Ronald R Breaker

List of Publications by Year
in descending order

218	33,840	93	178
papers	citations	h-index	g-index
		2970	3822

232	232	232	13238
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	DIMPL: a bioinformatics pipeline for the discovery of structured noncoding RNA motifs in bacteria. Bioinformatics, 2022, 38, 533-535.	1.8	3
2	The Biochemical Landscape of Riboswitch Ligands. Biochemistry, 2022, 61, 137-149.	1.2	47
3	Variants of the guanine riboswitch class exhibit altered ligand specificities for xanthine, guanine, or 2â€²-deoxyguanosine. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .	3.3	13
4	The case of the missing allosteric ribozymes. Nature Chemical Biology, 2021, 17, 375-382.	3.9	11
5	Structure of a bacterial OapB protein with its OLE RNA target gives insights into the architecture of the OLE ribonucleoprotein complex. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	1
6	Comprehensive discovery of novel structured noncoding RNAs in 26 bacterial genomes. RNA Biology, 2021, 18, 2417-2432.	1.5	13
7	Witnessing the structural evolution of an RNA enzyme. ELife, 2021, 10, .	2.8	14
8	A second riboswitch class for the enzyme cofactor NAD⁺. Rna, 2021, 27, 99-105.	1.6	23
9	Natural circularly permuted group II introns in bacteria produce RNA circles. IScience, 2021, 24, 103431.	1.9	7
10	Imaginary Ribozymes. ACS Chemical Biology, 2020, 15, 2020-2030.	1.6	19
11	Biochemical Validation of a Fourth Guanidine Riboswitch Class in Bacteria. Biochemistry, 2020, 59, 4654-4662.	1.2	32
12	Bacillus halodurans OapB forms a high-affinity complex with the P13 region of the noncoding RNA OLE. Journal of Biological Chemistry, 2020, 295, 9326-9334.	1.6	6
13	A rare bacterial RNA motif is implicated in the regulation of thepurFgene whose encoded enzyme synthesizes phosphoribosylamine. Rna, 2020, 26, 1838-1846.	1.6	5
14	A bacterial riboswitch class senses xanthine and uric acid to regulate genes associated with purine oxidation. Rna, 2020, 26, 960-968.	1.6	24
15	Former orphan riboswitches reveal unexplored areas of bacterial metabolism, signaling, and gene control processes. Rna, 2020, 26, 675-693.	1.6	34
16	Variant Bacterial Riboswitches Associated with Nucleotide Hydrolase Genes Sense Nucleoside Diphosphates. Biochemistry, 2019, 58, 401-410.	1.2	34
17	Biochemical validation of a second class of tetrahydrofolate riboswitches in bacteria. Rna, 2019, 25, 1091-1097.	1.6	17
18	Employing a ZTP Riboswitch to Detect Bacterial Folate Biosynthesis Inhibitors in a Small Molecule High-Throughput Screen. ACS Chemical Biology, 2019, 14, 2841-2850.	1.6	13

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19	Disruption of the OLE ribonucleoprotein complex causes magnesium toxicity in Bacillus halodurans. Molecular Microbiology, 2019, 112, 1552-1563.	1.2	6
20	Evidence that the <i>nadA</i> motif is a bacterial riboswitch for the ubiquitous enzyme cofactor NAD⁺. Rna, 2019, 25, 1616-1627.	1.6	30
21	Genome-wide discovery of structured noncoding RNAs in bacteria. BMC Microbiology, 2019, 19, 66.	1.3	40
22	Rare variants of the FMN riboswitch class in <i>Clostridium difficile</i> and other bacteria exhibit altered ligand specificity. Rna, 2019, 25, 23-34.	1.6	18
23	Genomeâ€wide Discovery of Rare Riboswitches in Bacteria. FASEB Journal, 2019, 33, 778.8.	0.2	1
24	A bacterial riboswitch class for the thiamin precursor HMP-PP employs a terminator-embedded aptamer. ELife, 2019, 8, .	2.8	33
25	SAM-VI RNAs selectively bind <i>S</i>-adenosylmethionine and exhibit similarities to SAM-III riboswitches. RNA Biology, 2018, 15, 371-378.	1.5	42
26	Challenges of ligand identification for the second wave of orphan riboswitch candidates. RNA Biology, 2018, 15, 377-390.	1.5	33
27	Large Noncoding RNAs in Bacteria. , 2018, , 515-526.		3
28	Riboswitches and Translation Control. Cold Spring Harbor Perspectives in Biology, 2018, 10, a032797.	2.3	147
29	Riboswitches for the alarmone ppGpp expand the collection of RNA-based signaling systems. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 6052-6057.	3.3	94
30	A glutamine riboswitch is a key element for the regulation of glutamine synthetase in cyanobacteria. Nucleic Acids Research, 2018, 46, 10082-10094.	6.5	51
31	Tandem riboswitches form a natural Boolean logic gate to control purine metabolism in bacteria. ELife, 2018, 7, .	2.8	59
32	Large Noncoding RNAs in Bacteria. Microbiology Spectrum, 2018, 6, .	1.2	39
33	A second RNA-binding protein is essential for ethanol tolerance provided by the bacterial OLE ribonucleoprotein complex. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E6319-E6328.	3.3	9
34	High Throughput Validation of Orphan Riboswitch Candidates. FASEB Journal, 2018, 32, lb18.	0.2	0
35	PROSPECTS FOR RIBOZYME DISCOVERY AND ANALYSIS. , 2018, , .		0
36	The Biology of Free Guanidine As Revealed by Riboswitches. Biochemistry, 2017, 56, 345-347.	1.2	31

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37	Search for 5â€²-leader regulatory RNA structures based on gene annotation aided by the RiboGap database. Methods, 2017, 117, 3-13.	1.9	8
38	Bioinformatic analysis of riboswitch structures uncovers variant classes with altered ligand specificity. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E2077-E2085.	3.3	75
39	Mechanistic Debris Generated by Twister Ribozymes. ACS Chemical Biology, 2017, 12, 886-891.	1.6	25
40	Riboswitch diversity and distribution. Rna, 2017, 23, 995-1011.	1.6	374
41	The lost language of the RNA World. Science Signaling, 2017, 10, .	1.6	95
42	Biochemical Validation of a Second Guanidine Riboswitch Class in Bacteria. Biochemistry, 2017, 56, 352-358.	1.2	87
43	Biochemical Validation of a Third Guanidine Riboswitch Class in Bacteria. Biochemistry, 2017, 56, 359-363.	1.2	70
44	Metabolism of Free Guanidine in Bacteria Is Regulated by a Widespread Riboswitch Class. Molecular Cell, 2017, 65, 220-230.	4.5	129
45	Detection of 224 candidate structured RNAs by comparative analysis of specific subsets of intergenic regions. Nucleic Acids Research, 2017, 45, 10811-10823.	6.5	116
46	Numerous small hammerhead ribozyme variants associated with Penelope-like retrotransposons cleave RNA as dimers. RNA Biology, 2017, 14, 1499-1507.	1.5	17
47	Identification of 15 candidate structured noncoding RNA motifs in fungi by comparative genomics. BMC Genomics, 2017, 18, 785.	1.2	16
48	Singlet glycine riboswitches bind ligand as well as tandem riboswitches. Rna, 2016, 22, 1728-1738.	1.6	19
49	The <i>yjdF</i> riboswitch candidate regulates gene expression by binding diverse azaaromatic compounds. Rna, 2016, 22, 530-541.	1.6	33
50	Biochemical analysis of hatchet self-cleaving ribozymes. Rna, 2015, 21, 1845-1851.	1.6	36
51	Biochemical analysis of pistol self-cleaving ribozymes. Rna, 2015, 21, 1852-1858.	1.6	59
52	An Ancient Riboswitch Class in Bacteria Regulates Purine Biosynthesis and One-Carbon Metabolism. Molecular Cell, 2015, 57, 317-328.	4.5	102
53	Novel Riboswitch-Binding Flavin Analog That Protects Mice against Clostridium difficile Infection without Inhibiting Cecal Flora. Antimicrobial Agents and Chemotherapy, 2015, 59, 5736-5746.	1.4	75
54	New classes of self-cleaving ribozymes revealed by comparative genomics analysis. Nature Chemical Biology, 2015, 11, 606-610.	3.9	174

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55	Small Molecule Fluoride Toxicity Agonists. Chemistry and Biology, 2015, 22, 527-534.	6.2	21
56	Bacterial Riboswitches Cooperatively Bind Ni 2+ or Co 2+ Ions and Control Expression of Heavy Metal Transporters. Molecular Cell, 2015, 57, 1088-1098.	4.5	147
57	Control of bacterial exoelectrogenesis by c-AMP-GMP. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 5389-5394.	3.3	98
58	A widespread self-cleaving ribozyme class is revealed by bioinformatics. Nature Chemical Biology, 2014, 10, 56-60.	3.9	217
59	Gramicidin D enhances the antibacterial activity of fluoride. Bioorganic and Medicinal Chemistry Letters, 2014, 24, 2969-2971.	1.0	10
60	The Expanding View of RNA and DNA Function. Chemistry and Biology, 2014, 21, 1059-1065.	6.2	87
61	Structural, Functional, and Taxonomic Diversity of Three PreQ1 Riboswitch Classes. Chemistry and Biology, 2014, 21, 880-889.	6.2	78
62	Riboswitches That Sense Cyclic Di-GMP. , 2014, , 215-229.		4
63	In Vitro Selection of Allosteric Ribozymes that Sense the Bacterial Second Messenger c-di-GMP. Methods in Molecular Biology, 2014, 1111, 209-220.	0.4	9
64	Riboswitches in eubacteria sense the second messenger c-di-AMP. Nature Chemical Biology, 2013, 9, 834-839.	3.9	247
65	Integron attI1 Sites, Not Riboswitches, Associate with Antibiotic Resistance Genes. Cell, 2013, 153, 1417-1418.	13.5	19
66	Small, Highly Active DNAs That Hydrolyze DNA. Journal of the American Chemical Society, 2013, 135, 9121-9129.	6.6	134
67	Eukaryotic TPP riboswitch regulation of alternative splicing involving long-distance base pairing. Nucleic Acids Research, 2013, 41, 3022-3031.	6.5	96
68	Eukaryotic resistance to fluoride toxicity mediated by a widespread family of fluoride export proteins. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 19018-19023.	3.3	108
69	Production of single-stranded DNAs by self-cleavage of rolling-circle amplification products. BioTechniques, 2013, 54, 337-343.	0.8	23
70	OLE RNA protects extremophilic bacteria from alcohol toxicity. Nucleic Acids Research, 2012, 40, 6898-6907.	6.5	23
71	Ancient, giant riboswitches at atomic resolution. Nature Structural and Molecular Biology, 2012, 19, 1208-1209.	3.6	4
72	Riboswitches and the RNA World. Cold Spring Harbor Perspectives in Biology, 2012, 4, a003566-a003566.	2.3	529

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73	Engineered Allosteric Ribozymes That Sense the Bacterial Second Messenger Cyclic Diguanosyl 5â€²-Monophosphate. Analytical Chemistry, 2012, 84, 4935-4941.	3.2	45
74	Widespread Genetic Switches and Toxicity Resistance Proteins for Fluoride. Science, 2012, 335, 233-235.	6.0	356
75	A highly specialized flavin mononucleotide riboswitch responds differently to similar ligands and confers roseoflavin resistance to Streptomyces davawensis. Nucleic Acids Research, 2012, 40, 8662-8673.	6.5	75
76	Identification of Ligand Analogues that Control c-di-GMP Riboswitches. ACS Chemical Biology, 2012, 7, 1436-1443.	1.6	41
77	Fluoride enhances the activity of fungicides that destabilize cell membranes. Bioorganic and Medicinal Chemistry Letters, 2012, 22, 3317-3322.	1.0	31
78	Mechanism and Distribution of glmS Ribozymes. Methods in Molecular Biology, 2012, 848, 113-129.	0.4	22
79	New Insight on the Response of Bacteria to Fluoride. Caries Research, 2012, 46, 78-81.	0.9	47
80	Prospects for Riboswitch Discovery and Analysis. Molecular Cell, 2011, 43, 867-879.	4.5	445
81	Association of OLE RNA with bacterial membranes via an RNAâ€"protein interaction. Molecular Microbiology, 2011, 79, 21-34.	1.2	28
82	Improved genetic transformation methods for the model alkaliphile Bacillus halodurans C-125. Letters in Applied Microbiology, 2011, 52, 430-432.	1.0	11
83	R2R - software to speed the depiction of aesthetic consensus RNA secondary structures. BMC Bioinformatics, 2011, 12, 3.	1.2	226
84	An expanded collection and refined consensus model of <i>glmS</i> ribozymes. Rna, 2011, 17, 728-736.	1.6	50
85	Mechanism for gene control by a natural allosteric group I ribozyme. Rna, 2011, 17, 1967-1972.	1.6	55
86	New families of human regulatory RNA structures identified by comparative analysis of vertebrate genomes. Genome Research, 2011, 21, 1929-1943.	2.4	100
87	Challenges of ligand identification for riboswitch candidates. RNA Biology, 2011, 8, 5-10.	1.5	61
88	Bacterial aptamers that selectively bind glutamine. RNA Biology, 2011, 8, 82-89.	1.5	85
89	Identification of Hammerhead Ribozymes in All Domains of Life Reveals Novel Structural Variations. PLoS Computational Biology, 2011, 7, e1002031.	1.5	124
90	A Eubacterial Riboswitch Class That Senses the Coenzyme Tetrahydrofolate. Chemistry and Biology, 2010, 17, 681-685.	6.2	86

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91	Evidence for Widespread Gene Control Function by the <i>ydaO</i> Riboswitch Candidate. Journal of Bacteriology, 2010, 192, 3983-3989.	1.0	60
92	RNA Switches Out in the Cold. Molecular Cell, 2010, 37, 1-2.	4.5	21
93	An Allosteric Self-Splicing Ribozyme Triggered by a Bacterial Second Messenger. Science, 2010, 329, 845-848.	6.0	309
94	Comparative genomics reveals 104 candidate structured RNAs from bacteria, archaea, and their metagenomes. Genome Biology, 2010, 11, R31.	13.9	348
95	RNA Second Messengers and Riboswitches: Relics from the RNA World?. Microbe Magazine, 2010, 5, 13-20.	0.4	7
96	The large, noncoding OLE RNA is associated with membrane biochemistry. FASEB Journal, 2010, 24, 493.2.	0.2	0
97	Roseoflavin is a natural antibacterial compound that binds to FMN riboswitches and regulates gene expression. RNA Biology, 2009, 6, 187-194.	1.5	202
98	A variant riboswitch aptamer class for <i>S</i>-adenosylmethionine common in marine bacteria. Rna, 2009, 15, 2046-2056.	1.6	96
99	Riboswitches: from ancient gene-control systems to modern drug targets. Future Microbiology, 2009, 4, 771-773.	1.0	60
100	Identification of candidate structured RNAs in the marine organism 'Candidatus Pelagibacter ubique'. BMC Genomics, 2009, 10, 268.	1.2	56
101	Engineering ligand-responsive gene-control elements: lessons learned from natural riboswitches. Gene Therapy, 2009, 16, 1189-1201.	2.3	68
102	Exceptional structured noncoding RNAs revealed by bacterial metagenome analysis. Nature, 2009, 462, 656-659.	13.7	102
103	A plant 5S ribosomal RNA mimic regulates alternative splicing of transcription factor IIIA pre-mRNAs. Nature Structural and Molecular Biology, 2009, 16, 541-549.	3.6	43
104	Structural basis of ligand binding by a c-di-GMP riboswitch. Nature Structural and Molecular Biology, 2009, 16, 1218-1223.	3.6	257
105	Unique glycineâ€activated riboswitch linked to glycineâ€"serine auxotrophy in SAR11. Environmental Microbiology, 2009, 11, 230-238.	1.8	90
106	The Structural and Functional Diversity of Metabolite-Binding Riboswitches. Annual Review of Biochemistry, 2009, 78, 305-334.	5.0	506
107	Design and Antimicrobial Action of Purine Analogues That Bind Guanine Riboswitches. ACS Chemical Biology, 2009, 4, 915-927.	1.6	113
108	FINDING NON-CODING RNAs THROUGH GENOME-SCALE CLUSTERING. Journal of Bioinformatics and Computational Biology, 2009, 07, 373-388.	0.3	28

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109	In Vitro Selection of glmS Ribozymes. Methods in Molecular Biology, 2009, 540, 349-364.	0.4	1
110	A plant 5S rRNA mimic regulates alternative splicing of transcription factor IIIA preâ€mRNAs. FASEB Journal, 2009, 23, 665.4.	0.2	0
111	A widespread riboswitch candidate that controls bacterial genes involved in molybdenum cofactor and tungsten cofactor metabolism. Molecular Microbiology, 2008, 68, 918-932.	1.2	142
112	In-Line Probing Analysis of Riboswitches. Methods in Molecular Biology, 2008, 419, 53-67.	0.4	289
113	Purine sensing by riboswitches. Biology of the Cell, 2008, 100, 1-11.	0.7	87
114	Riboswitches in Eubacteria Sense the Second Messenger Cyclic Di-GMP. Science, 2008, 321, 411-413.	6.0	654
115	Riboswitches that Sense S-adenosylhomocysteine and Activate Genes Involved in Coenzyme Recycling. Molecular Cell, 2008, 29, 691-702.	4.5	153
116	Riboswitches that sense <i>S</i>-adenosylmethionine and <i>S</i>-adenosylhomocysteineThis paper is one of a selection of papers published in this Special Issue, entitled CSBMCB â€" Systems and Chemical Biology, and has undergone the Journal's usual peer review process.. Biochemistry and Cell Biology, 2008, 86, 157-168.	0.9	105
117	Complex Riboswitches. Science, 2008, 319, 1795-1797.	6.0	105
118	The aptamer core of SAM-IV riboswitches mimics the ligand-binding site of SAM-I riboswitches. Rna, 2008, 14, 822-828.	1.6	103
119	Confirmation of a second natural preQ₁ aptamer class in Streptococcaceae bacteria. Rna, 2008, 14, 685-695.	1.6	102
120	In Vitro Selection and Characterization of Cellulose-Binding RNA Aptamers Using isothermal Amplification. Nucleosides, Nucleotides and Nucleic Acids, 2008, 27, 949-966.	0.4	18
121	Riboswitches as new antibiotics targets. FASEB Journal, 2008, 22, 264.3.	0.2	0
122	Ligand binding and gene control characteristics of tandem riboswitches in Bacillus anthracis. Rna, 2007, 13, 573-582.	1.6	110
123	A Computational Pipeline for High- Throughput Discovery of cis-Regulatory Noncoding RNA in Prokaryotes. PLoS Computational Biology, 2007, 3, e126.	1.5	77
124	Identification of 22 candidate structured RNAs in bacteria using the CMfinder comparative genomics pipeline. Nucleic Acids Research, 2007, 35, 4809-4819.	6.5	292
125	Guanine riboswitch variants from <i>Mesoplasma florum</i> selectively recognize 2â€²-deoxyguanosine. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 16092-16097.	3.3	129
126	In vitro selection and characterization of cellulose-binding DNA aptamers. Nucleic Acids Research, 2007, 35, 6378-6388.	6.5	46

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127	Engineering high-speed allosteric hammerhead ribozymes. Biological Chemistry, 2007, 388, 779-786.	1.2	50
128	The distributions, mechanisms, and structures of metabolite-binding riboswitches. Genome Biology, 2007, 8, R239.	13.9	414
129	Riboswitch Control of Gene Expression in Plants by Splicing and Alternative 3â€² End Processing of mRNAs. Plant Cell, 2007, 19, 3437-3450.	3.1	281
130	Importance of the Debye Screening Length on Nanowire Field Effect Transistor Sensors. Nano Letters, 2007, 7, 3405-3409.	4.5	716
131	Antibacterial lysine analogs that target lysine riboswitches. , 2007, 3, 44-49.		205
132	A riboswitch selective for the queuosine precursor preQ1 contains an unusually small aptamer domain. Nature Structural and Molecular Biology, 2007, 14, 308-317.	3.6	224
133	Control of alternative RNA splicing and gene expression by eukaryotic riboswitches. Nature, 2007, 447, 497-500.	13.7	377
134	Tandem Riboswitch Architectures Exhibit Complex Gene Control Functions. Science, 2006, 314, 300-304.	6.0	232
135	Riboswitches: Natural Metabolite-binding RNAs Controlling Gene Expression. , 2006, , 191-207.		0
136	Riboswitches as antibacterial drug targets. Nature Biotechnology, 2006, 24, 1558-1564.	9.4	419
137	Structural basis for gene regulation by a thiamine pyrophosphate-sensing riboswitch. Nature, 2006, 441, 1167-1171.	13.7	404
138	Molecular-Recognition Characteristics of SAM-Binding Riboswitches. Angewandte Chemie - International Edition, 2006, 45, 964-968.	7.2	51
139	Characteristics of Ligand Recognition by aglmS Self-Cleaving Ribozyme. Angewandte Chemie - International Edition, 2006, 45, 6689-6693.	7.2	58
140	Development and Application of a High-Throughput Assay for glmS Riboswitch Activators. RNA Biology, 2006, 3, 77-81.	1.5	71
141	Examination of the structural and functional versatility of glmS ribozymes by using in vitro selection. Nucleic Acids Research, 2006, 34, 4968-4975.	6.5	21
142	Characteristics of the glmS ribozyme suggest only structural roles for divalent metal ions. Rna, 2006, 12, 607-619.	1.6	102
143	Identification of a large noncoding RNA in extremophilic eubacteria. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 19490-19495.	3.3	34
144	Riboswitches as Genetic Control Elements. , 2006, , 89-106.		0

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145	Genetic control by riboswitches and ribozymes. FASEB Journal, 2006, 20, A455.	0.2	0
146	Riboswitches: Regulators of modern and ancient metabolism. Biochemist, 2006, 28, 11-15.	0.2	1
147	Gene expression control: Harnessing RNA switches. Gene Therapy, 2005, 12, 725-726.	2.3	6
148	Thiamine Pyrophosphate Riboswitches Are Targets for the Antimicrobial Compound Pyrithiamine. Chemistry and Biology, 2005, 12, 1325-1335.	6.2	237
149	Computational design and experimental validation of oligonucleotide-sensing allosteric ribozymes. Nature Biotechnology, 2005, 23, 1424-1433.	9.4	199
150	Riboswitches as versatile gene control elements. Current Opinion in Structural Biology, 2005, 15, 342-348.	2.6	503
151	Engineered allosteric ribozymes that respond to specific divalent metal ions. Nucleic Acids Research, 2005, 33, 622-631.	6.5	59
152	REGULATION OF BACTERIAL GENE EXPRESSION BY RIBOSWITCHES. Annual Review of Microbiology, 2005, 59, 487-517.	2.9	687
153	6S RNA is a widespread regulator of eubacterial RNA polymerase that resembles an open promoter. Rna, 2005, 11, 774-784.	1.6	210
154	The Kinetics of Ligand Binding by an Adenine-Sensing Riboswitch. Biochemistry, 2005, 44, 13404-13414.	1.2	264
155	The Speed of RNA Transcription and Metabolite Binding Kinetics Operate an FMN Riboswitch. Molecular Cell, 2005, 18, 49-60.	4.5	430
156	Evidence for a second class of S-adenosylmethionine riboswitches and other regulatory RNA motifs in alpha-proteobacteria. Genome Biology, 2005, 6, R70.	13.9	213
157	New RNA motifs suggest an expanded scope for riboswitches in bacterial genetic control. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 6421-6426.	3.3	432
158	Selection In Vitro<I> </I>of Allosteric Ribozymes. , 2004, 252, 145-164.		19
159	Coenzyme B12 riboswitches are widespread genetic control elements in prokaryotes. Nucleic Acids Research, 2004, 32, 143-150.	6.5	292
160	Gene regulation by riboswitches. Nature Reviews Molecular Cell Biology, 2004, 5, 451-463.	16.1	799
161	Adenine riboswitches and gene activation by disruption of a transcription terminator. Nature Structural and Molecular Biology, 2004, 11, 29-35.	3.6	471
162	Control of gene expression by a natural metabolite-responsive ribozyme. Nature, 2004, 428, 281-286.	13.7	847

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163	Natural and engineered nucleic acids as tools to explore biology. Nature, 2004, 432, 838-845.	13.7	336
164	Structural Basis for Discriminative Regulation of Gene Expression by Adenine- and Guanine-Sensing mRNAs. Chemistry and Biology, 2004, 11, 1729-1741.	6.2	505
165	Ligating DNA with DNA. Journal of the American Chemical Society, 2004, 126, 3454-3460.	6.6	100
166	A Glycine-Dependent Riboswitch That Uses Cooperative Binding to Control Gene Expression. Science, 2004, 306, 275-279.	6.0	491
167	Genetic Control by Metabolite-Binding Riboswitches. ChemInform, 2003, 34, no.	0.1	0
168	Genetic Control by Metabolite-Binding Riboswitches. ChemBioChem, 2003, 4, 1024-1032.	1.3	254
169	An mRNA structure that controls gene expression by binding S-adenosylmethionine. Nature Structural and Molecular Biology, 2003, 10, 701-707.	3.6	406
170	Riboswitches Control Fundamental Biochemical Pathways in Bacillus subtilis and Other Bacteria. Cell, 2003, 113, 577-586.	13.5	665
171	An mRNA structure in bacteria that controls gene expression by binding lysine. Genes and Development, 2003, 17, 2688-2697.	2.7	303
172	A common speed limit for RNA-cleaving ribozymes and deoxyribozymes. Rna, 2003, 9, 949-957.	1.6	119
173	Substrate specificity and reaction kinetics of an X-motif ribozyme. Rna, 2003, 9, 688-697.	1.6	17
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176	An mRNA structure that controls gene expression by binding FMN. Proceedings of the National Academy of Sciences of the United States of America, 2002, 99, 15908-15913.	3.3	599
177	Genetic Control by a Metabolite Binding mRNA. Chemistry and Biology, 2002, 9, 1043-1049.	6.2	686
178	Thiamine derivatives bind messenger RNAs directly to regulate bacterial gene expression. Nature, 2002, 419, 952-956.	13.7	1,075
179	Engineered allosteric ribozymes as biosensor components. Current Opinion in Biotechnology, 2002, 13, 31-39.	3.3	270
180	In Vitro Selection of Kinase and Ligase Deoxyribozymes. Methods, 2001, 23, 179-190.	1.9	18

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181	Generating new ligand-binding RNAs by affinity maturation and disintegration of allosteric ribozymes. Rna, 2001, 7, 524-536.	1.6	71
182	Characterization of a DNA-Cleaving deoxyribozyme. Bioorganic and Medicinal Chemistry, 2001, 9, 2589-2600.	1.4	108
183	Immobilized RNA switches for the analysis of complex chemical and biological mixtures. Nature Biotechnology, 2001, 19, 336-341.	9.4	214
184	Selection for Catalytic Function with Nucleic Acids. Current Protocols in Nucleic Acid Chemistry, 2000, 00, Unit 9.4.	0.5	1
185	Allosteric nucleic acid catalysts. Current Opinion in Structural Biology, 2000, 10, 318-325.	2.6	139
186	Structural diversity of self-cleaving ribozymes. Proceedings of the National Academy of Sciences of the United States of America, 2000, 97, 5784-5789.	3.3	93
187	Altering molecular recognition of RNA aptamers by allosteric selection. Journal of Molecular Biology, 2000, 298, 623-632.	2.0	119
188	MOLECULAR BIOLOGY: Making Catalytic DNAs. Science, 2000, 290, 2095-2096.	6.0	123
189	Capping DNA with DNAâ€. Biochemistry, 2000, 39, 3106-3114.	1.2	131
190	Molecular Recognition of cAMP by an RNA Aptamerâ€. Biochemistry, 2000, 39, 8983-8992.	1.2	66
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