# Simon N Stacey

List of Publications by Year
in descending order



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|---|---|---|---|
| 38 papers | 10,142 citations | 136740 32 h-index | 315357 38 g-index |
| 38 all docs | 38 docs citations | 38 times ranked | 16316 citing authors |

| # | Article | IF | Citations |
|---|---|---|---|
| 1 | Loss-of-Function Variants in the Tumor-Suppressor Gene PTPN14 Confer Increased Cancer Risk. Cancer Research, 2021, 81, 1954-1964. | 0.4 | 15 |
| 2 | Large-scale integration of the plasma proteome with genetics and disease. Nature Genetics, 2021, 53, 1712-1721. | 9.4 | 340 |
| 3 | Genome-wide meta-analysis identifies eight new susceptibility loci for cutaneous squamous cell carcinoma. Nature Communications, 2020, 11, 820. | 5.8 | 30 |
| 4 | Characterizing mutagenic effects of recombination through a sequence-level genetic map. Science, 2019, 363, . | 6.0 | 252 |
| 5 | Genome-wide associations for benign prostatic hyperplasia reveal a genetic correlation with serum levels of PSA. Nature Communications, 2018, 9, 4568. | 5.8 | 44 |
| 6 | Multiple transmissions of de novo mutations in families. Nature Genetics, 2018, 50, 1674-1680. | 9.4 | 89 |
| 7 | Insights into imprinting from parent-of-origin phased methylomes and transcriptomes. Nature Genetics, 2018, 50, 1542-1552. | 9.4 | 94 |
| 8 | Variants associating with uterine leiomyoma highlight genetic background shared by various cancers and hormone-related traits. Nature Communications, 2018, 9, 3636. | 5.8 | 74 |
| 9 | Association of BRCA2 K3326* With Small Cell Lung Cancer and Squamous Cell Cancer of the Skin. Journal of the National Cancer Institute, 2018, 110, 967-974. | 3.0 | 29 |
| 10 | Whole genome characterization of sequence diversity of 15,220 Icelanders. Scientific Data, 2017, 4, 170115. | 2.4 | 98 |
| 11 | Parental influence on human germline de novo mutations in 1,548 trios from Iceland. Nature, 2017, 549, 519-522. | 13.7 | 410 |
| 12 | Cross-Cancer Genome-Wide Analysis of Lung, Ovary, Breast, Prostate, and Colorectal Cancer Reveals Novel Pleiotropic Associations. Cancer Research, 2016, 76, 5103-5114. | 0.4 | 100 |
| 13 | Insertion of an SVA-E retrotransposon into theCASP8gene is associated with protection against prostate cancer. Human Molecular Genetics, 2016, 25, 1008-1018. | 1.4 | 22 |
| 14 | Loss-of-function variants in ATM confer risk of gastric cancer. Nature Genetics, 2015, 47, 906-910. | 9.4 | 155 |
| 15 | Large-scale whole-genome sequencing of the Icelandic population. Nature Genetics, 2015, 47, 435-444. | 9.4 | 663 |
| 16 | New basal cell carcinoma susceptibility loci. Nature Communications, 2015, 6, 6825. | 5.8 | 59 |
| 17 | Germline sequence variants in TGM3 and RGS22 confer risk of basal cell carcinoma. Human Molecular Genetics, 2014, 23, 3045-3053. | 1.4 | 48 |
| 18 | A Polymorphism in IRF4 Affects Human Pigmentation through a Tyrosinase-Dependent MITF/TFAP2A Pathway. Cell, 2013, 155, 1022-1033. | 13.5 | 184 |

| # | Article | IF | Citations |
|---|---|---|---|
| 19 | A common variant at 8q24.21 is associated with renal cell cancer. Nature Communications, 2013, 4, 2776. | 5.8 | 56 |
| 20 | A study based on whole-genome sequencing yields a rare variant at 8q24 associated with prostate cancer. Nature Genetics, 2012, 44, 1326-1329. | 9.4 | 178 |
| 21 | Discovery of common variants associated with low TSH levels and thyroid cancer risk. Nature Genetics, 2012, 44, 319-322. | 9.4 | 208 |
| 22 | A germline variant in the TP53 polyadenylation signal confers cancer susceptibility. Nature Genetics, 2011, 43, 1098-1103. | 9.4 | 251 |
| 23 | Genetic Correction of PSA Values Using Sequence Variants Associated with PSA Levels. Science Translational Medicine, 2010, 2, 62ra92. | 5.8 | 140 |
| 24 | Ancestry-Shift Refinement Mapping of the C6orf97-ESR1 Breast Cancer Susceptibility Locus. PLoS Genetics, 2010, 6, e1001029. | 1.5 | 82 |
| 25 | Evaluation of Association of HNF1B Variants with Diverse Cancers: Collaborative Analysis of Data from 19 Genome-Wide Association Studies. PLoS ONE, 2010, 5, e10858. | 1.1 | 28 |
| 26 | Parental origin of sequence variants associated with complex diseases. Nature, 2009, 462, 868-874. | 13.7 | 521 |
| 27 | Sequence variants at the TERT-CLPTM1L locus associate with many cancer types. Nature Genetics, 2009, 41, 221-227. | 9.4 | 572 |
| 28 | New common variants affecting susceptibility to basal cell carcinoma. Nature Genetics, 2009, 41, 909-914. | 9.4 | 303 |
| 29 | A variant associated with nicotine dependence, lung cancer and peripheral arterial disease. Nature, 2008, 452, 638-642. | 13.7 | 1,399 |
| 30 | Common variants on chromosome 5p12 confer susceptibility to estrogen receptorâ€“positive breast cancer. Nature Genetics, 2008, 40, 703-706. | 9.4 | 412 |
| 31 | Two newly identified genetic determinants of pigmentation in Europeans. Nature Genetics, 2008, 40, 835-837. | 9.4 | 331 |
| 32 | ASIP and TYR pigmentation variants associate with cutaneous melanoma and basal cell carcinoma. Nature Genetics, 2008, 40, 886-891. | 9.4 | 306 |
| 33 | Sequence variant on 8q24 confers susceptibility to urinary bladder cancer. Nature Genetics, 2008, 40, 1307-1312. | 9.4 | 377 |
| 34 | Common variants on 1p36 and 1q42 are associated with cutaneous basal cell carcinoma but not with melanoma or pigmentation traits. Nature Genetics, 2008, 40, 1313-1318. | 9.4 | 111 |
| 35 | Genetic determinants of hair, eye and skin pigmentation in Europeans. Nature Genetics, 2007, 39, 1443-1452. | 9.4 | 659 |
| 36 | Two variants on chromosome 17 confer prostate cancer risk, and the one in TCF2 protects against type 2 diabetes. Nature Genetics, 2007, 39, 977-983. | 9.4 | 670 |

| # | Article | IF | Citations |
|---|---|---|---|
| 37 | Common variants on chromosomes 2q35 and 16q12 confer susceptibility to estrogen receptorâ€“positive breast cancer. Nature Genetics, 2007, 39, 865-869. | 9.4 | 774 |
| 38 | The BARD1 Cys557Ser Variant and Breast Cancer Risk in Iceland. PLoS Medicine, 2006, 3, e217. | 3.9 | 58 |