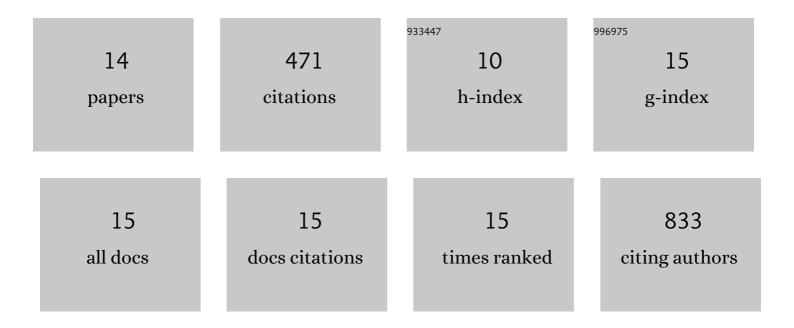
## Servetas, Sl

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4753018/publications.pdf Version: 2024-02-01



SEDVETAS SI

#	Article	IF	CITATIONS
1	Evolution of FMT â $\in$ " From early clinical to standardized treatments. Biologicals, 2022, , .	1.4	3
2	Certhrax Is an Antivirulence Factor for the Anthrax-Like Organism Bacillus cereus Strain G9241. Infection and Immunity, 2018, 86, .	2.2	2
3	Comparative analysis of the Hom family of outer membrane proteins in isolates from two geographically distinct regions: The United States and South Korea. Helicobacter, 2018, 23, e12461.	3.5	18
4	Helicobacter pylori Biofilm Formation Is Differentially Affected by Common Culture Conditions, and Proteins Play a Central Role in the Biofilm Matrix. Applied and Environmental Microbiology, 2018, 84, .	3.1	27
5	Helicobacter pylori Biofilm Formation and Its Potential Role in Pathogenesis. Microbiology and Molecular Biology Reviews, 2018, 82, .	6.6	113
6	ArsRS-Dependent Regulation of homB Contributes to Helicobacter pylori Biofilm Formation. Frontiers in Microbiology, 2018, 9, 1497.	3.5	21
7	Structure-function analyses of metal-binding sites of HypA reveal residues important for hydrogenase maturation in Helicobacter pylori. PLoS ONE, 2017, 12, e0183260.	2.5	16
8	Molecular mechanisms of gastric cancer initiation and progression by Helicobacter pylori. Current Opinion in Infectious Diseases, 2016, 29, 304-310.	3.1	24
9	Characterization of Key Helicobacter pylori Regulators Identifies a Role for ArsRS in Biofilm Formation. Journal of Bacteriology, 2016, 198, 2536-2548.	2.2	34
10	Helicobacter pylori outer membrane protein, HomC, shows geographic dependent polymorphism that is influenced by the Bab family. Journal of Microbiology, 2016, 54, 846-852.	2.8	10
11	Crosstalk between the HpArsRS two-component system and HpNikR is necessary for maximal activation of urease transcription. Frontiers in Microbiology, 2015, 6, 558.	3.5	23
12	Analysis of a single Helicobacter pylori strain over a 10-year period in a primate model. International Journal of Medical Microbiology, 2015, 305, 392-403.	3.6	13
13	Helicobacter pylori bab Paralog Distribution and Association with cagA, vacA, and homA/B Genotypes in American and South Korean Clinical Isolates. PLoS ONE, 2015, 10, e0137078.	2.5	23
14	Comparison of three next-generation sequencing platforms for metagenomic sequencing and identification of pathogens in blood. BMC Genomics, 2014, 15, 96.	2.8	143