## Claudia Silva

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7231041/publications.pdf

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		1163117	1058476
15	209	8	14
papers	citations	h-index	g-index
15	15	15	396
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Association of virulence plasmid and antibiotic resistance determinants with chromosomal multilocus genotypes in Mexican Salmonella enterica serovar Typhimurium strains. BMC Microbiology, 2009, 9, 131.	3.3	59
2	Salmonella Typhimurium ST213 is associated with two types of $IncA/C$ plasmids carrying multiple resistance determinants. BMC Microbiology, 2011, 11, 9.	3.3	25
3	Genome analysis of Salmonella enterica subsp. diarizonae isolates from invasive human infections reveals enrichment of virulence-related functions in lineage ST1256. BMC Genomics, 2019, 20, 99.	2.8	24
4	Conjugative transfer of an IncA/C plasmid-borne blaCMY-2gene through genetic re-arrangements with an IncX1 plasmid. BMC Microbiology, 2013, 13, 264.	3.3	20
5	Characterization of Salmonella enterica isolates causing bacteremia in Lima, Peru, using multiple typing methods. PLoS ONE, 2017, 12, e0189946.	2.5	16
6	Complete Genome Sequencing of a Multidrug-Resistant and Human-Invasive Salmonella enterica Serovar Typhimurium Strain of the Emerging Sequence Type 213 Genotype. Genome Announcements, 2015, 3, .	0.8	14
7	A multi-drug resistant Salmonella Typhimurium ST213 human-invasive strain (33676) containing the bla CMY-2 gene on an IncF plasmid is attenuated for virulence in BALB/c mice. BMC Microbiology, 2016, 16, 18.	3.3	13
8	Draft Genome Sequence of Salmonella enterica subsp. <i>enterica</i> Serovar Infantis Strain SPE101, Isolated from a Chronic Human Infection. Genome Announcements, 2017, 5, .	0.8	10
9	Complete Genome Sequence of a Human-Invasive Salmonella enterica Serovar Typhimurium Strain of the Emerging Sequence Type 213 Harboring a Multidrug Resistance IncA/C Plasmid and a <i>bla</i> <sub>CMY-2</sub> -Carrying IncF Plasmid. Genome Announcements, 2015, 3, .	0.8	8
10	IS200 and multilocus sequence typing for the identification of Salmonella enterica serovar Typhi strains from Indonesia. International Microbiology, 2015, 18, 99-104.	2.4	6
11	Complete Genome Sequence of <i>Salmonella enterica</i> Serovar Typhimurium Strain SO3 (Sequence) Tj ETQq1	1.0.7843	14 rgBT /0v
12	Complete Genome Sequence of Salmonella enterica Serovar Typhimurium Strain YU15 (Sequence Type) Tj ETQq0 2016, 4, .	0 0 rgBT /0 0.8	Overlock 10 4
13	Complete Genome Sequence of <i>Salmonella enterica</i> Serovar Typhimurium Strain SO2 (Sequence) Tj ETQq1	1.0.7843	14 rgBT /0v
14	Population analysis of D6-like plasmid prophage variants associated with specific IncC plasmid types in the emerging Salmonella Typhimurium ST213 genotype. PLoS ONE, 2019, 14, e0223975.	2.5	2
15	Self-Conjugation of the EnteropathogenicEscherichia coliAdherence Factor Plasmid of Four Typical EPEC Isolates. BioMed Research International, 2017, 2017, 1-7.	1.9	О