

# Sarah J Coulthurst

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2781720/publications.pdf>

Version: 2024-02-01

51  
papers

3,979  
citations

147801

31  
h-index

182427

51  
g-index

58  
all docs

58  
docs citations

58  
times ranked

3741  
citing authors

#	ARTICLE	IF	CITATIONS
1	Aim, Load, Fire: The Type VI Secretion System, a Bacterial Nanoweapon. <i>Trends in Microbiology</i> , 2016, 24, 51-62.	7.7	366
2	Quorum Sensing Coordinates Brute Force and Stealth Modes of Infection in the Plant Pathogen <i>Pectobacterium atrosepticum</i> . <i>PLoS Pathogens</i> , 2008, 4, e1000093.	4.7	216
3	The Type VI secretion system: a versatile bacterial weapon. <i>Microbiology (United Kingdom)</i> , 2019, 165, 503-515.	1.8	216
4	The Opportunistic Pathogen <i>Serratia marcescens</i> Utilizes Type VI Secretion To Target Bacterial Competitors. <i>Journal of Bacteriology</i> , 2011, 193, 6057-6069.	2.2	203
5	The type VI secretion system deploys antifungal effectors against microbial competitors. <i>Nature Microbiology</i> , 2018, 3, 920-931.	13.3	199
6	VgrG and PAAR Proteins Define Distinct Versions of a Functional Type VI Secretion System. <i>PLoS Pathogens</i> , 2016, 12, e1005735.	4.7	184
7	The Type VI secretion system “a widespread and versatile cell targeting system. <i>Research in Microbiology</i> , 2013, 164, 640-654.	2.1	177
8	Intraspecies Competition in <i>Serratia marcescens</i> Is Mediated by Type VI-Secreted Rhs Effectors and a Conserved Effector-Associated Accessory Protein. <i>Journal of Bacteriology</i> , 2015, 197, 2350-2360.	2.2	165
9	Genome Evolution and Plasticity of <i>Serratia marcescens</i> , an Important Multidrug-Resistant Nosocomial Pathogen. <i>Genome Biology and Evolution</i> , 2014, 6, 2096-2110.	2.5	155
10	Molecular weaponry: diverse effectors delivered by the Type VI secretion system. <i>Cellular Microbiology</i> , 2015, 17, 1742-1751.	2.1	150
11	Quorum sensing, virulence and secondary metabolite production in plant soft-rotting bacteria. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2007, 362, 1165-1183.	4.0	140
12	Regulation and biosynthesis of carbapenem antibiotics in bacteria. <i>Nature Reviews Microbiology</i> , 2005, 3, 295-306.	28.6	135
13	New secreted toxins and immunity proteins encoded within the <i>type VI secretion system gene cluster of <i>Serratia marcescens</i></i> . <i>Molecular Microbiology</i> , 2012, 86, 921-936.	2.5	121
14	The regulation of virulence in phytopathogenic <i>Erwinia</i> species: quorum sensing, antibiotics and ecological considerations. <i>Antonie Van Leeuwenhoek</i> , 2002, 81, 223-231.	1.7	110
15	Stable-isotope labeling with amino acids in nematodes. <i>Nature Methods</i> , 2011, 8, 849-851.	19.0	108
16	Type VI secretion system effector proteins: Effective weapons for bacterial competitiveness. <i>Cellular Microbiology</i> , 2020, 22, e13241.	2.1	93
17	<i>luxS</i> mutants of <i>Serratia</i> defective in autoinducer-2-dependent “quorum sensing” show strain-dependent impacts on virulence and production of carbapenem and prodigiosin. <i>Microbiology (United Kingdom)</i> , 2004, 150, 1901-1910.	1.8	91
18	Proteomic Identification of Novel Secreted Antibacterial Toxins of the <i>Serratia marcescens</i> Type VI Secretion System. <i>Molecular and Cellular Proteomics</i> , 2013, 12, 2735-2749.	3.8	81

#	ARTICLE	IF	CITATIONS
19	Metabolic and regulatory engineering of <i>Serratia marcescens</i> : mimicking phage-mediated horizontal acquisition of antibiotic biosynthesis and quorum-sensing capacities. <i>Microbiology (United Kingdom)</i> , 2006, 152, 1899-1911.	1.8	79
20	A multi-repeat adhesin of the phytopathogen, <i>Pectobacterium atrosepticum</i> , is secreted by a Type I pathway and is subject to complex regulation involving a non-canonical diguanylate cyclase. <i>Molecular Microbiology</i> , 2011, 82, 719-733.	2.5	64
21	Visualization of the <i>Serratia</i> Type VI Secretion System Reveals Unprovoked Attacks and Dynamic Assembly. <i>Cell Reports</i> , 2015, 12, 2131-2142.	6.4	63
22	Genetic and proteomic analysis of the role of luxS in the enteric phytopathogen, <i>Erwinia carotovora</i> . <i>Molecular Plant Pathology</i> , 2006, 7, 31-45.	4.2	57
23	A family of Type VI secretion system effector proteins that form ion-selective pores. <i>Nature Communications</i> , 2019, 10, 5484.	12.8	57
24	The Insect Pathogen <i>Serratia marcescens</i> Db10 Uses a Hybrid Non-Ribosomal Peptide Synthetase-Polyketide Synthase to Produce the Antibiotic Althiomycin. <i>PLoS ONE</i> , 2012, 7, e44673.	2.5	54
25	Biochemical analysis of TssK, a core component of the bacterial Type VI secretion system, reveals distinct oligomeric states of TssK and identifies a TssK-TssFG subcomplex. <i>Biochemical Journal</i> , 2014, 461, 291-304.	3.7	53
26	DsbA Plays a Critical and Multifaceted Role in the Production of Secreted Virulence Factors by the Phytopathogen <i>Erwinia carotovora</i> subsp. <i>atroseptica</i> . <i>Journal of Biological Chemistry</i> , 2008, 283, 23739-23753.	3.4	48
27	N-Acetylglucosamine-dependent biofilm formation in <i>Pectobacterium atrosepticum</i> is cryptic and activated by elevated c-di-GMP levels. <i>Microbiology (United Kingdom)</i> , 2011, 157, 3340-3348.	1.8	47
28	A holin and an endopeptidase are essential for chitinolytic protein secretion in <i>Serratia marcescens</i> . <i>Journal of Cell Biology</i> , 2014, 207, 615-626.	5.2	47
29	The ecological impact of a bacterial weapon: microbial interactions and the Type VI secretion system. <i>FEMS Microbiology Reviews</i> , 2021, 45, .	8.6	45
30	Structure-activity relationships of <i>Erwinia carotovora</i> quorum sensing signaling molecules. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2005, 15, 4235-4238.	2.2	37
31	Structural basis for type VI secreted peptidoglycan-endopeptidase function, specificity and neutralization in <i>Serratia marcescens</i> . <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2013, 69, 2468-2482.	2.5	37
32	Remnant signal peptides on non-exported enzymes: implications for the evolution of prokaryotic respiratory chains. <i>Microbiology (United Kingdom)</i> , 2009, 155, 3992-4004.	1.8	36
33	Quorum sensing has an unexpected role in virulence in the model pathogen <i>Citrobacter rodentium</i> . <i>EMBO Reports</i> , 2007, 8, 698-703.	4.5	32
34	Dual Role for DsbA in Attacking and Targeted Bacterial Cells during Type VI Secretion System-Mediated Competition. <i>Cell Reports</i> , 2018, 22, 774-785.	6.4	31
35	Killing with proficiency: Integrated post-translational regulation of an offensive Type VI secretion system. <i>PLoS Pathogens</i> , 2018, 14, e1007230.	4.7	30
36	Two mobile <i>Pectobacterium atrosepticum</i> prophages modulate virulence. <i>FEMS Microbiology Letters</i> , 2010, 304, 195-202.	1.8	29

#	ARTICLE	IF	CITATIONS
37	The structure of <i>Serratia marcescens</i> Lip, a membrane-bound component of the type VI secretion system. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2011, 67, 1065-1072.	2.5	27
38	Signal peptide etiquette during assembly of a complex respiratory enzyme. <i>Molecular Microbiology</i> , 2013, 90, 400-414.	2.5	27
39	A synthetic system for expression of components of a bacterial microcompartment. <i>Microbiology (United Kingdom)</i> , 2013, 159, 2427-2436.	1.8	26
40	Can boron get bacteria talking?. <i>Trends in Biochemical Sciences</i> , 2002, 27, 217-219.	7.5	25
41	Conserved Signal Peptide Recognition Systems across the Prokaryotic Domains. <i>Biochemistry</i> , 2012, 51, 1678-1686.	2.5	25
42	The archetype <i>Pseudomonas aeruginosa</i> proteins TssB and TagJ form a novel subcomplex in the bacterial type VI secretion system. <i>Molecular Microbiology</i> , 2012, 86, 437-456.	2.5	22
43	Role of the phosphopantetheinyltransferase enzyme, PswP, in the biosynthesis of antimicrobial secondary metabolites by <i>Serratia marcescens</i> Db10. <i>Microbiology (United Kingdom)</i> , 2014, 160, 1609-1617.	1.8	20
44	A New Front in Microbial Warfare—Delivery of Antifungal Effectors by the Type VI Secretion System. <i>Journal of Fungi (Basel, Switzerland)</i> , 2019, 5, 50.	3.5	17
45	The plant pathogen <i>Pectobacterium atrosepticum</i> contains a functional formate hydrogenlyase complex. <i>Molecular Microbiology</i> , 2019, 112, 1440-1452.	2.5	8
46	A new way out: protein localization on the bacterial cell surface via Tat and a novel Type II secretion system. <i>Molecular Microbiology</i> , 2008, 69, 1331-1335.	2.5	6
47	Activation of a [NiFe]-hydrogenase-4 isoenzyme by maturation proteases. <i>Microbiology (United Kingdom)</i> , 2014, 158, 1784-1794.	1.8	5
48	Communication, Cooperation, and Social Interactions: a Report from the Third Young Microbiologists Symposium on Microbe Signalling, Organisation, and Pathogenesis. <i>Journal of Bacteriology</i> , 2014, 196, 3527-3533.	2.2	2
49	Quantitative Determination of Anti-bacterial Activity During Bacterial Co-culture. <i>Methods in Molecular Biology</i> , 2017, 1615, 517-524.	0.9	2
50	Quorum Sensing in the Soft-Rot <i>Erwinias</i> . , 0, , 185-199.		1
51	A Snapshot of the Extraordinary World of Social Microbiology. <i>Journal of Molecular Biology</i> , 2015, 427, 3625-3627.	4.2	0