

# JÄdrzej M MaÅ,ecki

## List of Publications by Year in descending order

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18  
papers

766  
citations

643344

15  
h-index

939365

18  
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18  
all docs

18  
docs citations

18  
times ranked

1062  
citing authors

#	ARTICLE	IF	CITATIONS
1	Protein methylation in mitochondria. <i>Journal of Biological Chemistry</i> , 2022, 298, 101791.	1.6	18
2	The methyltransferase METTL9 mediates pervasive 1-methylhistidine modification in mammalian proteomes. <i>Nature Communications</i> , 2021, 12, 891.	5.8	54
3	Human METTL18 is a histidine-specific methyltransferase that targets RPL3 and affects ribosome biogenesis and function. <i>Nucleic Acids Research</i> , 2021, 49, 3185-3203.	6.5	34
4	The mutational landscape of cold agglutinin disease: <sc><i>CARD11</i></sc> and <sc><i>CXCR4</i></sc> mutations are correlated with lower hemoglobin levels. <i>American Journal of Hematology</i> , 2021, 96, E279-E283.	2.0	7
5	The human methyltransferase ZCCHC4 catalyses N6-methyladenosine modification of 28S ribosomal RNA. <i>Nucleic Acids Research</i> , 2020, 48, 830-846.	6.5	88
6	Human FAM173A is a mitochondrial lysine-specific methyltransferase that targets adenine nucleotide translocase and affects mitochondrial respiration. <i>Journal of Biological Chemistry</i> , 2019, 294, 11654-11664.	1.6	18
7	Lysine methylation by the mitochondrial methyltransferase FAM173B optimizes the function of mitochondrial ATP synthase. <i>Journal of Biological Chemistry</i> , 2019, 294, 1128-1141.	1.6	18
8	Regulation of eukaryotic elongation factor 1 alpha (eEF1A) by dynamic lysine methylation. <i>RNA Biology</i> , 2018, 15, 314-319.	1.5	37
9	Frequent somatic mutations of <i><sc>KMT</sc>2D</i> (<i><sc>MLL</sc>2</i>) and <i><sc>CARD</sc>11</i> genes in primary cold agglutinin disease. <i>British Journal of Haematology</i> , 2018, 183, 838-842.	1.2	53
10	The dual methyltransferase METTL13 targets N terminus and Lys55 of eEF1A and modulates codon-specific translation rates. <i>Nature Communications</i> , 2018, 9, 3411.	5.8	81
11	Identification of FAM173B as a protein methyltransferase promoting chronic pain. <i>PLoS Biology</i> , 2018, 16, e2003452.	2.6	22
12	The novel lysine specific methyltransferase METTL21B affects mRNA translation through inducible and dynamic methylation of Lys-165 in human eukaryotic elongation factor 1 alpha (eEF1A). <i>Nucleic Acids Research</i> , 2017, 45, gkx002.	6.5	64
13	Methylation of human eukaryotic elongation factor alpha (eEF1A) by a member of a novel protein lysine methyltransferase family modulates mRNA translation. <i>Nucleic Acids Research</i> , 2017, 45, 8239-8254.	6.5	44
14	Uncovering human METTL12 as a mitochondrial methyltransferase that modulates citrate synthase activity through metabolite-sensitive lysine methylation. <i>Journal of Biological Chemistry</i> , 2017, 292, 17950-17962.	1.6	27
15	Protein lysine methylation by seven-Î²-strand methyltransferases. <i>Biochemical Journal</i> , 2016, 473, 1995-2009.	1.7	92
16	The METTL20 Homologue from <i>Agrobacterium tumefaciens</i> Is a Dual Specificity Protein-lysine Methyltransferase That Targets Ribosomal Protein L7/L12 and the Î² Subunit of Electron Transfer Flavoprotein (ETFÎ²). <i>Journal of Biological Chemistry</i> , 2016, 291, 9581-9595.	1.6	14
17	Human METTL20 Is a Mitochondrial Lysine Methyltransferase That Targets the Î² Subunit of Electron Transfer Flavoprotein (ETFÎ²) and Modulates Its Activity. <i>Journal of Biological Chemistry</i> , 2015, 290, 423-434.	1.6	48
18	<i>Saccharomyces cerevisiae</i> Eukaryotic Elongation Factor 1A (eEF1A) Is Methylated at Lys-390 by a METTL21-Like Methyltransferase. <i>PLoS ONE</i> , 2015, 10, e0131426.	1.1	47