## Sylviane Olschwang

List of Publications by Year in descending order

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118 papers 14,426 citations

50276 46 h-index 118 g-index

134 all docs

134 docs citations

times ranked

134

15985 citing authors

#	Article	IF	CITATIONS
1	Variation in the risk of colorectal cancer in families with Lynch syndrome: a retrospective cohort study. Lancet Oncology, The, 2021, 22, 1014-1022.	10.7	58
2	Characterisation of heterozygous <i>PMS2</i> variants in French patients with Lynch syndrome. Journal of Medical Genetics, 2020, 57, 487-499.	3.2	10
3	The Balance Between Cytotoxic T-cell Lymphocytes and Immune Checkpoint Expression in the Prognosis of Colon Tumors. Journal of the National Cancer Institute, 2018, 110, 68-77.	6.3	89
4	Improving Mutation Screening in Patients with Colorectal Cancer Predisposition Using Next-Generation Sequencing. Journal of Molecular Diagnostics, 2017, 19, 589-601.	2.8	11
5	Molecular combing reveals complex 4q35 rearrangements in Facioscapulohumeral dystrophy. Human Mutation, 2017, 38, 1432-1441.	2.5	39
6	Point Mutations in Exon 1B of APC Reveal Gastric Adenocarcinoma and Proximal Polyposis of the Stomach as a Familial Adenomatous Polyposis Variant. American Journal of Human Genetics, 2016, 98, 830-842.	6.2	201
7	Actionable Genes, Core Databases, and Locus-Specific Databases. Human Mutation, 2016, 37, 1299-1307.	2.5	6
8	PIK3CA mutations predict recurrence in localized microsatellite stable colon cancer. Cancer Medicine, 2015, 4, 371-382.	2.8	25
9	Vascular Endothelial Growth Factor A c.*237C>T polymorphism is associated with bevacizumab efficacy and related hypertension in metastatic colorectal cancer. Digestive and Liver Disease, 2015, 47, 331-337.	0.9	23
10	Frequent intragenic rearrangements of DPYD in colorectal tumours. Pharmacogenomics Journal, 2015, 15, 211-218.	2.0	3
11	Obesity, Aspirin, and Risk of Colorectal Cancer in Carriers of Hereditary Colorectal Cancer: A Prospective Investigation in the CAPP2 Study. Journal of Clinical Oncology, 2015, 33, 3591-3597.	1.6	91
12	Genomic variations integrated database for <i>MUTYH</i> -associated adenomatous polyposis. Journal of Medical Genetics, 2015, 52, 25-27.	3.2	2
13	Clinical Utility Gene Card for: Familial adenomatous polyposis (FAP) and attenuated FAP (AFAP) - update 2014. European Journal of Human Genetics, 2015, 23, 889-889.	2.8	9
14	Gastric Adenocarcinoma in Familial Adenomatous Polyposis Can Occur Without Previous Lesions. Journal of Gastrointestinal Cancer, 2014, 45, 377-379.	1.3	4
15	The UMD-APC Database, a Model of Nation-Wide Knowledge Base: Update with Data from 3,581 Variations. Human Mutation, 2014, 35, 532-536.	2.5	15
16	Molecular patterns in deficient mismatch repair colorectal tumours: results from a French prospective multicentric biological and genetic study. British Journal of Cancer, 2014, 110, 2728-2737.	6.4	20
17	High-resolution analysis of DNA copy number alterations in rectal cancer. Strahlentherapie Und Onkologie, 2014, 190, 1028-1036.	2.0	4
18	A Multicenter Blinded Study Evaluating EGFR and KRAS Mutation Testing Methods in the Clinical Non–Small Cell Lung Cancer Setting—IFCT/ERMETIC2 Project Part 1. Journal of Molecular Diagnostics, 2014, 16, 45-55.	2.8	31

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19	Abstract 5555: Pharmacogenetics-pharmacokinetics study of bevacizumab in mCRC patients treated with Avastin-Folfiri regimen: Search for predictive markers of response and study of the pharmacokinetics variability. , 2014, , .		1
20	Semiparametric inference on the penetrances of rare genetic mutations based on a case-family design. Journal of Statistical Planning and Inference, 2013, 143, 368-377.	0.6	0
21	Clinical utility gene card for: Lynch syndrome (MLH1, MSH2, MSH6, PMS2, EPCAM) - update 2012. European Journal of Human Genetics, 2013, 21, 118-118.	2.8	16
22	Gene Expression Classification of Colon Cancer into Molecular Subtypes: Characterization, Validation, and Prognostic Value. PLoS Medicine, 2013, 10, e1001453.	8.4	1,064
23	UMD-MLH1/MSH2/MSH6 databases: description and analysis of genetic variations in French Lynch syndrome families. Database: the Journal of Biological Databases and Curation, 2013, 2013, bat036-bat036.	3.0	15
24	Design of a Core Classification Process for DNA Mismatch Repair Variations of A Priori Unknown Functional Significance. Human Mutation, 2013, 34, 920-922.	2.5	3
25	A Seven-Gene Signature Aggregates a Subgroup of Stage II Colon Cancers with Stage III. OMICS A Journal of Integrative Biology, 2012, 16, 560-565.	2.0	69
26	KRAS Mutation Spectrum Notably Diverges between Non-small Cell Lung and Colorectal Carcinomas. Journal of Thoracic Oncology, 2012, 7, 773-774.	1.1	4
27	Long-term effect of resistant starch on cancer risk in carriers of hereditary colorectal cancer: an analysis from the CAPP2 randomised controlled trial. Lancet Oncology, The, 2012, 13, 1242-1249.	10.7	95
28	Expression Profiles in Stage II Colon Cancer According to APC Gene Status. Translational Oncology, 2012, 5, 72-76.	3.7	18
29	Is the controversy on breast cancer as part of the Lynch-related tumor spectrum still open?. Familial Cancer, 2012, 11, 681-683.	1.9	11
30	8q24 Cancer Risk Allele Associated with Major Metastatic Risk in Inflammatory Breast Cancer. PLoS ONE, 2012, 7, e37943.	2.5	34
31	Cancer Risks Associated With Germline Mutations in <emph type="ital">MLH1</emph> , <emph type="ital">MSH2</emph> , and <emph type="ital">MSH6</emph> Genes in Lynch Syndrome. JAMA - Journal of the American Medical Association, 2011, 305, 2304.	7.4	878
32	Long-term effect of aspirin on cancer risk in carriers of hereditary colorectal cancer: an analysis from the CAPP2 randomised controlled trial. Lancet, The, 2011, 378, 2081-2087.	13.7	849
33	Cross-Validation Study for Epidermal Growth Factor Receptor and RRAS Mutation Detection in 74 Blinded Non-small Cell Lung Carcinoma Samples: A Total of 5550 Exons Sequenced by 15 Molecular French Laboratories (Evaluation of the EGFR Mutation Status for the Administration of EGFR-TKIs in) Tj ETQq1 1	0.7 <b>81</b> 314	rg <b>B∑</b>  Overlo
34	1006-1015. Clinical utility gene card for: Familial adenomatous polyposis (FAP) and attenuated FAP (AFAP). European Journal of Human Genetics, 2011, 19, 832-832.	2.8	10
35	The LKB1 complex-AMPK pathway: the tree that hides the forest. Familial Cancer, 2011, 10, 415-424.	1.9	27
36	SMAD4 protein expression and cell proliferation in colorectal adenocarcinomas. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2011, 459, 511-519.	2.8	7

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37	SMARCB1/INI1 germline mutations contribute to 10% of sporadic schwannomatosis. BMC Neurology, 2011, 11, 9.	1.8	76
38	The educational role of external quality assessment in genetic testing: a 7-year experience of the European Molecular Genetics Quality Network (EMQN) in Lynch syndrome. Human Mutation, 2011, 32, 696-697.	2.5	6
39	Contribution of Ultrasonography to Endometrial Cancer Screening in Patients With Hereditary Nonpolyposis Colorectal Cancer/Lynch Syndrome. International Journal of Gynecological Cancer, 2010, 20, 583-587.	2.5	17
40	Combined mutations of ASXL1, CBL, FLT3, IDH1, IDH2, JAK2, KRAS, NPM1, NRAS, RUNX1, TET2 and WT1 genes in myelodysplastic syndromes and acute myeloid leukemias. BMC Cancer, 2010, 10, 401.	2.6	140
41	Leiden open variation database of the MUTYH gene. Human Mutation, 2010, 31, 1205-1215.	2.5	72
42	Cancer occurrence during follow-up of the CAPP2 study -aspirin use for up to four years significantly reduces Lynch syndrome cancers for up to several years after completion of therapy. Hereditary Cancer in Clinical Practice, 2010, 8, O5.	1.5	2
43	Clinical utility gene card for: Lynch syndrome (MLH1, MSH2, MSH6, PMS2). European Journal of Human Genetics, 2010, 18, 1071-1071.	2.8	10
44	Statistical inference on the penetrances of rare genetic mutations based on a case-family design. Biostatistics, 2010, 11, 519-532.	1.5	5
45	Germline APC mutation spectrum derived from 863 genomic variations identified through a 15-year medical genetics service to French patients with FAP. Journal of Medical Genetics, 2010, 47, 721-722.	3.2	32
46	Chronic Ironâ€deficiency Anemia Caused by a Jejunojejunal Intussusception on a Solitary Hamartomatous Polyp. Journal of Pediatric Gastroenterology and Nutrition, 2010, 50, 450-452.	1.8	1
47	iCOMET, projet national de recherche de gÃ"nes de prédisposition au développement des métastases chez les patients traités pour un cancer colique sporadique de stade II. Cancéro Digest, 2009, , 7.	0.0	O
48	Analysis of Candidate Genes in Occurrence and Growth of Colorectal Adenomas. Journal of Oncology, 2009, 2009, 1-9.	1.3	2
49	Age-Dependent Cancer Risk Is Not Different in between <i>MSH2</i> and <i>MLH1</i> Mutation Carriers. Journal of Cancer Epidemiology, 2009, 2009, 1-6.	1.1	4
50	Quantitative PCR high-resolution melting (qPCR-HRM) curve analysis, a new approach to simultaneously screen point mutations and large rearrangements: application to <i>MLH1</i> germline mutations in Lynch syndrome. Human Mutation, 2009, 30, 867-875.	2.5	58
51	Mutations of polycombâ€associated gene <i>ASXL1</i> in myelodysplastic syndromes and chronic myelomonocytic leukaemia. British Journal of Haematology, 2009, 145, 788-800.	2.5	537
52	A large fraction of unclassified variants of the mismatch repair genes <i>MLH1</i> and <i>MSH2</i> is associated with splicing defects. Human Mutation, 2008, 29, 1412-1424.	2.5	150
53	Mitochondrial D310 mutations in colorectal adenomas: An early but not causative genetic event during colorectal carcinogenesis. International Journal of Cancer, 2008, 122, 2242-2248.	5.1	19
54	Y253H mutation appearing in a $\hat{1}\frac{1}{4}$ -BCR-ABL (e19a2) CML. Leukemia Research, 2008, 32, 361-362.	0.8	13

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55	A polymorphism of EGFR extracellular domain is associated with progression free-survival in metastatic colorectal cancer patients receiving cetuximab-based treatment. BMC Cancer, 2008, 8, 169.	2.6	65
56	Genome profiling of chronic myelomonocytic leukemia: frequent alterations of RAS and RUNX1genes. BMC Cancer, 2008, 8, 299.	2.6	109
57	Identification in Daily Practice of Patients With Lynch Syndrome (Hereditary Nonpolyposis Colorectal) Tj ETQq1 I Journal of Gastroenterology, 2008, 103, 2825-2835.	0.78431 0.4	4 rgBT /Over 118
58	Usefulness of Prophylactic Gastrectomy in a Novel Large Hereditary Diffuse Gastric Cancer (HDGC) Family. American Journal of Gastroenterology, 2008, 103, 2160-2161.	0.4	28
59	Effect of Aspirin or Resistant Starch on Colorectal Neoplasia in the Lynch Syndrome. New England Journal of Medicine, 2008, 359, 2567-2578.	27.0	273
60	Multicenter study of ZAP-70 expression in patients with B-cell chronic lymphocytic leukemia using an optimized flow cytometry method. Haematologica, 2008, 93, 215-223.	3.5	32
61	High-resolution Genotyping of Chromosome 8 in Colon Adenocarcinomas Reveals Recurrent Break Point But No Gene Mutation in the 8p21 Region. Diagnostic Molecular Pathology, 2008, 17, 90-93.	2.1	9
62	Duodenal adenocarcinoma and Mut Y human homologue-associated polyposis. European Journal of Gastroenterology and Hepatology, 2008, 20, 1024-1027.	1.6	9
63	The Thorough Screening of the MUTYH Gene in a Large French Cohort of Sporadic Colorectal Cancers. Genetic Testing and Molecular Biomarkers, 2007, 11, 373-380.	1.7	15
64	Founder and Recurrent CDH1 Mutations in Families With Hereditary Diffuse Gastric Cancer. JAMA - Journal of the American Medical Association, 2007, 297, 2360.	7.4	394
65	New types of MYST3-CBP and CBP-MYST3 fusion transcripts in t(8;16)(p11;p13) acute myeloid leukemias. Haematologica, 2007, 92, 262-263.	3.5	12
66	Similar Colorectal Cancer Risk in Patients with Monoallelic and Biallelic Mutations in the MYH Gene Identified in a Population with Adenomatous Polyposis. Genetic Testing and Molecular Biomarkers, 2007, 11, 315-320.	1.7	37
67	Colonoscopic Screening of First-Degree Relatives of Patients With Large Adenomas: Increased Risk of Colorectal Tumors. Gastroenterology, 2007, 133, 1086-1092.	1.3	87
68	Genome-wide association scan identifies a colorectal cancer susceptibility locus on chromosome 8q24. Nature Genetics, 2007, 39, 989-994.	21.4	676
69	Partial duplications of the MSH2 and MLH1 genes in hereditary nonpolyposis colorectal cancer. European Journal of Human Genetics, 2007, 15, 383-386.	2.8	14
70	Rearrangements involving 12q in myeloproliferative disorders: possible role of HMGA2 and SOCS2 genes. Cancer Genetics and Cytogenetics, 2007, 176, 80-88.	1.0	26
71	Hysteroscopic findings in women at risk of HNPCC. Results of a prospective observational study. Familial Cancer, 2007, 6, 295-299.	1.9	24
72	High Frequency of Chromosome 14 Deletion in Early-Onset Colon Cancer. Diseases of the Colon and Rectum, 2007, 50, 1881-1886.	1.3	22

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73	Genetic polymorphisms of MMP1, MMP3 and MMP7gene promoter and risk of colorectal adenoma. BMC Cancer, 2006, 6, 270.	2.6	56
74	Retention of chromosome arm 5q in stage II colon cancers identifies 83% of liver metastasis occurrences. Genes Chromosomes and Cancer, 2006, 45, 94-102.	2.8	8
75	Frequency and Spectrum of Cancers in the Peutz-Jeghers Syndrome. Clinical Cancer Research, 2006, 12, 3209-3215.	7.0	746
76	SMAD4 Germinal Mosaicism in a Family with Juvenile Polyposis and Hypertrophic Osteoarthropathy. Journal of Pediatric Gastroenterology and Nutrition, 2005, 41, 117-120.	1.8	6
77	Value of microsatellite instability typing in detecting hereditary non-polyposis colorectal cancer. Gastroenterologie Clinique Et Biologique, 2005, 29, 667-675.	0.9	11
78	The $5\hat{a} \in \mathbb{R}^2$ region of the MSH2 gene involved in hereditary non-polyposis colorectal cancer contains a high density of recombinogenic sequences. Human Mutation, 2005, 26, 255-261.	2.5	31
79	Vessels' morphology inSMAD4 andBMPR1A-related juvenile polyposis. American Journal of Medical Genetics, Part A, 2005, 138A, 113-117.	1.2	45
80	Functional analysis of Peutz–Jeghers mutations reveals that the LKB1 C-terminal region exerts a crucial role in regulating both the AMPK pathway and the cell polarity. Human Molecular Genetics, 2005, 14, 1283-1292.	2.9	131
81	Analysis of the allele-specific expression of the mismatch repair geneMLH1using a simple DHPLC-Based Method. Human Mutation, 2004, 23, 379-384.	2.5	24
82	Genotype-phenotype correlation in von Hippel-Lindau families with renal lesions. Human Mutation, 2004, 24, 215-224.	2.5	81
83	Relative frequency and morphology of cancers in STK11 mutation carriers1 â~†. Gastroenterology, 2004, 126, 1788-1794.	1.3	228
84	Mucinous colon carcinomas with microsatellite instability have a lower microvessel density and lower vascular endothelial growth factor expression. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2003, 442, 111-117.	2.8	22
85	Tissue microarray technology: validation in colorectal carcinoma and analysis of p53, hMLH1, and hMSH2 immunohistochemical expression. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2003, 443, 115-121.	2.8	109
86	Association between mutations in the CARD15 (NOD2) gene and Crohn's disease in Israeli Jewish patients. American Journal of Medical Genetics Part A, 2003, 121A, 240-244.	2.4	29
87	Microsatellite Instability in Colorectal Carcinoma. Archives of Pathology and Laboratory Medicine, 2003, 127, 694-700.	2.5	81
88	MSH2 in contrast to MLH1 and MSH6 is frequently inactivated by exonic and promoter rearrangements in hereditary nonpolyposis colorectal cancer. Cancer Research, 2002, 62, 848-53.	0.9	86
89	Kirsten ras mutations in patients with colorectal cancer: the â€~RASCAL II' study. British Journal of Cancer, 2001, 85, 692-696.	6.4	790
90	Familial Adenomatous Polyposis: Prevalence of Adenomas in the Ileal Pouch After Restorative Proctocolectomy. Annals of Surgery, 2001, 233, 360-364.	4.2	139

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91	Peutz-Jeghers families unlinked toSTK11/LKB1 gene mutations are highly predisposed to primitive biliary adenocarcinoma. Journal of Medical Genetics, 2001, 38, 356-360.	3.2	81
92	Germ-line mutations of the APC gene in patients with FAP associated hepatoblastoma are located out of the mutation cluster region (codons 1286–1513). Gastroenterology, 2000, 118, A45.	1.3	2
93	Deletion mapping of the tumor suppressor locus involved in colorectal cancer on chromosome band 8p21., 1999, 25, 147-153.		26
94	Fine deletion mapping of chromosome 8p in non-small-cell lung carcinoma. , 1999, 81, 854-858.		40
95	A serine/threonine kinase gene defective in Peutz–Jeghers syndrome. Nature, 1998, 391, 184-187.	27.8	1,451
96	PTEN germ-line mutations in juvenile polyposis coli. Nature Genetics, 1998, 18, 12-14.	21.4	153
97	Somatically acquired genetic alterations in flat colorectal neoplasias. , 1998, 77, 366-369.		34
98	Germline mutation profile of the VHL gene in von Hippel-Lindau disease and in sporadic hemangioblastoma. Human Mutation, 1998, 12, 424-430.	2.5	89
99	Genetic Alterations in Thyroid Carcinoma Associated with Familial Adenomatous Polyposis: Clinical Implications and Suggestions for Early Detection. World Journal of Surgery, 1998, 22, 1231-1236.	1.6	75
100	Alternative genetic pathways in colorectal carcinogenesis. Proceedings of the National Academy of Sciences of the United States of America, 1997, 94, 12122-12127.	7.1	209
101	Germline mutations in the Von Hippel-Lindau disease (VHL) gene in families from North America, Europe, and Japan. Human Mutation, 1996, 8, 348-357.	2.5	436
102	Germline mutations in the Von Hippelâ€Lindau disease (VHL) gene in families from North America, Europe, and Japan. Human Mutation, 1996, 8, 348-357.	2.5	33
103	High resolution genetic map of the adenomatous polyposis coli gene (APC) region. American Journal of Medical Genetics Part A, 1995, 56, 413-419.	2.4	15
104	Predominance of normal karyotype in colorectal tumors from hereditary non-polyposis colorectal cancer patients. Genes Chromosomes and Cancer, 1995, 14, 223-226.	2.8	24
105	Association of p53 mutations with short survival in colorectal cancer. Gastroenterology, 1994, 106, 42-48.	1.3	348
106	Expression phénotypique de la polypose adénomateuse familiale. Medecine/Sciences, 1994, 10, 454.	0.2	0
107	Alleles of the APC gene: An attenuated form of familial polyposis. Cell, 1993, 75, 951-957.	28.9	611
108	Restriction of ocular fundus lesions to a specific subgroup of APC mutations in adenomatous polyposis coli patients. Cell, 1993, 75, 959-968.	28.9	307

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109	Is the multiple endocrine neoplasia type 1 gene a suppressor for fundic argyrophil tumors in the Zollinger-Ellison syndrome?. Gastroenterology, 1993, 105, 579-582.	1.3	79
110	Survival and acquired genetic alterations in colorectal cancer. Gastroenterology, 1992, 102, 1136-1141.	1.3	182
111	Detection by DGGE of a new polymorphism closely linked to the adenomatous polyposis coli region. Human Genetics, 1992, 88, 658-660.	3.8	6
112	Reliability of presymptomatic test for adenomatous polyposis coli. Lancet, The, 1991, 337, 1171-1172.	13.7	5
113	Genetic characterization of the APC locus involved in familial adenomatous polyposis. Gastroenterology, 1991, 101, 154-160.	1.3	17
114	Association of ki-ras mutation with differentiation and tumor-formation pathways in colorectal carcinoma. International Journal of Cancer, 1991, 49, 220-223.	5.1	61
115	Cytogenetic and molecular approaches of polyploidization in colorectal adenocarcinomas. Cancer Genetics and Cytogenetics, 1990, 44, 107-118.	1.0	41
116	Preservation of chromosome and DNA characteristics of human colorectal adenocarcinomas after passage in nude mice. International Journal of Cancer, 1989, 44, 871-878.	5.1	21
117	Chromosomes in Ewing's sarcoma. I. An evaluation of 85 cases and remarkable consistency of $t(11;22)(q24;q12)$ . Cancer Genetics and Cytogenetics, 1988, 32, 229-238.	1.0	488
118	RFLP identified by the anonymous DNA segment OL VII E10 at 18q21.3 (HGM do. D18S8). Nucleic Acids Research, 1987, 15, 1348-1348.	14.5	19