## Konstantin S Vassilenko

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Abstract P-27: The 30S Ribosomal Subunit Assembly Factor Rbfa Plays a Key Role in the Formation of the Central Pseudoknot and in the Correct Docking of Helix 44 of the Decoding Center. International Journal of Biomedicine, 2021, 11, S23-S24.	0.2	0
2	RbfA Is Involved in Two Important Stages of 30S Subunit Assembly: Formation of the Central Pseudoknot and Docking of Helix 44 to the Decoding Center. International Journal of Molecular Sciences, 2021, 22, 6140.	4.1	14
3	Non-Canonical Translation Initiation Mechanisms Employed by Eukaryotic Viral mRNAs. Biochemistry (Moscow), 2021, 86, 1060-1094.	1.5	22
4	Functional Cyclization of Eukaryotic mRNAs. International Journal of Molecular Sciences, 2020, 21, 1677.	4.1	31
5	Unusual dicistronic expression from closely spaced initiation codons in an umbravirus subgenomic RNA. Nucleic Acids Research, 2018, 46, 11726-11742.	14.5	12
6	Concerted action of two 3′ cap-independent translation enhancers increases the competitive strength of translated viral genomes. Nucleic Acids Research, 2017, 45, 9558-9572.	14.5	23
7	Identification of a Mg2+-sensitive ORF in the 5′-leader of TRPM7 magnesium channel mRNA. Nucleic Acids Research, 2014, 42, 12779-12788.	14.5	16
8	Translation initiation in eukaryotes: Versatility of the scanning model. Biochemistry (Moscow), 2012, 77, 1465-1477.	1.5	20
9	Unidirectional constant rate motion of the ribosomal scanning particle during eukaryotic translation initiation. Nucleic Acids Research, 2011, 39, 5555-5567.	14.5	71
10	Quantitative analysis of ribosome–mRNA complexes at different translation stages. Nucleic Acids Research, 2010, 38, e15-e15.	14.5	32
11	Translation of non-capped mRNAs in a eukaryotic cell-free system: acceleration of initiation rate in the course of polysome formation. Nucleic Acids Research, 2007, 35, 6547-6559.	14.5	28
12	Compilation of tRNA sequences and sequences of tRNA genes. Nucleic Acids Research, 2004, 33, D139-D140.	14.5	1,121
13	Native-like secondary structure of molten globules. BBA - Proteins and Proteomics, 2002, 1594, 168-177.	2.1	69
14	Structure and stability of recombinant protein depend on the extra N-terminal methionine residue: S6 permutein from direct and fusion expression systems. BBA - Proteins and Proteomics, 1999, 1432, 324-332.	2.1	7
15	Effect of Self-Association on the Structural Organization of Partially Folded Proteins: Inactivated Actin. Biophysical Journal, 1999, 77, 2788-2800.	0.5	45
16	Preparation of active tRNA gene transcripts devoid of 3'-extended products and dimers. Nucleic Acids Research, 1998, 26, 2500-2501.	14.5	20
17	Mg2+ binding and structural stability of mature and in vitro synthesized unmodified Escherichia coli tRNAPhe. Nucleic Acids Research, 1998, 26, 2723-2728.	14.5	44
18	Circularly permuted dihydrofolate reductase possesses all the properties of the molten globule state, but can resume functional tertiary structure by interaction with its ligands. Protein Science, 1996, 5, 1844-1851.	7.6	66

#	Article	IF	CITATIONS
19	The Major Protein of Messenger Ribonucleoprotein Particles in Somatic Cells Is a Member of the Y-box Binding Transcription Factor Family. Journal of Biological Chemistry, 1995, 270, 3186-3192.	3.4	163
20	Determination of Protein Tertiary Structure Class from Circular Dichroism Spectra. Analytical Biochemistry, 1994, 222, 176-184.	2.4	124