

Cheng Qing

List of Publications by Year in descending order

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12
papers

336
citations

1040056

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1199594

12
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12
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docs citations

12
times ranked

585
citing authors

#	ARTICLE	IF	CITATIONS
1	DNA Methylation-based Diagnostic and Prognostic Biomarkers for Glioblastoma. <i>Cell Transplantation</i> , 2020, 29, 096368972093324.	2.5	11
2	Distinct expression and prognostic value of members of SMAD family in non-small cell lung cancer. <i>Medicine (United States)</i> , 2020, 99, e19451.	1.0	13
3	Expression profile and diagnostic value of circRNAs in peripheral blood from patients with systemic lupus erythematosus. <i>Molecular Medicine Reports</i> , 2020, 23, 1-1.	2.4	27
4	Mitochondrial DNA plays an important role in lung injury induced by sepsis. <i>Journal of Cellular Biochemistry</i> , 2019, 120, 8547-8560.	2.6	7
5	Serum PGLYRP α 1 is a highly discriminatory biomarker for the diagnosis of rheumatoid arthritis. <i>Molecular Medicine Reports</i> , 2019, 19, 589-594.	2.4	16
6	The Expression of lncRNA NEAT1 in Human Tuberculosis and Its Antituberculosis Effect. <i>BioMed Research International</i> , 2018, 2018, 1-8.	1.9	39
7	Plasma Circular RNAs hsa_circ_0001953 and hsa_circ_0009024 as Diagnostic Biomarkers for Active Tuberculosis. <i>Frontiers in Microbiology</i> , 2018, 9, 2010.	3.5	33
8	Overexpression of CD64 on CD14 ⁺⁺ CD16 ⁺ and CD14 ⁺⁺ CD16 ⁺ monocytes of rheumatoid arthritis patients correlates with disease activity. <i>Experimental and Therapeutic Medicine</i> , 2018, 16, 2703-2711.	1.8	13
9	Decreased expression of TIGIT in NK cells correlates negatively with disease activity in systemic lupus erythematosus. <i>International Journal of Clinical and Experimental Pathology</i> , 2018, 11, 2408-2418.	0.5	5
10	Diagnostic Value of the lncRNA <i>NEAT1</i> in Peripheral Blood Mononuclear Cells of Patients with Sepsis. <i>Disease Markers</i> , 2017, 2017, 1-6.	1.3	41
11	Identification of Differentially Expressed Long Non-coding RNAs in Polarized Macrophages. <i>Scientific Reports</i> , 2016, 6, 19705.	3.3	63
12	The long non-coding RNA CCAT2 is up-regulated in ovarian cancer and associated with poor prognosis. <i>Diagnostic Pathology</i> , 2016, 11, 49.	2.0	68