Sven Rossel

List of Publications by Year in descending order

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1162367 1199166 12 161 8 12 citations h-index g-index papers 12 12 12 120 citing authors all docs docs citations times ranked

#	Article	IF	Citations
1	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
2	Correct Species Identification and Its Implications for Conservation Using Haploniscidae (Crustacea,) Tj ETQq0 0	0 rgBT /C	overlock 10 Tf
3	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
4	Rapid species level identification of fish eggs by proteome fingerprinting using MALDI-TOF MS. Journal of Proteomics, 2021, 231, 103993.	1,2	13
5	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
6	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
7	First insights into the phylogeography and demographic history of the common hermit crab Pagurus bernhardus (Linnaeus, 1758) (Decapoda: Anomura: Paguridae) across the Eastern Atlantic and North Sea. Journal of Crustacean Biology, 2020, 40, 435-449.	0.3	1
8	Unsupervised biodiversity estimation using proteomic fingerprints from MALDIâ€₹OF MS data. Limnology and Oceanography: Methods, 2020, 18, 183-195.	1.0	4
9	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
10	Comparison of Rapid Biodiversity Assessment of Meiobenthos Using MALDI-TOF MS and Metabarcoding. Frontiers in Marine Science, $2019, 6, .$	1.2	18
11	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
12	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