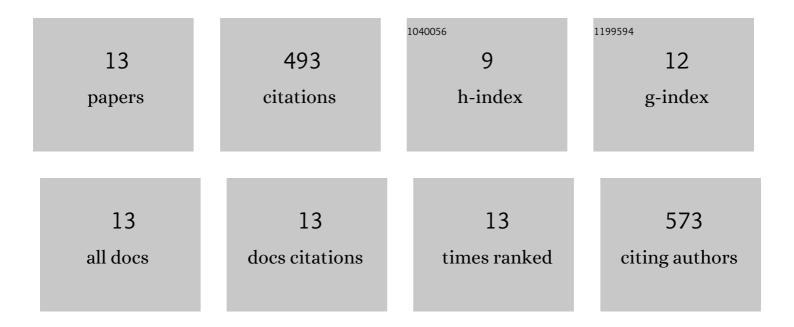
Albert T Young

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4353277/publications.pdf Version: 2024-02-01



ALBERT T YOUNG

#	Article	IF	CITATIONS
1	The role of technology in melanoma screening and diagnosis. Pigment Cell and Melanoma Research, 2021, 34, 288-300.	3.3	22
2	Stress testing reveals gaps in clinic readiness of image-based diagnostic artificial intelligence models. Npj Digital Medicine, 2021, 4, 10.	10.9	25
3	Idiopathic pure sudomotor failure: A review and two cases. International Journal of Women's Dermatology, 2021, 7, 276-279.	2.0	3
4	Association of Wildfire Air Pollution and Health Care Use for Atopic Dermatitis and Itch. JAMA Dermatology, 2021, 157, 658.	4.1	56
5	An abdominal skin lesion: to lump or split? a case presentation. Dermatology Online Journal, 2021, 27, .	0.5	0
6	Patient and general public attitudes towards clinical artificial intelligence: a mixed methods systematic review. The Lancet Digital Health, 2021, 3, e599-e611.	12.3	88
7	Multi-omic regulatory networks capture downstream effects of kinase inhibition in Mycobacterium tuberculosis. Npj Systems Biology and Applications, 2021, 7, 8.	3.0	3
8	Machine learning for the prediction of antimicrobial stewardship intervention in hospitalized patients receiving broad-spectrum agents. Infection Control and Hospital Epidemiology, 2020, 41, 1022-1027.	1.8	10
9	Artificial Intelligence in Dermatology: A Primer. Journal of Investigative Dermatology, 2020, 140, 1504-1512.	0.7	100
10	Artificial Intelligence in Teledermatology. Current Dermatology Reports, 2019, 8, 85-90.	2.1	12
11	Multisystem Analysis of <i>Mycobacterium tuberculosis</i> Reveals Kinase-Dependent Remodeling of the Pathogen-Environment Interface. MBio, 2018, 9, .	4.1	57
12	Development and Validation of an Electronic Health Record–Based Machine Learning Model to Estimate Delirium Risk in Newly Hospitalized Patients Without Known Cognitive Impairment. JAMA Network Open, 2018, 1, e181018.	5.9	102
13	Estimating gene regulatory networks with pandaR. Bioinformatics, 2017, 33, 2232-2234.	4.1	15