Sara Larriba

List of Publications by Year
in descending order

37 papers	1,546 citations	361296 20 h-index	360920 35 g-index
37 all docs	37 docs citations	37 times ranked	2071 citing authors

#	Article	IF	Citations
1	Common Variation in the PIN1 Locus Increases the Genetic Risk to Suffer from Sertoli Cell-Only Syndrome. Journal of Personalized Medicine, 2022, 12, 932.	1.1	0
2	Common genetic variation in KATNAL1 nonâ€coding regions is involved in the susceptibility to severe phenotypes of male infertility. Andrology, 2022, 10, 1339-1350.	1.9	5
3	Extracellular vesicle ncRNAs in seminal plasma as biomarkers for nonobstructive azoospermia. Human Reproduction, 2021, 36, 1452-1452.	0.4	2
4	Effect and in silico characterization of genetic variants associated with severe spermatogenic disorders in a large Iberian cohort. Andrology, 2021, 9, 1151-1165.	1.9	12
5	Evaluation of Male Fertility-Associated Loci in a European Population of Patients with Severe Spermatogenic Impairment. Journal of Personalized Medicine, 2021, 11, 22.	1.1	10
6	Intronic variation of the SOHLH2 gene confers risk to male reproductive impairment. Fertility and Sterility, 2020, 114, 398-406.	0.5	9
7	Impact of Extracellular Vesicle Isolation Methods on Downstream miRNA Analysis in Semen: A Comparative Study. International Journal of Molecular Sciences, 2020, 21, 5949.	1.8	27
8	Seminal plasma microRNAs improve diagnosis/prognosis of prostate cancer in men with moderately altered prostate-specific antigen. American Journal of Translational Research (discontinued), 2020, 12, 2041-2051.	0.0	6
9	Semen miRNAs Contained in Exosomes as Non-Invasive Biomarkers for Prostate Cancer Diagnosis. Scientific Reports, 2019, 9, 13772.	1.6	92
10	Exosomal microRNAs in seminal plasma are markers of the origin of azoospermia and can predict the presence of sperm in testicular tissue. Human Reproduction, 2018, 33, 1087-1098.	0.4	131
11	Altered Transcriptomic Profiles Associated With Male Infertility. , 2018, , 125-145.		0
12	Altered miRNA Signature of Developing Germ-cells in Infertile Patients Relates to the Severity of Spermatogenic Failure and Persists in Spermatozoa. Scientific Reports, 2015, 5, 17991.	1.6	72
13	Aberrant DNA methylation patterns of spermatozoa in men with unexplained infertility. Human Reproduction, 2015, 30, 1014-1028.	0.4	144
14	Epigenetic loss of the PIWI/piRNA machinery in human testicular tumorigenesis. Epigenetics, 2014, 9, 113-118.	1.3	87
15	Assessing the residual CFTR gene expression in human nasal epithelium cells bearing CFTR splicing mutations causing cystic fibrosis. European Journal of Human Genetics, 2014, 22, 784-791.	1.4	24
16	Altered gene expression signature of early stages of the germ line supports the preâ€meiotic origin of human spermatogenic failure. Andrology, 2014, 2, 596-606.	1.9	13
17	Association of PIWIL4 genetic variants with germ cell maturation arrest in infertile Spanish men. Asian Journal of Andrology, 2014, 16, 931.	0.8	9
18	GUSB and ATP2B4 are suitable reference genes for CFTR gene expression data normalization in nasal epithelium cells. Journal of Cystic Fibrosis, 2012, 11, 398-404.	0.3	4

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19	Sperm gene expression profile is related to pregnancy rate after insemination and is predictive of low fecundity in normozoospermic men. Human Reproduction, 2012, 27, 1556-1567.	0.4	46
20	Epigenetic Disruption of the PIWI Pathway in Human Spermatogenic Disorders. PLoS ONE, 2012, 7, e47892.	1.1	94
21	Changes in the Expression Profile of the Meiosisâ€Involved Mismatch Repair Genes in Impaired Human Spermatogenesis. Journal of Andrology, 2010, 31, 346-357.	2.0	45
22	N-terminal CFTR missense variants severely affect the behavior of the CFTR chloride channel. Human Mutation, 2008, 29, 738-749.	1.1	25
23	Mutations of SYCP3 are rare in infertile Spanish men with meiotic arrest. Fertility and Sterility, 2007, 88, 988-989.	0.5	23
24	Single nucleotide polymorphisms in succinate dehydrogenase subunits and citrate synthase genes: association results for impaired spermatogenesis. Journal of Developmental and Physical Disabilities, 2007, 30, 144-152.	3.6	8
25	NF1mutation rather than individual genetic variability is the main determinant of theNF1-transcriptional profile of mutations affecting splicing. Human Mutation, 2006, 27, 1104-1114.	1.1	23
26	Molecular evaluation of CFTR sequence variants in male infertility of testicular origin. Journal of Developmental and Physical Disabilities, 2005, 28, 284-290.	3.6	26
27	Suitability of oligonucleotide-mediated cystic fibrosis gene repair in airway epithelial cells. Journal of Gene Medicine, 2003, 5, 625-639.	1.4	5
28	Non-viral vector-mediated uptake, distribution, and stability of chimeraplasts in human airway epithelial cells. Journal of Gene Medicine, 2002, 4, 308-322.	1.4	15
29	Quantitative assessment of chimeraplast stability in biological fluids by polyacrylamide gel electrophoresis and laser-assisted fluorescence analysis. Pharmaceutical Research, 2002, 19, 914-918.	1.7	1
30	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.	1.4	10
31	Adenosine Triphosphate-Binding Cassette Superfamily Transporter Gene Expression in Severe Male Infertility1. Biology of Reproduction, 2001, 65, 394-400.	1.2	20
32	Heterogeneity for mutations in the CFTR gene and clinical correlations in patients with congenital absence of the vas deferens. Human Reproduction, 2000, 15, 1476-1483.	0.4	131
33	Detection of a cystic fibrosis modifier locus for meconium ileus on human chromosome 19q13. Nature Genetics, 1999, 22, 128-129.	9.4	216
34	Testicular CFTR Splice Variants in Patients with Congenital Absence of the Vas Deferens. Human Molecular Genetics, 1998, 7, 1739-1744.	1.4	74
35	Sertoli cell-specific expression of rat androgen-binding protein in transgenic mice: effects on somatic cell lineages. Molecular and Cellular Endocrinology, 1997, 132, 127-136.	1.6	18
36	High heterogeneity for cystic fibrosis in Spanish families: 75 mutations account for 90% of chromosomes. Human Genetics, 1997, 101, 365-370.	1.8	94

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37	Androgen binding protein is tissue-specifically expressed and biologically active in transgenic mice. Journal of Steroid Biochemistry and Molecular Biology, 1995, 53, 573-578.	1.2	25