

# Gene W Yeo

## List of Publications by Citations

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230  
papers

21,950  
citations

78  
h-index

146  
g-index

277  
ext. papers

28,237  
ext. citations

15.4  
avg, IF

6.95  
L-index

#	Paper	IF	Citations
230	Maximum entropy modeling of short sequence motifs with applications to RNA splicing signals. <i>Journal of Computational Biology</i> , <b>2004</b> , 11, 377-94	1.7	1253
229	A model for neural development and treatment of Rett syndrome using human induced pluripotent stem cells. <i>Cell</i> , <b>2010</b> , 143, 527-39	56.2	997
228	Long pre-mRNA depletion and RNA missplicing contribute to neuronal vulnerability from loss of TDP-43. <i>Nature Neuroscience</i> , <b>2011</b> , 14, 459-68	25.5	827
227	Divergent transcription from active promoters. <i>Science</i> , <b>2008</b> , 322, 1849-51	33.3	695
226	Robust transcriptome-wide discovery of RNA-binding protein binding sites with enhanced CLIP (eCLIP). <i>Nature Methods</i> , <b>2016</b> , 13, 508-14	21.6	634
225	L1 retrotransposition in human neural progenitor cells. <i>Nature</i> , <b>2009</b> , 460, 1127-31	50.4	613
224	Systematic identification and analysis of exonic splicing silencers. <i>Cell</i> , <b>2004</b> , 119, 831-45	56.2	522
223	Divergent roles of ALS-linked proteins FUS/TLS and TDP-43 intersect in processing long pre-mRNAs. <i>Nature Neuroscience</i> , <b>2012</b> , 15, 1488-97	25.5	483
222	L1 retrotransposition in neurons is modulated by MeCP2. <i>Nature</i> , <b>2010</b> , 468, 443-6	50.4	479
221	Wnt-mediated activation of NeuroD1 and retro-elements during adult neurogenesis. <i>Nature Neuroscience</i> , <b>2009</b> , 12, 1097-105	25.5	474
220	An RNA code for the FOX2 splicing regulator revealed by mapping RNA-protein interactions in stem cells. <i>Nature Structural and Molecular Biology</i> , <b>2009</b> , 16, 130-7	17.6	447
219	Targeted degradation of sense and antisense C9orf72 RNA foci as therapy for ALS and frontotemporal degeneration. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2013</b> , 110, E4530-9	11.5	420
218	Context-Dependent and Disease-Specific Diversity in Protein Interactions within Stress Granules. <i>Cell</i> , <b>2018</b> , 172, 590-604.e13	56.2	411
217	Variation in alternative splicing across human tissues. <i>Genome Biology</i> , <b>2004</b> , 5, R74	18.3	378
216	Expanded encyclopaedias of DNA elements in the human and mouse genomes. <i>Nature</i> , <b>2020</b> , 583, 699-710	30.4	360
215	Programmable RNA Tracking in Live Cells with CRISPR/Cas9. <i>Cell</i> , <b>2016</b> , 165, 488-96	56.2	358
214	Noncoding RNAs in the mammalian central nervous system. <i>Annual Review of Neuroscience</i> , <b>2006</b> , 29, 77-103	17	346

213	Genome-wide analysis of PTB-RNA interactions reveals a strategy used by the general splicing repressor to modulate exon inclusion or skipping. <i>Molecular Cell</i> , <b>2009</b> , 36, 996-1006	17.6	338
212	Integrative genome-wide analysis reveals cooperative regulation of alternative splicing by hnRNP proteins. <i>Cell Reports</i> , <b>2012</b> , 1, 167-78	10.6	322
211	ALS-linked TDP-43 mutations produce aberrant RNA splicing and adult-onset motor neuron disease without aggregation or loss of nuclear TDP-43. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2013</b> , 110, E736-45	11.5	284
210	Complex Oscillatory Waves Emerging from Cortical Organoids Model Early Human Brain Network Development. <i>Cell Stem Cell</i> , <b>2019</b> , 25, 558-569.e7	18	266
209	RNA sequence analysis defines DicerB role in mouse embryonic stem cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2007</b> , 104, 18097-102	11.5	261
208	Comprehensive discovery of endogenous Argonaute binding sites in <i>Caenorhabditis elegans</i> . <i>Nature Structural and Molecular Biology</i> , <b>2010</b> , 17, 173-9	17.6	252
207	Transcriptional signature and memory retention of human-induced pluripotent stem cells. <i>PLoS ONE</i> , <b>2009</b> , 4, e7076	3.7	247
206	The Clothes Make the mRNA: Past and Present Trends in mRNP Fashion. <i>Annual Review of Biochemistry</i> , <b>2015</b> , 84, 325-54	29.1	235
205	Identification and analysis of alternative splicing events conserved in human and mouse. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2005</b> , 102, 2850-5	11.5	227
204	The EJC factor eIF4AIII modulates synaptic strength and neuronal protein expression. <i>Cell</i> , <b>2007</b> , 130, 179-91	56.2	223
203	Plant-derived flavanol (-)epicatechin enhances angiogenesis and retention of spatial memory in mice. <i>Journal of Neuroscience</i> , <b>2007</b> , 27, 5869-78	6.6	217
202	Sequence, Structure, and Context Preferences of Human RNA Binding Proteins. <i>Molecular Cell</i> , <b>2018</b> , 70, 854-867.e9	17.6	212
201	RESCUE-ESE identifies candidate exonic splicing enhancers in vertebrate exons. <i>Nucleic Acids Research</i> , <b>2004</b> , 32, W187-90	20.1	212
200	Genome-wide analysis reveals SR protein cooperation and competition in regulated splicing. <i>Molecular Cell</i> , <b>2013</b> , 50, 223-35	17.6	208
199	Rbfox proteins regulate alternative mRNA splicing through evolutionarily conserved RNA bridges. <i>Nature Structural and Molecular Biology</i> , <b>2013</b> , 20, 1434-42	17.6	206
198	Circadian oscillations of protein-coding and regulatory RNAs in a highly dynamic mammalian liver epigenome. <i>Cell Metabolism</i> , <b>2012</b> , 16, 833-45	24.6	199
197	Pairing beyond the Seed Supports MicroRNA Targeting Specificity. <i>Molecular Cell</i> , <b>2016</b> , 64, 320-333	17.6	199
196	A large-scale binding and functional map of human RNA-binding proteins. <i>Nature</i> , <b>2020</b> , 583, 711-719	50.4	198

195	LIN28 binds messenger RNAs at GGAGA motifs and regulates splicing factor abundance. <i>Molecular Cell</i> , <b>2012</b> , 48, 195-206	17.6	193
194	Variation in sequence and organization of splicing regulatory elements in vertebrate genes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2004</b> , 101, 15700-5	11.5	184
193	The PIWI proteins SMEDWI-2 and SMEDWI-3 are required for stem cell function and piRNA expression in planarians. <i>Rna</i> , <b>2008</b> , 14, 1174-86	5.8	174
192	Differential L1 regulation in pluripotent stem cells of humans and apes. <i>Nature</i> , <b>2013</b> , 503, 525-529	50.4	170
191	ALS-causative mutations in FUS/TLS confer gain and loss of function by altered association with SMN and U1-snRNP. <i>Nature Communications</i> , <b>2015</b> , 6, 6171	17.4	162
190	Elimination of Toxic Microsatellite Repeat Expansion RNA by RNA-Targeting Cas9. <i>Cell</i> , <b>2017</b> , 170, 899-913.e10	155	155
189	Identification of novel long noncoding RNAs underlying vertebrate cardiovascular development. <i>Circulation</i> , <b>2015</b> , 131, 1278-1290	16.7	146
188	The Ro60 autoantigen binds endogenous retroelements and regulates inflammatory gene expression. <i>Science</i> , <b>2015</b> , 350, 455-9	33.3	142
187	Early specification of CD8+ T lymphocyte fates during adaptive immunity revealed by single-cell gene-expression analyses. <i>Nature Immunology</i> , <b>2014</b> , 15, 365-372	19.1	137
186	Emergence and rapid transmission of SARS-CoV-2 B.1.1.7 in the United States. <i>Cell</i> , <b>2021</b> , 184, 2587-2594.e7	132	132
185	A combinatorial code for splicing silencing: UAGG and GGGG motifs. <i>PLoS Biology</i> , <b>2005</b> , 3, e158	9.7	129
184	How RNA-Binding Proteins Interact with RNA: Molecules and Mechanisms. <i>Molecular Cell</i> , <b>2020</b> , 78, 9-29	17.6	126
183	Misregulated RNA processing in amyotrophic lateral sclerosis. <i>Brain Research</i> , <b>2012</b> , 1462, 3-15	3.7	124
182	Th17 Lymphocytes Induce Neuronal Cell Death in a Human iPSC-Based Model of Parkinson's Disease. <i>Cell Stem Cell</i> , <b>2018</b> , 23, 123-131.e6	18	119
181	High-resolution profiling and analysis of viral and host small RNAs during human cytomegalovirus infection. <i>Journal of Virology</i> , <b>2012</b> , 86, 226-35	6.6	113
180	Genetic mutations in RNA-binding proteins and their roles in ALS. <i>Human Genetics</i> , <b>2017</b> , 136, 1193-1214	6.3	111
179	Nonclassical splicing mutations in the coding and noncoding regions of the ATM Gene: maximum entropy estimates of splice junction strengths. <i>Human Mutation</i> , <b>2004</b> , 23, 67-76	4.7	111
178	Discovery and analysis of evolutionarily conserved intronic splicing regulatory elements. <i>PLoS Genetics</i> , <b>2007</b> , 3, e85	6	110

177	Early transcriptional and epigenetic regulation of CD8 T cell differentiation revealed by single-cell RNA sequencing. <i>Nature Immunology</i> , <b>2017</b> , 18, 422-432	19.1	109
176	LIN-28 co-transcriptionally binds primary let-7 to regulate miRNA maturation in <i>Caenorhabditis elegans</i> . <i>Nature Structural and Molecular Biology</i> , <b>2011</b> , 18, 302-8	17.6	108
175	NEAT1 scaffolds RNA-binding proteins and the Microprocessor to globally enhance pri-miRNA processing. <i>Nature Structural and Molecular Biology</i> , <b>2017</b> , 24, 816-824	17.6	106
174	Short poly(A) tails are a conserved feature of highly expressed genes. <i>Nature Structural and Molecular Biology</i> , <b>2017</b> , 24, 1057-1063	17.6	106
173	SONAR Discovers RNA-Binding Proteins from Analysis of Large-Scale Protein-Protein Interactomes. <i>Molecular Cell</i> , <b>2016</b> , 64, 282-293	17.6	105
172	Alternative splicing events identified in human embryonic stem cells and neural progenitors. <i>PLoS Computational Biology</i> , <b>2007</b> , 3, 1951-67	5	103
171	Pervasive Chromatin-RNA Binding Protein Interactions Enable RNA-Based Regulation of Transcription. <i>Cell</i> , <b>2019</b> , 178, 107-121.e18	56.2	101
170	Single-Cell Alternative Splicing Analysis with Expedition Reveals Splicing Dynamics during Neuron Differentiation. <i>Molecular Cell</i> , <b>2017</b> , 67, 148-161.e5	17.6	99
169	Protein-RNA Networks Regulated by Normal and ALS-Associated Mutant HNRNPA2B1 in the Nervous System. <i>Neuron</i> , <b>2016</b> , 92, 780-795	13.9	94
168	Distinct and shared functions of ALS-associated proteins TDP-43, FUS and TAF15 revealed by multisystem analyses. <i>Nature Communications</i> , <b>2016</b> , 7, 12143	17.4	94
167	iPSCORE: A Resource of 222 iPSC Lines Enabling Functional Characterization of Genetic Variation across a Variety of Cell Types. <i>Stem Cell Reports</i> , <b>2017</b> , 8, 1086-1100	8	93
166	Advances and challenges in the detection of transcriptome-wide protein-RNA interactions. <i>Wiley Interdisciplinary Reviews RNA</i> , <b>2018</b> , 9, e1436	9.3	93
165	Pseudotemporal Ordering of Single Cells Reveals Metabolic Control of Postnatal Cell Proliferation. <i>Cell Metabolism</i> , <b>2017</b> , 25, 1160-1175.e11	24.6	92
164	Disruption of RNA Metabolism in Neurological Diseases and Emerging Therapeutic Interventions. <i>Neuron</i> , <b>2019</b> , 102, 294-320	13.9	91
163	Predicting the functional states of human iPSC-derived neurons with single-cell RNA-seq and electrophysiology. <i>Molecular Psychiatry</i> , <b>2016</b> , 21, 1573-1588	15.1	90
162	Small-Molecule Modulation of TDP-43 Recruitment to Stress Granules Prevents Persistent TDP-43 Accumulation in ALS/FTD. <i>Neuron</i> , <b>2019</b> , 103, 802-819.e11	13.9	88
161	Resources for the Comprehensive Discovery of Functional RNA Elements. <i>Molecular Cell</i> , <b>2016</b> , 61, 903-1017	17.6	87
160	Determination of tag density required for digital transcriptome analysis: application to an androgen-sensitive prostate cancer model. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2008</b> , 105, 20179-84	11.5	84

159	A regulator of Dscam mutually exclusive splicing fidelity. <i>Nature Structural and Molecular Biology</i> , <b>2007</b> , 14, 1134-40	17.6	84
158	Widespread RNA editing dysregulation in brains from autistic individuals. <i>Nature Neuroscience</i> , <b>2019</b> , 22, 25-36	25.5	84
157	Systematic Discovery of RNA Binding Proteins that Regulate MicroRNA Levels. <i>Molecular Cell</i> , <b>2018</b> , 69, 1005-1016.e7	17.6	80
156	MEDU-44. MUSASHI-1 IS A MASTER REGULATOR OF ABERRANT TRANSLATION IN GROUP 3 MEDULLOBLASTOMA. <i>Neuro-Oncology</i> , <b>2019</b> , 21, ii112-ii113	1	78
155	TMOD-28. AUTHENTIC HUMAN GLIOMA MODELING USING GENETICALLY ENGINEERED INDUCED PLURIPOTENT STEM CELLS. <i>Neuro-Oncology</i> , <b>2019</b> , 21, vi268-vi269	1	78
154	Direct RNA sequencing enables mA detection in endogenous transcript isoforms at base-specific resolution. <i>Rna</i> , <b>2020</b> , 26, 19-28	5.8	78
153	OTEH-9. scRNA sequencing of proneural GBM avatar model reveals acquisition of oncogenic transcriptional programming and infers a developmental path towards a genomically unstable state. <i>Neuro-Oncology Advances</i> , <b>2021</b> , 3, ii12-ii12	0.9	78
152	Interaction Landscape of Inherited Polymorphisms with Somatic Events in Cancer. <i>Cancer Discovery</i> , <b>2017</b> , 7, 410-423	24.4	77
151	Enhanced CLIP Uncovers IMP Protein-RNA Targets in Human Pluripotent Stem Cells Important for Cell Adhesion and Survival. <i>Cell Reports</i> , <b>2016</b> , 15, 666-679	10.6	75
150	RNA-binding proteins in neurodegeneration: Seq and you shall receive. <i>Trends in Neurosciences</i> , <b>2015</b> , 38, 226-36	13.3	73
149	Hippocampus-dependent learning is associated with adult neurogenesis in MRL/MpJ mice. <i>Hippocampus</i> , <b>2009</b> , 19, 658-69	3.5	73
148	Musashi-2 attenuates AHR signalling to expand human haematopoietic stem cells. <i>Nature</i> , <b>2016</b> , 532, 508-511	50.4	71
147	Target Discrimination in Nonsense-Mediated mRNA Decay Requires Upf1 ATPase Activity. <i>Molecular Cell</i> , <b>2015</b> , 59, 413-25	17.6	65
146	Inference of splicing regulatory activities by sequence neighborhood analysis. <i>PLoS Genetics</i> , <b>2006</b> , 2, e191	6	65
145	PPAR- $\delta$ s repressed in Huntington's disease, is required for normal neuronal function and can be targeted therapeutically. <i>Nature Medicine</i> , <b>2016</b> , 22, 37-45	50.5	64
144	Base editing: advances and therapeutic opportunities. <i>Nature Reviews Drug Discovery</i> , <b>2020</b> , 19, 839-859	64.1	60
143	Context-dependent functional compensation between Ythdf mA reader proteins. <i>Genes and Development</i> , <b>2020</b> , 34, 1373-1391	12.6	59
142	Regulation of RNA editing by RNA-binding proteins in human cells. <i>Communications Biology</i> , <b>2019</b> , 2, 19	6.7	59

141	Evidence for premature aging due to oxidative stress in iPSCs from Cockayne syndrome. <i>Human Molecular Genetics</i> , <b>2012</b> , 21, 3825-34	5.6	58
140	Heterogenous Populations of Tissue-Resident CD8 T Cells Are Generated in Response to Infection and Malignancy. <i>Immunity</i> , <b>2020</b> , 52, 808-824.e7	32.3	57
139	A Gene Regulatory Network Cooperatively Controlled by Pdx1 and Sox9 Governs Lineage Allocation of Foregut Progenitor Cells. <i>Cell Reports</i> , <b>2015</b> , 13, 326-36	10.6	56
138	A Small RNA-Catalytic Argonaute Pathway Tunes Germline Transcript Levels to Ensure Embryonic Divisions. <i>Cell</i> , <b>2016</b> , 165, 396-409	56.2	56
137	Intracellular mRNA regulation with self-assembled locked nucleic acid polymer nanoparticles. <i>Journal of the American Chemical Society</i> , <b>2014</b> , 136, 7615-8	16.4	54
136	Principles of RNA processing from analysis of enhanced CLIP maps for 150 RNA binding proteins. <i>Genome Biology</i> , <b>2020</b> , 21, 90	18.3	54
135	Early precursors and molecular determinants of tissue-resident memory CD8 T lymphocytes revealed by single-cell RNA sequencing. <i>Science Immunology</i> , <b>2020</b> , 5,	28	50
134	MicroRNA-101 Regulates Multiple Developmental Programs to Constrain Excitation in Adult Neural Networks. <i>Neuron</i> , <b>2016</b> , 92, 1337-1351	13.9	50
133	Overriding FUS autoregulation in mice triggers gain-of-toxic dysfunctions in RNA metabolism and autophagy-lysosome axis. <i>ELife</i> , <b>2019</b> , 8,	8.9	49
132	High-Throughput and Cost-Effective Characterization of Induced Pluripotent Stem Cells. <i>Stem Cell Reports</i> , <b>2017</b> , 8, 1101-1111	8	48
131	Human cytomegalovirus infection of human embryonic stem cell-derived primitive neural stem cells is restricted at several steps but leads to the persistence of viral DNA. <i>Journal of Virology</i> , <b>2014</b> , 88, 4021-39	6.6	48
130	The dsRBP and inactive editor ADR-1 utilizes dsRNA binding to regulate A-to-I RNA editing across the <i>C. elegans</i> transcriptome. <i>Cell Reports</i> , <b>2014</b> , 6, 599-607	10.6	45
129	Biallelic mutations in the 3' exonuclease TOE1 cause pontocerebellar hypoplasia and uncover a role in snRNA processing. <i>Nature Genetics</i> , <b>2017</b> , 49, 457-464	36.3	43
128	Deep sequencing identifies new and regulated microRNAs in <i>Schmidtea mediterranea</i> . <i>Rna</i> , <b>2009</b> , 15, 1483-91	5.8	42
127	Genome-wide approaches to dissect the roles of RNA binding proteins in translational control: implications for neurological diseases. <i>Frontiers in Neuroscience</i> , <b>2012</b> , 6, 144	5.1	40
126	Reactivation of fetal splicing programs in diabetic hearts is mediated by protein kinase C signaling. <i>Journal of Biological Chemistry</i> , <b>2013</b> , 288, 35372-86	5.4	40
125	A distinct microRNA signature for definitive endoderm derived from human embryonic stem cells. <i>Stem Cells and Development</i> , <b>2010</b> , 19, 797-807	4.4	40
124	GC Repeat RNA Initiates a POM121-Mediated Reduction in Specific Nucleoporins in C9orf72 ALS/FTD. <i>Neuron</i> , <b>2020</b> , 107, 1124-1140.e11	13.9	40

123	Transcriptome-pathology correlation identifies interplay between TDP-43 and the expression of its kinase CK1E in sporadic ALS. <i>Acta Neuropathologica</i> , <b>2018</b> , 136, 405-423	14.3	38
122	The RNA Helicase DDX6 Controls Cellular Plasticity by Modulating P-Body Homeostasis. <i>Cell Stem Cell</i> , <b>2019</b> , 25, 622-638.e13	18	35
121	Dysregulation of RBFOX2 Is an Early Event in Cardiac Pathogenesis of Diabetes. <i>Cell Reports</i> , <b>2016</b> , 15, 2200-2213	10.6	35
120	A Large-Scale Binding and Functional Map of Human RNA Binding Proteins		35
119	Pooled CRISPR screens with imaging on microarray reveals stress granule-regulatory factors. <i>Nature Methods</i> , <b>2020</b> , 17, 636-642	21.6	33
118	Coordinate Nodal and BMP inhibition directs Baf60c-dependent cardiomyocyte commitment. <i>Genes and Development</i> , <b>2013</b> , 27, 2332-44	12.6	32
117	Robust, Cost-Effective Profiling of RNA Binding Protein Targets with Single-end Enhanced Crosslinking and Immunoprecipitation (seCLIP). <i>Methods in Molecular Biology</i> , <b>2017</b> , 1648, 177-200	1.4	32
116	Functional genomic analysis of the let-7 regulatory network in <i>Caenorhabditis elegans</i> . <i>PLoS Genetics</i> , <b>2013</b> , 9, e1003353	6	31
115	Immunoglobulin light chain (IgL) genes in zebrafish: Genomic configurations and inversional rearrangements between (V(L)-J(L)-C(L)) gene clusters. <i>Developmental and Comparative Immunology</i> , <b>2008</b> , 32, 421-34	3.2	31
114	Transcriptome-wide profiles of circular RNA and RNA-binding protein interactions reveal effects on circular RNA biogenesis and cancer pathway expression. <i>Genome Medicine</i> , <b>2020</b> , 12, 112	14.4	31
113	Retrotransposon long interspersed nucleotide element-1 (LINE-1) is activated during salamander limb regeneration. <i>Development Growth and Differentiation</i> , <b>2012</b> , 54, 673-85	3	30
112	Heterogeneity and clonal relationships of adaptive immune cells in ulcerative colitis revealed by single-cell analyses. <i>Science Immunology</i> , <b>2020</b> , 5,	28	30
111	Regulation of asymmetric division and CD8+ T lymphocyte fate specification by protein kinase C $\delta$ and protein kinase C $\zeta$ . <i>Journal of Immunology</i> , <b>2015</b> , 194, 2249-59	5.3	29
110	Reintroduction of the archaic variant of in cortical organoids alters neurodevelopment. <i>Science</i> , <b>2021</b> , 371,	33.3	28
109	Active Protein Neddylation or Ubiquitylation Is Dispensable for Stress Granule Dynamics. <i>Cell Reports</i> , <b>2019</b> , 27, 1356-1363.e3	10.6	27
108	Applications of Cas9 as an RNA-programmed RNA-binding protein. <i>BioEssays</i> , <b>2015</b> , 37, 732-9	4.1	27
107	Rbfox2 function in RNA metabolism is impaired in hypoplastic left heart syndrome patient hearts. <i>Scientific Reports</i> , <b>2016</b> , 6, 30896	4.9	27
106	CRISPR/Cas9-mediated integration enables TAG-eCLIP of endogenously tagged RNA binding proteins. <i>Methods</i> , <b>2017</b> , 118-119, 50-59	4.6	24



105	RNA-binding protein CPEB1 remodels host and viral RNA landscapes. <i>Nature Structural and Molecular Biology</i> , <b>2016</b> , 23, 1101-1110	17.6	24
104	Genomic analysis of the molecular neuropathology of tuberous sclerosis using a human stem cell model. <i>Genome Medicine</i> , <b>2016</b> , 8, 94	14.4	24
103	RBP-Maps enables robust generation of splicing regulatory maps. <i>Rna</i> , <b>2019</b> , 25, 193-204	5.8	24
102	Longitudinal assessment of tumor development using cancer avatars derived from genetically engineered pluripotent stem cells. <i>Nature Communications</i> , <b>2020</b> , 11, 550	17.4	23
101	RNA-targeting CRISPR systems from metagenomic discovery to transcriptomic engineering. <i>Nature Cell Biology</i> , <b>2020</b> , 22, 143-150	23.4	23
100	The Calcineurin Variant CnA $\beta$ Controls Mouse Embryonic Stem Cell Differentiation by Directing mTORC2 Membrane Localization and Activation. <i>Cell Chemical Biology</i> , <b>2016</b> , 23, 1372-1382	8.2	22
99	Genome-wide approaches in the study of microRNA biology. <i>Wiley Interdisciplinary Reviews: Systems Biology and Medicine</i> , <b>2011</b> , 3, 491-512	6.6	21
98	Patch-Seq Protocol to Analyze the Electrophysiology, Morphology and Transcriptome of Whole Single Neurons Derived From Human Pluripotent Stem Cells. <i>Frontiers in Molecular Neuroscience</i> , <b>2018</b> , 11, 261	6.1	21
97	Integrin Activation Controls Regulatory T Cell-Mediated Peripheral Tolerance. <i>Journal of Immunology</i> , <b>2018</b> , 200, 4012-4023	5.3	20
96	Nol12 is a multifunctional RNA binding protein at the nexus of RNA and DNA metabolism. <i>Nucleic Acids Research</i> , <b>2017</b> , 45, 12509-12528	20.1	20
95	Self-Transfecting Micellar RNA: Modulating Nanoparticle Cell Interactions via High Density Display of Small Molecule Ligands on Micelle Coronas. <i>Bioconjugate Chemistry</i> , <b>2018</b> , 29, 126-135	6.3	19
94	Tethered Function Assays as Tools to Elucidate the Molecular Roles of RNA-Binding Proteins. <i>Advances in Experimental Medicine and Biology</i> , <b>2016</b> , 907, 61-88	3.6	19
93	The neural editome reveals an ADAR target mRNA required for proper chemotaxis. <i>ELife</i> , <b>2017</b> , 6,	8.9	18
92	AMPK regulation of Raptor and TSC2 mediate metformin effects on transcriptional control of anabolism and inflammation. <i>Genes and Development</i> , <b>2020</b> , 34, 1330-1344	12.6	18
91	An important class of intron retention events in human erythroblasts is regulated by cryptic exons proposed to function as splicing decoys. <i>Rna</i> , <b>2018</b> , 24, 1255-1265	5.8	18
90	Allele-specific binding of RNA-binding proteins reveals functional genetic variants in the RNA. <i>Nature Communications</i> , <b>2019</b> , 10, 1338	17.4	17
89	Blurred Boundaries: The RNA Binding Protein Lin28A Is Also an Epigenetic Regulator. <i>Molecular Cell</i> , <b>2016</b> , 61, 1-2	17.6	17
88	A novel splice-site mutation in ALS2 establishes the diagnosis of juvenile amyotrophic lateral sclerosis in a family with early onset anarthria and generalized dystonias. <i>PLoS ONE</i> , <b>2014</b> , 9, e113258	3.7	17

87	Inhibition of YTHDF2 triggers proteotoxic cell death in MYC-driven breast cancer. <i>Molecular Cell</i> , <b>2021</b> , 81, 3048-3064.e9	17.6	17
86	An in vivo genome-wide CRISPR screen identifies the RNA-binding protein Staufen2 as a key regulator of myeloid leukemia. <i>Nature Cancer</i> , <b>2020</b> , 1, 410-422	15.4	16
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