Prescott L Deininger

List of Publications by Year
in descending order

165 papers	18,129 citations	20817 60 h-index	14208 128 g-index
177 all docs	177 docs citations	177 times ranked	11091 citing authors

#	Article	IF	Citations
1	DNA sequence and expression of the B95-8 Epsteinâ€"Barr virus genome. Nature, 1984, 310, 207-211.	27.8	2,339
2	Alu repeats and human genomic diversity. Nature Reviews Genetics, 2002, 3, 370-379.	16.3	1,245
3	Alu Repeats and Human Disease. Molecular Genetics and Metabolism, 1999, 67, 183-193.	1.1	825
4	Base sequence studies of 300 nucleotide renatured repeated human DNA clones. Journal of Molecular Biology, 1981, 151, 17-33.	4.2	629
5	Random subcloning of sonicated DNA: Application to shotgun DNA sequence analysis. Analytical Biochemistry, 1983, 129, 216-223.	2.4	618
6	Alu elements: know the SINEs. Genome Biology, 2011, 12, 236.	9.6	465
7	The Human LINE-1 Retrotransposon Creates DNA Double-strand Breaks. Journal of Molecular Biology, 2006, 357, 1383-1393.	4.2	431
8	Mobile elements and mammalian genome evolution. Current Opinion in Genetics and Development, 2003, 13, 651-658.	3.3	422
9	Dominant and recessive deafness caused by mutations of a novel gene, TMC1, required for cochlear hair-cell function. Nature Genetics, 2002, 30, 277-284.	21.4	395
10	Partial nucleotide sequence of the 300-nucleotide interspersed repeated human DNA sequences. Nature, 1980, 284, 372-374.	27.8	351
11	AluInsertion Polymorphisms and Human Evolution: Evidence for a Larger Population Size inâ€‰Africa. Genome Research, 1997, 7, 1061-1071.	5.5	311
12	Mammalian Retroelements. Genome Research, 2002, 12, 1455-1465.	5.5	309
13	Evolution of the master Alu gene(s). Journal of Molecular Evolution, 1991, 33, 311-320.	1.8	287
14	Master genes in mammalian repetitive DNA amplification. Trends in Genetics, 1992, 8, 307-311.	6.7	287
15	Sequence organization of the human genome. Cell, 1975, 6, 345-358.	28.9	285
16	Mammalian non-LTR retrotransposons: For better or worse, in sickness and in health. Genome Research, 2008, 18, 343-358.	5.5	285
17	Inviting instability: Transposable elements, double-strand breaks, and the maintenance of genome integrity. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2007, 616, 46-59.	1.0	270
18	Standardized nomenclature for Alu repeats. Journal of Molecular Evolution, 1996, 42, 3-6.	1.8	261

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19	A human-specific subfamily of Alu sequences. Genomics, 1991, 9, 481-487.	2.9	260
20	RNA truncation by premature polyadenylation attenuates human mobile element activity. Nature Genetics, 2003, 35, 363-366.	21.4	241
21	Mutation near the polyoma DNA replication origin permits productive infection of F9 embryonal carcinoma cells. Cell, 1981, 23, 809-814.	28.9	240
22	Repeat sequence families derived from mammalian tRNA genes. Nature, 1985, 317, 819-822.	27.8	238
23	The nucleotide sequence and genome organization of the polyoma early region: Extensive nucleotide and amino acid homology with SV40. Cell, 1979, 17, 715-724.	28.9	210
24	Somatic expression of LINE-1 elements in human tissues. Nucleic Acids Research, 2010, 38, 3909-3922.	14.5	206
25	Identification of a New Subclass of Alu DNA Repeats Which Can Function as Estrogen Receptor-dependent Transcriptional Enhancers. Journal of Biological Chemistry, 1995, 270, 22777-22782.	3.4	205
26	Alu Repeats: A Source for the Genesis of Primate Microsatellites. Genomics, 1995, 29, 136-144.	2.9	205
27	Genetic variation of recent Alu insertions in human populations. Journal of Molecular Evolution, 1996, 42, 22-29.	1.8	194
28	Structure and variability of recently inserted Alu family members. Nucleic Acids Research, 1990, 18, 6793-6798.	14.5	193
29	LINE-1 RNA splicing and influences on mammalian gene expression. Nucleic Acids Research, 2006, 34, 1512-1521.	14.5	180
30	The recent evolution of mammalian repetitive DNA elements. Trends in Genetics, 1986, 2, 76-80.	6.7	166
31	All yâ€™all need to know â€˜bout retroelements in cancer. Seminars in Cancer Biology, 2010, 20, 200-210.	9.6	166
32	Large-scale analysis of the Alu Ya5 and Yb8 subfamilies and their contribution to human genomic diversity. Journal of Molecular Biology, 2001, 311, 17-40.	4.2	152
33	Emergence of primate genes by retrotransposon-mediated sequence transduction. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 17608-17613.	7.1	141
34	Sequence, structure and promoter characterization of the human thymidine kinase gene. Gene, 1987, 52, 267-277.	2.2	133
35	L1 mobile element expression causes multiple types of toxicity. Gene, 2008, 419, 75-81.	2.2	128
36	Active Alu Element â€œA-Tailsâ€: Size Does Matter. Genome Research, 2002, 12, 1333-1344.	5.5	127

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37	Alu Insertion Polymorphisms for the Study of Human Genomic Diversity. Genetics, 2001, 159, 279-290.	2.9	127
38	Analysis of transcription of the human Alu family ubiquitous repeating element by eukaryotic RNA polymerase III. Nucleic Acids Research, 1981, 9, 6439-6456.	14.5	125
39	An electron microscope study of the DNA sequence organization of the human genome. Journal of Molecular Biology, 1976, 106, 773-790.	4.2	123
40	Evolution of Retroposons. , 1993, , 157-196.		121
41	Amplification dynamics of human-specific (HS) alu family members. Nucleic Acids Research, 1991, 19, 3619-3623.	14.5	120
42	LINE dancing in the human genome: transposable elements and disease. Genome Medicine, 2009, 1, 97.	8.2	118
43	Renaturation rate studies of a single family of interspersed repeated sequences in human deoxyribonucleic acid. Biochemistry, 1981, 20, 3003-3010.	2.5	110
44	Potential Gene Conversion and Source Genes for Recently Integrated Alu Elements. Genome Research, 2000, 10, 1485-1495.	5.5	108
45	Heavy Metal Exposure Influences Double Strand Break DNA Repair Outcomes. PLoS ONE, 2016, 11, e0151367.	2.5	107
46	Dispersion and Insertion Polymorphism in Two Small Subfamilies of Recently Amplified HumanAluRepeats. Journal of Molecular Biology, 1995, 247, 418-427.	4.2	105
47	Integration site preferences of the Alu family and similar repetitive DNA sequences. Nucleic Acids Research, 1985, 13, 8939-8954.	14.5	104
48	Alu elements: an intrinsic source of human genome instability. Current Opinion in Virology, 2013, 3, 639-645.	5.4	95
49	ERCC1/XPF limits L1 retrotransposition. DNA Repair, 2008, 7, 983-989.	2.8	90
50	Non-traditional Alu evolution and primate genomic diversity. Journal of Molecular Biology, 2002, 316, 1033-1040.	4.2	87
51	A comprehensive approach to expression of L1 loci. Nucleic Acids Research, 2017, 45, e31-e31.	14.5	86
52	LINE-1 ORF1 protein enhances Alu SINE retrotransposition. Gene, 2008, 419, 1-6.	2.2	84
53	Recently integrated human Alu repeats: finding needles in the haystack. Genetica, 1999, 107, 149-161.	1.1	82
54	The impact of multiple splice sites in human L1 elements. Gene, 2008, 411, 38-45.	2.2	82

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55	8-Hydroxy-2â€²-deoxyguanosine (8-OH-dG) as a potential survival biomarker in patients with nonsmall-cell lung cancer. Cancer, 2007, 109, 574-580.	4.1	80
56	A second major class of Alu family repeated DNA sequences in a primate genome. Nucleic Acids Research, 1983, 11, 7595-7610.	14.5	76
57	Human retroelements may introduce intragenic polyadenylation signals. Cytogenetic and Genome Research, 2005, 110, 365-371.	1.1	75
58	Upstream flanking sequences and transcription of SINEs 1 1Edited by M. Gottesman. Journal of Molecular Biology, 2000, 302, 17-25.	4.2	74
59	Nucleotide sequence and genetic organization of the polyoma late region: Features common to the polyoma early region and SV40. Cell, 1979, 18, 771-779.	28.9	72
60	The Contribution of Alu Elements to Mutagenic DNA Double-Strand Break Repair. PLoS Genetics, 2015, 11, e1005016.	3.5	71
61	Diverse cis factors controlling Alu retrotransposition: What causes Alu elements to die?. Genome Research, 2009, 19, 545-555.	5.5	70
62	Approaches to rapid DNA sequence analysis. Analytical Biochemistry, 1983, 135, 247-263.	2.4	68
63	A study of the evolution of repeated DNA sequences in primates and the existence of a new class of repetitive sequences in primates. Journal of Molecular Biology, 1979, 127, 437-460.	4.2	63
64	Shared Protein Components of SINE RNPs. Journal of Molecular Biology, 2002, 321, 423-432.	4.2	61
65	Nickel Stimulates L1 Retrotransposition by a Post-transcriptional Mechanism. Journal of Molecular Biology, 2005, 354, 246-257.	4.2	59
66	Novel variant of the P2X2 ATP receptor from the guinea pig organ of Corti. Hearing Research, 1998, 121, 62-70.	2.0	58
67	cDNAs derived from primary and small cytoplasmic Alu (scAlu) transcripts. Journal of Molecular Biology, 1997, 271, 222-234.	4.2	56
68	Rare mutations ofFGFR2causing apert syndrome: identification of the first partial gene deletion, and anAluelement insertion from a new subfamily. Human Mutation, 2009, 30, 204-211.	2.5	55
69	Species-specific homogeneity of the primate Alu family of repeated DNA sequences. Nucleic Acids Research, 1983, 11, 7579-7593.	14.5	53
70	Heavy Metals Stimulate Human LINE-1 Retrotransposition. International Journal of Environmental Research and Public Health, 2005, 2, 14-23.	2.6	53
71	Characterization of a third major SINE family of repetitive sequences in the galago genome. Nucleic Acids Research, 1991, 19, 1649-1656.	14.5	47
72	Sporadic amplification of ID elements in rodents. Journal of Molecular Evolution, 1996, 42, 7-14.	1.8	47

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73	Polymorphic human specific Alu insertions as markers for human identification. Electrophoresis, 1995, 16, 1596-1601.	2.4	44
74	Identification and analysis of a â€˜youngâ€™ polymorphic Alu element. Biochimica Et Biophysica Acta Gene Regulatory Mechanisms, 1995, 1263, 99-102.	2.4	42
75	Analysis of CAG Repeat of the Machado-Joseph Gene in Human, Chimpanzee and Monkey Populations: A Variant Nucleotide is Associated with the Number of CAG Repeats. Human Molecular Genetics, 1996, 5, 207-213.	2.9	42
76	AluY insertion (IVS4-52ins316alu) in the glycerol kinase gene from an individual with benign glycerol kinase deficiency. Human Mutation, 2000, 15, 316-323.	2.5	42
77	Jerzy Jurka â€“ 1950â€“2014. Mobile DNA, 2015, 6, 1.	3.6	42
78	Sequence diversity and chromosomal distribution of â€œyoungâ€•Alu repeats. Gene, 1995, 163, 273-278.	2.2	40
79	A dimer satellite sequence in bonnet monkey DNA consists of distinct monomer subunits. Journal of Molecular Biology, 1980, 136, 151-167.	4.2	39
80	Recent Amplification of Rat ID Sequences. Journal of Molecular Biology, 1996, 261, 322-327.	4.2	39
81	Alu distribution and mutation types of cancer genes. BMC Genomics, 2011, 12, 157.	2.8	39
82	The aging clock and circadian control of metabolism and genome stability. Frontiers in Genetics, 2014, 5, 455.	2.3	38
83	Molecular cloning: A laboratory manual. Analytical Biochemistry, 1990, 186, 182-183.	2.4	36
84	Temporal changes in gene expression following cryogenic rat brain injury. Molecular Brain Research, 1998, 55, 9-19.	2.3	36
85	HPV 5 and 8 E6 expression reduces ATM protein levels and attenuates LINE-1 retrotransposition. Virology, 2013, 443, 69-79.	2.4	35
86	Enhanced evolutionary PCR using oligonucleotides with inosine at the 3â€²-terminus. Nucleic Acids Research, 1991, 19, 5081-5081.	14.5	33
87	Sequencing, identification and mapping of primed L1 elements (SIMPLE) reveals significant variation in full length L1 elements between individuals. BMC Genomics, 2015, 16, 220.	2.8	33
88	Transcription and processing of the rodent ID repeat family in germline and somatic cells. Nucleic Acids Research, 1995, 23, 2245-2251.	14.5	32
89	Evolution, Expression, and Possible Function of a Master Gene for Amplification of an Interspersed Repeated DNA Family in Rodents. Progress in Molecular Biology and Translational Science, 1996, 52, 67-88.	1.9	32
90	Identification and mutation analysis of a cochlear-expressed, zinc finger protein gene at the DFNB7/11 and dn hearing-loss-loci on human chromosome 9q and mouse chromosome 19. Gene, 1998, 215, 461-469.	2.2	32

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91	Functional Expression of Three P2X2 Receptor Splice Variants From Guinea Pig Cochlea. Journal of Neurophysiology, 2000, 83, 1502-1509.	1.8	32
92	Evolution of B2 repeats: the muroid explosion. Genetica, 1997, 99, 1-13.	1.1	31
93	Aluinsertion polymorphisms in Native Americans and related Asian populations. Annals of Human Biology, 2006, 33, 142-160.	1.0	31
94	Ush1c216A knock-in mouse survives Katrina. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2007, 616, 139-144.	1.0	31
95	The Nucleotide Excision Repair Pathway Limits L1 Retrotransposition. Genetics, 2017, 205, 139-153.	2.9	31
96	BC1 RNA, the Transcript from A Master Gene for ID Element Amplification, Is Able to Prime Its Own Reverse Transcription. Nucleic Acids Research, 1997, 25, 1641-1648.	14.5	30
97	The USH1C 216G?A splice-site mutation results in a 35-base-pair deletion. Human Genetics, 2005, 116, 225-227.	3.8	30
98	Applications of computational algorithm tools to identify functional SNPs in cytokine genes. Cytokine, 2006, 35, 62-66.	3.2	29
99	Requirements for polyadenylation at the 3â€² end of LINE-1 elements. Gene, 2007, 390, 98-107.	2.2	29
100	Characterization of pre-insertion loci of de novo L1 insertions. Gene, 2007, 390, 190-198.	2.2	28
101	A consensus Alu repeat probe for physical mapping. Genetic Analysis, Techniques and Applications, 1994, 11, 34-38.	1.5	27
102	DNA sequences of Alu elements indicate a recent replacement of the human autosomal genetic complement.. Proceedings of the National Academy of Sciences of the United States of America, 1996, 93, 4360-4364.	7.1	27
103	Nucleotide sequence and structure of integrated bovine leukemia virus long terminal repeats. Virology, 1985, 141, 162-166.	2.4	25
104	[16] Evolutionary analyses of repetitive DNA sequences. Methods in Enzymology, 1993, 224, 213-232.	1.0	25
105	Identification of a human specificAlu insertion in the factor XIIIB gene. Genetica, 1994, 94, 1-8.	1.1	25
106	Phylogenetic Analysis of the Friedreich Ataxia GAA Trinucleotide Repeat. Journal of Molecular Evolution, 2001, 52, 232-238.	1.8	24
107	Insertion of Retrotransposons at Chromosome Ends: Adaptive Response to Chromosome Maintenance. Frontiers in Genetics, 2015, 6, 358.	2.3	24
108	The Mobile Genetic Element "Alu" in the Human Genome. BioScience, 1996, 46, 32-41.	4.9	22

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109	Potential for Retroposition by Old Alu Subfamilies. Journal of Molecular Evolution, 2003, 56, 658-664.	1.8	22
110	Regulation of rodent myelin proteolipid protein gene expression. Neuroscience Letters, 1992, 137, 56-60.	2.1	21
111	Regions of the polyoma genome coding for T antigens. Nucleic Acids Research, 1979, 7, 2275-2288.	14.5	20
112	The role and amplification of the HS Alu subfamily founder gene. Journal of Molecular Evolution, 1996, 42, 15-21.	1.8	20
113	Near-IR single fluorophore quenching system based on phthalocyanine (Pc) aggregation and its application for monitoring inhibitor/activator action on a therapeutic target: L1-EN. Analyst, The, 2011, 136, 1103.	3.5	18
114	Alu elements and DNA double-strand break repair. Mobile Genetic Elements, 2015, 5, 81-85.	1.8	18
115	The USH1C 216Gâ†'A mutation and the 9-repeat VNTR(t,t) allele are in complete linkage disequilibrium in the Acadian population. Human Genetics, 2002, 110, 95-97.	3.8	17
116	Worldwide Genetic Variation at the 3â€²â€UTR Region of theLDLRGene: Possible Influence of Natural Selection. Annals of Human Genetics, 2005, 69, 389-400.	0.8	17
117	Tandem insertions of Alu elements. Cytogenetic and Genome Research, 2005, 108, 58-62.	1.1	16
118	Cross-Talk-Free Dual-Color Fluorescence Cross-Correlation Spectroscopy for the Study of Enzyme Activity. Analytical Chemistry, 2010, 82, 1401-1410.	6.5	16
119	miRNA-Mediated Relationships between Cis-SNP Genotypes and Transcript Intensities in Lymphocyte Cell Lines. PLoS ONE, 2012, 7, e31429.	2.5	15
120	[41] Full-Length cDNA clones: Vector-primed cDNA synthesis. Methods in Enzymology, 1987, 152, 371-389.	1.0	14
121	In vivotranscription of a cloned prosimian primate SINE sequence. Nucleic Acids Research, 1989, 17, 8669-8682.	14.5	14
122	Identification and characterization of two polymorphic Ya5 Alu repeats. Mutation Research - Mutation Research Genomics, 1997, 382, 5-11.	1.1	14
123	Predicting Mammalian SINE Subfamily Activity from A-tail Length. Molecular Biology and Evolution, 2004, 21, 2140-2148.	8.9	14
124	Feedback inhibition of L1 and alu retrotransposition through altered double strand break repair kinetics. Mobile DNA, 2010, 1, 22.	3.6	14
125	Protein binding sites within the human thymidine kinase promoter. Gene, 1992, 111, 249-254.	2.2	13
126	Assembly of a high-resolution map of the Acadian Usher syndrome region and localization of the nuclear EF-hand acidic gene. Biochimica Et Biophysica Acta - Molecular Basis of Disease, 1998, 1407, 84-91.	3.8	13

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127	Comparative analysis on the expression of L1 loci using various RNA-Seq preparations. Mobile DNA, 2020, 11, 2.	3.6	12
128	Organ-, sex-Âand age-dependent patterns of endogenous L1 mRNA expression at a single locus resolution. Nucleic Acids Research, 2021, 49, 5813-5831.	14.5	12
129	A mutation increasing the size of the polyoma virion proteins, VP2 and VP3. Virology, 1981, 109, 35-46.	2.4	11
130	An in vivo assay for measuring the recombination potential between DNA sequences in mammalian cells. Analytical Biochemistry, 1992, 205, 83-89.	2.4	11
131	Characterization and population diversity of interspersed repeat sequence variants (IRS-morphs). Genome, 1996, 39, 688-696.	2.0	11
132	Comparative studies of the CAG repeats in the spinocerebellar ataxia type 1 (SCA1) gene. , 1997, 74, 488-493.		11
133	Alu-linked hairpins efficiently mediate RNA interference with less toxicity than do H1-expressed short hairpin RNAs. Analytical Biochemistry, 2006, 349, 41-48.	2.4	11
134	Detection of LINE-1 RNAs by Northern Blot. Methods in Molecular Biology, 2016, 1400, 223-236.	0.9	11
135	Characterization and Phylogenetic Significance of a Repetitive DNA Sequence from Whooping Cranes (Grus americana). Auk, 1992, 109, 73-79.	1.4	9
136	Recent B2 Element Insertions in the Mouse Genome. DNA Sequence, 1998, 8, 343-348.	0.7	9
137	RNA Next-Generation Sequencing and a Bioinformatics Pipeline to Identify Expressed LINE-1s at the Locus-Specific Level. Journal of Visualized Experiments, 2019, , .	0.3	9
138	Altered DNA repair creates novel Alu/Alu repeatâ€mediated deletions. Human Mutation, 2021, 42, 600-613.	2.5	9
139	Analysis of epigenetic features characteristic of L1 loci expressed in human cells. Nucleic Acids Research, 2022, 50, 1888-1907.	14.5	9
140	LINE-1 and Alu retrotransposition exhibit clonal variation. Mobile DNA, 2013, 4, 16.	3.6	8
141	Sequence analysis and in vitro transcription of portions of the epstein-barr virus genome. Journal of Cellular Biochemistry, 1982, 19, 267-274.	2.6	7
142	Abasic sites and survival in resected patients with non-small cell lung cancer. Cancer Letters, 2007, 246, 47-53.	7.2	7
143	Simultaneous analysis of multiple gene expression patterns as a function of development, injury or senescence. Brain Research Protocols, 1998, 3, 1-6.	1.6	6
144	Breaking the computational barrier: a divide-conquer and aggregate based approach for Alu insertion site characterisation. International Journal of Computational Biology and Drug Design, 2009, 2, 302.	0.3	6

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145	Inferring the expression variability of human transposable element-derived exons by linear model analysis of deep RNA sequencing data. BMC Genomics, 2013, 14, 584.	2.8	6
146	The mouse deafness locus (dn) is associated with an inversion on chromosome 19. Biochimica Et Biophysica Acta - Molecular Basis of Disease, 1998, 1407, 257-262.	3.8	5
147	Long-Distance Relationships: Suppression of Repeat-Mediated Deletions. Trends in Genetics, 2018, 34, 572-574.	6.7	5
148	Response. Journal of Molecular Evolution, 1997, 45, 7-8.	1.8	3
149	Evolution of a Hypervariable Region of the Low Density Lipoprotein Receptor (LDLR) Gene in Humans and other Hominoids. Genetica, 2004, 121, 187-193.	1.1	3
150	Transcription coupled repair and biased insertion of human retrotransposon L1 in transcribed genes. Mobile DNA, 2017, 8, 18.	3.6	3
151	Alu Elements. , 2006, , 21-34.		3
152	A critical examination of possible fractionations of human DNA according to base composition. Nucleic Acids and Protein Synthesis, 1978, 520, 21-37.	1.7	2
153	The rat thymidine kinase gene 5â€² region: evolution of a promoter. DNA Sequence, 1991, 2, 129-131.	0.7	2
154	Structure and variability of recently inserted Alu family members. Nucleic Acids Research, 1991, 19, 698-698.	14.5	2
155	Evolution of B2 repeats: the muroid explosion. Genetica, 1997, 99, 1-13.	1.1	2
156	SINEs, LINEs and retrotransposable elements: Functional implications. Journal of Molecular Evolution, 1996, 42, 1-1.	1.8	1
157	Interspersed Repeat Insertion Polymorphisms for Studies of Human Molecular Anthropology. , 1999, , 201-212.		1
158	Structure, diversity, and evolution of the 45-bp VNTR in intron 5 of the USH1C gene. Genomics, 2004, 83, 439-444.	2.9	1
159	ANALYSIS OF CLONED HUMAN UBIQUITOUS REPEATED DNA SEQUENCES. , 1980, , 369-378.		1
160	Gene Transfer into Glial Cells using Herpes Simplex Virus Vectors. Annals of the New York Academy of Sciences, 1990, 605, 346-349.	3.8	0
161	Meeting Report for Mobile DNA 2010. Mobile DNA, 2010, 1, 20.	3.6	0
162	Alu-Alu Recombinations in Genetic Diseases. , 2017, , 239-257.		0

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163	2057 L1 expression analysis in adipose-derived stem cells. Journal of Clinical and Translational Science, 2018, 2, 16-16.	0.6	0
164	Recently integrated human Alu repeats: finding needles in the haystack. , 2000, , 149-161.		0
165	The Paracaspase MALT1 Acts Independently of Pre-B-Cell Receptor Signaling As a Key Factor in Leukemic Cell Survival in Precursor B-Cell Acute Lymphoblastic Leukemia. Blood, 2019, 134, 1288-1288.	1.4	0