Alexei A Sleptcov

List of Publications by Year in descending order

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| # | Article | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | Human exome sequence data in support of somatic mosaicism in carotid atherosclerosis. Data in Brief, 2021, 39, 107656. | 1.0 | 3 |
| 2 | Deoxyribonucleic acid methylation in the enhancer region of the CDKN2A/2B and CDKN2B-AS1 genes in blood vessels and cells in patients with carotid atherosclerosis. Russian Journal of Cardiology, 2020, 25, 4060. | 1.4 | 1 |
| 3 | Comparative Analysis of Gene Expression in Vascular Cells of Patients with Advanced Atherosclerosis. Biochemistry (Moscow) Supplement Series B: Biomedical Chemistry, 2019, 13, 74-80. | 0.4 | 0 |
| 4 | Genomic alterations in cells involved in the atherosclerotic process. Atherosclerosis, 2018, 275, e134. | 0.8 | 0 |
| 5 | Mitochondrial DNA polymorphism study in patients with carotid atherosclerosis suggests protective effect of haplogroup J. Atherosclerosis, 2018, 275, e187-e188. | 0.8 | 0 |
| 6 | Genomic structural variations for cardiovascular and metabolic comorbidity. Scientific Reports, 2017, 7, 41268. | 3.3 | 29 |
| 7 | Copy number variations in patients with advanced coronary artery disease. Atherosclerosis, 2017, 263, e90. | 0.8 | 0 |
| 8 | Analysis of genomic rearrangements in macrophages dissected from human atherosclerotic plaques. Atherosclerosis, 2017, 263, e278. | 0.8 | 0 |
| 9 | Clinically relevant morphological structures in breast cancer represent transcriptionally distinct tumor cell populations with varied degrees of epithelial-mesenchymal transition and CD44+CD24-stemness. Oncotarget, 2017, 8, 61163-61180. | 1.8 | 22 |
| 10 | STRUCTURAL VARIABLITY OF LEUCOCYTE GENOME AND ARTERIAL CELLS IN HUMAN ATHEROSCLEROSIS. Russian Journal of Cardiology, 2017, , 140-146. | 1.4 | 0 |
| 11 | IDENTIFICATION OF DIFFERENTLY METYLATED GENES POTENTIALLY RELATED TO HUMAN ATHEROSCLEROSIS. Russian Journal of Cardiology, 2017, , 42-48. | 1.4 | 12 |
| 12 | Genes for fibrogenesis in the determination of susceptibility to myocardial infarction. Molecular Biology, 2016, 50, 81-90. | 1.3 | 6 |
| 13 | DNA methylation and copy number events in atherosclerotic lesions. Atherosclerosis, 2016, 252, e83. | 0.8 | 1 |
| 14 | Genomic rearrangements in human atherosclerotic vascular tissues. Atherosclerosis, 2016, 252, e167-e168. | 0.8 | 0 |
| 15 | Analysis of heteroplasmy in the major noncoding region of mitochondrial DNA in the blood and atherosclerotic plaques of carotid arteries. Russian Journal of Genetics, 2016, 52, 436-440. | 0.6 | 1 |
| 16 | A Comparison of Genome-Wide DNA Methylation Patterns between Different Vascular Tissues from Patients with Coronary Heart Disease. PLoS ONE, 2015, 10, e0122601. | 2.5 | 54 |
| 17 | Somatic genome variations in vascular tissues and peripheral blood leukocytes in patients with atherosclerosis. Russian Journal of Genetics, 2014, 50, 870-878. | 0.6 | 6 |
| 18 | DNA methylation profiling of the vascular tissues in the setting of atherosclerosis. Molecular Biology, 2013, 47, 352-357. | 1.3 | 10 |

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|----|--|-----|-----------|
| 19 | Methylation profile of INK4B-ARF-INK4A locus in atherosclerosis. Russian Journal of Genetics, 2013, 49, 681-684. | 0.6 | 0 |