Emmanuelle Viré

List of Publications by Year in descending order

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EMMANHELLE VIDÃO

#	Article	IF	CITATIONS
1	The Polycomb group protein EZH2 directly controls DNA methylation. Nature, 2006, 439, 871-874.	27.8	1,964
2	Myc represses transcription through recruitment of DNA methyltransferase corepressor. EMBO Journal, 2005, 24, 336-346.	7.8	375
3	Role of the Polycomb Repressive Complex 2 in Acute Promyelocytic Leukemia. Cancer Cell, 2007, 11, 513-525.	16.8	228
4	Epigenetic Regulation of Tumor Endothelial Cell Anergy: Silencing of Intercellular Adhesion Molecule-1 by Histone Modifications. Cancer Research, 2006, 66, 10770-10777.	0.9	139
5	Identification of Epigenetically Silenced Genes in Tumor Endothelial Cells. Cancer Research, 2007, 67, 4138-4148.	0.9	126
6	Angiostatic activity of DNA methyltransferase inhibitors. Molecular Cancer Therapeutics, 2006, 5, 467-475.	4.1	93
7	Inhibition of the acetyltransferase NAT10 normalizes progeric and aging cells by rebalancing the Transportin-1 nuclear import pathway. Science Signaling, 2018, 11, .	3.6	57
8	The Breast Cancer Oncogene EMSY Represses Transcription of Antimetastatic microRNA miR-31. Molecular Cell, 2014, 53, 806-818.	9.7	55
9	White matter DNA methylation profiling reveals deregulation of HIP1, LMAN2, MOBP, and other loci in multiple system atrophy. Acta Neuropathologica, 2020, 139, 135-156.	7.7	42
10	Identification of novel risk loci and causal insights for sporadic Creutzfeldt-Jakob disease: a genome-wide association study. Lancet Neurology, The, 2020, 19, 840-848.	10.2	42
11	Synthetic viability genomic screening defines Sae2 function in DNA repair. EMBO Journal, 2015, 34, 1509-1522.	7.8	37
12	Altered DNA methylation profiles in blood from patients with sporadic Creutzfeldt–Jakob disease. Acta Neuropathologica, 2020, 140, 863-879.	7.7	18
13	MOBP and HIP1 in multiple system atrophy: New αâ€synuclein partners in glial cytoplasmic inclusions implicated in the disease pathogenesis. Neuropathology and Applied Neurobiology, 2021, 47, 640-652.	3.2	11
14	Epigenomics and transcriptomics analyses of multiple system atrophy brain tissue supports a role for inflammatory processes in disease pathogenesis. Acta Neuropathologica Communications, 2020, 8, 71.	5.2	5
15	Prion protein gene mutation detection using long-read Nanopore sequencing. Scientific Reports, 2022, 12, 8284.	3.3	4