ClÃ¡udia Sofia Ferreira Raposo De Maga

List of Publications by Year
in descending order

12 papers	195 citations	1307594 7 h-index	1372567 10 g-index
12 all docs	12 docs citations	12 times ranked	248 citing authors

#	ARTICLE	IF	CITATIONS
1	Proteomics for Quality and Safety in Fishery Products. , 2022, , 45-78.		0
2	Effect of creatine and EDTA supplemented diets on European seabass (Dicentrarchus labrax) allergenicity, fish muscle quality and omics fingerprint. Comparative Biochemistry and Physiology Part D: Genomics and Proteomics, 2022, 41, 100941.	1.0	5
3	Fish Allergenicity Modulation Using Tailored Enriched Dietsâ€"Where Are We?. Frontiers in Physiology, 2022, 13, .	2.8	4
4	Metabolic Plasticity of Gilthead Seabream Under Different Stressors: Analysis of the Stress Responsive Hepatic Proteome and Gene Expression. Frontiers in Marine Science, 2021, 8, .	2.5	10
5	Fish Pathology Research and Diagnosis in Aquaculture of Farmed Fish; a Proteomics Perspective. Animals, 2021, 11, 125.	2.3	23
6	A Proteomics and other Omics approach in the context of farmed fish welfare and biomarker discovery. Reviews in Aquaculture, 2020, 12, 122-144.	9.0	43
7	Effect of EDTA enriched diets on farmed fish allergenicity and muscle quality; a proteomics approach. Food Chemistry, 2020, 305, 125508.	8.2	15
8	Mid-infrared spectroscopic screening of metabolic alterations in stress-exposed gilthead seabream (Sparus aurata). Scientific Reports, 2020, 10, 16343.	3.3	4
9	Protein changes as robust signatures of fish chronic stress: a proteomics approach to fish welfare research. BMC Genomics, 2020, 21, 309.	2.8	49
10	Dietary Creatine Supplementation in Gilthead Seabream (Sparus aurata): Comparative Proteomics Analysis on Fish Allergens, Muscle Quality, and Liver. Frontiers in Physiology, 2018, 9, 1844.	2.8	31
11	Improvement of the cryopreservation protocols for the dusky grouper, Epinephelus marginatus. Aquaculture, 2017, 470, 207-213.	3.5	11
12	Modulation of fish allergenicity towards the production of a low allergen farmed fish: A proteomics approach. Journal of Proteomics and Bioinformatics, 2017, 10, .	0.4	0