

## List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Postâ€transcriptional regulation is involved in the coldâ€active methanolâ€based methanogenic pathway of a psychrophilic methanogen. Environmental Microbiology, 2021, 23, 3773-3788.	3.8	6
2	aCPSF1 cooperates with terminator U-tract to dictate archaeal transcription termination efficacy. ELife, 2021, 10, .	6.0	12
3	Characterization of the Methanomicrobial Archaeal RNase Zs for Processing the CCA-Containing tRNA Precursors. Frontiers in Microbiology, 2020, 11, 1851.	3.5	2
4	The conserved ribonuclease aCPSF1 triggers genome-wide transcription termination of Archaea via a 3′-end cleavage mode. Nucleic Acids Research, 2020, 48, 9589-9605.	14.5	31
5	A newly identified duplex RNA unwinding activity of archaeal RNase J depends on processive exoribonucleolysis coupled steric occlusion by its structural archaeal loops. RNA Biology, 2020, 17, 1480-1491.	3.1	3
6	Comprehensive analysis of the pre-ribosomal RNA maturation pathway in a methanoarchaeon exposes the conserved circularization and linearization mode in archaea. RNA Biology, 2020, 17, 1427-1441.	3.1	15
7	The archaeal RNA chaperone TRAM0076 shapes the transcriptome and optimizes the growth of Methanococcus maripaludis. PLoS Genetics, 2019, 15, e1008328.	3.5	14
8	Genome-wide mRNA processing in methanogenic archaea reveals post-transcriptional regulation of ribosomal protein synthesis. Nucleic Acids Research, 2017, 45, 7285-7298.	14.5	35
9	Conserved TRAM Domain Functions as an Archaeal Cold Shock Protein via RNA Chaperone Activity. Frontiers in Microbiology, 2017, 8, 1597.	3.5	18
10	Global mapping transcriptional start sites revealed both transcriptional and post-transcriptional regulation of cold adaptation in the methanogenic archaeon Methanolobus psychrophilus. Scientific Reports, 2015, 5, 9209.	3.3	58