## Ã~yvind Kommedal

List of Publications by Year in descending order

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759233 677142 22 630 12 22 h-index g-index citations papers 23 23 23 937 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Characterization of abscesses from liver, pancreas and kidney using deep sequencing of the 16S rRNA gene. Diagnostic Microbiology and Infectious Disease, 2021, 99, 115277.	1.8	5
2	The cultivable microbiota of the human distal ileum. Clinical Microbiology and Infection, 2021, 27, 912.e7-912.e13.	6.0	11
3	Managing Contamination and Diverse Bacterial Loads in 16S rRNA Deep Sequencing of Clinical Samples: Implications of the Law of Small Numbers. MBio, 2021, 12, e0059821.	4.1	8
4	Repeated bronchoscopy in health and obstructive lung disease: is the airway microbiome stable?. BMC Pulmonary Medicine, 2021, 21, 342.	2.0	4
5	Bacteria and fungi in acute cholecystitis. A prospective study comparing next generation sequencing to culture. Journal of Infection, 2020, 80, 16-23.	3.3	22
6	The first report on epidemiology of oropharyngeal <i>Kingella kingae</i> carriage in Scandinavian children: <i>K.Âkingae</i> carriage is very common in children attending day care facilities in Western Norway. Apmis, 2020, 128, 35-40.	2.0	7
7	Horizontal Plasmid Transfer among Klebsiella pneumoniae Isolates Is the Key Factor for Dissemination of Extended-Spectrum $\hat{l}^2$ -Lactamases among Children in Tanzania. MSphere, 2020, 5, .	2.9	9
8	Eikenella exigua sp. nov., isolated from brain abscess and blood. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 1478-1488.	1.7	9
9	Validation and Retrospective Clinical Evaluation of a Quantitative 16S rRNA Gene Metagenomic Sequencing Assay for Bacterial Pathogen Detection in Body Fluids. Journal of Molecular Diagnostics, 2019, 21, 913-923.	2.8	21
10	Species Level Description of the Human Ileal Bacterial Microbiota. Scientific Reports, 2018, 8, 4736.	3.3	80
11	Infective endocarditis in Western Norway: a 20-year retrospective survey. Infectious Diseases, 2018, 50, 757-763.	2.8	18
12	HPV Genotyping of Modified General Primer-Amplicons Is More Analytically Sensitive and Specific by Sequencing than by Hybridization. PLoS ONE, 2017, 12, e0169074.	2.5	9
13	Genetic antimicrobial susceptibility testing in Gramâ€negative sepsis – impact on time to results in a routine laboratory. Apmis, 2016, 124, 603-610.	2.0	8
14	High Prevalence of Faecal Carriage of ESBL-Producing Enterobacteriaceae among Children in Dar es Salaam, Tanzania. PLoS ONE, 2016, 11, e0168024.	2.5	77
15	Genetic Diversity of Circulating Rotavirus Strains in Tanzania Prior to the Introduction of Vaccination. PLoS ONE, 2014, 9, e97562.	2.5	36
16	Massive Parallel Sequencing Provides New Perspectives on Bacterial Brain Abscesses. Journal of Clinical Microbiology, 2014, 52, 1990-1997.	3.9	65
17	Genetic diversity of norovirus in hospitalised diarrhoeic children and asymptomatic controls in Dar es Salaam, Tanzania. Infection, Genetics and Evolution, 2014, 26, 340-347.	2.3	22
18	Dual Priming Oligonucleotides for Broad-Range Amplification of the Bacterial 16S rRNA Gene Directly from Human Clinical Specimens. Journal of Clinical Microbiology, 2012, 50, 1289-1294.	3.9	52

## $\tilde{\mathsf{A}}^{\sim}\mathsf{YVIND}$ Kommedal

#	Article	IF	CITATIONS
19	Characterization of polybacterial clinical samples using a set of group-specific broad-range primers targeting the 16S rRNA gene followed by DNA sequencing and RipSeq analysis. Journal of Medical Microbiology, 2011, 60, 927-936.	1.8	36
20	Direct 16S rRNA Gene Sequencing from Clinical Specimens, with Special Focus on Polybacterial Samples and Interpretation of Mixed DNA Chromatograms. Journal of Clinical Microbiology, 2009, 47, 3562-3568.	3.9	64
21	Analysis of Mixed Sequencing Chromatograms and Its Application in Direct 16S rRNA Gene Sequencing of Polymicrobial Samples. Journal of Clinical Microbiology, 2008, 46, 3766-3771.	3.9	60
22	Antibiotic susceptibility of blood culture isolates of anaerobic bacteria at a Norwegian university hospital. Apmis, 2007, 115, 956-961.	2.0	7