Sergio Alonso

List of Publications by Year in descending order

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361413 377865 1,227 44 20 34 citations h-index g-index papers 45 45 45 2132 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Global DNA demethylation in gastrointestinal cancer is age dependent and precedes genomic damage. Cancer Cell, 2006, 9, 199-207.	16.8	245
2	Genetic and Functional Analysis of the Styrene Catabolic Cluster of <i>Pseudomonas</i> sp. Strain Y2. Journal of Bacteriology, 1998, 180, 1063-1071.	2.2	138
3	Lung tumorspheres reveal cancer stem cell-like properties and a score with prognostic impact in resected non-small-cell lung cancer. Cell Death and Disease, 2019, 10, 660.	6.3	68
4	DNA demethylation in normal colon tissue predicts predisposition to multiple cancers. Oncogene, 2012, 31, 5029-5037.	5.9	52
5	Liver CPT1A gene therapy reduces dietâ€induced hepatic steatosis in mice and highlights potential lipid biomarkers for human NAFLD. FASEB Journal, 2020, 34, 11816-11837.	0.5	44
6	Novel miRNA-mRNA interactions conserved in essential cancer pathways. Scientific Reports, 2017, 7, 46101.	3.3	38
7	Characterization of a second functional gene cluster for the catabolism of phenylacetic acid in Pseudomonas sp. strain Y2. Gene, 2004, 341, 167-179.	2.2	37
8	DNA methylation and chromatin modifiers in colorectal cancer. Molecular Aspects of Medicine, 2019, 69, 73-92.	6.4	34
9	A novel long non-coding RNA from NBL2 pericentromeric macrosatellite forms a perinucleolar aggregate structure in colon cancer. Nucleic Acids Research, 2018, 46, 5504-5524.	14.5	30
10	Coregulation by Phenylacetyl-Coenzyme A-Responsive PaaX Integrates Control of the Upper and Lower Pathways for Catabolism of Styrene by Pseudomonas sp. Strain Y2. Journal of Bacteriology, 2006, 188, 4812-4821.	2.2	29
11	Methylation of <i>MGMT</i> and <i>ADAMTS14</i> iin normal colon mucosa: biomarkers of a field defect for cancerization preferentially targeting elder African-Americans. Oncotarget, 2015, 6, 3420-3431.	1.8	29
12	Genetic characterization of the styrene lower catabolic pathway of Pseudomonas sp. strain Y2. Gene, 2003, 319, 71-83.	2.2	28
13	MicroRNA profiling associated with non-small cell lung cancer: next generation sequencing detection, experimental validation, and prognostic value. Oncotarget, 2017, 8, 56143-56157.	1.8	28
14	A Colorectal Cancer Susceptibility New Variant at 4q26 in the Spanish Population Identified by Genome-Wide Association Analysis. PLoS ONE, 2014, 9, e101178.	2.5	26
15	Somatic DNA Hypomethylation in H. pylori -Associated High-Risk Gastritis and Gastric Cancer: Enhanced Somatic Hypomethylation Associates with Advanced Stage Cancer. Clinical and Translational Gastroenterology, 2015, 6, e85.	2.5	26
16	Construction of a bacterial biosensor for styrene. Journal of Biotechnology, 2003, 102, 301-306.	3.8	25
17	DNA methylation alterations of AXIN2 in serrated adenomas and colon carcinomas with microsatellite instability. BMC Cancer, 2014, 14, 466.	2.6	25
18	Epigenetic inactivation of the extracellular matrix metallopeptidase ADAMTS19 gene and the metastatic spread in colorectal cancer. Clinical Epigenetics, 2015, 7, 124.	4.1	24

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19	Clinical characteristics of synchronous colorectal cancers in Japan. World Journal of Surgical Oncology, 2016, 14, 272.	1.9	24
20	Requirement of RIZ1 for Cancer Prevention by Methyl-Balanced Diet. PLoS ONE, 2008, 3, e3390.	2.5	24
21	Array-based identification of common DNA methylation alterations in ulcerative colitis. International Journal of Oncology, 2012, 40, 983-994.	3.3	21
22	DNA fingerprinting techniques for the analysis of genetic and epigenetic alterations in colorectal cancer. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2010, 693, 61-76.	1.0	19
23	Helicase Lymphoid-Specific Enzyme Contributes to the Maintenance of Methylation of SST1 Pericentromeric Repeats That Are Frequently Demethylated in Colon Cancer and Associate with Genomic Damage. Epigenomes, 2017, 1, 2.	1.8	19
24	New insights into the genome of Rhodococcus ruber strain Chol-4. BMC Genomics, 2019, 20, 332.	2.8	19
25	Tumor size is an independent risk predictor for metachronous colorectal cancer. Oncotarget, 2016, 7, 17896-17904.	1.8	18
26	Down-regulation of <i>Epidermal Growth Factor Receptor</i> by Selective Expansion of a 5′-End Regulatory Dinucleotide Repeat in Colon Cancer with Microsatellite Instability. Clinical Cancer Research, 2009, 15, 4531-4537.	7.0	16
27	Interplay between post-translational cyclooxygenase-2 modifications and the metabolic and proteomic profile in a colorectal cancer cohort. World Journal of Gastroenterology, 2019, 25, 433-446.	3.3	16
28	Microsatellite instability and ploidy status define three categories with distinctive prognostic impact in endometrioid endometrial cancer. Oncotarget, 2014, 5, 6206-6217.	1.8	16
29	Design of catabolic cassettes for styrene biodegradation. Antonie Van Leeuwenhoek, 2003, 84, 17-24.	1.7	15
30	Quantification of Unmethylated Alu (QUAlu): a tool to assess global hypomethylation in routine clinical samples. Oncotarget, 2016, 7, 10536-10546.	1.8	14
31	Curative resection for leiomyosarcoma of the descending mesocolon with metachronous liver metastasis: A case report and literature review. Molecular and Clinical Oncology, 2016, 5, 53-56.	1.0	12
32	HOXD8 hypermethylation as a fully sensitive and specific biomarker for biliary tract cancer detectable in tissue and bile samples. British Journal of Cancer, 2022, 126, 1783-1794.	6.4	12
33	Frequent somatic demethylation of RAPGEF1/C3G intronic sequences in gastrointestinal and gynecological cancer. International Journal of Oncology, 2011, 38, 1575-7.	3.3	11
34	Malignant, but not benign, intraductal papillary mucinous neoplasm preferentially associates with prior extrapancreatic malignancies. Oncology Reports, 2016, 35, 3236-3240.	2.6	11
35	Draft Genome Sequence of the Steroid Degrader Rhodococcus ruber Strain Chol-4. Genome Announcements, 2013, $1,\dots$	0.8	8
36	Epigenetic and transcriptional dysregulation of VWA2 associated with a MYC-driven oncogenic program in colorectal cancer. Scientific Reports, 2018, 8, 11097.	3.3	7

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37	Production of a recombinant form of the propeptide NH2-terminal of the precursor of pulmonary surfactant protein B. Enzyme and Microbial Technology, 2006, 40, 85-92.	3.2	5
38	Somatic Hypomethylation of Pericentromeric SST1 Repeats and Tetraploidization in Human Colorectal Cancer Cells. Cancers, 2021, 13, 5353.	3.7	5
39	Methylation-Sensitive Amplification Length Polymorphism (MS-AFLP) Microarrays for Epigenetic Analysis of Human Genomes. Methods in Molecular Biology, 2018, 1766, 137-156.	0.9	3
40	Memory B-cell like chronic lymphocytic leukaemia is associated with specific methylation profile of <i>WNT5A</i> promoter and undetectable expression of <i>WNT5A</i> gene. Epigenetics, 2022, 17, 1628-1635.	2.7	3
41	Analysis of Somatic DNA Methylation Alterations of Genes Encoding Cell Surface Metallopeptidases in Colorectal Cancer. Methods in Molecular Biology, 2018, 1731, 271-294.	0.9	2
42	Abstract 2792: Pericentromeric repeats undergo genome-wide DNA hypomethylation associated with older colorectal cancer patients without mutant p53 and are target for a reverse epigenetic switching. , 2011 , , .		0
43	Abstract 1381: Somatic DNA demethylation and epigenetic reprogramming of SST1 pericentromeric repeats associate with genomic damage in colorectal cancer. , 2014, , .		0
44	Improving metachronous cancer risk assessment using clinicopathological information in colorectal cancer patients Journal of Clinical Oncology, 2016, 34, e15092-e15092.	1.6	0