

# Florence Baudin

## List of Publications by Year in descending order

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45  
papers

4,659  
citations

201674

27  
h-index

233421

45  
g-index

51  
all docs

51  
docs citations

51  
times ranked

4854  
citing authors

#	ARTICLE	IF	CITATIONS
1	Probing the structure of RNAs in solution. Nucleic Acids Research, 1987, 15, 9109-9128.	14.5	751
2	The cap-snatching endonuclease of influenza virus polymerase resides in the PA subunit. Nature, 2009, 458, 914-918.	27.8	630
3	Functional Sites in the 5' Region of Human Immunodeficiency Virus Type 1 RNA Form Defined Structural Domains. Journal of Molecular Biology, 1993, 229, 382-397.	4.2	326
4	Structure and nuclear import function of the C-terminal domain of influenza virus polymerase PB2 subunit. Nature Structural and Molecular Biology, 2007, 14, 229-233.	8.2	275
5	Dimerization of human immunodeficiency virus (type 1) RNA: stimulation by cations and possible mechanism. Nucleic Acids Research, 1991, 19, 2349-2357.	14.5	202
6	Nuclear traffic of influenza virus proteins and ribonucleoprotein complexes. Virus Research, 2007, 124, 12-21.	2.2	197
7	Pervasive Protein Thermal Stability Variation during the Cell Cycle. Cell, 2018, 173, 1495-1507.e18.	28.9	183
8	Complex Interdependence Regulates Heterotypic Transcription Factor Distribution and Coordinates Cardiogenesis. Cell, 2016, 164, 999-1014.	28.9	179
9	Crystal structure of the M1 protein-binding domain of the influenza A virus nuclear export protein (NEP/NS2). EMBO Journal, 2003, 22, 4646-4655.	7.8	174
10	In Vitro Dissection of the Membrane and RNP Binding Activities of Influenza Virus M1 Protein. Virology, 2001, 281, 102-108.	2.4	141
11	Influenza Polymerase Can Adopt an Alternative Configuration Involving a Radical Repacking of PB2 Domains. Molecular Cell, 2016, 61, 125-137.	9.7	123
12	Architecture of CRM1/Exportin1 Suggests How Cooperativity Is Achieved during Formation of a Nuclear Export Complex. Molecular Cell, 2004, 16, 761-775.	9.7	119
13	Combined Results from Solution Studies on Intact Influenza Virus M1 Protein and from a New Crystal Form of Its N-Terminal Domain Show That M1 Is an Elongated Monomer. Virology, 2001, 279, 439-446.	2.4	112
14	Use of Lead(II) to Probe the Structure of Large RNA's. Conformation of the 3' Terminal Domain of E. coli 16S rRNA and its Involvement in Building the tRNA Binding Sites. Journal of Biomolecular Structure and Dynamics, 1989, 6, 971-984.	3.5	94
15	HMGB1 Protein Binds to Influenza Virus Nucleoprotein and Promotes Viral Replication. Journal of Virology, 2012, 86, 9122-9133.	3.4	94
16	Monomeric Nucleoprotein of Influenza A Virus. PLoS Pathogens, 2013, 9, e1003275.	4.7	89
17	Structural Basis of Lytic Cycle Activation by the Epstein-Barr Virus ZEBRA Protein. Molecular Cell, 2006, 21, 565-572.	9.7	82
18	Mutations in 5S DNA and 5S RNA have different effects on the binding of Xenopus transcription factor IIIA. Biochemistry, 1991, 30, 2495-2500.	2.5	68

#	ARTICLE	IF	CITATIONS
19	Chromatin-modifying Complex Component Nurf55/p55 Associates with Histones H3 and H4 and Polycomb Repressive Complex 2 Subunit Su(z)12 through Partially Overlapping Binding Sites. <i>Journal of Biological Chemistry</i> , 2011, 286, 23388-23396.	3.4	61
20	Structural basis for tRNA modification by Elp3 from <i>Dehalococcoides mccartyi</i> . <i>Nature Structural and Molecular Biology</i> , 2016, 23, 794-802.	8.2	59
21	Cryo-EM structures of human RNA polymerase III in its unbound and transcribing states. <i>Nature Structural and Molecular Biology</i> , 2021, 28, 210-219.	8.2	59
22	Structure of the RNA inside the vesicular stomatitis virus nucleocapsid. <i>Rna</i> , 2000, 6, 270-281.	3.5	55
23	Interaction of influenza virus proteins with nucleosomes. <i>Virology</i> , 2005, 332, 329-336.	2.4	54
24	Chemical modification of nucleotide bases and mRNA editing depend on hexamer or nucleoprotein phase in Sendai virus nucleocapsids. <i>Rna</i> , 2002, 8, 1056-1067.	3.5	51
25	Higher-order structure of domain III in <i>Escherichia coli</i> 16S ribosomal RNA, 30S subunit and 70S ribosome. <i>Biochimie</i> , 1987, 69, 1081-1096.	2.6	50
26	A difference in the importance of bulged nucleotides and their parent base pairs in the binding of transcription factor IIIA to <i>Xenopus</i> 5S RNA and 5S RNA genes. <i>Nucleic Acids Research</i> , 1989, 17, 2043-2056.	14.5	41
27	Involvement of "hinge" nucleotides of <i>Xenopus laevis</i> 5 S rRNA in the RNA structural organization and in the binding of transcription factor TFIIIA. <i>Journal of Molecular Biology</i> , 1991, 218, 69-81.	4.2	41
28	Structural basis for RNA polymerase III transcription repression by Maf1. <i>Nature Structural and Molecular Biology</i> , 2020, 27, 229-232.	8.2	37
29	Molecular insight into RNA polymerase I promoter recognition and promoter melting. <i>Nature Communications</i> , 2019, 10, 5543.	12.8	33
30	Cryo-EM structures of human RNA polymerase I. <i>Nature Structural and Molecular Biology</i> , 2021, 28, 997-1008.	8.2	28
31	Human importin alpha and RNA do not compete for binding to influenza A virus nucleoprotein. <i>Virology</i> , 2011, 409, 84-90.	2.4	27
32	Structure of a knockout mutant of influenza virus M1 protein that has altered activities in membrane binding, oligomerisation and binding to NEP (NS2). <i>Virus Research</i> , 2004, 99, 115-119.	2.2	24
33	Functional determinants of the Epstein-Barr virus protease. <i>Journal of Molecular Biology</i> , 2001, 311, 217-228.	4.2	23
34	RNA polymerase III-specific general transcription factor IIIC contains a heterodimer resembling TFIIIF Rap30/Rap74. <i>Nucleic Acids Research</i> , 2013, 41, 9183-9196.	14.5	23
35	Structural snapshots of La Crosse virus polymerase reveal the mechanisms underlying Peribunyaviridae replication and transcription. <i>Nature Communications</i> , 2022, 13, 902.	12.8	23
36	Structural studies on site-directed mutants of domain 3 of <i>Xenopus laevis</i> oocyte 5 S ribosomal RNA. <i>Journal of Molecular Biology</i> , 1991, 219, 243-255.	4.2	22

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37	Crosslinking of transcription factor TFIIIA to ribosomal 5S RNA from <i>X. laevis</i> by trans-diamminedichloroplatinum (II). <i>Nucleic Acids Research</i> , 1989, 17, 10035-10046.	14.5	20
38	Structure-based design of NS2 mutants for attenuated influenza A virus vaccines. <i>Virus Research</i> , 2011, 155, 240-248.	2.2	20
39	Structure of the TFIIIC subcomplex $\beta_1\beta_2$ provides insights into RNA polymerase III pre-initiation complex formation. <i>Nature Communications</i> , 2020, 11, 4905.	12.8	16
40	Human initiation factor eIF3 subunit b interacts with HCV IRES RNA through its N-terminal RNA recognition motif. <i>FEBS Letters</i> , 2009, 583, 70-74.	2.8	14
41	Role of influenza virus M1 protein in the viral budding process. <i>International Congress Series</i> , 2001, 1219, 397-404.	0.2	11
42	Vault RNA1 riboregulates the autophagic function of p62 by binding to lysine 7 and arginine 21, both of which are critical for p62 oligomerization. <i>Rna</i> , 2022, 28, 742-755.	3.5	9
43	Mechanism of RNA polymerase I selection by transcription factor UAF. <i>Science Advances</i> , 2022, 8, eabn5725.	10.3	9
44	Bacterial Expression, Purification, and Crystallization of Tyrosine Phosphorylated STAT Proteins. <i>Methods in Molecular Biology</i> , 2013, 967, 301-317.	0.9	3
45	Structure of the RNA inside influenza virus RNPs. <i>International Congress Series</i> , 2001, 1219, 451-456.	0.2	1