Malte Bayer

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Substance P Serum Degradation in Complex Regional Pain Syndrome – Another Piece of the Puzzle?. Journal of Pain, 2022, 23, 501-507.	0.7	5
2	Are formalinâ€fixed and paraffinâ€embedded tissues fit for proteomic analysis?. Journal of Mass Spectrometry, 2020, 55, e4347.	0.7	13
3	Validation data for the use of bradykinin and substance P protease activity assays with capillary blood and blood cards. Data in Brief, 2020, 28, 104873.	0.5	3
4	Labeled substance P as a neuropeptide reporter substance for enzyme activity. Journal of Pharmaceutical and Biomedical Analysis, 2020, 178, 112953.	1.4	5
5	Fast oxidation of αâ€melanocyteâ€stimulating hormone and derived peptides under laboratory conditions causes irreproducible results—Insights from studies of prolylcarboxypeptidase in human cell types. Pigment Cell and Melanoma Research, 2020, 33, 378-382.	1.5	1
6	Derivatization of the amino acids glycine and valine causes peptide formation—relevance for the analysis of prebiotic oligomerization. Rapid Communications in Mass Spectrometry, 2020, 34, e8912.	0.7	0
7	The hypertrehalosaemic neuropeptide conformational twins of cicadas consist of only l-amino acids: are they cis–trans isomers?. Amino Acids, 2019, 51, 1023-1028.	1.2	6
8	The serum protease network—one key to understand complex regional pain syndrome pathophysiology. Pain, 2019, 160, 1402-1409.	2.0	14
9	A vote for robustness: Monitoring serum enzyme activity by thin-layer chromatography of dabsylated bradykinin products. Journal of Pharmaceutical and Biomedical Analysis, 2017, 143, 199-203.	1.4	9
10	Pyryliumâ€based dye and charge tagging in proteomics. Electrophoresis, 2016, 37, 2953-2958.	1.3	10
11	Abundant cysteine side reactions in traditional buffers interfere with the analysis of posttranslational modifications and protein quantification – How to compromise. Rapid Communications in Mass Spectrometry, 2016, 30, 1823-1828.	0.7	8