Isabel Barragan

List of Publications by Year
in descending order

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37 papers	1,483 citations	471509 17 h-index	395702 33 g-index
39 all docs	39 docs citations	39 times ranked	2899 citing authors

#	Article	IF	Citations
1	Emerging noninvasive methylation biomarkers of cancer prognosis and drug response prediction. Seminars in Cancer Biology, 2022, 83, 584-595.	9.6	18
2	CCT3-LINC00326axis regulates hepatocarcinogenic lipid metabolism. Gut, 2022, 71, 2081-2092.	12.1	32
3	1326P Combination of stereotactic ablative radiotherapy with anti-PD-1 in oligoprogressive disease: Final results of a prospective multicenter study. Annals of Oncology, 2021, 32, S1016-S1017.	1.2	1
4	Epigenetic biomarkers of disease. , 2021, , 117-141.		0
5	Liquid Biopsy as a Tool for the Characterisation and Early Detection of the Field Cancerization Effect in Patients with Oral Cavity Carcinoma. Biomedicines, 2021, 9, 1478.	3.2	3
6	EDIL3 promotes epithelialâ€“mesenchymal transition and paclitaxel resistance through its interaction with integrin Î±VÎ²3 in cancer cells. Cell Death Discovery, 2020, 6, 86.	4.7	29
7	Epigenetics modulates the complexity of the response to Immune Checkpoint Blockade. EBioMedicine, 2020, 60, 103005.	6.1	2
8	Resistance to Neoadjuvant Treatment in Breast Cancer: Clinicopathological and Molecular Predictors. Cancers, 2020, 12, 2012.	3.7	13
9	Genetic and Epigenetic Biomarkers of Immune Checkpoint Blockade Response. Journal of Clinical Medicine, 2020, 9, 286.	2.4	50
10	CYP3A5 is unlikely to mediate anticancer drug resistance in hepatocellular carcinoma. Pharmacogenomics, 2019, 20, 1085-1092.	1.3	2
11	Pharmacoepigenetics and Toxicoepigenetics: Novel Mechanistic Insights and Therapeutic Opportunities. Annual Review of Pharmacology and Toxicology, 2018, 58, 161-185.	9.4	45
12	Epigenetic prediction of response to anti-PD-1 treatment in non-small-cell lung cancer: a multicentre, retrospective analysis. Lancet Respiratory Medicine,the, 2018, 6, 771-781.	10.7	167
13	Single base resolution analysis of 5-hydroxymethylcytosine in 188 human genes: implications for hepatic gene expression. Nucleic Acids Research, 2016, 44, 6756-6769.	14.5	15
14	Cytostatic Effect of Repeated Exposure to Simvastatin: A Mechanism for Chronic Myotoxicity Revealed by the Use of Mesodermal Progenitors Derived from Human Pluripotent Stem Cells. Stem Cells, 2015, 33, 2936-2948.	3.2	11
15	Genetic and epigenetic regulation of gene expression in fetal and adult human livers. BMC Genomics, 2014, 15, 860.	2.8	124
16	Long-Term Chronic Toxicity Testing Using Human Pluripotent Stem Cellâ€“Derived Hepatocytes. Drug Metabolism and Disposition, 2014, 42, 1401-1406.	3.3	87
17	Epigenetic mechanisms of importance for drug treatment. Trends in Pharmacological Sciences, 2014, 35, 384-396.	8.7	129
18	CYP2W1 polymorphism: functional aspects and relation to risk for colorectal cancer. Pharmacogenomics, 2013, 14, 1615-1622.	1.3	9

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19	Ontogeny, distribution and potential roles of 5-hydroxymethylcytosine in human liver function. Genome Biology, 2013, 14, R83.	9.6	61
20	Strong effects of environmental factors on prevalence and course of major depressive disorder are not moderated by 5-HTTLPR polymorphisms in a large Dutch sample. Journal of Affective Disorders, 2013, 146, 91-99.	4.1	26
21	Mutation Screening of Multiple Genes in Spanish Patients with Autosomal Recessive Retinitis Pigmentosa by Targeted Resequencing. PLoS ONE, 2011, 6, e27894.	2.5	36
22	Copy-Number Variations inEYS:A Significant Event in the Appearance of arRP. , 2011, 52, 5625.		40
23	EYS is a major gene for rod-cone dystrophies in France. Human Mutation, 2010, 31, E1406-E1435.	2.5	86
24	Mutation spectrum of EYS in Spanish patients with autosomal recessive retinitis pigmentosa. Human Mutation, 2010, 31, E1772-E1800.	2.5	69
25	Identification of Novel Mutations in the Ortholog ofDrosophilaEyes Shut Gene (EYS) Causing Autosomal Recessive Retinitis Pigmentosa. , 2010, 51, 4266.		57
26	Microarray-Based Mutation Analysis of 183 Spanish Families with Usher Syndrome. , 2010, 51, 1311.		57
27	Genetic Analysis of FAM46A in Spanish Families with Autosomal Recessive Retinitis Pigmentosa: Characterisation of Novel VNTRs. Annals of Human Genetics, 2008, 72, 26-34.	0.8	29
28	EYS, encoding an ortholog of Drosophila spacemaker, is mutated in autosomal recessive retinitis pigmentosa. Nature Genetics, 2008, 40, 1285-1287.	21.4	175
29	Linkage Validation of RP25 Using the 10K GeneChip Array and Further Refinement of the Locus by New Linked Families. Annals of Human Genetics, 2008, 72, 454-462.	0.8	17
30	Large-scale Molecular Analysis of a 34 Mb Interval on Chromosome 6q: Major Refinement of the RP25 Interval. Annals of Human Genetics, 2008, 72, 463-477.	0.8	6
31	A Novel Genetic Study of Chinese Families with Autosomal Recessive Retinitis Pigmentosa. Annals of Human Genetics, 2007, 71, 281-294.	0.8	16
32	Effects of incorporated drugs on degradation of novel 2,2â€²-bis(2-oxazoline) linked poly(lactic acid) films. International Journal of Pharmaceutics, 2006, 310, 162-167.	5.2	13
33	Exclusion of Four Candidate Genes, KHDRBS2, PTP4A1, KIAA1411 and OGFRL1, as Causative of Autosomal Recessive Retinitis Pigmentosa. Ophthalmic Research, 2006, 38, 19-23.	1.9	10
34	Molecular Analysis of <i>RIM1</i> in Autosomal Recessive Retinitis pigmentosa. Ophthalmic Research, 2005, 37, 89-93.	1.9	18
35	Mutation screening of three candidate genes, ELOVL5, SMAP1 and GLULD1 in autosomal recessive retinitis pigmentosa. International Journal of Molecular Medicine, 2005, 16, 1163.	4.0	5
36	Molecular Genetic Analysis of Two Functional Candidate Genes in the Autosomal Recessive Retinitis Pigmentosa, RP25, Locus. Current Eye Research, 2005, 30, 1081-1087.	1.5	8

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37	Mutation screening of three candidate genes, ELOVL5, SMAP1 and GLULD1 in autosomal recessive retinitis pigmentosa. International Journal of Molecular Medicine, 2005, 16, 1163-7.	4.0	17