siranoush manoukian

List of Publications by Year in Descending Order

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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.

13,562 50 115 143 h-index g-index citations papers 15,183 7.8 4.35 144 avg, IF L-index ext. citations ext. papers

| # | Paper | IF | Citations |
|-----|---|------|-----------|
| 143 | Malignant salivary gland tumours in families with breast cancer susceptibility. <i>Virchows Archiv Fur</i> Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2021 , 479, 221-226 | 5.1 | |
| 142 | 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 | 4.4 | 15 |
| 141 | Whole-exome sequencing and targeted gene sequencing provide insights into the role of PALB2 as a male breast cancer susceptibility gene. <i>Cancer</i> , 2017 , 123, 210-218 | 6.4 | 22 |
| 140 | Revertant mosaicism for family mutations is not observed in BRCA1/2 phenocopies. <i>PLoS ONE</i> , 2017 , 12, e0171663 | 3.7 | 6 |
| 139 | Haplotype analyses of the c.1027C>T and c.2167_2168delAT recurrent truncating mutations in the breast cancer-predisposing gene PALB2. <i>Breast Cancer Research and Treatment</i> , 2016 , 160, 121-129 | 4.4 | 7 |
| 138 | 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 | 8.3 | 25 |
| 137 | Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. <i>Nature Communications</i> , 2016 , 7, 11375 | 17.4 | 64 |
| 136 | Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast-ovarian cancer susceptibility locus. <i>Nature Communications</i> , 2016 , 7, 12675 | 17.4 | 53 |
| 135 | Fine scale mapping of the 17q22 breast cancer locus using dense SNPs, genotyped within the Collaborative Oncological Gene-Environment Study (COGs). <i>Scientific Reports</i> , 2016 , 6, 32512 | 4.9 | 16 |
| 134 | Cardio-Oncology: The Carney Complex Type I. <i>Journal of the American College of Cardiology</i> , 2016 , 68, 1921-1923 | 15.1 | 2 |
| 133 | Multiple primary melanomas (MPMs) and criteria for genetic assessment: MultiMEL, a multicenter study of the Italian Melanoma Intergroup. <i>Journal of the American Academy of Dermatology</i> , 2016 , 74, 325-32 | 4.5 | 23 |
| 132 | Treatment of infertility does not increase the risk of ovarian cancer among women with a BRCA1 or BRCA2 mutation. <i>Fertility and Sterility</i> , 2016 , 105, 781-785 | 4.8 | 28 |
| 131 | The incidence of leukaemia in women with BRCA1 and BRCA2 mutations: an International Prospective Cohort Study. <i>British Journal of Cancer</i> , 2016 , 114, 1160-4 | 8.7 | 19 |
| 130 | No evidence that protein truncating variants in BRIP1 are associated with breast cancer risk: implications for gene panel testing. <i>Journal of Medical Genetics</i> , 2016 , 53, 298-309 | 5.8 | 83 |
| 129 | Breast cancer risk variants at 6q25 display different phenotype associations and regulate ESR1, RMND1 and CCDC170. <i>Nature Genetics</i> , 2016 , 48, 374-86 | 36.3 | 93 |
| 128 | BRCA2 Polymorphic Stop Codon K3326X and the Risk of Breast, Prostate, and Ovarian Cancers. Journal of the National Cancer Institute, 2016 , 108, | 9.7 | 65 |
| 127 | Triple-Negative versus Non-Triple-Negative Breast Cancers in High-Risk Women: Phenotype Features and Survival from the HIBCRIT-1 MRI-Including Screening Study. <i>Clinical Cancer Research</i> , 2016 , 22, 895-904 | 12.9 | 25 |

(2015-2016)

| 126 | A targeted approach to genetic counseling in breast cancer patients: the experience of an Italian local project. <i>Tumori</i> , 2016 , 102, 45-50 | 1.7 | 3 |
|-----|--|------|-----|
| 125 | Fine-scale mapping of 8q24 locus identifies multiple independent risk variants for breast cancer. <i>International Journal of Cancer</i> , 2016 , 139, 1303-1317 | 7.5 | 26 |
| 124 | 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 | 8.3 | 25 |
| 123 | Mutation detection rates associated with specific selection criteria for BRCA1/2 testing in 1854 high-risk families: A monocentric Italian study. <i>European Journal of Internal Medicine</i> , 2016 , 32, 65-71 | 3.9 | 18 |
| 122 | Association of genetic susceptibility variants for type 2 diabetes with breast cancer risk in women of European ancestry. <i>Cancer Causes and Control</i> , 2016 , 27, 679-93 | 2.8 | 15 |
| 121 | Male breast cancer in BRCA1 and BRCA2 mutation carriers: pathology data from the Consortium of Investigators of Modifiers of BRCA1/2. <i>Breast Cancer Research</i> , 2016 , 18, 15 | 8.3 | 58 |
| 120 | Identification of six new susceptibility loci for invasive epithelial ovarian cancer. <i>Nature Genetics</i> , 2015 , 47, 164-71 | 36.3 | 177 |
| 119 | Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. <i>Nature Genetics</i> , 2015 , 47, 373-80 | 36.3 | 406 |
| 118 | Novel and known genetic variants for male breast cancer risk at 8q24.21, 9p21.3, 11q13.3 and 14q24.1: results from a multicenter study in Italy. <i>European Journal of Cancer</i> , 2015 , 51, 2289-95 | 7.5 | 20 |
| 117 | FANCM c.5791C>T nonsense mutation (rs144567652) induces exon skipping, affects DNA repair activity and is a familial breast cancer risk factor. <i>Human Molecular Genetics</i> , 2015 , 24, 5345-55 | 5.6 | 68 |
| 116 | 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> , 2015 , 313, 1347-61 | 27.4 | 286 |
| 115 | Height and Breast Cancer Risk: Evidence From Prospective Studies and Mendelian Randomization. Journal of the National Cancer Institute, 2015 , 107, | 9.7 | 74 |
| 114 | Fine-scale mapping of the 4q24 locus identifies two independent loci associated with breast cancer risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015 , 24, 1680-91 | 4 | 17 |
| 113 | Estimate of the penetrance of BRCA mutation and the COS software for the assessment of BRCA mutation probability. <i>Familial Cancer</i> , 2015 , 14, 117-28 | 3 | 12 |
| 112 | Identification and characterization of novel associations in the CASP8/ALS2CR12 region on chromosome 2 with breast cancer risk. <i>Human Molecular Genetics</i> , 2015 , 24, 285-98 | 5.6 | 35 |
| 111 | Candidate genetic modifiers for breast and ovarian cancer risk in BRCA1 and BRCA2 mutation carriers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015 , 24, 308-16 | 4 | 20 |
| 110 | Constitutional de novo deletion of the FBXW7 gene in a patient with focal segmental glomerulosclerosis and multiple primitive tumors. <i>Scientific Reports</i> , 2015 , 5, 15454 | 4.9 | 10 |
| 109 | An original phylogenetic approach identified mitochondrial haplogroup T1a1 as inversely associated with breast cancer risk in BRCA2 mutation carriers. <i>Breast Cancer Research</i> , 2015 , 17, 61 | 8.3 | 16 |

| 108 | Common germline polymorphisms associated with breast cancer-specific survival. <i>Breast Cancer Research</i> , 2015 , 17, 58 | 8.3 | 24 |
|-----|---|--------------------|-----|
| 107 | Assessing associations between the AURKA-HMMR-TPX2-TUBG1 functional module and breast cancer risk in BRCA1/2 mutation carriers. <i>PLoS ONE</i> , 2015 , 10, e0120020 | 3.7 | 26 |
| 106 | Prediction of breast cancer risk based on profiling with common genetic variants. <i>Journal of the National Cancer Institute</i> , 2015 , 107, | 9.7 | 324 |
| 105 | PALB2 sequencing in Italian familial breast cancer cases reveals a high-risk mutation recurrent in the province of Bergamo. <i>Genetics in Medicine</i> , 2014 , 16, 688-94 | 8.1 | 21 |
| 104 | A randomized controlled trial of diet and physical activity in BRCA mutation carriers. <i>Familial Cancer</i> , 2014 , 13, 181-7 | 3 | 12 |
| 103 | The psychological impact of breast and ovarian cancer preventive options in BRCA1 and BRCA2 mutation carriers. <i>Clinical Genetics</i> , 2014 , 85, 7-15 | 4 | 26 |
| 102 | FGF receptor genes and breast cancer susceptibility: results from the Breast Cancer Association Consortium. <i>British Journal of Cancer</i> , 2014 , 110, 1088-100 | 8.7 | 20 |
| 101 | Breast-cancer risk in families with mutations in PALB2. New England Journal of Medicine, 2014, 371, 497 | '- 5 96 | 576 |
| 100 | Duration of tamoxifen use and the risk of contralateral breast cancer in BRCA1 and BRCA2 mutation carriers. <i>Breast Cancer Research and Treatment</i> , 2014 , 146, 421-7 | 4.4 | 29 |
| 99 | A large de novo 9p21.3 deletion in a girl affected by astrocytoma and multiple melanoma. <i>BMC Medical Genetics</i> , 2014 , 15, 59 | 2.1 | 8 |
| 98 | Genetic variation in mitotic regulatory pathway genes is associated with breast tumor grade. <i>Human Molecular Genetics</i> , 2014 , 23, 6034-46 | 5.6 | 11 |
| 97 | DNA glycosylases involved in base excision repair may be associated with cancer risk in BRCA1 and BRCA2 mutation carriers. <i>PLoS Genetics</i> , 2014 , 10, e1004256 | 6 | 33 |
| 96 | Association of SULT1A1 ArgIIIHis polymorphism with male breast cancer risk: results from a multicenter study in Italy. <i>Breast Cancer Research and Treatment</i> , 2014 , 148, 623-8 | 4.4 | 4 |
| 95 | Refined histopathological predictors of BRCA1 and BRCA2 mutation status: a large-scale analysis of breast cancer characteristics from the BCAC, CIMBA, and ENIGMA consortia. <i>Breast Cancer Research</i> , 2014 , 16, 3419 | 8.3 | 82 |
| 94 | Associations of common breast cancer susceptibility alleles with risk of breast cancer subtypes in BRCA1 and BRCA2 mutation carriers. <i>Breast Cancer Research</i> , 2014 , 16, 3416 | 8.3 | 46 |
| 93 | Characterization of an Italian founder mutation in the RING-finger domain of BRCA1. <i>PLoS ONE</i> , 2014 , 9, e86924 | 3.7 | 18 |
| 92 | miR-342 regulates BRCA1 expression through modulation of ID4 in breast cancer. <i>PLoS ONE</i> , 2014 , 9, e87039 | 3.7 | 54 |
| 91 | First description of an acinic cell carcinoma of the breast in a BRCA1 mutation carrier: a case report. BMC Cancer, 2013 , 13, 46 | 4.8 | 28 |

(2012-2013)

| 90 | 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, 371-84, 384e1-2 | 36.3 | 422 |
|----|---|----------------|-----|
| 89 | Association of low-penetrance alleles with male breast cancer risk and clinicopathological characteristics: results from a multicenter study in Italy. <i>Breast Cancer Research and Treatment</i> , 2013 , 138, 861-8 | 4.4 | 26 |
| 88 | The impact of pregnancy on breast cancer survival in women who carry a BRCA1 or BRCA2 mutation. <i>Breast Cancer Research and Treatment</i> , 2013 , 142, 177-85 | 4.4 | 40 |
| 87 | X chromosome inactivation pattern in BRCA gene mutation carriers. <i>European Journal of Cancer</i> , 2013 , 49, 1136-41 | 7.5 | 8 |
| 86 | Large-scale genotyping identifies 41 new loci associated with breast cancer risk. <i>Nature Genetics</i> , 2013 , 45, 353-61, 361e1-2 | 36.3 | 813 |
| 85 | Identification of a BRCA2-specific modifier locus at 6p24 related to breast cancer risk. <i>PLoS Genetics</i> , 2013 , 9, e1003173 | 6 | 90 |
| 84 | Genome-wide association study in BRCA1 mutation carriers identifies novel loci associated with breast and ovarian cancer risk. <i>PLoS Genetics</i> , 2013 , 9, e1003212 | 6 | 209 |
| 83 | Whole exome sequencing suggests much of non-BRCA1/BRCA2 familial breast cancer is due to moderate and low penetrance susceptibility alleles. <i>PLoS ONE</i> , 2013 , 8, e55681 | 3.7 | 77 |
| 82 | Comparative in vitro and in silico analyses of variants in splicing regions of BRCA1 and BRCA2 genes and characterization of novel pathogenic mutations. <i>PLoS ONE</i> , 2013 , 8, e57173 | 3.7 | 46 |
| 81 | Breastfeeding and the risk of breast cancer in BRCA1 and BRCA2 mutation carriers. <i>Breast Cancer Research</i> , 2012 , 14, R42 | 8.3 | 68 |
| 80 | What is specific in hereditary breast cancer? High T2 signal intensity as a new semeiotic pattern?. <i>European Journal of Radiology</i> , 2012 , 81 Suppl 1, S165-70 | 4.7 | 1 |
| 79 | Genome-wide association analysis identifies three new breast cancer susceptibility loci. <i>Nature Genetics</i> , 2012 , 44, 312-8 | 36.3 | 237 |
| 78 | Rare variants in XRCC2 as breast cancer susceptibility alleles. <i>Journal of Medical Genetics</i> , 2012 , 49, 618- | - 250 8 | 37 |
| 77 | 9q31.2-rs865686 as a susceptibility locus for estrogen receptor-positive breast cancer: evidence from the Breast Cancer Association Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012 , 21, 1783-91 | 4 | 17 |
| 76 | Sequencing analysis of SLX4/FANCP gene in Italian familial breast cancer cases. <i>PLoS ONE</i> , 2012 , 7, e310 | 038 | 9 |
| 75 | 11q13 is a susceptibility locus for hormone receptor positive breast cancer. <i>Human Mutation</i> , 2012 , 33, 1123-32 | 4.7 | 33 |
| 74 | Identification of fifteen novel germline variants in the BRCA1 3RJTR reveals a variant in a breast cancer case that introduces a functional miR-103 target site. <i>Human Mutation</i> , 2012 , 33, 1665-75 | 4.7 | 42 |
| 73 | Clinical and pathologic characteristics of BRCA-positive and BRCA-negative male breast cancer patients: results from a collaborative multicenter study in Italy. <i>Breast Cancer Research and Treatment</i> , 2012 , 134, 411-8 | 4.4 | 58 |

| 72 | The SNP rs895819 in miR-27a is not associated with familial breast cancer risk in Italians. <i>Breast Cancer Research and Treatment</i> , 2012 , 133, 805-7 | 4.4 | 28 |
|----|---|----------------------------------|-----|
| 71 | 19p13.1 is a triple-negative-specific breast cancer susceptibility locus. <i>Cancer Research</i> , 2012 , 72, 1795- | -8 03 .1 | 93 |
| 7º | Common variants at the 19p13.1 and ZNF365 loci are associated with ER subtypes of breast cancer and ovarian cancer risk in BRCA1 and BRCA2 mutation carriers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012 , 21, 645-57 | 4 | 44 |
| 69 | 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> , 2012 , 21, 134-47 | 4 | 411 |
| 68 | Comparison of 6q25 breast cancer hits from Asian and European Genome Wide Association Studies in the Breast Cancer Association Consortium (BCAC). <i>PLoS ONE</i> , 2012 , 7, e42380 | 3.7 | 49 |
| 67 | Exploring the link between MORF4L1 and risk of breast cancer. <i>Breast Cancer Research</i> , 2011 , 13, R40 | 8.3 | 16 |
| 66 | Common breast cancer susceptibility alleles are associated with tumour subtypes in BRCA1 and BRCA2 mutation carriers: results from the Consortium of Investigators of Modifiers of BRCA1/2. Breast Cancer Research, 2011, 13, R110 | 8.3 | 62 |
| 65 | Multicenter surveillance of women at high genetic breast cancer risk using mammography, ultrasonography, and contrast-enhanced magnetic resonance imaging (the high breast cancer risk italian 1 study): final results. <i>Investigative Radiology</i> , 2011 , 46, 94-105 | 10.1 | 243 |
| 64 | Evaluation of the XRCC1 gene as a phenotypic modifier in BRCA1/2 mutation carriers. Results from the consortium of investigators of modifiers of BRCA1/BRCA2. <i>British Journal of Cancer</i> , 2011 , 104, 135 | 56 ⁸ 6 ⁷ 1 | 6 |
| 63 | Serum levels of IGF-I and BRCA penetrance: a case control study in breast cancer families. <i>Familial Cancer</i> , 2011 , 10, 521-8 | 3 | 25 |
| 62 | The CASP8 rs3834129 polymorphism and breast cancer risk in BRCA1 mutation carriers. <i>Breast Cancer Research and Treatment</i> , 2011 , 125, 855-60 | 4.4 | 10 |
| 61 | No evidence for an association between the earwax-associated polymorphism in ABCC11 and breast cancer risk in Caucasian women. <i>Breast Cancer Research and Treatment</i> , 2011 , 126, 235-9 | 4.4 | 16 |
| 60 | PALB2 germline mutations in familial breast cancer cases with personal and family history of pancreatic cancer. <i>Breast Cancer Research and Treatment</i> , 2011 , 126, 825-8 | 4.4 | 32 |
| 59 | Evidence for a link between TNFRSF11A and risk of breast cancer. <i>Breast Cancer Research and Treatment</i> , 2011 , 129, 947-54 | 4.4 | 11 |
| 58 | Two new CHEK2 germ-line variants detected in breast cancer/sarcoma families negative for BRCA1, BRCA2, and TP53 gene mutations. <i>Breast Cancer Research and Treatment</i> , 2011 , 130, 207-15 | 4.4 | 11 |
| 57 | Confirmation of 5p12 as a susceptibility locus for progesterone-receptor-positive, lower grade breast cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011 , 20, 2222-31 | 4 | 27 |
| 56 | Associations of breast cancer risk factors with tumor subtypes: a pooled analysis from the Breast Cancer Association Consortium studies. <i>Journal of the National Cancer Institute</i> , 2011 , 103, 250-63 | 9.7 | 513 |
| 55 | Common alleles at 6q25.1 and 1p11.2 are associated with breast cancer risk for BRCA1 and BRCA2 mutation carriers. <i>Human Molecular Genetics.</i> 2011 . 20. 3304-21 | 5.6 | 62 |

(2009-2011)

| 54 | Associations of common variants at 1p11.2 and 14q24.1 (RAD51L1) with breast cancer risk and heterogeneity by tumor subtype: findings from the Breast Cancer Association Consortium. <i>Human Molecular Genetics</i> , 2011 , 20, 4693-706 | 5.6 | 66 |
|----|--|------|-----|
| 53 | The rs12975333 variant in the miR-125a and breast cancer risk in Germany, Italy, Australia and Spain. <i>Journal of Medical Genetics</i> , 2011 , 48, 703-4 | 5.8 | 13 |
| 52 | Common variants of the BRCA1 wild-type allele modify the risk of breast cancer in BRCA1 mutation carriers. <i>Human Molecular Genetics</i> , 2011 , 20, 4732-47 | 5.6 | 21 |
| 51 | Genetic variation at 9p22.2 and ovarian cancer risk for BRCA1 and BRCA2 mutation carriers. <i>Journal of the National Cancer Institute</i> , 2011 , 103, 105-16 | 9.7 | 37 |
| 50 | Evaluation of variation in the phosphoinositide-3-kinase catalytic subunit alpha oncogene and breast cancer risk. <i>British Journal of Cancer</i> , 2011 , 105, 1934-9 | 8.7 | 4 |
| 49 | A locus on 19p13 modifies risk of breast cancer in BRCA1 mutation carriers and is associated with hormone receptor-negative breast cancer in the general population. <i>Nature Genetics</i> , 2010 , 42, 885-92 | 36.3 | 276 |
| 48 | Common breast cancer susceptibility alleles and the risk of breast cancer for BRCA1 and BRCA2 mutation carriers: implications for risk prediction. <i>Cancer Research</i> , 2010 , 70, 9742-54 | 10.1 | 147 |
| 47 | A BRCA1 promoter variant (rs11655505) and breast cancer risk. <i>Journal of Medical Genetics</i> , 2010 , 47, 268-70 | 5.8 | 5 |
| 46 | Common genetic variants and modification of penetrance of BRCA2-associated breast cancer. <i>PLoS Genetics</i> , 2010 , 6, e1001183 | 6 | 74 |
| 45 | Single-nucleotide polymorphisms inside microRNA target sites influence tumor susceptibility. <i>Cancer Research</i> , 2010 , 70, 2789-98 | 10.1 | 314 |
| 44 | HMGA1 protein expression in familial breast carcinoma patients. <i>European Journal of Cancer</i> , 2010 , 46, 332-9 | 7.5 | 16 |
| 43 | Four new cases of double heterozygosity for BRCA1 and BRCA2 gene mutations: clinical, pathological, and family characteristics. <i>Breast Cancer Research and Treatment</i> , 2010 , 124, 251-8 | 4.4 | 19 |
| 42 | Evaluation of SNPs in miR-146a, miR196a2 and miR-499 as low-penetrance alleles in German and Italian familial breast cancer cases. <i>Human Mutation</i> , 2010 , 31, E1052-7 | 4.7 | 138 |
| 41 | An unusual BRCA2 allele carrying two splice site mutations. <i>Annals of Oncology</i> , 2009 , 20, 1143-4 | 10.3 | 7 |
| 40 | SNPs in ultraconserved elements and familial breast cancer risk. <i>Carcinogenesis</i> , 2009 , 30, 544-5; author reply 546 | 4.6 | 22 |
| 39 | Analysis of a set of missense, frameshift, and in-frame deletion variants of BRCA1. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2009 , 660, 1-11 | 3.3 | 31 |
| 38 | Evidences for association of the CASP8 -652 6N del promoter polymorphism with age at diagnosis in familial breast cancer cases. <i>Breast Cancer Research and Treatment</i> , 2009 , 113, 607-8 | 4.4 | 18 |
| 37 | Cutaneous melanoma in childhood and adolescence shows frequent loss of INK4A and gain of KIT. Journal of Investigative Dermatology, 2009, 129, 1759-68 | 4.3 | 46 |

| 36 | The TP53 Arg72Pro and MDM2 309G>T polymorphisms are not associated with breast cancer risk in BRCA1 and BRCA2 mutation carriers. <i>British Journal of Cancer</i> , 2009 , 101, 1456-60 | 8.7 | 17 |
|----|--|-----|-----|
| 35 | Evaluation of a candidate breast cancer associated SNP in ERCC4 as a risk modifier in BRCA1 and BRCA2 mutation carriers. Results from the Consortium of Investigators of Modifiers of BRCA1/BRCA2 (CIMBA). <i>British Journal of Cancer</i> , 2009 , 101, 2048-54 | 8.7 | 13 |
| 34 | Clinical genetic testing for familial melanoma in Italy: a cooperative study. <i>Journal of the American Academy of Dermatology</i> , 2009 , 61, 775-82 | 4.5 | 39 |
| 33 | Cyclin D1 expression analysis in familial breast cancers may discriminate BRCAX from BRCA2-linked cases. <i>Modern Pathology</i> , 2008 , 21, 1262-70 | 9.8 | 15 |
| 32 | The BOADICEA model of genetic susceptibility to breast and ovarian cancers: updates and extensions. <i>British Journal of Cancer</i> , 2008 , 98, 1457-66 | 8.7 | 358 |
| 31 | Rapid progression of prostate cancer in men with a BRCA2 mutation. <i>British Journal of Cancer</i> , 2008 , 99, 371-4 | 8.7 | 112 |
| 30 | The p53 Arg72Pro and Ins16bp polymorphisms and their haplotypes are not associated with breast cancer risk in BRCA-mutation negative familial cases. <i>Cancer Detection and Prevention</i> , 2008 , 32, 140-3 | | 16 |
| 29 | BRCA1 p.Val1688del is a deleterious mutation that recurs in breast and ovarian cancer families from Northeast Italy. <i>Journal of Clinical Oncology</i> , 2008 , 26, 26-31 | 2.2 | 39 |
| 28 | Predictors of contralateral prophylactic mastectomy in women with a BRCA1 or BRCA2 mutation: the Hereditary Breast Cancer Clinical Study Group. <i>Journal of Clinical Oncology</i> , 2008 , 26, 1093-7 | 2.2 | 141 |
| 27 | Re: Molecular basis for estrogen receptor alpha deficiency in BRCA1-linked breast cancer. <i>Journal of the National Cancer Institute</i> , 2008 , 100, 752-3; author reply 753-4 | 9.7 | 2 |
| 26 | Hormone therapy and the risk of breast cancer in BRCA1 mutation carriers. <i>Journal of the National Cancer Institute</i> , 2008 , 100, 1361-7 | 9.7 | 152 |
| 25 | Genomic rearrangements of the CDKN2A locus are infrequent in Italian malignant melanoma families without evidence of CDKN2A/CDK4 point mutations. <i>Melanoma Research</i> , 2008 , 18, 431-7 | 3.3 | 7 |
| 24 | Infertility, treatment of infertility, and the risk of breast cancer among women with BRCA1 and BRCA2 mutations: a case-control study. <i>Cancer Causes and Control</i> , 2008 , 19, 1111-9 | 2.8 | 63 |
| 23 | Indications for breast magnetic resonance imaging. Consensus document "Attualit[in senologia", Florence 2007. <i>Radiologia Medica</i> , 2008 , 113, 1085-95 | 6.5 | 34 |
| 22 | International variation in rates of uptake of preventive options in BRCA1 and BRCA2 mutation carriers. <i>International Journal of Cancer</i> , 2008 , 122, 2017-22 | 7.5 | 268 |
| 21 | Common breast cancer-predisposition alleles are associated with breast cancer risk in BRCA1 and BRCA2 mutation carriers. <i>American Journal of Human Genetics</i> , 2008 , 82, 937-48 | 11 | 218 |
| 20 | New Trends of MRI in Breast Cancer Diagnosis 2008 , 127-144 | | |
| 19 | Methyl group metabolism gene polymorphisms as modifier of breast cancer risk in Italian BRCA1/2 carriers. <i>Breast Cancer Research and Treatment</i> , 2007 , 103, 29-36 | 4.4 | 22 |

| 18 | A CDKN2A mutation in familial melanoma that abrogates binding of p16INK4a to CDK4 but not CDK6. <i>Cancer Research</i> , 2007 , 67, 9134-41 | 10.1 | 38 | |
|----|--|------|------|---|
| 17 | Multicenter comparative multimodality surveillance of women at genetic-familial high risk for breast cancer (HIBCRIT study): interim results. <i>Radiology</i> , 2007 , 242, 698-715 | 20.5 | 287 | |
| 16 | Determination of cancer risk associated with germ line BRCA1 missense variants by functional analysis. <i>Cancer Research</i> , 2007 , 67, 1494-501 | 10.1 | 98 | |
| 15 | Age at menarche and menopause and breast cancer risk in the International BRCA1/2 Carrier Cohort Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007 , 16, 740-6 | 4 | 56 | |
| 14 | Germline mutations of TP53 and BRCA2 genes in breast cancer/sarcoma families. <i>European Journal of Cancer</i> , 2007 , 43, 601-6 | 7.5 | 38 | |
| 13 | Incidental carcinomas in prophylactic specimens in BRCA1 and BRCA2 germ-line mutation carriers, with emphasis on fallopian tube lesions: report of 6 cases and review of the literature. <i>American Journal of Surgical Pathology</i> , 2006 , 30, 1222-30 | 6.7 | 112 | |
| 12 | Mutational screening of the RB1 gene in Italian patients with retinoblastoma reveals 11 novel mutations. <i>Journal of Human Genetics</i> , 2006 , 51, 209-216 | 4.3 | 23 | |
| 11 | Bilateral preaxial polydactyly in a WAGR syndrome patient. <i>American Journal of Medical Genetics,</i> Part A, 2005 , 134, 426-9 | 2.5 | 4 | |
| 10 | Breast and ovarian cancer risks to carriers of the BRCA1 5382insC and 185delAG and BRCA2 6174delT mutations: a combined analysis of 22 population based studies. <i>Journal of Medical Genetics</i> , 2005 , 42, 602-3 | 5.8 | 107 | |
| 9 | A comparison of bilateral breast cancers in BRCA carriers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2005 , 14, 1534-8 | 4 | 42 | |
| 8 | Classification of BRCA1 missense variants of unknown clinical significance. <i>Journal of Medical Genetics</i> , 2005 , 42, 138-46 | 5.8 | 73 | |
| 7 | The CHEK2 c.1100delC mutation plays an irrelevant role in breast cancer predisposition in Italy. Human Mutation, 2004 , 24, 100-1 | 4.7 | 36 | |
| 6 | Atypical epithelial proliferation in fallopian tubes in prophylactic salpingo-oophorectomy specimens from BRCA1 and BRCA2 germline mutation carriers. <i>International Journal of Gynecological Pathology</i> , 2004 , 23, 35-40 | 3.2 | 104 | |
| 5 | Different expressivity of BRCA1 and BRCA2: analysis of 179 Italian pedigrees with identified mutation. <i>Breast Cancer Research and Treatment</i> , 2003 , 81, 71-9 | 4.4 | 22 | |
| 4 | Average risks of breast and ovarian cancer associated with BRCA1 or BRCA2 mutations detected in case Series unselected for family history: a combined analysis of 22 studies. <i>American Journal of Human Genetics</i> , 2003 , 72, 1117-30 | 11 | 2643 | |
| 3 | Characterization of a cytogenetic 17q11.2 deletion in an NF1 patient with a contiguous gene syndrome. <i>Human Genetics</i> , 1996 , 98, 646-50 | 6.3 | 23 | |
| 2 | Kaufman oculocerebrofacial syndrome in a girl of 15 years. <i>American Journal of Medical Genetics Part A</i> , 1995 , 58, 21-3 | | 6 | |
| | Prenatal testing in a fetus at risk for autosomal dominant polycystic kidney disease and autosomal | | | 1 |