

Wasfi Fares

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

24
papers

242
citations

9
h-index

15
g-index

27
ext. papers

392
ext. citations

5.8
avg, IF

2.68
L-index

#	Paper	IF	Citations
24	A year of genomic surveillance reveals how the SARS-CoV-2 pandemic unfolded in Africa. <i>Science</i> , 2021 , 374, 423-431	33.3	35
23	An integrated overview of the midgut bacterial flora composition of <i>Phlebotomus perniciosus</i> , a vector of zoonotic visceral leishmaniasis in the Western Mediterranean Basin. <i>PLoS Neglected Tropical Diseases</i> , 2017 , 11, e0005484	4.8	27
22	Changes of Sand Fly Populations and <i>Leishmania infantum</i> Infection Rates in an Irrigated Village Located in Arid Central Tunisia. <i>International Journal of Environmental Research and Public Health</i> , 2016 , 13,	4.6	24
21	Sero-epidemiological survey of Crimean-Congo hemorrhagic fever virus in Tunisia. <i>Parasite</i> , 2016 , 23, 10	3	18
20	Isolation, full genomic characterization and neutralization-based human seroprevalence of Medjerda Valley virus, a novel sandfly-borne phlebovirus belonging to the Salehabad virus complex in northern Tunisia. <i>Journal of General Virology</i> , 2016 , 97, 602-610	4.9	16
19	Infection of sand flies collected from different bio-geographical areas of Tunisia with phleboviruses. <i>Acta Tropica</i> , 2015 , 141, 1-6	3.2	15
18	Presence of sandfly-borne phleboviruses of two antigenic complexes (Sandfly fever Naples virus and Sandfly fever Sicilian virus) in two different bio-geographical regions of Tunisia demonstrated by a microneutralisation-based seroprevalence study in dogs. <i>Parasites and Vectors</i> , 2014 , 7, 476	4	15
17	Phylogenetic analysis of complete VP1 sequences of echoviruses 11 and 6: high genetic diversity and circulation of genotypes with a wide geographical and temporal range. <i>Journal of Medical Microbiology</i> , 2011 , 60, 1017-1025	3.2	14
16	Serologic evidence of exposure to Rift Valley fever virus detected in Tunisia. <i>New Microbes and New Infections</i> , 2016 , 9, 1-7	4.1	13
15	Phleboviruses associated with sand flies in arid bio-geographical areas of Central Tunisia. <i>Acta Tropica</i> , 2016 , 158, 13-19	3.2	9
14	Update on molecular characterization of coxsackievirus B5 strains. <i>Journal of Medical Virology</i> , 2011 , 83, 1247-54	19.7	9
13	<i>Ixodes inopinatus</i> and <i>Ixodes ricinus</i> (Acari: Ixodidae) Are Sympatric Ticks in North Africa. <i>Journal of Medical Entomology</i> , 2020 , 57, 952-956	2.2	9
12	Co-circulation of Toscana virus and <i>Leishmania infantum</i> in a focus of zoonotic visceral leishmaniasis from Central Tunisia. <i>Acta Tropica</i> , 2020 , 204, 105342	3.2	6
11	The Impact of Illegal Waste Sites on the Transmission of Zoonotic Cutaneous Leishmaniasis in Central Tunisia. <i>International Journal of Environmental Research and Public Health</i> , 2020 , 18,	4.6	5
10	Tick-borne encephalitis virus in <i>Ixodes ricinus</i> (Acari: Ixodidae) ticks, Tunisia. <i>Ticks and Tick-borne Diseases</i> , 2021 , 12, 101606	3.6	5
9	Multiplexed Magnetofluorescent Bioplatfrom for the Sensitive Detection of SARS-CoV-2 Viral RNA without Nucleic Acid Amplification. <i>Analytical Chemistry</i> , 2021 , 93, 11225-11232	7.8	5
8	Sequencing Using a Two-Step Strategy Reveals High Genetic Diversity in the S Gene of SARS-CoV-2 after a High-Transmission Period in Tunis, Tunisia. <i>Microbiology Spectrum</i> , 2021 , e0063921	8.9	4

7	Absence of Crimean-Congo haemorrhagic fever virus in the tick <i>Hyalomma aegyptium</i> parasitizing the spur-thighed tortoise (<i>Testudo graeca</i>) in Tunisia. <i>Parasite</i> , 2019 , 26, 35	3	3
6	Genetic characterization of West Nile Virus strains during neuroinvasives infection outbreak in Tunisia, 2018. <i>Transboundary and Emerging Diseases</i> , 2021 , 68, 2414-2421	4.2	3
5	Molecular Epidemiology of SARS-CoV-2 in Tunisia (North Africa) through Several Successive Waves of COVID-19.. <i>Viruses</i> , 2022 , 14,	6.2	2
4	The value of West Nile virus RNA detection by real-time RT-PCR in urine samples from patients with neuroinvasive forms.. <i>Archives of Microbiology</i> , 2022 , 204, 238	3	2
3	Risk Assessment of the Role of the Ecotones in the Transmission of Zoonotic Cutaneous Leishmaniasis in Central Tunisia. <i>International Journal of Environmental Research and Public Health</i> , 2021 , 18,	4.6	1
2	SARS-CoV2 RT-PCR assays: In vitro comparison of 4 WHO approved protocols on clinical specimens and its implications for real laboratory practice through variant emergence.. <i>Virology Journal</i> , 2022 , 19, 54	6.1	1
1	Whole genome sequencing and phylogenetic analysis of six SARS-CoV-2 strains isolated during COVID-19 pandemic in Tunisia, North Africa. <i>BMC Genomics</i> , 2021 , 22, 540	4.5	0