# Gord Glendon

List of Publications by Year
in descending order

| 56 | 6,466 | 136950 | 144013 |
|---|---|---|---|
| papers | citations | 32 | 57 |
| | | h-index | g-index |

| 58 | 58 | 58 | 9322 |
|---|---|---|---|
| all docs | docs citations | times ranked | citing authors |

| # | Article | IF | Citations |
|---|---------|-----|-----------|
| 1 | Association analysis identifies 65 new breast cancer risk loci. Nature, 2017, 551, 92-94. | 27.8 | 1,099 |
| 2 | Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes. American Journal of Human Genetics, 2019, 104, 21-34. | 6.2 | 711 |
| 3 | Breast Cancer Risk Genes â€" Association Analysis in More than 113,000 Women. New England Journal of Medicine, 2021, 384, 428-439. | 27.0 | 532 |
| 4 | Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. Nature Genetics, 2015, 47, 373-380. | 21.4 | 513 |
| 5 | Prediction of Breast Cancer Risk Based on Profiling With Common Genetic Variants. Journal of the National Cancer Institute, 2015, 107, . | 6.3 | 428 |
| 6 | Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. Nature Genetics, 2017, 49, 1767-1778. | 21.4 | 289 |
| 7 | Identification of six new susceptibility loci for invasive epithelial ovarian cancer. Nature Genetics, 2015, 47, 164-171. | 21.4 | 221 |
| 8 | *PALB2*, *CHEK2* and *ATM* rare variants and cancer risk: data from COGS. Journal of Medical Genetics, 2016, 53, 800-811. | 3.2 | 174 |
| 9 | Common Breast Cancer Susceptibility Alleles and the Risk of Breast Cancer for *BRCA1* and *BRCA2* Mutation Carriers: Implications for Risk Prediction. Cancer Research, 2010, 70, 9742-9754. | 0.9 | 169 |
| 10 | Prediction of Breast and Prostate Cancer Risks in Male *BRCA1* and *BRCA2* Mutation Carriers Using Polygenic Risk Scores. Journal of Clinical Oncology, 2017, 35, 2240-2250. | 1.6 | 152 |
| 11 | Breast cancer risk variants at 6q25 display different phenotype associations and regulate ESR1, RMND1 and CCDC170. Nature Genetics, 2016, 48, 374-386. | 21.4 | 125 |
| 12 | Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. Nature Genetics, 2020, 52, 56-73. | 21.4 | 120 |
| 13 | 10-year performance of four models of breast cancer risk: a validation study. Lancet Oncology, The, 2019, 20, 504-517. | 10.7 | 116 |
| 14 | Combined genetic and splicing analysis of BRCA1 c.[594-2A&gt;C; 641A&gt;G] highlights the relevance of naturally occurring in-frame transcripts for developing disease gene variant classification algorithms. Human Molecular Genetics, 2016, 25, 2256-2268. | 2.9 | 106 |
| 15 | Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. Nature Communications, 2014, 5, 4999. | 12.8 | 105 |
| 16 | Height and Breast Cancer Risk: Evidence From Prospective Studies and Mendelian Randomization. Journal of the National Cancer Institute, 2015, 107, djv219. | 6.3 | 99 |
| 17 | No evidence that protein truncating variants in *BRIP1* are associated with breast cancer risk: implications for gene panel testing. Journal of Medical Genetics, 2016, 53, 298-309. | 3.2 | 94 |
| 18 | Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. Nature Communications, 2016, 7, 11375. | 12.8 | 93 |

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|---|---------|----|-----------|
| 19 | Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer. Nature Communications, 2019, 10, 1741. | 12.8 | 90 |
| 20 | Cancer Risks Associated With *BRCA1* and *BRCA2* Pathogenic Variants. Journal of Clinical Oncology, 2022, 40, 1529-1541. | 1.6 | 90 |
| 21 | Male breast cancer in BRCA1 and BRCA2 mutation carriers: pathology data from the Consortium of Investigators of Modifiers of BRCA1/2. Breast Cancer Research, 2016, 18, 15. | 5.0 | 88 |
| 22 | Polygenic risk scores and breast and epithelial ovarian cancer risks for carriers of BRCA1 and BRCA2 pathogenic variants. Genetics in Medicine, 2020, 22, 1653-1666. | 2.4 | 82 |
| 23 | Fine-Scale Mapping of the 5q11.2 Breast Cancer Locus Reveals at Least Three Independent Risk Variants Regulating MAP3K1. American Journal of Human Genetics, 2015, 96, 5-20. | 6.2 | 76 |
| 24 | *BRCA2* Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. Cancer Research, 2017, 77, 2789-2799. | 0.9 | 75 |
| 25 | Identification of Novel Genetic Markers of Breast Cancer Survival. Journal of the National Cancer Institute, 2015, 107, . | 6.3 | 56 |
| 26 | Common non-synonymous SNPs associated with breast cancer susceptibility: findings from the Breast Cancer Association Consortium. Human Molecular Genetics, 2014, 23, 6096-6111. | 2.9 | 53 |
| 27 | Genome-wide association study of germline variants and breast cancer-specific mortality. British Journal of Cancer, 2019, 120, 647-657. | 6.4 | 52 |
| 28 | Age-specific breast cancer risk by body mass index and familial risk: prospective family study cohort (ProF-SC). Breast Cancer Research, 2018, 20, 132. | 5.0 | 51 |
| 29 | Characterization of the Cancer Spectrum in Men With Germline *BRCA1* and *BRCA2* Pathogenic Variants. JAMA Oncology, 2020, 6, 1218. | 7.1 | 48 |
| 30 | Body mass index and breast cancer survival: a Mendelian randomization analysis. International Journal of Epidemiology, 2017, 46, 1814-1822. | 1.9 | 45 |
| 31 | Genetic Predisposition to In Situ and Invasive Lobular Carcinoma of the Breast. PLoS Genetics, 2014, 10, e1004285. | 3.5 | 39 |
| 32 | Association of Genomic Domains in *BRCA1* and *BRCA2* with Prostate Cancer Risk and Aggressiveness. Cancer Research, 2020, 80, 624-638. | 0.9 | 39 |
| 33 | Recreational Physical Activity Is Associated with Reduced Breast Cancer Risk in Adult Women at High Risk for Breast Cancer: A Cohort Study of Women Selected for Familial and Genetic Risk. Cancer Research, 2020, 80, 116-125. | 0.9 | 37 |
| 34 | Transcriptome-wide association study of breast cancer risk by estrogen-receptor status. Genetic Epidemiology, 2020, 44, 442-468. | 1.3 | 32 |
| 35 | Height and Body Mass Index as Modifiers of Breast Cancer Risk in *BRCA1*/*2* Mutation Carriers: A Mendelian Randomization Study. Journal of the National Cancer Institute, 2019, 111, 350-364. | 6.3 | 30 |
| 36 | The FANCM:p.Arg658* truncating variant is associated with risk of triple-negative breast cancer. Npj Breast Cancer, 2019, 5, 38. | 5.2 | 28 |

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|---|---------|-----|-----------|
| 37 | Common germline polymorphisms associated with breast cancer-specific survival. Breast Cancer Research, 2015, 17, 58. | 5.0 | 26 |
| 38 | *BRCA1* and *BRCA2* pathogenic sequence variants in women of African origin or ancestry. Human Mutation, 2019, 40, 1781-1796. | 2.5 | 26 |
| 39 | RAD51B in Familial Breast Cancer. PLoS ONE, 2016, 11, e0153788. | 2.5 | 26 |
| 40 | Polygenic risk modeling for prediction of epithelial ovarian cancer risk. European Journal of Human Genetics, 2022, 30, 349-362. | 2.8 | 23 |
| 41 | Mendelian randomisation study of height and body mass index as modifiers of ovarian cancer risk in 22,588 BRCA1 and BRCA2 mutation carriers. British Journal of Cancer, 2019, 121, 180-192. | 6.4 | 19 |
| 42 | Breast and Prostate Cancer Risks for Male *BRCA1* and *BRCA2* Pathogenic Variant Carriers Using Polygenic Risk Scores. Journal of the National Cancer Institute, 2022, 114, 109-122. | 6.3 | 19 |
| 43 | Breast cancer risks associated with missense variants in breast cancer susceptibility genes. Genome Medicine, 2022, 14, 51. | 8.2 | 19 |
| 44 | No clinical utility of KRAS variant rs61764370 for ovarian or breast cancer. Gynecologic Oncology, 2016, 141, 386-401. | 1.4 | 18 |
| 45 | Association of breast cancer risk in BRCA1 and BRCA2 mutation carriers with genetic variants showing differential allelic expression: identification of a modifier of breast cancer risk at locus 11q22.3. Breast Cancer Research and Treatment, 2017, 161, 117-134. | 2.5 | 18 |
| 46 | 2q36.3 is associated with prognosis for oestrogen receptor-negative breast cancer patients treated with chemotherapy. Nature Communications, 2014, 5, 4051. | 12.8 | 16 |
| 47 | Genetic variation at CYP3A is associated with age at menarche and breast cancer risk: a case-control study. Breast Cancer Research, 2014, 16, R51. | 5.0 | 14 |
| 48 | Inherited variants in the inner centromere protein (INCENP) gene of the chromosomal passenger complex contribute to the susceptibility of ER-negative breast cancer. Carcinogenesis, 2015, 36, 256-271. | 2.8 | 14 |
| 49 | TP53-based interaction analysis identifies cis-eQTL variants for TP53BP2, FBXO28, and FAM53A that associate with survival and treatment outcome in breast cancer. Oncotarget, 2017, 8, 18381-18398. | 1.8 | 14 |
| 50 | Psychosocial Adjustment in School-age Girls With a Family History of Breast Cancer. Pediatrics, 2015, 136, 927-937. | 2.1 | 13 |
| 51 | Does perceived risk predict breast cancer screening use? Findings from a prospective cohort study of female relatives from the Ontario site of the Breast Cancer Family Registry. Breast, 2014, 23, 482-488. | 2.2 | 10 |
| 52 | A randomized controlled trial of a supportive expressive group intervention for women with a family history of breast cancer. Psycho-Oncology, 2018, 27, 2645-2653. | 2.3 | 10 |
| 53 | Mendelian randomisation study of smoking exposure in relation to breast cancer risk. British Journal of Cancer, 2021, 125, 1135-1145. | 6.4 | 9 |
| 54 | Genetic variation in the immunosuppression pathway genes and breast cancer susceptibility: a pooled analysis of 42,510 cases and 40,577 controls from the Breast Cancer Association Consortium. Human Genetics, 2016, 135, 137-154. | 3.8 | 8 |

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| 55 | rs2735383, located at a microRNA binding site in the 3â€™UTR of NBS1, is not associated with breast cancer risk. Scientific Reports, 2016, 6, 36874. | 3.3 | 2 |
| 56 | Recreational Physical Activity and Outcomes After Breast Cancer in Women at High Familial Risk. JNCI Cancer Spectrum, 2021, 5, pkab090. | 2.9 | 1 |