## David Melka

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/12008779/publications.pdf

Version: 2024-02-01

11 papers	771 citations	933447 10 h-index	1281871 11 g-index
11	11	11	862 citing authors
all docs	docs citations	times ranked	

#	Article	IF	CITATIONS
1	Survey of <i>Salmonella</i> in raw tree nuts at retail in the United States. Journal of Food Science, 2021, 86, 495-504.	3.1	6
2	Whole genome sequencing uses for foodborne contamination and compliance: Discovery of an emerging contamination event in an ice cream facility using whole genome sequencing. Infection, Genetics and Evolution, 2019, 73, 214-220.	2.3	32
3	Survey of Foodborne Pathogens, Aerobic Plate Counts, Total Coliform Counts, and Escherichia coli Counts in Leafy Greens, Sprouts, and Melons Marketed in the United States. Journal of Food Protection, 2018, 81, 400-411.	1.7	34
4	Singleton Sequence Type 382, an Emerging Clonal Group of Listeria monocytogenes Associated with Three Multistate Outbreaks Linked to Contaminated Stone Fruit, Caramel Apples, and Leafy Green Salad. Journal of Clinical Microbiology, 2017, 55, 931-941.	3.9	56
5	Whole Genome and Core Genome Multilocus Sequence Typing and Single Nucleotide Polymorphism Analyses of Listeria monocytogenes Isolates Associated with an Outbreak Linked to Cheese, United States, 2013. Applied and Environmental Microbiology, 2017, 83, .	3.1	93
6	Prevalence of Salmonella in 11 Spices Offered for Sale from Retail Establishments and in Imported Shipments Offered for Entry to the United States. Journal of Food Protection, 2017, 80, 1791-1805.	1.7	31
7	Prevalence of Salmonella in Cashews, Hazelnuts, Macadamia Nuts, Pecans, Pine Nuts, and Walnuts in the United States. Journal of Food Protection, 2017, 80, 459-466.	1.7	40
8	Assessing the genome level diversity of Listeria monocytogenes from contaminated ice cream and environmental samples linked to a listeriosis outbreak in the United States. PLoS ONE, 2017, 12, e0171389.	2.5	72
9	Listeria monocytogenes in Stone Fruits Linked to a Multistate Outbreak: Enumeration of Cells and Whole-Genome Sequencing. Applied and Environmental Microbiology, 2016, 82, 7030-7040.	3.1	83
10	Practical Value of Food Pathogen Traceability through Building a Whole-Genome Sequencing Network and Database. Journal of Clinical Microbiology, 2016, 54, 1975-1983.	3.9	278
11	Notes from the field: listeriosis associated with stone fruit-United States, 2014. Morbidity and Mortality Weekly Report, 2015, 64, 282-3.	15.1	46