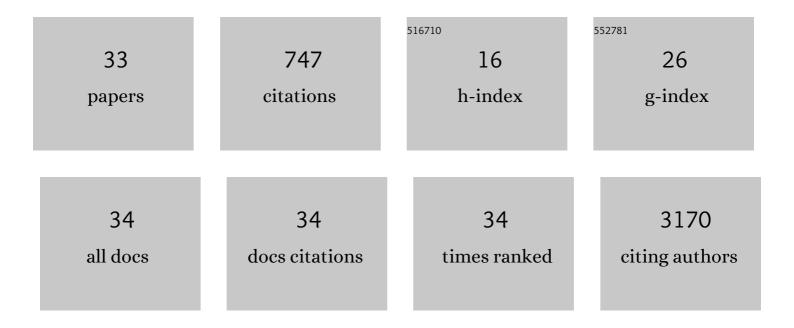
Karel Harant

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Proteomic Analysis of Trichomonas vaginalis Phagolysosome, Lysosomal Targeting, and Unconventional Secretion of Cysteine Peptidases. Molecular and Cellular Proteomics, 2022, 21, 100174.	3.8	6
2	Honey proteome of the bumblebee Bombus terrestris: similarities, differences, and exceptionality compared to honey bee honey as signatures of eusociality evolution. Apidologie, 2022, 53, 1.	2.0	2
3	Double-Stranded RNA Viruses Are Released From Trichomonas vaginalis Inside Small Extracellular Vesicles and Modulate the Exosomal Cargo. Frontiers in Microbiology, 2022, 13, .	3.5	10
4	Proteogenomic insight into the basis of the insecticide tolerance/resistance of the pollen beetle Brassicogethes (Meligethes) aeneus. Journal of Proteomics, 2021, 233, 104086.	2.4	1
5	A single honey proteome dataset for identifying adulteration by foreign amylases and mining various protein markers natural to honey. Journal of Proteomics, 2021, 239, 104157.	2.4	15
6	DIOXYGENASE FOR AUXIN OXIDATION 1 catalyzes the oxidation of IAA amino acid conjugates. Plant Physiology, 2021, 187, 103-115.	4.8	22
7	Analysis of diverse eukaryotes suggests the existence of an ancestral mitochondrial apparatus derived from the bacterial type II secretion system. Nature Communications, 2021, 12, 2947.	12.8	19
8	Label-free proteomic analysis reveals differentially expressed Wolbachia proteins in Tyrophagus putrescentiae: Mite allergens and markers reflecting population-related proteome differences. Journal of Proteomics, 2021, 249, 104356.	2.4	10
9	Proteogenomics of the house dust mite, Dermatophagoides farinae: Allergen repertoire, accurate allergen identification, isoforms, and sex-biased proteome differences. Journal of Proteomics, 2020, 210, 103535.	2.4	13
10	Anaerobic peroxisomes in <i>Mastigamoeba balamuthi</i> . Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 2065-2075.	7.1	19
11	Systematic analysis of the <scp>IL</scp> â€17 receptor signalosome reveals a robust regulatory feedback loop. EMBO Journal, 2020, 39, e104202.	7.8	16
12	High-throughput transcriptomic and proteomic profiling of mesenchymal-amoeboid transition in 3D collagen. Scientific Data, 2020, 7, 160.	5.3	15
13	Varroa destructor parasitism has a greater effect on proteome changes than the deformed wing virus and activates TGF-1 ² signaling pathways. Scientific Reports, 2019, 9, 9400.	3.3	27
14	A three-pronged "Pitchfork―strategy enables an extensive description of the human membrane proteome and the identification of missing proteins. Journal of Proteomics, 2019, 204, 103411.	2.4	3
15	The Unique Protein Composition of Honey Revealed by Comprehensive Proteomic Analysis: Allergens, Venom-like Proteins, Antibacterial Properties, Royal Jelly Proteins, Serine Proteases, and Their Inhibitors. Journal of Natural Products, 2019, 82, 1217-1226.	3.0	42
16	Chronic exposure of bumblebees to neonicotinoid imidacloprid suppresses the entire mevalonate pathway and fatty acid synthesis. Journal of Proteomics, 2019, 196, 69-80.	2.4	29
17	The effect of ω-3 polyunsaturated fatty acids on the liver lipidome, proteome and bile acid profile: parenteral versus enteral administration. Scientific Reports, 2019, 9, 19097.	3.3	11
18	Dynamic secretome of Trichomonas vaginalis: Case study of Î ² -amylases. Molecular and Cellular Proteomics, 2018, 17, 304-320.	3.8	40

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19	The Influence of Metabolic Surgery and Endoscopy on Serum Proteome in Subjects with Obesity and Type 2 Diabetes Mellitus. Diabetes, 2018, 67, 1974-P.	0.6	Ο
20	Detailed two-dimensional gel proteomic mapping of the feces of the house dust mite Dermatophagoides pteronyssinus and comparison with D. farinae: Reduced trypsin protease content in D. pteronyssinus and different isoforms. Journal of Proteomics, 2017, 162, 11-19.	2.4	28
21	Lateral gene transfer of <i>p</i> â€eresol―and indoleâ€producing enzymes from environmental bacteria to <scp><i>M</i></scp> <i>astigamoeba balamuthi</i> . Environmental Microbiology, 2017, 19, 1091-1102.	3.8	10
22	Beyond the survival and death of the deltamethrin-threatened pollen beetle Meligethes aeneus : An in-depth proteomic study employing a transcriptome database. Journal of Proteomics, 2017, 150, 281-289.	2.4	21
23	Feces Derived Allergens of Tyrophagus putrescentiae Reared on Dried Dog Food and Evidence of the Strong Nutritional Interaction between the Mite and Bacillus cereus Producing Protease Bacillolysins and Exo-chitinases. Frontiers in Physiology, 2016, 7, 53.	2.8	42
24	Detailed proteome mapping of newly emerged honeybee worker hemolymph and comparison with the red-eye pupal stage. Apidologie, 2016, 47, 805-817.	2.0	17
25	Minimal cytosolic ironâ€sulfur cluster assembly machinery of <i>Giardia intestinalis</i> is partially associated with mitosomes. Molecular Microbiology, 2016, 102, 701-714.	2.5	19
26	Large-scale identification of membrane proteins based on analysis of trypsin-protected transmembrane segments. Journal of Proteomics, 2016, 149, 15-22.	2.4	12
27	In-depth proteomic analysis of Varroa destructor: Detection of DWV-complex, ABPV, VdMLV and honeybee proteins in the mite. Scientific Reports, 2015, 5, 13907.	3.3	42
28	Giardia intestinalis Incorporates Heme into Cytosolic Cytochrome <i>b</i> ₅ . Eukaryotic Cell, 2014, 13, 231-239.	3.4	19
29	Two-dimensional gel proteome analysis of honeybee, Apis mellifera, worker red-eye pupa hemolymph. Apidologie, 2014, 45, 53-72.	2.0	15
30	NIF-type iron-sulfur cluster assembly system is duplicated and distributed in the mitochondria and cytosol of <i>Mastigamoeba balamuthi</i> . Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 7371-7376.	7.1	60
31	Cell Differentiation within a Yeast Colony: Metabolic and Regulatory Parallels with a Tumor-Affected Organism. Molecular Cell, 2012, 46, 436-448.	9.7	112
32	Secondary alcohol dehydrogenase catalyzes the reduction of exogenous acetone to 2â€propanol in <i>Trichomonas vaginalis</i> . FEBS Journal, 2012, 279, 2768-2780.	4.7	15
33	Putative role for ABC multidrug exporters in yeast quorum sensing. FEBS Letters, 2009, 583, 1107-1113.	2.8	34