List of Publications by Year in descending order

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#	Article	lF	CITATIONS
1	Analysis of inflammation-related microRNA expression in patients with ankylosing spondylitis. Immunologic Research, 2022, 70, 23-32.	2.9	4
2	Dense fine speckled immunofluorescence pattern in a Chinese population: Prevalence and clinical association. Journal of Clinical Laboratory Analysis, 2022, 36, e24173.	2.1	3
3	Association of genetic polymorphisms in IL-23R and IL-17A with the susceptibility to IgA nephropathy in a Chinese Han population. Genes and Immunity, 2022, 23, 33-41.	4.1	4
4	ETS1 polymorphism rs73013527 in relation to serum RANKL levels among patients with RA. Medicine (United States), 2021, 100, e24562.	1.0	5
5	Antiâ€eentrosome antibodies: Prevalence and disease association in Chinese population. Scandinavian Journal of Immunology, 2019, 90, e12803.	2.7	7
6	Prevalence of foodâ€allergen and aeroallergen sensitization among people in Sichuan, Western China: An 8â€year observational study. Journal of Clinical Laboratory Analysis, 2019, 33, e22723.	2.1	8
7	Genetic polymorphism of rs9277535 in HLA-DP associated with rheumatoid arthritis and anti-CCP production in a Chinese population. Clinical Rheumatology, 2018, 37, 1799-1805.	2.2	11
8	IL- $1 < i > \hat{1}^2 < /i >$ and IL-6 Are Highly Expressed in RF+lgE+ Systemic Lupus Erythematous Subtype. Journal of Immunology Research, 2017, 2017, 1-8.	2.2	21
9	IL-23R and IL-17A polymorphisms correlate with susceptibility of ankylosing spondylitis in a Southwest Chinese population. Oncotarget, 2017, 8, 70310-70316.	1.8	15
10	The Expression of LIGHT Was Increased and the Expression of HVEM and BTLA Were Decreased in the T Cells of Patients with Rheumatoid Arthritis. PLoS ONE, 2016, 11, e0155345.	2.5	11
11	Association of E26 Transformation Specific Sequence 1 Variants with Rheumatoid Arthritis in Chinese Han Population. PLoS ONE, 2015, 10, e0134875.	2.5	8
12	Association of polymorphisms in pre-miRNA with inflammatory biomarkers in rheumatoid arthritis in the Chinese Han population. Human Immunology, 2012, 73, 101-106.	2.4	77
13	Association of pre-microRNAs genetic variants with susceptibility in systemic lupus erythematosus. Molecular Biology Reports, 2011, 38, 1463-1468.	2.3	42
14	Systemic Lupus Erythematosus (SLE) Risk Factors: Novel Proteins Detected From Familial SLE Using Proteomics. Laboratory Medicine, 2009, 40, 408-411.	1.2	4