Sheryl M Gough

List of Publications by Year
in descending order

12 papers	431 citations	1478505 6 h-index	1474206 9 g-index
12 all docs	12 docs citations	12 times ranked	925 citing authors

#	Article	IF	Citations
1	Mechanistic insights into chromatin targeting by leukemic NUP98-PHF23 fusion. Nature Communications, 2020, 11, 3339.	12.8	15
2	Somatic mutations in murine models of leukemia and lymphoma: Disease specificity and clinical relevance. Genes Chromosomes and Cancer, 2017, 56, 472-483.	2.8	9
3	Progenitor B-1 B-cell acute lymphoblastic leukemia is associated with collaborative mutations in 3 critical pathways. Blood Advances, 2017, 1, 1749-1759.	5.2	19
4	Oxidative stress leads to increased mutation frequency in a murine model of myelodysplastic syndrome. Leukemia Research, 2014, 38, 95-102.	0.8	44
5	NUP98â€"PHF23 Is a Chromatin-Modifying Oncoprotein That Causes a Wide Array of Leukemias Sensitive to Inhibition of PHD Histone Reader Function. Cancer Discovery, 2014, 4, 564-577.	9.4	66
6	Depletion of Cytotoxic T-Cells Does Not Protect NUP98-HOXD13 Mice from Myelodysplastic Syndrome but Reveals a Modest Tumor Immunosurveillance Effect. PLoS ONE, 2012, 7, e36876.	2.5	6
7	Illegitimate V(D)J Recombination Involving Notch1 and Bcl11b in Precursor T-Cell Lymphoblastic Leukemia/Lymphoma. Blood, 2012, 120, 1323-1323.	1.4	0
8	NUP98 gene fusions and hematopoietic malignancies: common themes and new biologic insights. Blood, 2011, 118, 6247-6257.	1.4	263
9	Increased Mutation Frequency Induced by Oxidative Stress in the NUP98-HOXD13 MDS Mouse Model. Blood, 2011, 118, 1705-1705.	1.4	0
10	Rag1 Deficiency Does Not Affect Myelodysplasia but Leads to More Rapid Leukemic Transformation in a Mouse Model of MDS.. Blood, 2009, 114, 2784-2784.	1.4	0
11	Translocation (5;10)(q22;q24) in a case of acute lymphoblastic leukemia. Cancer Genetics and Cytogenetics, 2006, 165, 36-40.	1.0	3
12	Refined physical map of the human PAX2/HOX11/NFKB2 cancer gene region at 10q24 and relocalization of the HPV6AI1 viral integration site to 14q13.3-q21.1. BMC Genomics, 2003, 4, 9.	2.8	6