Sahal A Al-Hajoj

List of Publications by Year
in descending order

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43 papers	1,751 citations	567281 15 h-index	276875 41 g-index
44 all docs	44 docs citations	44 times ranked	2225 citing authors

#	Article	IF	Citations
1	Sub-Lineage Specific Phenolic Glycolipid Patterns in the Mycobacterium tuberculosis Complex Lineage 1. Frontiers in Microbiology, 2022, 13, 832054.	3.5	3
2	Origin and Global Expansion of Mycobacterium tuberculosis Complex Lineage 3. Genes, 2022, 13, 990.	2.4	13
3	Family cluster of multi-drug resistant tuberculosis in Kingdom of Saudi Arabia. Journal of Infection and Public Health, 2020, 13, 154-157.	4.1	3
4	Mycobacterium tuberculosis DNA in living donor transplanted livers and donorâ€related tuberculosis in recipients: A retrospective longitudinal cohort study. Transplant Infectious Disease, 2020, 22, e13212.	1.7	3
5	Burden of non-tuberculous mycobacterial diseases in Saudi Arabian children: The first nationwide experience. Journal of Infection and Public Health, 2019, 12, 803-808.	4.1	6
6	Demographic risk factors for extra-pulmonary tuberculosis among adolescents and adults in Saudi Arabia. PLoS ONE, 2019, 14, e0213846.	2.5	13
7	Clinical Management of Drug-resistant Mycobacterium tuberculosis Strains: Pathogen-targeted Versus Host-directed Treatment Approaches. Current Pharmaceutical Biotechnology, 2019, 20, 272-284.	1.6	3
8	Impact of Mycobacterium tuberculosis complex lineages as a determinant of disease phenotypes from an immigrant rich moderate tuberculosis burden country. Respiratory Research, 2018, 19, 259.	3.6	9
9	Drug-resistance profiling and transmission dynamics of multidrug-resistant Mycobacterium tuberculosis in Saudi Arabia revealed by whole genome sequencing. Infection and Drug Resistance, 2018, Volume 11, 2219-2229.	2.7	17
10	The first Saudi Arabian national inventory study revealed the upcoming challenges of highly diverse non-tuberculous mycobacterial diseases. PLoS Neglected Tropical Diseases, 2018, 12, e0006515.	3.0	6
11	QuantiFERON-TB Gold In-Tube in Saudi Arabia benchmarked with other sites of the Middle East: A meta-analysis review. Journal of Infection in Developing Countries, 2018, 12, 687-699.	1.2	0
12	Diagnostic potential of interferonâ€gamma release assay to detect latent tuberculosis infection in kidney transplant recipients. Transplant Infectious Disease, 2017, 19, e12675.	1.7	20
13	Mycobacterium riyadhensein Saudi Arabia. Emerging Infectious Diseases, 2017, 23, 1732-1734.	4.3	7
14	Nontuberculous Mycobacteria in Saudi Arabia and Gulf Countries: A Review. Canadian Respiratory Journal, 2017, 2017, 1-13.	1.6	15
15	Diversity and evolution of drug resistance mechanisms in Mycobacterium tuberculosis. Infection and Drug Resistance, 2017, Volume 10, 333-342.	2.7	31
16	Emergence of Rare Species of Nontuberculous Mycobacteria as Potential Pathogens in Saudi Arabian Clinical Setting. PLoS Neglected Tropical Diseases, 2017, 11, e0005288.	3.0	21
17	First Insight Into the Fluoroquinolone and Aminoglycoside Resistance of Multidrug-Resistant Mycobacterium tuberculosis in Saudi Arabia. American Journal of Tropical Medicine and Hygiene, 2017, 96, 1066-1070.	1.4	4
18	Risk factors for tuberculosis and beyond. International Journal of Mycobacteriology, 2017, 6, 326.	0.6	1

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19	Drug-resistant tuberculosis viewed from bacterial and host genomes. International Journal of Antimicrobial Agents, 2016, 48, 353-360.	2.5	9
20	Mycobacterium tuberculosis lineage 4 comprises globally distributed and geographically restricted sublineages. Nature Genetics, 2016, 48, 1535-1543.	21.4	326
21	Interferon Gamma Release Assay versus Tuberculin Skin Testing among Healthcare Workers of Highly Diverse Origin in a Moderate Tuberculosis Burden Country. PLoS ONE, 2016, 11, e0154803.	2.5	19
22	Tuberculosis in Saudi Arabia: the journey across time. Journal of Infection in Developing Countries, 2015, 9, 222-231.	1.2	24
23	Mapping the epidemiology and trends of extra-pulmonary tuberculosis in Saudi Arabia. International Journal of Mycobacteriology, 2015, 4, 261-269.	0.6	17
24	Exploring the Sociodemographic and Clinical Features of Extrapulmonary Tuberculosis in Saudi Arabia. PLoS ONE, 2015, 10, e0101667.	2.5	26
25	Occurrence of Diverse Mutations in Isoniazid- and Rifampicin-ResistantMycobacterium tuberculosisIsolates from Autochthonous and Immigrant Populations of Saudi Arabia. Microbial Drug Resistance, 2014, 20, 623-631.	2.0	9
26	Molecular Confirmation of Bacillus Calmette Guerin Vaccine Related Adverse Events among Saudi Arabian Children. PLoS ONE, 2014, 9, e113472.	2.5	12
27	Current trends of Mycobacterium tuberculosis molecular epidemiology in Saudi Arabia and associated demographical factors. Infection, Genetics and Evolution, 2013, 16, 362-368.	2.3	23
28	Inconsistencies in drug susceptibility testing of Mycobacterium tuberculosis: Current riddles and recommendations. International Journal of Mycobacteriology, 2013, 2, 14-17.	0.6	3
29	Endogenous reactivation followed by exogenous re-infection with drug resistant strains, a new challenge for tuberculosis control inÂSaudi Arabia. Tuberculosis, 2013, 93, 246-249.	1.9	17
30	Qualitative research: Is this a missing link to control tuberculosis in Saudi Arabia?. International Journal of Mycobacteriology, 2013, 2, 126-127.	0.6	2
31	Emergence of Clinically Relevant Non-Tuberculous Mycobacterial Infections in Saudi Arabia. PLoS Neglected Tropical Diseases, 2013, 7, e2234.	3.0	43
32	Epidemiology of Antituberculosis Drug Resistance in Saudi Arabia: Findings of the First National Survey. Antimicrobial Agents and Chemotherapy, 2013, 57, 2161-2166.	3.2	44
33	Tuberculosis Transmission among Immigrants and Autochthonous Populations of the Eastern Province of Saudi Arabia. PLoS ONE, 2013, 8, e77635.	2.5	23
34	Admixed Phylogenetic Distribution of Drug Resistant Mycobacterium tuberculosis in Saudi Arabia. PLoS ONE, 2013, 8, e55598.	2.5	26
35	Mycobacterium riyadhense overlooked: we can only find what we are looking for. Journal of Infection in Developing Countries, 2013, 7, 293-294.	1.2	2
36	New insight into the molecular characterization of isoniazid and rifampicin resistant Mycobacterium tuberculosis strains from Saudi Arabia. Infection, Genetics and Evolution, 2012, 12, 549-556.	2.3	13

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37	First case report of chronic pulmonary lung disease caused by Mycobacterium abscessus in two immunocompetent patients in Saudi Arabia. Annals of Saudi Medicine, 2012, 32, 312-314.	1.1	18
38	The emergence of Beijing genotype of mycobacterium tuberculosis in the Kingdom of Saudi Arabia. Annals of Thoracic Medicine, 2010, 5, 149.	1.8	12
39	Molecular strain typing of Mycobacterium tuberculosis isolates to detect cross-contamination events. Proposed modifications to prevent its recurrence. Journal of King Abdulaziz University, Islamic Economics, 2009, 30, 1515-9.	1.1	1
40	Usefulness of molecular techniques to identify ongoing tuberculosis transmission in Saudi Arabia. Journal of King Abdulaziz University, Islamic Economics, 2007, 28, 268-70.	1.1	3
41	Is Saudi Arabia a fertile land for exchanging infectious diseases?. Journal of King Abdulaziz University, Islamic Economics, 2007, 28, 803-4.	1.1	0
42	Mycobacterium tuberculosis complex genetic diversity: mining the fourth international spoligotyping database (SpolDB4) for classification, population genetics and epidemiology. BMC Microbiology, 2006, 6, 23.	3.3	900
43	Role of tuberculosis laboratories in Saudi Arabia. A call to implement standardized procedures. Journal of King Abdulaziz University, Islamic Economics, 2004, 25, 1545-8.	1.1	4