## Sven Rossel

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Revealing higher than expected diversity of Harpacticoida (Crustacea:Copepoda) in the North Sea using MALDI-TOF MS and molecular barcoding. Scientific Reports, 2019, 9, 9182.	1.6	31
2	Automatic specimen identification of Harpacticoids (Crustacea:Copepoda) using Random Forest and <scp>MALDI</scp> â€ <scp>TOF</scp> mass spectra, including a post hoc test for false positive discovery. Methods in Ecology and Evolution, 2018, 9, 1421-1434.	2.2	22
3	Comparison of Rapid Biodiversity Assessment of Meiobenthos Using MALDI-TOF MS and Metabarcoding. Frontiers in Marine Science, 2019, 6, .	1.2	18
4	Effects of Sample Fixation on Specimen Identification in Biodiversity Assemblies Based on Proteomic Data (MALDI-TOF). Frontiers in Marine Science, 2018, 5, .	1.2	16
5	Toward a reliable assessment of potential ecological impacts of deepâ€sea polymetallic nodule mining on abyssal infauna. Limnology and Oceanography: Methods, 2021, 19, 626-650.	1.0	16
6	Recent speciation and hybridization in Icelandic deepâ€sea isopods: An integrative approach using genomics and proteomics. Molecular Ecology, 2022, 31, 313-330.	2.0	15
7	Rapid species level identification of fish eggs by proteome fingerprinting using MALDI-TOF MS. Journal of Proteomics, 2021, 231, 103993.	1.2	13

8 Correct Species Identification and Its Implications for Conservation Using Haploniscidae (Crustacea,) Tj ETQq0 0 0 rgBT /Overlock 10 Tf

9	Proteomic fingerprinting facilitates biodiversity assessments in understudied ecosystems: A case study on integrated taxonomy of deep sea copepods. Molecular Ecology Resources, 2021, 21, 1936-1951.	2.2	8
10	Unsupervised biodiversity estimation using proteomic fingerprints from MALDIâ€TOF MS data. Limnology and Oceanography: Methods, 2020, 18, 183-195.	1.0	4
11	Species Delimitation of Hexacorallia and Octocorallia Around Iceland Using Nuclear and Mitochondrial DNA and Proteome Fingerprinting. Frontiers in Marine Science, 2022, 9, .	1.2	4