

# Norihiro Okada

## List of Publications by Year in descending order

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134  
papers

9,921  
citations

57681

46  
h-index

43601

95  
g-index

136  
all docs

136  
docs citations

136  
times ranked

8288  
citing authors

#	ARTICLE	IF	CITATIONS
1	Kampo formulas alleviate aging-related emotional disturbances and neuroinflammation in male senescence-accelerated mouse prone 8 mice. <i>Aging</i> , 2022, 14, 109-142.	1.4	5
2	User pairing using laser chaos decision maker for NOMA systems. <i>Nonlinear Theory and Its Applications IEICE</i> , 2022, 13, 72-83.	0.4	9
3	Data of RNA-seq transcriptomes of liver, bone, heart, kidney and blood in klotho mice at a pre-symptomatic state and the effect of a traditional Japanese multi-herbal medicine, juzentaihoto. <i>Data in Brief</i> , 2022, 42, 108197.	0.5	2
4	Intron retention is a stress response in sensor genes and is restored by Japanese herbal medicines: A basis for future clinical applications. <i>Gene</i> , 2022, 830, 146496.	1.0	0
5	SINEs as Credible Signs to Prove Common Ancestry in the Tree of Life: A Brief Review of Pioneering Case Studies in Retroposon Systematics. <i>Genes</i> , 2022, 13, 989.	1.0	3
6	Analysis on Effectiveness of Surrogate Data-Based Laser Chaos Decision Maker. <i>Complexity</i> , 2021, 2021, 1-9.	0.9	9
7	Genomic Signatures for Species-Specific Adaptation in Lake Victoria Cichlids Derived from Large-Scale Standing Genetic Variation. <i>Molecular Biology and Evolution</i> , 2021, 38, 3111-3125.	3.5	9
8	A unique neurogenomic state emerges after aggressive confrontations in males of the fish <i>Betta splendens</i> . <i>Gene</i> , 2021, 784, 145601.	1.0	3
9	Intron retention as a new pre-symptomatic marker of aging and its recovery to the normal state by a traditional Japanese multi-herbal medicine. <i>Gene</i> , 2021, 794, 145752.	1.0	10
10	Data of RNA-seq transcriptomes in the brain associated with aggression in males of the fish <i>Betta splendens</i> . <i>Data in Brief</i> , 2021, 38, 107448.	0.5	0
11	Alternative splicing plays key roles in response to stress across different stages of fighting in the fish <i>Betta splendens</i> . <i>BMC Genomics</i> , 2021, 22, .	1.2	2
12	Implication of a new function of human tDNAs in chromatin organization. <i>Scientific Reports</i> , 2020, 10, 17440.	1.6	6
13	Behavioral and brain- transcriptomic synchronization between the two opponents of a fighting pair of the fish <i>Betta splendens</i> . <i>PLoS Genetics</i> , 2020, 16, e1008831.	1.5	22
14	Title is missing!. , 2020, 16, e1008831.		0
15	Title is missing!. , 2020, 16, e1008831.		0
16	Title is missing!. , 2020, 16, e1008831.		0
17	Title is missing!. , 2020, 16, e1008831.		0
18	Title is missing!. , 2020, 16, e1008831.		0

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19	Title is missing!. , 2020, 16, e1008831.		0
20	Patterns of genomic differentiation between two Lake Victoria cichlid species, <i>Haplochromis pyrrhocephalus</i> and <i>H. sp. "macula"</i> . <i>BMC Evolutionary Biology</i> , 2019, 19, 68.	3.2	4
21	Evolution of <i>Shh</i> endoderm enhancers during morphological transition from ventral lungs to dorsal gas bladder. <i>Nature Communications</i> , 2017, 8, 14300.	5.8	19
22	Laterality is Universal Among Fishes but Increasingly Cryptic Among Derived Groups. <i>Zoological Science</i> , 2017, 34, 267.	0.3	17
23	Solution structure of a reverse transcriptase recognition site of a LINE RNA from zebrafish. <i>Journal of Biochemistry</i> , 2017, 162, 279-285.	0.9	6
24	Visual adaptation in Lake Victoria cichlid fishes: depth-related variation of color and scotopic opsins in species from sand/mud bottoms. <i>BMC Evolutionary Biology</i> , 2017, 17, 200.	3.2	28
25	Molecular developmental mechanism in polypterid fish provides insight into the origin of vertebrate lungs. <i>Scientific Reports</i> , 2016, 6, 30580.	1.6	41
26	MetaSINES: Broad Distribution of a Novel SINE Superfamily in Animals. <i>Genome Biology and Evolution</i> , 2016, 8, 528-539.	1.1	22
27	Determining the Position of Storks on the Phylogenetic Tree of Waterbirds by Retroposon Insertion Analysis. <i>Genome Biology and Evolution</i> , 2015, 7, 3180-3189.	1.1	16
28	Distinct functions of two olfactory marker protein genes derived from teleost-specific whole genome duplication. <i>BMC Evolutionary Biology</i> , 2015, 15, 245.	3.2	16
29	A microsatellite-based genetic linkage map and putative sex-determining genomic regions in Lake Victoria cichlids. <i>Gene</i> , 2015, 560, 156-164.	1.0	12
30	LINE retrotransposition and host DNA repair machinery. <i>Mobile Genetic Elements</i> , 2015, 5, 92-97.	1.8	3
31	Mechanism by which a LINE protein recognizes its 3' tail RNA. <i>Nucleic Acids Research</i> , 2014, 42, 10605-10617.	6.5	22
32	Integrated mechanism for the generation of the 5' junctions of LINE inserts. <i>Nucleic Acids Research</i> , 2014, 42, 13269-13279.	6.5	4
33	Multiple Episodic Evolution Events in V1R Receptor Genes of East-African Cichlids. <i>Genome Biology and Evolution</i> , 2014, 6, 1135-1144.	1.1	22
34	Evolutionary Changes in Vertebrate Genome Signatures with Special Focus on Coelacanth. <i>DNA Research</i> , 2014, 21, 459-467.	1.5	8
35	The genomic substrate for adaptive radiation in African cichlid fish. <i>Nature</i> , 2014, 513, 375-381.	13.7	874
36	Sensory drive speciation and patterns of variation at selectively neutral genes. <i>Evolutionary Ecology</i> , 2014, 28, 591-609.	0.5	4

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37	High prevalence of non-synonymous substitutions in mtDNA of cichlid fishes from Lake Victoria. <i>Gene</i> , 2014, 552, 239-245.	1.0	7
38	Coelacanth genomes reveal signatures for evolutionary transition from water to land. <i>Genome Research</i> , 2013, 23, 1740-1748.	2.4	108
39	Lineage-Specific Expansion of Vomeronasal Type 2 Receptor-Like (OlfC) Genes in Cichlids May Contribute to Diversification of Amino Acid Detection Systems. <i>Genome Biology and Evolution</i> , 2013, 5, 711-722.	1.1	26
40	Genetic Structure of Pelagic and Littoral Cichlid Fishes from Lake Victoria. <i>PLoS ONE</i> , 2013, 8, e74088.	1.1	11
41	Correlation between Nuptial Colors and Visual Sensitivities Tuned by Opsins Leads to Species Richness in Sympatric Lake Victoria Cichlid Fishes. <i>Molecular Biology and Evolution</i> , 2012, 29, 3281-3296.	3.5	45
42	A new mechanism to ensure integration during LINE retrotransposition: A suggestion from analyses of the 5' extra nucleotides. <i>Gene</i> , 2012, 505, 345-351.	1.0	6
43	Characterization of V1R receptor ( <i>ora</i> ) genes in Lake Victoria cichlids. <i>Gene</i> , 2012, 499, 273-279.	1.0	18
44	Reverse Evolution in RH1 for Adaptation of Cichlids to Water Depth in Lake Tanganyika. <i>Molecular Biology and Evolution</i> , 2011, 28, 1769-1776.	3.5	33
45	B Chromosomes Have a Functional Effect on Female Sex Determination in Lake Victoria Cichlid Fishes. <i>PLoS Genetics</i> , 2011, 7, e1002203.	1.5	134
46	Speciation of Cichlid Fishes by Sensory Drive. <i>Primate Monographs</i> , 2011, , 311-328.	0.8	6
47	Emergence of mammals by emergency: exaptation. <i>Genes To Cells</i> , 2010, 15, 801-812.	0.5	27
48	Vertebrate Rhodopsin Adaptation to Dim Light via Rapid Meta-II Intermediate Formation. <i>Molecular Biology and Evolution</i> , 2010, 27, 506-519.	3.5	34
49	Genetic variation and demographic history of the <i>Haplochromis laparogramma</i> group of Lake Victoria—An analysis based on SINEs and mitochondrial DNA. <i>Gene</i> , 2010, 450, 39-47.	1.0	17
50	Characterization of a novel SINE superfamily from invertebrates: “Ceph-SINEs” from the genomes of squids and cuttlefish. <i>Gene</i> , 2010, 454, 8-19.	1.0	28
51	mRNA Retrotransposition Coupled with 5' Inversion as a Possible Source of New Genes. <i>Molecular Biology and Evolution</i> , 2009, 26, 1405-1420.	3.5	11
52	Genetic Evidence That the Non-Homologous End-Joining Repair Pathway Is Involved in LINE Retrotransposition. <i>PLoS Genetics</i> , 2009, 5, e1000461.	1.5	121
53	Retroposons of salmonoid fishes (Actinopterygii: Salmonoidei) and their evolution. <i>Gene</i> , 2009, 434, 16-28.	1.0	24
54	Extensive analysis of EST sequences reveals that all cichlid species in Lake Victoria share almost identical transcript sets. <i>Gene</i> , 2009, 441, 187-191.	1.0	14

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55	The evolution of two partner LINE/SINE families and a full-length chromodomain-containing Ty3/Gypsy LTR element in the first reptilian genome of <i>Anolis carolinensis</i> . <i>Gene</i> , 2009, 441, 111-118.	1.0	41
56	Retroposon analysis and recent geological data suggest near-simultaneous divergence of the three superorders of mammals. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 5235-5240.	3.3	162
57	Speciation through sensory drive in cichlid fish. <i>Nature</i> , 2008, 455, 620-626.	13.7	947
58	Newly discovered young CORE-SINEs in marsupial genomes. <i>Gene</i> , 2008, 407, 176-185.	1.0	23
59	Retroposons: Genetic Footprints on the Evolutionary Paths of Life. <i>Methods in Molecular Biology</i> , 2008, 422, 201-225.	0.4	20
60	Novel SINE Families from Salmon Validate Parahucho ( <i>Salmonidae</i> ) as a Distinct Genus and Give Evidence that SINEs Can Incorporate LINE-related 3' Tails of Other SINEs. <i>Molecular Biology and Evolution</i> , 2007, 24, 1656-1666.	3.5	27
61	Poxviruses as possible vectors for horizontal transfer of retroposons from reptiles to mammals. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 12046-12051.	3.3	119
62	Functional splice sites in a zebrafish LINE and their influence on zebrafish gene expression. <i>Gene</i> , 2007, 390, 221-231.	1.0	11
63	A new system for analyzing LINE retrotransposition in the chicken DT40 cell line widely used for reverse genetics. <i>Gene</i> , 2007, 395, 116-124.	1.0	6
64	MyrSINEs: A novel SINE family in the anteater genomes. <i>Gene</i> , 2007, 400, 98-103.	1.0	11
65	Rooting the eutherian tree: the power and pitfalls of phylogenomics. <i>Genome Biology</i> , 2007, 8, R199.	13.9	82
66	Toothed whale monophyly reassessed by SINE insertion analysis: The absence of lineage sorting effects suggests a small population of a common ancestral species. <i>Molecular Phylogenetics and Evolution</i> , 2007, 43, 216-224.	1.2	43
67	Genomic expansion of the Bov-A2 retroposon relating to phylogeny and breed management. <i>Mammalian Genome</i> , 2007, 18, 187-196.	1.0	11
68	Pegasoferae, an unexpected mammalian clade revealed by tracking ancient retroposon insertions. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 9929-9934.	3.3	207
69	Isolation and characterization of retrotransposition-competent LINEs from zebrafish. <i>Gene</i> , 2006, 365, 74-82.	1.0	42
70	Probing the secondary structure of salmon Smal SINE RNA. <i>Gene</i> , 2006, 365, 67-73.	1.0	11
71	The rise and fall of the CR1 subfamily in the lineage leading to penguins. <i>Gene</i> , 2006, 365, 57-66.	1.0	39
72	magp4 gene may contribute to the diversification of cichlid morphs and their speciation. <i>Gene</i> , 2006, 373, 126-133.	1.0	23

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73	Genomic alterations upon integration of zebrafish L1 elements revealed by the TANT method. <i>Gene</i> , 2006, 383, 108-116.	1.0	7
74	<i>Balaenoptera omurai</i> is a newly discovered baleen whale that represents an ancient evolutionary lineage. <i>Molecular Phylogenetics and Evolution</i> , 2006, 41, 40-52.	1.2	84
75	Sauria SINEs: Novel Short Interspersed Retroposable Elements That Are Widespread in Reptile Genomes. <i>Journal of Molecular Evolution</i> , 2006, 62, 630-644.	0.8	55
76	Solution structure and functional importance of a conserved RNA hairpin of eel LINE UnaL2. <i>Nucleic Acids Research</i> , 2006, 34, 5184-5193.	6.5	18
77	Functional noncoding sequences derived from SINEs in the mammalian genome. <i>Genome Research</i> , 2006, 16, 864-874.	2.4	207
78	Baleen Whale Phylogeny and a Past Extensive Radiation Event Revealed by SINE Insertion Analysis. <i>Molecular Biology and Evolution</i> , 2006, 23, 866-873.	3.5	69
79	Divergent Selection on Opsins Drives Incipient Speciation in Lake Victoria Cichlids. <i>PLoS Biology</i> , 2006, 4, e433.	2.6	167
80	Extensive Morphological Convergence and Rapid Radiation in the Evolutionary History of the Family Geoemydidae (Old World Pond Turtles) Revealed by SINE Insertion Analysis. <i>Systematic Biology</i> , 2006, 55, 912-927.	2.7	29
81	Isolation and Characterization of Active LINE and SINEs from the Eel. <i>Molecular Biology and Evolution</i> , 2005, 22, 673-682.	3.5	29
82	A Retroposon Analysis of Afrotherian Phylogeny. <i>Molecular Biology and Evolution</i> , 2005, 22, 1823-1833.	3.5	88
83	Retroposon Mapping in Molecular Systematics. , 2004, 260, 189-226.		47
84	Solution structure of an RNA stem-loop derived from the 3' conserved region of eel LINE UnaL2. <i>Rna</i> , 2004, 10, 1380-1387.	1.6	24
85	First Application of the SINE (Short Interspersed Repetitive Element) Method to Infer Phylogenetic Relationships in Reptiles: An Example from the Turtle Superfamily Testudinoidea. <i>Molecular Biology and Evolution</i> , 2004, 21, 705-715.	3.5	46
86	Phylogenetic Relationships Among East African Haplochromine Fish as Revealed by Short Interspersed Elements (SINEs). <i>Journal of Molecular Evolution</i> , 2004, 58, 64-78.	0.8	38
87	SINEs of speciation: tracking lineages with retroposons. <i>Trends in Ecology and Evolution</i> , 2004, 19, 545-553.	4.2	143
88	Mitochondrial phylogeny of hedgehogs and monophyly of Eulipotyphla. <i>Molecular Phylogenetics and Evolution</i> , 2003, 28, 276-284.	1.2	61
89	Whole-genome screening indicates a possible burst of formation of processed pseudogenes and Alu repeats by particular L1 subfamilies in ancestral primates. <i>Genome Biology</i> , 2003, 4, R74.	13.9	172
90	Ancient SINEs from African Endemic Mammals. <i>Molecular Biology and Evolution</i> , 2003, 20, 522-527.	3.5	81

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91	Using SINEs to Probe Ancient Explosive Speciation: "Hidden" Radiation of African Cichlids?. <i>Molecular Biology and Evolution</i> , 2003, 20, 924-930.	3.5	50
92	Unique Mammalian tRNA-Derived Repetitive Elements in Dermopterans: The t-SINE Family and Its Retrotransposition Through Multiple Sources. <i>Molecular Biology and Evolution</i> , 2003, 20, 1659-1668.	3.5	34
93	Nonlinear partial differential equations and applications: The effect of selection on a long wavelength-sensitive (LWS) opsin gene of Lake Victoria cichlid fishes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 15501-15506.	3.3	118
94	Mosaic Structure and Retropositional Dynamics During Evolution of Subfamilies of Short Interspersed Elements in African Cichlids. <i>Molecular Biology and Evolution</i> , 2002, 19, 1303-1312.	3.5	25
95	Characterization of Novel Alu- and tRNA-Related SINEs from the Tree Shrew and Evolutionary Implications of Their Origins. <i>Molecular Biology and Evolution</i> , 2002, 19, 1964-1972.	3.5	76
96	V-SINEs: A New Superfamily of Vertebrate SINEs That Are Widespread in Vertebrate Genomes and Retain a Strongly Conserved Segment within Each Repetitive Unit. <i>Genome Research</i> , 2002, 12, 316-324.	2.4	97
97	The Evolution of the Pro-Domain of Bone Morphogenetic Protein 4 (Bmp4) in an Explosively Speciated Lineage of East African Cichlid Fishes. <i>Molecular Biology and Evolution</i> , 2002, 19, 1628-1632.	3.5	52
98	LINES Mobilize SINEs in the Eel through a Shared 3' Sequence. <i>Cell</i> , 2002, 111, 433-444.	13.5	295
99	Intra- and Interfamily Relationships of Vespertilionidae Inferred by Various Molecular Markers Including SINE Insertion Data. <i>Journal of Molecular Evolution</i> , 2002, 55, 284-301.	0.8	60
100	Éléments mobiles SINE en phylogénie. <i>Medecine/Sciences</i> , 2002, 18, 1276-1281.	0.0	1
101	Retroposon analysis of major cetacean lineages: The monophyly of toothed whales and the paraphyly of river dolphins. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2001, 98, 7384-7389.	3.3	239
102	Retroposition of the AFC Family of SINEs (Short Interspersed Repetitive Elements) Before and During the Adaptive Radiation of Cichlid Fishes in Lake Malawi and Related Inferences About Phylogeny. <i>Journal of Molecular Evolution</i> , 2001, 53, 496-507.	0.8	35
103	Maximum Likelihood Analysis of the Complete Mitochondrial Genomes of Eutherians and a Reevaluation of the Phylogeny of Bats and Insectivores. <i>Journal of Molecular Evolution</i> , 2001, 53, 508-516.	0.8	128
104	Evolution of CHR-2 SINEs in cetartiodactyl genomes: possible evidence for the monophyletic origin of toothed whales. <i>Mammalian Genome</i> , 2001, 12, 909-915.	1.0	34
105	Phylogenetic Relationships and Ancient Incomplete Lineage Sorting Among Cichlid Fishes in Lake Tanganyika as Revealed by Analysis of the Insertion of Retroposons. <i>Molecular Biology and Evolution</i> , 2001, 18, 2057-2066.	3.5	191
106	SINE insertions: powerful tools for molecular systematics. <i>BioEssays</i> , 2000, 22, 148-160.	1.2	336
107	CetSINEs and AREs are not SINEs but are parts of cetartiodactyl L1. <i>Mammalian Genome</i> , 2000, 11, 1123-1126.	1.0	7
108	Monophyletic Origin of the Order Chiroptera and Its Phylogenetic Position Among Mammalia, as Inferred from the Complete Sequence of the Mitochondrial DNA of a Japanese Megabat, the Ryukyu Flying Fox ( <i>Pteropus dasymallus</i> ). <i>Journal of Molecular Evolution</i> , 2000, 51, 318-328.	0.8	62

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109	Consistency of SINE Insertion Topology and Flanking Sequence Tree: Quantifying Relationships Among Cetartiodactyls. <i>Molecular Biology and Evolution</i> , 2000, 17, 1417-1424.	3.5	35
110	SINE Evolution, Missing Data, and the Origin of Whales. <i>Systematic Biology</i> , 2000, 49, 808-817.	2.7	54
111	Towards Resolving the Interordinal Relationships of Placental Mammals. <i>Systematic Biology</i> , 1999, 48, 1-5.	2.7	306
112	Phylogenetic relationships among cetartiodactyls based on insertions of short and long interspersed elements: Hippopotamuses are the closest extant relatives of whales. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1999, 96, 10261-10266.	3.3	402
113	Conflict Among Individual Mitochondrial Proteins in Resolving the Phylogeny of Eutherian Orders. <i>Journal of Molecular Evolution</i> , 1998, 47, 307-322.	0.8	208
114	Molecular evidence from short interspersed elements (SINEs) that <i>Oncorhynchus masou</i> (cherry) Tj ETQq0 0 0 rgBT/Overlock 10 Tf 50 5	0.7	11
115	Detection of the Ongoing Sorting of Ancestrally Polymorphic SINEs Toward Fixation or Loss in Populations of Two Species of Charr During Speciation. <i>Genetics</i> , 1998, 150, 301-311.	1.2	34
116	SINEs and LINEs share common 3' ends sequences: a review. <i>Gene</i> , 1997, 205, 229-243.	1.0	256
117	Molecular evidence from retroposons that whales form a clade within even-toed ungulates. <i>Nature</i> , 1997, 388, 666-670.	13.7	413
118	The 3' ends of tRNA-derived SINEs originated from the 3' ends of LINEs: A new example from the bovine genome. <i>Journal of Molecular Evolution</i> , 1997, 44, S52-S56.	0.8	85
119	The Origin of Chlorarachniophyte Plastids, as Inferred from Phylogenetic Comparisons of Amino Acid Sequences of EF-Tu. <i>Journal of Molecular Evolution</i> , 1997, 45, 682-687.	0.8	72
120	A Newly Isolated Family of Short Interspersed Repetitive Elements (SINEs) in Coregonid Fishes (Whitefish) With Sequences That Are Almost Identical To Those of the Smal Family of Repeats: Possible Evidence for the Horizontal Transfer of SINEs. <i>Genetics</i> , 1997, 146, 355-367.	1.2	49
121	The Salmon <i>Sma</i> Family of Short Interspersed Repetitive Elements (SINEs): Interspecific and Intraspecific Variation of the Insertion of SINEs in the Genomes of Chum and Pink Salmon. <i>Genetics</i> , 1997, 146, 369-380.	1.2	34
122	Characterization of species-specifically amplified SINEs in three salmonid species—Chum salmon, pink salmon, and kokanee: The local environment of the genome may be important for the generation of a dominant source gene at a newly retroposed locus. <i>Journal of Molecular Evolution</i> , 1996, 42, 103-116.	0.8	36
123	Protein phylogeny of translation elongation factor EF-1 $\alpha$ suggests microsporidians are extremely ancient eukaryotes. <i>Journal of Molecular Evolution</i> , 1996, 42, 257-263.	0.8	110
124	Details of Retropositional Genome Dynamics That Provide a Rationale for a Generic Division: The Distinct Branching of All the Pacific Salmon and Trout ( <i>Oncorhynchus</i> ) From the Atlantic Salmon and Trout ( <i>Salmo</i> ). <i>Genetics</i> , 1996, 142, 915-926.	1.2	71
125	Evolution of the active sequences of the HpaI short interspersed elements. <i>Journal of Molecular Evolution</i> , 1995, 41, 986-995.	0.8	29
126	A Novel tRNA Species as an Origin of Short Interspersed Repetitive Elements (SINEs). <i>Journal of Molecular Biology</i> , 1994, 239, 731-735.	2.0	36



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127	Amplification of Distinct Subfamilies of Short Interspersed Elements During Evolution of the Salmonidae. <i>Journal of Molecular Biology</i> , 1994, 241, 633-644.	2.0	44
128	Generality of the tRNA Origin of Short Interspersed Repetitive Elements (SINEs). <i>Journal of Molecular Biology</i> , 1994, 243, 25-37.	2.0	33
129	Determination of the phylogenetic relationships among Pacific salmonids by using short interspersed elements (SINEs) as temporal landmarks of evolution.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1993, 90, 6995-6999.	3.3	188
130	SINEs: Short interspersed repeated elements of the eukaryotic genome. <i>Trends in Ecology and Evolution</i> , 1991, 6, 358-361.	4.2	167
131	SINEs. <i>Current Opinion in Genetics and Development</i> , 1991, 1, 498-504.	1.5	200
132	A highly repetitive and transcribable sequence in the tortoise genome is probably a retroposon. <i>FEBS Journal</i> , 1990, 189, 25-31.	0.2	35
133	Rodent type 2 Alu family, rat identifier sequence, rabbit C family, and bovine or goat 73-bp repeat may have evolved from tRNA genes. <i>Journal of Molecular Evolution</i> , 1985, 22, 134-140.	0.8	111
134	SINE insertions: powerful tools for molecular systematics. , 0, .		3