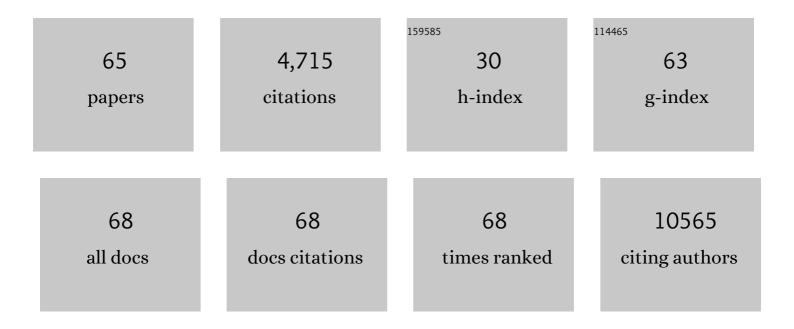
Lauro Sumoy

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Genomewide Association Study of Severe Covid-19 with Respiratory Failure. New England Journal of Medicine, 2020, 383, 1522-1534.	27.0	1,548
2	A β-catenin/XTcf-3 complex binds to the <i>siamois</i> promoter to regulate dorsal axis specification in <i>Xenopus</i> . Genes and Development, 1997, 11, 2359-2370.	5.9	494
3	The MAPK Hog1 recruits Rpd3 histone deacetylase to activate osmoresponsive genes. Nature, 2004, 427, 370-374.	27.8	295
4	A myriad of miRNA variants in control and Huntington's disease brain regions detected by massively parallel sequencing. Nucleic Acids Research, 2010, 38, 7219-7235.	14.5	270
5	Local Tissue Interactions across the Dorsal Midline of the Forebrain Establish CNS Laterality. Neuron, 2003, 39, 423-438.	8.1	175
6	Hnf1α (MODY3) Controls Tissue-Specific Transcriptional Programs and Exerts Opposed Effects on Cell Growth in Pancreatic Islets and Liver. Molecular and Cellular Biology, 2009, 29, 2945-2959.	2.3	122
7	Expression of the chicken homeobox-containing gene GHox-8 during embryonic chick limb development. Mechanisms of Development, 1991, 34, 143-154.	1.7	104
8	Curcumin mediates oxaliplatin-acquired resistance reversion in colorectal cancer cell lines through modulation of CXC-Chemokine/NF-κB signalling pathway. Scientific Reports, 2016, 6, 24675.	3.3	103
9	A role for notochord in axial vascular development revealed by analysis of phenotype and the expression of VEGR-2 in zebrafish flh and ntl mutant embryos. Mechanisms of Development, 1997, 63, 15-27.	1.7	93
10	The expression pattern of the Distal-less homeoâ~•containing gene Dlx-5 in the developing chick limb bud suggests its involvement in apical ectodermal ridge activity, pattern formation, and cartilage differentiation. Mechanisms of Development, 1995, 52, 257-264.	1.7	92
11	MacroH2A1 Regulates the Balance between Self-Renewal and Differentiation Commitment in Embryonic and Adult Stem Cells. Molecular and Cellular Biology, 2012, 32, 1442-1452.	2.3	86
12	Epigenetic footprint enables molecular risk stratification of hepatoblastoma with clinical implications. Journal of Hepatology, 2020, 73, 328-341.	3.7	82
13	LRRN6A/LERN1 (leucine-rich repeat neuronal protein 1), a novel gene with enriched expression in limbic system and neocortex. European Journal of Neuroscience, 2003, 18, 3167-3182.	2.6	76
14	Genome-wide differences between microsatellite stable and unstable colorectal tumors. Carcinogenesis, 2006, 27, 419-428.	2.8	66
15	Regulation of GABAA and Glutamate Receptor Expression, Synaptic Facilitation and Long-Term Potentiation in the Hippocampus of Prion Mutant Mice. PLoS ONE, 2009, 4, e7592.	2.5	60
16	Additional Complexity on Human Chromosome 15q: Identification of a Set of Newly Recognized Duplicons (LCR15) on 15q11–q13, 15q24, and 15q26. Genome Research, 2001, 11, 98-111.	5.5	60
17	Human vitamin K-dependentGAS6: Gene structure, allelic variation, and association with stroke. Human Mutation, 2004, 23, 506-512.	2.5	54
18	GHox-7: A chicken homeobox-containing gene expressed in a fashion consistent with a role in patterning events during embryonic chick limb development. Differentiation, 1992, 49, 85-92.	1.9	46

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19	A comprehensive custom panel design for routine hereditary cancer testing: preserving control, improving diagnostics and revealing a complex variation landscape. Scientific Reports, 2017, 7, 39348.	3.3	45
20	Developmental analysis of Lingoâ€1/Lern1 protein expression in the mouse brain: Interaction of its intracellular domain with Myt11. Developmental Neurobiology, 2008, 68, 521-541.	3.0	44
21	Identification of a Spatially Specific Enhancer Element in the Chicken Msx-2 Gene That Regulates Its Expression in the Apical Ectodermal Ridge of the Developing Limb Buds of Transgenic Mice. Developmental Biology, 1995, 170, 230-242.	2.0	42
22	Transcriptional regulation by Poly(ADP-ribose) polymerase-1 during T cell activation. BMC Genomics, 2008, 9, 171.	2.8	42
23	Cyclic AMP signaling restricts activation and promotes maturation and antioxidant defenses in astrocytes. BMC Genomics, 2016, 17, 304.	2.8	42
24	Role of <i>Saccharomyces cerevisiae</i> Oxidoreductases Bdh1p and Ara1p in the Metabolism of Acetoin and 2,3-Butanediol. Applied and Environmental Microbiology, 2010, 76, 670-679.	3.1	38
25	HMG20A and HMG20B map to human chromosomes 15q24 and 19p13.3 and constitute a distinct class of HMG-box genes with ubiquitous expression. Cytogenetic and Genome Research, 2000, 88, 62-67.	1.1	36
26	CD40: an upstream master switch for endothelial cell activation uncovered by RNAi-coupled transcriptional profiling. Blood, 2008, 112, 3624-3637.	1.4	36
27	Expression of CD11c Is Associated with Unconventional Activated T Cell Subsets with High Migratory Potential. PLoS ONE, 2016, 11, e0154253.	2.5	36
28	Conservation of intracellular Wnt signaling components in dorsal-ventral axis formation in zebrafish. Development Genes and Evolution, 1999, 209, 48-58.	0.9	34
29	Gene expression profiling distinguishes JAK2V617F-negative from JAK2V617F-positive patients in essential thrombocythemia. Leukemia, 2008, 22, 1368-1376.	7.2	34
30	Microarray and deep sequencing cross-platform analysis of the mirRNome and isomiR variation in response to epidermal growth factor. BMC Genomics, 2013, 14, 371.	2.8	33
31	GCAT Genomes for life: a prospective cohort study of the genomes of Catalonia. BMJ Open, 2018, 8, e018324.	1.9	31
32	MicroRNA 22 Regulates Cell Cycle Length in Cerebellar Granular Neuron Precursors. Molecular and Cellular Biology, 2013, 33, 2706-2717.	2.3	29
33	Developmental Expression of the Oligodendrocyte Myelin Glycoprotein in the Mouse Telencephalon. Cerebral Cortex, 2010, 20, 1769-1779.	2.9	28
34	Multitrait genome association analysis identifies new susceptibility genes for human anthropometric variation in the GCAT cohort. Journal of Medical Genetics, 2018, 55, 765-778.	3.2	28
35	PACSIN 3 is a novel SH3 domain cytoplasmic adapter protein of the pacsin-syndapin-FAP52 gene family. Gene, 2001, 262, 199-205.	2.2	25
36	Hypothalamus transcriptome profile suggests an anorexia-cachexia syndrome in the anx/anx mouse model. Physiological Genomics, 2008, 35, 341-350.	2.3	22

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37	Partially Degraded RNA from Bladder Washing is a Suitable Sample for Studying Gene Expression Profiles in Bladder Cancer. European Urology, 2006, 50, 1347-1356.	1.9	19
38	Multiple platform assessment of the EGF dependent transcriptome by microarray and deep tag sequencing analysis. BMC Genomics, 2011, 12, 326.	2.8	19
39	Cloning, expression, and mapping of PDCD9, the human homolog of <i>Gallus gallus</i> pro-apoptotic protein p52. Cytogenetic and Genome Research, 1999, 87, 85-88.	1.1	18
40	Genomic imbalances in Schistosoma-associated and non–Schistosoma-associated bladder carcinoma. An array comparative genomic hybridization analysis. Cancer Genetics and Cytogenetics, 2007, 177, 16-19.	1.0	18
41	Cell cycle control pathways act as conditioning factors for TK/GCV sensitivity in pancreatic cancer cells. Biochimica Et Biophysica Acta - Molecular Cell Research, 2010, 1803, 1175-1185.	4.1	18
42	Identification and characterization of BTBD1 , a novel BTB domain containing gene on human chromosome 15q24. Gene, 2001, 262, 275-281.	2.2	15
43	Identification and characterization of UBXD1, a novel UBX domain-containing gene on human chromosome 19p13, and its mouse ortholog. Biochimica Et Biophysica Acta Gene Regulatory Mechanisms, 2001, 1517, 298-301.	2.4	14
44	Loss of HDAC11 accelerates skeletal muscle regeneration in mice. FEBS Journal, 2021, 288, 1201-1223.	4.7	14
45	Aberrant brain microRNA target and miRISC gene expression in the anx/anx anorexia mouse model. Gene, 2012, 497, 181-190.	2.2	12
46	Draft Genome Sequences of Mycobacterium setense Type Strain DSM-45070 and the Nonpathogenic Strain Manresensis, Isolated from the Bank of the Cardener River in Manresa, Catalonia, Spain. Genome Announcements, 2015, 3, .	0.8	11
47	PALML, a novel paralemmin-related gene mapping on human chromosome 1p21. Gene, 2001, 278, 33-40.	2.2	10
48	ATB0/SLC1A5 gene. Fine localisation and exclusion of association with the intestinal phenotype of cystic fibrosis. European Journal of Human Genetics, 2001, 9, 860-866.	2.8	10
49	Genomic profiling in advanced stage non-small-cell lung cancer patients with platinum-based chemotherapy identifies germline variants with prognostic value in SMYD2. Cancer Treatment and Research Communications, 2018, 15, 21-31.	1.7	9
50	Identification and Expression Analysis of C3orf1, a Novel Human Gene Homologous to the Drosophila RP140—Upstream Gene. DNA Sequence, 2000, 11, 335-338.	0.7	8
51	Characterization of human FSD1, a novel brain specific gene on chromosome 19 with paralogy to 9q31. Biochimica Et Biophysica Acta Gene Regulatory Mechanisms, 2001, 1518, 200-203.	2.4	8
52	Large-scale screening of circulating microRNAs in individuals with HIV-1 mono-infections reveals specific liver damage signatures. Antiviral Research, 2018, 155, 106-114.	4.1	8
53	The yeast ζâ€crystallin/NADPH:quinone oxidoreductase (Zta1p) is under nutritional control by the target of rapamycin pathway and is involved in the regulation of argininosuccinate lyase <scp>mRNA</scp> halfâ€life. FEBS Journal, 2015, 282, 1953-1964.	4.7	6
54	GCAT Panel, a comprehensive structural variant haplotype map of the Iberian population from high-coverage whole-genome sequencing. Nucleic Acids Research, 2022, 50, 2464-2479.	14.5	6

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55	Msx2 Expression in the Apical Ectoderm Ridge Is Regulated by an Msx2 and Dlx5 Binding Site. Biochemical and Biophysical Research Communications, 2002, 290, 955-961.	2.1	5
56	Gene expression changes in blood RNA after swimming in a chlorinated pool. Journal of Environmental Sciences, 2017, 58, 250-261.	6.1	5
57	<psa as="" in="" kinetics="" markers="" metastatic<br="" of="" overall="" patients="" prognostic="" survival="" with="">Castration-Resistant Prostate Cancer Treated with Abiraterone Acetate. Cancer Management and Research, 2020, Volume 12, 10251-10260.</psa>	1.9	5
58	Paired-end small RNA sequencing reveals a possible overestimation in the isomiR sequence repertoire previously reported from conventional single read data analysis. BMC Bioinformatics, 2021, 22, 215.	2.6	5
59	Characterization of Acute HCV Infection and Transmission Networks in People Who Currently Inject Drugs in Catalonia: Usefulness of Dried Blood Spots. Hepatology, 2021, 74, 591-606.	7.3	5
60	A likelihood ratio approach for identifying three-quarter siblings in genetic databases. Heredity, 2021, 126, 537-547.	2.6	5
61	Adhesion Molecules Associated with Female Genital Tract Infection. PLoS ONE, 2016, 11, e0156605.	2.5	4
62	Cloning, Mapping and Expression Analysis of C15 or f4, a Novel Human Gene with Homology to the Yeast Mitochondrial Ribosomal Protein Yml30 Gene. DNA Sequence, 2001, 12, 91-96.	0.7	2
63	Identification of C15orf5, a Heart-Enriched Transcript on Chromosome 15q23-q24. DNA Sequence, 2001, 12, 67-69.	0.7	1
64	P6: Detection ofÂpathologic andÂpolymorphic cryptic chromosomal imbalances byÂarray-CGH using aÂBAC-array targeting regions between segmental duplications. European Journal of Medical Genetics, 2005, 48, 451-452.	1.3	0
65	Gene Expression Profiling Distinguishes Essential Thrombocythemia from Polycythemia Vera Patients and Identifies a Common Expressed Set of Genes in Relation to JAK2V617F Status. Blood, 2008, 112,	1.4	0