

# Xavier Rambout

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4859907/publications.pdf>

Version: 2024-02-01

11  
papers

1,412  
citations

1163117

8  
h-index

1281871

11  
g-index

11  
all docs

11  
docs citations

11  
times ranked

3430  
citing authors

#	ARTICLE	IF	CITATIONS
1	A Proteome-Scale Map of the Human Interactome Network. <i>Cell</i> , 2014, 159, 1212-1226.	28.9	1,199
2	Beyond Transcription: Roles of Transcription Factors in Pre-mRNA Splicing. <i>Chemical Reviews</i> , 2018, 118, 4339-4364.	47.7	50
3	The nuclear cap-binding complex as choreographer of gene transcription and pre-mRNA processing. <i>Genes and Development</i> , 2020, 34, 1113-1127.	5.9	41
4	The transcription factor ERG recruits CCR4â€“NOT to control mRNA decay and mitotic progression. <i>Nature Structural and Molecular Biology</i> , 2016, 23, 663-672.	8.2	35
5	The Inositol Phosphatase SHIP-1 Inhibits NOD2-Induced NF-Î±B Activation by Disturbing the Interaction of XIAP with RIP2. <i>PLoS ONE</i> , 2012, 7, e41005.	2.5	26
6	Transcriptional coactivator PGC-1Î± contains a novel CBP80-binding motif that orchestrates efficient target gene expression. <i>Genes and Development</i> , 2018, 32, 555-567.	5.9	18
7	Varicella-Zoster Virus ORF9p Binding to Cellular Adaptor Protein Complex 1 Is Important for Viral Infectivity. <i>Journal of Virology</i> , 2018, 92, .	3.4	13
8	ERG transcription factors have a splicing regulatory function involving RBFOX2 that is altered in the EWS-FLI1 oncogenic fusion. <i>Nucleic Acids Research</i> , 2021, 49, 5038-5056.	14.5	11
9	3â€²READS + RIP defines differential Staufen1 binding to alternative 3â€²UTR isoforms and reveals structures and sequence motifs influencing binding and polysome association. <i>Rna</i> , 2020, 26, 1621-1636.	3.5	8
10	NCBP3: A Multifaceted Adaptive Regulator of Gene Expression. <i>Trends in Biochemical Sciences</i> , 2021, 46, 87-96.	7.5	7
11	Transcriptional Coactivator PGC-1Î± Binding to Newly Synthesized RNA via CBP80: A Nexus for Co- and Posttranscriptional Gene Regulation. <i>Cold Spring Harbor Symposia on Quantitative Biology</i> , 2019, 84, 47-54.	1.1	4