## Giuseppe Biamonti

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Transcription of Satellite III non-coding RNAs is a general stress response in human cells. Nucleic Acids Research, 2008, 36, 423-434.	14.5	353
2	Cell Motility Is Controlled by SF2/ASF through Alternative Splicing of the Ron Protooncogene. Molecular Cell, 2005, 20, 881-890.	9.7	339
3	A novel procedure for quantitative polymerase chain reaction by coamplification of competitive templates. Gene, 1992, 122, 313-320.	2.2	260
4	Cellular response to etoposide treatment. Cancer Letters, 2007, 252, 9-18.	7.2	205
5	The roles of heterogeneous nuclear ribonucleoproteins (hnRNP) in RNA metabolism. BioEssays, 1996, 18, 747-756.	2.5	199
6	Nuclear Stress Bodies. Cold Spring Harbor Perspectives in Biology, 2010, 2, a000695-a000695.	5.5	188
7	Cellular stress and RNA splicing. Trends in Biochemical Sciences, 2009, 34, 146-153.	7.5	181
8	Fine mapping of a replication origin of human DNA Proceedings of the National Academy of Sciences of the United States of America, 1994, 91, 7119-7123.	7.1	180
9	DNA ligasel is recruited to sites of DNA replication by an interaction with proliferating cell nuclear antigen: identification of a common targeting mechanism for the assembly of replication factories. EMBO Journal, 1998, 17, 3786-3795.	7.8	179
10	hnRNP A1 Selectively Interacts Through its Gly-rich Domain with Different RNA-binding Proteins. Journal of Molecular Biology, 1996, 259, 337-348.	4.2	172
11	Molecular mechanisms of etoposide. EXCLI Journal, 2015, 14, 95-108.	0.7	172
12	Start Sites of Bidirectional DNA Synthesis at the Human Lamin B2 Origin. Science, 2000, 287, 2023-2026.	12.6	171
13	Transcriptional Activation of a Constitutive Heterochromatic Domain of the Human Genome in Response to Heat Shock. Molecular Biology of the Cell, 2004, 15, 543-551.	2.1	170
14	Stress-induced Nuclear Bodies Are Sites of Accumulation of Pre-mRNA Processing Factors. Molecular Biology of the Cell, 2001, 12, 3502-3514.	2.1	155
15	Alternative Splicing and Tumor Progression. Current Genomics, 2008, 9, 556-570.	1.6	152
16	Sam68 regulates EMT through alternative splicing–activated nonsense-mediated mRNA decay of the SF2/ASF proto-oncogene. Journal of Cell Biology, 2010, 191, 87-99.	5.2	146
17	cDNA cloning of human hnRNP protein Al reveals the existence of multiple mRNA isoforms. Nucleic Acids Research, 1988, 16, 3751-3770.	14.5	120
18	Structural and Functional Characterization of Noncoding Repetitive RNAs Transcribed in Stressed Human Cells. Molecular Biology of the Cell, 2005, 16, 2597-2604.	2.1	115

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19	The alternative splicing side of cancer. Seminars in Cell and Developmental Biology, 2014, 32, 30-36.	5.0	93
20	Human Chromosomes 9, 12, and 15 Contain the Nucleation Sites of Stress-Induced Nuclear Bodies. Molecular Biology of the Cell, 2002, 13, 2069-2079.	2.1	89
21	Oncogenic Alternative Splicing Switches: Role in Cancer Progression and Prospects for Therapy. International Journal of Cell Biology, 2013, 2013, 1-17.	2.5	89
22	Early mitotic degradation of the homeoprotein HOXC10 is potentially linked to cell cycle progression. EMBO Journal, 2003, 22, 3715-3724.	7.8	86
23	Modular Structure of the Human Lamin B2 Replicator. Molecular and Cellular Biology, 2004, 24, 2958-2967.	2.3	82
24	Nuclear stress bodies: a heterochromatin affair?. Nature Reviews Molecular Cell Biology, 2004, 5, 493-498.	37.0	82
25	Isolation of an active gene encoding human hnRNP protein A1. Journal of Molecular Biology, 1989, 207, 491-503.	4.2	81
26	Functional characterization of the T4 DNA ligase: a new insight into the mechanism of action. Nucleic Acids Research, 1997, 25, 2106-2113.	14.5	78
27	HnRNP A1 controls a splicing regulatory circuit promoting mesenchymal-to-epithelial transition. Nucleic Acids Research, 2013, 41, 8665-8679.	14.5	77
28	New insights into the auxiliary domains of eukaryotic RNA binding proteins. FEBS Letters, 1994, 340, 1-8.	2.8	73
29	The specific binding of nuclear protein(s) to the cAMP responsive element (CRE) sequence (TGACGTCA) is reduced by the misincorporation of U and increased by the deamination of C. Nucleic Acids Research, 1990, 18, 5775-5780.	14.5	72
30	Recombinant hnRNP protein A1 and its N-terminal domain show preferential affinity for oligodeoxynucleotides homologous to intron/exon acceptor sites. Nucleic Acids Research, 1990, 18, 6595-6600.	14.5	70
31	The replication factory targeting sequence/PCNA-binding site is required in G1 to control the phosphorylation status of DNA ligase I. EMBO Journal, 1999, 18, 5745-5754.	7.8	68
32	Pro-metastatic splicing of Ron proto-oncogene mRNA can be reversed: Therapeutic potential of bifunctional oligonucleotides and indole derivatives. RNA Biology, 2010, 7, 495-503.	3.1	65
33	From "Cellular―RNA to "Smart―RNA: Multiple Roles of RNA in Genome Stability and Beyond. Chemical Reviews, 2018, 118, 4365-4403.	47.7	63
34	Two homologous genes, originated by duplication, encode the human hnRNP proteins A2 and A1. Nucleic Acids Research, 1994, 22, 1996-2002.	14.5	61
35	Cell cycle modulation of protein-DNA interactions at a human replication origin. EMBO Journal, 1998, 17, 2961-2969.	7.8	58
36	Pre-mRNA processing factors meet the DNA damage response. Frontiers in Genetics, 2013, 4, 102.	2.3	58

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37	Making alternative splicing decisions during epithelial-to-mesenchymal transition (EMT). Cellular and Molecular Life Sciences, 2012, 69, 2515-2526.	5.4	56
38	Constitutive heterochromatin: a surprising variety of expressed sequences. Chromosoma, 2009, 118, 419-435.	2.2	55
39	Human hnRNP Protein A1 Gene Expression. Journal of Molecular Biology, 1993, 230, 77-89.	4.2	53
40	Mammalian single-stranded DNA binding proteins and heterogeneous nuclear RNA proteins have common antigenic determinants. Nucleic Acids Research, 1985, 13, 337-346.	14.5	52
41	DNA ligase I gene expression during differentiation and cell proliferation. Nucleic Acids Research, 1992, 20, 6209-6214.	14.5	51
42	In vivo protein-DNA interactions at human DNA replication origin Proceedings of the National Academy of Sciences of the United States of America, 1996, 93, 1498-1503.	7.1	50
43	The alternative splicing factor Nova2 regulates vascular development and lumen formation. Nature Communications, 2015, 6, 8479.	12.8	50
44	A single polypyrimidine tract binding protein (PTB) binding site mediates splicing inhibition at mouse IgM exons M1 and M2. Rna, 2004, 10, 787-794.	3.5	48
45	Transcription of Satellite DNAs in Mammals. Progress in Molecular and Subcellular Biology, 2011, 51, 95-118.	1.6	47
46	Subnuclear distribution of the largest subunit of the human origin recognition complex during the cell cycle. Journal of Cell Science, 2004, 117, 5221-5231.	2.0	46
47	Single stranded DNA binding proteins derive from hnRNP proteins by proteolysis in mammalian cells. Nucleic Acids Research, 1985, 13, 6577-6590.	14.5	45
48	Functional interactions of DNA topoisomerases with a human replication origin. EMBO Journal, 2007, 26, 998-1009.	7.8	45
49	A human DNA replication origin: localization and transcriptional characterization. Chromosoma, 1992, 102, S24-S31.	2.2	44
50	RNA recognition motif 2 directs the recruitment of SF2/ASF to nuclear stress bodies. Nucleic Acids Research, 2004, 32, 4127-4136.	14.5	44
51	Characterization of human DNA sequences synthesized at the onset of S-phase. Nucleic Acids Research, 1987, 15, 10211-10232.	14.5	43
52	Selection of homeotic proteins for binding to a human DNA replication origin 1 1Edited by M. Yaniv. Journal of Molecular Biology, 2000, 299, 667-680.	4.2	43
53	Cell cycle-dependent dynamic association of cyclin/Cdk complexes with human DNA replication proteins. EMBO Journal, 2002, 21, 2485-2495.	7.8	42
54	The Dispersal of Replication Proteins after Etoposide Treatment Requires the Cooperation of Nbs1 with the Ataxia Telangiectasia Rad3-Related/Chk1 Pathway. Cancer Research, 2006, 66, 1675-1683.	0.9	41

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55	DNA Ligase I Deficiency Leads to Replication-Dependent DNA Damage and Impacts Cell Morphology without Blocking Cell Cycle Progression. Molecular and Cellular Biology, 2009, 29, 2032-2041.	2.3	41
56	Fate of exogenous recombinant plasmids introduced into mouse and human cells. Nucleic Acids Research, 1985, 13, 5545-5561.	14.5	39
57	AlternativeÂsplicing in Alzheimer's disease. Aging Clinical and Experimental Research, 2021, 33, 747-758.	2.9	37
58	Cell Cycle-dependent Phosphorylation of Human DNA Ligase I at the Cyclin-dependent Kinase Sites. Journal of Biological Chemistry, 2003, 278, 37761-37767.	3.4	36
59	First dual AK/GSK-3β inhibitors endowed with antioxidant properties as multifunctional, potential neuroprotective agents. European Journal of Medicinal Chemistry, 2017, 138, 438-457.	5.5	33
60	Large-scale purification of hnRNP proteins from HeLa cells by affinity chromatography on ssDNA-cellulose. FEBS Journal, 1987, 162, 213-220.	0.2	31
61	Late induction of human DNA ligase I after UV-C irradiation. Nucleic Acids Research, 1995, 23, 962-966.	14.5	31
62	Phosphorylation of SRSF1 is modulated by replicational stress. Nucleic Acids Research, 2012, 40, 1106-1117.	14.5	31
63	Studies on the ATP Binding Site of Fyn Kinase for the Identification of New Inhibitors and Their Evaluation as Potential Agents against Tauopathies and Tumors. Journal of Medicinal Chemistry, 2015, 58, 4590-4609.	6.4	31
64	Etoposide Induces the Dispersal of DNA Ligase I from Replication Factories. Molecular Biology of the Cell, 2001, 12, 2109-2118.	2.1	29
65	Molecular and Structural Transactions at Human DNA Replication Origins. Cell Cycle, 2007, 6, 1705-1712.	2.6	28
66	Presence of transcription signals in two putative DNA replication origins of human cells. Biochimica Et Biophysica Acta Gene Regulatory Mechanisms, 1988, 951, 430-442.	2.4	27
67	Identification of autoantibodies to the I protein of the heterogeneous nuclear ribonucleoprotein complex in patients with systemic sclerosis. Arthritis and Rheumatism, 1996, 39, 1669-1676.	6.7	26
68	ls DNA sequence sufficient to specify DNA replication origins in metazoan cells?. Chromosome Research, 2003, 11, 403-412.	2.2	24
69	Homeotic proteins participate in the function of human-DNA replication origins. Nucleic Acids Research, 2010, 38, 8105-8119.	14.5	23
70	Sequence Determinants for hnRNP I Protein Nuclear Localization. Experimental Cell Research, 1997, 235, 300-304.	2.6	22
71	Growth-Dependent and Growth-Independent Translation of Messengers for Heterogeneous Nuclear Ribonucleoproteins. Nucleic Acids Research, 1997, 25, 3950-3954.	14.5	21
72	SRSF2 promotes splicing and transcription of exon 11 included isoform in Ron proto-oncogene. Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms, 2014, 1839, 1132-1140.	1.9	21

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73	An Intricate Connection between Alternative Splicing and Phenotypic Plasticity in Development and Cancer. Cells, 2020, 9, 34.	4.1	21
74	SAFB re-distribution marks steps of the apoptotic process. Experimental Cell Research, 2007, 313, 3914-3923.	2.6	20
75	A missense MT-ND5 mutation in differentiated Parkinson Disease cytoplasmic hybrid induces ROS-dependent DNA Damage Response amplified by DROSHA. Scientific Reports, 2017, 7, 9528.	3.3	20
76	hnRNP L inhibits CD44 V10 exon splicing through interacting with its upstream intron. Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms, 2015, 1849, 743-750.	1.9	19
77	Hybridization of oligodeoxynucleotide probes to RNA molecules: specificity and stability ofduplexes. Nucleic Acids Research, 1987, 15, 9091-9091.	14.5	18
78	Cloning and sequence analysis of a cDNA coding for the murine DNA ligase I enzyme. Gene, 1994, 144, 253-257.	2.2	17
79	DNA replication, development and cancer: a homeotic connection?. Critical Reviews in Biochemistry and Molecular Biology, 2010, 45, 14-22.	5.2	17
80	DNA ligase I is dephosphorylated during the execution step of etoposide-induced apoptosis. Cell Death and Differentiation, 2002, 9, 89-90.	11.2	16
81	Heat Shock Affects Mitotic Segregation of Human Chromosomes Bound to Stress-Induced Satellite III RNAs. International Journal of Molecular Sciences, 2020, 21, 2812.	4.1	15
82	Structural and functional heterogeneity of single-stranded DNA-binding proteins from calf thymus. Biochimica Et Biophysica Acta Gene Regulatory Mechanisms, 1984, 782, 147-155.	2.4	14
83	Assignment of the human heterogeneous nuclear ribonucleoprotein A1 gene (HNRPA1) to chromosome 12q13.1 by cDNA competitive in situ hybridization. Genomics, 1992, 12, 171-174.	2.9	14
84	Searching for replication origins in mammalian DNA. Gene, 1993, 135, 125-135.	2.2	14
85	Human hnRNP protein A1: A model polypeptide for a structural and genetic investigation of a broad family of RNA binding proteins. Genetica, 1994, 94, 101-114.	1.1	14
86	The Krebs Cycle Connection: Reciprocal Influence Between Alternative Splicing Programs and Cell Metabolism. Frontiers in Oncology, 2018, 8, 408.	2.8	14
87	A protein target site in an early replicated human DNA sequence: A highly conserved binding motif. Biochemical and Biophysical Research Communications, 1989, 165, 956-965.	2.1	13
88	DNA ligase I and Nbs1 proteins associate in a complex and colocalize at replication factories. Cell Cycle, 2009, 8, 2600-2607.	2.6	10
89	Reduced levels of prostaglandin I <sub>2</sub> synthase: a distinctive feature of the cancer-free trichothiodystrophy. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	8
90	A DNA dependent ATPase from HeLa cells. Biochemical and Biophysical Research Communications, 1982, 104, 402-409.	2.1	7

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91	Alternative Splicing and Cancer. Journal of Nucleic Acids, 2012, 2012, 1-2.	1.2	6
92	DNA-protein interaction dynamics at the Lamin B2 replication origin. Cell Cycle, 2015, 14, 64-73.	2.6	6
93	Alternative Splicing of Tumor Suppressors and Oncogenes. Cancer Treatment and Research, 2013, 158, 95-117.	0.5	6
94	CorrelaGenes: a new tool for the interpretation of the human transcriptome. BMC Bioinformatics, 2014, 15, S6.	2.6	4
95	Chronic Replication Problems Impact Cell Morphology and Adhesion of DNA Ligase I Defective Cells. PLoS ONE, 2015, 10, e0130561.	2.5	4
96	A 2-nt RNA enhancer on exon 11 promotes exon 11 inclusion of the Ron proto-oncogene. Oncology Reports, 2014, 31, 450-455.	2.6	3
97	A repeated element in the human lamin B2 gene covers most of an intron and reiterates the exon/intron junction. Gene, 1997, 196, 267-277.	2.2	1
98	CorrelaGenes: a new tool for the interpretation of the human transcriptome. EMBnet Journal, 2012, 18, 103.	0.6	1