Anthony R Dallosso

List of Publications by Year
in descending order

19	737	687363 13	839539 18
papers	citations	h-index	g-index

20	20	20	1563
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Frequent Long-Range Epigenetic Silencing of Protocadherin Gene Clusters on Chromosome 5q31 in Wilms' Tumor. PLoS Genetics, 2009, 5, e1000745.	3.5	129
2	Multiple Rare Nonsynonymous Variants in the Adenomatous Polyposis Coli Gene Predispose to Colorectal Adenomas. Cancer Research, 2008, 68, 358-363.	0.9	77
3	Long-range epigenetic silencing of chromosome 5q31 protocadherins is involved in early and late stages of colorectal tumorigenesis through modulation of oncogenic pathways. Oncogene, 2012, 31, 4409-4419.	5.9	77
4	Alternately spliced WT1 antisense transcripts interact with WT1 sense RNA and show epigenetic and splicing defects in cancer. Rna, 2007, 13, 2287-2299.	3.5	71
5	Genomic imprinting at the WT1 gene involves a novel coding transcript (AWT1) that shows deregulation in Wilms' tumours. Human Molecular Genetics, 2003, 13, 405-415.	2.9	69
6	Inherited predisposition to colorectal adenomas caused by multiple rare alleles of MUTYH but not OGG1, NUDT1, NTH1 or NEIL 1, 2 or 3. Gut, 2008, 57, 1252-1255.	12.1	51
7	Hypomethylation and Aberrant Expression of the Glioma Pathogenesis-Related 1 Gene in Wilms Tumors. Neoplasia, 2007, 9, 970-978.	5.3	40
8	A CTCF-binding silencer regulates the imprinted genes AWT1 and WT1-AS and exhibits sequential epigenetic defects during Wilms' tumourigenesis. Human Molecular Genetics, 2007, 16, 343-354.	2.9	36
9	Control of epigenetic states by WT1 via regulation of de novo DNA methyltransferase 3A. Human Molecular Genetics, 2013, 22, 74-83.	2.9	36
10	Perilobar Nephrogenic Rests Are Nonobligate Molecular Genetic Precursor Lesions of Insulin-Like Growth Factor-II-Associated Wilms Tumors. Clinical Cancer Research, 2008, 14, 7635-7644.	7.0	30
11	The parathyroid hormone-responsive B1 gene is interrupted by a t(1;7)(q42;p15) breakpoint associated with Wilms' tumour. Oncogene, 2003, 22, 1371-1380.	5.9	28
12	Genomeâ€wide DNA methylation analysis identifies MEGF10 as a novel epigenetically repressed candidate tumor suppressor gene in neuroblastoma. Molecular Carcinogenesis, 2017, 56, 1290-1301.	2.7	23
13	Epigenetic deregulation of GATA3 in neuroblastoma is associated with increased GATA3 protein expression and with poor outcomes. Scientific Reports, 2019, 9, 18934.	3.3	17
14	Characterization of 17.94, a novel anaplastic Wilms' tumor cell line. Cancer Genetics, 2012, 205, 319-326.	0.4	16
15	Increased Efficacy of Histone Methyltransferase G9a Inhibitors Against MYCN-Amplified Neuroblastoma. Frontiers in Oncology, 2020, 10, 818.	2.8	14
16	A human importin-Î²-related disorder: Syndromic thoracic aortic aneurysm caused by bi-allelic loss-of-function variants in IPO8. American Journal of Human Genetics, 2021, 108, 1115-1125.	6.2	10
17	TheAPCVariant p.Glu1317Gln predisposes to colorectal adenomas by a novel mechanism of relaxing the target for tumorigenic somaticAPCmutations. Human Mutation, 2009, 30, 1412-1418.	2.5	8
18	Rapid recognition of aberrant dHPLC elution profiles using the Transgenomic NavigatorTM software. Human Mutation, 2005, 26, 165-165.	2.5	5

#	Article	IF	Citations
19	Severe Congenital Myopathy and Neuropathy with Congenital Cataracts due to GFER Variant: A Neuropathological Study. Journal of Pediatric Neurology, 0, , .	0.2	0