Practical Hereditary Gynecologic Cancers

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Passenger Mutations and Driver Mutations



Accumulation of Driver and Passenger mutations

Passengers and Drivers

Driver mutations

Passenger mutations

- Contribute to oncogenesis; provide growth advantage; selected for in microniche
- Occur in
 - oncogenes (gain of function mutations) or
 - tumour-suppressor genes (loss of function mutations)

- Neutral mutations
- Carried along for the ride
 - somatic mutations without functional consequences often occur during cell division
- PLUS huge increase in mutation rate with loss of genome repair mechanisms



Proto-oncogene to oncogene

1st mutation **(leads** to accelerated cell division)

Tumor suppressor genes Active oncogene 1st mutation (susceptible carrier)

2nd mutation or loss deads

to cancer}





UREU

Active oncogene

http://cisncancer.org/



Cancer Risk is increased

KNOW THE RED FLAGS ASSOCIATED WITH HEREDITARY CANCER

An individual with a personal or family history of any ONE of the following:

MULTIPLE CANCERS A combination of cancers on the same side of the family	 2 or more: breast / ovarian / prostate / pancreatic cancer 2 or more: colorectal / endometrial / ovarian / gastric / pancreatic / other cancers (i.e., ureter/renal pelvis, billary tract, small bowel, brain, sebaceous adenomas) 2 or more: melanoma / pancreatic cancer
YOUNG CANCERS Any 1 of the following cancers at age 50 or younger	Breast cancer Colorectal cancer Endometrial cancer
RARE CANCERS Any 1 of these rare presentations at any age	 Ovarian cancer Breast: male breast cancer or triple negative breast cancer Colorectal cancer with abnormal MSI/IHC, MSI associated histology" Endometrial cancer with abnormal MSI/IHC 10 or more gastrointestinal polyps*

Certain ancestries may have greater risk for hereditary cancer syndromes (e.g., Ashkenazi Jewish ancestry)

Assessment onteria based on medical society guidelines. For these individual medical society guidelines, go to www.MyriadPro.com/guidelines. Family members include first-, second-, and third-degree blood relatives on both your mother's and father's sides.

"Aderomations type. "Presence of tumor infiltrating tymphocytes. Only's likely mphocytic maction, much each type tring different attion, or medullary growth pattern

Hereditary Gynecologic Cancer Syndromes

- 1. Hereditary Breast/Ovarian Cancer Syndrome (HBOC)
- 2. Hereditary Site-Specific Ovarian Cancer Syndrome
- 3. Hereditary Nonpolyposis Colon Cancer Syndrome (HNPCC or Lynch II)

- Cowden disease
- DICER1 syndrome
- Peutz-Jeghers syndrome
- Hereditary diffuse gastric cancer syndrome
- Li-Fraumeni syndrome
- Small cell CA of ovary, hypercalcemia type (SCCOHT)
- Neurofibromatosis type 1

Rare Causes of Hereditary Gynecologic Cancer

- Breast cancer
- Male breast cancer
- Pancreatic cancer
- Prostate cancer
- Laryngeal cancer
- Colon cancer
- Endometrial cancer
- Urinary tract cancer
- GI tract cancer
- Sarcoma
- Brain cancer
- Leukemia





Ovarian cancer- associated cancers

Cancer Type	General Population Risk	Lynch Syndrome (MLH1 and MSH2 heterozygotes)	
		Risk	Mean Age of Onset
Colon	5.5%	52%-82%	44-61 years
Endometrium	2.7%	25%-60%	48-62 years
Stomach	<1%	6%-13%	56 years
Ovary	1.6%	4 <mark>%-12</mark> %	42.5 years
Hepatobiliary tract	<1%	1.4%-4%%	Not reported
Urinary tract	<1%	1%-4%	~55 years
Small bowel	<1%	3%-6%	49 years
Brain/central nervous system	<1%	1%-3%	~50 years
Sebaceous neoplasms	<1%	1%-9%	Not reported



MSI TESTING

- Identify MMR proteins
- Normally present
- If protein is absent, gene is not being expressed (mutation or methylation)
- Helps direct gene testing by predicting likely involved gene
- If abnormal IHC (absent), MSI+



MLH1





IHC FOR MMR PROTEIN







-Familial breast/ovarian cancer: Multiple shared genes and environmental factors

BRCA1/BRCA2 mutations in ovarian cancer (UW, Seattle, USA)

Ovarian cancer: BRCA1/BRCA2 mutations in 63/360 (18%) patients not selected for family history or age at onset



Courtesy of MC King, UW

DIFFERENT HISTOLOGICAL SUBTYPES OF OVARIAN CANCER INDICATE DIFFERENT THERAPEUTIC OPPORTUNITIES



Low grade serous KRAS

Clear Cell PIK3CA mutations

Endometrioid ARID1A mutation and deletion

Mucinous Probably metastatic colon

High grade serous p53, BRCA1/2 copy number long tail of actionable mutations

Targeted agents Bevaczumib PARP inhibitors have now been FDA approved



Kurman and Shih, Am J Pathol 2016

Ovarian cancer paradox

- More than 90% of BRCA carrier who became symptomatic had ovarian CA
- More than 70-80% of SO specimen from women opted for prophylaxis had FT cancer



Cancer Profiling

Progression-Free Survival



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ESENTED AT: ASCO Annual 15 Meeting

Presented By Dung Le at 2015 ASCO Annual Meeting

MSS vs MSI survival CRC



Nature Clin Prac Oncol 2008



OVARIAN CANCER EXPRESSION PROFILING





Jarboe et al. Pathology of BRCA related ovarian cancer. 2008



- To whom the test should be offered ?
- Which test to offer ?
- When and How to offer DNA test ?
- What are pros and cons of DNA testing in each patient ?
- What to do once the test result is available ?

Translating DNA test to Clinics

• DIANGOSTIC TESTING IN AFFECTED MEMBER

• DIAGNOSTIC TESTING IN ASYMPTOMATIC MEMBER

The Use of Genetic Testing in Ovarian CA







predictive genetic testing

tests in people who are well

To whom the test should be offered ?





Data from Myriad Inc.

- GERMLINE MUTATION TESTING
 - SINGLE GENE TESTING
 - NGS PANEL GENE TESTING
 - FOUNDER MUTATION TESTING
- SOMATIC MUTATION TESTING
 - SINGLE GENE TESTING
 - NGS PANEL TESTING

Which Test to Offer ?

Evolution of Hereditary Cancer Risk Testing... from 2 Genes to Multi-Gene Panel Testing

Early 90's

-BRCA 1 and 2 Discovered by several scientists (Mary Claire King)

-Myriad granted a patent on the BRCA 1-2 Genes and has monopoly on market

-Myriad launched BRCAnalysis, a predictive medicine product for hereditary breast and ovarian cancer (testing BRCA 1-2) in 1996

-Supreme Court Overturns Myriad BRCA 1-2 Patent in June 2013 opening market to other labs (Ambry, GeneDx, LabCorp, Quest, etc)

-Move FROM just testing for BRCA 1&2 to Multiple Gene Testing









Pathogenic variant



- Variant of uncertain significance (VUS)
- Likely benign variant



Benign variant

VARIANT CLASSIFICATION RESULT

NCCN NCCN Network*

Comprehensive NCCN Guidelines Version 1.2017 Cancer Genetic/Familial High-Risk Assessment: Breast and Ovarian

NCCN Guidelines Index Table of Contents Discussion

BREAST AND OVARIAN MANAGEMENT BASED ON GENETIC TEST RESULTS^a

The inclusion of a gene on this table below does not imply the endorsement either for or against multi-gene testing for moderatepenetrance genes.

Gene	Breast Cancer Risk and Management	Ovarian Cancer Risk and Management	Other Cancer Risks and Management		
PALB2	Increased risk of BC • Screening: Annual mammogram and consider breast MRI with contrast at 30 y • RRM: Consider based on family history.	Unknown or insufficient evidence for OC risk	Unknown or insufficient evidence		
	Comments: Counsel for risk of autosomal recessive condition in offspring.				
PTEN	Increased risk of BC See Cowden Syndrome Management	No increased risk of OC	See Cowden Syndrome Management		
	Unknown or insufficient evidence for BC risk	Increased risk of OC • Consider RRSO at 45-50 y	N/A		
RAD51C	Comments: Counsel for risk of autosomal rece cancer in carriers of mutations in RAD51C app firm recommendation as to the optimal age for held around age 45–50 y or earlier based on a	ssive condition in offspring. Based on estimates fr ears to be sufficient to justify consideration of RRS this procedure. Based on the current, limited evide specific family history of an earlier onset ovarian of	om available studies, the lifetime risk of ovarian SO. The current evidence is insufficient to make a ence base, a discussion about surgery should be cancer.		
	Unknown or insufficient evidence for BC risk	Increased risk of OC • Consider RRSO at 45–50 y	N/A		
RAD51D	Comments: Based on estimates from available to justify consideration of RRSO. The current e on the current, limited evidence base, a discus an earlier onset ovarian cancer.	studies, the lifetime risk of ovarian cancer in carri vidence is insufficient to make a firm recommenda sion about surgery should be held around age 45-	ers of mutations in <i>RAD51D</i> appears to be sufficient tion as to the optimal age for this procedure. Based -50 y or earlier based on a specific family history of		
	Increased risk of BC • Screening: See NCCN Guidelines for	Increased risk of non-epithelial OC	See NCON Quidelines for Cenetic/Esmilial High		
STK11	Genetic/Familial High-Risk Assessment: <u>Colorectal</u> RRM: Evidence insufficient, manage based on family history.	High-Risk Assessment: Colorectal	Risk Assessment: Colorectal		
STK11 TP53	Genetic/Familial High-Risk Assessment: <u>Colorectal</u> • RRM: Evidence insufficient, manage based on family history. Increased risk of BC • See Li-Fraumeni Syndrome Management	See Nocing Guidelines for Generic'r aninal. High-Risk Assessment: Colorectal No increased risk of OC	See Li-Fraumeni Syndrome Management		

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- OF 24 DISORDERS THAT ARE DEEM TO BE APPROPRIATE FOR REPORTING INCIDENTAL VARIANT FINDING FROM EXOME SEQUENCING, 16 ARE HEREDITARY CANCER SYNDROMES
- HBOC, FAP, LFS, PJS, LS, MYHAP, VHL, MEN1, MEN2, FMTC, PTEN-HAMARTOMA TUMOR S, RETINOBLASTOMA, HEREDITARY
 PARAGANGLIOMA/PHEOCHROMOCYTOMA, TS, T1-WILMS TUMOR

ACMG RECOMMENDATION FOR REPORTING OF INCIDENTAL FINDING IN CLINICAL EXOME AND GENOME SEQUENCING 2013

 The process by which a counsellor provides relevant information (informative) to counselee about disease burden, etiology, and investigative and therapeutic options (advocacy) in order for the counselee to be able to make informed decision regarding his or her own health

Genetic counseling

- Valid medical evaluation and test result
- Understandable information
- Clear layout of pros ad cons of each options
- Ability of counselee to make unbiased non-directive choices
- Co-ordinated care path
- Good follow up plan

Factor influencing Successful genetic counseling

- First genetic evaluation
 - Pedigree construction, Risk estimation
- Pre-test counselling
 - Pros and cons of undergoing DNA testing
- Post-test counselling
 - Result and prophylactic options

Steps of genetic counselling for ovarian cancer

- 3 generation pedigree
- History of relevant cancer : ovarian, breast, prostate, pancreatic, melanoma, colon
- Patient perception regarding the likelihood of her cancer being hereditary
- Patient readiness to discuss about genetics
- Initial opinion regarding the risk of "being hereditary"

Obtaining and giving information in the first visit

- When and How to offer DNA test ?
- To whom the test should be offered ?
- Which test to offer ?
- What are pros and cons of DNA testing in each patient?
- What to do once the test result is available ?

Pre-test counseling consideration

- When : Suspicion by family history Suspicion by personal history Indication for target therapy Population testing
- How : By oncologist with or without detailed counseling
 By clinical geneticist/ counselor with detailed counseling

When and How to offer DNA testing ?

- DNA test guided target therapy : Olaparib
- Synthetic lethality concept
- Somatic (tumor) vs Germline (blood) testing
- Patients with positive result of somatic testing may need to undergo germline testing
- Offering somatic testing first have benefits over germline first

Special circumstance counseling





positive



Test	Use	Strengths	Limitations
Capillary (Sanger) sequencing	Sequencing of small genomic regions, e.g. individual exons	Highly accurate	Low throughput, labour intensive, expensive
Panel testing using next-generation sequencing	Simultaneous sequencing of genes causing a particular phenotype (up to several hundred genes)	Allows multipanel gene testing Useful in heterogeneous conditions	Needs adjusting when new genes are discovered, and coverage of each gene may not be as good as capillary sequencing
Array CGH	Detection of large structural chromosome rearrangements	Highly accurate, high throughput	
Exome sequencing	Simultaneous sequencing of all coding regions of the genome	Streamlines lab workflow and useful extension of the panel test	Coverage of some genes is inadequate, no information on structural rearrangements
Genome sequencing	Sequencing of the whole genome	More even coverage of all genes	Expensive, data storage and analysis costs are high, and non-coding regions hard to interpret

LITY	HIGH RISK	 Penetrance: high; causes a well known cancer syndrome with well defined cancer risks by site (i.e., <i>BRCA1/2</i> and hereditary breast and ovarian cancer syndrome) Actionability: high; evidence based risk reducing national guidelines exist for at least one organ system (i.e., Tamoxifen therapy/salpingo-oophorectomy for <i>BRCA1/BRCA2</i> carriers (23), colectomy for <i>APC</i> carriers (18) Implications for other family members: straightforward
NICAL UTI	MODERATE RISK	 Penetrance: moderate; organ specific cancer risks are fairly well defined for at least one cancer site(i.e., ATM causes an increased risk for breast cancer, however, pancreatic risks remain unclear) Actionability: moderate; enough evidence exists to supersede empiric risk (if necessary) for enhanced surveillance for at least one at risk site (i.e., enhanced breast cancer surveillance for PALB2 carriers is justified even in the absence of a family history of breast cancer (54) Implications for other family members: may not be straightforward
CLII	LOW RISK	 Penetrance: low or uncertain; vague organ specific cancer risks (i.e., MRE11A carriers have currently unclear organ specific cancer risks) Actionability: low; due to lack of established evidence based guidelines. Screening and management recommendations are provided based on empiric risk estimates and case-by-case literature and laboratory data review Implications for other family members: not well defined

High risk (odds^{θ} >5.0) Low risk (<2.0 odds^{θ} >1.0 or Moderate risk (>2.0 odds⁶ <5.0) Cancer site growing evidence of association) Breast (female) BRCA1 (20), BRCA2 (20), CDH1 (21), PTEN ATM (26, 27), BRIP1 (28), BAP1 (33), BARD1 (34, 35), RAD50 (36, 37), (22), STK11 (23, 24), TP53 (25) CHEK2 (29, 30), PALB2 (31, 32) RAD51C (38), RAD51D (39, 40), MRE11A (36), MUTYH (41), NBN (42, 43), XRCC2 (44, 45) Colorectal APC (46), BMPR1A (47), [¥]EPCAM (48), MLH1 CHEK2 (54, 55), PTEN (56), CDH1 (57, 58), EXO1 (59), GALNT12 (60, 61), (49), MSH2 (49), MSH6 (49, 50), *MUTYH (51), MUTYH (62, 63), POLD1 (64), POLE (64) TP53 (25) PMS2 (52), SMAD4 (47), STK11 (53) BRCA1 (65), BRCA2 (65), MLH1 (66), MSH2 MSH6 (66), PALB2 (32, 65), BARD1 (65, 68), BRIP1 (65), CHEK2 (65), MRE11A Ovary (66), STK11 (24) RAD51C (65, 67), RAD51D (39) (65), MUTYH (69), NBN (65), RAD50 (65), TP53 (65)

Due to study design variation, genetic risk categorization was extrapolated from odds ratios, relative risks, cumulative, or absolute cancer risks and presented as an estimate of the generalized odds (θ) over the baseline population for organ specific cancer risk. Genes in each category are in alphabetical order. Please see individual key reference for specific risk estimate method used. When study discrepancy, or wide reported confidence intervals were reported, expert opinion was used for the final risk categorization. The list is not exhaustive for breast, colorectal, and ovarian cancer predisposition. More studies, especially on moderate and low risk category genes will be needed to better clarify the associated cancer risks and penetrance. Single nucleotide polymorphism studies, which could add hundreds of gene and locus associations to the low risk category, were not included (70). Penetrance and expressivity can widely vary with specific mutations. Asterisk (*) denotes MUTYH biallelic mutation. (*) denotes deletions only affecting transcription of MSH2.

RELATIVE RISK GENE PANEL TESTING

Positive information

• Negative information

Ability to know Availability of prophylaxis Ability to make autonomous health choice Decrease uncertainty Increase anxiety Survivor guilt Potential discrimination Unwanted medical certainty Secondary patients in family

Pros and Cons of DNA testing

- Detailed upfront counseling strategy : follow set plan for personal treatment and offer pre-test presymptomatic counseling to at risk family members
- Limited upfront counseling strategy : offer detailed counseling about germline mutation status and make plan according to client's wishes

What to do once the test result is available ?

Mutation positive : Increased risk (life time risk OvCa 50%, BrCa 70%)

Increased risk of recurrence Prophylactic measures are available Presymptomatic test is available for family members

 Mutation negative : non-familial cancer -> risk is not significantly increased

Familial cancer -> risk may be increased

• VUS found : risk can not be accurately quantified by test alone but may be increased by family history

Post-test counseling

 Ov Ca : TAH/BSO -> 90%, BSO -> 90%, BO -> no reduction TM -> no reduction OCP -> 50% (>3-5 yrs use) but increase BrCa risk
 Br Ca : TM -> 99%-100%, Nipple sparing reconstruction -> 95% TAH/BSO, BSO -> 50% Tamoxifen /Raloxifene -> 50% but increase endometrial Ca risk

Risk reduction strategy

- Molecular genetics of Ovarian Cancer is much better understood
- Causative single genes as well as risk alleles are being increasingly indentified albeit not all with actionable guidelines
- Panel gene testing is likely to be more useful than single gene testing
- Molecular profiling can lead to better individualized targeted therapy
- How to translate DNA testing into clinics needs to be carefully considered

Conclusion

- Counseling for BRCA testing must be done by oncologist or clinical geneticist familiar with benefit as well as limitation of test
- Panel gene testing is likely to be more useful than single gene testing in usual scenario with exception of target therapy decision
- Somatic tumor testing is preferred for target therapy decision and could be given in conjuction with limited genetic counseling
- Germline blood testing is preferred for familial Br/Ov cancer patients

Conclusion