

Yael Laitman

List of Publications by Year in descending order

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Version: 2024-02-01

73
papers

3,411
citations

257450

24
h-index

155660

55
g-index

73
all docs

73
docs citations

73
times ranked

6946
citing authors

#	ARTICLE	IF	CITATIONS
1	Re-evaluating cancer risks associated with the CHEK2 p.Ser428Phe Ashkenazi Jewish founder pathogenic variant. <i>Familial Cancer</i> , 2022, 21, 305-308.	1.9	4
2	Population-based screening of Uruguayan Ashkenazi Jews for recurrent <i>BRCA1</i> and <i>BRCA2</i> pathogenic sequence variants. <i>Molecular Genetics & Genomic Medicine</i> , 2022, , e1928.	1.2	1
3	Common Susceptibility Loci for Male Breast Cancer. <i>Journal of the National Cancer Institute</i> , 2021, 113, 453-461.	6.3	12
4	Locoregional Treatments and Ipsilateral Breast Cancer Recurrence Rates in <i>BRCA1/2</i> Mutation Carriers. <i>International Journal of Radiation Oncology Biology Physics</i> , 2021, 109, 1332-1340.	0.8	15
5	Time trends in uptake rates of risk-reducing mastectomy in Israeli asymptomatic <i>BRCA1</i> and <i>BRCA2</i> mutation carriers. <i>Breast Cancer Research and Treatment</i> , 2021, 185, 391-399.	2.5	7
6	Double heterozygosity for <i>TP53</i> and <i>BRCA1</i> mutations: clinical implications in populations with founder mutations. <i>Breast Cancer Research and Treatment</i> , 2021, 186, 259-263.	2.5	2
7	Re-evaluating the pathogenicity of the c.783+2T>C <i>BAP1</i> germline variant. <i>Human Mutation</i> , 2021, 42, 592-599.	2.5	3
8	Yield of targeted genotyping for the recurring pathogenic variants in cancer susceptibility genes in a healthy, multiethnic Israeli population. <i>Cancer</i> , 2021, 127, 3599-3604.	4.1	1
9	The spectrum of tumors harboring <i>BAP1</i> gene alterations. <i>Cancer Genetics</i> , 2021, 256-257, 31-35.	0.4	12
10	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020, 52, 56-73.	21.4	120
11	De novo pathogenic germline variant in <i>PALB2</i> in a patient with pancreatic cancer. <i>Familial Cancer</i> , 2020, 19, 193-196.	1.9	5
12	Breast cancer surveillance for <i>BRCA1/2</i> mutation carriers – is “early detection” early enough?. <i>Breast</i> , 2020, 49, 81-86.	2.2	10
13	Polygenic risk scores and breast and epithelial ovarian cancer risks for carriers of <i>BRCA1</i> and <i>BRCA2</i> pathogenic variants. <i>Genetics in Medicine</i> , 2020, 22, 1653-1666.	2.4	82
14	Age at diagnosis of cancer in 185 <i>BRCA1</i> mutation carriers of diverse ethnicities: tentative evidence for modifier factors. <i>Familial Cancer</i> , 2020, 20, 189-194.	1.9	1
15	Clinical Characteristics and Prognosis of Gastric Cancer Patients with <i>BRCA1/2</i> Germline Mutations: Report of Ten Cases and a Literature Review. <i>OncoTargets and Therapy</i> , 2020, Volume 13, 11637-11644.	2.0	7
16	Histology results of systematic prostate biopsies by in-bore magnetic resonance imaging vs. transrectal ultrasound. <i>Canadian Urological Association Journal</i> , 2020, 15, E244-E247.	0.6	3
17	Radiation-Associated Secondary Malignancies in <i>BRCA</i> Mutation Carriers Treated for Breast Cancer. <i>International Journal of Radiation Oncology Biology Physics</i> , 2020, 107, 353-359.	0.8	17
18	Transcriptome-wide association study of breast cancer risk by estrogen receptor status. <i>Genetic Epidemiology</i> , 2020, 44, 442-468.	1.3	32

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19	Germline variant in REXO2 is a novel candidate gene in familial pheochromocytoma. <i>Genetical Research</i> , 2020, 102, e3.	0.9	10
20	Circulating cell-free DNA (cfDNA) levels in BRCA1 and BRCA2 mutation carriers: A preliminary study. <i>Cancer Biomarkers</i> , 2020, 28, 269-273.	1.7	4
21	Activating genomic alterations in the Gs alpha gene (<sc><i>GNAS</i></sc>) in 274â€™694 tumors. <i>Genes Chromosomes and Cancer</i> , 2020, 59, 503-516.	2.8	14
22	Diagnostic yield of multigene panel testing in an Israeli cohort: enrichment of low-penetrance variants. <i>Breast Cancer Research and Treatment</i> , 2020, 181, 445-453.	2.5	7
23	Effects of â€™real lifeâ€™ prostate MRI inter-observer variability on total needle samples and indication for biopsy. <i>Urologic Oncology: Seminars and Original Investigations</i> , 2020, 38, 793.e13-793.e18.	1.6	9
24	The spectrum of <i>BRCA1</i> and <i>BRCA2</i> pathogenic sequence variants in Middle Eastern, North African, and South European countries. <i>Human Mutation</i> , 2019, 40, e1-e23.	2.5	34
25	Mendelian randomisation study of height and body mass index as modifiers of ovarian cancer risk in 22,588 BRCA1 and BRCA2 mutation carriers. <i>British Journal of Cancer</i> , 2019, 121, 180-192.	6.4	19
26	Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer. <i>Nature Communications</i> , 2019, 10, 1741.	12.8	90
27	Multigene panel testing in unselected Israeli breast cancer cases: mutational spectrum and use of BRCA1/2 mutation prediction algorithms. <i>Breast Cancer Research and Treatment</i> , 2019, 176, 165-170.	2.5	7
28	Height and Body Mass Index as Modifiers of Breast Cancer Risk in <i>BRCA1</i>/<i>2</i> Mutation Carriers: A Mendelian Randomization Study. <i>Journal of the National Cancer Institute</i> , 2019, 111, 350-364.	6.3	30
29	Uterine cancer in Jewish Israeli <i>BRCA1/2</i> mutation carriers. <i>Cancer</i> , 2019, 125, 698-703.	4.1	28
30	The rate of the recurrent MSH6 mutations in Ashkenazi Jewish breast cancer patients. <i>Cancer Causes and Control</i> , 2019, 30, 97-101.	1.8	5
31	Mutational spectrum in a worldwide study of 29,700 families with<i>BRCA1</i> or<i>BRCA2</i> mutations. <i>Human Mutation</i> , 2018, 39, 593-620.	2.5	224
32	Abnormal Findings Detected by Multi-modality Breast Imaging and Biopsy Results in a High-risk Clinic. <i>Clinical Breast Cancer</i> , 2018, 18, e695-e698.	2.4	2
33	Phenotypic characteristics of colorectal cancer in BRCA1/2 mutation carriers. <i>European Journal of Human Genetics</i> , 2018, 26, 382-386.	2.8	11
34	Are VNTRs co-localizing with breast cancer-associated SNPs?. <i>Breast Cancer Research and Treatment</i> , 2018, 168, 277-281.	2.5	2
35	Mutational analysis of candidate genes in Israeli male breast cancer cases. <i>Breast Cancer Research and Treatment</i> , 2018, 170, 399-404.	2.5	5
36	The yield of targeted genotyping for the recurring mutations in BRCA1/2 in Israel. <i>Breast Cancer Research and Treatment</i> , 2018, 167, 697-702.	2.5	9

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37	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. <i>Nature Genetics</i> , 2017, 49, 680-691.	21.4	356
38	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	21.4	289
39	Inherited predisposition to breast and ovarian cancer in non-Jewish populations in Israel. <i>Breast Cancer Research and Treatment</i> , 2017, 166, 881-885.	2.5	12
40	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. <i>Breast Cancer Research and Treatment</i> , 2017, 161, 117-134.	2.5	18
41	De novo mutation in MEN1 is not associated with parental somatic mosaicism. <i>Endocrine-Related Cancer</i> , 2017, 24, L1-L3.	3.1	3
42	Primary Peritoneal Serous Carcinoma in Men: A Rare and Non-BRCA-associated Entity. <i>Anticancer Research</i> , 2017, 37, 3069-3072.	1.1	6
43	Colorectal and Endometrial Cancer Risk and Age at Diagnosis in BLMash Mutation Carriers. <i>Israel Medical Association Journal</i> , 2017, 19, 365-367.	0.1	8
44	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> , 2016, 11, e0158801.	2.5	10
45	Identification of independent association signals and putative functional variants for breast cancer risk through fine-scale mapping of the 12p11 locus. <i>Breast Cancer Research</i> , 2016, 18, 64.	5.0	31
46	An international survey of surveillance schemes for unaffected BRCA1 and BRCA2 mutation carriers. <i>Breast Cancer Research and Treatment</i> , 2016, 157, 319-327.	2.5	26
47	Inheritance of deleterious mutations at both BRCA1 and BRCA2 in an international sample of 32,295 women. <i>Breast Cancer Research</i> , 2016, 18, 112.	5.0	42
48	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. <i>Nature Communications</i> , 2016, 7, 11375.	12.8	93
49	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast-ovarian cancer susceptibility locus. <i>Nature Communications</i> , 2016, 7, 12675.	12.8	78
50	The rate of recurrent BRCA1, BRCA2, and TP53 mutations in the general population, and unselected ovarian cancer cases, in Belo Horizonte, Brazil. <i>Cancer Genetics</i> , 2016, 209, 283-284.	0.4	2
51	The risk for developing cancer in Israeli ATM, BLM, and FANCC heterozygous mutation carriers. <i>Cancer Genetics</i> , 2016, 209, 70-74.	0.4	29
52	Breast cancer risk variants at 6q25 display different phenotype associations and regulate ESR1, RMND1 and CCDC170. <i>Nature Genetics</i> , 2016, 48, 374-386.	21.4	125
53	The rate of recurrent BRCA1, BRCA2, and TP53 mutations in the general population, and unselected ovarian cancer cases, in Belo Horizonte, Brazil. <i>Cancer Genetics</i> , 2016, 209, 50-52.	0.4	18
54	No clinical utility of KRAS variant rs61764370 for ovarian or breast cancer. <i>Gynecologic Oncology</i> , 2016, 141, 386-401.	1.4	18

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55	Genome Sequencing of Multiple Primary Tumors Reveals a Novel <i>PALB2</i> Variant. <i>Journal of Clinical Oncology</i> , 2016, 34, e61-e67.	1.6	6
56	<i>GREM1</i> germline mutation screening in Ashkenazi Jewish patients with familial colorectal cancer. <i>Genetical Research</i> , 2015, 97, e11.	0.9	17
57	An original phylogenetic approach identified mitochondrial haplogroup T1a1 as inversely associated with breast cancer risk in <i>BRCA2</i> mutation carriers. <i>Breast Cancer Research</i> , 2015, 17, 61.	5.0	26
58	Assessing Associations between the AURKA-HMMR-TPX2-TUBG1 Functional Module and Breast Cancer Risk in <i>BRCA1/2</i> Mutation Carriers. <i>PLoS ONE</i> , 2015, 10, e0120020.	2.5	34
59	Identification of six new susceptibility loci for invasive epithelial ovarian cancer. <i>Nature Genetics</i> , 2015, 47, 164-171.	21.4	221
60	Cancer risks in Jewish male <i>BRCA1</i> and <i>BRCA2</i> mutation carriers. <i>Breast Cancer Research and Treatment</i> , 2015, 150, 631-635.	2.5	14
61	Association of Type and Location of <i>BRCA1</i> and <i>BRCA2</i> Mutations With Risk of Breast and Ovarian Cancer. <i>JAMA - Journal of the American Medical Association</i> , 2015, 313, 1347.	7.4	390
62	Candidate Genetic Modifiers for Breast and Ovarian Cancer Risk in <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 308-316.	2.5	22
63	DNA Glycosylases Involved in Base Excision Repair May Be Associated with Cancer Risk in <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers. <i>PLoS Genetics</i> , 2014, 10, e1004256.	3.5	47
64	<i>FMR1</i> CGG allele length in Israeli <i>BRCA1/BRCA2</i> mutation carriers and the general population display distinct distribution patterns. <i>Genetical Research</i> , 2014, 96, e11.	0.9	5
65	Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. <i>Nature Genetics</i> , 2013, 45, 371-384.	21.4	493
66	Haplotype analysis of the 185delAG <i>BRCA1</i> mutation in ethnically diverse populations. <i>European Journal of Human Genetics</i> , 2013, 21, 212-216.	2.8	44
67	A genetic variant of 5-hydroxytryptamine receptor 3C (<i>HTR3C</i>): A novel link to chemotherapy-induced side effects.. <i>Journal of Clinical Oncology</i> , 2013, 31, 9630-9630.	1.6	0
68	The founder Ashkenazi Jewish mutations in the <i>MSH2</i> and <i>MSH6</i> genes in Israeli patients with gastric and pancreatic cancer. <i>Familial Cancer</i> , 2012, 11, 243-247.	1.9	9
69	The KL-VS sequence variant of <i>Klotho</i> and cancer risk in <i>BRCA1</i> and <i>BRCA2</i> mutation carriers. <i>Breast Cancer Research and Treatment</i> , 2012, 132, 1119-1126.	2.5	8
70	Recurrent germline mutations in <i>BRCA1</i> and <i>BRCA2</i> genes in high risk families in Israel. <i>Breast Cancer Research and Treatment</i> , 2012, 133, 1153-1157.	2.5	18
71	Germline mutations in <i>BRCA1</i> and <i>BRCA2</i> genes in ethnically diverse high risk families in Israel. <i>Breast Cancer Research and Treatment</i> , 2011, 127, 489-495.	2.5	25
72	Haplotype of the C61G <i>BRCA1</i> Mutation in Polish and Jewish Individuals. <i>Genetic Testing and Molecular Biomarkers</i> , 2009, 13, 465-469.	0.7	13

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73	Germline CHEK2 mutations in Jewish Ashkenazi women at high risk for breast cancer. Israel Medical Association Journal, 2007, 9, 791-6.	0.1	11