Rowaida Z Taha

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Transcriptomic Profiling of Circulating HLA-DR [–] Myeloid Cells, Compared with HLA-DR ⁺ Myeloid Antigen-presenting Cells. Immunological Investigations, 2021, 50, 952-963.	2.0	2
2	Differential gene expression of tumor-infiltrating CD33+ myeloid cells in advanced- versus early-stage colorectal cancer. Cancer Immunology, Immunotherapy, 2021, 70, 803-815.	4.2	15
3	Tâ€cell responses and therapies against SARSâ€CoVâ€2 infection. Immunology, 2021, 162, 30-43.	4.4	159
4	Transcriptome of Tumor-Infiltrating T Cells in Colorectal Cancer Patients Uncovered a Unique Gene Signature in CD4+ T Cells Associated with Poor Disease-Specific Survival. Vaccines, 2021, 9, 334.	4.4	5
5	Family-Based Genome-Wide Association Study of Autism Spectrum Disorder in Middle Eastern Families. Genes, 2021, 12, 761.	2.4	7
6	Epigenetic regulation of immune checkpoints and TÂcell exhaustion markers in tumor-infiltrating T cells of colorectal cancer patients. Epigenomics, 2020, 12, 1871-1882.	2.1	11
7	Differential gene expression of tumor-infiltrating CD4 ⁺ T cells in advanced versus early stage colorectal cancer and identification of a gene signature of poor prognosis. Oncolmmunology, 2020, 9, 1825178.	4.6	6
8	Differential gene expression of tumor-infiltrating CD8 ⁺ T cells in advanced versus early-stage colorectal cancer and identification of a gene signature of poor prognosis. , 2020, 8, e001294.		25
9	DNA methylation in the promoters of PD-L1, MMP9, ARG1, galectin-9, TIM-3, VISTA and TGF-β genes in HLA-DR [–] myeloid cells, compared with HLA-DR ⁺ antigen-presenting cells. Epigenetics, 2020, 15, 1275-1288.	2.7	21
10	Expression of immune checkpoints and T cell exhaustion markers in early and advanced stages of colorectal cancer. Cancer Immunology, Immunotherapy, 2020, 69, 1989-1999.	4.2	75
11	Transcriptomic Profiling of Tumor-Infiltrating CD4+TIM-3+ T Cells Reveals Their Suppressive, Exhausted, and Metastatic Characteristics in Colorectal Cancer Patients. Vaccines, 2020, 8, 71.	4.4	19
12	Transcriptomic profiling disclosed the role of DNA methylation and histone modifications in tumor-infiltrating myeloid-derived suppressor cell subsets in colorectal cancer. Clinical Epigenetics, 2020, 12, 13.	4.1	52
13	PD-L1 Blockade by Atezolizumab Downregulates Signaling Pathways Associated with Tumor Growth, Metastasis, and Hypoxia in Human Triple Negative Breast Cancer. Cancers, 2019, 11, 1050.	3.7	50
14	DNA methylation of immune checkpoints in the peripheral blood of breast and colorectal cancer patients. Oncolmmunology, 2019, 8, e1542918.	4.6	54
15	Multiplex epithelium dysfunction due to CLDN10 mutation: the HELIX syndrome. Genetics in Medicine, 2018, 20, 190-201.	2.4	75
16	DNA methylation and repressive H3K9 and H3K27 trimethylation in the promoter regions of PD-1, CTLA-4, TIM-3, LAG-3, TIGIT, and PD-L1 genes in human primary breast cancer. Clinical Epigenetics, 2018, 10, 78.	4.1	103
17	DNA methylation and repressive histones in the promoters of PD-1, CTLA-4, TIM-3, LAG-3, TIGIT, PD-L1, and galectin-9 genes in human colorectal cancer. Clinical Epigenetics, 2018, 10, 104.	4.1	68
18	The M694I/M694I genotype: A genetic risk factor of AA-amyloidosis in a group of Algerian patients with familial Mediterranean fever. European Journal of Medical Genetics, 2017, 60, 149-153.	1.3	14

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19	A chromosomal microdeletion of 15q in a female patient with epilepsy, <scp>ID</scp> , and autism spectrum disorder: a case report. Clinical Case Reports (discontinued), 2017, 5, 1013-1017.	0.5	11
20	Chronic recurrent multifocal osteomyelitis in a patient with Familial Mediterranean Fever. Pediatric Rheumatology, 2015, 13, .	2.1	4
21	Overlap of Familial Mediterranean Fever and Hyper-IgD Syndrome in an Arabic Kindred. Journal of Clinical Immunology, 2015, 35, 249-253.	3.8	15