Yannick Labreuche

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/4916515/publications.pdf

Version: 2024-02-01

10	770	933447 10	1372567 10
papers	citations	h-index	g-index

11	11	11	718
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Cellular and molecular hemocyte responses of the Pacific oyster, Crassostrea gigas, following bacterial infection with Vibrio aestuarianus strain 01/32. Microbes and Infection, 2006, 8, 2715-2724.	1.9	160
2	Crassostrea gigas mortality in France: the usual suspect, a herpes virus, may not be the killer in this polymicrobial opportunistic disease. Frontiers in Microbiology, 2015, 6, 686.	3.5	135
3	Populations, not clones, are the unit of vibrio pathogenesis in naturally infected oysters. ISME Journal, 2015, 9, 1523-1531.	9.8	126
4	Vibrio crassostreae, a benign oyster colonizer turned into a pathogen after plasmid acquisition. ISME Journal, 2017, 11, 1043-1052.	9.8	116
5	Vibrio aestuarianus zinc metalloprotease causes lethality in the Pacific oyster Crassostrea gigas and impairs the host cellular immune defenses. Fish and Shellfish Immunology, 2010, 29, 753-758.	3.6	69
6	Species-specific mechanisms of cytotoxicity toward immune cells determine the successful outcome ofVibrioinfections. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 14238-14247.	7.1	62
7	Ancestral gene acquisition as the key to virulence potential in environmental Vibrio populations. ISME Journal, 2018, 12, 2954-2966.	9.8	37
8	Selection of Vibrio crassostreae relies on a plasmid expressing a type 6 secretion system cytotoxic for host immune cells. Environmental Microbiology, 2020, 22, 4198-4211.	3.8	26
9	Ontogeny of osmoregulation in the Pacific blue shrimp, Litopenaeus stylirostris (Decapoda,) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 5 Biochemistry and Molecular Biology, 2016, 196-197, 27-37.	1.6	17
10	Vibrio splendidus Oâ€antigen structure: a tradeâ€off between virulence to oysters and resistance to grazers. Environmental Microbiology, 2020, 22, 4264-4278.	3.8	14