Christophe Hirtz

List of Publications by Year in descending order

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#	Article	lF	CITATIONS
1	Plasma and CSF biomarkers for the diagnosis of Alzheimer's disease in adults with Down syndrome: a cross-sectional study. Lancet Neurology, The, 2018, 17, 860-869.	10.2	140
2	Cerebrospinal fluid phospho-tau T217 outperforms T181 as a biomarker for the differential diagnosis of Alzheimer's disease and PET amyloid-positive patient identification. Alzheimer's Research and Therapy, 2020, 12, 26.	6.2	138
3	Differential Mass Spectrometry Profiles of Tau Protein in the Cerebrospinal Fluid of Patients with Alzheimer's Disease, Progressive Supranuclear Palsy, and Dementia with Lewy Bodies. Journal of Alzheimer's Disease, 2016, 51, 1033-1043.	2.6	104
4	Current and future use of "dried blood spot―analyses in clinical chemistry. Clinical Chemistry and Laboratory Medicine, 2013, 51, 1897-1909.	2.3	102
5	Tau Protein Quantification in Human Cerebrospinal Fluid by Targeted Mass Spectrometry at High Sequence Coverage Provides Insights into Its Primary Structure Heterogeneity. Journal of Proteome Research, 2016, 15, 667-676.	3.7	91
6	Cerebrospinal fluid levels of orexin-A and histamine, and sleep profile within the Alzheimer process. Neurobiology of Aging, 2017, 53, 59-66.	3.1	76
7	MS characterization of multiple forms of alpha-amylase in human saliva. Proteomics, 2005, 5, 4597-4607.	2.2	70
8	Impact of harmonization of collection tubes on Alzheimer's disease diagnosis. , 2014, 10, S390-S394.e2.		58
9	Proteins and proteolysis in pre-term and term human milk and possible implications for infant formulae. International Dairy Journal, 2010, 20, 715-723.	3.0	56
10	Use of Reducing/Nonreducing Two-Dimensional Electrophoresis for the Study of Disulfide-Mediated Interactions between Proteins in Raw and Heated Bovine Milk. Journal of Agricultural and Food Chemistry, 2009, 57, 5948-5955.	5.2	54
11	Quantitative Clinical Chemistry Proteomics (qCCP) using mass spectrometry: general characteristics and application. Clinical Chemistry and Laboratory Medicine, 2013, 51, 919-35.	2.3	47
12	Alzheimer's Disease: Advances in Drug Development. Journal of Alzheimer's Disease, 2018, 65, 3-13.	2.6	45
13	SILK studies — capturing the turnover of proteins linked to neurodegenerative diseases. Nature Reviews Neurology, 2019, 15, 419-427.	10.1	37
14	From radioimmunoassay to mass spectrometry: a new method to quantify orexin-A (hypocretin-1) in cerebrospinal fluid. Scientific Reports, 2016, 6, 25162.	3.3	36
15	Peripheral Blood and Salivary Biomarkers of Blood–Brain Barrier Permeability and Neuronal Damage: Clinical and Applied Concepts. Frontiers in Neurology, 2020, 11, 577312.	2.4	36
16	Antibody-free quantification of seven tau peptides in human CSF using targeted mass spectrometry. Frontiers in Neuroscience, 2015, 9, 302.	2.8	34
17	Cerebrospinal fluid A beta 1–40 peptides increase in Alzheimer's disease and are highly correlated with phospho-tau in control individuals. Alzheimer's Research and Therapy, 2020, 12, 123.	6.2	33
18	Quantitative detection of amyloid-β peptides by mass spectrometry: state of the art and clinical applications. Clinical Chemistry and Laboratory Medicine, 2015, 53, 1483-93.	2.3	30

CHRISTOPHE HIRTZ

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19	Use of plasma biomarkers for AT(N) classification of neurodegenerative dementias. Journal of Neurology, Neurosurgery and Psychiatry, 2021, 92, 1206-1214.	1.9	30
20	Clinical mass spectrometry proteomics (cMSP) for medical laboratory: What does the future hold?. Clinica Chimica Acta, 2017, 467, 51-58.	1.1	29
21	Association between serum hepcidin level and restless legs syndrome. Movement Disorders, 2018, 33, 618-627.	3.9	25
22	Detection of amyloid beta peptides in body fluids for the diagnosis of alzheimer's disease: Where do we stand?. Critical Reviews in Clinical Laboratory Sciences, 2020, 57, 99-113.	6.1	24
23	Proteomic Studies of Saliva: A Proposal for a Standardized Handling of Clinical Samples. Clinical Proteomics, 2007, 3, 13-21.	2.1	23
24	Cerebrospinal Fluid Alzheimer's Disease Biomarkers in Cerebral Amyloid Angiopathy-Related Inflammation. Journal of Alzheimer's Disease, 2016, 50, 759-764.	2.6	23
25	Absolute quantification of 35 plasma biomarkers in human saliva using targeted MS. Bioanalysis, 2016, 8, 43-53.	1.5	22
26	Hepcidin and ferritin levels in restless legs syndrome: a case–control study. Scientific Reports, 2020, 10, 11914.	3.3	21
27	Sample Pooling and Inflammation Linked to the False Selection of Biomarkers for Neurodegenerative Diseases in Top–Down Proteomics: A Pilot Study. Frontiers in Molecular Neuroscience, 2018, 11, 477.	2.9	20
28	The prognostic value of theÂTau protein serum level in metastatic breast cancer patients and its correlation with brain metastases. BMC Cancer, 2019, 19, 110.	2.6	20
29	Clinical measurement of Hepcidin-25 in human serum: Is quantitative mass spectrometry up to the job?. EuPA Open Proteomics, 2014, 3, 60-67.	2.5	19
30	Development of new quantitative mass spectrometry and semi-automatic isofocusing methods for the determination of Apolipoprotein E typing. Clinica Chimica Acta, 2016, 454, 33-38.	1.1	19
31	Clinical perspectives of dried blood spot protein quantification using mass spectrometry methods. Critical Reviews in Clinical Laboratory Sciences, 2017, 54, 173-184.	6.1	19
32	Salivary protein profiling in type I diabetes using two-dimensional electrophoresis and mass spectrometry. Clinical Proteomics, 2006, 2, 117-127.	2.1	18
33	Towards a routine application of Top-Down approaches for label-free discovery workflows. Journal of Proteomics, 2018, 175, 12-26.	2.4	17
34	Natural history, dynamics, and ecology of human papillomaviruses in genital infections of young women: protocol of the PAPCLEAR cohort study. BMJ Open, 2019, 9, e025129.	1.9	17
35	Cerebrospinal Fluid Alzheimer's Disease Biomarkers in Isolated Supratentorial Cortical Superficial Siderosis. Journal of Alzheimer's Disease, 2016, 54, 1291-1295.	2.6	16
36	Impurity determination for hepcidin by liquid chromatography-high resolution and ion mobility mass spectrometry for the value assignment of candidate primary calibrators. Analytical and Bioanalytical Chemistry, 2017, 409, 2559-2567.	3.7	16

CHRISTOPHE HIRTZ

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37	Comparison of HbA1c detection in whole blood and dried blood spots using an automated ion-exchange HPLC system. Bioanalysis, 2017, 9, 427-434.	1.5	14
38	Quantifying RNA modifications by mass spectrometry: a novel source of biomarkers in oncology. Critical Reviews in Clinical Laboratory Sciences, 2022, 59, 1-18.	6.1	14
39	Quantification of hepcidin-25 in human cerebrospinal fluid using LC–MS/MS. Bioanalysis, 2017, 9, 337-347.	1.5	12
40	Assessing a multiplex-targeted proteomics approach for the clinical diagnosis of periodontitis using saliva samples. Bioanalysis, 2018, 10, 35-45.	1.5	12
41	Identification of multiple proteoforms biomarkers on clinical samples by routine Top-Down approaches. Data in Brief, 2018, 18, 1013-1021.	1.0	12
42	What sample preparation should be chosen for targeted MS monoclonal antibody quantification in human serum?. Bioanalysis, 2018, 10, 723-735.	1.5	12
43	Stable Isotope Labeling by Amino acid <i>in Vivo</i> (SILAV): a new method to explore protein metabolism. Rapid Communications in Mass Spectrometry, 2015, 29, 1917-1925.	1.5	10
44	Impact of iron deficiency diagnosis using hepcidin mass spectrometry dosage methods on hospital stay and costs after a prolonged ICU stay: Study protocol for a multicentre, randomised, single-blinded medico-economic trial. Anaesthesia, Critical Care & Pain Medicine, 2017, 36, 391-396.	1.4	9
45	Regulatory context and validation of assays for clinical mass spectrometry proteomics (cMSP) methods. Critical Reviews in Clinical Laboratory Sciences, 2018, 55, 346-358.	6.1	9
46	Cerebrospinal Fluid and Plasma Biomarkers do not Differ in the Presenile and Late-Onset Behavioral Variants of Frontotemporal Dementia. Journal of Alzheimer's Disease, 2020, 74, 903-911.	2.6	9
47	Gravitational Transitions Increase Posterior Cerebral Perfusion and Systemic Oxidative-nitrosative Stress: Implications for Neurovascular Unit Integrity. Neuroscience, 2020, 441, 142-160.	2.3	9
48	Development and validation of dried matrix spot sampling for the quantitative determination of amyloid β peptides in cerebrospinal fluid. Clinical Chemistry and Laboratory Medicine, 2014, 52, 649-55.	2.3	8
49	Biochemical markers of time since death in cerebrospinal fluid: A first step towards"Forensomics― Critical Reviews in Clinical Laboratory Sciences, 2019, 56, 274-286.	6.1	8
50	Serum glial fibrillary acidic protein is a predictor of brain metastases in patients with metastatic breast cancer. International Journal of Cancer, 2021, 149, 1605-1618.	5.1	8
51	From "Clinical Proteomics―to "Clinical Chemistry Proteomics― considerations using quantitative mass-spectrometry as a model approach. Clinical Chemistry and Laboratory Medicine, 2012, 50, 235-42.	2.3	7
52	Variation of human salivary alpha-amylase proteoforms in three stimulation models. Clinical Oral Investigations, 2020, 24, 475-486.	3.0	7
53	Tau protein in cerebrospinal fluid: a novel biomarker of the time of death?. International Journal of Legal Medicine, 2021, 135, 2081-2089.	2.2	7
54	Concussion history in rugby union players is associated with depressed cerebrovascular reactivity and cognition. Scandinavian Journal of Medicine and Science in Sports, 2021, 31, 2291-2299.	2.9	7

CHRISTOPHE HIRTZ

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55	Blood amyloid and tau biomarkers as predictors of cerebrospinal fluid profiles. Journal of Neural Transmission, 2022, 129, 231-237.	2.8	7
56	Nano-flow vs standard-flow: Which is the more suitable LC/MS method for quantifying hepcidin-25 in human serum in routine clinical settings?. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2018, 1086, 110-117.	2.3	6
57	In Vivo Large-Scale Mapping of Protein Turnover in Human Cerebrospinal Fluid. Analytical Chemistry, 2019, 91, 15500-15508.	6.5	6
58	Analytical comparison of ELISA and mass spectrometry for quantification of serum hepcidin in critically ill patients. Bioanalysis, 2021, 13, 1029-1035.	1.5	6
59	Impact of biological matrix on inflammatory protein biomarker quantification based on targeted mass spectrometry. Bioanalysis, 2018, 10, 1383-1399.	1.5	5
60	Changes Occurring on the Activity of Salivary Alpha-Amylase Proteoforms in Two Naturalistic Situations Using a Spectrophotometric Assay. Biology, 2021, 10, 227.	2.8	5
61	Cytokine response following perturbation of the cervicovaginal milieu during HPV genital infection. Immunologic Research, 2021, 69, 255-263.	2.9	5
62	Efficient extraction of intact HSA-A \hat{l}^2 peptide complexes from sera: Toward albuminome biomarker identification. Talanta, 2020, 216, 121002.	5.5	4
63	Soluble Intercellular Adhesion Molecule- (sICAM-) 1, Thrombospondin-1, and Vinculin for the Identification of Septic Shock Patients Suffering from an Invasive Fungal Infection. Mediators of Inflammation, 2020, 2020, 1-13.	3.0	2
64	What is the potential of dried matrix spot sampling for cerebrospinal fluid analysis?. Bioanalysis, 2015, 7, 2849-2851.	1.5	1
65	Characterizing Deep White Matter Hyperintensities in Patients with Symptomatic Isolated Cortical Superficial Siderosis. Journal of Stroke and Cerebrovascular Diseases, 2017, 26, 465-469.	1.6	1
66	Stable Isotope Labeling Kinetics in CNS Translational Medicine: Introduction to SILK Technology. Handbook of Behavioral Neuroscience, 2019, 29, 173-190.	0.7	0
67	Intact Protein Analysis by LC-MS for Characterizing Biomarkers in Cerebrospinal Fluid. Methods in Molecular Biology, 2019, 1959, 163-172.	0.9	0