

Jane S Green

List of Publications by Year in descending order

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Version: 2024-02-01

25
papers

3,241
citations

758635

12
h-index

676716

22
g-index

25
all docs

25
docs citations

25
times ranked

2319
citing authors

#	ARTICLE	IF	CITATIONS
1	A dominant <i>RAD51C</i> pathogenic splicing variant predisposes to breast and ovarian cancer in the Newfoundland population due to founder effect. <i>Molecular Genetics & Genomic Medicine</i> , 2020, 8, e1070.	0.6	6
2	The genetic architecture of Stargardt macular dystrophy (STGD1): a longitudinal 40-year study in a genetic isolate. <i>European Journal of Human Genetics</i> , 2020, 28, 925-937.	1.4	10
3	A novel pathogenic missense ADAMTS17 variant that impairs secretion causes Weill-Marchesani Syndrome with variably dysmorphic hand features. <i>Scientific Reports</i> , 2020, 10, 10827.	1.6	13
4	The long-term survival characteristics of a cohort of colorectal cancer patients and baseline variables associated with survival outcomes with or without time-varying effects. <i>BMC Medicine</i> , 2019, 17, 150.	2.3	32
5	Novel Usher syndrome pathogenic variants identified in cases with hearing and vision loss. <i>BMC Medical Genetics</i> , 2019, 20, 68.	2.1	10
6	Screening of BMPR1a for pathogenic mutations in familial colorectal cancer type X families from Newfoundland. <i>Familial Cancer</i> , 2018, 17, 205-208.	0.9	2
7	Impact of colonoscopic screening in Familial Colorectal Cancer Type X. <i>Molecular Genetics & Genomic Medicine</i> , 2018, 6, 1021-1030.	0.6	10
8	Evidence for <i>GALNT12</i> as a moderate penetrance gene for colorectal cancer. <i>Human Mutation</i> , 2018, 39, 1092-1101.	1.1	20
9	Germline <i>INDEL</i> s and <i>CNV</i> s in a cohort of colorectal cancer patients: their characteristics, associations with relapse-free survival time, and potential time-varying effects on the risk of relapse. <i>Cancer Medicine</i> , 2017, 6, 1220-1232.	1.3	14
10	Novel 25 kb Deletion of MERTK Causes Retinitis Pigmentosa With Severe Progression. , 2017, 58, 1736.		17
11	Structured assessment and followup for patients with hereditary kidney tumour syndromes. <i>Canadian Urological Association Journal</i> , 2016, 10, 214.	0.3	12
12	No associations of a set of SNPs in the Vascular Endothelial Growth Factor (VEGF) and Matrix Metalloproteinase (MMP) genes with survival of colorectal cancer patients. <i>Cancer Medicine</i> , 2016, 5, 2221-2231.	1.3	6
13	Genetic structure of the Newfoundland and Labrador population: founder effects modulate variability. <i>European Journal of Human Genetics</i> , 2016, 24, 1063-1070.	1.4	22
14	MG-100...Review of a large family with schwannomatosis identifies an expanded phenotype. <i>Journal of Medical Genetics</i> , 2015, 52, A1.1-A1.	1.5	0
15	A Survival Association Study of 102 Polymorphisms Previously Associated with Survival Outcomes in Colorectal Cancer. <i>BioMed Research International</i> , 2015, 2015, 1-9.	0.9	5
16	Community engagement with genetics: public perceptions and expectations about genetics research. <i>Health Expectations</i> , 2015, 18, 1413-1425.	1.1	30
17	A genome wide association study on Newfoundland colorectal cancer patients's™ survival outcomes. <i>Biomarker Research</i> , 2015, 3, 6.	2.8	17
18	Public Attitudes About Genetic Testing in the Newborn Period. <i>JOGNN - Journal of Obstetric, Gynecologic, and Neonatal Nursing</i> , 2012, 41, 191-200.	0.2	28

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19	Impact of Gender and Parent of Origin on the Phenotypic Expression of Hereditary Nonpolyposis Colorectal Cancer in a Large Newfoundland Kindred With a Common MSH2 Mutation. <i>Diseases of the Colon and Rectum</i> , 2002, 45, 1223-1232.	0.7	48
20	Mutations in MKKS cause Bardet-Biedl syndrome. <i>Nature Genetics</i> , 2000, 26, 15-16.	9.4	256
21	Mutations in MKKS cause obesity, retinal dystrophy and renal malformations associated with Bardet-Biedl syndrome. <i>Nature Genetics</i> , 2000, 26, 67-70.	9.4	311
22	Canadian Bardet-Biedl syndrome family reduces the critical region of BBS3 (3p) and presents with a variable phenotype. , 1998, 78, 461-467.		49
23	Common ancestral mutations in the MEN1 gene is likely responsible for the prolactinoma variant of MEN1 (MEN1Burin) in four kindreds from Newfoundland. <i>Human Mutation</i> , 1998, 11, 264-269.	1.1	120
24	Common ancestral mutations in the MEN1 gene is likely responsible for the prolactinoma variant of MEN1 (MEN1Burin) in four kindreds from Newfoundland. <i>Human Mutation</i> , 1998, 11, 264-269.	1.1	8
25	Mutations of a mutS homolog in hereditary nonpolyposis colorectal cancer. <i>Cell</i> , 1993, 75, 1215-1225.	13.5	2,195