

Sarah L Spain

List of Publications by Year in descending order

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Version: 2024-02-01

37
papers

11,815
citations

159585

30
h-index

315739

38
g-index

43
all docs

43
docs citations

43
times ranked

19860
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|------|-----------|
| 1 | Transcriptomic effects of rs4845604, an IBD and allergy-associated RORC variant, in stimulated ex vivo CD4+ T cells. PLoS ONE, 2021, 16, e0258316. | 2.5 | 0 |
| 2 | Cross-disorder analysis of schizophrenia and 19 immune-mediated diseases identifies shared genetic risk. Human Molecular Genetics, 2019, 28, 3498-3513. | 2.9 | 65 |
| 3 | Fine-mapping inflammatory bowel disease loci to single-variant resolution. Nature, 2017, 547, 173-178. | 27.8 | 473 |
| 4 | Exome-wide association study reveals novel psoriasis susceptibility locus at TNFSF15 and rare protective alleles in genes contributing to type I IFN signalling. Human Molecular Genetics, 2017, 26, 4301-4313. | 2.9 | 41 |
| 5 | Genome-Wide Association Studies Suggest Limited Immune Gene Enrichment in Schizophrenia Compared to 5 Autoimmune Diseases. Schizophrenia Bulletin, 2016, 42, 1176-1184. | 4.3 | 62 |
| 6 | Analysis of five chronic inflammatory diseases identifies 27 new associations and highlights disease-specific patterns at shared loci. Nature Genetics, 2016, 48, 510-518. | 21.4 | 617 |
| 7 | Enhanced meta-analysis and replication studies identify five new psoriasis susceptibility loci. Nature Communications, 2015, 6, 7001. | 12.8 | 156 |
| 8 | Strategies for fine-mapping complex traits. Human Molecular Genetics, 2015, 24, R111-R119. | 2.9 | 191 |
| 9 | Pooled Sequencing of 531 Genes in Inflammatory Bowel Disease Identifies an Associated Rare Variant in BTLN2 and Implicates Other Immune Related Genes. PLoS Genetics, 2015, 11, e1004955. | 3.5 | 59 |
| 10 | Do Genetic Susceptibility Variants Associate with Disease Severity in Early Active Rheumatoid Arthritis?. Journal of Rheumatology, 2015, 42, 1131-1140. | 2.0 | 18 |
| 11 | Investigating the genetic variation underlying episodicity in major depressive disorder: Suggestive evidence for a bipolar contribution. Journal of Affective Disorders, 2014, 155, 81-89. | 4.1 | 15 |
| 12 | Germline mutations affecting the proofreading domains of POLE and POLD1 predispose to colorectal adenomas and carcinomas. Nature Genetics, 2013, 45, 136-144. | 21.4 | 851 |
| 13 | An In-Depth Characterization of the Major Psoriasis Susceptibility Locus Identifies Candidate Susceptibility Alleles within an HLA-C Enhancer Element. PLoS ONE, 2013, 8, e71690. | 2.5 | 45 |
| 14 | Conditional analysis identifies three novel major histocompatibility complex loci associated with psoriasis. Human Molecular Genetics, 2012, 21, 5185-5192. | 2.9 | 58 |
| 15 | Refinement of the associations between risk of colorectal cancer and polymorphisms on chromosomes 1q41 and 12q13.13. Human Molecular Genetics, 2012, 21, 934-946. | 2.9 | 19 |
| 16 | Common variation near CDKN1A, POLD3 and SHROOM2 influences colorectal cancer risk. Nature Genetics, 2012, 44, 770-776. | 21.4 | 210 |
| 17 | Host-microbe interactions have shaped the genetic architecture of inflammatory bowel disease. Nature, 2012, 491, 119-124. | 27.8 | 4,038 |
| 18 | Identification of 15 new psoriasis susceptibility loci highlights the role of innate immunity. Nature Genetics, 2012, 44, 1341-1348. | 21.4 | 848 |

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|----|---|------|-----------|
| 19 | Combined Analysis of Genome-wide Association Studies for Crohn Disease and Psoriasis Identifies Seven Shared Susceptibility Loci. <i>American Journal of Human Genetics</i> , 2012, 90, 636-647. | 6.2 | 290 |
| 20 | Methotrexate polyglutamates as a marker of patient compliance and clinical response in psoriasis: a single-centre prospective study. <i>British Journal of Dermatology</i> , 2012, 167, 165-173. | 1.5 | 21 |
| 21 | Mutations in IL36RN/IL1F5 Are Associated with the Severe Episodic Inflammatory Skin Disease Known as Generalized Pustular Psoriasis. <i>American Journal of Human Genetics</i> , 2011, 89, 432-437. | 6.2 | 468 |
| 22 | Meta-analysis of three genome-wide association studies identifies susceptibility loci for colorectal cancer at 1q41, 3q26.2, 12q13.13 and 20q13.33. <i>Nature Genetics</i> , 2010, 42, 973-977. | 21.4 | 335 |
| 23 | Common variation at the adiponectin locus is not associated with colorectal cancer risk in the UK. <i>Human Molecular Genetics</i> , 2009, 18, 1889-1892. | 2.9 | 31 |
| 24 | Colorectal Cancer Risk Is Not Associated with Increased Levels of Homozygosity in a Population from the United Kingdom. <i>Cancer Research</i> , 2009, 69, 7422-7429. | 0.9 | 36 |
| 25 | Low penetrance breast cancer predisposition SNPs are site specific. <i>Breast Cancer Research and Treatment</i> , 2009, 117, 151-159. | 2.5 | 37 |
| 26 | A mitotic recombination map proximal to the APC locus on chromosome 5q and assessment of influences on colorectal cancer risk. <i>BMC Medical Genetics</i> , 2009, 10, 54. | 2.1 | 18 |
| 27 | Deciphering the genetics of hereditary non-syndromic colorectal cancer. <i>European Journal of Human Genetics</i> , 2008, 16, 1477-1486. | 2.8 | 31 |
| 28 | A genome-wide association study identifies colorectal cancer susceptibility loci on chromosomes 10p14 and 8q23.3. <i>Nature Genetics</i> , 2008, 40, 623-630. | 21.4 | 514 |
| 29 | Common genetic variants at the CRAC1 (HMPS) locus on chromosome 15q13.3 influence colorectal cancer risk. <i>Nature Genetics</i> , 2008, 40, 26-28. | 21.4 | 277 |
| 30 | Meta-analysis of genome-wide association data identifies four new susceptibility loci for colorectal cancer. <i>Nature Genetics</i> , 2008, 40, 1426-1435. | 21.4 | 498 |
| 31 | Refinement of the basis and impact of common 11q23.1 variation to the risk of developing colorectal cancer. <i>Human Molecular Genetics</i> , 2008, 17, 3720-3727. | 2.9 | 61 |
| 32 | Pregnancy does not influence colonic polyp multiplicity but may modulate upper gastrointestinal disease in patients with FAP. <i>Journal of Medical Genetics</i> , 2007, 44, 541-544. | 3.2 | 1 |
| 33 | A genome-wide association study shows that common alleles of SMAD7 influence colorectal cancer risk. <i>Nature Genetics</i> , 2007, 39, 1315-1317. | 21.4 | 463 |
| 34 | A genome-wide association scan of tag SNPs identifies a susceptibility variant for colorectal cancer at 8q24.21. <i>Nature Genetics</i> , 2007, 39, 984-988. | 21.4 | 754 |
| 35 | Disease severity and genetic pathways in attenuated familial adenomatous polyposis vary greatly but depend on the site of the germline mutation. <i>Gut</i> , 2006, 55, 1440-1448. | 12.1 | 87 |
| 36 | Evidence for a colorectal cancer susceptibility locus on chromosome 3q21-q24 from a high-density SNP genome-wide linkage scan. <i>Human Molecular Genetics</i> , 2006, 15, 2903-2910. | 2.9 | 52 |

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|----|--|-----|-----------|
| 37 | Evidence of Linkage to Chromosome 9q22.33 in Colorectal Cancer Kindreds from the United Kingdom. Cancer Research, 2006, 66, 5003-5006. | 0.9 | 51 |