

Svenja Heesch

List of Publications by Year in descending order

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Version: 2024-02-01

32
papers

1,850
citations

471477

17
h-index

414395

32
g-index

33
all docs

33
docs citations

33
times ranked

2618
citing authors

#	ARTICLE	IF	CITATIONS
1	The <i>Ectocarpus</i> genome and the independent evolution of multicellularity in brown algae. <i>Nature</i> , 2010, 465, 617-621.	27.8	774
2	Amino acid composition, protein content, and nitrogen-to-protein conversion factors of 21 seaweed species from Norwegian waters. <i>Journal of Applied Phycology</i> , 2017, 29, 1001-1009.	2.8	128
3	Chemical characterization of 21 species of marine macroalgae common in Norwegian waters: benefits of and limitations to their potential use in food and feed. <i>Journal of the Science of Food and Agriculture</i> , 2018, 98, 2035-2042.	3.5	98
4	<i>Ulva</i> , <i>Umbraulva</i> and <i>Gemina</i> : genetic survey of New Zealand taxa reveals diversity and introduced species. <i>European Journal of Phycology</i> , 2009, 44, 143-154.	2.0	83
5	Saturating light and not increased carbon dioxide under ocean acidification drives photosynthesis and growth in <i>Ulva rigida</i> (Chlorophyta). <i>Ecology and Evolution</i> , 2015, 5, 874-888.	1.9	80
6	In-depth metabolic profiling of marine macroalgae confirms strong biochemical differences between brown, red and green algae. <i>Algal Research</i> , 2017, 26, 240-249.	4.6	67
7	Multiple gene movements into and out of haploid sex chromosomes. <i>Genome Biology</i> , 2017, 18, 104.	8.8	63
8	Sequencing type material resolves the identity and distribution of the genotype <i>Lithophyllum incrustans</i> , and related European species <i>L. Åhibernicum</i> and <i>L. Åbathyporum</i> (Corallinales, Rhodophyta). <i>Journal of Phycology</i> , 2015, 51, 791-807.	2.3	62
9	A sequence-tagged genetic map for the brown alga <i>Ectocarpus siliculosus</i> provides large-scale assembly of the genome sequence. <i>New Phytologist</i> , 2010, 188, 42-51.	7.3	59
10	Transitions between marine and freshwater environments provide new clues about the origins of multicellular plants and algae. <i>Journal of Phycology</i> , 2017, 53, 731-745.	2.3	54
11	Assessment and Characterisation of Ireland's Green Tides (<i>Ulva</i> Species). <i>PLoS ONE</i> , 2017, 12, e0169049.	2.5	51
12	Evolution of life cycles and reproductive traits: Insights from the brown algae. <i>Journal of Evolutionary Biology</i> , 2021, 34, 992-1009.	1.7	35
13	Genetic diversity and biogeography of native and introduced populations of <i>Ulva pertusa</i> (<i>Ulvales</i> , <i>Chlorophyta</i>). <i>Phycological Research</i> , 2016, 64, 102-109.	1.6	26
14	Spatial and temporal variability of biomass and composition of green tides in Ireland. <i>Harmful Algae</i> , 2019, 81, 94-105.	4.8	25
15	Prasiolales (Trebouxiophyceae, Chlorophyta) of the Svalbard Archipelago: diversity, biogeography and description of the new genera <i>Prasionella</i> and <i>Prasionema</i> . <i>European Journal of Phycology</i> , 2016, 51, 171-187.	2.0	23
16	The CCAP KnowledgeBase: linking protistan and cyanobacterial biological resources with taxonomic and molecular data. <i>Systematics and Biodiversity</i> , 2013, 11, 407-413.	1.2	20
17	The <i>Ectocarpus</i> Genome and Brown Algal Genomics. <i>Advances in Botanical Research</i> , 2012, 64, 141-184.	1.1	18
18	Scanning electron microscopy observation of host entry by two brown algae endophytic in <i>Laminaria saccharina</i> (Laminariales, Phaeophyceae). <i>Phycological Research</i> , 1999, 47, 1-5.	1.6	18

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19	Diversity, biogeography and host specificity of kelp endophytes with a focus on the genera <i>Laminarionema</i> and <i>Laminariocolax</i> (Ectocarpales, Phaeophyceae). <i>European Journal of Phycology</i> , 2019, 54, 39-51.	2.0	17
20	Intracellular eukaryotic pathogens in brown macroalgae in the Eastern Mediterranean, including LSU rRNA data for the oomycete <i>Eurychasma dicksonii</i> . <i>Diseases of Aquatic Organisms</i> , 2013, 104, 1-11.	1.0	17
21	Chemical profiling of <i>Ulva</i> species for food applications: What is in a name?. <i>Food Chemistry</i> , 2021, 361, 130084.	8.2	16
22	Affiliation of the parasite <i>Herpodiscus durvillaeae</i> (Phaeophyceae) with the Sphacelariales based on DNA sequence comparisons and morphological observations. <i>European Journal of Phycology</i> , 2008, 43, 283-295.	2.0	15
23	The arrival of a red invasive seaweed to a nutrient over-enriched estuary increases the spatial extent of macroalgal blooms. <i>Marine Environmental Research</i> , 2020, 158, 104944.	2.5	15
24	Latitudinal trends in stable isotope signatures and carbon-concentrating mechanisms of northeast Atlantic rhodoliths. <i>Biogeosciences</i> , 2018, 15, 6139-6149.	3.3	14
25	Marine Prasiolales (Trebouxiophyceae, Chlorophyta) from New Zealand and the Balleny Islands, with descriptions of <i>Prasiola novaezelandiae</i> sp. nov. and <i>Rosenvingiella australis</i> sp. nov.. <i>Phycologia</i> , 2012, 51, 217-227.	1.4	13
26	Red algal extracts from <i>Plocamium lyngbyanum</i> and <i>Ceramium secundatum</i> stimulate osteogenic activities in vitro and bone growth in zebrafish larvae. <i>Scientific Reports</i> , 2018, 8, 7725.	3.3	12
27	Screening for osteogenic activity in extracts from Irish marine organisms: The potential of <i>Ceramium pallidum</i> . <i>PLoS ONE</i> , 2018, 13, e0207303.	2.5	11
28	Development and Validation of an HPLC Method for the Quantitative Analysis of Bromophenolic Compounds in the Red Alga <i>Vertebrata lanosa</i> . <i>Marine Drugs</i> , 2019, 17, 675.	4.6	10
29	Unravelling the complexity of salt marsh <i>Fucus cottonii</i> forms (Phaeophyceae, Fucales). <i>European Journal of Phycology</i> , 2017, 52, 360-370.	2.0	9
30	Looks can be deceiving: contrasting temperature characteristics of two morphologically similar kelp species co-occurring in the Arctic. <i>Botanica Marina</i> , 2021, 64, 163-175.	1.2	9
31	Molecular phylogeny and taxonomic reassessment of the genus <i>Cladostephus</i> (Sphacelariales.) Tj ETQq1 1 0,784314 rgBT /Ov	2.0	8
32	Providing a phylogenetic framework for trait-based analyses in brown algae: Phylogenomic tree inferred from 32 nuclear protein-coding sequences. <i>Molecular Phylogenetics and Evolution</i> , 2022, 168, 107408.	2.7	2