

HEREDITARY CANCER REFERENCE LIST

Note: This reference list is updated bi-annually and this version is current as of August 1, 2018.

Gene	PMID	Reference Text
ALK	18724359	Mosse YP et al. Identification of ALK as a major familial neuroblastoma predisposition gene. <i>Nature</i> . 2008 Oct 16;455(7215):930-5. http://www.ncbi.nlm.nih.gov/pubmed/18724359
	18923503	Eng C. Cancer: A ringleader identified. <i>Nature</i> . 2008 Oct 16;455(7215):883-4. http://www.ncbi.nlm.nih.gov/pubmed/18923503
	18923523	Janoueix-Lerosey I et al. Somatic and germline activating mutations of the ALK kinase receptor in neuroblastoma. <i>Nature</i> . 2008 Oct 16 455(7215):967-70. http://www.ncbi.nlm.nih.gov/pubmed/18923523
	18923524	Chen Y et al. Oncogenic mutations of ALK kinase in neuroblastoma. <i>Nature</i> . 2008 Oct 16 455(7215):971-4. http://www.ncbi.nlm.nih.gov/pubmed/18923524
	18923525	George RE et al. Activating mutations in ALK provide a therapeutic target in neuroblastoma. <i>Nature</i> . 2008 Oct 16;455(7215):975-8. http://www.ncbi.nlm.nih.gov/pubmed/18923525
	20301782	Greengard EG, Park JR. ALK-Related Neuroblastic Tumor Susceptibility. 2010 Jan 5 [Updated 2015 Apr 9]. In: Pagon RA, Adam MP, Ardinger HH, et al., editors. <i>GeneReviews</i> ® [Internet]. Seattle (WA): University of Washington, Seattle; 1993-2016. http://www.ncbi.nlm.nih.gov/books/NBK24599/
	21972109	De Pontual et al. Germline gain-of-function mutations of ALK disrupt central nervous system development. <i>Hum Mutat</i> . 2011 Mar;32(3):272-6. http://www.ncbi.nlm.nih.gov/pubmed/21972109
	22071890	Bourdeaut F et al. ALK germline mutations in patients with neuroblastoma: a rare and weakly penetrant syndrome. <i>Eur J Hum Genet</i> . 2012 Mar;20(3):291-7. http://www.ncbi.nlm.nih.gov/pubmed/22071890
	ANKRD26	21211618
24030261		Noris P et al. ANKRD26-related thrombocytopenia and myeloid malignancies. <i>Blood</i> . 2013 Sep 12;122(11):1987-9. http://www.ncbi.nlm.nih.gov/pubmed/24030261
24628296		Marquez R et al. A new family with a germline ANKRD26 mutation and predisposition to myeloid malignancies. <i>Leuk Lymphoma</i> . 2014 Dec;55(12):2945-6. http://www.ncbi.nlm.nih.gov/pubmed/24628296
28109976		Averina M et al. A novel ANKRD26 gene variant causing inherited thrombocytopenia in a family of Finnish origin: Another brick in the wall? <i>Thromb Res</i> . 2017 Mar;151:41-43. http://www.ncbi.nlm.nih.gov/pubmed/28109976
28277066		Ferrari S et al. Spectrum of 5'UTR mutations in ANKRD26 gene in patients with inherited thrombocytopenia: c.-140C>G mutation is more frequent than expected. <i>Platelets</i> . 2017 Sep;28(6):621-624. http://www.ncbi.nlm.nih.gov/pubmed/28277066
28600339		Godley LA and Shimamura A. Genetic predisposition to hematologic malignancies: management and surveillance. <i>Blood</i> . 2017 Jul 27;130(4):424-432. http://www.ncbi.nlm.nih.gov/pubmed/28600339
APC		15300576
	17064931	Nieuwenhuis MH and Vasen HF. Correlations between mutation site in APC and phenotype of familial adenomatous polyposis (FAP): a review of the literature. <i>Crit Rev Oncol Hematol</i> . 2007 Feb;61(2):153-61. http://www.ncbi.nlm.nih.gov/pubmed/17064931
	17258512	Herraiz M et al. Prevalence of thyroid cancer in familial adenomatous polyposis syndrome and the role of screening ultrasound examinations. <i>Clinical Gastroenterology And Hepatology: The Official Clinical Practice Journal Of The American Gastroenterological Association</i> . 2007 Mar 5(3):367-73. http://www.ncbi.nlm.nih.gov/pubmed/17258512

Gene	PMID	Reference Text
APC	20420945	Jasperson KW et al. Hereditary and Familial Colon Cancer. Gastroenterology. 2010 Jun;138(6):2044-58 http://www.ncbi.nlm.nih.gov/pubmed/20420945
	20808249	Alkhouri N et al. Familial Adenomatous Polyposis in Children and Adolescents. Gastroenterology. 2010, 51 (6): 727-732. http://www.ncbi.nlm.nih.gov/pubmed/20808249
	22425061	Steinhagen E et al. The prevalence of thyroid cancer and benign thyroid disease in patients with familial adenomatous polyposis may be higher than previously recognized. Clinical Colorectal Cancer. 2012 Dec 11(4):304-8. http://www.ncbi.nlm.nih.gov/pubmed/22425061
	24549056	Ibrahim A et al. Attenuated familial adenomatous polyposis manifests as autosomal dominant late-onset colorectal cancer. EJHG 2014 Feb 19: 1-4. http://www.ncbi.nlm.nih.gov/pubmed/24549056
	25501924	Samadder NJ et al. Hereditary and Common Familial Colorectal Cancer: Evidence for Colorectal Screening. Digestive Diseases And Sciences. 2014 Dec 12. http://www.ncbi.nlm.nih.gov/pubmed/25501924
	25931827	Leoz ML et al. The genetic basis of familial adenomatous polyposis and its implications for clinical practice and risk management. Appl Clin Genet. 2015 Apr 16;8:95-107. http://www.ncbi.nlm.nih.gov/pubmed/25931827
	27623068	Uchino S et al. Age- and Gender-Specific Risk of Thyroid Cancer in Patients With Familial Adenomatous Polyposis. The Journal Of Clinical Endocrinology And Metabolism. 2016 Dec 101(12):4611-4617. http://www.ncbi.nlm.nih.gov/pubmed/27623068
	28185118	Mankaney G et al. Gastric cancer in FAP: a concerning rise in incidence. Familial Cancer. 2017 Jul 16(3):371-376. http://www.ncbi.nlm.nih.gov/pubmed/28185118
	9038672	Giardiello FM et al. Hepatoblastoma and APC gene mutation in familial adenomatous polyposis. Gut. 1996 Dec;39(6):867-9. http://www.ncbi.nlm.nih.gov/pubmed/9038672
	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Colorectal. http://www.nccn.org/clinical.asp
ATM	15928302	Thompson D et al. Cancer risks and mortality in heterozygous ATM mutation carriers. J Natl Cancer Inst. 2005 Jun 1; 97(11):813-22. http://www.ncbi.nlm.nih.gov/pubmed/15928302
	16832357	Renwick A et al. ATM mutations that cause ataxia-telangiectasia are breast cancer susceptibility alleles. Nat Genet. 2006 Aug; 38(8):873-5. http://www.ncbi.nlm.nih.gov/pubmed/16832357
	19781682	Tavtigian SV et al. Rare, evolutionary unlikely missense substitutions in ATM confer increased risk of breast cancer. Am J Hum Genet. 2009 Oct;85(4):427-46. http://www.ncbi.nlm.nih.gov/pubmed/19781682
	22585167	Roberts NJ et al. ATM Mutations in Patients with Hereditary Pancreatic Cancer. Cancer Discov. 2012 Jan;2(1):41-6. http://www.ncbi.nlm.nih.gov/pubmed/22585167
	24556621	Leongamornlert D et al. Frequent germline deleterious mutations in DNA repair genes in familial prostate cancer cases are associated with advanced disease. Br J Cancer. 2014 Mar 18;110(6):1663-72. http://www.ncbi.nlm.nih.gov/pubmed/24556621
	26014596	Easton DF et al. Gene-Panel Sequencing and the Prediction of Breast-Cancer Risk. N Engl J Med. 2015 Jun 4;372(23):2243-57. http://www.ncbi.nlm.nih.gov/pubmed/26014596
	26098866	Helgason H et al. Loss-of-function variants in ATM confer risk of gastric cancer. Nat Genet. 2015 Aug;47(8):906-10. http://www.ncbi.nlm.nih.gov/pubmed/26098866
	26182300	Hansford S et al. Hereditary diffuse gastric cancer syndrome: CDH1 mutations and beyond. JAMA Oncol. 2015 Apr;1(1):23-32. http://www.ncbi.nlm.nih.gov/pubmed/26182300
	26506520	Huang DS et al. Prevalence of deleterious ATM germline mutations in gastric cancer patients. Oncotarget. 2015 Dec 1;6(38):40953-8. http://www.ncbi.nlm.nih.gov/pubmed/26506520
	26658419	Roberts NJ et al. Whole Genome Sequencing Defines the Genetic Heterogeneity of Familial Pancreatic Cancer. Cancer Discovery. 2016 Feb 6(2):166-75. http://www.ncbi.nlm.nih.gov/pubmed/26658419

Gene	PMID	Reference Text	
ATM	27112364	Marabelli M et al. Penetrance of ATM gene mutations in breast cancer: a meta-analysis of different measures of risk. <i>Genet Epidemiol.</i> 2016 Jul;40(5):425-31. http://www.ncbi.nlm.nih.gov/pubmed/27112364	
	27433846	Pritchard CC et al. Inherited DNA-Repair Gene Mutations in Men with Metastatic Prostate Cancer. <i>N Engl J Med.</i> 2016 Aug 4;375(5):443-53. http://www.ncbi.nlm.nih.gov/pubmed/27433846	
	27989354	Na R et al. Germline mutations in ATM and BRCA1/2 distinguish risk for lethal and indolent prostate cancer and are associated with early age at death. <i>Eur Urol.</i> 2016 Dec 9. S0302-2838(16)30885-5. http://www.ncbi.nlm.nih.gov/pubmed/27989354	
	28418444	Couch FJ et al. Associations Between Cancer Predisposition Testing Panel Genes and Breast Cancer. <i>JAMA Oncol.</i> 2017 Apr 13. http://www.ncbi.nlm.nih.gov/pubmed/28418444	
	28649662	Slavin TP et al. The contribution of pathogenic variants in breast cancer susceptibility genes to familial breast cancer risk. <i>NPJ Breast Cancer.</i> 2017 Jun 9;3:22. http://www.ncbi.nlm.nih.gov/pubmed/28649662	
	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Breast and Ovarian. http://www.nccn.org/clinical.asp	
	AXIN2	15042511	Lammi L et al. Mutations in AXIN2 Cause Familial Tooth Agenesis and Predispose to Colorectal Cancer. <i>Am J Hum Genet.</i> 2004 May;74(5):1043-50. http://www.ncbi.nlm.nih.gov/pubmed/15042511
16110024		Renkonen ET et al. Adenomatous polyposis families that screen APC mutation-negative by conventional methods are genetically heterogeneous. <i>J Clin Oncol.</i> 2005 Aug 20;23(24):5651-9. http://www.ncbi.nlm.nih.gov/pubmed/16110024	
18708403		Wang X et al. Association of genetic variation in genes implicated in the beta-catenin destruction complex with risk of breast cancer. <i>Cancer Epidemiol Biomarkers Prev.</i> 2008 Aug;17(8):2101-8. http://www.ncbi.nlm.nih.gov/pubmed/18708403	
19119171		Menezes R et al. AXIS inhibition protein 2, orofacial clefts and a family history of cancer. <i>J Am Dent Assoc.</i> 2009 January ; 140(1):80-84. http://www.ncbi.nlm.nih.gov/pubmed/19119171	
21416598		Marvin ML et al. AXIN2-associated autosomal dominant ectodermal dysplasia and neoplastic syndrome. <i>Am J Med Genet A.</i> 2011 Apr;155A(4):898-902. http://www.ncbi.nlm.nih.gov/pubmed/21416598	
21476993		Mongin C et al. Unexplained polyposis: a challenge for geneticists, pathologists and gastroenterologists. <i>Clin Genet.</i> 2012 Jan;81(1):38-46. http://www.ncbi.nlm.nih.gov/pubmed/21476993	
23516639		Alanazi MS et al. Association of single nucleotide polymorphisms in Wnt signaling pathway genes with breast cancer in Saudi patients. <i>PLoS One.</i> 2013;8(3):e59555. http://www.ncbi.nlm.nih.gov/pubmed/23516639	
23838596		Rivera B et al. A novel AXIN2 germline variant associated with attenuated FAP without signs of oligodontia or ectodermal dysplasia. <i>Eur J Hum Genet.</i> 2014 Mar;22(3):423-6. http://www.ncbi.nlm.nih.gov/pubmed/23838596	
27090353		Yeu H et al. Functional analysis of a novel missense mutation in AXIN2 associated with non-syndromic tooth agenesis. <i>Eur J Oral Sci.</i> 2016 Jun;124(3):228-33. http://www.ncbi.nlm.nih.gov/pubmed/27090353	
27696107		Rohlin A et al. Expanding the genotype-phenotype spectrum in hereditary colorectal cancer by gene panel testing. <i>Fam Cancer.</i> 2017 Apr;16(2):195-203. http://www.ncbi.nlm.nih.gov/pubmed/27696107	
NCCN		NCCN Guidelines. Genetic/Familial High-Risk Assessment: Colorectal. (http://www.nccn.org/clinical.asp) http://www.nccn.org/clinical.asp	
BAP1		21874000	Testa JR et al. Germline BAP1 mutations predispose to malignant mesothelioma. <i>Nat Genet.</i> 2011 Aug 28;43(10):1022-5. http://www.ncbi.nlm.nih.gov/pubmed/21874000
		21874003	Wiesner T et al. Germline mutations in BAP1 predispose to melanocytic tumors. <i>Nat Genet.</i> 2011 Aug 28;43(10):1018-21. http://www.ncbi.nlm.nih.gov/pubmed/21874003

Gene	PMID	Reference Text
BAP1	22935333	Carbone M et al. BAP1 cancer syndrome: malignant mesothelioma, uveal and cutaneous melanoma, and MBAITs. <i>J Transl Med.</i> 2012 Aug 30;10:179. http://www.ncbi.nlm.nih.gov/pubmed/22935333
	23171164	Aoude LG et al. Prevalence of germline BAP1 mutation in a population-based sample of uveal melanoma cases. <i>Pigment Cell Melanoma Res.</i> 2013 Mar;26(2):278-9. http://www.ncbi.nlm.nih.gov/pubmed/23171164
	23684012	Popova T et al. Germline BAP1 mutations predispose to renal cell carcinomas. <i>Am J Hum Genet.</i> 2013 Jun 6;92(6):974-80. http://www.ncbi.nlm.nih.gov/pubmed/23684012
	24243779	Pilarski R et al. Expanding the clinical phenotype of hereditary BAP1 cancer predisposition syndrome, reporting three new cases. <i>Genes Chromosomes Cancer.</i> 2014 Feb;53(2):177-82. http://www.ncbi.nlm.nih.gov/pubmed/24243779
	26096145	Rai K et al. Comprehensive review of BAP1 tumor predisposition syndrome with report of two new cases. <i>Clin Genet.</i> 2015 Jun 22. http://www.ncbi.nlm.nih.gov/pubmed/26096145
	27748099	Pilarski R, Rai K, Cebulla C, et al. BAP1 Tumor Predisposition Syndrome. 2016 Oct 13. In: Pagon RA, Adam MP, Ardinger HH, et al., editors. <i>GeneReviews</i> ® [Internet]. Seattle (WA): University of Washington, Seattle; 1993-2017. https://www.ncbi.nlm.nih.gov/books/NBK390611
	28283772	Leachman SA et al. Identification, genetic testing, and management of hereditary melanoma. <i>Cancer Metastasis Rev.</i> 2017 Mar;36(1):77-90. http://www.ncbi.nlm.nih.gov/pubmed/28283772
	BARD1	15342711
16768547		Stacey SN et al. The BARD1 Cys557Ser variant and breast cancer risk in Iceland. <i>PLoS Med.</i> 2006 Jul;3(7):e217. http://www.ncbi.nlm.nih.gov/pubmed/16768547
17972171		Gorringe KL et al. BARD1 variants are not associated with breast cancer risk in Australian familial breast cancer. <i>Breast Cancer Res Treat.</i> 2008 Oct;111(3):505-9. http://www.ncbi.nlm.nih.gov/pubmed/17972171
18481171		Johnatty SE et al. The BARD1 Cys557Ser polymorphism and breast cancer risk: an Australian case-control and family analysis. <i>Breast Cancer Res Treat.</i> 2009 May;115(1):145-50. http://www.ncbi.nlm.nih.gov/pubmed/18481171
20077502		DeBrakeleer S et al. Cancer predisposing missense and protein truncating BARD1 mutations in non-BRCA1 or BRCA2 breast cancer families. <i>Hum Mutat.</i> 2010 Mar;31(3):E1175-85. http://www.ncbi.nlm.nih.gov/pubmed/20077502
21344236		Ratajska M et al. Cancer predisposing BARD1 mutations in breast-ovarian cancer families. <i>Breast Cancer Res Treat.</i> 2012 Jan;131(1):89-97. http://www.ncbi.nlm.nih.gov/pubmed/21344236
21809034		Ding DP et al. Lack of association between BARD1 Cys557Ser variant and breast cancer risk: a meta-analysis of 11,870 cases and 7,687 controls. <i>J Cancer Res Clin Oncol.</i> 2011 Oct;137(10):1463-8 http://www.ncbi.nlm.nih.gov/pubmed/21809034
23334666		Pugh TJ et al. The genetic landscape of high-risk neuroblastoma. <i>Nat Genet.</i> 2013 Mar;45(3):279-84. http://www.ncbi.nlm.nih.gov/pubmed/23334666
24240112		Pennington KP et al. Germline and somatic mutations in homologous recombination genes predict platinum response and survival in ovarian, fallopian tube, and peritoneal carcinomas. <i>Clin Cancer Res.</i> 2014 Feb 1;20(3):764-75. http://www.ncbi.nlm.nih.gov/pubmed/24240112
25452441		Couch FJ et al. Inherited mutations in 17 breast cancer susceptibility genes among a large triple-negative breast cancer cohort unselected for family history of breast cancer. <i>J Clin Oncol.</i> 2015 Feb 1;33(4):304-11. http://www.ncbi.nlm.nih.gov/pubmed/25452441
26010302		De Brakeleer S et al. Frequent incidence of BARD1 truncating mutations in germline DNA from triple negative breast cancer patients. <i>Clinical Genetics.</i> 2015 May 22. http://www.ncbi.nlm.nih.gov/pubmed/26010302

Gene	PMID	Reference Text
<i>BARD1</i>	26315354	Ramus SJ et al. Germline Mutations in the BRIP1, BARD1, PALB2, and NBN Genes in Women With Ovarian Cancer. <i>J Natl Cancer Inst.</i> 2015 Aug 27;107(11). http://www.ncbi.nlm.nih.gov/pubmed/26315354
	26483394	Hu C et al. Prevalence of pathogenic mutations in cancer predisposition genes among pancreatic cancer patients. <i>Cancer Epidemiol Biomarkers Prev.</i> 2016 Jan;25(1):207-11. http://www.ncbi.nlm.nih.gov/pubmed/26483394
	26720728	Norquist BM et al. Inherited Mutations in Women With Ovarian Carcinoma. <i>Jama Oncology.</i> 2015 Dec 30:1-9. http://www.ncbi.nlm.nih.gov/pubmed/26720728
	27443514	Ring KL et al. Germline multi-gene hereditary cancer panel testing in an unselected endometrial cancer cohort. <i>Mod Pathol.</i> 2016 Nov;29(11):1381-1389. http://www.ncbi.nlm.nih.gov/pubmed/27443514
<i>BLM</i>	12242432	Gruber SB et al. BLM heterozygosity and the risk of colorectal cancer. <i>Science.</i> 2002 Sep 20;297(5589):2013. http://www.ncbi.nlm.nih.gov/pubmed/12242432
	12702560	Cleary SP et al. Heterozygosity for the BLMAsh Mutation and Cancer Risk. <i>Cancer Res.</i> 2003 Apr 15;63(8):1769-71. http://www.ncbi.nlm.nih.gov/pubmed/12702560
	21815139	Sokolenko AP et al. High prevalence and breast cancer predisposing role of the BLM c.1642C>T (Q548X) mutation in Russia. <i>Int J Cancer.</i> 2012 Jun 15;130(12):2867-73. http://www.ncbi.nlm.nih.gov/pubmed/21815139
	9837821	Ellis NA et al. The Ashkenazic Jewish Bloom syndrome mutation blmAsh is present in non-Jewish Americans of Spanish ancestry. <i>Am J Hum Genet.</i> 1998 Dec;63(6):1685-93. http://www.ncbi.nlm.nih.gov/pubmed/9837821
<i>BMPR1A</i>	16436638	Pyatt RE et al. Mutation screening in juvenile polyposis syndrome. <i>The Journal Of Molecular Diagnostics : Jmd.</i> 2006 8(1):84-8. http://www.ncbi.nlm.nih.gov/pubmed/16436638
	16525031	Cao X et al. Mapping of hereditary mixed polyposis syndrome (HMPS) to chromosome 10q23 by genomewide high-density single nucleotide polymorphism (SNP) scan and identification of BMPR1A loss of function. <i>J Med Genet.</i> 2006 Mar;43(3):e13. http://www.ncbi.nlm.nih.gov/pubmed/16525031
	17303595	Brosens LAA et al. Risk of colorectal cancer in juvenile polyposis. <i>Gut.</i> 2007;56:965-7. http://www.ncbi.nlm.nih.gov/pubmed/17303595
	19438883	O'Riordan JM et al. Hereditary mixed polyposis syndrome due to a BMPR1A mutation. <i>Colorectal Disease : The Official Journal Of The Association Of Coloproctology Of Great Britain And Ireland.</i> 2010 12(6):570-3. http://www.ncbi.nlm.nih.gov/pubmed/19438883
	19773747	Cheah PY. Germline bone morphogenesis protein receptor 1A mutation causes colorectal tumorigenesis in hereditary mixed polyposis syndrome. <i>Am J Gastroenterol.</i> 2009 Dec;104(12):3027-33. http://www.ncbi.nlm.nih.gov/pubmed/19773747
	21640116	Nieminen TT et al. BMPR1A mutations in hereditary nonpolyposis colorectal cancer without mismatch repair deficiency. <i>Gastroenterology.</i> 2011 Jul 141(1):e23-6. http://www.ncbi.nlm.nih.gov/pubmed/21640116
	22965402	Latchford AR et al. Juvenile polyposis syndrome: a study of genotype, phenotype, and long-term outcome. <i>Diseases Of The Colon And Rectum.</i> 2012 Oct 55(10):1038-43. http://www.ncbi.nlm.nih.gov/pubmed/22965402
	23057600	Fernandez-Rozadilla C et al. BMPR1A mutations in early-onset colorectal cancer with mismatch repair proficiency. <i>Clinical Genetics.</i> 2013 Jul 84(1):94-6. http://www.ncbi.nlm.nih.gov/pubmed/23057600
	25848489	Campos FG et al. Colorectal cancer risk in hamartomatous polyposis syndromes. <i>World Journal Of Gastrointestinal Surgery.</i> 2015 Mar 27 7(3):25-32. http://www.ncbi.nlm.nih.gov/pubmed/25848489
	26363537	Jasperson K and Burt RW. The Genetics of Colorectal Cancer. <i>Surgical Oncology Clinics Of North America.</i> 2015 Oct 24(4):683-703. http://www.ncbi.nlm.nih.gov/pubmed/26363537
27326320	Shenoy S. Genetic risks and familial associations of small bowel carcinoma. <i>World Journal Of Gastrointestinal Oncology.</i> 2016 Jun 15 8(6):509-19. http://www.ncbi.nlm.nih.gov/pubmed/27326320	

Gene	PMID	Reference Text
BMPR1A	27696107	Rohlin A et al. Expanding the genotype-phenotype spectrum in hereditary colorectal cancer by gene panel testing. <i>Fam Cancer</i> . 2017 Apr;16(2):195-203. http://www.ncbi.nlm.nih.gov/pubmed/27696107
	9643289	Desai DC et al. A survey of phenotypic features in juvenile polyposis. <i>Journal Of Medical Genetics</i> . 1998 Jun 35(6):476-81. http://www.ncbi.nlm.nih.gov/pubmed/9643289
	9869523	Howe JR et al. The risk of gastrointestinal carcinoma in familial juvenile polyposis. <i>Ann Surg Oncol</i> . 1998 Dec;5(8):751-6. http://www.ncbi.nlm.nih.gov/pubmed/9869523
	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Colorectal. http://www.nccn.org/clinical.asp
BRCA1	12237281	Thompson D, Easton DF, and the Breast Cancer Linkage Consortium. Cancer Incidence in BRCA1 mutation carriers. <i>J Natl Cancer Inst</i> . 2002 Sep;94(18):1358-65. http://www.ncbi.nlm.nih.gov/pubmed/12237281
	12237282	Brose MS et al. Cancer risk estimates for BRCA1 mutation carriers identified in a risk evaluation program. <i>J Natl Cancer Inst</i> . 2002 Sep 18;94(18):1365-72. http://www.ncbi.nlm.nih.gov/pubmed/12237282
	12677558	Antoniou A et al. 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>Am J Hum Genet</i> . 2003 May;72(5):1117-30. http://www.ncbi.nlm.nih.gov/pubmed/12677558
	14576434	King MC et al. Breast and ovarian cancer risks due to inherited mutations in BRCA1 and BRCA2. <i>Science</i> . 2003 Oct;302(5645):643-6. http://www.ncbi.nlm.nih.gov/pubmed/14576434
	14615451	Levine DA et al. Fallopian Tube and Primary Peritoneal Carcinomas Associated With BRCA Mutations. <i>J Clin Oncol</i> . 2003 Nov 15;21(22):4222-7. http://www.ncbi.nlm.nih.gov/pubmed/14615451
	14966099	Liede A et al. Cancer Risks for Male Carriers of Germline Mutations in BRCA1 or BRCA2: A Review of the Literature. <i>J Clin Oncol</i> . 2004 Feb 15;22(4):735-42 http://www.ncbi.nlm.nih.gov/pubmed/14966099
	17416853	Chen S and Parmigiani G. Meta-analysis of BRCA1 and BRCA2 penetrance. <i>J Clin Oncol</i> . 2007 Apr;25(11):1329-33. http://www.ncbi.nlm.nih.gov/pubmed/17416853
	18042939	Tai YC et al. Breast cancer risk among male BRCA1 and BRCA2 mutation carriers. <i>J Natl Cancer Inst</i> . 2007 Dec;99(23):1811-4. http://www.ncbi.nlm.nih.gov/pubmed/18042939
	19858402	Graeser MK et al. Contralateral Breast Cancer Risk in BRCA1 and BRCA2 Mutation Carriers. <i>J Clin Oncol</i> . 2009 Dec 10;27(35): 5887-92. http://www.ncbi.nlm.nih.gov/pubmed/19858402
	22010008	Lee E et al. Characteristics of triple-negative breast cancer in patients with a BRCA1 mutation: results from a population-based study of young women. <i>J Clin Oncol</i> . 2011 Nov 20;29(33):4373-80. http://www.ncbi.nlm.nih.gov/pubmed/22010008
	22516946	Leongamornlert D et al. Germline BRCA1 mutations increase prostate cancer risk. <i>Br J Cancer</i> . 2012 May 8;106(10):1697-701. http://www.ncbi.nlm.nih.gov/pubmed/22516946
	22811390	Pennington KP et al. BRCA1, TP53, and CHEK2 germline mutations in uterine serous carcinoma. <i>Cancer</i> . 2013 Jan;119(2):332-8. http://www.ncbi.nlm.nih.gov/pubmed/22811390
	23099806	Iqbal J et al. The incidence of pancreatic cancer in BRCA1 and BRCA2 mutation carriers. <i>British Journal Of Cancer</i> . 2012 Dec 04 107(12):2005-9. http://www.ncbi.nlm.nih.gov/pubmed/23099806
	23628597	Mavaddat N et al. Cancer risks for BRCA1 and BRCA2 mutation carriers: results from prospective analysis of EMBRACE. <i>Journal Of The National Cancer Institute</i> . 2013 105(11):812-22. http://www.ncbi.nlm.nih.gov/pubmed/23628597
	28049106	de Jonge MM et al. Linking uterine serous carcinoma to BRCA1/2-associated cancer syndrome: A meta-analysis and case report. <i>European Journal Of Cancer (Oxford, England : 1990)</i> . 2017 Feb 72:215-225. http://www.ncbi.nlm.nih.gov/pubmed/28049106

Gene	PMID	Reference Text
BRCA1	28632866	Kuchenbaecker KB et al. Risks of Breast, Ovarian, and Contralateral Breast Cancer for BRCA1 and BRCA2 Mutation Carriers. JAMA. 2017 Jun 20;317(23):2402-2416. http://www.ncbi.nlm.nih.gov/pubmed/28632866
	7907678	Ford D et al. Risks of cancer in BRCA1-mutation carriers. Breast Cancer Linkage Consortium. Lancet. 1994 343(8899):692-5. http://www.ncbi.nlm.nih.gov/pubmed/7907678
	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Breast and Ovarian. http://www.nccn.org/clinical.asp
BRCA2	10433620	The Breast Cancer Linkage Consortium. Cancer risks in BRCA2 mutation carriers. J Natl Cancer Inst. 1999 Aug;91(15):1310-6. http://www.ncbi.nlm.nih.gov/pubmed/10433620
	11170890	Thompson D, Easton D, Breast Cancer Linkage Consortium. Variation in cancer risks, by mutation position, in BRCA2 mutation carriers. Am J Hum Genet. 2001 Feb;68(2):410-9. http://www.ncbi.nlm.nih.gov/pubmed/11170890
	12556369	Hearle N et al. Contribution of germline mutations in BRCA2, P16(INK4A), P14(ARF) and P15 to uveal melanoma. Invest Ophthalmol Vis Sci. 2003 Feb;44(2):458-62. http://www.ncbi.nlm.nih.gov/pubmed/12556369
	12677558	Antoniou A et al. 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. Am J Hum Genet. 2003 May;72(5):1117-30. http://www.ncbi.nlm.nih.gov/pubmed/12677558
	14615451	Levine DA et al. Fallopian Tube and Primary Peritoneal Carcinomas Associated With BRCA Mutations. J Clin Oncol. 2003 Nov 15;21(22):4222-7. http://www.ncbi.nlm.nih.gov/pubmed/14615451
	14966099	Liede A et al. Cancer Risks for Male Carriers of Germline Mutations in BRCA1 or BRCA2: A Review of the Literature. J Clin Oncol. 2004 Feb 15;22(4):735-42 http://www.ncbi.nlm.nih.gov/pubmed/14966099
	16650962	Biron-Shental T et al. High incidence of BRCA1-2 germline mutations, previous breast cancer and familial cancer history in Jewish patients with uterine serous papillary carcinoma. Eur J Surg Oncol. 2006 Dec;32(10):1097-100. http://www.ncbi.nlm.nih.gov/pubmed/16650962
	18042939	Tai YC et al. Breast cancer risk among male BRCA1 and BRCA2 mutation carriers. J Natl Cancer Inst. 2007 Dec;99(23):1811-4. http://www.ncbi.nlm.nih.gov/pubmed/18042939
	19858402	Graeser MK et al. Contralateral Breast Cancer Risk in BRCA1 and BRCA2 Mutation Carriers. J Clin Oncol. 2009 Dec 10;27(35): 5887-92. http://www.ncbi.nlm.nih.gov/pubmed/19858402
	20301575	Mehta PA, Tolar J. Fanconi Anemia. 2002 Feb 14 [Updated 2016 Sep 22]. In: Pagon RA, Adam MP, Ardinger HH, et al., editors. GeneReviews® [Internet]. Seattle (WA): University of Washington, Seattle; 1993-2017. http://www.ncbi.nlm.nih.gov/pubmed/20301575
	20587410	Evans DG et al. Risk of breast cancer in male BRCA2 carriers. J Med Genet. 2010 Oct;47(10):710-1. http://www.ncbi.nlm.nih.gov/pubmed/20587410
	21952622	Kote-Jarai Z et al. BRCA2 is a moderate penetrance gene contributing to young-onset prostate cancer: implications for genetic testing in prostate cancer patients. Br J Cancer. 2011 Oct 11;105(8):1230-4. http://www.ncbi.nlm.nih.gov/pubmed/21952622
	23099806	Iqbal J et al. The incidence of pancreatic cancer in BRCA1 and BRCA2 mutation carriers. British Journal Of Cancer. 2012 Dec 04 107(12):2005-9. http://www.ncbi.nlm.nih.gov/pubmed/23099806
	28632866	Kuchenbaecker KB et al. Risks of Breast, Ovarian, and Contralateral Breast Cancer for BRCA1 and BRCA2 Mutation Carriers. JAMA. 2017 Jun 20;317(23):2402-2416. http://www.ncbi.nlm.nih.gov/pubmed/28632866
	9140390	Ozcelik H et al. Germline BRCA2 6174delT mutations in Ashkenazi Jewish pancreatic cancer patients. Nat Genet. 1997 May;16(1):17-8. http://www.ncbi.nlm.nih.gov/pubmed/9140390
	9497246	Ford D et al. Genetic heterogeneity and penetrance analysis of the BRCA1 and BRCA2 genes in breast cancer families. The Breast Cancer Linkage Consortium. Am J Hum Genet. 1998 Mar;62(3):676-89. http://www.ncbi.nlm.nih.gov/pubmed/9497246

Gene	PMID	Reference Text	
BRCA2	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Breast and Ovarian. http://www.nccn.org/clinical.asp	
BRIP1	17033622	Seal S et al. Truncating mutations in the Fanconi anemia J gene BRIP1 are low-penetrance breast cancer susceptibility alleles. Nat Genet. 2006 Nov;38(11):1239-41. http://www.ncbi.nlm.nih.gov/pubmed/17033622	
	21964575	Rafnar T et al. Mutations in BRIP1 confer high risk of ovarian cancer. Nat Genet. 2011 Oct 2;43(11):1104-7. http://www.ncbi.nlm.nih.gov/pubmed/21964575	
	23586058	Apostolou P and Fostira F. Hereditary breast cancer: the era of new susceptibility genes. BioMed Res Int. 2013;747318. PMID 23586058. http://www.ncbi.nlm.nih.gov/pubmed/23586058	
	24556621	Leongamornlert D et al. Frequent germline deleterious mutations in DNA repair genes in familial prostate cancer cases are associated with advanced disease. Br J Cancer. 2014 Mar 18;110(6):1663-72. http://www.ncbi.nlm.nih.gov/pubmed/24556621	
	26315354	Ramus SJ et al. Germline Mutations in the BRIP1, BARD1, PALB2, and NBN Genes in Women With Ovarian Cancer. J Natl Cancer Inst. 2015 Aug 27;107(11). http://www.ncbi.nlm.nih.gov/pubmed/26315354	
	26720728	Norquist BM et al. Inherited Mutations in Women With Ovarian Carcinoma. Jama Oncology. 2015 Dec 30:1-9. http://www.ncbi.nlm.nih.gov/pubmed/26720728	
	26921362	Easton DF et al. No evidence that protein truncating variants in BRIP1 are associated with breast cancer risk: implications for gene panel testing. Journal Of Medical Genetics. 2016 Feb 26. http://www.ncbi.nlm.nih.gov/pubmed/26921362	
	26968956	Ghazwani Y et al. Clinical characteristics and genetic subtypes of Fanconi anemia in Saudi patients. Cancer Genet. 2016 Apr;209(4):171-6. http://www.ncbi.nlm.nih.gov/pubmed/26968956	
	27433846	Pritchard CC et al. Inherited DNA-Repair Gene Mutations in Men with Metastatic Prostate Cancer. N Engl J Med. 2016 Aug 4;375(5):443-53. http://www.ncbi.nlm.nih.gov/pubmed/27433846	
	28418444	Couch FJ et al. Associations Between Cancer Predisposition Testing Panel Genes and Breast Cancer. JAMA Oncol. 2017 Apr 13. http://www.ncbi.nlm.nih.gov/pubmed/28418444	
	28657667	Pilié PG et al. Germline genetic variants in men with prostate cancer and one or more additional cancers. Cancer. 2017 Jun 28. http://www.ncbi.nlm.nih.gov/pubmed/28657667	
	28888541	Lilyquist J et al. Frequency of mutations in a large series of clinically ascertained ovarian cancer cases tested on multi-gene panels compared to reference controls. Gynecol Oncol. 2017 Nov;147(2):375-380. 28888541 http://www.ncbi.nlm.nih.gov/pubmed/28888541	
	29356034	Beebe-Dimmer JL et al. Rare germline mutations in African American men diagnosed with early-onset prostate cancer. The Prostate. 2018 Apr 78(5):321-326. http://www.ncbi.nlm.nih.gov/pubmed/29356034	
	29368626	Weber-Lassalle N et al. BRIP1 loss-of-function mutations confer high risk for familial ovarian cancer, but not familial breast cancer. Breast Cancer Research : Bcr. 2018 Jan 24 20(1):7. http://www.ncbi.nlm.nih.gov/pubmed/29368626	
	NCCN	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Breast and Ovarian. http://www.nccn.org/clinical.asp
CASR	11013439	Hendy GN et al. Mutations of the calcium-sensing receptor (CASR) in familial hypocalciuric hypercalcemia, neonatal severe hyperparathyroidism, and autosomal dominant hypocalcemia. Human Mutation. 2000 Oct 16(4):281-96. http://www.ncbi.nlm.nih.gov/pubmed/11013439	
	16497624	Felderbauer P et al. Mutations in the calcium-sensing receptor: a new genetic risk factor for chronic pancreatitis?. Scandinavian Journal Of Gastroenterology. 2006 Mar 41(3):343-8. http://www.ncbi.nlm.nih.gov/pubmed/16497624	
	18680227	Muddana V et al. Association between calcium sensing receptor gene polymorphisms and chronic pancreatitis in a US population: role of serine protease inhibitor Kazal 1 type and alcohol. World Journal Of Gastroenterology. 2008 Jul 28 14(28):4486-91. http://www.ncbi.nlm.nih.gov/pubmed/18680227	

Gene	PMID	Reference Text
CASR	18751724	Obermannova B et al. Unusually severe phenotype of neonatal primary hyperparathyroidism due to a heterozygous inactivating mutation in the CASR gene. <i>European Journal Of Pediatrics</i> . 2009 168(5):569-73. http://www.ncbi.nlm.nih.gov/pubmed/18751724
	18938753	Murugaian EE et al. Novel mutations in the calcium sensing receptor gene in tropical chronic pancreatitis in India. <i>Scandinavian Journal Of Gastroenterology</i> . 2008 Jan 43(1):117-21. http://www.ncbi.nlm.nih.gov/pubmed/18938753
	21844754	LaRusch J and Whitcomb DC. Genetics of pancreatitis. <i>Current Opinion In Gastroenterology</i> . 2011 Sep 27(5):467-74. http://www.ncbi.nlm.nih.gov/pubmed/21844754
	24624459	LaRusch J, Solomon S, Whitcomb DC. Pancreatitis Overview. 2014 Mar 13. In: Adam MP, Ardinger HH, Pagon RA, et al., editors. <i>GeneReviews</i> ® [Internet]. Seattle (WA): University of Washington, Seattle; 1993-2017. http://www.ncbi.nlm.nih.gov/pubmed/24624459
	26166472	Masson E et al. Overrepresentation of Rare CASR Coding Variants in a Sample of Young French Patients With Idiopathic Chronic Pancreatitis. <i>Pancreas</i> . 2015 Aug 44(6):996-8. http://www.ncbi.nlm.nih.gov/pubmed/26166472
	28122587	Vahe C et al. Diseases associated with calcium-sensing receptor. <i>Orphanet Journal Of Rare Diseases</i> . 2017 Jan 25 12(1):19. http://www.ncbi.nlm.nih.gov/pubmed/28122587
	9039332	Pearce SH et al. Calcium-sensing receptor mutations in familial hypocalciuric hypercalcaemia with recurrent pancreatitis. <i>Clinical Endocrinology</i> . 1996 45(6):675-80. http://www.ncbi.nlm.nih.gov/pubmed/9039332
	CDC73	12434154
12755959		Chen JD et al. Hyperparathyroidism-jaw tumour syndrome. <i>J Intern Med</i> . 2003 Jun;253(6):634-42. http://www.ncbi.nlm.nih.gov/pubmed/12755959
15579037		Tan MH et al. Renal neoplasia in the hyperparathyroidism-jaw tumor syndrome. <i>Curr Mol Med</i> . 2004 Dec;4(8):895-7. http://www.ncbi.nlm.nih.gov/pubmed/15579037
15606373		Bradley KJ et al. Uterine tumours are a phenotypic manifestation of the hyperparathyroidism-jaw tumour syndrome. <i>J Intern Med</i> . 2005 Jan;257(1):18-26. http://www.ncbi.nlm.nih.gov/pubmed/15606373
20052758		Newey PJ et al. Cell division cycle protein 73 homolog (CDC73) mutations in the hyperparathyroidism-jaw tumor syndrome (HPT-JT) and parathyroid tumors. <i>Hum Mutat</i> . 2010 Mar;31(3):295-307. http://www.ncbi.nlm.nih.gov/pubmed/20052758
20301744		Jackson MA et al. CDC73-related disorders. 2008 Dec 31 [updated 2015 Jan 15]. In: Pagon RA, Adam MP, Ardinger HH, Wallace SE, Amemiya A, Bean LJH, Bird TD, Ledbetter N, Mefford HC, Smith RJH, Stephens K, editors. <i>SourceGeneReviews</i> ® [Internet]. Seattle (WA): University of Washington, Seattle; 1993-2017. http://www.ncbi.nlm.nih.gov/pubmed/20301744
21652691		Frank-Raue K et al. CDC73-related hereditary hyperparathyroidism: five new mutations and the clinical spectrum. <i>Eur J Endocrinol</i> . 2011 Sep;165(3):477-83. http://www.ncbi.nlm.nih.gov/pubmed/21652691
23293331		Bricaire L et al. Frequent large germline HRPT2 deletions in a French National cohort of patients with primary hyperparathyroidism. <i>J Clin Endocrinol Metab</i> . 2013 Feb;98(2):E403-8. http://www.ncbi.nlm.nih.gov/pubmed/23293331
26650250		Davidson JT et al. Parathyroid Cancer in the Pediatric Patient. <i>J Pediatr Hematol Oncol</i> . 2016 Jan;38(1):32-7. http://www.ncbi.nlm.nih.gov/pubmed/26650250
CDH1		10072428
	11729114	Pharoah PD et al. Incidence of gastric cancer and breast cancer in CDH1 (E-cadherin) mutation carriers from hereditary diffuse gastric cancer families. <i>Gastroenterology</i> . 2001 Dec;121(6):1348-53. http://www.ncbi.nlm.nih.gov/pubmed/11729114

Gene	PMID	Reference Text
CDH1	11968083	Oliveira C et al. Screening E-cadherin in gastric cancer families reveals germline mutations only in hereditary diffuse gastric cancer kindred. Hum Mutat. 2002 May;19(5):510-7. http://www.ncbi.nlm.nih.gov/pubmed/11968083
	17221870	More H et al. Identification of seven novel germline mutations in the human E-cadherin (CDH1) gene. Hum Mutat. 2007 Feb;28(2):203. http://www.ncbi.nlm.nih.gov/pubmed/17221870
	17545690	Kaurah P et al. Founder and recurrent CDH1 mutations in families with hereditary diffuse gastric cancer. JAMA. 2007 Jun 6;297(21):2360-72. http://www.ncbi.nlm.nih.gov/pubmed/17545690
	19031083	van der Post RS et al. Immunohistochemistry is not an accurate first step towards the molecular diagnosis of MUTYH-associated polyposis. Virchows Arch. 2009 Jan;454(1):25-9. http://www.ncbi.nlm.nih.gov/pubmed/19031083
	23709761	Benusiglio P et al. CDH1 germline mutations and the hereditary diffuse gastric and lobular breast cancer syndrome: a multicentre study. J Med Genet. 2013 Jul;50(7):486-9 http://www.ncbi.nlm.nih.gov/pubmed/23709761
	24366306	Petridis C et al. Germline CDH1 mutations in bilateral lobular carcinoma in situ. Br J Cancer. 2014 Feb 18;110(4):1053-7. http://www.ncbi.nlm.nih.gov/pubmed/24366306
	26072394	van der Post RS et al. Accuracy of Hereditary Diffuse Gastric Cancer Testing Criteria and Outcomes in Patients With a Germline Mutation in CDH1. Gastroenterology. 2015 Jun 11. pii: S0016-5085(15)00819-7. http://www.ncbi.nlm.nih.gov/pubmed/26072394
	26182300	Hansford S et al. Hereditary diffuse gastric cancer syndrome: CDH1 mutations and beyond. JAMA Oncol. 2015 Apr;1(1):23-32. http://www.ncbi.nlm.nih.gov/pubmed/26182300
	9537325	Guilford P. E-cadherin germline mutations in familial gastric cancer. Nature. 1998 Mar 26;392(6674):402-5. http://www.ncbi.nlm.nih.gov/pubmed/9537325
	NCCN	NCCN Guidelines. Gastric Cancer. http://www.nccn.org/clinical.asp
	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Breast and Ovarian. http://www.nccn.org/clinical.asp
	CDK4	17047042
23384855		Puntovoll HE et al. Melanoma prone families with CDK4 germline mutation: phenotypic profile and associations with MC1R variants. J Med Genet. 2013 Apr;50(4):264-70. http://www.ncbi.nlm.nih.gov/pubmed/23384855
28283772		Leachman SA et al. Identification, genetic testing, and management of hereditary melanoma. Cancer Metastasis Rev. 2017 Mar;36(1):77-90. http://www.ncbi.nlm.nih.gov/pubmed/28283772
CDKN2A	10506626	Kefford RF et al. Counseling and DNA testing for individuals perceived to be genetically predisposed to melanoma: A consensus statement of the Melanoma Genetics Consortium. J Clin Oncol. 1999 Oct;17(10):3245-51. http://www.ncbi.nlm.nih.gov/pubmed/10506626
	11136714	Randerson-Moor JA et al. A germline deletion of p14(ARF) but not CDKN2A in a melanoma-neural system tumour syndrome family. Hum Mol Genet. 2001 Jan 1;10(1):55-62. http://www.ncbi.nlm.nih.gov/pubmed/11136714
	12072543	Bishop DT et al. Geographical variation in the penetrance of CDKN2A mutations for melanoma. J Natl Cancer Inst. 2002 Jun 19;94(12):894-903. http://www.ncbi.nlm.nih.gov/pubmed/12072543
	14871223	Czajkowski R et al. FAMMM syndrome: pathogenesis and management. Dermatol Surg. 2004 Feb;30(2 Pt 2):291-6. http://www.ncbi.nlm.nih.gov/pubmed/14871223
	16234564	Begg CB et al. Lifetime risk of melanoma in CDKN2A mutation carriers in a population-based sample. J Natl Cancer Inst. 2005 Oct 19;97(20):1507-15. http://www.ncbi.nlm.nih.gov/pubmed/16234564

Gene	PMID	Reference Text
CDKN2A	18981015	de Snoo FA et al. Increased risk of cancer other than melanoma in CDKN2A founder mutation (p16-Leiden)-positive melanoma families. Clin Cancer Res. 2008 Nov 1;14(21):7151-7. http://www.ncbi.nlm.nih.gov/pubmed/18981015
	23135763	Canto MI et al. International Cancer of the Pancreas Screening (CAPS) Consortium summit on the management of patients with increased risk for familial pancreatic cancer. Gut. 2013 Mar;62(3):339-47. http://www.ncbi.nlm.nih.gov/pubmed/23135763
	24935963	Helgadóttir H et al. High risk of tobacco-related cancers in CDKN2A mutation-positive melanoma families. J Med Genet. 2014 Aug;51(8):545-52. http://www.ncbi.nlm.nih.gov/pubmed/24935963
	25227142	Potjer TP et al. Prospective risk of cancer and the influence of tobacco use in carriers of the p16-Leiden germline variant. Eur J Hum Genet. 2015 May;23(5):711-4. http://www.ncbi.nlm.nih.gov/pubmed/25227142
	25803691	Wadt KA et al. Molecular characterization of melanoma cases in Denmark suspected of genetic predisposition. Plo S One. 2015 Mar 24;10(3):e0122662 http://www.ncbi.nlm.nih.gov/pubmed/25803691
	26892651	Soura E et al. Hereditary melanoma: Update on syndromes and management: Emerging melanoma cancer complexes and genetic counseling. J Am Acad Dermatol. 2016 Mar;74(3):411-20. http://www.ncbi.nlm.nih.gov/pubmed/26892651
	27114589	Vasen H et al. Benefit of Surveillance for Pancreatic Cancer in High-Risk Individuals: Outcome of Long-Term Prospective Follow-Up Studies From Three European Expert Centers. J Clin Oncol. 2016 Jun 10;34(17):2010-9. http://www.ncbi.nlm.nih.gov/pubmed/27114589
	27978560	Pearlman R et al. Prevalence and Spectrum of Germline Cancer Susceptibility Gene Mutations Among Patients With Early-Onset Colorectal Cancer. JAMA Oncol. 2017 Apr 1;3(4):464-471. http://www.ncbi.nlm.nih.gov/pubmed/27978560
	28135145	Yurgelun MB et al. Cancer Susceptibility Gene Mutations in Individuals With Colorectal Cancer. J Clin Oncol. 2017 Apr 1;35(10):1086-1095. http://www.ncbi.nlm.nih.gov/pubmed/28135145
	CFTR	15379964
20301428		Adam MP et al. Cystic Fibrosis and Congenital Absence of the Vas Deferens. 1993. http://www.ncbi.nlm.nih.gov/pubmed/20301428
21594800		Cai Z et al. Application of high-resolution single-channel recording to functional studies of cystic fibrosis mutants. Methods In Molecular Biology (Clifton, N.J.). 2011 741:419-41. http://www.ncbi.nlm.nih.gov/pubmed/21594800
21658649		Bombieri C et al. Recommendations for the classification of diseases as CFTR-related disorders. Journal Of Cystic Fibrosis : Official Journal Of The European Cystic Fibrosis Society. 2011 Jun 10 Suppl 2:S86-102. http://www.ncbi.nlm.nih.gov/pubmed/21658649
24624459		LaRusch J, Solomon S, Whitcomb DC. Pancreatitis Overview. 2014 Mar 13. In: Adam MP, Ardinger HH, Pagon RA, et al., editors. GeneReviews® [Internet]. Seattle (WA): University of Washington, Seattle; 1993-2017. http://www.ncbi.nlm.nih.gov/pubmed/24624459
25033378		LaRusch J et al. Mechanisms of CFTR functional variants that impair regulated bicarbonate permeation and increase risk for pancreatitis but not for cystic fibrosis. P Lo S Genetics. 2014 Jul 10(7):e1004376. http://www.ncbi.nlm.nih.gov/pubmed/25033378
CHEK2	14612911	Seppala EH et al. CHEK2 variants associate with hereditary prostate cancer. Br J Cancer. 2003 Nov 17;89(10):1966-70. http://www.ncbi.nlm.nih.gov/pubmed/14612911
	14648718	Neuhausen S et al. Role of CHEK2*1100delC in unselected series of non-BRCA1/2 male breast cancers. Int J Cancer. 2004 Jan 20;108(3):477-8. http://www.ncbi.nlm.nih.gov/pubmed/14648718
	14648719	Ohayon T et al. CHEK2*1100delC and male breast cancer risk in Israel. Int J Cancer. 2004 Jan 20;108(3):479-80. http://www.ncbi.nlm.nih.gov/pubmed/14648719

Gene	PMID	Reference Text
CHEK2	15087378	Cybulski C et al. A novel founder CHEK2 mutation is associated with increased prostate cancer risk. <i>Cancer Res.</i> 2004 Apr 15;64(8):2677-9. http://www.ncbi.nlm.nih.gov/pubmed/15087378
	15122511	The CHEK2 Breast Cancer Consortium. CHEK2*1100delC and susceptibility to breast cancer: a collaborative analysis involving 10,860 breast cancer cases and 9,065 controls from 10 studies. <i>Am J Hum Genet.</i> 2004 Jun;74(6):1175-82. http://www.ncbi.nlm.nih.gov/pubmed/15122511
	15492928	Cybulski C et al. CHEK2 is a multiorgan cancer susceptibility gene. <i>Am J Hum Genet.</i> 2004 Dec;75(6):1131-5. http://www.ncbi.nlm.nih.gov/pubmed/15492928
	16816021	Kilpivaara O et al. CHEK2 I157T associates with familial and sporadic colorectal cancer. <i>J Med Genet.</i> 2006 Jul;43(7):e34. http://www.ncbi.nlm.nih.gov/pubmed/16816021
	17661168	Falchetti M et al. BRCA1/BRCA2 rearrangements and CHEK2 common mutations are infrequent in Italian male breast cancer cases. <i>Breast Cancer Res Treat.</i> 2008 Jul;110(1):161-7. http://www.ncbi.nlm.nih.gov/pubmed/17661168
	18759107	Wasielewski M et al. CHEK2 1100delC and male breast cancer in the Netherlands. <i>Breast Cancer Res Treat.</i> 2009 Jul;116(2):397-400. http://www.ncbi.nlm.nih.gov/pubmed/18759107
	19876921	Suchy J et al. CHEK2 mutations and HNPCC-related colorectal cancer. <i>Int J Cancer.</i> 2010 Jun 15;126(12):3005-9. http://www.ncbi.nlm.nih.gov/pubmed/19876921
	21244692	Le Calvez-Kelm et al. Rare, evolutionarily unlikely missense substitutions in CHEK2 contribute to breast cancer susceptibility: results from a breast cancer family registry case-control mutation-screening study. <i>Breast Cancer Res.</i> 2011 Jan 18;13(1):R6. http://www.ncbi.nlm.nih.gov/pubmed/21244692
	22799331	Liu C et al. The CHEK2 I157T variant and breast cancer susceptibility: a systematic review and meta-analysis. <i>Asian Pac J Cancer Prev.</i> 2012;13(4):1355-60. http://www.ncbi.nlm.nih.gov/pubmed/22799331
	23109706	Weischer M et al. CHEK2 1100delC heterozygosity in women with breast cancer associated with early death, breast cancer-specific death, and increased risk of a second breast cancer. <i>J Clin Oncol.</i> 2012 Dec 10;30(35):4308-16. http://www.ncbi.nlm.nih.gov/pubmed/23109706
	23149842	Cybulski C et al. An inherited NBN mutation is associated with poor prognosis prostate cancer. <i>Br J Cancer.</i> 2013 Feb 5;108(2):461-8. http://www.ncbi.nlm.nih.gov/pubmed/23149842
	23296741	Teodorczyk U et al. The risk of gastric cancer in carriers of CHEK2 mutations. <i>Fam Cancer.</i> 2013 Sep;12(3):473-8 http://www.ncbi.nlm.nih.gov/pubmed/23296741
	23713947	Han FF et al. The effect of CHEK2 variant I157T on cancer susceptibility: evidence from a meta-analysis. <i>DNA Cell Biol.</i> 2013 Jun;32(6):329-35. http://www.ncbi.nlm.nih.gov/pubmed/23713947
	25431674	Hale V et al. CHEK2 (*) 1100delC Mutation and Risk of Prostate Cancer. <i>Prostate Cancer.</i> 2014;2014:294575. http://www.ncbi.nlm.nih.gov/pubmed/25431674
	25583358	Siolek M et al. CHEK2 mutations and the risk of papillary thyroid cancer. <i>Int J Cancer.</i> 2015 Aug 1;137(3):548-52. http://www.ncbi.nlm.nih.gov/pubmed/25583358
	26884562	Näslund-Koch C et al. Increased Risk for Other Cancers in Addition to Breast Cancer for CHEK2*1100delC Heterozygotes Estimated From the Copenhagen General Population Study. <i>J Clin Oncol.</i> 2016 Apr 10;34(11):1208-16. http://www.ncbi.nlm.nih.gov/pubmed/26884562
	27433846	Pritchard CC et al. Inherited DNA-Repair Gene Mutations in Men with Metastatic Prostate Cancer. <i>N Engl J Med.</i> 2016 Aug 4;375(5):443-53. http://www.ncbi.nlm.nih.gov/pubmed/27433846
	27716369	Muranen TA et al. Patient survival and tumor characteristics associated with CHEK2:p.I157T - findings from the Breast Cancer Association Consortium. <i>Breast Cancer Res.</i> 2016 Oct 3;18(1):98.27716369 http://www.ncbi.nlm.nih.gov/pubmed/27716369
	27751358	Leedom TP et al. Breast cancer risk is similar for CHEK2 founder and non-founder mutation carriers. <i>Cancer Genet.</i> 2016 Sep;209(9):403-407. http://www.ncbi.nlm.nih.gov/pubmed/27751358

Gene	PMID	Reference Text
CHEK2	28008555	Pritzlaff M et al. Male breast cancer in a multi-gene panel testing cohort: insights and unexpected results. <i>Breast Cancer Res Treat.</i> 2017 Feb;161(3):575-586. http://www.ncbi.nlm.nih.gov/pubmed/28008555
	28418444	Couch FJ et al. Associations Between Cancer Predisposition Testing Panel Genes and Breast Cancer. <i>JAMA Oncol.</i> 2017 Apr 13. http://www.ncbi.nlm.nih.gov/pubmed/28418444
	28649662	Slavin TP et al. The contribution of pathogenic variants in breast cancer susceptibility genes to familial breast cancer risk. <i>NPJ Breast Cancer.</i> 2017 Jun 9;3:22. http://www.ncbi.nlm.nih.gov/pubmed/28649662
	28874143	Hallamies S et al. CHEK2 c.1100delC mutation is associated with an increased risk for male breast cancer in Finnish patient population. <i>BMC Cancer.</i> 2017 Sep 5;17(1):620. http://www.ncbi.nlm.nih.gov/pubmed/28874143
	29978187	Carlo MI et al. Prevalence of Germline Mutations in Cancer Susceptibility Genes in Patients With Advanced Renal Cell Carcinoma. <i>JAMA Oncol.</i> 2018 Jul 5. http://www.ncbi.nlm.nih.gov/pubmed/29978187
	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Breast and Ovarian. http://www.nccn.org/clinical.asp
	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Breast and Ovarian. http://www.nccn.org/clinical.asp
	CTRC	18059268
18172691		Masson E et al. Association of rare chymotrypsinogen C (CTRC) gene variations in patients with idiopathic chronic pancreatitis. <i>Human Genetics.</i> 2008 Feb 123(1):83-91. http://www.ncbi.nlm.nih.gov/pubmed/18172691
22427236		Rosendahl J et al. CFTR, SPINK1, CTRC and PRSS1 variants in chronic pancreatitis: is the role of mutated CFTR overestimated?. <i>Gut.</i> 2013 Apr 62(4):582-92. http://www.ncbi.nlm.nih.gov/pubmed/22427236
22942235		Beer S et al. Comprehensive functional analysis of chymotrypsin C (CTRC) variants reveals distinct loss-of-function mechanisms associated with pancreatitis risk. <i>Gut.</i> 2013 Nov 62(11):1616-24. http://www.ncbi.nlm.nih.gov/pubmed/22942235
23622139		Whitcomb DC. Genetic risk factors for pancreatic disorders. <i>Gastroenterology.</i> 2013 Jun 144(6):1292-302. http://www.ncbi.nlm.nih.gov/pubmed/23622139
24624459		LaRusch J, Solomon S, Whitcomb DC. Pancreatitis Overview. 2014 Mar 13. In: Adam MP, Ardinger HH, Pagon RA, et al., editors. <i>GeneReviews</i> ® [Internet]. Seattle (WA): University of Washington, Seattle; 1993-2017. http://www.ncbi.nlm.nih.gov/pubmed/24624459
DICER1		19556464
	20822816	Priest JR et al. Nasal chondromesenchymal hamartoma in children with pleuropulmonary blastoma--A report from the International Pleuropulmonary Blastoma Registry registry. <i>Int J Pediatr Otorhinolaryngol.</i> 2010 Nov;74(11):1240-4. 20822816 http://www.ncbi.nlm.nih.gov/pubmed/20822816
	21036787	Bahubeshi A et al. Germline DICER1 mutations and familial cystic nephroma. <i>J Med Genet.</i> 2010 Dec;47(12):863-6. 21036787 http://www.ncbi.nlm.nih.gov/pubmed/21036787
	21156700	Priest JR et al. Ciliary body medulloepithelioma: four cases associated with pleuropulmonary blastoma--a report from the International Pleuropulmonary Blastoma Registry. <i>Br J Ophthalmol.</i> 2011 Jul;95(7):1001-5. 21156700 http://www.ncbi.nlm.nih.gov/pubmed/21156700
	21205968	Rio Frio et al. DICER1 mutations in familial multinodular goiter with and without ovarian Sertoli-Leydig cell tumors. <i>JAMA.</i> 2011 Jan 5;305(1):68-77. 21205968 http://www.ncbi.nlm.nih.gov/pubmed/21205968
	21266384	Slade I et al. DICER1 syndrome: clarifying the diagnosis, clinical features and management implications of a pleiotropic tumour predisposition syndrome. <i>J Med Genet.</i> 2011 Apr;48(4):273-8. 21266384 http://www.ncbi.nlm.nih.gov/pubmed/21266384

Gene	PMID	Reference Text	
DICER1	21501861	Schultz KA et al. Ovarian sex cord-stromal tumors, pleuropulmonary blastoma and DICER1 mutations: a report from the International Pleuropulmonary Blastoma Registry. <i>Gynecol Oncol</i> . 2011 Aug;122(2):246-50. 21501861 http://www.ncbi.nlm.nih.gov/pubmed/21501861	
	21882293	Foulkes WD et al. Extending the phenotypes associated with DICER1 mutations. <i>Hum Mutat</i> . 2011 Dec;32(12):1381-4. 21882293 http://www.ncbi.nlm.nih.gov/pubmed/21882293	
	22180160	Doros L et al. DICER1 mutations in embryonal rhabdomyosarcomas from children with and without familial PPB-tumor predisposition syndrome. <i>Pediatr Blood Cancer</i> . 2012 Sep;59(3):558-60. 22180160 http://www.ncbi.nlm.nih.gov/pubmed/22180160	
	24676357	Klein S et al. Expanding the phenotype of mutations in DICER1: mosaic missense mutations in the RNase IIIb domain of DICER1 cause GLOW syndrome. <i>J Med Genet</i> . 2014 May;51(5):294-302. 24676357 http://www.ncbi.nlm.nih.gov/pubmed/24676357	
	24761742	Doros et al. DICER1-Related Disorders. 2014 Apr 24. In: Pagon RA, Bird TD, Dolan CR, et al., editors. <i>GeneReviews</i> [Internet]. Seattle (WA): University of Washington, Seattle; 1993-. http://www.ncbi.nlm.nih.gov/books/NBK196157/	
	24839956	de Kock et al. Pituitary blastoma: a pathognomonic feature of germ-line DICER1 mutations. <i>Acta Neuropathol</i> . 2014b Jul;128(1):111-22. 24839956 http://www.ncbi.nlm.nih.gov/pubmed/24839956	
	25022261	de Kock et al. Germ-line and somatic DICER1 mutations in pineoblastoma. <i>Acta Neuropathol</i> . 2014a Oct;128(4):583-95. 25022261 http://www.ncbi.nlm.nih.gov/pubmed/25022261	
	25176334	Foulkes WD et al. DICER1: mutations, microRNAs and mechanisms. <i>Nat Rev Cancer</i> . 2014 Oct;14(10):662-72. 25176334 http://www.ncbi.nlm.nih.gov/pubmed/25176334	
	26475046	de Kock et al. High-sensitivity sequencing reveals multi-organ somatic mosaicism causing DICER1 syndrome. <i>J Med Genet</i> . 2016 Jan;53(1):43-52. 26475046 http://www.ncbi.nlm.nih.gov/pubmed/26475046	
	26555935	Rutter MM et al. DICER1 Mutations and Differentiated Thyroid Carcinoma: Evidence of a Direct Association. <i>J Clin Endocrinol Metab</i> . 2016 Jan;101(1):1-5. 26555935 http://www.ncbi.nlm.nih.gov/pubmed/26555935	
	26566882	Palculict TB et al. Identification of germline DICER1 mutations and loss of heterozygosity in familial Wilms tumour. <i>J Med Genet</i> . 2015 Nov 13. pii: jmedgenet-2015-103311. 26566882 http://www.ncbi.nlm.nih.gov/pubmed/26566882	
	EPCAM	17448233	Goulet O et al. Intestinal epithelial dysplasia (tufting enteropathy). <i>Orphanet J Rare Dis</i> . 2007 Apr 20;2:20.0 http://www.ncbi.nlm.nih.gov/pubmed/17448233
		18572020	Sivagnanam M et al. Identification of EpCAM as the gene for congenital tufting enteropathy. <i>Gastroenterology</i> . 2008 Aug;135(2):429-37. http://www.ncbi.nlm.nih.gov/pubmed/18572020
19098912		Ligtenberg MJ et al. Heritable somatic methylation and inactivation of MSH2 in families with Lynch syndrome due to deletion of the 3' exons of TACSTD1. <i>Nat Genet</i> . 2009 Jan;41(1):112-7. http://www.ncbi.nlm.nih.gov/pubmed/19098912	
19177550		Kovacs ME et al. Deletions removing the last exon of TACSTD1 constitute a distinct class of mutations predisposing to Lynch syndrome. <i>Hum Mutat</i> . 2009 Feb;30(2):197-203. http://www.ncbi.nlm.nih.gov/pubmed/19177550	
20442441		Wimmer K and Kratz CP. Constitutional mismatch repair-deficiency syndrome. <i>Haematologica</i> . 2010 May; 95(5): 699-701. http://www.ncbi.nlm.nih.gov/pubmed/20442441	
20531397		Durno CA et al. The gastrointestinal phenotype of germline biallelic mismatch repair gene mutations. <i>Am J Gastroenterol</i> . 2010 Nov;105(11):2449-56. http://www.ncbi.nlm.nih.gov/pubmed/20531397	
21145788		Kempers MJ et al. Risk of colorectal and endometrial cancers in EPCAM deletion-positive Lynch syndrome: a cohort study. <i>Lancet Oncol</i> . 2011 Jan;12(1):49-55. http://www.ncbi.nlm.nih.gov/pubmed/21145788	
21227399		Rumilla K et al. Frequency of deletions of EPCAM (TACSTD1) in MSH2-associated Lynch syndrome cases. <i>J Mol Diagn</i> . 2011 Jan;13(1):93-9. http://www.ncbi.nlm.nih.gov/pubmed/21227399	

Gene	PMID	Reference Text
EPCAM	21769135	Lynch HT et al. Lynch syndrome-associated extracolonic tumors are rare in two extended families with the same EPCAM deletion. <i>Am J Gastroenterol.</i> 2011 Oct;106(10):1829-36. http://www.ncbi.nlm.nih.gov/pubmed/21769135
	23264089	Ligtenberg MJ et al. EPCAM deletion carriers constitute a unique subgroup of Lynch syndrome patients. <i>Fam Cancer.</i> 2013 Jun;12(2):169-74. http://www.ncbi.nlm.nih.gov/pubmed/23264089
	23454724	Li-Chang HH et al. Colorectal cancer in a 9-year-old due to combined EPCAM and MSH2 germline mutations: case report of a unique genotype and immunophenotype. <i>J Clin Pathol.</i> 2013 Jul;66(7):631-3 http://www.ncbi.nlm.nih.gov/pubmed/23454724
	24425144	Ryan S et al. Risk of prostate cancer in Lynch syndrome: a systematic review and meta-analysis. <i>Cancer Epidemiology, Biomarkers & Prevention : A Publication Of The American Association For Cancer Research, Cosponsored By The American Society Of Preventive Oncology.</i> 2014 Mar 23(3):437-49. http://www.ncbi.nlm.nih.gov/pubmed/24425144
	24440087	Bakry D et al. Genetic and clinical determinants of constitutional mismatch repair deficiency syndrome: report from the constitutional mismatch repair deficiency consortium. <i>Eur J Cancer.</i> 2014 Mar;50(5):987-96. http://www.ncbi.nlm.nih.gov/pubmed/24440087
	24737826	Wimmer K et al. Diagnostic criteria for constitutional mismatch repair deficiency syndrome: suggestions of the European consortium 'care for CMMRD' (C4CMMRD). <i>Journal Of Medical Genetics.</i> 2014 Jun 51(6):355-65. http://www.ncbi.nlm.nih.gov/pubmed/24737826
	26657901	Møller P et al. Cancer incidence and survival in Lynch syndrome patients receiving colonoscopic and gynaecological surveillance: first report from the prospective Lynch syndrome database. <i>Gut.</i> 2017 Mar 66(3):464-472. http://www.ncbi.nlm.nih.gov/pubmed/26657901
	27261338	Møller P et al. Incidence of and survival after subsequent cancers in carriers of pathogenic MMR variants with previous cancer: a report from the prospective Lynch syndrome database. <i>Gut.</i> 2016 Jun 03. http://www.ncbi.nlm.nih.gov/pubmed/27261338
	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Colorectal. http://www.nccn.org/clinical.asp
	EPCAM-MSH2	17448233
18572020		Sivagnanam M et al. Identification of EpCAM as the gene for congenital tufting enteropathy. <i>Gastroenterology.</i> 2008 Aug;135(2):429-37. http://www.ncbi.nlm.nih.gov/pubmed/18572020
19098912		Ligtenberg MJ et al. Heritable somatic methylation and inactivation of MSH2 in families with Lynch syndrome due to deletion of the 3' exons of TACSTD1. <i>Nat Genet.</i> 2009 Jan;41(1):112-7. http://www.ncbi.nlm.nih.gov/pubmed/19098912
20442441		Wimmer K and Kratz CP. Constitutional mismatch repair-deficiency syndrome. <i>Haematologica.</i> 2010 May; 95(5): 699-701. http://www.ncbi.nlm.nih.gov/pubmed/20442441
20531397		Durno CA et al. The A1:D785 phenotype of germline biallelic mismatch repair gene mutations. <i>Am J Gastroenterol.</i> 2010 Nov;105(11):2449-56. http://www.ncbi.nlm.nih.gov/pubmed/20531397
21145788		Kempers MJ et al. Risk of colorectal and endometrial cancers in EPCAM deletion-positive Lynch syndrome: a cohort study. <i>Lancet Oncol.</i> 2011 Jan;12(1):49-55. http://www.ncbi.nlm.nih.gov/pubmed/21145788
21791569		Perez-Cabornero L et al. Frequency of rearrangements in Lynch syndrome cases associated with MSH2: characterization of a new deletion involving both EPCAM and the 5' part of MSH2. <i>Cancer Prev Res (Phila).</i> 2011 Oct;4(10):1556-62 http://www.ncbi.nlm.nih.gov/pubmed/21791569
23264089		Ligtenberg MJ et al. EPCAM deletion carriers constitute a unique subgroup of Lynch syndrome patients. <i>Fam Cancer.</i> 2013 Jun;12(2):169-74. http://www.ncbi.nlm.nih.gov/pubmed/23264089
23454724		Li-Chang HH et al. Colorectal cancer in a 9-year-old due to combined EPCAM and MSH2 germline mutations: case report of a unique genotype and immunophenotype. <i>J Clin Pathol.</i> 2013 Jul;66(7):631-3 http://www.ncbi.nlm.nih.gov/pubmed/23454724

Gene	PMID	Reference Text
<i>EPCAM-MSH2</i>	24425144	Ryan S et al. Risk of prostate cancer in Lynch syndrome: a systematic review and meta-analysis. <i>Cancer Epidemiology, Biomarkers & Prevention</i> : A Publication Of The American Association For Cancer Research, Cosponsored By The American Society Of Preventive Oncology. 2014 Mar 23(3):437-49. http://www.ncbi.nlm.nih.gov/pubmed/24425144
	24440087	Bakry D et al. Genetic and clinical determinants of constitutional mismatch repair deficiency syndrome: report from the constitutional mismatch repair deficiency consortium. <i>Eur J Cancer</i> . 2014 Mar;50(5):987-96. http://www.ncbi.nlm.nih.gov/pubmed/24440087
	24737826	Wimmer K et al. Diagnostic criteria for constitutional mismatch repair deficiency syndrome: suggestions of the European consortium 'care for CMMRD' (C4CMMRD). <i>Journal Of Medical Genetics</i> . 2014 Jun 51(6):355-65. http://www.ncbi.nlm.nih.gov/pubmed/24737826
	26657901	Møller P et al. Cancer incidence and survival in Lynch syndrome patients receiving colonoscopic and gynaecological surveillance: first report from the prospective Lynch syndrome database. <i>Gut</i> . 2017 Mar 66(3):464-472. http://www.ncbi.nlm.nih.gov/pubmed/26657901
	27261338	Møller P et al. Incidence of and survival after subsequent cancers in carriers of pathogenic MMR variants with previous cancer: a report from the prospective Lynch syndrome database. <i>Gut</i> . 2016 Jun 03. http://www.ncbi.nlm.nih.gov/pubmed/27261338
	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Colorectal. http://www.nccn.org/clinical.asp
	<i>EPCAM-MSH2-MSH6</i>	24425144
<i>FAM175A</i>	22357538	Solyon S et al. Breast cancer-associated Abraxas mutation disrupts nuclear localization and DNA damage response functions. <i>Sci Transl Med</i> . 2012 Feb 22;4(122):122ra23. http://www.ncbi.nlm.nih.gov/pubmed/22357538
	24240112	Pennington KP et al. Germline and somatic mutations in homologous recombination genes predict platinum response and survival in ovarian, fallopian tube, and peritoneal carcinomas. <i>Clin Cancer Res</i> . 2014 Feb 1;20(3):764-75. http://www.ncbi.nlm.nih.gov/pubmed/24240112
<i>FANCC</i>	17909071	Berwick M et al. Genetic heterogeneity among Fanconi anemia heterozygotes and risk of cancer. <i>Cancer Res</i> . 2007 Oct 1;67(19):9591-6. http://www.ncbi.nlm.nih.gov/pubmed/17909071
	23028338	Thompson ER et al. Exome sequencing identifies rare deleterious mutations in DNA repair genes FANCC and BLM as potential breast cancer susceptibility alleles. <i>PLoS Genet</i> . 2012 Sep;8(9):e1002894. http://www.ncbi.nlm.nih.gov/pubmed/23028338
<i>FH</i>	11248088	Launonen V et al. Inherited susceptibility to uterine leiomyomas and renal cell cancer. <i>Proc Natl Acad Sci U S A</i> . 2001 Mar 13;98(6):3387-92. http://www.ncbi.nlm.nih.gov/pubmed/11248088
	11865300	Tomlinson IP et al. Germline mutations in FH predispose to dominantly inherited uterine fibroids, skin leiomyomata and papillary renal cell cancer. <i>Nature Genetics</i> . 2002 30(4):406-10. http://www.ncbi.nlm.nih.gov/pubmed/11865300
	12761039	Alam NA et al. Genetic and functional analyses of FH mutations in multiple cutaneous and uterine leiomyomatosis, hereditary leiomyomatosis and renal cancer, and fumarate hydratase deficiency. <i>Hum Mol Genet</i> . 2003 Jun 1;12(11):1241-52. http://www.ncbi.nlm.nih.gov/pubmed/12761039
	12772087	Toro JR et al. Mutations in the fumarate hydratase gene cause hereditary leiomyomatosis and renal cell cancer in families in North America. <i>Am J Hum Genet</i> . 2003 Jul;73(1):95-106. http://www.ncbi.nlm.nih.gov/pubmed/12772087
	15937070	Wei MH et al. Novel mutations in FH and expansion of the spectrum of phenotypes expressed in families with hereditary leiomyomatosis and renal cell cancer. <i>J Med Genet</i> . 2006 Jan;43(1):18-27. http://www.ncbi.nlm.nih.gov/pubmed/15937070
	16155190	Lehtonen HJ et al. Increased risk of cancer in patients with fumarate hydratase germline mutation. <i>J Med Genet</i> . 2006 Jun;43(6):523-6. http://www.ncbi.nlm.nih.gov/pubmed/16155190

Gene	PMID	Reference Text
FH	16477632	Ylisaukko-oja SK et al. Analysis of fumarate hydratase mutations in a population-based series of early onset uterine leiomyosarcoma patients. <i>Int J Cancer</i> . 2006 Jul 15;119(2):283-7. http://www.ncbi.nlm.nih.gov/pubmed/16477632
	17392716	Refae MA et al. Hereditary leiomyomatosis and renal cell cancer: an unusual and aggressive form of hereditary renal carcinoma. <i>Nat Clin Pract Oncol</i> . 2007 Apr;4(4):256-61. http://www.ncbi.nlm.nih.gov/pubmed/17392716
	19967458	Alrashdi I et al. Hereditary leiomyomatosis and renal cell carcinoma: very early diagnosis of renal cancer in a paediatric patient. <i>Fam Cancer</i> . 2010 Jun;9(2):239-43. http://www.ncbi.nlm.nih.gov/pubmed/19967458
	20301430	Pithukpakorn M, Toro JR. Hereditary Leiomyomatosis and Renal Cell Cancer. 2006 Jul 31 [Updated 2015 Aug 6]. In: Adam MP, Ardinger HH, Pagon RA, et al., editors. <i>GeneReviews</i> ® [Internet]. Seattle (WA): University of Washington, Seattle; 1993-2018. Available from: https://www.ncbi.nlm.nih.gov/books/NBK1252/ http://www.ncbi.nlm.nih.gov/pubmed/20301430
	21398687	Gardie B et al. Novel FH mutations in families with hereditary leiomyomatosis and renal cell cancer (HLRCC) and patients with isolated type 2 papillary renal cell carcinoma. <i>J Med Genet</i> . 2011 Apr;48(4):226-34. http://www.ncbi.nlm.nih.gov/pubmed/21398687
	22069215	Mroch AR et al. Detection of a novel FH whole gene deletion in the proband leading to subsequent prenatal diagnosis in a sibship with fumarase deficiency. <i>Am J Med Genet A</i> . 2012 Jan;158A(1):155-8. http://www.ncbi.nlm.nih.gov/pubmed/22069215
	2314594	Gellera C et al. Fumarase deficiency is an autosomal recessive encephalopathy affecting both the mitochondrial and the cytosolic enzymes. <i>Neurology</i> . 1990 Mar;40(3 Pt 1):495-9. http://www.ncbi.nlm.nih.gov/pubmed/2314594
	23707781	Letouzé E et al. SDH mutations establish a hypermethylator phenotype in paraganglioma. <i>Cancer Cell</i> . 2013 Jun 10;23(6):739-52. http://www.ncbi.nlm.nih.gov/pubmed/23707781
	24334767	Castro-Vega LJ et al. Germline mutations in FH confer predisposition to malignant pheochromocytomas and paragangliomas. <i>Hum Mol Genet</i> . 2014 May 1;23(9):2440-6. http://www.ncbi.nlm.nih.gov/pubmed/24334767
	24441663	Chen YB et al. Hereditary leiomyomatosis and renal cell carcinoma syndrome-associated renal cancer: recognition of the syndrome by pathologic features and the utility of detecting aberrant succination by immunohistochemistry. <i>Am J Surg Pathol</i> . 2014 May;38(5):627-37. http://www.ncbi.nlm.nih.gov/pubmed/24441663
	24526232	Wong MH et al. Potential genetic anticipation in hereditary leiomyomatosis-renal cell cancer (HLRCC). <i>Fam Cancer</i> . 2014 Jun;13(2):281-9. http://www.ncbi.nlm.nih.gov/pubmed/24526232
	25004247	Clark GR et al. Germline FH mutations presenting with pheochromocytoma. <i>J Clin Endocrinol Metab</i> . 2014 Oct;99(10):E2046-50. http://www.ncbi.nlm.nih.gov/pubmed/25004247
	25012257	Menko FH et al. Hereditary leiomyomatosis and renal cell cancer (HLRCC): renal cancer risk, surveillance and treatment. <i>Fam Cancer</i> . 2014 Dec;13(4):637-44. http://www.ncbi.nlm.nih.gov/pubmed/25012257
	25018647	Schmidt LS and Linehan WM2. Hereditary leiomyomatosis and renal cell carcinoma. <i>Int J Nephrol Renovasc Dis</i> . 2014 Jun 20;7:253-60. http://www.ncbi.nlm.nih.gov/pubmed/25018647
	27635946	Smith SC et al. Tubulocystic Carcinoma of the Kidney With Poorly Differentiated Foci: A Frequent Morphologic Pattern of Fumarate Hydratase-deficient Renal Cell Carcinoma. <i>The American Journal Of Surgical Pathology</i> . 2016 Nov 40(11):1457-1472. http://www.ncbi.nlm.nih.gov/pubmed/27635946
	28300276	Muller M et al. Reassessing the clinical spectrum associated with Hereditary Leiomyomatosis and Renal Cell Carcinoma syndrome in French FH mutation carriers. <i>Clinical Genetics</i> . 2017 Mar 16. http://www.ncbi.nlm.nih.gov/pubmed/28300276
	28314682	Patel VM et al. Hereditary leiomyomatosis and renal cell cancer syndrome: An update and review. <i>Journal Of The American Academy Of Dermatology</i> . 2017 Mar 14. http://www.ncbi.nlm.nih.gov/pubmed/28314682
	9635293	Coughlin EM et al. Molecular analysis and prenatal diagnosis of human fumarase deficiency. <i>Mol Genet Metab</i> . 1998 Apr;63(4):254-62. http://www.ncbi.nlm.nih.gov/pubmed/9635293

Gene	PMID	Reference Text
<i>FH</i>	HLRCC Handbook	Lovitt G., ed. The HLRCC Handbook, v 2.0. HLRCC Family Alliance. 2013.
		http://www.hlrccinfo.org/handbook/HLRCC-HANDBOOK.pdf
<i>FLCN</i>	11927500	Zbar B et al. Risk of renal and colonic neoplasms and spontaneous pneumothorax in the Birt-Hogg-Dubé syndrome. <i>Cancer Epidemiol Biomarkers Prev.</i> 2002 Apr;11(4):393-400.
		http://www.ncbi.nlm.nih.gov/pubmed/11927500
	15821464	Pavlovich CP et al. Evaluation and management of renal tumors in the Birt-Hogg-Dubé syndrome. <i>J Urol.</i> 2005 May;173(5):1482-6.
		http://www.ncbi.nlm.nih.gov/pubmed/15821464
	15852235	Schmidt LS et al. Germline BHD-mutation spectrum and phenotype analysis of a large cohort of families with Birt-Hogg-Dubé syndrome. <i>Am J Hum Genet.</i> 2005 Jun;76(6):1023-33.
		http://www.ncbi.nlm.nih.gov/pubmed/15852235
	18234728	Toro JR et al. BHD mutations, clinical and molecular genetic investigations of Birt-Hogg-Dubé syndrome: a new series of 50 families and a review of published reports. <i>J Med Genet.</i> 2008 Jun;45(6):321-31.
		http://www.ncbi.nlm.nih.gov/pubmed/18234728
19785621	Kluger N et al. Birt-Hogg-Dubé syndrome: clinical and genetic studies of 10 French families. <i>Br J Dermatol.</i> 2010 Mar;162(3):527-37.	
	http://www.ncbi.nlm.nih.gov/pubmed/19785621	
19959076	Menko FH et al. Birt-Hogg-Dubé syndrome: diagnosis and management. <i>Lancet Oncol.</i> 2009 Dec;10(12):1199-206.	
	http://www.ncbi.nlm.nih.gov/pubmed/19959076	
25519458	Benusiglio PR et al. Renal cell tumour characteristics in patients with the Birt-Hogg-Dubé cancer susceptibility syndrome: a retrospective, multicentre study. <i>Orphanet J Rare Dis.</i> 2014 Oct 29;9:163.	
	http://www.ncbi.nlm.nih.gov/pubmed/25519458	
<i>HOXB13</i>	22236224	Ewing CM et al. Germline mutations in HOXB13 and prostate-cancer risk. <i>N Engl J Med.</i> 2012 Jan 12;366(2):141-9.
		http://www.ncbi.nlm.nih.gov/pubmed/22236224
	22781434	Akbari MR et al. Association between germline HOXB13 G84E mutation and risk of prostate cancer. <i>J Natl Cancer Inst.</i> 2012 Aug 22;104(16):1260-2.
	http://www.ncbi.nlm.nih.gov/pubmed/22781434	
22853031	Alane S et al. Association of a HOXB13 variant with breast cancer. <i>N Engl J Med.</i> 2012 Aug 2;367(5):480-1.	
	http://www.ncbi.nlm.nih.gov/pubmed/22853031	
<i>MAX</i>	21685915	Comino-Mendez I et al. Exome sequencing identifies MAX mutations as a cause of hereditary pheochromocytoma. <i>Nature Genetics.</i> 2011 Jul 43(7):663-7.
		http://www.ncbi.nlm.nih.gov/pubmed/21685915
	22452945	Burnichon N et al. MAX mutations cause hereditary and sporadic pheochromocytoma and paraganglioma. <i>Clinical Cancer Research : An Official Journal Of The American Association For Cancer Research.</i> 2012 May 15 18(10):2828-37.
		http://www.ncbi.nlm.nih.gov/pubmed/22452945
	24893135	Lenders JW et al. Pheochromocytoma and paraganglioma: an endocrine society clinical practice guideline. <i>The Journal Of Clinical Endocrinology And Metabolism.</i> 2014 Jun 99(6):1915-42.24893135
	http://www.ncbi.nlm.nih.gov/pubmed/24893135	
24899893	Martins R and Bugalho MJ. Paragangliomas/Pheochromocytomas: clinically oriented genetic testing. <i>International Journal Of Endocrinology.</i> 2014 2014:794187.	
	http://www.ncbi.nlm.nih.gov/pubmed/24899893	
28384794	Bausch B et al. Clinical Characterization of the Pheochromocytoma and Paraganglioma Susceptibility Genes SDHA, TMEM127, MAX, and SDHAF2 for Gene-Informed Prevention. <i>Jama Oncology.</i> 2017 Apr 06.	
	http://www.ncbi.nlm.nih.gov/pubmed/28384794	
<i>MEN1</i>	11579199	Guo SS, Sawicki MP. Molecular and genetic mechanisms of tumorigenesis in multiple endocrine neoplasia type-1. <i>Mol Endocrinol.</i> 2001;15:1653-64.
		http://www.ncbi.nlm.nih.gov/pubmed/11579199
	11739416	Brandi ML et al. Guidelines for diagnosis and therapy of MEN type 1 and type 2. <i>J Clin Endocrinol Metab.</i> 2001 Dec;86(12):5658-71.
	http://www.ncbi.nlm.nih.gov/pubmed/11739416	
12016472	Langer P et al. Adrenal involvement in multiple endocrine neoplasia type 1. <i>World J Surg.</i> 2002 Aug;26(8):891-6.	
	http://www.ncbi.nlm.nih.gov/pubmed/12016472	

Gene	PMID	Reference Text
<i>MEN1</i>	14747767	Gibril F et al. Multiple endocrine neoplasia type 1 and Zollinger-Ellison syndrome: a prospective study of 107 cases and comparison with 1009 cases from the literature. <i>Medicine (Baltimore)</i> . 2004 Jan;83(1):43-83. http://www.ncbi.nlm.nih.gov/pubmed/14747767
	20301710	Giusti F, Marini F, Brandi ML. Multiple Endocrine Neoplasia Type 1. 2005 Aug 31 [Updated 2015 Feb 12]. In: Pagon RA, Adam MP, Ardinger HH, et al., editors. <i>GeneReviews®</i> [Internet]. Seattle (WA): University of Washington, Seattle; 1993-2016. http://www.ncbi.nlm.nih.gov/books/NBK1538/
	21249756	Marini F et al. Multiple Endocrine Neoplasia Type 1 (MEN1) Syndrome. 2008 Jul 18 [Updated 2008 Aug 09]. In: Reigert-Johnson DL, Boardman LA, Hefferon T, Roberts M, editors. <i>Cancer Syndromes</i> [Internet]. Bethesda (MD): National Center for Biotechnology Information (US); 2009-. http://www.ncbi.nlm.nih.gov/pubmed/21249756
	22723327	Thakker et al. Clinical Practice Guidelines for Multiple Endocrine Neoplasia Type 1 (MEN1). <i>The Journal of Clinical Endocrinology and Metabolism</i> . 2012;97(9):2990-3011. http://www.ncbi.nlm.nih.gov/pubmed/22723327
	25494863	Dénes J et al. Heterogeneous genetic background of the association of pheochromocytoma/paraganglioma and pituitary adenoma: results from a large patient cohort. <i>J Clin Endocrinol Metab</i> . 2015 Mar;100(3):E531-41. http://www.ncbi.nlm.nih.gov/pubmed/25494863
	26363542	Norton JA et al. Multiple Endocrine Neoplasia: Genetics and Clinical Management. <i>Surgical Oncology Clinics Of North America</i> . 2015 Oct 24(4):795-832. http://www.ncbi.nlm.nih.gov/pubmed/26363542
	26813904	De Sousa SM and McCormack AI. Cutaneous lichen amyloidosis in multiple endocrine neoplasia. <i>Internal Medicine Journal</i> . 2016 Jan 46(1):116-7. http://www.ncbi.nlm.nih.gov/pubmed/26813904
	<i>MET</i>	10417759
10433944		Lubensky IA et al. Hereditary and sporadic papillary renal carcinomas with c-met mutations share a distinct morphological phenotype. <i>Am J Pathol</i> . 1999 Aug;155(2):517-26. http://www.ncbi.nlm.nih.gov/pubmed/10433944
10647647		Ornstein DK et al. Prevalence of microscopic tumors in normal appearing renal parenchyma of patients with hereditary papillary renal cancer. <i>J Urol</i> . 2000 Feb;163(2):431-3. http://www.ncbi.nlm.nih.gov/pubmed/10647647
15371818		Schmidt LS et al. Early onset hereditary papillary renal carcinoma: germline missense mutations in the tyrosine kinase domain of the met proto-oncogene. <i>J Urol</i> . 2004 Oct;172(4 Pt 1):1256-61. http://www.ncbi.nlm.nih.gov/pubmed/15371818
23882344		Mustafa S. et al. Case of hereditary papillary renal cell carcinoma. <i>J Community Hosp Intern Med Perspect</i> . 2012 Jan 26;1(4). http://www.ncbi.nlm.nih.gov/pubmed/23882344
22012259		Bertolotto C et al. A SUMOylation-defective MITF germline mutation predisposes to melanoma and renal carcinoma. <i>Nature</i> . 2011 Dec 1 480(7375):94-8. http://www.ncbi.nlm.nih.gov/pubmed/22012259
<i>MITF</i>		22080950
	23167872	Ghiorzo P et al. Prevalence of the E318K MITF germline mutation in Italian melanoma patients: associations with histological subtypes and family cancer history. <i>Pigment Cell & Melanoma Research</i> . 2013 Mar 26(2):259-62. http://www.ncbi.nlm.nih.gov/pubmed/23167872
	24767713	Gromowski T et al. Prevalence of the E318K and V320I MITF germline mutations in Polish cancer patients and multiorgan cancer risk-a population-based study. <i>Cancer Genetics</i> . 2014 Apr 207(4):128-32. http://www.ncbi.nlm.nih.gov/pubmed/24767713
	25803691	Wadt KA et al. Molecular characterization of melanoma cases in Denmark suspected of genetic predisposition. <i>Plo S One</i> . 2015 Mar 24;10(3):e0122662 http://www.ncbi.nlm.nih.gov/pubmed/25803691

Gene	PMID	Reference Text
MITF	26650189	Potrony M et al. Prevalence of MITF p.E318K in Patients With Melanoma Independent of the Presence of CDKN2A Causative Mutations. <i>Jama Dermatology</i> . 2016 Apr 1 152(4):405-12. http://www.ncbi.nlm.nih.gov/pubmed/26650189
	28283772	Leachman SA et al. Identification, genetic testing, and management of hereditary melanoma. <i>Cancer Metastasis Rev</i> . 2017 Mar;36(1):77-90. http://www.ncbi.nlm.nih.gov/pubmed/28283772
MLH1	15937084	Quehenberger F et al. Risk of colorectal and endometrial cancer for carriers of mutations of the hMLH1 and hMSH2 gene: correction for ascertainment. <i>J Med Genet</i> .2005 Jun;42(6):491-6. http://www.ncbi.nlm.nih.gov/pubmed/15937084
	19125127	Palomaki GE et al. EGAPP supplementary evidence review: DNA testing strategies aimed at reducing morbidity and mortality from Lynch syndrome. <i>Genet Med</i> . 2009 Jan;11(1):42-65. http://www.ncbi.nlm.nih.gov/pubmed/19125127
	19861671	Kastrinos F et al. Risk of pancreatic cancer in families with Lynch syndrome. <i>Jama</i> . 2009 Oct 28 302(16):1790-5. http://www.ncbi.nlm.nih.gov/pubmed/19861671
	20301390	Kohlmann W, Gruber SB. Lynch Syndrome. 2004 Feb 5 [Updated 2014 May 22]. In: Pagon RA, Adam MP, Ardinger HH, et al., editors. <i>GeneReviews</i> ® [Internet]. Seattle (WA): University of Washington, Seattle; 1993-2016. http://www.ncbi.nlm.nih.gov/books/NBK1211
	20442441	Wimmer K and Kratz CP. Constitutional mismatch repair-deficiency syndrome. <i>Haematologica</i> . 2010 May; 95(5): 699–701. http://www.ncbi.nlm.nih.gov/pubmed/20442441
	20531397	Durno CA et al. The gastrointestinal phenotype of germline biallelic mismatch repair gene mutations. <i>Am J Gastroenterol</i> . 2010 Nov;105(11):2449-56. http://www.ncbi.nlm.nih.gov/pubmed/20531397
	21642682	Bonadona V et al. Cancer risks associated with germline mutations in MLH1, MSH2, and MSH6 genes in Lynch syndrome. <i>JAMA</i> . 2011 Jun;305(22):2304-10. http://www.ncbi.nlm.nih.gov/pubmed/21642682
	23255516	Dowty JG et al. Cancer risks for MLH1 and MSH2 mutation carriers. <i>Hum Mutat</i> . 2013 Mar;34(3):490-7. http://www.ncbi.nlm.nih.gov/pubmed/23255516
	23385444	Win AK et al. Risks of colorectal and other cancers after endometrial cancer for women with Lynch syndrome. <i>J Natl Cancer Inst</i> . 2013 Feb;105(4):274-9. http://www.ncbi.nlm.nih.gov/pubmed/23385444
	24425144	Ryan S et al. Risk of prostate cancer in Lynch syndrome: a systematic review and meta-analysis. <i>Cancer Epidemiology, Biomarkers & Prevention : A Publication Of The American Association For Cancer Research, Cosponsored By The American Society Of Preventive Oncology</i> . 2014 Mar 23(3):437-49. http://www.ncbi.nlm.nih.gov/pubmed/24425144
	24440087	Bakry D et al. Genetic and clinical determinants of constitutional mismatch repair deficiency syndrome: report from the constitutional mismatch repair deficiency consortium. <i>Eur J Cancer</i> . 2014 Mar;50(5):987-96. http://www.ncbi.nlm.nih.gov/pubmed/24440087
	24737826	Wimmer K et al. Diagnostic criteria for constitutional mismatch repair deficiency syndrome: suggestions of the European consortium 'care for CMMRD' (C4CMMRD). <i>Journal Of Medical Genetics</i> . 2014 Jun 51(6):355-65. http://www.ncbi.nlm.nih.gov/pubmed/24737826
	26657901	Møller P et al. Cancer incidence and survival in Lynch syndrome patients receiving colonoscopic and gynaecological surveillance: first report from the prospective Lynch syndrome database. <i>Gut</i> . 2017 Mar 66(3):464-472. http://www.ncbi.nlm.nih.gov/pubmed/26657901
	27241104	Mills AM and Longacre TA2. Lynch Syndrome: Female Genital Tract Cancer Diagnosis and Screening. <i>Surg Pathol Clin</i> . 2016 Jun;9(2):201-14. http://www.ncbi.nlm.nih.gov/pubmed/26657901
	27261338	Møller P et al. Incidence of and survival after subsequent cancers in carriers of pathogenic MMR variants with previous cancer: a report from the prospective Lynch syndrome database. <i>Gut</i> . 2016 Jun 03. http://www.ncbi.nlm.nih.gov/pubmed/27261338
	8612988	Vasen HF et al. Cancer risk in families with hereditary nonpolyposis colorectal cancer diagnosed by mutation analysis. <i>Gastroenterology</i> . 1996 Apr;110(4):1020-7. http://www.ncbi.nlm.nih.gov/pubmed/8612988

Gene	PMID	Reference Text
<i>MLH1</i>	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Colorectal. http://www.nccn.org/clinical.asp
<i>MRE11A</i>	10612394	Stewart GS et al. The DNA double-strand break repair gene hMRE11 is mutated in individuals with an ataxia-telangiectasia-like disorder. <i>Cell</i> . 1999 Dec 10;99(6):577-87. http://www.ncbi.nlm.nih.gov/pubmed/10612394
	14684699	Heikkinen K et al. Mutation screening of Mre11 complex genes: indication of RAD50 involvement in breast and ovarian cancer susceptibility. <i>J Med Genet</i> . 2003 Dec;40(12):e131. http://www.ncbi.nlm.nih.gov/pubmed/14684699
	15574463	Fernet M et al. Identification and functional consequences of a novel MRE11 mutation affecting 10 Saudi Arabian patients with the ataxia telangiectasia-like disorder. <i>Hum Mol Genet</i> . 2005 Jan 15;14(2):307-18. http://www.ncbi.nlm.nih.gov/pubmed/15574463
	19383352	Bartkova J et al. Aberrations of the MRE11-RAD50-NBS1 DNA damage sensor complex in human breast cancer: MRE11 as a candidate familial cancer-predisposing gene. <i>Mol Oncol</i> . 2008 Dec; 2(4):296-316. http://www.ncbi.nlm.nih.gov/pubmed/19383352
	8445618	Hernandez D et al. A family showing no evidence of linkage between the ataxia telangiectasia gene and chromosome 11q22-23. <i>J Med Genet</i> . 1993 Feb;30(2):135-40. http://www.ncbi.nlm.nih.gov/pubmed/8445618
<i>MSH2</i>	15937084	Quehenberger F et al. Risk of colorectal and endometrial cancer for carriers of mutations of the hMLH1 and hMSH2 gene: correction for ascertainment. <i>J Med Genet</i> . 2005 Jun;42(6):491-6. http://www.ncbi.nlm.nih.gov/pubmed/15937084
	19125127	Palomaki GE et al. EGAPP supplementary evidence review: DNA testing strategies aimed at reducing morbidity and mortality from Lynch syndrome. <i>Genet Med</i> . 2009 Jan;11(1):42-65. http://www.ncbi.nlm.nih.gov/pubmed/19125127
	20301390	Kohlmann W, Gruber SB. Lynch Syndrome. 2004 Feb 5 [Updated 2014 May 22]. In: Pagon RA, Adam MP, Ardinger HH, et al., editors. <i>GeneReviews</i> ® [Internet]. Seattle (WA): University of Washington, Seattle; 1993-2016. http://www.ncbi.nlm.nih.gov/books/NBK1211
	20442441	Wimmer K and Kratz CP. Constitutional mismatch repair-deficiency syndrome. <i>Haematologica</i> . 2010 May; 95(5): 699–701. http://www.ncbi.nlm.nih.gov/pubmed/20442441
	20531397	Durno CA et al. The gastrointestinal phenotype of germline biallelic mismatch repair gene mutations. <i>Am J Gastroenterol</i> . 2010 Nov;105(11):2449-56. http://www.ncbi.nlm.nih.gov/pubmed/20531397
	21642682	Bonadona V et al. Cancer risks associated with germline mutations in MLH1, MSH2, and MSH6 genes in Lynch syndrome. <i>JAMA</i> . 2011 Jun;305(22):2304-10. http://www.ncbi.nlm.nih.gov/pubmed/21642682
	23255516	Dowty JG et al. Cancer risks for MLH1 and MSH2 mutation carriers. <i>Hum Mutat</i> . 2013 Mar;34(3):490-7. http://www.ncbi.nlm.nih.gov/pubmed/23255516
	23385444	Win AK et al. Risks of colorectal and other cancers after endometrial cancer for women with Lynch syndrome. <i>J Natl Cancer Inst</i> . 2013 Feb;105(4):274-9. http://www.ncbi.nlm.nih.gov/pubmed/23385444
	24425144	Ryan S et al. Risk of prostate cancer in Lynch syndrome: a systematic review and meta-analysis. <i>Cancer Epidemiology, Biomarkers & Prevention : A Publication Of The American Association For Cancer Research, Cosponsored By The American Society Of Preventive Oncology</i> . 2014 Mar 23(3):437-49. http://www.ncbi.nlm.nih.gov/pubmed/24425144
	24440087	Bakry D et al. Genetic and clinical determinants of constitutional mismatch repair deficiency syndrome: report from the constitutional mismatch repair deficiency consortium. <i>Eur J Cancer</i> . 2014 Mar;50(5):987-96. http://www.ncbi.nlm.nih.gov/pubmed/24440087
	24737826	Wimmer K et al. Diagnostic criteria for constitutional mismatch repair deficiency syndrome: suggestions of the European consortium 'care for CMMRD' (C4CMMRD). <i>Journal Of Medical Genetics</i> . 2014 Jun 51(6):355-65. http://www.ncbi.nlm.nih.gov/pubmed/24737826
	26657901	Møller P et al. Cancer incidence and survival in Lynch syndrome patients receiving colonoscopic and gynaecological surveillance: first report from the prospective Lynch syndrome database. <i>Gut</i> . 2017 Mar 66(3):464-472. http://www.ncbi.nlm.nih.gov/pubmed/26657901

Gene	PMID	Reference Text
MSH2	27241104	Mills AM and Longacre TA2. Lynch Syndrome: Female Genital Tract Cancer Diagnosis and Screening. Surg Pathol Clin. 2016 Jun;9(2):201-14. http://www.ncbi.nlm.nih.gov/pubmed/27241104
	27261338	Møller P et al. Incidence of and survival after subsequent cancers in carriers of pathogenic MMR variants with previous cancer: a report from the prospective Lynch syndrome database. Gut. 2016 Jun 03. http://www.ncbi.nlm.nih.gov/pubmed/27261338
	8612988	Vasen HF et al. Cancer risk in families with hereditary nonpolyposis colorectal cancer diagnosed by mutation analysis. Gastroenterology. 1996 Apr;110(4):1020-7. http://www.ncbi.nlm.nih.gov/pubmed/8612988
	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Colorectal. http://www.nccn.org/clinical.asp
MSH6	19125127	Palomaki GE et al. EGAPP supplementary evidence review: DNA testing strategies aimed at reducing morbidity and mortality from Lynch syndrome. Genet Med. 2009 Jan;11(1):42-65. http://www.ncbi.nlm.nih.gov/pubmed/19125127
	20028993	Baglietto L et al. Risks of Lynch syndrome cancers for MSH6 mutation carriers. J Natl Cancer Inst. 2010 Feb;102(3):193-201. http://www.ncbi.nlm.nih.gov/pubmed/20028993
	20442441	Wimmer K and Kratz CP. Constitutional mismatch repair-deficiency syndrome. Haematologica. 2010 May; 95(5): 699-701. http://www.ncbi.nlm.nih.gov/pubmed/20442441
	20487569	Talseth-Palmer BA et al. MSH6 and PMS2 mutation positive Australian Lynch syndrome families: novel mutations, cancer risk and age of diagnosis of colorectal cancer. Hered Cancer Clin Pract. 2010 May 21;8(1):5. http://www.ncbi.nlm.nih.gov/pubmed/20487569
	20531397	Durno CA et al. The gastrointestinal phenotype of germline biallelic mismatch repair gene mutations. Am J Gastroenterol. 2010 Nov;105(11):2449-56. http://www.ncbi.nlm.nih.gov/pubmed/20531397
	22331944	Win AK et al. Colorectal and other cancer risks for carriers and noncarriers from families with a DNA mismatch repair gene mutation: a prospective cohort study. J Clin Oncol. 2012 Mar;30(9):958-64. http://www.ncbi.nlm.nih.gov/pubmed/22331944
	23385444	Win AK et al. Risks of colorectal and other cancers after endometrial cancer for women with Lynch syndrome. J Natl Cancer Inst. 2013 Feb;105(4):274-9. http://www.ncbi.nlm.nih.gov/pubmed/23385444
	24425144	Ryan S et al. Risk of prostate cancer in Lynch syndrome: a systematic review and meta-analysis. Cancer Epidemiology, Biomarkers & Prevention : A Publication Of The American Association For Cancer Research, Cosponsored By The American Society Of Preventive Oncology. 2014 Mar 23(3):437-49. http://www.ncbi.nlm.nih.gov/pubmed/24425144
	24440087	Bakry D et al. Genetic and clinical determinants of constitutional mismatch repair deficiency syndrome: report from the constitutional mismatch repair deficiency consortium. Eur J Cancer. 2014 Mar;50(5):987-96. http://www.ncbi.nlm.nih.gov/pubmed/24440087
	24737826	Wimmer K et al. Diagnostic criteria for constitutional mismatch repair deficiency syndrome: suggestions of the European consortium 'care for CMMRD' (C4CMMRD). Journal Of Medical Genetics. 2014 Jun 51(6):355-65. http://www.ncbi.nlm.nih.gov/pubmed/24737826
	27241104	Mills AM and Longacre TA2. Lynch Syndrome: Female Genital Tract Cancer Diagnosis and Screening. Surg Pathol Clin. 2016 Jun;9(2):201-14. http://www.ncbi.nlm.nih.gov/pubmed/27241104
	27261338	Møller P et al. Incidence of and survival after subsequent cancers in carriers of pathogenic MMR variants with previous cancer: a report from the prospective Lynch syndrome database. Gut. 2016 Jun 03. http://www.ncbi.nlm.nih.gov/pubmed/27261338
	29345684	Roberts ME et al. MSH6 and PMS2 germ-line pathogenic variants implicated in Lynch syndrome are associated with breast cancer. Genetics In Medicine : Official Journal Of The American College Of Medical Genetics. 2018 Jan 18. http://www.ncbi.nlm.nih.gov/pubmed/29345684
	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Colorectal. http://www.nccn.org/clinical.asp

Gene	PMID	Reference Text
MUTYH	12606733	Sieber OM. Multiple colorectal adenomas, classic adenomatous polyposis, and germ-line in MYH. N Engl J Med. 2003 Feb 27;348(9):791-9. http://www.ncbi.nlm.nih.gov/pubmed/12606733
	16492921	Jenkins MA et al. Risk of colorectal cancer in monoallelic and biallelic carriers of MYH mutations: a population-based case-family study. Cancer Epidemiol Biomarkers Prev. 2006;15:312-14. http://www.ncbi.nlm.nih.gov/pubmed/16492921
	17956577	Barnetson RA. Germline mutation prevalence in the base excision repair gene, MYH, in patients with endometrial cancer. Clin Genet. 2007 Dec;72(6):551-5. http://www.ncbi.nlm.nih.gov/pubmed/17956577
	19013464	Boparai KS et al. Hyperplastic polyps and sessile serrated adenomas as a phenotypic expression of MYH-associated polyposis. Gastroenterology. 2008 Dec;135(6):2014-8. http://www.ncbi.nlm.nih.gov/pubmed/19013464
	19338676	Ashton KA et al. Genetic variants in MUTYH are not associated with endometrial cancer risk. Hereditary Cancer In Clinical Practice. 2009 7(1):3. http://www.ncbi.nlm.nih.gov/pubmed/19338676
	19620482	Lubbe SJ et al. Clinical implications of the colorectal cancer risk associated with MUTYH mutations. J Clin Oncol. 2009;27:3975-80. http://www.ncbi.nlm.nih.gov/pubmed/19620482
	19732775	Vogt S et al. Expanded extracolonic tumor spectrum in MUTYH-associated polyposis. Gastroenterology. 2009 Dec;137(6):1976-85.e1-10. http://www.ncbi.nlm.nih.gov/pubmed/19732775
	20512164	Aretz S and Hes FJ. Clinical utility gene card for: MUTYH-associated polyposis (MAP), autosomal recessive colorectal adenomatous polyposis. Eur J Hum Genet. 2010 Sep;18(9). http://www.ncbi.nlm.nih.gov/pubmed/20512164
	21061173	Win AK et al. Association between monoallelic MUTYH mutation and colorectal cancer risk: a meta-regression analysis. Familial Cancer. 2011 Mar 10(1):1-9. http://www.ncbi.nlm.nih.gov/pubmed/21061173
	21952991	Rennert G et al. MutYH mutation carriers have increased breast cancer risk. Cancer. 2012 Apr;118(8):1989-93. http://www.ncbi.nlm.nih.gov/pubmed/21952991
	22297469	Out AA et al. MUTYH gene variants and breast cancer in a Dutch case-control study. Breast Cancer Res Treat. 2012 Jul;134(1):219-27. http://www.ncbi.nlm.nih.gov/pubmed/22297469
	24444654	Win AK et al. Risk of colorectal cancer for carriers of mutations in MUTYH, with and without a family history of cancer. Gastroenterology. 2014 May;146(5):1208-11.e1-5. http://www.ncbi.nlm.nih.gov/pubmed/24444654
	24518836	Morak M et al. Biallelic MUTYH mutations can mimic Lynch syndrome. European Journal Of Human Genetics : Ejhg. 2014. http://www.ncbi.nlm.nih.gov/pubmed/24518836
	24953332	Castillejo A et al. Prevalence of germline MUTYH mutations among Lynch-like syndrome patients. European Journal Of Cancer (Oxford, England : 1990). 2014 Sep 50(13):2241-50. http://www.ncbi.nlm.nih.gov/pubmed/24953332
	27194394	Win AK et al. Risk of extracolonic cancers for people with biallelic and monoallelic mutations in MUTYH. International Journal Of Cancer. 2016 May 19. http://www.ncbi.nlm.nih.gov/pubmed/27194394
	28649662	Slavin TP et al. The contribution of pathogenic variants in breast cancer susceptibility genes to familial breast cancer risk. NPJ Breast Cancer. 2017 Jun 9;3:22. http://www.ncbi.nlm.nih.gov/pubmed/28649662
	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Colorectal. http://www.nccn.org/clinical.asp
MUTYH homoz	28649662	Slavin TP et al. The contribution of pathogenic variants in breast cancer susceptibility genes to familial breast cancer risk. NPJ Breast Cancer. 2017 Jun 9;3:22. http://www.ncbi.nlm.nih.gov/pubmed/28649662
MUTYH-single	28649662	Slavin TP et al. The contribution of pathogenic variants in breast cancer susceptibility genes to familial breast cancer risk. NPJ Breast Cancer. 2017 Jun 9;3:22. http://www.ncbi.nlm.nih.gov/pubmed/28649662

Gene	PMID	Reference Text	
NBN	15185344	Steffen J et al. Increased cancer risk of heterozygotes with NBS1 germline mutations in Poland. <i>Int J Cancer</i> . 2004 Aug 10;111(1):67-71. http://www.ncbi.nlm.nih.gov/pubmed/15185344	
	15578693	Buslov KG et al. NBS1 657del5 mutation may contribute only to a limited fraction of breast cancer cases in Russia. <i>Int J Cancer</i> . 2005 Apr 20;114(4):585-9. http://www.ncbi.nlm.nih.gov/pubmed/15578693	
	16770759	Steffen J et al. Germline mutations 657del5 of the NBS1 gene contribute significantly to the incidence of breast cancer in Central Poland. <i>Int J Cancer</i> . 2006 Jul 15;119(2):472-5. http://www.ncbi.nlm.nih.gov/pubmed/16770759	
	16998789	Steffen J et al. Increased risk of gastrointestinal lymphoma in carriers of the 657del5 NBS1 gene mutation. <i>Int J Cancer</i> . 2006 Dec 15;119(12):2970-3. http://www.ncbi.nlm.nih.gov/pubmed/16998789	
	17695489	Kanka C et al. Germline NBS1 mutations in families with aggregation of Breast and/or ovarian cancer from north-east Poland. <i>Anticancer Res</i> . 2007 Jul-Aug;27(4C):3015-8. http://www.ncbi.nlm.nih.gov/pubmed/17695489	
	18328813	di Masi A et al. The R215W mutation in NBS1 impairs gamma-H2AX binding and affects DNA repair: molecular bases for the severe phenotype of 657del5/R215W Nijmegen breakage syndrome patients. <i>Biochem Biophys Res Commun</i> . 2008 May 9;369(3):835-40. http://www.ncbi.nlm.nih.gov/pubmed/18328813	
	19452044	di Masi A and Antoccia A. NBS1 Heterozygosity and Cancer Risk. <i>Current Genomics</i> . 2008 Jun 9(4):275-81. http://www.ncbi.nlm.nih.gov/pubmed/19452044	
	22491912	Mateju M et al. Germline mutations 657del5 and 643C>T (R215W) in NBN are not likely to be associated with increased risk of breast cancer in Czech women. <i>Breast Cancer Res Treat</i> . 2012 Jun;133(2):809-11. http://www.ncbi.nlm.nih.gov/pubmed/22491912	
	23149842	Cybulski C et al. An inherited NBN mutation is associated with poor prognosis prostate cancer. <i>Br J Cancer</i> . 2013 Feb 5;108(2):461-8. http://www.ncbi.nlm.nih.gov/pubmed/23149842	
	23979977	He YZ et al. NBS1 Glu185Gln polymorphism and cancer risk: update on current evidence. <i>Tumour Biol</i> . 2014 Jan;35(1):675-87. http://www.ncbi.nlm.nih.gov/pubmed/23979977	
	24113799	Gao P et al. Functional variants in NBS1 and cancer risk: evidence from a meta-analysis of 60 publications with 111 individual studies. <i>Mutagenesis</i> . 2013 Nov;28(6):683-97. http://www.ncbi.nlm.nih.gov/pubmed/24113799	
	26250988	Aloraifi F et al. Protein-truncating variants in moderate-risk breast cancer susceptibility genes: A meta-analysis of high-risk case-control screening studies. <i>Cancer Genetics</i> . 2015 Sep 20(8):455-63. http://www.ncbi.nlm.nih.gov/pubmed/26250988	
	26822949	Lhota F et al. Hereditary truncating mutations of DNA repair and other genes in BRCA1/BRCA2/PALB2-negatively tested breast cancer patients. <i>Clin Genet</i> . 2016 Jan 29. http://www.ncbi.nlm.nih.gov/pubmed/26822949	
	27038244	Lener MR et al. Do founder mutations characteristic of some cancer sites also predispose to pancreatic cancer? <i>Int J Cancer</i> . 2016 Aug 1;139(3):601-6. http://www.ncbi.nlm.nih.gov/pubmed/27038244	
	27150568	Borecka M et al. The c.657del5 variant in the NBN gene predisposes to pancreatic cancer. <i>Gene</i> . 2016 Aug 10 587(2):169-72. http://www.ncbi.nlm.nih.gov/pubmed/27150568	
	28418444	Couch FJ et al. Associations Between Cancer Predisposition Testing Panel Genes and Breast Cancer. <i>JAMA Oncol</i> . 2017 Apr 13. http://www.ncbi.nlm.nih.gov/pubmed/28418444	
	28649662	Slavin TP et al. The contribution of pathogenic variants in breast cancer susceptibility genes to familial breast cancer risk. <i>NPJ Breast Cancer</i> . 2017 Jun 9;3:22. http://www.ncbi.nlm.nih.gov/pubmed/28649662	
	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Breast and Ovarian. http://www.nccn.org/clinical.asp	
	NF1	10524872	Walther MM et al. von Recklinghausen's disease and pheochromocytomas. <i>J Urol</i> . 1999 Nov;162(5):1582-6. http://www.ncbi.nlm.nih.gov/pubmed/10524872

Gene	PMID	Reference Text
NF1	12011145	Evans DG et al. Malignant peripheral nerve sheath tumours in neurofibromatosis 1. J Med Genet. 2002 May;39(5):311-4. http://www.ncbi.nlm.nih.gov/pubmed/12011145
	14759722	Wilson CH et al. Gynaecomastia, neurofibromatosis and breast cancer. Breast. 2004 Feb;13(1):77-9. http://www.ncbi.nlm.nih.gov/pubmed/14759722
	15655144	Lammert M et al. Prevalence of Neurofibromatosis 1 in German Children at Elementary School Enrollment. Arch Dermatol. 2005 Jan;141(1):71-4. http://www.ncbi.nlm.nih.gov/pubmed/15655144
	16330947	Miettinen M et al. Gastrointestinal stromal tumors in patients with neurofibromatosis 1: a clinicopathologic and molecular genetic study of 45 cases. Am J Surg Pathol. 2006;30:90-6. http://www.ncbi.nlm.nih.gov/pubmed/16330947
	16786042	Walker L et al. A prospective study of neurofibromatosis type 1 cancer incidence in the UK. Br J Cancer. 2006 Jul 17;95(2):233-8. http://www.ncbi.nlm.nih.gov/pubmed/16786042
	16790714	Bosch B et al. Clinical and genetic characteristics of patients with neurofibromatosis type 1 and pheochromocytoma. N Engl J Med. 2006 Jun 22;354(25):2729-31. http://www.ncbi.nlm.nih.gov/pubmed/16790714
	17105749	Ferner RE et al. Guidelines for the diagnosis and management of individuals with neurofibromatosis 1. J Med Genet. 2007 Feb;44(2):81-8. http://www.ncbi.nlm.nih.gov/pubmed/17105749
	17327286	McCaughan JA et al. Further evidence of the increased risk for malignant peripheral nerve sheath tumour from a Scottish cohort of patients with neurofibromatosis type 1. J Med Genet. 2007 Jul;44(7):463-6. http://www.ncbi.nlm.nih.gov/pubmed/17327286
	17369502	Sharif S et al. Women with neurofibromatosis 1 are at a moderately increased risk of developing breast cancer and should be considered for early screening. J Med Genet. 2007 Aug;44(8):481-4. http://www.ncbi.nlm.nih.gov/pubmed/17369502
	17514588	Zelinka T et al. Pheochromocytoma as a catecholamine producing tumor: implications for clinical practice. Stress. 2007 Jun;10(2):195-203. http://www.ncbi.nlm.nih.gov/pubmed/17514588
	18341492	Toumpanakis CG & Caplin ME. Molecular genetics of gastroenteropancreatic neuroendocrine tumors. The American Journal of Gastroenterology 2008 Mar; 103(3): 729-732. http://www.ncbi.nlm.nih.gov/pubmed/18341492
	20034338	Yoshimi A et al. Juvenile myelomonocytic leukemia: epidemiology, etiopathogenesis, diagnosis, and management considerations. Paediatr Drugs. 2010;12(1):11-21. http://www.ncbi.nlm.nih.gov/pubmed/20034338
	22965642	Wang X et al. Breast cancer and other neoplasms in women with neurofibromatosis type 1: a retrospective review of cases in the Detroit metropolitan area. Am J Med Genet A. 2012 Dec;158A(12):3061-4. http://www.ncbi.nlm.nih.gov/pubmed/22965642
	23036231	Evans DG et al. Malignant peripheral nerve sheath tumours in inherited disease. Clin Sarcoma Res. 2012 Oct 4;2(1):17. http://www.ncbi.nlm.nih.gov/pubmed/23036231
	23165953	Madanikia SA et al. Increased risk of breast cancer in women with NF1. Am J Med Genet A 2012; 158A: 3056-60. http://www.ncbi.nlm.nih.gov/pubmed/23165953
	23931823	Ferner RE & Gutmann DH. Neurofibromatosis type 1 (NF1): diagnosis and management. Handb Clin Neurol. 2013;115:939-55. http://www.ncbi.nlm.nih.gov/pubmed/23931823
	25742481	Seminog OO and Goldacre MJ. Age-specific risk of breast cancer in women with neurofibromatosis type 1. Br J Cancer 2015; 112: 1546-8. http://www.ncbi.nlm.nih.gov/pubmed/25742481
	26014596	Easton DF et al. Gene-Panel Sequencing and the Prediction of Breast-Cancer Risk. N Engl J Med. 2015 Jun 4;372(23):2243-57. http://www.ncbi.nlm.nih.gov/pubmed/26014596
	26511941	Nishida T et al. Gastrointestinal stromal tumors in Japanese patients with neurofibromatosis type I. J Gastroenterol. 2016 Jun; 51(6):571-8. http://www.ncbi.nlm.nih.gov/pubmed/26511941

Gene	PMID	Reference Text
NF1	26926675	Uusitalo E et al. Distinctive Cancer Associations in Patients With Neurofibromatosis Type 1. <i>J Clin Oncol</i> . 2016 Jun 10;34(17):1978-86. http://www.ncbi.nlm.nih.gov/pubmed/26926675
	27460956	Gruber LM et al. Pheochromocytoma and paraganglioma in patients with neurofibromatosis type 1. <i>Clin Endocrinol (Oxf)</i> . 2017 Jan;86(1):141-149. http://www.ncbi.nlm.nih.gov/pubmed/27460956
	7947106	Stillier CA et al. Neurofibromatosis and childhood leukaemia/lymphoma: a population-based UKCCSG study. <i>Br J Cancer</i> . 1994 Nov;70(5):969-72. http://www.ncbi.nlm.nih.gov/pubmed/7947106
	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Breast and Ovarian. http://www.nccn.org/clinical.asp
NF2	10569966	Evans DG et al. Paediatric presentation of type 2 neurofibromatosis. <i>Arch Dis Child</i> . 1999 Dec;81(6):496-9. http://www.ncbi.nlm.nih.gov/pubmed/10569966
	15190457	Baser ME et al. Genotype-phenotype correlations for nervous system tumors in neurofibromatosis 2: a population-based study. <i>Am J Hum Genet</i> . 2004 Aug;75(2):231-9. http://www.ncbi.nlm.nih.gov/pubmed/15190457
	15994874	Baser ME et al. The location of constitutional neurofibromatosis 2 (NF2) splice site mutations is associated with the severity of NF2. <i>J Med Genet</i> . 2005 Jul;42(7):540-6. http://www.ncbi.nlm.nih.gov/pubmed/15994874
	19476995	Asthagiri AR et al. Neurofibromatosis type 2. <i>Lancet</i> . 2009 Jun 6;373(9679):1974-86. http://www.ncbi.nlm.nih.gov/pubmed/19476995
	20082463	Evans DG et al. Birth incidence and prevalence of tumor-prone syndromes: estimates from a UK family genetic register service. <i>Am J Med Genet A</i> . 2010 Feb;152A(2):327-32. http://www.ncbi.nlm.nih.gov/pubmed/20082463
	21278391	Smith MJ et al. Cranial meningiomas in 411 neurofibromatosis type 2 (NF2) patients with proven gene mutations: clear positional effect of mutations, but absence of female severity effect on age at onset. <i>J Med Genet</i> . 2011 Apr;48(4):261-5. http://www.ncbi.nlm.nih.gov/pubmed/21278391
	22098617	Evans DG et al. Genetic testing and screening of individuals at risk of NF2. <i>Clin Genet</i> . 2012 Nov;82(5):416-24. http://www.ncbi.nlm.nih.gov/pubmed/22098617
	23931824	Lloyd SK et al. Neurofibromatosis type 2 (NF2): diagnosis and management. <i>Handb Clin Neurol</i> . 2013;115:957-67. http://www.ncbi.nlm.nih.gov/pubmed/23931824
	26043141	Slattery WH. Neurofibromatosis type 2. <i>Otolaryngol Clin North Am</i> . 2015 Jun;48(3):443-60. http://www.ncbi.nlm.nih.gov/pubmed/26043141
	26275417	Hexter A et al. Clinical and molecular predictors of mortality in neurofibromatosis 2: a UK national analysis of 1192 patients. <i>J Med Genet</i> . 2015 Oct;52(10):699-705. http://www.ncbi.nlm.nih.gov/pubmed/26275417
	26564072	Evans DG. Neurofibromatosis type 2. <i>Handb Clin Neurol</i> . 2015;132:87-96. http://www.ncbi.nlm.nih.gov/pubmed/26564072
	27655473	Ardern-Holmes S et al. Neurofibromatosis Type 2. <i>J Child Neurol</i> . 2017 Jan;32(1):9-22. http://www.ncbi.nlm.nih.gov/pubmed/27655473
	28620005	Evans DGR et al. Cancer and Central Nervous System Tumor Surveillance in Pediatric Neurofibromatosis 2 and Related Disorders. <i>Clin Cancer Res</i> . 2017 Jun 15;23(12):e54-e61. http://www.ncbi.nlm.nih.gov/pubmed/28620005
	28848060	Halliday D et al. Genetic Severity Score predicts clinical phenotype in NF2. <i>J Med Genet</i> . 2017 Oct;54(10):657-664. http://www.ncbi.nlm.nih.gov/pubmed/28848060
	9643284	Evans DG et al. Genotype/phenotype correlations in type 2 neurofibromatosis (NF2): evidence for more severe disease associated with truncating mutations. <i>J Med Genet</i> . 1998 Jun;35(6):450-5. http://www.ncbi.nlm.nih.gov/pubmed/9643284
	NTHL1	25938944

Gene	PMID	Reference Text
NTHL1	26559593	Rivera B et al. Biallelic NTHL1 Mutations in a Woman with Multiple Primary Tumors. The New England Journal Of Medicine. 2015 Nov 12 373(20):1985-6. http://www.ncbi.nlm.nih.gov/pubmed/26559593
	27713038	Broderick P et al. Validation of Recently Proposed Colorectal Cancer Susceptibility Gene Variants in an Analysis of Families and Patients-a Systematic Review. Gastroenterology. 2016 Oct 3. http://www.ncbi.nlm.nih.gov/pubmed/27713038
	27720914	Belhadj S et al. Delineating the phenotypic spectrum of the NTHL1-associated polyposis. Clinical Gastroenterology And Hepatology : The Official Clinical Practice Journal Of The American Gastroenterological Association. 2016 Oct 5. http://www.ncbi.nlm.nih.gov/pubmed/27720914
	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Colorectal. http://www.nccn.org/clinical.asp
PALB2	17200668	Rahman N et al. PALB2, which encodes a BRCA2-interacting protein, is a breast cancer susceptibility gene. Nat Genet. 2007 Feb;39(2):165-7. http://www.ncbi.nlm.nih.gov/pubmed/17200668
	17200671	Reid S et al. Biallelic mutations in PALB2 cause Fanconi anemia subtype FA-N and predispose to childhood cancer. Nat Genet. 2007 Feb;39(2):162-4. http://www.ncbi.nlm.nih.gov/pubmed/17200671
	17287723	Erkko H et al. A recurrent mutation in PALB2 in Finnish cancer families. Nature. 2007 Mar 15;446(7133):316-9. http://www.ncbi.nlm.nih.gov/pubmed/17287723
	19264984	Jones S et al. Exomic sequencing identifies PALB2 as a pancreatic cancer susceptibility gene. Science. 2009 Apr 10;324(5924):217. http://www.ncbi.nlm.nih.gov/pubmed/19264984
	20412113	Slater EP et al. PALB2 mutations in European familial pancreatic cancer families. Clin Genet. 2010 Nov;78(5):490-4. http://www.ncbi.nlm.nih.gov/pubmed/20412113
	20927582	Ding Y et al. Mutations in BRCA2 and PALB2 in male breast cancer cases from the United States. Breast Cancer Res Treat. 2011 Apr;126(3):771-8 http://www.ncbi.nlm.nih.gov/pubmed/20927582
	21285249	Casadei S et al. Contribution of inherited mutations in the BRCA2-interacting protein PALB2 to familial breast cancer. Cancer Res. 2011 Mar 15;71(6):2222-9. http://www.ncbi.nlm.nih.gov/pubmed/21285249
	23135763	Canto MI et al. International Cancer of the Pancreas Screening (CAPS) Consortium summit on the management of patients with increased risk for familial pancreatic cancer. Gut. 2013 Mar;62(3):339-47. http://www.ncbi.nlm.nih.gov/pubmed/23135763
	23935836	Blanco A et al. Analysis of PALB2 gene in BRCA1/BRCA2 negative Spanish hereditary breast/ovarian cancer families with pancreatic cancer cases. PLoS One. 2013 Jul 23;8(7):e67538. http://www.ncbi.nlm.nih.gov/pubmed/23935836
	24240112	Pennington KP et al. Germline and somatic mutations in homologous recombination genes predict platinum response and survival in ovarian, fallopian tube, and peritoneal carcinomas. Clin Cancer Res. 2014 Feb 1;20(3):764-75. http://www.ncbi.nlm.nih.gov/pubmed/24240112
	24556621	Leongamornlert D et al. Frequent germline deleterious mutations in DNA repair genes in familial prostate cancer cases are associated with advanced disease. Br J Cancer. 2014 Mar 18;110(6):1663-72. http://www.ncbi.nlm.nih.gov/pubmed/24556621
	25099575	Antoniou A et al. Breast-cancer risk in families with mutations in PALB2. N Engl J Med. 2014 Aug 7;371(6):497-506. http://www.ncbi.nlm.nih.gov/pubmed/25099575
	25186627	Tung N et al. Frequency of mutations in individuals with breast cancer referred for BRCA1 and BRCA2 testing using next-generation sequencing with a 25-gene panel. Cancer. 2015 Jan 1;121(1):25-33. http://www.ncbi.nlm.nih.gov/pubmed/25186627
	25356972	Zhen DB et al. BRCA1, BRCA2, PALB2, and CDKN2A mutations in familial pancreatic cancer: a PACGENE study. Genet Med. 2015 Jul;17(7):569-77. http://www.ncbi.nlm.nih.gov/pubmed/25356972
	25959805	Cybulski C et al. Clinical outcomes in women with breast cancer and a PALB2 mutation: a prospective cohort analysis. Lancet Oncol. 2015 Jun;16(6):638-44. http://www.ncbi.nlm.nih.gov/pubmed/25959805

Gene	PMID	Reference Text
PALB2	26315354	Ramus SJ et al. Germline Mutations in the BRIP1, BARD1, PALB2, and NBN Genes in Women With Ovarian Cancer. J Natl Cancer Inst. 2015 Aug 27;107(11). http://www.ncbi.nlm.nih.gov/pubmed/26315354
	26440929	Salo-Mullen EE et al. Identification of germline genetic mutations in patients with pancreatic cancer. Cancer. 2015 Dec 15;121(24):4382-8. http://www.ncbi.nlm.nih.gov/pubmed/26440929
	26786923	Thompson ER et al. Panel Testing for Familial Breast Cancer: Calibrating the Tension Between Research and Clinical Care. Journal Of Clinical Oncology : Official Journal Of The American Society Of Clinical Oncology. 2016 Jan 19. http://www.ncbi.nlm.nih.gov/pubmed/26786923
	27038244	Lener MR et al. Do founder mutations characteristic of some cancer sites also predispose to pancreatic cancer? Int J Cancer. 2016 Aug 1;139(3):601-6. http://www.ncbi.nlm.nih.gov/pubmed/27038244
	27099641	Southey MC et al. PALB2: research reaching to clinical outcomes for women with breast cancer. Hereditary Cancer In Clinical Practice. 2016 14:9. http://www.ncbi.nlm.nih.gov/pubmed/27099641
	27433846	Pritchard CC et al. Inherited DNA-Repair Gene Mutations in Men with Metastatic Prostate Cancer. N Engl J Med. 2016 Aug 4;375(5):443-53. http://www.ncbi.nlm.nih.gov/pubmed/27433846
	27648926	Silvestri V et al. Whole-exome sequencing and targeted gene sequencing provide insights into the role of PALB2 as a male breast cancer susceptibility gene. Cancer. 2017 Jan 01 123(2):210-218. http://www.ncbi.nlm.nih.gov/pubmed/27648926
	28008555	Pritzlaff M et al. Male breast cancer in a multi-gene panel testing cohort: insights and unexpected results. Breast Cancer Res Treat. 2017 Feb;161(3):575-586. http://www.ncbi.nlm.nih.gov/pubmed/28008555
	28259476	Annala M et al. Treatment Outcomes and Tumor Loss of Heterozygosity in Germline DNA Repair-deficient Prostate Cancer. Eur Urol. 2017 Mar 1. [Epub ahead of print] http://www.ncbi.nlm.nih.gov/pubmed/28259476
	28418444	Couch FJ et al. Associations Between Cancer Predisposition Testing Panel Genes and Breast Cancer. JAMA Oncol. 2017 Apr 13. http://www.ncbi.nlm.nih.gov/pubmed/28418444
	28649662	Slavin TP et al. The contribution of pathogenic variants in breast cancer susceptibility genes to familial breast cancer risk. NPJ Breast Cancer. 2017 Jun 9;3:22. http://www.ncbi.nlm.nih.gov/pubmed/28649662
	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Breast and Ovarian. http://www.nccn.org/clinical.asp
	PALLD	17194196
17415588		Zogopoulos G et al. The P239S palladin variant does not account for a significant fraction of hereditary or early onset pancreas cancer. Hum Genet. 2007 Jun;121(5):635-7. http://www.ncbi.nlm.nih.gov/pubmed/17415588
19336541		Klein AP et al. Absence of deleterious palladin mutations in patients with familial pancreatic cancer. Cancer Epidemiol Biomarkers Prev. 2009 Apr;18(4):1328-30. http://www.ncbi.nlm.nih.gov/pubmed/19336541
PHOX2B	14608649	Weese-Mayer DE et al. Idiopathic congenital central hypoventilation syndrome: analysis of genes pertinent to early autonomic nervous system embryologic development and identification of mutations in PHOX2b. Am J Med Genet A. 2003 Dec 15;123A(3):267-78.14608649 http://www.ncbi.nlm.nih.gov/pubmed/14608649
	15024693	Trochet D et al. Germline mutations of the paired-like homeobox 2B (PHOX2B) gene in neuroblastoma. Am J Hum Genet. 2004 Apr;74(4):761-4.15024693 http://www.ncbi.nlm.nih.gov/pubmed/15024693
	15516980	van Limpt et al. The Phox2B homeobox gene is mutated in sporadic neuroblastomas. Oncogene. 2004 Dec 9;23(57):9280-8.15516980 http://www.ncbi.nlm.nih.gov/pubmed/15516980

Gene	PMID	Reference Text
PHOX2B	15657873	Trochet D et al. PHOX2B genotype allows for prediction of tumor risk in congenital central hypoventilation syndrome. <i>Am J Hum Genet.</i> 2005 Mar;76(3):421-6.15657873 http://www.ncbi.nlm.nih.gov/pubmed/15657873
	16691592	McConville C et al. PHOX2B analysis in non-syndromic neuroblastoma cases shows novel mutations and genotype-phenotype associations. <i>Am J Med Genet A.</i> 2006 Jun 15;140(12):1297-301.16691592 http://www.ncbi.nlm.nih.gov/pubmed/16691592
	16888290	Berry-Kravis EM et al. Congenital central hypoventilation syndrome: PHOX2B mutations and phenotype. <i>Am J Respir Crit Care Med.</i> 2006 Nov 15;174(10):1139-44.16888290 http://www.ncbi.nlm.nih.gov/pubmed/16888290
	19058226	Trochet D et al. In Vitro studies of non poly alanine PHOX2B mutations argue against a loss-of-function mechanism for congenital central hypoventilation. <i>Hum Mutat.</i> 2009 Feb;30(2):E421-31. 19058226 http://www.ncbi.nlm.nih.gov/pubmed/19058226
	20208042	Weese-Mayer DE et al. An official ATS clinical policy statement: Congenital central hypoventilation syndrome: genetic basis, diagnosis, and management. <i>Am J Respir Crit Care Med.</i> 2010 Mar 15;181(6):626-44. 20208042 http://www.ncbi.nlm.nih.gov/pubmed/20208042
	20301600	Weese-Mayer DE, Marazita ML, Rand CM, et al. Congenital Central Hypoventilation Syndrome. 2004 Jan 28 [Updated 2014 Jan 30]. In: Pagon RA, Adam MP, Ardinger HH, et al., editors. <i>GeneReviews</i> ® [Internet]. Seattle (WA): University of Washington, Seattle; 1993-2016. http://www.ncbi.nlm.nih.gov/pubmed/20301600
	21830319	Jennings LJ et al. Variable human phenotype associated with novel deletions of the PHOX2B gene. <i>Pediatr Pulmonol.</i> 2012 Feb;47(2):153-61. http://www.ncbi.nlm.nih.gov/pubmed/21830319
	26011159	Armstrong AE et al. Treatment of neuroblastoma in congenital central hypoventilation syndrome with a PHOX2B polyalanine repeat expansion mutation: New twist on a neurocristopathy syndrome. <i>Pediatric Blood & Cancer.</i> 2015 Nov 62(11):2007-10. http://www.ncbi.nlm.nih.gov/pubmed/26011159
	26375764	Heide S et al. Oncologic Phenotype of Peripheral Neuroblastic Tumors Associated With PHOX2B Non-Polyalanine Repeat Expansion Mutations. <i>Pediatr Blood Cancer.</i> 2016 Jan;63(1):71-7. 26375764 http://www.ncbi.nlm.nih.gov/pubmed/26375764
	27485184	Klaskova E et al. Significant phenotype variability of congenital central hypoventilation syndrome in a family with polyalanine expansion mutation of the PHOX2B gene. <i>Biomedical Papers Of The Medical Faculty Of The University Palacky, Olomouc, Czechoslovakia.</i> 2016 Aug 2. http://www.ncbi.nlm.nih.gov/pubmed/27485184
	28633714	Kasi AS et al. Three-Generation Family With Congenital Central Hypoventilation Syndrome and Novel PHOX2B Gene Non-Polyalanine Repeat Mutation. <i>Journal Of Clinical Sleep Medicine : Jcsm : Official Publication Of The American Academy Of Sleep Medicine.</i> 2017 Jul 15 13(7):925-927. http://www.ncbi.nlm.nih.gov/pubmed/28633714
	PMS2	18602922
19125127		Palomaki GE et al. EGAPP supplementary evidence review: DNA testing strategies aimed at reducing morbidity and mortality from Lynch syndrome. <i>Genet Med.</i> 2009 Jan;11(1):42-65. http://www.ncbi.nlm.nih.gov/pubmed/19125127
20442441		Wimmer K and Kratz CP. Constitutional mismatch repair-deficiency syndrome. <i>Haematologica.</i> 2010 May; 95(5): 699–701. http://www.ncbi.nlm.nih.gov/pubmed/20442441
20531397		Durno CA et al. The gastrointestinal phenotype of germline biallelic mismatch repair gene mutations. <i>Am J Gastroenterol.</i> 2010 Nov;105(11):2449-56. http://www.ncbi.nlm.nih.gov/pubmed/20531397
23385444		Win AK et al. Risks of colorectal and other cancers after endometrial cancer for women with Lynch syndrome. <i>J Natl Cancer Inst.</i> 2013 Feb;105(4):274-9. http://www.ncbi.nlm.nih.gov/pubmed/23385444
24425144		Ryan S et al. Risk of prostate cancer in Lynch syndrome: a systematic review and meta-analysis. <i>Cancer Epidemiology, Biomarkers & Prevention : A Publication Of The American Association For Cancer Research, Cosponsored By The American Society Of Preventive Oncology.</i> 2014 Mar 23(3):437-49. http://www.ncbi.nlm.nih.gov/pubmed/24425144

Gene	PMID	Reference Text
PMS2	24440087	Bakry D et al. Genetic and clinical determinants of constitutional mismatch repair deficiency syndrome: report from the constitutional mismatch repair deficiency consortium. <i>Eur J Cancer</i> . 2014 Mar;50(5):987-96. http://www.ncbi.nlm.nih.gov/pubmed/24440087
	24737826	Wimmer K et al. Diagnostic criteria for constitutional mismatch repair deficiency syndrome: suggestions of the European consortium 'care for CMMRD' (C4CMMRD). <i>Journal Of Medical Genetics</i> . 2014 Jun 51(6):355-65. http://www.ncbi.nlm.nih.gov/pubmed/24737826
	25512458	Broeke SW et al. Lynch Syndrome Caused by Germline PMS2 Mutations: Delineating the Cancer Risk. <i>Journal of clinical oncology : official journal of the American Society of Clinical Oncology</i> . 2015 Feb 1 33(4):319-25 http://www.ncbi.nlm.nih.gov/pubmed/25512458
	27241104	Mills AM and Longacre TA2. Lynch Syndrome: Female Genital Tract Cancer Diagnosis and Screening. <i>Surg Pathol Clin</i> . 2016 Jun;9(2):201-14. http://www.ncbi.nlm.nih.gov/pubmed/27241104
	29345684	Roberts ME et al. MSH6 and PMS2 germ-line pathogenic variants implicated in Lynch syndrome are associated with breast cancer. <i>Genetics In Medicine : Official Journal Of The American College Of Medical Genetics</i> . 2018 Jan 18. http://www.ncbi.nlm.nih.gov/pubmed/29345684
	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Colorectal. http://www.nccn.org/clinical.asp
POLD1	23263490	Palles C et al. Germline mutations affecting the proofreading domains of POLE and POLD1 predispose to colorectal adenomas and carcinomas. <i>Nat Genet</i> . 2013 Feb;45(2):136-44. http://www.ncbi.nlm.nih.gov/pubmed/23263490
	24861832	Henninger EE et al. DNA polymerase e and its roles in genome stability. <i>IUBMB Life</i> . 2014 May;66(5):339-51. http://www.ncbi.nlm.nih.gov/pubmed/24861832
	26133394	Bellido F et al. POLE and POLD1 mutations in 529 kindred with familial colorectal cancer and/or polyposis: review of reported cases and recommendations for genetic testing and surveillance. <i>Genet Med</i> . 2015 Jul 2. http://www.ncbi.nlm.nih.gov/pubmed/26133394
	26344056	Arora S et al. Genetic Variants That Predispose to DNA Double-Strand Breaks in Lymphocytes From a Subset of Patients With Familial Colorectal Carcinomas. <i>Gastroenterology</i> . 2015 Dec;149(7):1872-1883.e9. http://www.ncbi.nlm.nih.gov/pubmed/26344056
	28306219	Ferrer-Avargues R et al. Characterization of a novel POLD1 missense founder mutation in a Spanish population. <i>J Gene Med</i> . 2017 Apr;19(4). http://www.ncbi.nlm.nih.gov/pubmed/28306219
	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Colorectal. http://www.nccn.org/clinical.asp
POLE	23230001	Pachlopnik Schmid J et al. Polymerase e1 mutation in a human syndrome with facial dysmorphism, immunodeficiency, livedo, and short stature ("FILS syndrome"). <i>J Exp Med</i> . 2012 Dec 17;209(13):2323-30. http://www.ncbi.nlm.nih.gov/pubmed/23230001
	23263490	Palles C et al. Germline mutations affecting the proofreading domains of POLE and POLD1 predispose to colorectal adenomas and carcinomas. <i>Nat Genet</i> . 2013 Feb;45(2):136-44. http://www.ncbi.nlm.nih.gov/pubmed/23263490
	24501277	Valle L et al. New insights into POLE and POLD1 germline mutations in familial colorectal cancer and polyposis. <i>Hum Mol Genet</i> . 2014 Jul 1;23(13):3506-12. http://www.ncbi.nlm.nih.gov/pubmed/24501277
	24788313	Rohlin A et al. A mutation in POLE predisposing to a multi-tumour phenotype. <i>Int J Oncol</i> . 2014 Jul;45(1):77-81. http://www.ncbi.nlm.nih.gov/pubmed/24788313
	24861832	Henninger EE et al. DNA polymerase e and its roles in genome stability. <i>IUBMB Life</i> . 2014 May;66(5):339-51. http://www.ncbi.nlm.nih.gov/pubmed/24861832
	25370038	Elsayed FA et al. Germline variants in POLE are associated with early onset mismatch repair deficient colorectal cancer. <i>Eur J Hum Genet</i> . 2015 Aug;23(8):1080-4. http://www.ncbi.nlm.nih.gov/pubmed/25370038
	25529843	Spier I et al. Frequency and phenotypic spectrum of germline mutations in POLE and seven other polymerase genes in 266 patients with colorectal adenomas and carcinomas. <i>Int J Cancer</i> . 2015 Jul 15;137(2):320-31 http://www.ncbi.nlm.nih.gov/pubmed/25529843

Gene	PMID	Reference Text
POLE	25860647	Hansen MF et al. A novel POLE mutation associated with cancers of colon, pancreas, ovaries and small intestine. <i>Fam Cancer</i> . 2015 Sep;14(3):437-48. http://www.ncbi.nlm.nih.gov/pubmed/25860647
	26133394	Bellido F et al. POLE and POLD1 mutations in 529 kindred with familial colorectal cancer and/or polyposis: review of reported cases and recommendations for genetic testing and surveillance. <i>Genet Med</i> . 2015 Jul 2. http://www.ncbi.nlm.nih.gov/pubmed/26133394
	26493165	Rohlin A et al. GREM1 and POLE variants in hereditary colorectal cancer syndromes. <i>Genes, Chromosomes & Cancer</i> . 2016 Jan 55(1):95-106. http://www.ncbi.nlm.nih.gov/pubmed/26493165
	27573199	Wimmer K et al. A novel germline POLE mutation causes an early onset cancer prone syndrome mimicking constitutional mismatch repair deficiency. <i>Fam Cancer</i> . 2017 Jan;16(1):67-71. http://www.ncbi.nlm.nih.gov/pubmed/27573199
	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Colorectal. http://www.nccn.org/clinical.asp
	POT1	24686846
24686849		Robles-Espinoza CD et al. POT1 loss-of-function variants predispose to familial melanoma. <i>Nature Genetics</i> . 2014 May 46(5):478-81. http://www.ncbi.nlm.nih.gov/pubmed/24686849
25482530		Bainbridge MN et al. Germline mutations in shelterin complex genes are associated with familial glioma. <i>Journal Of The National Cancer Institute</i> . 2015 Jan 107(1):384. http://www.ncbi.nlm.nih.gov/pubmed/25482530
26403419		Calvete O et al. A mutation in the POT1 gene is responsible for cardiac angiosarcoma in TP53-negative Li-Fraumeni-like families. <i>Nature Communications</i> . 2015 6:8383. http://www.ncbi.nlm.nih.gov/pubmed/26403419
27528712		Speedy HE et al. Germline mutations in shelterin complex genes are associated with familial chronic lymphocytic leukemia. <i>Blood</i> . 2016 Aug 15. http://www.ncbi.nlm.nih.gov/pubmed/27528712
28283772		Leachman SA et al. Identification, genetic testing, and management of hereditary melanoma. <i>Cancer Metastasis Rev</i> . 2017 Mar;36(1):77-90. http://www.ncbi.nlm.nih.gov/pubmed/28283772
28389767		Wilson TL et al. A new POT1 germline mutation-expanding the spectrum of POT1-associated cancers. <i>Familial Cancer</i> . 2017 Apr 07. http://www.ncbi.nlm.nih.gov/pubmed/28389767
PRKAR1A		11549623
	19293268	Bertherat J et al. Mutations in regulatory subunit type 1A of cyclic adenosine 5'-monophosphate-dependent protein kinase (PRKAR1A): phenotype analysis in 353 patients and 80 different genotypes. <i>The Journal Of Clinical Endocrinology And Metabolism</i> . 2009 94(6):2085-91. http://www.ncbi.nlm.nih.gov/pubmed/19293268
	20301463	Stratakis CA, Salpea P, Raygada M. Carney Complex. 2003 Feb 5 [Updated 2015 Jan 29]. In: Pagon RA, Adam MP, Ardinger HH, et al., editors. <i>GeneReviews</i> ® [Internet]. Seattle (WA): University of Washington, Seattle; 1993-2016. Available from: https://www.ncbi.nlm.nih.gov/books/NBK1286/ http://www.ncbi.nlm.nih.gov/pubmed/20301463
	20833331	Rothenbuhler A and Stratakis CA. Clinical and molecular genetics of Carney complex. <i>Best Practice & Research. Clinical Endocrinology & Metabolism</i> . 2010 Jun 24(3):389-99. http://www.ncbi.nlm.nih.gov/pubmed/20833331
	21651393	Linglart A et al. Recurrent PRKAR1A mutation in acrodysostosis with hormone resistance. <i>The New England Journal Of Medicine</i> . 2011 364(23):2218-26. http://www.ncbi.nlm.nih.gov/pubmed/21651393
	25592387	Schernthaner-Reiter MH et al. MEN1, MEN4, and Carney Complex: Pathology and Molecular Genetics. <i>Neuroendocrinology</i> . 2015 Jan 9. http://www.ncbi.nlm.nih.gov/pubmed/25592387

Gene	PMID	Reference Text
PRKAR1A	26130139	Correa R et al. Carney complex: an update. <i>Eur J Endocrinol.</i> 2015 Oct;173(4):M85-97.
		http://www.ncbi.nlm.nih.gov/pubmed/26130139
PRSS1	15017610	Howes N et al. Clinical and genetic characteristics of hereditary pancreatitis in Europe. <i>Clinical Gastroenterology And Hepatology : The Official Clinical Practice Journal Of The American Gastroenterological Association.</i> 2004 Mar 2(3):252-61.
		http://www.ncbi.nlm.nih.gov/pubmed/15017610
	18184119	Rebours V et al. Risk of pancreatic adenocarcinoma in patients with hereditary pancreatitis: a national exhaustive series. <i>The American Journal Of Gastroenterology.</i> 2008 Jan 103(1):111-9.
		http://www.ncbi.nlm.nih.gov/pubmed/18184119
	19951905	Grocock CJ et al. The variable phenotype of the p.A16V mutation of cationic trypsinogen (PRSS1) in pancreatitis families. <i>Gut.</i> 2010 Mar 59(3):357-63.
		http://www.ncbi.nlm.nih.gov/pubmed/19951905
	21844754	LaRusch J and Whitcomb DC. Genetics of pancreatitis. <i>Current Opinion In Gastroenterology.</i> 2011 Sep 27(5):467-74.
		http://www.ncbi.nlm.nih.gov/pubmed/21844754
	22539344	Szabó A and Sahin-Tóth M. Increased activation of hereditary pancreatitis-associated human cationic trypsinogen mutants in presence of chymotrypsin C. <i>The Journal Of Biological Chemistry.</i> 2012 Jun 8 287(24):20701-10.
		http://www.ncbi.nlm.nih.gov/pubmed/22539344
	23686146	Masamune A et al. PRSS1 c.623G>C (p.G208A) variant is associated with pancreatitis in Japan. <i>Gut.</i> 2014 Feb 63(2):366.
http://www.ncbi.nlm.nih.gov/pubmed/23686146		
24458023	Németh BC and Sahin-Tóth M. Human cationic trypsinogen (PRSS1) variants and chronic pancreatitis. <i>American Journal Of Physiology. Gastrointestinal And Liver Physiology.</i> 2014 Mar 306(6):G466-73.	
	http://www.ncbi.nlm.nih.gov/pubmed/24458023	
24624459	LaRusch J, Solomon S, Whitcomb DC. Pancreatitis Overview. 2014 Mar 13. In: Adam MP, Ardinger HH, Pagon RA, et al., editors. <i>GeneReviews® [Internet].</i> Seattle (WA): University of Washington, Seattle; 1993-2017.	
	http://www.ncbi.nlm.nih.gov/pubmed/24624459	
9091646	Lowenfels AB et al. Hereditary pancreatitis and the risk of pancreatic cancer. International Hereditary Pancreatitis Study Group. <i>Journal Of The National Cancer Institute.</i> 1997 Mar 19 89(6):442-6.	
	http://www.ncbi.nlm.nih.gov/pubmed/9091646	
PTCH1	15545745	Kimonis VE et al. Radiological features in 82 patients with nevoid basal cell carcinoma (NBCC or Gorlin) syndrome. <i>Genet Med.</i> 2004 Nov-Dec;6(6):495-502.
		http://www.ncbi.nlm.nih.gov/pubmed/15545745
	15545751	Gorlin RJ. Nevoid basal cell carcinoma (Gorlin) syndrome. <i>Genet Med.</i> 2004 Nov-Dec;6(6):530-9.
		http://www.ncbi.nlm.nih.gov/pubmed/15545751
	16088933	Marsh A et al. DHPLC analysis of patients with Nevoid Basal Cell Carcinoma Syndrome reveals novel PTCH missense mutations in the sterol-sensing domain. <i>Hum Mutat.</i> 2005 Sep;26(3):283.
		http://www.ncbi.nlm.nih.gov/pubmed/16088933
	16301862	Klein RD et al. Clinical testing for the nevoid basal cell carcinoma syndrome in a DNA diagnostic laboratory. <i>Genetics In Medicine : Official Journal Of The American College Of Medical Genetics.</i> 2005 7(9):611-9.
		http://www.ncbi.nlm.nih.gov/pubmed/16301862
	16909134	Soufir N et al. PTCH mutations and deletions in patients with typical nevoid basal cell carcinoma syndrome and in patients with a suspected genetic predisposition to basal cell carcinoma: a French study. <i>Br J Cancer.</i> 2006 Aug 21;95(4):548-53.
		http://www.ncbi.nlm.nih.gov/pubmed/16909134
	19276247	Garrè ML et al. Medulloblastoma variants: age-dependent occurrence and relation to Gorlin syndrome--a new clinical perspective. <i>Clin Cancer Res.</i> 2009 Apr 1;15(7):2463-71.
http://www.ncbi.nlm.nih.gov/pubmed/19276247		
20082463	Evans DG et al. Birth incidence and prevalence of tumor-prone syndromes: estimates from a UK family genetic register service. <i>Am J Med Genet A.</i> 2010 Feb;152A(2):327-32.	
	http://www.ncbi.nlm.nih.gov/pubmed/20082463	
20301330	Evans DG, Farndon PA. Nevoid Basal Cell Carcinoma Syndrome. 2002 Jun 20 [Updated 2015 Oct 1]. In: Pagon RA, Adam MP, Ardinger HH, et al., editors. <i>GeneReviews® [Internet].</i> Seattle (WA): University of Washington, Seattle; 1993-2016.	
	http://www.ncbi.nlm.nih.gov/pubmed/20301330	
21152126	Jones EA et al. Basal cell carcinomas in gorlin syndrome: a review of 202 patients. <i>J Skin Cancer.</i> 2011;2011:217378.	
	http://www.ncbi.nlm.nih.gov/pubmed/21152126	

Gene	PMID	Reference Text
<i>PTCH1</i>	21210781	Nagao K et al. Entire <i>PTCH1</i> deletion is a common event in point mutation-negative cases with nevoid basal cell carcinoma syndrome in Japan. <i>Clin Genet.</i> 2011 Feb;79(2):196-8. http://www.ncbi.nlm.nih.gov/pubmed/21210781
	21834049	Bree AF et al. Consensus statement from the first international colloquium on basal cell nevus syndrome (BCNS). <i>Am J Med Genet A.</i> 2011 Sep;155A(9):2091-7. http://www.ncbi.nlm.nih.gov/pubmed/21834049
	22918513	Kimonis VE et al. Clinical and radiological features in young individuals with nevoid basal cell carcinoma syndrome. <i>Genet Med.</i> 2013 Jan;15(1):79-83. http://www.ncbi.nlm.nih.gov/pubmed/22918513
	25131638	Fujii K et al. Gorlin syndrome (nevoid basal cell carcinoma syndrome): update and literature review. <i>Pediatr Int.</i> 2014 Oct;56(5):667-74. http://www.ncbi.nlm.nih.gov/pubmed/25131638
	28620006	Foulkes WD et al. Cancer Surveillance in Gorlin Syndrome and Rhabdoid Tumor Predisposition Syndrome. <i>Clin Cancer Res.</i> 2017 Jun 15;23(12):e62-e67. http://www.ncbi.nlm.nih.gov/pubmed/28620006
	8326488	Evans DG et al. Complications of the naevoid basal cell carcinoma syndrome: results of a population based study. <i>J Med Genet.</i> 1993 Jun;30(6):460-4. http://www.ncbi.nlm.nih.gov/pubmed/8326488
	9096761	Kimonis VE et al. Clinical manifestations in 105 persons with nevoid basal cell carcinoma syndrome. <i>Am J Med Genet.</i> 1997 Mar 31;69(3):299-308. http://www.ncbi.nlm.nih.gov/pubmed/9096761
	9231911	Cowan R et al. The gene for the naevoid basal cell carcinoma syndrome acts as a tumour-suppressor gene in medulloblastoma. <i>Br J Cancer.</i> 1997;76(2):141-5. http://www.ncbi.nlm.nih.gov/pubmed/9231911
	9415689	Wicking C et al. De novo mutations of the Patched gene in nevoid basal cell carcinoma syndrome help to define the clinical phenotype. <i>Am J Med Genet.</i> 1997 Dec 19;73(3):304-7. http://www.ncbi.nlm.nih.gov/pubmed/9415689
<i>PTEN</i>	17526800	Lachlan KL et al. Cowden syndrome and Bannayan Riley Ruvalcaba syndrome represent one condition with variable expression and age-related penetrance: results of a clinical study of <i>PTEN</i> mutation carriers. <i>J Med Genet.</i> 2007 Sep;44(9):579-85. http://www.ncbi.nlm.nih.gov/pubmed/17526800
	17526801	Tan WH et al. The spectrum of vascular anomalies in patients with <i>PTEN</i> mutations: implications for diagnosis and management. <i>J Med Genet.</i> 2007 Sep;44(9):594-602. http://www.ncbi.nlm.nih.gov/pubmed/17526801
	18160807	Devi M et al. Testicular mixed germ cell tumor in an adolescent with cowden disease. <i>Oncology.</i> 2007;72(3-4):194-6. http://www.ncbi.nlm.nih.gov/pubmed/18160807
	18594467	Cho MY et al. First report of ovarian dysgerminoma in Cowden syndrome with germline <i>PTEN</i> mutation and <i>PTEN</i> -related 10q loss of tumor heterozygosity. <i>Am J Surg Pathol.</i> 2008 Aug;32(8):1258-64. http://www.ncbi.nlm.nih.gov/pubmed/18594467
	19265751	Varga EA et al. The prevalence of <i>PTEN</i> mutations in a clinical pediatric cohort with autism spectrum disorders, developmental delay, and macrocephaly. <i>Genet Med.</i> 2009 Feb;11(2):111-7. http://www.ncbi.nlm.nih.gov/pubmed/19265751
	20600018	Heald B et al. Frequent gastrointestinal polyps and colorectal adenocarcinomas in a prospective series of <i>PTEN</i> mutation carriers. <i>Gastroenterology.</i> 2010 Dec;139(6):1927-33. http://www.ncbi.nlm.nih.gov/pubmed/20600018
	20962022	Smith JR et al. Thyroid nodules and cancer in children with <i>PTEN</i> hamartoma tumor syndrome. <i>J Clin Endocrinol Metab.</i> 2011 Jan;96(1):34-7. http://www.ncbi.nlm.nih.gov/pubmed/20962022
	21194675	Tan MH et al. A clinical scoring system for selection of patients for <i>PTEN</i> mutation testing is proposed on the basis of a prospective study of 3042 probands. <i>Am J Hum Genet.</i> 2011 Jan 7;88(1):42-56. http://www.ncbi.nlm.nih.gov/pubmed/21194675
	21343951	Mester JL et al. Analysis of prevalence and degree of macrocephaly in patients with germline <i>PTEN</i> mutations and of brain weight in <i>Pten</i> knock-in murine model. <i>Eur J Hum Genet.</i> 2011 Jul;19(7):763-8. http://www.ncbi.nlm.nih.gov/pubmed/21343951

Gene	PMID	Reference Text
PTEN	22252256	Tan MH et al. Lifetime cancer risks in individuals with germline PTEN mutations. <i>Clin Cancer Res.</i> 2012 Jan 15;18(2):400-7. http://www.ncbi.nlm.nih.gov/pubmed/22252256
	22446940	Kurek KC et al. PTEN hamartoma of soft tissue: a distinctive lesion in PTEN syndromes. <i>Am J Surg Pathol.</i> 2012 May;36(5):671-87. http://www.ncbi.nlm.nih.gov/pubmed/22446940
	22595938	Mester J & Eng C. Estimate of de novo mutation frequency in probands with PTEN hamartoma tumor syndrome. <i>Genet Med.</i> 2012 Sep;14(9):819-22. http://www.ncbi.nlm.nih.gov/pubmed/22595938
	23335809	Bubien V et al. High cumulative risks of cancer in patients with PTEN hamartoma tumour syndrome. <i>J Med Genet.</i> 2013 Apr;50(4):255-63. http://www.ncbi.nlm.nih.gov/pubmed/23335809
	23934601	Nieuwenhuis MH et al. Cancer risk and genotype-phenotype correlations in PTEN hamartoma tumor syndrome. <i>Fam Cancer.</i> 2014 Mar;13(1):57-63. http://www.ncbi.nlm.nih.gov/pubmed/23934601
	26142096	Eng C. Cowden Syndrome. <i>J Genet Couns.</i> 1997 Jun;6(2):181-92. http://www.ncbi.nlm.nih.gov/pubmed/26142096
	3698331	Starink TM et al. The Cowden syndrome: a clinical and genetic study in 21 patients. <i>Clin Genet.</i> 1986 Mar;29(3):222-33. http://www.ncbi.nlm.nih.gov/pubmed/3698331
	657103	Brownstein MH et al. Cowden's disease: a cutaneous marker of breast cancer. <i>Cancer.</i> 1978 Jun;41(6):2393-8. http://www.ncbi.nlm.nih.gov/pubmed/657103
	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Breast and Ovarian. http://www.nccn.org/clinical.asp
	RAD50	16474176
19409520		Walters R et al. Human RAD50 deficiency in a Nijmegen breakage syndrome-like disorder. <i>Am J Hum Genet.</i> 2009 May;84(5):605-16. http://www.ncbi.nlm.nih.gov/pubmed/19409520
22006311		Walsh T et al. Mutations in 12 genes for inherited ovarian, fallopian tube, and peritoneal carcinoma identified by massively parallel sequencing. <i>Proc Natl Acad Sci U S A.</i> 2011 Nov 1;108(44):18032-7. http://www.ncbi.nlm.nih.gov/pubmed/22006311
RAD51C	20400963	Vaz F et al. Mutation of the RAD51C gene in a Fanconi anemia-like disorder. <i>Nat Genet.</i> 2010 May;42(5):406-9. http://www.ncbi.nlm.nih.gov/pubmed/20400963
	20400964	Meindl A et al. Germline mutations in breast and ovarian cancer pedigrees establish RAD51C as a human cancer susceptibility gene. <i>Nat Genet.</i> 2010 May;42(5):410-4. http://www.ncbi.nlm.nih.gov/pubmed/20400964
	20697805	Zheng Y et al. Screening RAD51C nucleotide alterations in patients with a family history of breast and ovarian cancer. <i>Breast Cancer Res Treat.</i> 2010 Dec;124(3):857-61. http://www.ncbi.nlm.nih.gov/pubmed/20697805
	20723205	Akbari MR et al. RAD51C germline mutations in breast and ovarian cancer patients. <i>Breast Cancer Res.</i> 2010;12(4):404. http://www.ncbi.nlm.nih.gov/pubmed/20723205
	21616938	Pelttari LM et al. RAD51C is a susceptibility gene for ovarian cancer. <i>Hum Mol Genet.</i> 2011 Aug 15;20(16):3278-88. http://www.ncbi.nlm.nih.gov/pubmed/21616938
	21990120	Thompson ER et al. Analysis of RAD51C germline mutations in high-risk breast and ovarian cancer families and ovarian cancer patients. <i>Hum Mutat.</i> 2012 Jan;33(1):95-9. http://www.ncbi.nlm.nih.gov/pubmed/21990120
	22370629	De Leener K et al. Evaluation of RAD51C as cancer susceptibility gene in a large breast-ovarian cancer patient population referred for genetic testing. <i>Breast Cancer Res Treat.</i> 2012 May;133(1):393-8. http://www.ncbi.nlm.nih.gov/pubmed/22370629
	22451500	Osorio A et al. Predominance of pathogenic missense variants in the RAD51C gene occurring in breast and ovarian cancer families. <i>Hum Mol Genet.</i> 2012 Jul 1;21(13):2889-98. http://www.ncbi.nlm.nih.gov/pubmed/22451500

Gene	PMID	Reference Text	
RAD51C	22538716	Loveday C et al. Germline RAD51C mutations confer susceptibility to ovarian cancer. <i>Nat Genet.</i> 2012 Apr 26;44(5):475-6 http://www.ncbi.nlm.nih.gov/pubmed/22538716	
	22725699	Coulet F et al. Germline RAD51C mutations in ovarian cancer susceptibility. <i>Clin Genet.</i> 2013 Apr;83(4):332-6. http://www.ncbi.nlm.nih.gov/pubmed/22725699	
	24240112	Pennington KP et al. Germline and somatic mutations in homologous recombination genes predict platinum response and survival in ovarian, fallopian tube, and peritoneal carcinomas. <i>Clin Cancer Res.</i> 2014 Feb 1;20(3):764-75. http://www.ncbi.nlm.nih.gov/pubmed/24240112	
	24359560	Schnurbein G et al. RAD51C deletion screening identifies a recurrent gross deletion in breast cancer and ovarian cancer families. <i>Breast Cancer Res.</i> 2013 Dec 20;15(6):R120. http://www.ncbi.nlm.nih.gov/pubmed/24359560	
	24504028	Cunningham JM et al. Clinical characteristics of ovarian cancer classified by BRCA1, BRCA2, and RAD51C status. <i>Sci Rep.</i> 2014 Feb 7;4:4026. http://www.ncbi.nlm.nih.gov/pubmed/24504028	
	24800917	Rashid MU et al. Deleterious RAD51C germline mutations rarely predispose to breast and ovarian cancer in Pakistan. <i>Breast Cancer Res Treat.</i> 2014 Jun;145(3):775-84. http://www.ncbi.nlm.nih.gov/pubmed/24800917	
	24993905	Gevensleben et al. Pathological features of breast and ovarian cancers in RAD51C germline mutation carriers. <i>Virchows Arch.</i> 2014 Sep;465(3):365-9. http://www.ncbi.nlm.nih.gov/pubmed/24993905	
	25086635	Blanco A et al. RAD51C germline mutations found in Spanish site-specific breast cancer and breast-ovarian cancer families. <i>Breast Cancer Res Treat.</i> 2014 Aug;147(1):133-43. http://www.ncbi.nlm.nih.gov/pubmed/25086635	
	25452441	Couch FJ et al. Inherited mutations in 17 breast cancer susceptibility genes among a large triple-negative breast cancer cohort unselected for family history of breast cancer. <i>J Clin Oncol.</i> 2015 Feb 1;33(4):304-11. http://www.ncbi.nlm.nih.gov/pubmed/25452441	
	25619955	Sokolenko AP et al. Candidate gene analysis of BRCA1/2 mutation-negative high-risk Russian breast cancer patients. <i>Cancer Lett.</i> 2015 Apr 10;359(2):259-61. http://www.ncbi.nlm.nih.gov/pubmed/25619955	
	26261251	Song H et al. Contribution of Germline Mutations in the RAD51B, RAD51C, and RAD51D Genes to Ovarian Cancer in the Population. <i>J Clin Oncol.</i> 2015 Sep 10;33(26):2901-7. http://www.ncbi.nlm.nih.gov/pubmed/26261251	
	26720728	Norquist BM et al. Inherited Mutations in Women With Ovarian Carcinoma. <i>Jama Oncology.</i> 2015 Dec 30:1-9. http://www.ncbi.nlm.nih.gov/pubmed/26720728	
	26740214	Jonson et al. Identification of six pathogenic RAD51C mutations via mutational screening of 1228 Danish individuals with increased risk of hereditary breast and/or ovarian cancer. <i>Breast Cancer Res Treat.</i> 2016 Jan;155(2):215-22. http://www.ncbi.nlm.nih.gov/pubmed/26740214	
	27433846	Pritchard CC et al. Inherited DNA-Repair Gene Mutations in Men with Metastatic Prostate Cancer. <i>N Engl J Med.</i> 2016 Aug 4;375(5):443-53. http://www.ncbi.nlm.nih.gov/pubmed/27433846	
	28888541	Lilyquist J et al. Frequency of mutations in a large series of clinically ascertained ovarian cancer cases tested on multi-gene panels compared to reference controls. <i>Gynecol Oncol.</i> 2017 Nov;147(2):375-380. 28888541 http://www.ncbi.nlm.nih.gov/pubmed/28888541	
	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Breast and Ovarian. http://www.nccn.org/clinical.asp	
	RAD51D	21822267	Loveday C et al. Germline mutations in RAD51D confer susceptibility to ovarian cancer. <i>Nat Genet.</i> 2011 Aug 7;43(9):879-82. http://www.ncbi.nlm.nih.gov/pubmed/21822267
		23372765	Thompson ER et al. Analysis of RAD51D in ovarian cancer patients and families with a history of ovarian or breast cancer. <i>PLoS One.</i> 2013;8(1):e54772. http://www.ncbi.nlm.nih.gov/pubmed/23372765
26261251		Song H et al. Contribution of Germline Mutations in the RAD51B, RAD51C, and RAD51D Genes to Ovarian Cancer in the Population. <i>J Clin Oncol.</i> 2015 Sep 10;33(26):2901-7. http://www.ncbi.nlm.nih.gov/pubmed/26261251	

Gene	PMID	Reference Text
RAD51D	26720728	Norquist BM et al. Inherited Mutations in Women With Ovarian Carcinoma. <i>Jama Oncology</i> . 2015 Dec 30;1-9. http://www.ncbi.nlm.nih.gov/pubmed/26720728
	27433846	Pritchard CC et al. Inherited DNA-Repair Gene Mutations in Men with Metastatic Prostate Cancer. <i>N Engl J Med</i> . 2016 Aug 4;375(5):443-53. http://www.ncbi.nlm.nih.gov/pubmed/27433846
	28418444	Couch FJ et al. Associations Between Cancer Predisposition Testing Panel Genes and Breast Cancer. <i>JAMA Oncol</i> . 2017 Apr 13. http://www.ncbi.nlm.nih.gov/pubmed/28418444
	28888541	Lilyquist J et al. Frequency of mutations in a large series of clinically ascertained ovarian cancer cases tested on multi-gene panels compared to reference controls. <i>Gynecol Oncol</i> . 2017 Nov;147(2):375-380. 28888541 http://www.ncbi.nlm.nih.gov/pubmed/28888541
	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Breast and Ovarian. http://www.nccn.org/clinical.asp
RB1		Lohmann DR, Gallie BL. Retinoblastoma. <i>Gene Reviews</i> 2000 Jul 18 [Updated 2015 Nov 19]. http://www.ncbi.nlm.nih.gov/books/NBK1452/
		Ries LAG, Smith MA, Gurney JG, Linet M, Tamra T, Young JL, Bunin GR (eds). <i>Cancer Incidence and Survival among Children and Adolescents: United States SEER Program 1975-1995</i> , National Cancer Institute, SEER Program. NIH Pub. No. 99-4649. Bethesda, MD, 1999.
	1637670	Draper GJ et al. Patterns of risk of hereditary retinoblastoma and applications to genetic counselling. <i>British Journal Of Cancer</i> . 1992 Jul 66(1):211-9. http://www.ncbi.nlm.nih.gov/pubmed/1637670
	17202110	Kleinerman RA et al. Risk of soft tissue sarcomas by individual subtype in survivors of hereditary retinoblastoma. <i>Journal Of The National Cancer Institute</i> . 2007 Jan 3 99(1):24-31. http://www.ncbi.nlm.nih.gov/pubmed/17202110
	18211953	Dimaras H et al. Loss of RB1 induces non-proliferative retinoma: increasing genomic instability correlates with progression to retinoblastoma. <i>Human Molecular Genetics</i> . 2008 May 15 17(10):1363-72. http://www.ncbi.nlm.nih.gov/pubmed/18211953
	19066271	Marees T et al. Risk of second malignancies in survivors of retinoblastoma: more than 40 years of follow-up. <i>Journal Of The National Cancer Institute</i> . 2008 Dec 17 100(24):1771-9. http://www.ncbi.nlm.nih.gov/pubmed/19066271
	1985763	Holladay DA et al. Clinical presentation, treatment, and outcome of trilateral retinoblastoma. <i>Cancer</i> . 1991 Feb 1 67(3):710-5. http://www.ncbi.nlm.nih.gov/pubmed/1985763
	20237571	National Retinoblastoma Strategy Canadian Guidelines for Care: Stratégie thérapeutique du rétinoblastome guide clinique canadien. <i>Canadian Journal Of Ophthalmology. Journal Canadien D'ophtalmologie</i> . 2009 44 Suppl 2:S1-88. http://www.ncbi.nlm.nih.gov/pubmed/20237571
	20301625	Lohmann DR, Gallie BL. Retinoblastoma. 2000 Jul 18 [Updated 2015 Nov 19]. In: Pagon RA, Adam MP, Ardinger HH, et al., editors. <i>GeneReviews</i> ® [Internet]. Seattle (WA): University of Washington, Seattle; 1993-2016. Available from: https://www.ncbi.nlm.nih.gov/books/NBK1452/ http://www.ncbi.nlm.nih.gov/pubmed/20301625
	22355046	Kleinerman RA et al. Variation of second cancer risk by family history of retinoblastoma among long-term survivors. <i>Journal Of Clinical Oncology : Official Journal Of The American Society Of Clinical Oncology</i> . 2012 Mar 20 30(9):950-7. http://www.ncbi.nlm.nih.gov/pubmed/22355046
	22909775	Castéra L et al. Fine mapping of whole RB1 gene deletions in retinoblastoma patients confirms PCDH8 as a candidate gene for psychomotor delay. <i>European Journal Of Human Genetics : Ejhg</i> . 2013 Apr 21(4):460-4. http://www.ncbi.nlm.nih.gov/pubmed/22909775
	25126964	de Jong MC et al. Trilateral retinoblastoma: a systematic review and meta-analysis. <i>The Lancet. Oncology</i> . 2014 Sep 15(10):1157-67. http://www.ncbi.nlm.nih.gov/pubmed/25126964
	25970657	Temming P et al. Pediatric second primary malignancies after retinoblastoma treatment. <i>Pediatric Blood & Cancer</i> . 2015 Oct 62(10):1799-804. http://www.ncbi.nlm.nih.gov/pubmed/25970657
	7927327	Lohmann DR et al. Distinct RB1 gene mutations with low penetrance in hereditary retinoblastoma. <i>Human Genetics</i> . 1994 Oct 94(4):349-54. http://www.ncbi.nlm.nih.gov/pubmed/7927327

Gene	PMID	Reference Text
RB1	9333268	Wong FL et al. Cancer incidence after retinoblastoma. Radiation dose and sarcoma risk. <i>Jama</i> . 1997 Oct 15 278(15):1262-7. http://www.ncbi.nlm.nih.gov/pubmed/9333268
	9497263	Sippel KC et al. Frequency of somatic and germ-line mosaicism in retinoblastoma: implications for genetic counseling. <i>American Journal Of Human Genetics</i> . 1998 Mar 62(3):610-9. http://www.ncbi.nlm.nih.gov/pubmed/9497263
	SEER	Young JL, Ries et al. Retinoblastoma. Cancer Incidence and Survival among Children and Adolescents: United States SEER Program 1975-1995, National Cancer Institute, SEER Program. NIH Pub. No. 99-4649. Bethesda, MD, 1999. http://seer.cancer.gov/archive/publications/childhood/childhood-monograph.pdf
RECQL	25915596	Cybulski C et al. Germline RECQL mutations are associated with breast cancer susceptibility. <i>Nature Genetics</i> . 2015 Apr 27. http://www.ncbi.nlm.nih.gov/pubmed/25915596
	25945795	Sun J et al. Mutations in RECQL Gene Are Associated with Predisposition to Breast Cancer. <i>P Lo S Genetics</i> . 2015 May 11(5):e1005228. http://www.ncbi.nlm.nih.gov/pubmed/25945795
	26960971	Lindor NM, Hopper J, Dowty J (2016) Estimating cumulative risks for breast cancer for carriers of variants in uncommon genes. <i>Fam Cancer</i> 15:367–370 http://www.ncbi.nlm.nih.gov/pubmed/26960971
	27832498	Bogdanova N et al. Analysis of a RECQL splicing mutation, c.1667_1667+3delAGTA, in breast cancer patients and controls from Central Europe. <i>Familial Cancer</i> . 2017 Apr 16(2):181-186. http://www.ncbi.nlm.nih.gov/pubmed/27832498
RET	10220148	Takahashi M et al. Co-segregation of MEN2 and Hirschsprung's disease: the same mutation of RET with both gain and loss-of-function? <i>Hum Mutat</i> . 1999;13(4):331-6. http://www.ncbi.nlm.nih.gov/pubmed/10220148
	11174835	Nguyen L et al. Pheochromocytoma in multiple endocrine neoplasia type 2: a prospective study. <i>Eur J Endocrinol</i> . 2001 Jan;144(1):37-44. http://www.ncbi.nlm.nih.gov/pubmed/11174835
	11739416	Brandi ML et al. Guidelines for diagnosis and therapy of MEN type 1 and type 2. <i>J Clin Endocrinol Metab</i> . 2001 Dec;86(12):5658-71. http://www.ncbi.nlm.nih.gov/pubmed/11739416
	11953745	Gabriel SB et al. Segregation at three loci explains familial and population risk in Hirschsprung disease. <i>Nat Genet</i> . 2002 May;31(1):89-93. http://www.ncbi.nlm.nih.gov/pubmed/11953745
	12686527	Yip L et al. Multiple endocrine neoplasia type 2: evaluation of the genotype-phenotype relationship. <i>Arch Surg</i> . 2003 Apr;138(4):409-16; discussion 416. http://www.ncbi.nlm.nih.gov/pubmed/12686527
	15827097	Machens A et al. Codon-specific development of pheochromocytoma in multiple endocrine neoplasia type 2. <i>J Clin Endocrinol Metab</i> . 2005 Jul;90(7):3999-4003. http://www.ncbi.nlm.nih.gov/pubmed/15827097
	17895320	Elisei R et al. RET genetic screening in patients with medullary thyroid cancer and their relatives: experience with 807 individuals at one center. <i>J Clin Endocrinol Metab</i> . 2007 Dec;92(12):4725-9. http://www.ncbi.nlm.nih.gov/pubmed/17895320
	17965226	Amiel J et al. Hirschsprung disease, associated syndromes and genetics: a review. <i>J Med Genet</i> . 2008 Jan;45(1):1-14. http://www.ncbi.nlm.nih.gov/pubmed/17965226
	19469690	American Thyroid Association Guidelines Task Force. Medullary thyroid cancer: management guidelines of the American Thyroid Association. <i>Thyroid : Official Journal Of The American Thyroid Association</i> . 2009 Jun 19(6):565-612. http://www.ncbi.nlm.nih.gov/pubmed/19469690
	20301434	Marquard J, Eng C. Multiple Endocrine Neoplasia Type 2. 1999 Sep 27 [Updated 2015 Jun 25]. In: Pagon RA, Adam MP, Ardinger HH, et al., editors. <i>GeneReviews</i> ® [Internet]. Seattle (WA): University of Washington, Seattle; 1993-2015. http://www.ncbi.nlm.nih.gov/books/NBK1257/#men2.REF.americanthyroidassocia.2009.565
	20301612	Parisi MA. Hirschsprung Disease Overview. 2002 Jul 12 [Updated 2015 Oct 1]. In: Pagon RA, Adam MP, Ardinger HH, et al., editors. <i>GeneReviews</i> ® [Internet]. Seattle (WA): University of Washington, Seattle; 1993-2017. Available from: http://www.ncbi.nlm.nih.gov/books/NBK1439/ http://www.ncbi.nlm.nih.gov/pubmed/20301612

Gene	PMID	Reference Text
RET	25440022	Romei C et al. Twenty years of lesson learning: how does the RET genetic screening test impact the clinical management of medullary thyroid cancer? Clin Endocrinol (Oxf). 2015 Jun;82(6):892-9. http://www.ncbi.nlm.nih.gov/pubmed/25440022
	2563193	Easton DF et al. The clinical and screening age-at-onset distribution for the MEN-2 syndrome. Am J Hum Genet. 1989 Feb;44(2):208-15. http://www.ncbi.nlm.nih.gov/pubmed/2563193
	25810047	Wells SA et al. Revised American Thyroid Association guidelines for the management of medullary thyroid carcinoma. Thyroid. 2015 Jun;25(6):567-610. http://www.ncbi.nlm.nih.gov/pubmed/25810047
	26920351	Scapinelli JO et al. MEN 2A-related cutaneous lichen amyloidosis: report of three kindred and systematic literature review of clinical, biochemical and molecular characteristics. Fam Cancer. 2016 Oct; 15(4):625-33. http://www.ncbi.nlm.nih.gov/pubmed/26920351
	28274275	Gui H et al. Whole exome sequencing coupled with unbiased functional analysis reveals new Hirschsprung disease genes. Genome Biol. 2017 Mar 8;18(1):48. http://www.ncbi.nlm.nih.gov/pubmed/28274275
	7581377	Attié T et al. Diversity of RET proto-oncogene mutations in familial and sporadic Hirschsprung disease. Hum Mol Genet. 1995 Aug;4(8):1381-6. http://www.ncbi.nlm.nih.gov/pubmed/7581377
	7595170	Mulligan LM et al. Genotype-phenotype correlation in multiple endocrine neoplasia type 2: report of the International RET Mutation Consortium. J Intern Med. 1995 Oct;238(4):343-6. http://www.ncbi.nlm.nih.gov/pubmed/7595170
	7874109	Schuffenecker I et al. RET proto-oncogene mutations in French MEN 2A and FMTC families. Hum Mol Genet. 1994 Nov;3(11):1939-43. http://www.ncbi.nlm.nih.gov/pubmed/7874109
	7907913	Mulligan LM et al. Specific mutations of the RET proto-oncogene are related to disease phenotype in MEN 2A and FMTC. Nat Genet. 1994 Jan;6(1):70-4. http://www.ncbi.nlm.nih.gov/pubmed/7907913
	7977365	Carlson KM et al. Parent-of-origin effects in multiple endocrine neoplasia type 2B. Am J Hum Genet. 1994 Dec;55(6):1076-82. http://www.ncbi.nlm.nih.gov/pubmed/7977365
	8632274	Skinner MA et al. Medullary thyroid carcinoma in children with multiple endocrine neoplasia types 2A and 2B. J Pediatr Surg. 1996 Jan;31(1):177-81; Discussion 181-2. http://www.ncbi.nlm.nih.gov/pubmed/8632274
	8880581	Morrison PJ and Nevin NC. Multiple endocrine neoplasia type 2B (mucosal neuroma syndrome, Wagenmann-Froboese syndrome). J Med Genet. 1996 Sep;33(9):779-82. http://www.ncbi.nlm.nih.gov/pubmed/8880581
	8918855	Eng C et al. The relationship between specific RET proto-oncogene mutations and disease phenotype in multiple endocrine neoplasia type 2. International RET mutation consortium analysis. JAMA. 1996 Nov 20;276(19):1575-9. http://www.ncbi.nlm.nih.gov/pubmed/8918855
	9294615	Smith DP et al. Germline mutation of RET codon 883 in two cases of de novo MEN 2B. Oncogene. 1997 Sep 4;15(10):1213-7. http://www.ncbi.nlm.nih.gov/pubmed/9294615
	NCCN	NCCN Guidelines. Thyroid Carcinoma. Version 2.2015 (URL: http://www.nccn.org/professionals/physician_gls/pdf/thyroid.pdf) November 2015 accessed. http://www.nccn.org/professionals/physician_gls/pdf/thyroid.pdf
SCG5/ GREM1	22561515	Jaeger E et al. Hereditary mixed polyposis syndrome is caused by a 40-kb upstream duplication that leads to increased and ectopic expression of the BMP antagonist GREM1. Nat Genet. 2012 May 6;44(6):699-703. http://www.ncbi.nlm.nih.gov/pubmed/22561515
	26493165	Rohlin A et al. GREM1 and POLE variants in hereditary colorectal cancer syndromes. Genes, Chromosomes & Cancer. 2016 Jan 55(1):95-106. http://www.ncbi.nlm.nih.gov/pubmed/26493165
	26947005	Ziai J et al. Defining the polyposis/colorectal cancer phenotype associated with the Ashkenazi GREM1 duplication: counselling and management recommendations. Genet Res (Camb). 2016 Mar 7;98:e5. http://www.ncbi.nlm.nih.gov/pubmed/26947005

Gene	PMID	Reference Text	
SCG5/ GREM1	27984123	Plesec T et al. Clinicopathological features of a kindred with SCG5-GREM1-associated hereditary mixed polyposis syndrome. Hum Pathol. 2017 Feb;60:75-81. http://www.ncbi.nlm.nih.gov/pubmed/27984123	
	28242209	Lieberman S et al. Features of Patients With Hereditary Mixed Polyposis Syndrome Caused by Duplication of GREM1 and Implications for Screening and Surveillance. Gastroenterology. 2017 Jun;152(8):1876-1880.e1. http://www.ncbi.nlm.nih.gov/pubmed/28242209	
	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Colorectal. http://www.nccn.org/clinical.asp	
SDHA	10746566	Parfait B et al. Compound heterozygous mutations in the flavoprotein gene of the respiratory chain complex II in a patient with Leigh syndrome. Hum Genet. 2000 Feb;106(2):236-43. http://www.ncbi.nlm.nih.gov/pubmed/10746566	
	20301715	Kirman S and Young WF. Hereditary Paraganglioma-Pheochromocytoma Syndromes. 2008 May 21 [Updated 2014 Nov 6]. In: Pagon RA, Bird TD, Dolan CR, et al., editors. GeneReviews [Internet]. Seattle (WA): University of Washington, Seattle; 1993-. http://www.ncbi.nlm.nih.gov/pubmed/20301715	
	20484225	Burnichon N et al. SDHA is a tumor suppressor gene causing paraganglioma. Hum Mol Genet. 2010 Aug 1;19(15):3011-20. http://www.ncbi.nlm.nih.gov/pubmed/20484225	
	20551992	Levitas A et al. Familial neonatal isolated cardiomyopathy caused by a mutation in the flavoprotein subunit of succinate dehydrogenase. Eur J Hum Genet. 2010 Oct;18(10):1160-5. http://www.ncbi.nlm.nih.gov/pubmed/20551992	
	21505157	Pantaleo MA et al. SDHA loss-of-function mutations in KIT-PDGFR α wild-type gastrointestinal stromal tumors identified by massively parallel sequencing. J Natl Cancer Inst. 2011 Jun 22;103(12):983-7. http://www.ncbi.nlm.nih.gov/pubmed/21505157	
	21752896	Korpershoek E et al. SDHA immunohistochemistry detects germline SDHA gene mutations in apparently sporadic paragangliomas and pheochromocytomas. J Clin Endocrinol Metab. 2011 Sep;96(9):E1472-6. http://www.ncbi.nlm.nih.gov/pubmed/21752896	
	22972948	Alston CL et al. Recessive germline SDHA and SDHB mutations causing leukodystrophy and isolated mitochondrial complex II deficiency. J Med Genet. 2012 Sep;49(9):569-77. http://www.ncbi.nlm.nih.gov/pubmed/22972948	
	22974104	Italiano A et al. SDHA loss of function mutations in a subset of young adult wild-type gastrointestinal stromal tumors. BMC Cancer. 2012 Sep 14;12:408. http://www.ncbi.nlm.nih.gov/pubmed/22974104	
	23174333	Hoekstra AS and Bayley JP. The role of complex II in disease. Biochimica Et Biophysica Acta. 2013 May 1827(5):543-51. http://www.ncbi.nlm.nih.gov/pubmed/23174333	
	24781757	Renkema GH et al. SDHA mutations causing a multisystem mitochondrial disease: novel mutations and genetic overlap with hereditary tumors. Eur J Hum Genet. 2015 Feb;23(2):202-9. http://www.ncbi.nlm.nih.gov/pubmed/24781757	
	24893135	Lenders JW et al. Pheochromocytoma and paraganglioma: an endocrine society clinical practice guideline. The Journal Of Clinical Endocrinology And Metabolism. 2014 Jun 99(6):1915-42.24893135 http://www.ncbi.nlm.nih.gov/pubmed/24893135	
	26642834	Helman G et al. Magnetic resonance imaging spectrum of succinate dehydrogenase-related infantile leukoencephalopathy. Annals Of Neurology. 2016 Mar 79(3):379-86. http://www.ncbi.nlm.nih.gov/pubmed/26642834	
	7550341	Bourgeron T et al. Mutation of a nuclear succinate dehydrogenase gene results in mitochondrial respiratory chain deficiency. Nat Genet. 1995 Oct;11(2):144-9. http://www.ncbi.nlm.nih.gov/pubmed/7550341	
	SDHAF2	19628817	Hao HX et al. SDH5, a gene required for flavination of succinate dehydrogenase, is mutated in paraganglioma. Science (New York, N.Y.). 2009 325(5944):1139-42. http://www.ncbi.nlm.nih.gov/pubmed/19628817
		20071235	Bayley JP et al. SDHAF2 mutations in familial and sporadic paraganglioma and pheochromocytoma. The Lancet Oncology. 2010 11(4):366-72. http://www.ncbi.nlm.nih.gov/pubmed/20071235

Gene	PMID	Reference Text
SDHAF2	20301715	Kirmani S and Young WF. Hereditary Paraganglioma-Pheochromocytoma Syndromes. 2008 May 21 [Updated 2014 Nov 6]. In: Pagon RA, Bird TD, Dolan CR, et al., editors. GeneReviews [Internet]. Seattle (WA): University of Washington, Seattle; 1993-. http://www.ncbi.nlm.nih.gov/pubmed/20301715
	21224366	Kunst HP et al. SDHAF2 (PGL2-SDH5) and hereditary head and neck paraganglioma. Clinical Cancer Research : An Official Journal Of The American Association For Cancer Research. 2011 17(2):247-54. http://www.ncbi.nlm.nih.gov/pubmed/21224366
	24893135	Lenders JW et al. Pheochromocytoma and paraganglioma: an endocrine society clinical practice guideline. The Journal Of Clinical Endocrinology And Metabolism. 2014 Jun 99(6):1915-42.24893135 http://www.ncbi.nlm.nih.gov/pubmed/24893135
	26273102	Benn DE et al. 15 YEARS OF PARAGANGLIOMA: Clinical manifestations of paraganglioma syndromes types 1-5. Endocrine Related Cancer. 2015 Aug 22(4):T91-103. http://www.ncbi.nlm.nih.gov/pubmed/26273102
	6286462	van Baars F et al. Genetic aspects of nonchromaffin paraganglioma. Human Genetics. 1982 60(4):305-9 http://www.ncbi.nlm.nih.gov/pubmed/6286462
	SDHB	11404820
14685938		Vanharanta S et al. Early-onset renal cell carcinoma as a novel extraparaganglial component of SDHB-associated heritable paraganglioma. Am J Hum Genet. 2004 Jan;74(1):153-9. http://www.ncbi.nlm.nih.gov/pubmed/14685938
15328326		Neumann HP et al. Distinct clinical features of paraganglioma syndromes associated with SDHB and SDHD gene mutations. JAMA. 2004 Aug 25;292(8):943-51. http://www.ncbi.nlm.nih.gov/pubmed/15328326
16314641		Amar L et al. Genetic testing in pheochromocytoma or functional paraganglioma. J Clin Oncol. 2005 Dec 1;23(34):8812-8. http://www.ncbi.nlm.nih.gov/pubmed/16314641
16317055		Benn DE et al. Clinical presentation and penetrance of pheochromocytoma/paraganglioma syndromes. J Clin Endocrinol Metab. 2006 Mar;91(3):827-36. http://www.ncbi.nlm.nih.gov/pubmed/16317055
16912137		Brouwers FM et al. High frequency of SDHB germline mutations in patients with malignant catecholamine-producing paragangliomas: implications for genetic testing. J Clin Endocrinol Metab. 2006 Nov;91(11):4505-9. http://www.ncbi.nlm.nih.gov/pubmed/16912137
17200167		Timmers HJ et al. Clinical presentations, biochemical phenotypes, and genotype-phenotype correlations in patients with succinate dehydrogenase subunit B-associated pheochromocytomas and paragangliomas. J Clin Endocrinol Metab. 2007 Mar;92(3):779-86. http://www.ncbi.nlm.nih.gov/pubmed/17200167
17652212		Amar L et al. Succinate dehydrogenase B gene mutations predict survival in patients with malignant pheochromocytomas or paragangliomas. J Clin Endocrinol Metab. 2007 Oct;92(10):3822-8. http://www.ncbi.nlm.nih.gov/pubmed/17652212
17667967		Pasini B et al. Clinical and molecular genetics of patients with the Carney-Stratakis syndrome and germline mutations of the genes coding for the succinate dehydrogenase subunits SDHB, SDHC, and SDHD. Eur J Hum Genet. 2008 Jan;16(1):79-88. http://www.ncbi.nlm.nih.gov/pubmed/17667967
18728283		Ricketts C et al. Germline SDHB mutations and familial renal cell carcinoma. J Natl Cancer Inst. 2008 Sep 3;100(17):1260-2. http://www.ncbi.nlm.nih.gov/pubmed/18728283
20301715		Kirmani S and Young WF. Hereditary Paraganglioma-Pheochromocytoma Syndromes. 2008 May 21 [Updated 2014 Nov 6]. In: Pagon RA, Bird TD, Dolan CR, et al., editors. GeneReviews [Internet]. Seattle (WA): University of Washington, Seattle; 1993-. http://www.ncbi.nlm.nih.gov/pubmed/20301715
22041710		Welander J et al. Genetics and clinical characteristics of hereditary pheochromocytomas and paragangliomas. Endocr Relat Cancer. 2011 Dec 1;18(6):R253-76. http://www.ncbi.nlm.nih.gov/pubmed/22041710
22972948		Alston CL et al. Recessive germline SDHA and SDHB mutations causing leukodystrophy and isolated mitochondrial complex II deficiency. J Med Genet. 2012 Sep;49(9):569-77. http://www.ncbi.nlm.nih.gov/pubmed/22972948

Gene	PMID	Reference Text
SDHB	23072324	Jafri M et al. Evaluation of SDHB, SDHD and VHL gene susceptibility testing in the assessment of individuals with non-syndromic pheochromocytoma, paraganglioma and head and neck paraganglioma. Clin Endocrinol (Oxf). 2013 Jun;78(6):898-906. http://www.ncbi.nlm.nih.gov/pubmed/23072324
	23099648	van Hulstijn et al. Risk of malignant paraganglioma in SDHB-mutation and SDHD-mutation carriers: a systematic review and meta-analysis. J Med Genet. 2012 Dec;49(12):768-76. http://www.ncbi.nlm.nih.gov/pubmed/23099648
	24893135	Lenders JW et al. Pheochromocytoma and paraganglioma: an endocrine society clinical practice guideline. The Journal Of Clinical Endocrinology And Metabolism. 2014 Jun 99(6):1915-42.24893135 http://www.ncbi.nlm.nih.gov/pubmed/24893135
	26925370	Ardisson A et al. Mitochondrial leukoencephalopathy and complex II deficiency associated with a recessive SDHB mutation with reduced penetrance. Molecular Genetics And Metabolism Reports. 2015 Dec 5:51-54. http://www.ncbi.nlm.nih.gov/pubmed/26925370
	27604842	Grønberg S et al. Leukoencephalopathy due to Complex II Deficiency and Bi-Allelic SDHB Mutations: Further Cases and Implications for Genetic Counselling. Jimd Reports. 2016 Sep 8. http://www.ncbi.nlm.nih.gov/pubmed/27604842
	27839933	Assadipour Y et al. SDHB mutation status and tumor size but not tumor grade are important predictors of clinical outcome in pheochromocytoma and abdominal paraganglioma. Surgery. 2017 Jan 161(1):230-239. http://www.ncbi.nlm.nih.gov/pubmed/27839933
SDHC	11062460	Niemann S and Müller U. Mutations in SDHC cause autosomal dominant paraganglioma, type 3. Nat Genet. 2000 Nov;26(3):268-70. http://www.ncbi.nlm.nih.gov/pubmed/11062460
	16249420	Schiavi F et al. Predictors and prevalence of paraganglioma syndrome associated with mutations of the SDHC gene. JAMA. 2005 Oct 26;294(16):2057-63. http://www.ncbi.nlm.nih.gov/pubmed/16249420
	17667967	Pasini B et al. Clinical and molecular genetics of patients with the Carney-Stratakis syndrome and germline mutations of the genes coding for the succinate dehydrogenase subunits SDHB, SDHC, and SDHD. Eur J Hum Genet. 2008 Jan;16(1):79-88. http://www.ncbi.nlm.nih.gov/pubmed/17667967
	18212813	Peczowska M et al. Extra-adrenal and adrenal pheochromocytomas associated with a germline SDHC mutation. Nat Clin Pract Endocrinol Metab. 2008 Feb;4(2):111-5. http://www.ncbi.nlm.nih.gov/pubmed/18212813
	20301715	Kirmani S and Young WF. Hereditary Paraganglioma-Pheochromocytoma Syndromes. 2008 May 21 [Updated 2014 Nov 6]. In: Pagon RA, Bird TD, Dolan CR, et al., editors. GeneReviews [Internet]. Seattle (WA): University of Washington, Seattle; 1993-. http://www.ncbi.nlm.nih.gov/pubmed/20301715
	23083876	Ricketts CJ et al. Succinate dehydrogenase kidney cancer: an aggressive example of the Warburg effect in cancer. J Urol. 2012 Dec;188(6):2063-71. http://www.ncbi.nlm.nih.gov/pubmed/23083876
	24758179	Eise T et al. The clinical phenotype of SDHC-associated hereditary paraganglioma syndrome (PGL3). J Clin Endocrinol Metab. 2014 Aug;99(8):E1482-6. http://www.ncbi.nlm.nih.gov/pubmed/24758179
	24886695	Miettinen M and Lasota J. Succinate dehydrogenase deficient gastrointestinal stromal tumors (GISTs) - a review. The International Journal Of Biochemistry & Cell Biology. 2014 Aug 53:514-9. http://www.ncbi.nlm.nih.gov/pubmed/24886695
	24893135	Lenders JW et al. Pheochromocytoma and paraganglioma: an endocrine society clinical practice guideline. The Journal Of Clinical Endocrinology And Metabolism. 2014 Jun 99(6):1915-42.24893135 http://www.ncbi.nlm.nih.gov/pubmed/24893135
SDHD	10657297	Baysal BE et al. Mutations in SDHD, a mitochondrial complex II gene, in hereditary paraganglioma. Science. 2000 Feb 4;287(5454):848-51. http://www.ncbi.nlm.nih.gov/pubmed/10657297
	15328326	Neumann HP et al. Distinct clinical features of paraganglioma syndromes associated with SDHB and SDHD gene mutations. JAMA. 2004 Aug 25;292(8):943-51. http://www.ncbi.nlm.nih.gov/pubmed/15328326
	16317055	Benn DE et al. Clinical presentation and penetrance of pheochromocytoma/paraganglioma syndromes. J Clin Endocrinol Metab. 2006 Mar;91(3):827-36. http://www.ncbi.nlm.nih.gov/pubmed/16317055

Gene	PMID	Reference Text
SDHD	20301715	Kirmani S and Young WF. Hereditary Paraganglioma-Pheochromocytoma Syndromes. 2008 May 21 [Updated 2014 Nov 6]. In: Pagon RA, Bird TD, Dolan CR, et al., editors. GeneReviews [Internet]. Seattle (WA): University of Washington, Seattle; 1993-. http://www.ncbi.nlm.nih.gov/pubmed/20301715
	21937622	Yeap PM et al. Molecular analysis of pheochromocytoma after maternal transmission of SDHD mutation elucidates mechanism of parent-of-origin effect. The Journal Of Clinical Endocrinology And Metabolism. 2011 Dec 96(12):E2009-13. http://www.ncbi.nlm.nih.gov/pubmed/21937622
	22041710	Welander J et al. Genetics and clinical characteristics of hereditary pheochromocytomas and paragangliomas. Endocr Relat Cancer. 2011 Dec 1;18(6):R253-76. http://www.ncbi.nlm.nih.gov/pubmed/22041710
	23072324	Jafri M et al. Evaluation of SDHB, SDHD and VHL gene susceptibility testing in the assessment of individuals with non-syndromic phaeochromocytoma, paraganglioma and head and neck paraganglioma. Clin Endocrinol (Oxf). 2013 Jun;78(6):898-906. http://www.ncbi.nlm.nih.gov/pubmed/23072324
	23083876	Ricketts CJ et al. Succinate dehydrogenase kidney cancer: an aggressive example of the Warburg effect in cancer. J Urol. 2012 Dec;188(6):2063-71. http://www.ncbi.nlm.nih.gov/pubmed/23083876
	23282968	Miettinen M et al. Immunohistochemical loss of succinate dehydrogenase subunit A (SDHA) in gastrointestinal stromal tumors (GISTs) signals SDHA germline mutation. The American Journal Of Surgical Pathology. 2013 37(2):234-40. http://www.ncbi.nlm.nih.gov/pubmed/23282968
	24096523	Papathomas TG et al. Non-pheochromocytoma (PCC)/paraganglioma (PGL) tumors in patients with succinate dehydrogenase-related PCC-PGL syndromes: a clinicopathological and molecular analysis. European Journal Of Endocrinology. 2014 Jan 170(1):1-12. http://www.ncbi.nlm.nih.gov/pubmed/24096523
	24367056	Jackson CB et al. Mutations in SDHD lead to autosomal recessive encephalomyopathy and isolated mitochondrial complex II deficiency. J Med Genet. 2014 Mar;51(3):170-5. 24367056 http://www.ncbi.nlm.nih.gov/pubmed/24367056
	24893135	Lenders JW et al. Pheochromocytoma and paraganglioma: an endocrine society clinical practice guideline. The Journal Of Clinical Endocrinology And Metabolism. 2014 Jun 99(6):1915-42.24893135 http://www.ncbi.nlm.nih.gov/pubmed/24893135
	25300370	Bayley JP et al. Paraganglioma and pheochromocytoma upon maternal transmission of SDHD mutations. BMC Medical Genetics. 2014 15:111. http://www.ncbi.nlm.nih.gov/pubmed/25300370
	26008905	Alston CL et al. A recessive homozygous p.Asp92Gly SDHD mutation causes prenatal cardiomyopathy and a severe mitochondrial complex II deficiency. Hum Genet. 2015 Aug;134(8):869-79. http://www.ncbi.nlm.nih.gov/pubmed/26008905
	26259135	Niemeijer ND et al. Succinate Dehydrogenase (SDH)-Deficient Pancreatic Neuroendocrine Tumor Expands the SDH-Related Tumor Spectrum. The Journal Of Clinical Endocrinology And Metabolism. 2015 Oct 100(10):E1386-93. http://www.ncbi.nlm.nih.gov/pubmed/26259135
	27856506	Burnichon N et al. Risk assessment of maternally inherited SDHD paraganglioma and phaeochromocytoma. Journal Of Medical Genetics. 2017 Feb 54(2):125-133. http://www.ncbi.nlm.nih.gov/pubmed/27856506
	SMAD4	16436638
16525031		Cao X et al. Mapping of hereditary mixed polyposis syndrome (HMPS) to chromosome 10q23 by genomewide high-density single nucleotide polymorphism (SNP) scan and identification of BMPR1A loss of function. J Med Genet. 2006 Mar;43(3):e13. http://www.ncbi.nlm.nih.gov/pubmed/16525031
17303595		Brosens LAA et al. Risk of colorectal cancer in juvenile polyposis. Gut. 2007;56:965-7. http://www.ncbi.nlm.nih.gov/pubmed/17303595
19773747		Cheah PY. Germline bone morphogenesis protein receptor 1A mutation causes colorectal tumorigenesis in hereditary mixed polyposis syndrome. Am J Gastroenterol. 2009 Dec;104(12):3027-33. http://www.ncbi.nlm.nih.gov/pubmed/19773747
20101697		Gallione C et al. Overlapping spectra of SMAD4 mutations in juvenile polyposis (JP) and JP-HHT syndrome. American Journal Of Medical Genetics. Part A. 2010 152A(2):333-9. http://www.ncbi.nlm.nih.gov/pubmed/20101697

Gene	PMID	Reference Text
SMAD4	20301525	McDonald J and Pyeritz RE. Hereditary Hemorrhagic Telangiectasia (2000 [updated 2012]). In Pagon RA, Adam MP, Bird TD, Dolan CR, Fong CT, and Stephens K (Eds), GeneReviews [Internet]. Seattle (WA): University of Washington, Seattle. http://www.ncbi.nlm.nih.gov/pubmed/20301525
	22810475	O'Malley M et al. The prevalence of hereditary hemorrhagic telangiectasia in juvenile polyposis syndrome. Diseases Of The Colon And Rectum. 2012 Aug 55(8):886-92. http://www.ncbi.nlm.nih.gov/pubmed/22810475
	22965402	Latchford AR et al. Juvenile polyposis syndrome: a study of genotype, phenotype, and long-term outcome. Diseases Of The Colon And Rectum. 2012 Oct 55(10):1038-43. http://www.ncbi.nlm.nih.gov/pubmed/22965402
	24525918	Wain KE et al. Appreciating the broad clinical features of SMAD4 mutation carriers: a multicenter chart review. Genetics In Medicine : Official Journal Of The American College Of Medical Genetics. 2014 Aug 16(8):588-93. http://www.ncbi.nlm.nih.gov/pubmed/24525918
	25389115	Aytac E et al. Genotype-defined cancer risk in juvenile polyposis syndrome. The British Journal Of Surgery. 2015 Jan 102(1):114-8. http://www.ncbi.nlm.nih.gov/pubmed/25389115
	25848489	Campos FG et al. Colorectal cancer risk in hamartomatous polyposis syndromes. World Journal Of Gastrointestinal Surgery. 2015 Mar 27 7(3):25-32. http://www.ncbi.nlm.nih.gov/pubmed/25848489
	25918283	Stoffel EM. Screening in GI Cancers: The Role of Genetics. Journal Of Clinical Oncology : Official Journal Of The American Society Of Clinical Oncology. 2015 Jun 01 33(16):1721-8. http://www.ncbi.nlm.nih.gov/pubmed/25918283
	25931195	Heald B et al. Prevalence of thoracic aortopathy in patients with juvenile Polyposis Syndrome-Hereditary Hemorrhagic Telangiectasia due to SMAD4. American Journal Of Medical Genetics. Part A. 2015 Aug 167(8):1758-62. http://www.ncbi.nlm.nih.gov/pubmed/25931195
	26159157	Soer E et al. Massive gastric polyposis associated with a germline SMAD4 gene mutation. Familial Cancer. 2015 Dec 14(4):569-73. http://www.ncbi.nlm.nih.gov/pubmed/26159157
	26363537	Jasperson K and Burt RW. The Genetics of Colorectal Cancer. Surgical Oncology Clinics Of North America. 2015 Oct 24(4):683-703. http://www.ncbi.nlm.nih.gov/pubmed/26363537
	27212857	Kadiyska T et al. Clinical and genetic challenges in a family with history of childhood polyp, aortopathy, and clinical diagnosis of hereditary hemorrhagic teleangiectasia (HHT). Annals Of Pediatric Cardiology. 2016 May-Aug 9(2):176-8. http://www.ncbi.nlm.nih.gov/pubmed/27212857
	27326320	Sheny S. Genetic risks and familial associations of small bowel carcinoma. World Journal Of Gastrointestinal Oncology. 2016 Jun 15 8(6):509-19. http://www.ncbi.nlm.nih.gov/pubmed/27326320
	9643289	Desai DC et al. A survey of phenotypic features in juvenile polyposis. Journal Of Medical Genetics. 1998 Jun 35(6):476-81. http://www.ncbi.nlm.nih.gov/pubmed/9643289
	9869523	Howe JR et al. The risk of gastrointestinal carcinoma in familial juvenile polyposis. Ann Surg Oncol. 1998 Dec;5(8):751-6. http://www.ncbi.nlm.nih.gov/pubmed/9869523
	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Colorectal. http://www.nccn.org/clinical.asp
	SMARCA4	20137775
21566516		Hasselblatt M et al. Nonsense mutation and inactivation of SMARCA4 (BRG1) in an atypical teratoid/rhabdoid tumor showing retained SMARCB1 (INI1) expression. Am J Surg Pathol. 2011 Jun;35(6):933-5. http://www.ncbi.nlm.nih.gov/pubmed/21566516
23556151		Schrier Vergano SA et al. Coffin-Siris Syndrome. 2013 Apr 4 [Updated 2013 Jul 11]. In: Pagon RA, Adam MP, Ardinger HH, et al., editors. GeneReviews® [Internet]. Seattle (WA): University of Washington, Seattle; 1993-2016. http://www.ncbi.nlm.nih.gov/books/NBK131811/
23775540		Witkowski L et al. Familial rhabdoid tumour 'avant la lettre'--from pathology review to exome sequencing and back again. J Pathol. 2013 Sep;231(1):35-43. http://www.ncbi.nlm.nih.gov/pubmed/23775540

Gene	PMID	Reference Text
SMARCA4	24658001	Ramos P et al. Small cell carcinoma of the ovary, hypercalcemic type, displays frequent inactivating germline and somatic mutations in SMARCA4. <i>Nat Genet.</i> 2014 May;46(5):427-9. http://www.ncbi.nlm.nih.gov/pubmed/24658001
	24658002	Witkowski L et al. Germline and somatic SMARCA4 mutations characterize small cell carcinoma of the ovary, hypercalcemic type. <i>Nat Genet.</i> 2014 May;46(5):438-43. http://www.ncbi.nlm.nih.gov/pubmed/24658002
	25060813	Hasselblatt M et al. SMARCA4-mutated atypical teratoid/rhabdoid tumors are associated with inherited germline alterations and poor prognosis. <i>Acta Neuropathol.</i> 2014 Sep;128(3):453-6. http://www.ncbi.nlm.nih.gov/pubmed/25060813
	26076152	Berchuck A et al. Prophylactic oophorectomy for hereditary small cell carcinoma of the ovary, hypercalcemic type. <i>Gynecol Oncol Rep.</i> 2015 Feb 25;12:20-2. http://www.ncbi.nlm.nih.gov/pubmed/26076152
	26975901	Witkowski L et al. The influence of clinical and genetic factors on patient outcome in small cell carcinoma of the ovary, hypercalcemic type. <i>Gynecol Oncol.</i> 2016 Jun;141(3):454-60. http://www.ncbi.nlm.nih.gov/pubmed/26975901
	27079213	Foulkes WD et al. Rare non-epithelial ovarian neoplasms: Pathology, genetics and treatment. <i>Gynecol Oncol.</i> 2016 Jul;142(1):190-8. http://www.ncbi.nlm.nih.gov/pubmed/27079213
SMARCB1	10739763	Taylor MD et al. Familial posterior fossa brain tumors of infancy secondary to germline mutation of the hSNF5 gene. <i>Am J Hum Genet.</i> 2000 Apr;66(4):1403-6. http://www.ncbi.nlm.nih.gov/pubmed/10739763
	16261613	Janson K et al. Predisposition to atypical teratoid/rhabdoid tumor due to an inherited INI1 mutation. <i>Pediatr Blood Cancer.</i> 2006 Sep;47(3):279-84. http://www.ncbi.nlm.nih.gov/pubmed/16261613
	18087273	Ammerlaan AC et al. Long-term survival and transmission of INI1-mutation via nonpenetrant males in a family with rhabdoid tumour predisposition syndrome. <i>Br J Cancer.</i> 2008 Jan 29;98(2):474-9. http://www.ncbi.nlm.nih.gov/pubmed/18087273
	19124645	Swensen JJ et al. Familial occurrence of schwannomas and malignant rhabdoid tumour associated with a duplication in SMARCB1. <i>J Med Genet.</i> 2009 Jan;46(1):68-72. http://www.ncbi.nlm.nih.gov/pubmed/19124645
	19912265	Hulsebos TJ et al. SMARCB1/INI1 maternal germ line mosaicism in schwannomatosis. <i>Clin Genet.</i> 2010 Jan;77(1):86-91. http://www.ncbi.nlm.nih.gov/pubmed/19912265
	20848638	Bruggers CS et al. Clinicopathologic comparison of familial versus sporadic atypical teratoid/rhabdoid tumors (AT/RT) of the central nervous system. <i>Pediatr Blood Cancer.</i> 2011 Jul 1;56(7):1026-31. http://www.ncbi.nlm.nih.gov/pubmed/20848638
	20930055	Christiaans I et al. Germline SMARCB1 mutation and somatic NF2 mutations in familial multiple meningiomas. <i>J Med Genet.</i> 2011 Feb;48(2):93-7. http://www.ncbi.nlm.nih.gov/pubmed/20930055
	21108436	Eaton KW et al. Spectrum of SMARCB1/INI1 mutations in familial and sporadic rhabdoid tumors. <i>Pediatr Blood Cancer.</i> 2011 Jan;56(1):7-15. http://www.ncbi.nlm.nih.gov/pubmed/21108436
	21208904	Bourdeaut F et al. Frequent hSNF5/INI1 germline mutations in patients with rhabdoid tumor. <i>Clin Cancer Res.</i> 2011 Jan 1;17(1):31-8. http://www.ncbi.nlm.nih.gov/pubmed/21208904
	21210147	Teplick A et al. Educational paper: Screening in cancer predisposition syndromes: guidelines for the general pediatrician. <i>Eur J Pediatr.</i> 2011 Mar;170(3):285-94. http://www.ncbi.nlm.nih.gov/pubmed/21210147
	22082606	Carter JM et al. Epithelioid malignant peripheral nerve sheath tumor arising in a schwannoma, in a patient with "neuroblastoma-like" schwannomatosis and a novel germline SMARCB1 mutation. <i>Am J Surg Pathol.</i> 2012 Jan;36(1):154-60. http://www.ncbi.nlm.nih.gov/pubmed/22082606
	24933152	Smith MJ et al. SMARCB1 mutations in schwannomatosis and genotype correlations with rhabdoid tumors. <i>Cancer Genet.</i> 2014 Sep;207(9):373-8. http://www.ncbi.nlm.nih.gov/pubmed/24933152

Gene	PMID	Reference Text
SMARCB1	25494491	Sredni ST & Tomita T. Rhabdoid tumor predisposition syndrome. <i>Pediatr Dev Pathol.</i> 2015 Jan-Feb;18(1):49-58. http://www.ncbi.nlm.nih.gov/pubmed/25494491
	25857641	Smith MJ. Germline and somatic mutations in meningiomas. <i>Cancer Genet.</i> 2015 Apr;208(4):107-14. http://www.ncbi.nlm.nih.gov/pubmed/25857641
	26001331	Paganini I et al. Broadening the spectrum of SMARCB1-associated malignant tumors: a case of uterine leiomyosarcoma in a patient with schwannomatosis. <i>Hum Pathol.</i> 2015 Aug;46(8):1226-31. http://www.ncbi.nlm.nih.gov/pubmed/26001331
	26342593	Gigante L et al. Rhabdoid tumor predisposition syndrome caused by SMARCB1 constitutional deletion: prenatal detection of new case of recurrence in siblings due to gonadal mosaicism. <i>Fam Cancer.</i> 2016 Jan;15(1):123-6. http://www.ncbi.nlm.nih.gov/pubmed/26342593
	26342709	Wu J et al. Identification of a novel germline SMARCB1 nonsense mutation in a family manifesting both schwannomatosis and unilateral vestibular schwannoma. <i>J Neurooncol.</i> 2015 Nov;125(2):439-41. http://www.ncbi.nlm.nih.gov/pubmed/26342709
	26364901	Gossai N et al. Report of a patient with a constitutional missense mutation in SMARCB1, Coffin-Siris phenotype, and schwannomatosis. <i>Am J Med Genet A.</i> 2015 Dec;167(12):3186-91. http://www.ncbi.nlm.nih.gov/pubmed/26364901
	27921248	Kehrer-Sawatzki H et al. The molecular pathogenesis of schwannomatosis, a paradigm for the co-involvement of multiple tumour suppressor genes in tumorigenesis. <i>Hum Genet.</i> 2016 Dec 5. [Epub ahead of print] http://www.ncbi.nlm.nih.gov/pubmed/27921248
	SPINK1	18414673
24624459		LaRusch J, Solomon S, Whitcomb DC. Pancreatitis Overview. 2014 Mar 13. In: Adam MP, Ardinger HH, Pagon RA, et al., editors. <i>GeneReviews</i> ® [Internet]. Seattle (WA): University of Washington, Seattle; 1993-2017. http://www.ncbi.nlm.nih.gov/pubmed/24624459
STK11	10982753	Pfützer RH et al. SPINK1/PSTI polymorphisms act as disease modifiers in familial and idiopathic chronic pancreatitis. <i>Gastroenterology.</i> 2000 Sep 119(3):615-23. http://www.ncbi.nlm.nih.gov/pubmed/10982753
	11113065	Giardiello FM et al. Very high risk of cancer in familial Peutz-Jeghers syndrome. <i>Gastroenterology.</i> 2000 Dec;119(6):1447-53. (PMID 11113065) http://www.ncbi.nlm.nih.gov/pubmed/11113065
	15121768	Amos CI et al. Genotype-phenotype correlations in Peutz-Jeghers syndrome. <i>J Med Genet.</i> 2004 May;41(5):327-33. http://www.ncbi.nlm.nih.gov/pubmed/15121768
	15502809	Young RH. Sex cord-stromal tumors of the ovary and testis: their similarities and differences with consideration of selected problems. <i>Modern Pathology : An Official Journal Of The United States And Canadian Academy Of Pathology, Inc.</i> 2005 Feb 18 Suppl 2:S81-98. http://www.ncbi.nlm.nih.gov/pubmed/15502809
	16582077 (b)	Hearle NC et al. Exonic STK11 deletions are not a rare cause of Peutz-Jeghers syndrome. <i>J Med Genet.</i> 2006 Apr;43(4):e15. http://www.ncbi.nlm.nih.gov/pubmed/16582077
	16707622 (a)	Hearle N et al. Frequency and spectrum of cancers in the Peutz-Jeghers syndrome. <i>Clin Cancer Res.</i> 2006 May;12(10):3209-15. http://www.ncbi.nlm.nih.gov/pubmed/16707622
	20051941	van Lier MG et al. High cancer risk in Peutz-Jeghers syndrome: a systematic review and surveillance recommendations. <i>The American Journal Of Gastroenterology.</i> 2010 Jun 105(6):1258-64. http://www.ncbi.nlm.nih.gov/pubmed/20051941
	20581245	Beggs AD et al. Peutz-Jeghers syndrome: a systematic review and recommendations for management. <i>Gut.</i> 2010 Jul;59(7):975-86. http://www.ncbi.nlm.nih.gov/pubmed/20581245
	21157440	van Lier MG et al. High cumulative risk of intussusception in patients with Peutz-Jeghers syndrome: time to update surveillance guidelines?. <i>The American Journal Of Gastroenterology.</i> 2011 May 106(5):940-5. http://www.ncbi.nlm.nih.gov/pubmed/21157440
	24179492	Banno K et al. Hereditary gynecological tumors associated with Peutz-Jeghers syndrome (Review). <i>Oncology Letters.</i> 2013 Nov 6(5):1184-1188. http://www.ncbi.nlm.nih.gov/pubmed/24179492

Gene	PMID	Reference Text	
STK11	9429144	Tomlinson IP and Houlston RS. Peutz-Jeghers syndrome. Journal Of Medical Genetics. 1997 Dec 34(12):1007-11. http://www.ncbi.nlm.nih.gov/pubmed/9429144	
	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Colorectal. http://www.nccn.org/clinical.asp	
SUFU	12068298	Taylor MD et al. Mutations in SUFU predispose to medulloblastoma. Nature Genetics. 2002 31(3):306-10. http://www.ncbi.nlm.nih.gov/pubmed/12068298	
	19533801	Pastorino L et al. Identification of a SUFU germline mutation in a family with Gorlin syndrome. American Journal Of Medical Genetics. Part A. 2009 Jul 149A(7):1539-43 http://www.ncbi.nlm.nih.gov/pubmed/19533801	
	19833601	Brugières L et al. Incomplete penetrance of the predisposition to medulloblastoma associated with germ-line SUFU mutations. Journal Of Medical Genetics. 2010 47(2):142-4. http://www.ncbi.nlm.nih.gov/pubmed/19833601	
	20301330	Evans DG, Farndon PA. Nevoid Basal Cell Carcinoma Syndrome. 2002 Jun 20 [Updated 2015 Oct 1]. In: Pagon RA, Adam MP, Ardinger HH, et al., editors. GeneReviews® [Internet]. Seattle (WA): University of Washington, Seattle; 1993-2016. http://www.ncbi.nlm.nih.gov/pubmed/20301330	
	21834049	Bree AF et al. Consensus statement from the first international colloquium on basal cell nevus syndrome (BCNS). Am J Med Genet A. 2011 Sep;155A(9):2091-7. http://www.ncbi.nlm.nih.gov/pubmed/21834049	
	22508808	Brugières L et al. High frequency of germline SUFU mutations in children with desmoplastic/nodular medulloblastoma younger than 3 years of age. Journal Of Clinical Oncology : Official Journal Of The American Society Of Clinical Oncology. 2012 30(17):2087-93. http://www.ncbi.nlm.nih.gov/pubmed/22508808	
	22829011	Kijima C et al. Two cases of nevoid basal cell carcinoma syndrome associated with meningioma caused by a PTCH1 or SUFU germline mutation. Familial Cancer. 2012 Dec 11(4):565-70. http://www.ncbi.nlm.nih.gov/pubmed/22829011	
	24651015	Kool M et al. Genome sequencing of SHH medulloblastoma predicts genotype-related response to smoothened inhibition. Cancer Cell. 2014 Mar 17;25(3):393-405. http://www.ncbi.nlm.nih.gov/pubmed/24651015	
	24659465	Bholah Z et al. Intronic splicing mutations in PTCH1 cause Gorlin syndrome. Familial Cancer. 2014 Sep 13(3):477-80. http://www.ncbi.nlm.nih.gov/pubmed/24659465	
	25287320	Mann K et al. Multiple skin hamartomata: a possible novel clinical presentation of SUFU neoplasia syndrome. Familial Cancer. 2015 Mar 14(1):151-5. http://www.ncbi.nlm.nih.gov/pubmed/25287320	
	25403219	Smith MJ et al. Germline mutations in SUFU cause Gorlin syndrome-associated childhood medulloblastoma and redefine the risk associated with PTCH1 mutations. J Clin Oncol. 2014 Dec 20;32(36):4155-61. http://www.ncbi.nlm.nih.gov/pubmed/25403219	
	TMEM127	20154675	Qin Y et al. Germline mutations in TMEM127 confer susceptibility to pheochromocytoma. Nat Genet. 2010 Mar;42(3):229-33. http://www.ncbi.nlm.nih.gov/pubmed/20154675
		21156949	Yao L et al. Spectrum and prevalence of FP/TMEM127 gene mutations in pheochromocytomas and paragangliomas. JAMA. 2010 Dec 15;304(23):2611-9. http://www.ncbi.nlm.nih.gov/pubmed/21156949
21613359		Neumann HP et al. Germline mutations of the TMEM127 gene in patients with paraganglioma of head and neck and extraadrenal abdominal sites. J Clin Endocrinol Metab. 2011 Aug;96(8):E1279-82. http://www.ncbi.nlm.nih.gov/pubmed/21613359	
24334765		Qin Y et al. The tumor susceptibility gene TMEM127 is mutated in renal cell carcinomas and modulates endolysosomal function. Hum Mol Genet. 2014 May 1;23(9):2428-39. http://www.ncbi.nlm.nih.gov/pubmed/24334765	
24893135		Lenders JW et al. Pheochromocytoma and paraganglioma: an endocrine society clinical practice guideline. The Journal Of Clinical Endocrinology And Metabolism. 2014 Jun 99(6):1915-42.24893135 http://www.ncbi.nlm.nih.gov/pubmed/24893135	
25389632		Toledo SP et al. Penetrance and clinical features of pheochromocytoma in a six-generation family carrying a germline TMEM127 mutation. J Clin Endocrinol Metab. 2015 Feb;100(2):E308-18. http://www.ncbi.nlm.nih.gov/pubmed/25389632	

Gene	PMID	Reference Text
TMEM127	25770152	King EE et al. Integrity of the pheochromocytoma susceptibility TMEM127 gene in patients with pediatric malignancies. <i>Endocr Relat Cancer</i> . 2015 Jun;22(3):L5-7. http://www.ncbi.nlm.nih.gov/pubmed/25770152
	26960314	Patócs A et al. Novel SDHB and TMEM127 Mutations in Patients with Pheochromocytoma/Paraganglioma Syndrome. <i>Pathol Oncol Res</i> . 2016 Oct;22(4):673-9. http://www.ncbi.nlm.nih.gov/pubmed/26960314
TP53	10864200	Chompret A et al. P53 germline mutations in childhood cancers and cancer risk for carrier individuals. <i>Br J Cancer</i> . 2000 Jun;82(12):1932-7. http://www.ncbi.nlm.nih.gov/pubmed/10864200
	14583457	Olivier M et al. Li-Fraumeni and related syndromes: correlation between tumor type, family structure, and TP53 genotype. <i>Cancer Res</i> . 2003 Oct 15;63(20):6643-50. http://www.ncbi.nlm.nih.gov/pubmed/14583457
	18511570	Bougeard G et al. Molecular basis of the Li-Fraumeni syndrome: an update from the French LFS families. <i>J Med Genet</i> . 2008 Aug;45(8):535-8. http://www.ncbi.nlm.nih.gov/pubmed/18511570
	19204208	Gonzalez PD et al. Beyond Li Fraumeni Syndrome: clinical characteristics of families with p53 germline mutations. <i>J Clin Oncol</i> . 2009 Mar 10;27(8):1250-6 http://www.ncbi.nlm.nih.gov/pubmed/19204208
	20522432	Ruijs MWG et al. TP53 germline mutation testing in 180 families suspected of Li Fraumeni syndrome: mutation detection rate and relative frequency of cancers in different familial phenotypes. <i>J Med Genet</i> . 2010 Jun;47(6):421-8. http://www.ncbi.nlm.nih.gov/pubmed/20522432
	23172776	Zerdoumi Y et al. Drastic effect of germline TP53 missense mutations in Li-Fraumeni patients. <i>Hum Mutat</i> . 2013 Mar;34(3):453-61. http://www.ncbi.nlm.nih.gov/pubmed/23172776
	260142920	Bougeard G et al. Revisiting Li-Fraumeni Syndrome From TP53 Mutation Carriers. <i>J Clin Oncol</i> . 2015 Jul 20;33(21):2345-52. http://www.ncbi.nlm.nih.gov/pubmed/26014290
	27496084	Mai PL et al. Risks of first and subsequent cancers among TP53 mutation carriers in the National Cancer Institute Li-Fraumeni syndrome cohort. <i>Cancer</i> . 2016 Aug 6. http://www.ncbi.nlm.nih.gov/pubmed/27496084
	9554443	Hisada M et al. Multiple primary cancers in families with Li-Fraumeni syndrome. <i>J Natl Cancer Inst</i> . 1998 Apr 15;90(8):606-11. http://www.ncbi.nlm.nih.gov/pubmed/9554443
	9764816	Birch JM et al. Cancer phenotype correlates with constitutional TP53 genotype in families with the Li-Fraumeni syndrome. <i>Oncogene</i> . 1998 Sep 3;17(9):1061-8. http://www.ncbi.nlm.nih.gov/pubmed/9764816
	NCCN	NCCN Guidelines. Genetic/Familial High-Risk Assessment: Breast and Ovarian. (http://www.nccn.org/clinical.asp) http://www.nccn.org/clinical.asp
	TSC1	10330349
17304050		Au KS et al. Genotype/phenotype correlation in 325 individuals referred for a diagnosis of tuberous sclerosis complex in the United States. <i>Genet Med</i> . 2007 Feb;9(2):88-100. http://www.ncbi.nlm.nih.gov/pubmed/17304050
24053982		Northrup H et al. Tuberous sclerosis complex diagnostic criteria update: recommendations of the 2012 International Tuberous Sclerosis Complex Consensus Conference. <i>Pediatr Neurol</i> . 2013 Oct;49(4):243-54. http://www.ncbi.nlm.nih.gov/pubmed/24053982
24053983		Krueger DA et al. Tuberous sclerosis complex surveillance and management: recommendations of the 2012 International Tuberous Sclerosis Complex Consensus Conference. <i>Pediatr Neurol</i> . 2013 Oct;49(4):255-65. http://www.ncbi.nlm.nih.gov/pubmed/24053983
24359990		Haas NB and Nathanson KL. Hereditary kidney cancer syndromes. <i>Adv Chronic Kidney Dis</i> . 2014 Jan;21(1):81-90. http://www.ncbi.nlm.nih.gov/pubmed/24359990
TSC2	10330349	Verhoef S et al. High rate of mosaicism in tuberous sclerosis complex. <i>Am J Hum Genet</i> . 1999 Jun;64(6):1632-7. http://www.ncbi.nlm.nih.gov/pubmed/10330349

Gene	PMID	Reference Text
TSC2	17304050	Au KS et al. Genotype/phenotype correlation in 325 individuals referred for a diagnosis of tuberous sclerosis complex in the United States. <i>Genet Med.</i> 2007 Feb;9(2):88-100. http://www.ncbi.nlm.nih.gov/pubmed/17304050
	24053982	Northrup H et al. Tuberous sclerosis complex diagnostic criteria update: recommendations of the 2012 International Tuberous Sclerosis Complex Consensus Conference. <i>Pediatr Neurol.</i> 2013 Oct;49(4):243-54. http://www.ncbi.nlm.nih.gov/pubmed/24053982
	24053983	Krueger DA et al. Tuberous sclerosis complex surveillance and management: recommendations of the 2012 International Tuberous Sclerosis Complex Consensus Conference. <i>Pediatr Neurol.</i> 2013 Oct;49(4):255-65. http://www.ncbi.nlm.nih.gov/pubmed/24053983
	24359990	Haas NB and Nathanson KL. Hereditary kidney cancer syndromes. <i>Adv Chronic Kidney Dis.</i> 2014 Jan;21(1):81-90. http://www.ncbi.nlm.nih.gov/pubmed/24359990
VHL	10205047	Stebbins CE et al. Structure of the VHL-ElonginC-ElonginB complex: implications for VHL tumor suppressor function. <i>Science.</i> 1999 Apr 16;284(5413):455-61. http://www.ncbi.nlm.nih.gov/pubmed/10205047
	11257110	Friedrich CA. Genotype-phenotype correlation in von Hippel-Lindau syndrome. <i>Hum Mol Genet.</i> 2001 Apr;10(7):763-7. http://www.ncbi.nlm.nih.gov/pubmed/11257110
	12202531	Dollfus H et al. Retinal hemangioblastoma in von Hippel-Lindau disease: a clinical and molecular study. <i>Invest Ophthalmol Vis Sci.</i> 2002 Sep;43(9):3067-74. http://www.ncbi.nlm.nih.gov/pubmed/12202531
	12461257	Marcos HB et al. Neuroendocrine tumors of the pancreas in von Hippel-Lindau disease: spectrum of appearances at CT and MR imaging with histopathologic comparison. <i>Radiology.</i> 2002 http://www.ncbi.nlm.nih.gov/pubmed/12461257
	12814730	Lonser RR et al. von Hippel-Lindau disease. <i>Lancet.</i> 2003 Jun;361(9374):2059-67. (PMID 12814730) http://www.ncbi.nlm.nih.gov/pubmed/12814730
	15796386	Kim HJ et al. Tumors of the endolymphatic sac in patients with von Hippel-Lindau disease: implications for their natural history, diagnosis, and treatment. <i>J Neurosurg.</i> 2005 Mar;102(3):503-12. http://www.ncbi.nlm.nih.gov/pubmed/15796386
	17024664	Ong KR et al. Genotype-phenotype correlations in von Hippel-Lindau disease. <i>Hum Mutat.</i> 2007 Feb;28(2):143-9. http://www.ncbi.nlm.nih.gov/pubmed/17024664
	17427103	Mannelli M et al. Genetics and biology of pheochromocytoma. <i>Experimental And Clinical Endocrinology & Diabetes.</i> 2007 115(3):160-5. http://www.ncbi.nlm.nih.gov/pubmed/17427103
	19017755	Jimenez C et al. Use of the tyrosine kinase inhibitor sunitinib in a patient with von Hippel-Lindau disease: targeting angiogenic factors in pheochromocytoma and other von Hippel-Lindau disease-related tumors. <i>J Clin Endocrinol Metab.</i> 2009 Feb;94(2):386-9 http://www.ncbi.nlm.nih.gov/pubmed/19017755
	20301636	Frantzen C et al. Von Hippel-Lindau Disease. (2000 [updated 2012]). In Pagon RA, Adam MP, Bird TD, Dolan CR, Fong CT, and Stephens K (Eds), <i>GeneReviews</i> [Internet]. Seattle (WA): University of Washington, Seattle. http://www.ncbi.nlm.nih.gov/pubmed/20301636
	21386872	Maher ER, Neumann HP, and Richard S. von Hippel-Lindau disease: a clinical and scientific review. <i>Eur J Hum Genet.</i> 2011 Jun;19(6):617-23. (PMID 21386872) http://www.ncbi.nlm.nih.gov/pubmed/21386872
	WT1	15150775
16199712		Fischbach BV et al. WAGR syndrome: a clinical review of 54 cases. <i>Pediatrics.</i> 2005 Oct 116(4):984-8. http://www.ncbi.nlm.nih.gov/pubmed/16199712
16857697		Scott RH et al. Surveillance for Wilms tumour in at-risk children: pragmatic recommendations for best practice. <i>Archives Of Disease In Childhood.</i> 2006 Dec 91(12):995-9. http://www.ncbi.nlm.nih.gov/pubmed/16857697
18618575		Royer-Pokora B et al. Clinical relevance of mutations in the Wilms tumor suppressor 1 gene WT1 and the cadherin-associated protein beta1 gene CTNBN1 for patients with Wilms tumors: results of long-term surveillance of 71 patients from International Society of Pediatric Oncology Study 9/Society for Pediatric Oncology. <i>Cancer.</i> 2008 Sep 1 113(5):1080-9. http://www.ncbi.nlm.nih.gov/pubmed/18618575

Gene	PMID	Reference Text
WT1	18816692	Auber F et al. Management of Wilms tumors in Drash and Frasier syndromes. <i>Pediatric Blood & Cancer</i> . 2009 Jan 52(1):55-9. http://www.ncbi.nlm.nih.gov/pubmed/18816692
	20301471	Dome JS, Huff V. Wilms Tumor Predisposition. 2003 Dec 19 [Updated 2016 Oct 20]. In: Pagon RA, Adam MP, Ardinger HH, et al., editors. <i>GeneReviews®</i> [Internet]. Seattle (WA): University of Washington, Seattle; 1993-2016. Available from: https://www.ncbi.nlm.nih.gov/books/NBK1294/ http://www.ncbi.nlm.nih.gov/books/NBK1294/
	25623218	Ezaki J et al. Gonadal tumor in Frasier syndrome: a review and classification. <i>Cancer Prevention Research (Philadelphia, Pa.)</i> . 2015 Apr 8(4):271-6. http://www.ncbi.nlm.nih.gov/pubmed/25623218
	25688735	Kaneko Y et al. A high incidence of WT1 abnormality in bilateral Wilms tumours in Japan, and the penetrance rates in children with WT1 germline mutation. <i>British Journal Of Cancer</i> . 2015 Mar 17 112(6):1121-33. http://www.ncbi.nlm.nih.gov/pubmed/25688735
	2984395	Eddy AA and Mauer SM. Pseudohermaphroditism, glomerulopathy, and Wilms tumor (Drash syndrome): frequency in end-stage renal failure. <i>The Journal Of Pediatrics</i> . 1985 Apr 106(4):584-7. http://www.ncbi.nlm.nih.gov/pubmed/2984395
	3039839	Dao DD et al. Genetic mechanisms of tumor-specific loss of 11p DNA sequences in Wilms tumor. <i>American Journal Of Human Genetics</i> . 1987 Aug 41(2):202-17. http://www.ncbi.nlm.nih.gov/pubmed/3039839
	8071974	Mueller RF. The Denys-Drash syndrome. <i>Journal Of Medical Genetics</i> . 1994 Jun 31(6):471-7. http://www.ncbi.nlm.nih.gov/pubmed/8071974
	9398852	Barbaux S et al. Donor splice-site mutations in WT1 are responsible for Frasier syndrome. <i>Nature Genetics</i> . 1997 17(4):467-70. http://www.ncbi.nlm.nih.gov/pubmed/9398852



207 Perry Parkway
Gaithersburg, MD 20877
T 1 888 729 1206 (Toll-free), 1 301 519 2100 • **F** 1 201 421 2010
E zebras@genedx.com • www.genedx.com

