## Olivier Vc Zemb

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9048465/publications.pdf

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#	Article	IF	CITATIONS
1	Microbiability and microbiome-wide association analyses of feed efficiency and performance traits in pigs. Genetics Selection Evolution, 2022, 54, 29.	3.0	9
2	Gut Microbiota-Derived Metabolite Signature in Suckling and Weaned Piglets. Journal of Proteome Research, 2021, 20, 982-994.	3.7	31
3	Genetic relationships between feed efficiency and gut microbiome in pig lines selected for residual feed intake. Journal of Animal Breeding and Genetics, 2021, 138, 491-507.	2.0	19
4	The fecal microbiota of piglets during weaning transition and its association with piglet growth across various farm environments. PLoS ONE, 2021, 16, e0250655.	2.5	21
5	Bacterial Consumption of T4 Phages. Microorganisms, 2021, 9, 1852.	3.6	0
6	Early Inoculation of Microbial Suspension in Suckling Piglets Affects the Transmission of Maternal Microbiota and the Associated Antibiotic Resistance Genes. Microorganisms, 2020, 8, 1576.	3.6	3
7	Microbiota Changes Due to Grape Seed Extract Diet Improved Intestinal Homeostasis and Decreased Fatness in Parental Broiler Hens. Microorganisms, 2020, 8, 1141.	3.6	8
8	Absolute quantitation of microbes using 16S rRNA gene metabarcoding: A rapid normalization of relative abundances by quantitative PCR targeting a 16S rRNA gene spikeâ€in standard. MicrobiologyOpen, 2020, 9, e977.	3.0	43
9	Gut microbiota derived metabolites contribute to intestinal barrier maturation at the suckling-to-weaning transition. Gut Microbes, 2020, 11, 1268-1286.	9.8	72
10	Effect of chronic and acute heat challenges on fecal microbiota composition, production, and thermoregulation traits in growing pigs1,2. Journal of Animal Science, 2019, 97, 3845-3858.	0.5	35
11	Isolation, characterization and bioaugmentation of an acidotolerant 1,2-dichloroethane respiring Desulfitobacterium species from a low pH aquifer. FEMS Microbiology Ecology, 2019, 95, .	2.7	10
12	Effects of a gut microbiota transfer on emotional reactivity in Japanese quails ( <i>Coturnix) Tj ETQq0 0 0 rgBT /O</i>	verlock 10 1.7	) Tf 50 302 T
13	Estimating direct genetic and maternal effects affecting rabbit growth and feed efficiency with a factorial design. Journal of Animal Breeding and Genetics, 2019, 136, 168-173.	2.0	8
14	Longitudinal Analysis of the Microbiota Composition and Enterotypes of Pigs from Post-Weaning to Finishing. Microorganisms, 2019, 7, 622.	3.6	20
15	Fecal microbial composition associated with variation in feed efficiency in pigs depends on diet and sex1. Journal of Animal Science, 2018, 96, 1405-1418.	0.5	50
16	Effect of dietary fiber content on nutrient digestibility and fecal microbiota composition in growing-finishing pigs. PLoS ONE, 2018, 13, e0206159.	2.5	54

17	Phage adsorption to bacteria in the light of the electrostatics: A case study using E. coli, T2 and flow cytometry. Journal of Virological Methods, 2013, 189, 283-289.	2.1	25

18Bacterial and fungal community composition over time in chicken litter with high or low moisture<br/>content. British Poultry Science, 2012, 53, 561-569.1.739

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#	Article	IF	CITATIONS
19	Improvement of RNA-SIP by pyrosequencing to identify putative 4-n-nonylphenol degraders in activated sludge. Water Research, 2012, 46, 601-610.	11.3	26
20	Capillary electrophoresis ribosomal RNA single-stranded conformation polymorphism: a new approach for characterization of low-diversity microbial communities. Analytical and Bioanalytical Chemistry, 2012, 404, 1897-1906.	3.7	4
21	Complete chloroform dechlorination by organochlorine respiration and fermentation. Environmental Microbiology, 2012, 14, 883-894.	3.8	94
22	Direct rRNA Fingerprinting, a Novel Method to Profile Low Diversity Microbial Communities. Microbial Ecology, 2011, 62, 177-187.	2.8	4
23	Reactive iron barriers: a niche enabling microbial dehalorespiration of 1,2-dichloroethane. Applied Microbiology and Biotechnology, 2010, 88, 319-325.	3.6	19
24	Effect of a transient perturbation on marine bacterial communities with contrasting history. Journal of Applied Microbiology, 2010, 109, 751-762.	3.1	11
25	Efficient method to isolate and purify viruses of bacteria from marine environments. Letters in Applied Microbiology, 2008, 47, 41-45.	2.2	2
26	Major differences of bacterial diversity and activity inside and outside of a natural ironâ€fertilized phytoplankton bloom in the Southern Ocean. Environmental Microbiology, 2008, 10, 738-756.	3.8	155
27	Dynamics of neutral biodiversity. Mathematical Biosciences, 2008, 212, 88-98.	1.9	17
28	safum: statistical analysis of SSCP fingerprints using PCA projections, dendrograms and diversity estimators. Molecular Ecology Notes, 2007, 7, 767-770.	1.7	40
29	Denaturing gradient electrophoresis (DGE) and single-strand conformation polymorphism (SSCP) molecular fingerprintings revisited by simulation and used as a tool to measure microbial diversity. Environmental Microbiology, 2006, 8, 720-731.	3.8	114