticity-Related Proteins. <i>Frontiers in Molecular Neuroscience</i> , 2018, 11, 364.                                      | 2.9  | 43        |
| 12 | Interaction proteomics of canonical Caspr2 (CNTNAP2) reveals the presence of two Caspr2 isoforms with overlapping interactomes. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2015, 1854, 827-833. | 2.3  | 32        |
| 13 | The proteome of granulovacuolar degeneration and neurofibrillary tangles in Alzheimer's disease. <i>Acta Neuropathologica</i> , 2021, 141, 341-358.   | 7.7  | 29        |
| 14 | Identifying true protein complex constituents in interaction proteomics: The example of the DMXL2 protein complex. <i>Proteomics</i> , 2012, 12, 2428-2432.   | 2.2  | 27        |
| 15 | Splice-dependent trans-synaptic PTP-IL-1-RAPL1 interaction regulates synapse formation and non-REM sleep. <i>EMBO Journal</i> , 2020, 39, e104150.  | 7.8  | 22        |
| 16 | Empirical Bayesian Random Censoring Threshold Model Improves Detection of Differentially Abundant Proteins. <i>Journal of Proteome Research</i> , 2014, 13, 3871-3880.  | 3.7  | 20        |
| 17 | SALM1 controls synapse development by promoting F-actin/PIP2-dependent Neurexin clustering. <i>EMBO Journal</i> , 2019, 38, e101289.  | 7.8  | 17        |
| 18 | A Fast and Economical Sample Preparation Protocol for Interaction Proteomics Analysis. <i>Proteomics</i> , 2019, 19, 1900027.   | 2.2  | 11        |

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|----|--|-----|-----------|
| 19 | Age-Dependent Hippocampal Proteomics in the APP/PS1 Alzheimer Mouse Model: A Comparative Analysis with Classical SWATH/DIA and directDIA Approaches. <i>Cells</i> , 2021, 10, 1588.                      | 4.1 | 11        |
| 20 | Systematic assessment of variability in the proteome of iPSC derivatives. <i>Stem Cell Research</i> , 2021, 56, 102512.  | 0.7 | 8         |
| 21 | Glycine Receptor Complex Analysis Using Immunoprecipitationâ€Blue Native Gel Electrophoresisâ€Mass Spectrometry. <i>Proteomics</i> , 2020, 20, e1900403.   | 2.2 | 7         |
| 22 | Data-Independent Acquisition (SWATH) Mass Spectrometry Analysis of Protein Content in Primary Neuronal Cultures. <i>Neuromethods</i> , 2019, , 119-127.  | 0.3 | 4         |
| 23 | Functional brain defects in a mouse model of a chromosomal t(1;11) translocation that disrupts DISC1 and confers increased risk of psychiatric illness. <i>Translational Psychiatry</i> , 2021, 11, 135. | 4.8 | 3         |
| 24 | Neuroproteomics of cognitively healthy centenarians in the context of aging and Alzheimer's disease.. <i>Alzheimer's and Dementia</i> , 2021, 17 Suppl 3, e053681.                                       | 0.8 | 0         |