

# Noralane M Lindor

## List of Publications by Citations

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**Version:** 2024-04-20

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

74  
papers

3,129  
citations

27  
h-index

55  
g-index

74  
ext. papers

3,839  
ext. citations

7.5  
avg, IF

4.48  
L-index

#	Paper	IF	Citations
74	Pathology of breast and ovarian cancers among BRCA1 and BRCA2 mutation carriers: results from the Consortium of Investigators of Modifiers of BRCA1/2 (CIMBA). <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2012</b> , 21, 134-47	4	411
73	Association of type and location of BRCA1 and BRCA2 mutations with risk of breast and ovarian cancer. <i>JAMA - Journal of the American Medical Association</i> , <b>2015</b> , 313, 1347-61	27.4	286
72	The Role of Risk-Reducing Surgery in Hereditary Breast and Ovarian Cancer. <i>New England Journal of Medicine</i> , <b>2016</b> , 374, 454-68	59.2	179
71	Identification of six new susceptibility loci for invasive epithelial ovarian cancer. <i>Nature Genetics</i> , <b>2015</b> , 47, 164-71	36.3	177
70	Point Mutations in Exon 1B of APC Reveal Gastric Adenocarcinoma and Proximal Polyposis of the Stomach as a Familial Adenomatous Polyposis Variant. <i>American Journal of Human Genetics</i> , <b>2016</b> , 98, 830-842	11	153
69	Mutational spectrum in a worldwide study of 29,700 families with BRCA1 or BRCA2 mutations. <i>Human Mutation</i> , <b>2018</b> , 39, 593-620	4.7	138
68	Association of aspirin and NSAID use with risk of colorectal cancer according to genetic variants. <i>JAMA - Journal of the American Medical Association</i> , <b>2015</b> , 313, 1133-42	27.4	135
67	Identification of susceptibility loci for colorectal cancer in a genome-wide meta-analysis. <i>Human Molecular Genetics</i> , <b>2014</b> , 23, 4729-37	5.6	107
66	A new GWAS and meta-analysis with 1000Genomes imputation identifies novel risk variants for colorectal cancer. <i>Scientific Reports</i> , <b>2015</b> , 5, 10442	4.9	94
65	Breast cancer risk variants at 6q25 display different phenotype associations and regulate ESR1, RMND1 and CCDC170. <i>Nature Genetics</i> , <b>2016</b> , 48, 374-86	36.3	93
64	Association analyses identify 31 new risk loci for colorectal cancer susceptibility. <i>Nature Communications</i> , <b>2019</b> , 10, 2154	17.4	81
63	Returning a Research Participant's Genomic Results to Relatives: Analysis and Recommendations. <i>Journal of Law, Medicine and Ethics</i> , <b>2015</b> , 43, 440-63	1.2	67
62	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. <i>Nature Communications</i> , <b>2016</b> , 7, 11375	17.4	64
61	Outcome of Whole Exome Sequencing for Diagnostic Odyssey Cases of an Individualized Medicine Clinic: The Mayo Clinic Experience. <i>Mayo Clinic Proceedings</i> , <b>2016</b> , 91, 297-307	6.4	63
60	BRCA1/2 sequence variants of uncertain significance: a primer for providers to assist in discussions and in medical management. <i>Oncologist</i> , <b>2013</b> , 18, 518-24	5.7	61
59	Pro-inflammatory fatty acid profile and colorectal cancer risk: A Mendelian randomisation analysis. <i>European Journal of Cancer</i> , <b>2017</b> , 84, 228-238	7.5	56
58	Mendelian Randomization Study of Body Mass Index and Colorectal Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2015</b> , 24, 1024-31	4	54

57	Mendelian randomisation implicates hyperlipidaemia as a risk factor for colorectal cancer. <i>International Journal of Cancer</i> , <b>2017</b> , 140, 2701-2708	7.5	50
56	PMS2 monoallelic mutation carriers: the known unknown. <i>Genetics in Medicine</i> , <b>2016</b> , 18, 13-9	8.1	42
55	Mendelian randomisation analysis strongly implicates adiposity with risk of developing colorectal cancer. <i>British Journal of Cancer</i> , <b>2016</b> , 115, 266-72	8.7	39
54	Assessment of the Clinical Relevance of BRCA2 Missense Variants by Functional and Computational Approaches. <i>American Journal of Human Genetics</i> , <b>2018</b> , 102, 233-248	11	38
53	Association between body mass index and mortality for colorectal cancer survivors: overall and by tumor molecular phenotype. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2015</b> , 24, 1229-38	4	38
52	Comprehensive annotation of BRCA1 and BRCA2 missense variants by functionally validated sequence-based computational prediction models. <i>Genetics in Medicine</i> , <b>2019</b> , 21, 71-80	8.1	36
51	Whole-Genome Sequencing in Healthy People. <i>Mayo Clinic Proceedings</i> , <b>2017</b> , 92, 159-172	6.4	33
50	DNA glycosylases involved in base excision repair may be associated with cancer risk in BRCA1 and BRCA2 mutation carriers. <i>PLoS Genetics</i> , <b>2014</b> , 10, e1004256	6	33
49	Genome-Wide Interaction Analyses between Genetic Variants and Alcohol Consumption and Smoking for Risk of Colorectal Cancer. <i>PLoS Genetics</i> , <b>2016</b> , 12, e1006296	6	30
48	Variation at 2q35 (PNKD and TMBIM1) influences colorectal cancer risk and identifies a pleiotropic effect with inflammatory bowel disease. <i>Human Molecular Genetics</i> , <b>2016</b> , 25, 2349-2359	5.6	27
47	Meta-analysis of genome-wide association studies identifies common susceptibility polymorphisms for colorectal and endometrial cancer near SH2B3 and TSHZ1. <i>Scientific Reports</i> , <b>2015</b> , 5, 17369	4.9	27
46	Assessing associations between the AURKA-HMMR-TPX2-TUBG1 functional module and breast cancer risk in BRCA1/2 mutation carriers. <i>PLoS ONE</i> , <b>2015</b> , 10, e0120020	3.7	26
45	A genome-wide association study for colorectal cancer identifies a risk locus in 14q23.1. <i>Human Genetics</i> , <b>2015</b> , 134, 1249-1262	6.3	25
44	Lack of association between screening interval and cancer stage in Lynch syndrome may be accounted for by over-diagnosis; a prospective Lynch syndrome database report. <i>Hereditary Cancer in Clinical Practice</i> , <b>2019</b> , 17, 8	2.3	24
43	Potential impact of family history-based screening guidelines on the detection of early-onset colorectal cancer. <i>Cancer</i> , <b>2020</b> , 126, 3013-3020	6.4	23
42	Preferences Regarding Return of Genomic Results to Relatives of Research Participants, Including after Participant Death: Empirical Results from a Cancer Biobank. <i>Journal of Law, Medicine and Ethics</i> , <b>2015</b> , 43, 464-75	1.2	21
41	Candidate genetic modifiers for breast and ovarian cancer risk in BRCA1 and BRCA2 mutation carriers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2015</b> , 24, 308-16	4	20
40	Molecular Biomarkers for the Evaluation of Colorectal Cancer: Guideline Summary From the American Society for Clinical Pathology, College of American Pathologists, Association for Molecular Pathology, and American Society of Clinical Oncology. <i>Journal of Oncology Practice</i> , <b>2017</b> , 13, 222-227	3.1	20

39	A preliminary investigation of genetic counselors' information needs when receiving a variant of uncertain significance result: a mixed methods study. <i>Genetics in Medicine</i> , <b>2015</b> , 17, 739-46	8.1	20
38	The Return of Actionable Variants Empirical (RAVE) Study, a Mayo Clinic Genomic Medicine Implementation Study: Design and Initial Results. <i>Mayo Clinic Proceedings</i> , <b>2018</b> , 93, 1600-1610	6.4	20
37	Targeted sequencing of established and candidate colorectal cancer genes in the Colon Cancer Family Registry Cohort. <i>Oncotarget</i> , <b>2017</b> , 8, 93450-93463	3.3	18
36	Non-BRCA familial breast cancer: review of reported pathology and molecular findings. <i>Pathology</i> , <b>2017</b> , 49, 363-370	1.6	17
35	Prediagnostic alcohol consumption and colorectal cancer survival: The Colon Cancer Family Registry. <i>Cancer</i> , <b>2017</b> , 123, 1035-1043	6.4	17
34	Survival by colon cancer stage and screening interval in Lynch syndrome: a prospective Lynch syndrome database report. <i>Hereditary Cancer in Clinical Practice</i> , <b>2019</b> , 17, 28	2.3	16
33	CYP24A1 variant modifies the association between use of oestrogen plus progestogen therapy and colorectal cancer risk. <i>British Journal of Cancer</i> , <b>2016</b> , 114, 221-9	8.7	16
32	Central pontine myelinolysis as a complication of partial ornithine carbamoyl transferase deficiency. <i>American Journal of Medical Genetics Part A</i> , <b>1995</b> , 60, 210-3		16
31	No clinical utility of KRAS variant rs61764370 for ovarian or breast cancer. <i>Gynecologic Oncology</i> , <b>2016</b> , 141, 386-401	4.9	15
30	Long-term weight loss after colorectal cancer diagnosis is associated with lower survival: The Colon Cancer Family Registry. <i>Cancer</i> , <b>2017</b> , 123, 4701-4708	6.4	15
29	Identifying Novel Susceptibility Genes for Colorectal Cancer Risk From a Transcriptome-Wide Association Study of 125,478 Subjects. <i>Gastroenterology</i> , <b>2021</b> , 160, 1164-1178.e6	13.3	15
28	DNA mismatch repair protein deficient non-neoplastic colonic crypts: a novel indicator of Lynch syndrome. <i>Modern Pathology</i> , <b>2018</b> , 31, 1608-1618	9.8	14
27	Developing a Process for Returning Medically Actionable Genomic Variants to Latino Patients in a Federally Qualified Health Center. <i>Public Health Genomics</i> , <b>2018</b> , 21, 77-84	1.9	14
26	Colorectal cancer and self-reported tooth agenesis. <i>Hereditary Cancer in Clinical Practice</i> , <b>2014</b> , 12, 7	2.3	13
25	Genomic medicine and incidental findings: balancing actionability and patient autonomy. <i>Mayo Clinic Proceedings</i> , <b>2014</b> , 89, 718-21	6.4	11
24	Myhre-LAPs syndrome and intubation related airway stenosis: keys to diagnosis and critical therapeutic interventions. <i>American Journal of Otolaryngology - Head and Neck Medicine and Surgery</i> , <b>2015</b> , 36, 636-41	2.8	9
23	Whole-Exome Sequencing of 10 Scientists: Evaluation of the Process and Outcomes. <i>Mayo Clinic Proceedings</i> , <b>2015</b> , 90, 1327-37	6.4	9
22	Transcriptome-wide association study of breast cancer risk by estrogen-receptor status. <i>Genetic Epidemiology</i> , <b>2020</b> , 44, 442-468	2.6	9

21	Association of a let-7 miRNA binding region of TGFBR1 with hereditary mismatch repair proficient colorectal cancer (MSS HNPCC). <i>Carcinogenesis</i> , <b>2016</b> , 37, 751-8	4.6	9
20	Physical activity and the risk of colorectal cancer in Lynch syndrome. <i>International Journal of Cancer</i> , <b>2018</b> , 143, 2250-2260	7.5	9
19	Leptin gene variants and colorectal cancer risk: Sex-specific associations. <i>PLoS ONE</i> , <b>2018</b> , 13, e0206519	3.7	9
18	Current Approaches to Cancer Genetic Counseling Services for Spanish-Speaking Patients. <i>Journal of Immigrant and Minority Health</i> , <b>2019</b> , 21, 434-437	2.2	8
17	How well do whole exome sequencing results correlate with medical findings? A study of 89 Mayo Clinic Biobank samples. <i>Frontiers in Genetics</i> , <b>2015</b> , 6, 244	4.5	8
16	Family history of colorectal cancer is not associated with colorectal cancer survival regardless of microsatellite instability status. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2014</b> , 23, 1700-4	4	8
15	Fine-Scale Mapping at 9p22.2 Identifies Candidate Causal Variants That Modify Ovarian Cancer Risk in BRCA1 and BRCA2 Mutation Carriers. <i>PLoS ONE</i> , <b>2016</b> , 11, e0158801	3.7	7
14	Germline miRNA DNA variants and the risk of colorectal cancer by subtype. <i>Genes Chromosomes and Cancer</i> , <b>2017</b> , 56, 177-184	5	6
13	Contribution of mRNA Splicing to Mismatch Repair Gene Sequence Variant Interpretation. <i>Frontiers in Genetics</i> , <b>2020</b> , 11, 798	4.5	5
12	Risks of Colorectal Cancer and Cancer-Related Mortality in Familial Colorectal Cancer Type X and Lynch Syndrome Families. <i>Journal of the National Cancer Institute</i> , <b>2019</b> , 111, 675-683	9.7	5
11	From the laboratory to the clinic: sharing BRCA VUS reclassification tools with practicing genetics professionals. <i>Journal of Community Genetics</i> , <b>2018</b> , 9, 209-215	2.5	5
10	Estimating cumulative risks for breast cancer for carriers of variants in uncommon genes. <i>Familial Cancer</i> , <b>2016</b> , 15, 367-70	3	3
9	Lynch syndrome 101 (years, that is). <i>American Society of Clinical Oncology Educational Book / ASCO American Society of Clinical Oncology Meeting</i> , <b>2014</b> , 27-32	7.1	2
8	Challenges in returning results in a genomic medicine implementation study: the Return of Actionable Variants Empirical (RAVE) study. <i>Npj Genomic Medicine</i> , <b>2020</b> , 5, 19	6.2	2
7	Genetic susceptibility markers for a breast-colorectal cancer phenotype: Exploratory results from genome-wide association studies. <i>PLoS ONE</i> , <b>2018</b> , 13, e0196245	3.7	2
6	Increasing access to individualized medicine: a matched-cohort study examining Latino participant experiences of genomic screening. <i>Genetics in Medicine</i> , <b>2021</b> , 23, 934-941	8.1	2
5	Clinical verification of genetic results returned to research participants: findings from a Colon Cancer Family Registry. <i>Molecular Genetics &amp; Genomic Medicine</i> , <b>2017</b> , 5, 700-708	2.3	1
4	Exploratory Genome-Wide Interaction Analysis of Nonsteroidal Anti-inflammatory Drugs and Predicted Gene Expression on Colorectal Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2020</b> , 29, 1800-1808	4	1

3	Penetrance and outcomes at 1-year following return of actionable variants identified by genome sequencing. <i>Genetics in Medicine</i> , <b>2021</b> , 23, 1192-1201	8.1	1
2	Experiences of Latino Participants Receiving Neutral Genomic Screening Results: A Qualitative Study. <i>Public Health Genomics</i> , <b>2021</b> , 24, 44-53	1.9	1
1	Response to ten Broeke and Nielsen. <i>Genetics in Medicine</i> , <b>2015</b> , 17, 684-5	8.1	