Keith W Brown

List of Publications by Year in descending order

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| # | Article | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | The epithelial splicing regulator <i>ESRP2</i> is epigenetically repressed by DNA hypermethylation in Wilms tumour and acts as a tumour suppressor. Molecular Oncology, 2022, 16, 630-647. | 4.6 | 3 |
| 2 | 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 |
| 3 | Genomeâ€wide DNA methylation analysis identifies <i>MEGF10</i> as a novel epigenetically repressed candidate tumor suppressor gene in neuroblastoma. Molecular Carcinogenesis, 2017, 56, 1290-1301. | 2.7 | 23 |
| 4 | Protein arginine methyltransferase 5 is a key regulator of the MYCN oncoprotein in neuroblastoma cells. Molecular Oncology, 2015, 9, 617-627. | 4.6 | 49 |
| 5 | MYCN is recruited to the <i>RASSF1A</i> promoter but is not critical for DNA hypermethylation in neuroblastoma. Molecular Carcinogenesis, 2014, 53, 413-420. | 2.7 | 6 |
| 6 | Insulin-like growth factor binding protein-3 (IGFBP-3) plays a role in the anti-tumorigenic effects of 5-Aza-2′-deoxycytidine (AZA) in breast cancer cells. Experimental Cell Research, 2013, 319, 2282-2295. | 2.6 | 16 |
| 7 | Control of epigenetic states by WT1 via regulation of de novo DNA methyltransferase 3A. Human Molecular Genetics, 2013, 22, 74-83. | 2.9 | 36 |
| 8 | Characterization of 17.94, a novel anaplastic Wilms' tumor cell line. Cancer Genetics, 2012, 205, 319-326. | 0.4 | 16 |
| 9 | DNA demethylation increases sensitivity of neuroblastoma cells to chemotherapeutic drugs. Biochemical Pharmacology, 2012, 83, 858-865. | 4.4 | 49 |
| 10 | Frequent Long-Range Epigenetic Silencing of Protocadherin Gene Clusters on Chromosome 5q31 in Wilms' Tumor. PLoS Genetics, 2009, 5, e1000745. | 3.5 | 129 |
| 11 | 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 |
| 12 | Frequency and Timing of Loss of Imprinting at 11p13 and 11p15 in Wilms' Tumor Development. Molecular Cancer Research, 2008, 6, 1114-1123. | 3.4 | 20 |
| 13 | 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 |
| 14 | Alternately spliced <i>WT1</i> antisense transcripts interact with <i>WT1</i> sense RNA and show epigenetic and splicing defects in cancer. Rna, 2007, 13, 2287-2299. | 3.5 | 71 |
| 15 | Hypomethylation and Aberrant Expression of the Glioma Pathogenesis-Related 1 Gene in Wilms Tumors. Neoplasia, 2007, 9, 970-978. | 5.3 | 40 |
| 16 | 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 |
| 17 | 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 |
| 18 | Low frequency of genetic lesions in Wilms tumors by representational difference analysis. Cancer Genetics and Cytogenetics, 2001, 127, 155-160. | 1.0 | 3 |

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|----|---|------|-----------|
| 19 | The molecular biology of Wilms' tumour. Expert Reviews in Molecular Medicine, 2001, 3, 1-16. | 3.9 | 24 |
| 20 | Transactivation of the WT1 antisense promoter is unique to the WT1[+/â^'] isoform. FEBS Letters, 1999, 456, 131-136. | 2.8 | 8 |
| 21 | Loss of WT1 function leads to ectopic myogenesis in Wilms' tumour. Nature Genetics, 1998, 18, 15-17. | 21.4 | 69 |
| 22 | Antisense WT1 transcription parallels sense mRNA and protein expression in fetal kidney and can elevate protein levelsin vitro. , 1998, 185, 352-359. | | 45 |
| 23 | Microdissecting the Genetic Events in Nephrogenic Rests and Wilms' Tumor Development. American Journal of Pathology, 1998, 153, 991-1000. | 3.8 | 78 |
| 24 | Localization of a novel t(1;7) translocation associated with Wilms' tumor predisposition and skeletal abnormalities. , 1996, 17, 151-155. | | 19 |
| 25 | Imprinting mutations in the Beckwith—Wiedemann syndrome suggested by an altered imprinting pattern in the IGF2–H19 domain. Human Molecular Genetics, 1995, 4, 2379-2385. | 2.9 | 235 |
| 26 | SSCP and incorporation of 7-deaza-2′ddGTP into PCR products. Trends in Genetics, 1994, 10, 225. | 6.7 | 0 |
| 27 | Germline and somatic abnormalities of chromosome 7 in Wilms' tumor. Cancer Genetics and Cytogenetics, 1994, 77, 93-98. | 1.0 | 54 |
| 28 | Autoregulation of the human WT1 gene promoter. FEBS Letters, 1994, 349, 75-78. | 2.8 | 38 |
| 29 | Somatic Allelic Loss at the DCC, APC, nm23-H1 and P53 Tumor Suppressor Gene Loci in Human Prostatic Carcinoma. Journal of Urology, 1994, 151, 1073-1077. | 0.4 | 91 |
| 30 | Low frequency of mutations in theWT1 coding region in Wilms' tumor. Genes Chromosomes and Cancer, 1993, 8, 74-79. | 2.8 | 42 |
| 31 | Alteration of the extracellular matrix of cultured human keratinocytes by transformation and during differentiation. International Journal of Cancer, 1985, 35, 799-807. | 5.1 | 32 |
| 32 | Extracellular matrix components produced by SV40-transformed human epidermal keratinocytes. International Journal of Cancer, 1984, 33, 257-263. | 5.1 | 29 |
| 33 | Radioimmunoassay of brain-type creatine kinase-BB isoenzyme in human tissues and in serum of patients with neurological disorders. Journal of the Neurological Sciences, 1980, 47, 241-254. | 0.6 | 70 |

Purification, radioimmuno assay, and distribution of human brain 14-3-2 protein (nervous-system) Tj ETQq0 0 0 rg $B_{2.4}^{T}$ (Overlock 10 Tf 50