

Bong Jun Kim

List of Publications by Year in descending order

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Version: 2024-02-01

20
papers

169
citations

1163117

8
h-index

1199594

12
g-index

20
all docs

20
docs citations

20
times ranked

130
citing authors

#	ARTICLE	IF	CITATIONS
1	Extracellular Mitochondrial Dysfunction in Cerebrospinal Fluid of Patients with Delayed Cerebral Ischemia after Aneurysmal Subarachnoid Hemorrhage. <i>Neurocritical Care</i> , 2020, 33, 422-428.	2.4	26
2	Genomic Variations in Susceptibility to Intracranial Aneurysm in the Korean Population. <i>Journal of Clinical Medicine</i> , 2019, 8, 275.	2.4	24
3	Correlation Between Altered DNA Methylation of Intergenic Regions of ITPR3 and Development of Delayed Cerebral Ischemia in Patients with Subarachnoid Hemorrhage. <i>World Neurosurgery</i> , 2019, 130, e449-e456.	1.3	17
4	Genome-wide blood DNA methylation analysis in patients with delayed cerebral ischemia after subarachnoid hemorrhage. <i>Scientific Reports</i> , 2020, 10, 11419.	3.3	16
5	Mitochondrial dysfunction associated with autophagy and mitophagy in cerebrospinal fluid cells of patients with delayed cerebral ischemia following subarachnoid hemorrhage. <i>Scientific Reports</i> , 2021, 11, 16512.	3.3	16
6	Study of Correlation Between Hp $\hat{\pm}$ 1 Expression of Haptoglobin 2-1 and Clinical Course in Aneurysmal Subarachnoid Hemorrhage. <i>World Neurosurgery</i> , 2018, 117, e221-e227.	1.3	11
7	A Preliminary Study of the Association between <i>SOX17</i> Gene Variants and Intracranial Aneurysms Using Exome Sequencing. <i>Journal of Korean Neurosurgical Society</i> , 2020, 63, 559-565.	1.2	11
8	Association of SOX17 Gene Polymorphisms and Intracranial Aneurysm: A Case-Control Study and Meta-Analysis. <i>World Neurosurgery</i> , 2018, 110, e823-e829.	1.3	8
9	The Role of Consecutive Plasma Copeptin Levels in the Screening of Delayed Cerebral Ischemia in Poor-Grade Subarachnoid Hemorrhage. <i>Life</i> , 2021, 11, 274.	2.4	7
10	Characterization of the TCR $\hat{\pm}$ 2 Chain CDR3 Repertoire in Subarachnoid Hemorrhage Patients with Delayed Cerebral Ischemia. <i>International Journal of Molecular Sciences</i> , 2020, 21, 3149.	4.1	6
11	Genome-Wide Association Study of Clinical Outcome After Aneurysmal Subarachnoid Haemorrhage: Protocol. <i>Translational Stroke Research</i> , 2022, 13, 565-576.	4.2	5
12	Recent Stem Cell Research on Hemorrhagic Stroke : An Update. <i>Journal of Korean Neurosurgical Society</i> , 2022, 65, 161-172.	1.2	4
13	Bioinformatics Analysis of Autophagy and Mitophagy Markers Associated with Delayed Cerebral Ischemia Following Subarachnoid Hemorrhage. <i>Journal of Korean Neurosurgical Society</i> , 2022, 65, 236-244.	1.2	4
14	Association of Haptoglobin Phenotype With Neurological and Cognitive Outcomes in Patients With Subarachnoid Hemorrhage. <i>Frontiers in Aging Neuroscience</i> , 2022, 14, 819628.	3.4	3
15	Genome-wide polygenic risk impact on intracranial aneurysms and acute ischemic stroke. <i>PLoS ONE</i> , 2022, 17, e0265581.	2.5	3
16	Identification of Differentially-Methylated Genes and Pathways in Patients with Delayed Cerebral Ischemia Following Subarachnoid Hemorrhage. <i>Journal of Korean Neurosurgical Society</i> , 2022, 65, 4-12.	1.2	2
17	Genome-Wide Association Study of the Relationship Between Matrix Metalloproteinases and		

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19	Profiling of T Cell Receptor β -Chain Complimentary Determining Regions 3 Repertoire in Subarachnoid Hemorrhage Patients Using High-Throughput Sequencing. <i>Journal of Korean Neurosurgical Society</i> , 2021, 64, 505-513.	1.2	1
20	Genome-Wide Association Study of the Relationship Between Matrix Metalloproteinases and		