## Evelyn Toh

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8585253/publications.pdf

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39	2,201	22	35
papers	citations	h-index	g-index
39	39	39	2733
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	24435 Pathogen-specific metabolic pathways and innate immune responses associated with Chlamydia trachomatis infection and other STIs. Journal of Clinical and Translational Science, 2021, 5, 87-88.	0.6	О
2	Evaluation of clinical, Gram stain, and microbiological cure outcomes in men receiving azithromycin for acute nongonococcal urethritis. Sexually Transmitted Diseases, 2021, Publish Ahead of Print, 67-75.	1.7	2
3	Aetiology and prevalence of mixed-infections and mono-infections in non-gonococcal urethritis in men: a case-control study. Sexually Transmitted Infections, 2020, 96, 306-311.	1.9	16
4	No Pathogen-Specific Sign or Symptom Predicts the Etiology of Monomicrobial Nongonococcal Urethritis in Men. Sexually Transmitted Diseases, 2020, 47, 329-331.	1.7	4
5	A Genital Infection-Attenuated Chlamydia muridarum Mutant Infects the Gastrointestinal Tract and Protects against Genital Tract Challenge. MBio, 2020, $11$ , .	4.1	16
6	Development of a SimpleProbe real-Time PCR Assay for rapid detection and identification of the US novel urethrotropic clade of Neisseria meningitidis ST-11 (US_NmUC). PLoS ONE, 2020, 15, e0228467.	2.5	6
7	Genetic Screen in Chlamydia muridarum Reveals Role for an Interferon-Induced Host Cell Death Program in Antimicrobial Inclusion Rupture. MBio, 2019, 10, .	4.1	19
8	P493â€Determination ofChlamydia trachomatisorganism load in men with Non-Gonococcal Urethritis (NGU). , 2019, , .		0
9	P794â€Signs and symptoms associated with single-pathogen nongonococcal urethritis in men. , 2019, , .		O
10	P795â€Prevalence and etiology of post-azithromycin persistent non-gonococcal urethritis (NGU) symptoms in men. , 2019, , .		0
11	Detection of Rectal Chlamydia trachomatis in Heterosexual Men Who Report Cunnilingus. Sexually Transmitted Diseases, 2019, 46, 440-445.	1.7	13
12	Characterization of Proximal Small Intestinal Microbiota in Patients With Suspected Small Intestinal Bacterial Overgrowth: A Cross-Sectional Study. Clinical and Translational Gastroenterology, 2019, 10, e00073.	2.5	13
13	Heteroresistance to the model antimicrobial peptide polymyxin B in the emerging <i>Neisseria meningitidis</i> lineage 11.2 urethritis clade: mutations in the <i>pilMNOPQ</i> operon. Molecular Microbiology, 2019, 111, 254-268.	2.5	15
14	Mutations in Sugar-Nucleotide Synthesis Genes Restore Holdfast Polysaccharide Anchoring to Caulobacter crescentus Holdfast Anchor Mutants. Journal of Bacteriology, 2018, 200, .	2.2	14
15	Cutaneous Burn Injury Promotes Shifts in the Bacterial Microbiome in Autologous Donor Skin. Shock, 2017, 48, 441-448.	2.1	36
16	Emergence of a new <i>Neisseria meningitidis</i> clonal complex 11 lineage 11.2 clade as an effective urogenital pathogen. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 4237-4242.	7.1	79
17	Cutaneous Burn Injury Modulates Urinary Antimicrobial Peptide Responses and the Urinary Microbiome. Critical Care Medicine, 2017, 45, e543-e551.	0.9	15
18	Quantity of alcohol drinking positively correlates with serum levels of endotoxin and markers of monocyte activation. Scientific Reports, 2017, 7, 4462.	3.3	44

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19	<i>Neisseria meningitidis</i> ST11 Complex Isolates Associated with Nongonococcal Urethritis, Indiana, USA, 2015–2016. Emerging Infectious Diseases, 2017, 23, 336-339.	4.3	31
20	Household air pollution and the lung microbiome of healthy adults in Malawi: a cross-sectional study. BMC Microbiology, 2016, 16, 182.	3.3	49
21	Interrogating Genes That Mediate Chlamydia trachomatis Survival in Cell Culture Using Conditional Mutants and Recombination. Journal of Bacteriology, 2016, 198, 2131-2139.	2.2	27
22	Effect of Advanced HIV Infection on the Respiratory Microbiome. American Journal of Respiratory and Critical Care Medicine, 2016, 194, 226-235.	5.6	83
23	Mutational Analysis of the Chlamydia muridarum Plasticity Zone. Infection and Immunity, 2015, 83, 2870-2881.	2.2	46
24	Similarities and seasonal variations in bacterial communities from the blood of rodents and from their flea vectors. ISME Journal, 2015, 9, 1662-1676.	9.8	31
25	Mo1202 Esophageal Microbiome in Healthy Children and Eosinophilic Esophagitis: A Prospective Study. Gastroenterology, 2015, 148, S-637-S-638.	1.3	2
26	The Human Skin Microbiome Associates with the Outcome of and Is Influenced by Bacterial Infection. MBio, 2015, 6, e01315-15.	4.1	94
27	Haemophilus ducreyi Cutaneous Ulcer Strains Are Nearly Identical to Class I Genital Ulcer Strains. PLoS Neglected Tropical Diseases, 2015, 9, e0003918.	3.0	26
28	Association of Host and Microbial Species Diversity across Spatial Scales in Desert Rodent Communities. PLoS ONE, 2014, 9, e109677.	2.5	21
29	Interplay between Bladder Microbiota and Urinary Antimicrobial Peptides: Mechanisms for Human Urinary Tract Infection Risk and Symptom Severity. PLoS ONE, 2014, 9, e114185.	2.5	106
30	Contingent interactions among biofilmâ€forming bacteria determine preservation or decay in the first steps toward fossilization of marine embryos. Evolution & Development, 2013, 15, 243-256.	2.0	33
31	The arthropod, but not the vertebrate host or its environment, dictates bacterial community composition of fleas and ticks. ISME Journal, 2013, 7, 221-223.	9.8	107
32	Evidence of Uncultivated Bacteria in the Adult Female Bladder. Journal of Clinical Microbiology, 2012, 50, 1376-1383.	3.9	543
33	Functional Characterization of UDP-Glucose:Undecaprenyl-Phosphate Glucose-1-Phosphate Transferases of Escherichia coli and Caulobacter crescentus. Journal of Bacteriology, 2012, 194, 2646-2657.	2.2	70
34	Bacterial Communities of the Coronal Sulcus and Distal Urethra of Adolescent Males. PLoS ONE, 2012, 7, e36298.	2.5	191
35	The Microbial Communities in Male First Catch Urine Are Highly Similar to Those in Paired Urethral Swab Specimens. PLoS ONE, 2011, 6, e19709.	2.5	170
36	A localized multimeric anchor attaches the Caulobacter holdfast to the cell pole. Molecular Microbiology, 2010, 76, 409-427.	2.5	64

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#	Article	IF	CITATION
37	Characterization of the <i>Caulobacter crescentus</i> Holdfast Polysaccharide Biosynthesis Pathway Reveals Significant Redundancy in the Initiating Glycosyltransferase and Polymerase Steps. Journal of Bacteriology, 2008, 190, 7219-7231.	2.2	76
38	Development of Surface Adhesion in Caulobacter crescentus. Journal of Bacteriology, 2004, 186, 1438-1447.	2.2	102
39	The HfaB and HfaD adhesion proteins of Caulobacter crescentus are localized in the stalk. Molecular Microbiology, 2003, 49, 1671-1683.	2.5	37