







Program (15. September 2022)



Thursday	y, 151	th S	eptember 2022								
8:30-9:00					REG	GISTRATION & WELCOME COFFEE			Main Hall		
9:00 - 9:15						OPENING					
9:15 - 10:00	KEYNOTE: "Where Art meets Science" Caroline Ngorobi from Jukwaa Arts Productions, Kenya										
10:00-10:45						PARALELL SESSION					
	Session 1.4				Session 2.3				Session 5.4 (part 1)		
	10:00 - 10:45		AVING INDIGENOUS KNOWLEDGE YOUNG MARINE RESEARCH hosted by Megan Ranapia & Natalie Pri	- 10:30		BIOBANKING, BIOPROSPECTING, BIODIVERSITY hosted by Darya Chernikhova	10:00 - 10:45	IMPA	HIRLS AND WAVES: EXPLORING THE CTS OF SMALL-SCALE MOTIONS IN THE OCEAN and by Nicolas Dettling & Simon Reifenberg		
		1.4.1	Weaving indigenous knowledg marine research, case studies fro Canada and Aotearoa New Zeala	om	2.3.1	S The benefits of long-term databasing cetaceans in Skjålfandi Bay, Iceland	2 01 of	5.4.1	S318 Of whirls and waves: Exploring the impacts of small-scale motions in the ocean		
			by Megan Ranapia & Nato	lie Prinz		by Charla Jean Bas	ran	l.	by Stephan Juricke		
		1.4.2	Toitū te Mauri - Designing and st mauri (life-force) in experimental research		2.3.2	Open up the black box of "The Unknows" creation of an open access platform for marine fungi holding polyphasic described model-organis		5.4.2	Scattering and Refraction of Low-Mode Internal Tides by Interaction with Mesoscale Eddies		
		1.4.3	The Hidden Gems for Conservati Indigenous and Local Knowledg Fishers in Fiji	e of		by Miriam Phoebe Ster	nel	5.4.3	Ouantifying Spectral Energy Transfers in the Eastern South Atlantic using Satellite Data		
			by Salanieta	Kitolelei					by Emelie Breunig		
10:45 - 11:00						COOFFE BREAK			Main Hall		
11:00-12:30						PARALELL SESSION					
	11:00 - 12:00	ı	Session 1.7 MARINE RESOURCES: ECONOMICA INTEREST AND POLLUTANTS	AL 11:00 -12:15	F	Session 2.5 LECULAR TOOLS IN MARINE BIOLOGN FROM METHODS TO APPLICATIONS	11 :00	' IMPA	Session 5.4 (part 2) HIRLS AND WAVES: EXPLORING THE CTS OF SMALL-SCALE MOTIONS IN THE OCEAN		
	hosted by Marta Moria		hosted by Marta Moriano Ortiz	Ortiz		sted by Paulina Urban, Lara Jansen, Yassine Kasmi & Anna Joelle Greife		hoste	ed by Nicolas Dettling & Simon Reifenberg		
		S207		S207			201		S318		
		1.7.1	Traceability implications for heav metal risks in commercial seafoc		2.5.1	Range expansions of scyphozoan jellyfish – the case study of <i>Periphylla</i> periphylla and Cyanea capillata		5.4.4	The Influence of Topography on Mesoscale Ocean Mixing		
			by Marta Pilar Moria	no Ortiz		by Niko Stei	ner	L	by Miriam Sterl		
		1.7.2	Integrated methodologies for the tracking of illegally traded glass		2.5.2	Who's there? A comprehensive eDNA metabarcoding survey of gelatinous zooplankton biodiversity in the Fram Strait		5.4.6	Modulation of a Dissipation Parameterization with time: Mixing over the Reykjanes Ridge		

00	INTEREST AND	POLLUTANTS	-12:15	FRO	M METHODS TO APPLICATIONS	-12:15	IMPAC	THE OCEAN
	hosted by Mart	a Moriano Ortiz		hosted	by Paulina Urban, Lara Jansen, Yassine Kasmi & Anna Joelle Greife		hosted	by Nicolas Dettling & Simon Reifenberg
		S207			S201			S318
1.7.1		plications for heavy ommercial seafood	2.	5.1	Range expansions of scyphozoan jellyfish – the case study of <i>Periphylla</i> <i>periphylla</i> and <i>Cyanea capillata</i>	5.4	1.4	The Influence of Topography on Mesoscale Ocean Mixing
	by	Marta Pilar Moriano Ortiz		by	Niko Steiner		by	Miriam Sterl
1.7.2		hodologies for the gally traded glass eels	2.	5.2	Who's there? A comprehensive eDNA metabarcoding survey of gelatinous zooplankton biodiversity in the Fram Strait	5.4	4.6	Modulation of a Dissipation Parameterization with time: Mixing over the Reykjanes Ridge
	by	Hugo Campillo Gancedo		by	Ayla Murray		,	2.5 "
1.7.3	Microplastics ir the Asturian co	n marine macrophytes in ast	2.	5.3	Utility of environmental DNA in biomonitoring of Tanzanian cryptobenthic fishes: Does the environmental DNA approach perform better than the traditional visual census method?	5.4	by 1.7	Peter Farrell Mixing along the Weddell Sea Gravity Current
	by	Amaia Kareaga Bilbao		by	Cretus Mtonga		by	
1.7.4	innovative entre for a conceptu	l ecosystem and epreneurship: elements al framework for red ng in Madagascar	2.	5.4	Genetic studies in the coral Parazoanthus axinellae for taxonomic determination	5.4	4.8 by	The Impact of Submesoscale Dynamics on the Air Sea Exchange Moritz Epke
	by	Mihary Rabearison		by	Alfredo Rosales Ruiz			
			2.	5.5	eDNA: Reality or Myth? Qualitative and quantitative approach			
				la.	Venning Venni			

Session 1.6 NORTHERN COASTAL COMMUNITIES: TRANSFORMING GOVERNANCE FOR A SUSTAINABLE FUTURE

hosted by Maria Wilke

Education for Sustainability Futures Research

12:00

12:15

1.8.1

Session 1.8

DOES TROPHIC-LEVEL MATTER?
AQUACULTURE OF MARINE LOW-TROPHIC 12:15 12:30 ORGANISMS

hosted by Beatrice Brix da Costa & Lara Elisabeth

Technical feasibility study for the cultivation of the red algae Halymenia durvillei in the South-West of

Madagascar Rakotonandrasana Santatriniaina Nambinintsoa

MOLECULAR TOOLS IN MARINE BIOLOGY: FROM METHODS TO APPLICATIONS

2.5.4

Genetic studies in the coral Parazoanthus axinellae for taxonomic determination

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Keywords: Zoanthid, Barcoding, Genome, Mitochondrial, Taxonomy

Parazoanthus axinellae is one of the most common corals in the Mediterranean Sea reefs ecosystems, characterizing upper circalittoral and cave-scyaphilic habitats. It can grow incrusting freely over rocky walls or as parasitic species over sponges, mainly the Axinella genus. This ecological variety and its high morphological plasticity led to taxonomic uncertainties. Therefore, a genetic characterization is crucial for a correct taxonomic assignment and for the development of conservation strategies. In this study, we have used two amplified DNA regions, cytochrome oxidase I (COI) and internal transcribed spacers from ribosomal genes (ITS) to elucidate the genetic differences between five morphotypes of P. axinellae present in the Alboran Sea ("stocky", "slender", "granatensis", "middling", and "cinnabar"). These morphotypes were initially identified on macrostructural and histological characteristics. The genetic analyses carried out in this study have shown that the COI gene is not a suitable marker for variability detection between the morphotypes, due to its low genetic variation, while ITS regions/primer revealed three genetically different clusters. To confirm these results, whole mitochondrial DNA was sequenced for determination of taxonomic status of these morphotypes. Our results suggest that "Stocky" or P. axinellae var. brevitentacularis and 'granatensis' can be separated from Parazoanthus and integrated in two new independent taxa due to the differences shown by morphological and molecular analyses. The other three morphotypes ('slender', 'middling', and 'cinnabar') could be included in P. axinellae sensu lato. In conclusion, our findings confirm that a deeper taxonomic re-evaluation for the morphotypes of P. axinellae based on morphologic and genetic characters is necessary.