## Ronny Mohren

List of Publications by Year in descending order

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PONNY MOHDEN

#	ARTICLE	IF	CITATIONS
1	Proteomics analysis of human intestinal organoids during hypoxia and reoxygenation as a model to study ischemia-reperfusion injury. Cell Death and Disease, 2021, 12, 95.	2.7	22
2	Inhibition of Extracellular Cathepsin D Reduces Hepatic Lipid Accumulation and Leads to Mild Changes in Inflammationin NASH Mice. Frontiers in Immunology, 2021, 12, 675535.	2.2	13
3	Insight into the Antioxidant Effect of Fermented and Non-Fermented Spirulina Water and Ethanol Extracts at the Proteome Level Using a Yeast Cell Model. Antioxidants, 2021, 10, 1366.	2.2	4
4	Aqueous humor proteome of primary open angle glaucoma: A combined dataset of mass spectrometry studies. Data in Brief, 2020, 32, 106327.	0.5	1
5	Comparative label-free proteomic analysis of equine osteochondrotic chondrocytes. Journal of Proteomics, 2020, 228, 103927.	1.2	5
6	Comparative proteomic analysis of human mesenchymal stromal cell behavior on calcium phosphate ceramics with different osteoinductive potential. Materials Today Bio, 2020, 7, 100066.	2.6	13
7	The aqueous humor proteome of primary open angle glaucoma: An extensive review. Experimental Eye Research, 2020, 197, 108077.	1.2	20
8	AntagomiR-103 and -107 Treatment Affects Cardiac Function and Metabolism. Molecular Therapy - Nucleic Acids, 2019, 14, 424-437.	2.3	25
9	Lipid surface modifications increase mesoporous silica nanoparticle labeling properties in mesenchymal stem cells. International Journal of Nanomedicine, 2018, Volume 13, 7711-7725.	3.3	27
10	Adipophilin protein expression in muscle – a possible protective role against insulin resistance. FEBS Journal, 2010, 277, 761-773.	2.2	23
11	An 8-Week High-Fat Diet Induces Obesity and Insulin Resistance with Small Changes in the Muscle Transcriptome of C57BL/6J Mice. Journal of Nutrigenetics and Nutrigenomics, 2009, 2, 280-291.	1.8	31
12	Short-term high fat-feeding results in morphological and metabolic adaptations in the skeletal muscle of C57BL/6J mice. Physiological Genomics, 2008, 32, 360-369.	1.0	114