

# Ivãjn D'Orso

## List of Publications by Year in descending order

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Version: 2024-02-01

26  
papers

1,591  
citations

471509

17  
h-index

552781

26  
g-index

30  
all docs

30  
docs citations

30  
times ranked

2862  
citing authors

#	ARTICLE	IF	CITATIONS
1	Decoding Human Genome Regulatory Features That Influence HIV-1 Proviral Expression and Fate Through an Integrated Genomics Approach. <i>Bioinformatics and Biology Insights</i> , 2022, 16, 117793222110723.	2.0	1
2	ADAP1 promotes latent HIV-1 reactivation by selectively tuning KRAS-ERK-AP-1 T cell signaling-transcriptional axis. <i>Nature Communications</i> , 2022, 13, 1109.	12.8	2
3	Cleavage and Polyadenylation Specificity Factor 6 Is Required for Efficient HIV-1 Latency Reversal. <i>MBio</i> , 2021, 12, e0109821.	4.1	2
4	Nascent RNA: Friend or foe of the chromatin bound?. <i>Molecular Cell</i> , 2021, 81, 2871-2872.	9.7	1
5	The ARF tumor suppressor targets PPM1G/PP2C $\beta$ to counteract NF- $\kappa$ B transcription tuning cell survival and the inflammatory response. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 32594-32605.	7.1	8
6	HIV-1 Proviral Transcription and Latency in the New Era. <i>Viruses</i> , 2020, 12, 555.	3.3	29
7	KAP1 Is a Chromatin Reader that Couples Steps of RNA Polymerase II Transcription to Sustain Oncogenic Programs. <i>Molecular Cell</i> , 2020, 78, 1133-1151.e14.	9.7	26
8	Transcriptional Circuit Fragility Influences HIV Proviral Fate. <i>Cell Reports</i> , 2019, 27, 154-171.e9.	6.4	24
9	CDK9: a signaling hub for transcriptional control. <i>Transcription</i> , 2019, 10, 57-75.	3.1	131
10	The HIV-1 Tat protein recruits a ubiquitin ligase to reorganize the 7SK snRNP for transcriptional activation. <i>ELife</i> , 2018, 7, .	6.0	29
11	Tandem Affinity Purification of Protein Complexes from Eukaryotic Cells. <i>Journal of Visualized Experiments</i> , 2017, . .	0.3	6
12	CIPHER: a flexible and extensive workflow platform for integrative next-generation sequencing data analysis and genomic regulatory element prediction. <i>BMC Bioinformatics</i> , 2017, 18, 363.	2.6	25
13	Genome-wide analysis of KAP1, the 7SK snRNP complex, and RNA polymerase II. <i>Genomics Data</i> , 2016, 7, 250-255.	1.3	12
14	Transcription elongation control by the 7SK snRNP complex: Releasing the pause. <i>Cell Cycle</i> , 2016, 15, 2115-2123.	2.6	34
15	7SKing on chromatin: Move globally, act locally. <i>RNA Biology</i> , 2016, 13, 545-553.	3.1	21
16	KAP1 Recruitment of the 7SK snRNP Complex to Promoters Enables Transcription Elongation by RNA Polymerase II. <i>Molecular Cell</i> , 2016, 61, 39-53.	9.7	109
17	HIV Tat controls RNA Polymerase II and the epigenetic landscape to transcriptionally reprogram target immune cells. <i>ELife</i> , 2015, 4, .	6.0	47
18	PPM1G Binds 7SK RNA and Hexim1 To Block P-TEFb Assembly into the 7SK snRNP and Sustain Transcription Elongation. <i>Molecular and Cellular Biology</i> , 2015, 35, 3810-3828.	2.3	40

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19	Transcription Factors Mediate the Enzymatic Disassembly of Promoter-Bound 7SK snRNP to Locally Recruit P-TEFb for Transcription Elongation. <i>Cell Reports</i> , 2013, 5, 1256-1268.	6.4	74
20	Transition Step during Assembly of HIV Tat:P-TEFb Transcription Complexes and Transfer to TAR RNA. <i>Molecular and Cellular Biology</i> , 2012, 32, 4780-4793.	2.3	46
21	Transcription control by long non-coding RNAs. <i>Transcription</i> , 2012, 3, 78-86.	3.1	23
22	Global landscape of HIV-human protein complexes. <i>Nature</i> , 2012, 481, 365-370.	27.8	651
23	Purification and characterization of HIV-human protein complexes. <i>Methods</i> , 2011, 53, 13-19.	3.8	56
24	RNA-mediated displacement of an inhibitory snRNP complex activates transcription elongation. <i>Nature Structural and Molecular Biology</i> , 2010, 17, 815-821.	8.2	115
25	HIV-1 Tat: Its Dependence on Host Factors is Crystal Clear. <i>Viruses</i> , 2010, 2, 2226-2234.	3.3	19
26	Tat acetylation modulates assembly of a viral-host RNA-protein transcription complex. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 3101-3106.	7.1	59