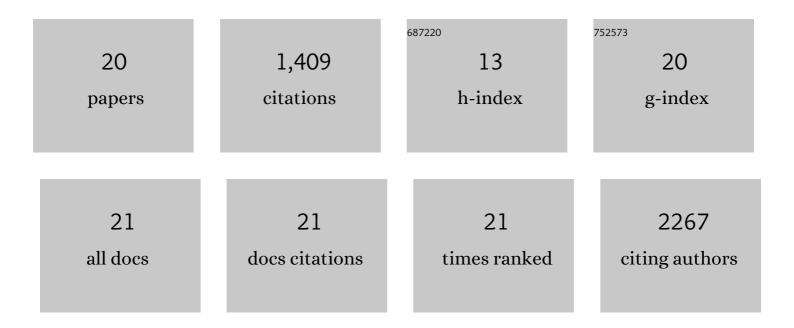
Peer-Hendrik Kuhn

List of Publications by Year in descending order

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DEED-HENDRIK KIIHN

#	Article	IF	CITATIONS
1	ADAM10 is the physiologically relevant, constitutive α-secretase of the amyloid precursor protein in primary neurons. EMBO Journal, 2010, 29, 3020-3032.	3.5	515
2	Secretome protein enrichment identifies physiological BACE1 protease substrates in neurons. EMBO Journal, 2012, 31, 3157-3168.	3.5	279
3	Systematic substrate identification indicates a central role for the metalloprotease ADAM10 in axon targeting and synapse function. ELife, 2016, 5, .	2.8	124
4	Seizure protein 6 and its homolog seizure 6-like protein are physiological substrates of BACE1 in neurons. Molecular Neurodegeneration, 2016, 11, 67.	4.4	90
5	Secretome Analysis Identifies Novel Signal Peptide Peptidase-Like 3 (SPPL3) Substrates and Reveals a Role of SPPL3 in Multiple Golgi Glycosylation Pathways*. Molecular and Cellular Proteomics, 2015, 14, 1584-1598.	2.5	74
6	Constitutive α- and β-secretase cleavages of the amyloid precursor protein are partially coupled in neurons, but not in frequently used cell lines. Neurobiology of Disease, 2013, 49, 137-147.	2.1	58
7	QUINT: Workflow for Quantification and Spatial Analysis of Features in Histological Images From Rodent Brain. Frontiers in Neuroinformatics, 2019, 13, 75.	1.3	51
8	Tumor Budding and Cell Nest Size Are Highly Prognostic in Laryngeal and Hypopharyngeal Squamous Cell Carcinoma. American Journal of Surgical Pathology, 2019, 43, 303-313.	2.1	41
9	Integration of innate into adaptive immune responses in ZAP-70–positive chronic lymphocytic leukemia. Blood, 2016, 127, 436-448.	0.6	25
10	Novel prognostic histopathological grading system in oral squamous cell carcinoma based on tumour budding and cell nest size shows high interobserver and intraobserver concordance. Journal of Clinical Pathology, 2019, 72, 285-294.	1.0	22
11	QARIP: a web server for quantitative proteomic analysis of regulated intramembrane proteolysis. Nucleic Acids Research, 2013, 41, W459-W464.	6.5	20
12	The immunologic tumor microenvironment in endometrioid endometrial cancer in the morphomolecular context: mutual correlations and prognostic impact depending on molecular alterations. Cancer Immunology, Immunotherapy, 2021, 70, 1679-1689.	2.0	18
13	Mouse brain proteomics establishes MDGA1 and CACHD1 as in vivo substrates of the Alzheimer protease BACE1. FASEB Journal, 2020, 34, 2465-2482.	0.2	16
14	Nonâ€ ϵ ellâ ϵ autonomous function of DR6 in Schwann cell proliferation. EMBO Journal, 2018, 37, .	3.5	14
15	Immunohistochemical Evidence from APP-Transgenic Mice for Glutaminyl Cyclase as Drug Target to Diminish pE-Abeta Formation. Molecules, 2018, 23, 924.	1.7	14
16	An optimised version of the secretome protein enrichment with click sugars (SPECS) method leads to enhanced coverage of the secretome. Proteomics, 2017, 17, 1600423.	1.3	12
17	Defined astrocytic expression of human amyloid precursor protein in Tg2576 mouse brain. Glia, 2019, 67, 393-403.	2.5	12
18	Evaluation of Disposable Trap Column nanoLC–FAIMS–MS/MS for the Proteomic Analysis of FFPE Tissue. Journal of Proteome Research, 2021, 20, 5402-5411.	1.8	12

#	Article	IF	CITATIONS
19	Iron-mediated aggregation and toxicity in a novel neuronal cell culture model with inducible alpha-synuclein expression. Scientific Reports, 2019, 9, 9100.	1.6	8
20	Cell Type-Specific Human APP Transgene Expression by Hippocampal Interneurons in the Tg2576 Mouse Model of Alzheimer's Disease. Frontiers in Neuroscience, 2019, 13, 137.	1.4	3