## Artem M Kiselev

## List of Publications by Year in descending order

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Version: 2024-02-01


Systematic dissection of biases in whole-exome and whole-genome sequencing reveals major
Systematic dissection of biases in whole-exome and whole-genome sequencing rever
determinants of coding sequence coverage. Scientific Reports, 2020, 10, 2057.
$3.3 \quad 72$

8 Dysregulation of Notch signaling in cardiac mesenchymal cells of patients with tetralogy of Fallot.
Pediatric Research, 2020, 88, 38-47.

Atomic Mechanisms of Timothy Syndrome-Associated Mutations in Calcium Channel Cav1.2. Frontiers
in Physiology, 2019, 10, 335.

Characterization of a novel SCN5A genetic variant A1294G associated with mixed clinical phenotype.
10 Biochemical and Biophysical Research Communications, 2019, 516, 777-783.
2.1

8

New germline BRCA2 gene variant in the Tuvinian Mongol breast cancer patients. Molecular Biology
Reports, 2019, 46, 5537-5541.
Truncating Variant in Myof Gene Is Associated With Limb-Girdle Type Muscular Dystrophy and
$12 \quad$ Cardiomyopathy. Frontiers in Cenetics, 2019, 10, 608.
2.3

10

Draft Genome Sequence of <i>Coxiella burnetii</i>Historical Strain Leningrad-2, Isolated from Blood
13 Draft Genome Sequence of <i>Coxielia burnetii\lli>Historical Strain Leningrad-2, Isolated from Batient with Acute Q Fever in Saint Petersburg, Russia. Genome Announcements, 2018, 6, .
0.8

1

De novo mutations in <i>FLNC</i> leading to early-onset restrictive cardiomyopathy and congenital myopathy. Human Mutation, 2018, 39, 1161-1172.
2.5

49
Rare Case of Ulnar-Mammary-Like Syndrome With Left Ventricular Tachycardia and Lack of TBX3 Mutation. Frontiers in Genetics, 2018, 9, 209.
2.3

Structural consequences of mutations associated with idiopathic restrictive cardiomyopathy. Amino
Acids, 2017, 49, 1815-1829.

The exon junction complex factor Y14 is dynamic in the nucleus of the beetle Tribolium castaneum
during late oogenesis. Molecular Cytogenetics, 2017, 10, 41. PLoS ONE, 2016, 11, e0163362.

