

# Edouard Bertrand

## List of Publications by Year in descending order

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133  
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15,342  
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19690

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150  
docs citations

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times ranked

14995  
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#	ARTICLE	IF	CITATIONS
1	The interaction between RPAP3 and TRBP reveals a possible involvement of the HSP90/R2TP chaperone complex in the regulation of miRNA activity. <i>Nucleic Acids Research</i> , 2022, 50, 2172-2189.	6.5	4
2	FISH-quant v2: a scalable and modular tool for smFISH image analysis. <i>Rna</i> , 2022, 28, 786-795.	1.6	45
3	GOLT1B Activation in Hepatitis C Virus-Infected Hepatocytes Links ER Trafficking and Viral Replication. <i>Pathogens</i> , 2022, 11, 46.	1.2	0
4	RNA transport from transcription to localized translation: a single molecule perspective. <i>RNA Biology</i> , 2021, 18, 1-17.	1.5	14
5	The box C/D snoRNP assembly factor Bcd1 interacts with the histone chaperone Rtt106 and controls its transcription dependent activity. <i>Nature Communications</i> , 2021, 12, 1859.	5.8	3
6	A choreography of centrosomal mRNAs reveals a conserved localization mechanism involving active polysome transport. <i>Nature Communications</i> , 2021, 12, 1352.	5.8	52
7	Exon junction complex dependent mRNA localization is linked to centrosome organization during ciliogenesis. <i>Nature Communications</i> , 2021, 12, 1351.	5.8	35
8	Non-canonical argonaute loading of extracellular vesicle-derived exogenous single-stranded miRNA in recipient cells. <i>Journal of Cell Science</i> , 2021, 134, .	1.2	14
9	Live-cell imaging reveals the spatiotemporal organization of endogenous RNA polymerase II phosphorylation at a single gene. <i>Nature Communications</i> , 2021, 12, 3158.	5.8	36
10	TSSC4 is a component of U5 snRNP that promotes tri-snRNP formation. <i>Nature Communications</i> , 2021, 12, 3646.	5.8	14
11	Stochastic pausing at latent HIV-1 promoters generates transcriptional bursting. <i>Nature Communications</i> , 2021, 12, 4503.	5.8	38
12	Quantitative imaging of transcription in living <i>Drosophila</i> embryos reveals the impact of core promoter motifs on promoter state dynamics. <i>Nature Communications</i> , 2021, 12, 4504.	5.8	39
13	The HSP90/R2TP assembly chaperone promotes cell proliferation in the intestinal epithelium. <i>Nature Communications</i> , 2021, 12, 4810.	5.8	7
14	Live single-cell transcriptional dynamics via RNA labelling during the phosphate response in plants. <i>Nature Plants</i> , 2021, 7, 1050-1064.	4.7	27
15	The kinesin KIF1C transports APC-dependent mRNAs to cell protrusions. <i>Rna</i> , 2021, 27, 1528-1544.	1.6	23
16	NOPCHAP1 is a PAQosome cofactor that helps loading NOP58 on RUVBL1/2 during box C/D snoRNP biogenesis. <i>Nucleic Acids Research</i> , 2021, 49, 1094-1113.	6.5	14
17	First Responders Shape a Prompt and Sharp NF- $\kappa$ B-Mediated Transcriptional Response to TNF- $\alpha$ . <i>IScience</i> , 2020, 23, 101529.	1.9	11
18	A Dual Protein-mRNA Localization Screen Reveals Compartmentalized Translation and Widespread Co-translational RNA Targeting. <i>Developmental Cell</i> , 2020, 54, 773-791.e5.	3.1	88

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19	New Generations of MS2 Variants and MCP Fusions to Detect Single mRNAs in Living Eukaryotic Cells. <i>Methods in Molecular Biology</i> , 2020, 2166, 121-144.	0.4	21
20	A Deep Learning Approach To Identify MRNA Localization Patterns. , 2019, , .		3
21	Live cell imaging reveals 3' UTR dependent mRNA sorting to synapses. <i>Nature Communications</i> , 2019, 10, 3178.	5.8	35
22	Meg3 Non-coding RNA Expression Controls Imprinting by Preventing Transcriptional Upregulation in cis. <i>Cell Reports</i> , 2018, 23, 337-348.	2.9	54
23	The PAQosome, an R2TP-Based Chaperone for Quaternary Structure Formation. <i>Trends in Biochemical Sciences</i> , 2018, 43, 4-9.	3.7	67
24	A computational framework to study sub-cellular RNA localization. <i>Nature Communications</i> , 2018, 9, 4584.	5.8	47
25	The RPAP3-Cterminal domain identifies R2TP-like quaternary chaperones. <i>Nature Communications</i> , 2018, 9, 2093.	5.8	59
26	A Growing Toolbox to Image Gene Expression in Single Cells: Sensitive Approaches for Demanding Challenges. <i>Molecular Cell</i> , 2018, 71, 468-480.	4.5	149
27	Deep Structural Analysis of RPAP3 and PIH1D1, Two Components of the HSP90 Co-chaperone R2TP Complex. <i>Structure</i> , 2018, 26, 1196-1209.e8.	1.6	36
28	The Role of Supercoiling in the Motor Activity of RNA Polymerases. <i>Methods in Molecular Biology</i> , 2018, 1805, 215-232.	0.4	14
29	The splicing factor SRSF3 is functionally connected to the nuclear RNA exosome for intronless mRNA decay. <i>Scientific Reports</i> , 2018, 8, 12901.	1.6	23
30	Assembly of the U5 snRNP component PRPF8 is controlled by the HSP90/R2TP chaperones. <i>Journal of Cell Biology</i> , 2017, 216, 1579-1596.	2.3	65
31	Mutually Exclusive CBC-Containing Complexes Contribute to RNA Fate. <i>Cell Reports</i> , 2017, 18, 2635-2650.	2.9	73
32	ARS2 is a general suppressor of pervasive transcription. <i>Nucleic Acids Research</i> , 2017, 45, 10229-10241.	6.5	53
33	P-Body Purification Reveals the Condensation of Repressed mRNA Regulons. <i>Molecular Cell</i> , 2017, 68, 144-157.e5.	4.5	581
34	Assembly and trafficking of box C/D and H/ACA snoRNPs. <i>RNA Biology</i> , 2017, 14, 680-692.	1.5	144
35	The in vivo dynamics of TCERG1, a factor that couples transcriptional elongation with splicing. <i>Rna</i> , 2016, 22, 571-582.	1.6	13
36	Visualization of single endogenous polysomes reveals the dynamics of translation in live human cells. <i>Journal of Cell Biology</i> , 2016, 214, 769-781.	2.3	158

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37	Imaging HIV-1 RNA dimerization in cells by multicolor super-resolution and fluctuation microscopies. <i>Nucleic Acids Research</i> , 2016, 44, 7922-7934.	6.5	35
38	smiFISH and FISH-quant – a flexible single RNA detection approach with super-resolution capability. <i>Nucleic Acids Research</i> , 2016, 44, e165-e165.	6.5	312
39	A single-molecule view of transcription reveals convoys of RNA polymerases and multi-scale bursting. <i>Nature Communications</i> , 2016, 7, 12248.	5.8	233
40	NUFIP and the HSP90/R2TP chaperone bind the SMN complex and facilitate assembly of U4-specific proteins. <i>Nucleic Acids Research</i> , 2015, 43, 8973-8989.	6.5	49
41	Long lasting control of viral rebound with a new drug ABX464 targeting Rev – mediated viral RNA biogenesis. <i>Retrovirology</i> , 2015, 12, 30.	0.9	78
42	SnoRNPs, ZNHIT proteins and the R2TP pathway. <i>Oncotarget</i> , 2015, 6, 41399-41400.	0.8	13
43	Hypermethylated-capped selenoprotein mRNAs in mammals. <i>Nucleic Acids Research</i> , 2014, 42, 8663-8677.	6.5	45
44	Characterization of the interaction between protein Snu13p/15.5K and the Rsa1p/NUFIP factor and demonstration of its functional importance for snoRNP assembly. <i>Nucleic Acids Research</i> , 2014, 42, 2015-2036.	6.5	34
45	Proteomic and 3D structure analyses highlight the C/D box snoRNP assembly mechanism and its control. <i>Journal of Cell Biology</i> , 2014, 207, 463-480.	2.3	57
46	Stable assembly of HIV-1 export complexes occurs cotranscriptionally. <i>Rna</i> , 2014, 20, 1-8.	1.6	33
47	MLN51 triggers P-body disassembly and formation of a new type of RNA granules. <i>Journal of Cell Science</i> , 2014, 127, 4692-701.	1.2	18
48	Drosophila Spag Is the Homolog of RNA Polymerase II-associated Protein 3 (RPAP3) and Recruits the Heat Shock Proteins 70 and 90 (Hsp70 and Hsp90) during the Assembly of Cellular Machineries. <i>Journal of Biological Chemistry</i> , 2014, 289, 6236-6247.	1.6	41
49	Genome-wide identification of mRNAs associated with the protein SMN whose depletion decreases their axonal localization. <i>Rna</i> , 2013, 19, 1755-1766.	1.6	52
50	The human cap-binding complex is functionally connected to the nuclear RNA exosome. <i>Nature Structural and Molecular Biology</i> , 2013, 20, 1367-1376.	3.6	199
51	FISH-quant: automatic counting of transcripts in 3D FISH images. <i>Nature Methods</i> , 2013, 10, 277-278.	9.0	338
52	CBC – ARS2 stimulates 3 – end maturation of multiple RNA families and favors cap-proximal processing. <i>Nature Structural and Molecular Biology</i> , 2013, 20, 1358-1366.	3.6	143
53	Retroviral GAG proteins recruit AGO2 on viral RNAs without affecting RNA accumulation and translation. <i>Nucleic Acids Research</i> , 2012, 40, 775-786.	6.5	32
54	CRM1 plays a nuclear role in transporting snoRNPs to nucleoli in higher eukaryotes. <i>Nucleus</i> , 2012, 3, 132-137.	0.6	25

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55	HSP90 and the R2TP co-chaperone complex: Building multi-protein machineries essential for cell growth and gene expression. <i>RNA Biology</i> , 2012, 9, 148-154.	1.5	57
56	Microprocessor dynamics and interactions at endogenous imprinted C19MC microRNA genes. <i>Journal of Cell Science</i> , 2012, 125, 2709-20.	1.2	18
57	Perispeckles are major assembly sites for the exon junction core complex. <i>Molecular Biology of the Cell</i> , 2012, 23, 1765-1782.	0.9	56
58	Nuclear Retention Prevents Premature Cytoplasmic Appearance of mRNA. <i>Molecular Cell</i> , 2012, 48, 145-152.	4.5	28
59	Real-time imaging of the HIV-1 transcription cycle in single living cells. <i>Methods</i> , 2011, 53, 62-67.	1.9	27
60	CRM1 controls the composition of nucleoplasmic pre-snoRNA complexes to licence them for nucleolar transport. <i>EMBO Journal</i> , 2011, 30, 2205-2218.	3.5	36
61	Real-time imaging of cotranscriptional splicing reveals a kinetic model that reduces noise: implications for alternative splicing regulation. <i>Journal of Cell Biology</i> , 2011, 193, 819-829.	2.3	104
62	Establishment of a Protein Frequency Library and Its Application in the Reliable Identification of Specific Protein Interaction Partners. <i>Molecular and Cellular Proteomics</i> , 2010, 9, 861-879.	2.5	63
63	Splicing-independent recruitment of U1 snRNP to a transcription unit in living cells. <i>Journal of Cell Science</i> , 2010, 123, 2085-2093.	1.2	59
64	RiboSys, a high-resolution, quantitative approach to measure the in vivo kinetics of pre-mRNA splicing and 3' end processing in <i>Saccharomyces cerevisiae</i> . <i>Rna</i> , 2010, 16, 2570-2580.	1.6	48
65	A Proteomic Screen for Nucleolar SUMO Targets Shows SUMOylation Modulates the Function of Nop5/Nop58. <i>Molecular Cell</i> , 2010, 39, 618-631.	4.5	72
66	HSP90 and Its R2TP/Prefoldin-like Cochaperone Are Involved in the Cytoplasmic Assembly of RNA Polymerase II. <i>Molecular Cell</i> , 2010, 39, 912-924.	4.5	246
67	Crosstalk between mRNA 3' End Processing and Transcription Initiation. <i>Molecular Cell</i> , 2010, 40, 410-422.	4.5	153
68	Processivity and Coupling in Messenger RNA Transcription. <i>PLoS ONE</i> , 2010, 5, e8845.	1.1	3
69	Endosomal Trafficking of HIV-1 Gag and Genomic RNAs Regulates Viral Egress. <i>Journal of Biological Chemistry</i> , 2009, 284, 19727-19743.	1.6	57
70	Assembly of an Export-Competent mRNP Is Needed for Efficient Release of the 3' End Processing Complex after Polyadenylation. <i>Molecular and Cellular Biology</i> , 2009, 29, 5327-5338.	1.1	58
71	DNA Damage Regulates Alternative Splicing through Inhibition of RNA Polymerase II Elongation. <i>Cell</i> , 2009, 137, 708-720.	13.5	267
72	DNA Damage Regulates Alternative Splicing through Inhibition of RNA Polymerase II Elongation. <i>Cell</i> , 2009, 139, 211.	13.5	1

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73	Mutations in a small region of the exportin Crm1p disrupt the daughter cell-specific nuclear localization of the transcription factor Ace2p in <i>Saccharomyces cerevisiae</i> . <i>Biology of the Cell</i> , 2008, 100, 343-354.	0.7	22
74	Dendrites of Mammalian Neurons Contain Specialized P-Body-Like Structures That Respond to Neuronal Activation. <i>Journal of Neuroscience</i> , 2008, 28, 13793-13804.	1.7	153
75	Characterization of a Short Isoform of Human Tgs1 Hypermethylase Associating with Small Nucleolar Ribonucleoprotein Core Proteins and Produced by Limited Proteolytic Processing. <i>Journal of Biological Chemistry</i> , 2008, 283, 2060-2069.	1.6	39
76	A Novel Role for PA28 <sup>γ</sup> -Proteasome in Nuclear Speckle Organization and SR Protein Trafficking. <i>Molecular Biology of the Cell</i> , 2008, 19, 1706-1716.	0.9	63
77	The Hsp90 chaperone controls the biogenesis of L7Ae RNPs through conserved machinery. <i>Journal of Cell Biology</i> , 2008, 180, 579-595.	2.3	196
78	Translationally Repressed mRNA Transiently Cycles through Stress Granules during Stress. <i>Molecular Biology of the Cell</i> , 2008, 19, 4469-4479.	0.9	197
79	Assembly and Traffic of Small Nuclear RNPs. <i>Progress in Molecular and Subcellular Biology</i> , 2008, 35, 79-97.	0.9	13
80	Imaging of Single mRNAs in the Cytoplasm of Living Cells. <i>Progress in Molecular and Subcellular Biology</i> , 2008, 35, 135-150.	0.9	16
81	Inhibition of nonsense-mediated mRNA decay (NMD) by a new chemical molecule reveals the dynamic of NMD factors in P-bodies. <i>Journal of Cell Biology</i> , 2007, 178, 1145-1160.	2.3	147
82	Bsr, a Nuclear-retained RNA with Monoallelic Expression. <i>Molecular Biology of the Cell</i> , 2007, 18, 2817-2827.	0.9	29
83	The transcriptional cycle of HIV-1 in real-time and live cells. <i>Journal of Cell Biology</i> , 2007, 179, 291-304.	2.3	174
84	The exon-junction-complex-component metastatic lymph node 51 functions in stress-granule assembly. <i>Journal of Cell Science</i> , 2007, 120, 2774-2784.	1.2	69
85	A Dynamic Scaffold of Pre-snoRNP Factors Facilitates Human Box C/D snoRNP Assembly. <i>Molecular and Cellular Biology</i> , 2007, 27, 6782-6793.	1.1	83
86	The Clathrin Adaptor Complex AP-1 Binds HIV-1 and MLV Gag and Facilitates Their Budding. <i>Molecular Biology of the Cell</i> , 2007, 18, 3193-3203.	0.9	89
87	A real-time view of the TAR:Tat:P-TEFb complex at HIV-1 transcription sites. <i>Retrovirology</i> , 2007, 4, 36.	0.9	48
88	Suv39H1 and HP1 <sup>γ</sup> are responsible for chromatin-mediated HIV-1 transcriptional silencing and post-integration latency. <i>EMBO Journal</i> , 2007, 26, 424-435.	3.5	281
89	The transcriptional cycle of HIV-1 in real-time and live cells. <i>Journal of Experimental Medicine</i> , 2007, 204, i25-i25.	4.2	0
90	Photobleaching of YFP does not produce a CFP-like species that affects FRET measurements. <i>Nature Methods</i> , 2006, 3, 492-493.	9.0	9

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91	The exonuclease ISG20 mainly localizes in the nucleolus and the Cajal (Coiled) bodies and is associated with nuclear SMN protein-containing complexes. <i>Journal of Cellular Biochemistry</i> , 2006, 98, 1320-1333.	1.2	30
92	Depletion of SMN by RNA interference in HeLa cells induces defects in Cajal body formation. <i>Nucleic Acids Research</i> , 2006, 34, 2925-2932.	6.5	59
93	Cell Cycle-dependent Recruitment of Telomerase RNA and Cajal Bodies to Human Telomeres. <i>Molecular Biology of the Cell</i> , 2006, 17, 944-954.	0.9	168
94	Photoconversion of YFP into a CFP-like species during acceptor photobleaching FRET experiments. <i>Nature Methods</i> , 2005, 2, 801-801.	9.0	108
95	Tsg101 and Alix Interact with Murine Leukemia Virus Gag and Cooperate with Nedd4 Ubiquitin Ligases during Budding. <i>Journal of Biological Chemistry</i> , 2005, 280, 27004-27012.	1.6	86
96	ADAR2-mediated editing of RNA substrates in the nucleolus is inhibited by C/D small nucleolar RNAs. <i>Journal of Cell Biology</i> , 2005, 169, 745-753.	2.3	223
97	Inhibition of Translational Initiation by Let-7 MicroRNA in Human Cells. <i>Science</i> , 2005, 309, 1573-1576.	6.0	1,247
98	The Packaging Signal of MLV is an Integrated Module that Mediates Intracellular Transport of Genomic RNAs. <i>Journal of Molecular Biology</i> , 2005, 354, 330-339.	2.0	28
99	A cotranscriptional model for 3'-end processing of the <i>Saccharomyces cerevisiae</i> pre-ribosomal RNA precursor. <i>Rna</i> , 2004, 10, 1572-1585.	1.6	23
100	Human Box H/ACA Pseudouridylation Guide RNA Machinery. <i>Molecular and Cellular Biology</i> , 2004, 24, 5797-5807.	1.1	180
101	Human telomerase RNA and box H/ACA scaRNAs share a common Cajal body-specific localization signal. <i>Journal of Cell Biology</i> , 2004, 164, 647-652.	2.3	248
102	Nuclear localization properties of a conserved protuberance in the Sm core complex. <i>Experimental Cell Research</i> , 2004, 299, 199-208.	1.2	27
103	PHAX and CRM1 Are Required Sequentially to Transport U3 snoRNA to Nucleoli. <i>Molecular Cell</i> , 2004, 16, 777-787.	4.5	157
104	From Silencing to Gene Expression. <i>Cell</i> , 2004, 116, 683-698.	13.5	658
105	Modification of Sm small nuclear RNAs occurs in the nucleoplasmic Cajal body following import from the cytoplasm. <i>EMBO Journal</i> , 2003, 22, 1878-1888.	3.5	213
106	A common sequence motif determines the Cajal body-specific localization of box H/ACA scaRNAs. <i>EMBO Journal</i> , 2003, 22, 4283-4293.	3.5	181
107	Single mRNA Molecules Demonstrate Probabilistic Movement in Living Mammalian Cells. <i>Current Biology</i> , 2003, 13, 161-167.	1.8	529
108	Interaction between the small nuclear RNA cap hypermethylase and the spinal muscular atrophy protein, survival of motor neuron. <i>EMBO Reports</i> , 2003, 4, 616-622.	2.0	96

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109	Retroviral Genomic RNAs Are Transported to the Plasma Membrane by Endosomal Vesicles. <i>Developmental Cell</i> , 2003, 5, 161-174.	3.1	138
110	Human let-7 stem-loop precursors harbor features of RNase III cleavage products. <i>Nucleic Acids Research</i> , 2003, 31, 6593-6597.	6.5	131
111	The RasGAP-associated endoribonuclease G3BP assembles stress granules. <i>Journal of Cell Biology</i> , 2003, 160, 823-831.	2.3	790
112	Exportin-5 Mediates Nuclear Export of Minihelix-containing RNAs. <i>Journal of Biological Chemistry</i> , 2003, 278, 5505-5508.	1.6	139
113	A Cajal body-specific pseudouridylation guide RNA is composed of two box H/ACA snoRNA-like domains. <i>Nucleic Acids Research</i> , 2002, 30, 4643-4649.	6.5	102
114	An active precursor in assembly of yeast nuclear ribonuclease P. <i>Rna</i> , 2002, 8, 1348-1360.	1.6	30
115	Hypermethylation of the Cap Structure of Both Yeast snRNAs and snoRNAs Requires a Conserved Methyltransferase that Is Localized to the Nucleolus. <i>Molecular Cell</i> , 2002, 9, 891-901.	4.5	222
116	Intra-nuclear RNA trafficking: insights from live cell imaging. <i>Biochimie</i> , 2002, 84, 805-813.	1.3	21
117	Mammalian and yeast U3 snoRNPs are matured in specific and related nuclear compartments. <i>EMBO Journal</i> , 2002, 21, 2736-2745.	3.5	167
118	Cajal body-specific small nuclear RNAs: a novel class of 2'-O-methylation and pseudouridylation guide RNAs. <i>EMBO Journal</i> , 2002, 21, 2746-2756.	3.5	417
119	Box C/D small nucleolar RNA trafficking involves small nucleolar RNP proteins, nucleolar factors and a novel nuclear domain. <i>EMBO Journal</i> , 2001, 20, 5480-5490.	3.5	98
120	A Well-Connected and Conserved Nucleoplasmic Helicase Is Required for Production of Box C/D and H/ACA snoRNAs and Localization of snoRNP Proteins. <i>Molecular and Cellular Biology</i> , 2001, 21, 7731-7746.	1.1	104
121	Terminal Minihelix, a Novel RNA Motif That Directs Polymerase III Transcripts to the Cell Cytoplasm. <i>Journal of Biological Chemistry</i> , 2001, 276, 25910-25918.	1.6	36
122	[33] Sensitive and high-resolution detection of RNA in situ. <i>Methods in Enzymology</i> , 2000, 318, 493-506.	0.4	51
123	mRNA localization signals can enhance the intracellular effectiveness of hammerhead ribozymes. <i>Rna</i> , 1999, 5, 1200-1209.	1.6	35
124	Localization of ASH1 mRNA Particles in Living Yeast. <i>Molecular Cell</i> , 1998, 2, 437-445.	4.5	1,475
125	3' End Modification of the Adenoviral VA1 Gene Affects Its Expression in Human Cells: Consequences for the Design of Chimeric VA1 RNA Ribozymes. <i>Oligonucleotides</i> , 1998, 8, 379-390.	4.4	4
126	Monitoring Retroviral RNA Dimerization In Vivo via Hammerhead Ribozyme Cleavage. <i>Journal of Virology</i> , 1998, 72, 8349-8353.	1.5	20



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127	Detection of Ribozyme Cleavage Products Using Reverse Ligation-Mediated PCR (RL-PCR). , 1997, 74, 311-324.		7
128	In Vivo Footprinting of the Interaction of Proteins with DNA and RNA. Advances in Molecular and Cell Biology, 1997, , 73-109.	0.1	2
129	In VivoFootprinting of the Interaction of Proteins with DNA and RNA. Methods, 1997, 11, 151-163.	1.9	36
130	Can hammerhead ribozymes be efficient tools to inactivate gene function?. Nucleic Acids Research, 1994, 22, 293-300.	6.5	141
131	Designing and Testing of Ribozymes as Therapeutic Agents. Methods, 1993, 5, 19-27.	1.9	4
132	A highly sensitive method for mapping the 5â€™ termini of mRNAs. Nucleic Acids Research, 1993, 21, 1683-1684.	6.5	98
133	A Localization Screen Reveals Translation Factories and Widespread Co-Translational Protein Targeting. SSRN Electronic Journal, 0, , .	0.4	1