

Louise Bundgaard

List of Publications by Year in descending order

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Version: 2024-02-01

14
papers

249
citations

933447

10
h-index

1058476

14
g-index

15
all docs

15
docs citations

15
times ranked

461
citing authors

#	ARTICLE	IF	CITATIONS
1	The Pig PeptideAtlas: A resource for systems biology in animal production and biomedicine. <i>Proteomics</i> , 2016, 16, 634-644.	2.2	47
2	Tendon response to matrix unloading is determined by the patho-physiological niche. <i>Matrix Biology</i> , 2020, 89, 11-26.	3.6	36
3	Regional disturbances in blood flow and metabolism in equine limb wound healing with formation of exuberant granulation tissue. <i>Wound Repair and Regeneration</i> , 2014, 22, 647-653.	3.0	25
4	Mass spectrometric analysis of the in vitro secretome from equine bone marrow-derived mesenchymal stromal cells to assess the effect of chondrogenic differentiation on response to interleukin-1 β treatment. <i>Stem Cell Research and Therapy</i> , 2020, 11, 187.	5.5	19
5	The Equine PeptideAtlas: A resource for developing proteomics-based veterinary research. <i>Proteomics</i> , 2014, 14, 763-773.	2.2	17
6	The occurrence of biofilm in an equine experimental wound model of healing by secondary intention. <i>Veterinary Microbiology</i> , 2017, 204, 90-95.	1.9	17
7	A selected reaction monitoring-based analysis of acute phase proteins in interstitial fluids from experimental equine wounds healing by secondary intention. <i>Wound Repair and Regeneration</i> , 2016, 24, 525-532.	3.0	16
8	The use of liquid chromatography tandem mass spectrometry to detect proteins in saliva from horses with and without systemic inflammation. <i>Veterinary Journal</i> , 2014, 202, 483-488.	1.7	15
9	Mapping of equine mesenchymal stromal cell surface proteomes for identification of specific markers using proteomics and gene expression analysis: an in vitro cross-sectional study. <i>Stem Cell Research and Therapy</i> , 2018, 9, 288.	5.5	14
10	Development of a Method for Absolute Quantification of Equine Acute Phase Proteins Using Concatenated Peptide Standards and Selected Reaction Monitoring. <i>Journal of Proteome Research</i> , 2014, 13, 5635-5647.	3.7	12
11	Effective protein extraction combined with data independent acquisition analysis reveals a comprehensive and quantifiable insight into the proteomes of articular cartilage and subchondral bone. <i>Osteoarthritis and Cartilage</i> , 2022, 30, 137-146.	1.3	11
12	Evaluation of Systemic and Local Inflammatory Parameters and Manifestations of Pain in an Equine Experimental Wound Model. <i>Journal of Equine Veterinary Science</i> , 2018, 68, 81-87.	0.9	9
13	Targeted mass spectrometry for Serum Amyloid A (SAA) isoform profiling in sequential blood samples from experimentally <i>Staphylococcus aureus</i> infected pigs. <i>Journal of Proteomics</i> , 2020, 227, 103904.	2.4	6
14	Mapping the N-Terminome in Tissue Biopsies by PCT-TAILS. <i>Methods in Molecular Biology</i> , 2020, 2043, 285-296.	0.9	5