Fabien Labroussaa

List of Publications by Year in descending order

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567281 477307 1,465 28 15 29 citations h-index g-index papers 35 35 35 2915 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	SARS-CoV-2 spike D614G change enhances replication and transmission. Nature, 2021, 592, 122-127.	27.8	440
2	Rapid reconstruction of SARS-CoV-2 using a synthetic genomics platform. Nature, 2020, 582, 561-565.	27.8	377
3	Enhanced fitness of SARS-CoV-2 variant of concern Alpha but not Beta. Nature, 2022, 602, 307-313.	27.8	79
4	Impact of donor–recipient phylogenetic distance on bacterial genome transplantation. Nucleic Acids Research, 2016, 44, 8501-8511.	14.5	60
5	Development of safe and highly protective live-attenuated SARS-CoV-2 vaccine candidates by genome recoding. Cell Reports, 2021, 36, 109493.	6.4	46
6	Contrasting Susceptibilities to Flavescence Dor \tilde{A} ©e in Vitis vinifera, Rootstocks and Wild Vitis Species. Frontiers in Plant Science, 2016, 7, 1762.	3.6	45
7	Cloning and Transplantation of the <i>Mesoplasma florum</i> Genome. ACS Synthetic Biology, 2018, 7, 209-217.	3.8	40
8	Specific Evolution of F1-Like ATPases in Mycoplasmas. PLoS ONE, 2012, 7, e38793.	2.5	40
9	Entry of <i>Spiroplasma citri</i> into <i>Circulifer haematoceps</i> Cells Involves Interaction between Spiroplasma Phosphoglycerate Kinase and Leafhopper Actin. Applied and Environmental Microbiology, 2010, 76, 1879-1886.	3.1	39
10	Removal of a Subset of Non-essential Genes Fully Attenuates a Highly Virulent Mycoplasma Strain. Frontiers in Microbiology, 2019, 10, 664.	3.5	31
11	Involvement of a Minimal Actin-Binding Region of Spiroplasma citri Phosphoglycerate Kinase in Spiroplasma Transmission by Its Leafhopper Vector. PLoS ONE, 2011, 6, e17357.	2.5	28
12	Reproduction of contagious caprine pleuropneumonia reveals the ability of convalescent sera to reduce hydrogen peroxide production in vitro. Veterinary Research, 2019, 50, 10.	3.0	24
13	Contagious Bovine and Caprine Pleuropneumonia: a research community's recommendations for the development of better vaccines. Npj Vaccines, 2020, 5, 66.	6.0	23
14	In vivo role of capsular polysaccharide in Mycoplasma mycoides. Journal of Infectious Diseases, 2019, 219, 1559-1563.	4.0	21
15	Genome Engineering of the Fast-Growing <i>Mycoplasma feriruminatoris</i> toward a Live Vaccine Chassis. ACS Synthetic Biology, 2022, 11, 1919-1930.	3.8	16
16	Blocking the Transmission of a Noncirculative Vector-Borne Plant Pathogenic Bacterium. Molecular Plant-Microbe Interactions, 2016, 29, 535-544.	2.6	15
17	A chitinase is required for Xylella fastidiosa colonization of its insect and plant hosts. Microbiology (United Kingdom), 2017, 163, 502-509.	1.8	15
18	Evidence for the Cytoplasmic Localization of the L-α-Glycerophosphate Oxidase in Members of the "Mycoplasma mycoides Cluster― Frontiers in Microbiology, 2019, 10, 1344.	3.5	12

#	Article	IF	CITATIONS
19	Genomics of Plant-Associated Bacteria. , 2014, , .		10
20	Attenuation of a Pathogenic <i>Mycoplasma</i> Strain by Modification of the <i>obg</i> Gene by Using Synthetic Biology Approaches. MSphere, 2019, 4, .	2.9	9
21	Minimalistic mycoplasmas harbor different functional toxin-antitoxin systems. PLoS Genetics, 2021, 17, e1009365.	3.5	7
22	SARS-CoV-2 nanobodies 2.0. Signal Transduction and Targeted Therapy, 2021, 6, 202.	17.1	6
23	In-Yeast Assembly of Coronavirus Infectious cDNA Clones Using a Synthetic Genomics Pipeline. Methods in Molecular Biology, 2020, 2203, 167-184.	0.9	5
24	Vaccination against CCPP in East Africa. Veterinary Record, 2019, 185, 272-272.	0.3	3
25	Establishment of caprine airway epithelial cells grown in an air-liquid interface system to study caprine respiratory viruses and bacteria. Veterinary Microbiology, 2021, 257, 109067.	1.9	3
26	Complete Genome Sequence of Mycoplasma feriruminatoris Strain IVB14/OD_0535, Isolated from an Alpine Ibex in a Swiss Zoo. Microbiology Resource Announcements, 2020, 9, .	0.6	2
27	In-yeast reconstruction of the African swine fever virus genome isolated from clinical samples. STAR Protocols, 2021, 2, 100803.	1.2	2
28	Complete Genome Sequences of the Methicillin-Resistant Strain Staphylococcus aureus 17Gst354 and Its Prophage Staphylococcus Phage vB_StaphS-IVBph354. Microbiology Resource Announcements, 2021, 10, e0058621.	0.6	1