

Disclosures

• I am a consultant for Color Genomics.

CW

Learning objectives

- 1. Recognize key features of common hereditary cancer syndromes
- 2. Evaluate your patient population for appropriate genetic counseling and testing referrals
- 3. Identify patients who may benefit from therapeutic agents linked to genetic markers
- 4. Determine when PRS and RNA testing can add clinical utility

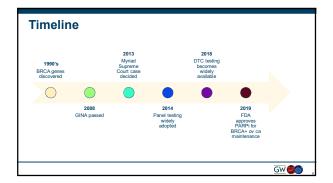
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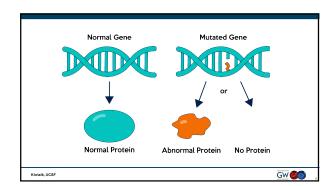
Why is this important Targeted Therapeutics Treatment decisions can and should be influenced by genetic testing results Prevention Secondary malignancies Cancers in family members

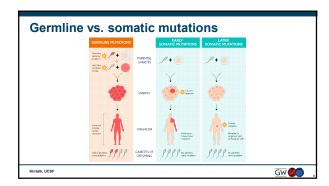
Every breast or ovarian cancer patient with a BRCA1 or BRCA2 mutation detected after diagnosis is a missed opportunity to prevent cancer.

No woman with a mutation in BRCA1 or BRCA2 should die of breast or ovarian cancer.

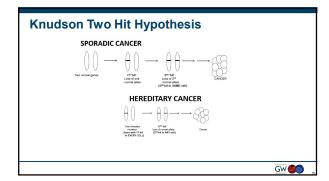
– Dr. Mary-Claire King







Different types of genetic testing	
Somatic Looks at <u>tumor</u> genetics only All tumors have genetic changes Primarily used to identify therapeutic targets, recurrence risk Most somatic mutations identified are <u>not</u> germline, BUT May suggest a higher risk of having a germline mutation	
Germline Present and constant from the time of conception Determines risk of developing a cancer (or a NEW primary) Now being used to identify therapeutic targets	
SNVs Single nucleotide variants (formerly polymorphism) Germline differences – the spice of life! Can have large impacts cumulatively Some are common (SNPs, >1% of population) and some are more rare	



Basic principles of hereditary cancer - About 15% of cancer is hereditary (higher for rare cancers) - Autosomal dominant or autosomal recessive inheritance - Men and women have the same chance to inherit a mutation - Each child is at risk to inherit the altered gene - Mutations can be inherited from either parent - Risk factor – not a diagnosis - Testing of affected family members is the most informative

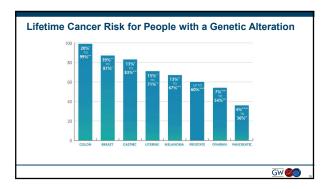
When to suspect a hereditary cancer syndrome

- Cancer in 2 or more family members (same side of family)
- Cancer diagnosis at early age
- Multiple primary tumors in the same individual
- Bilateral or multiple rare cancers
- Constellation of tumors consistent with hereditary cancer syndrome (eg, Lynch)
- Multiple affected generations
- Specific tumor histologies

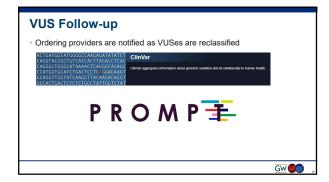






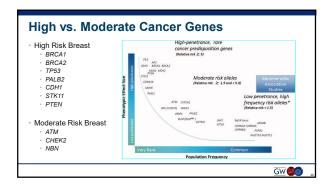


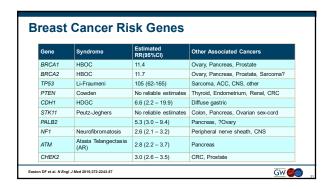
Positive Actionable Some modifications for a particularly strong family history Includes Pathogenic and Likely Pathogenic Follow NCCN guidelines Negative Does not always mean it's not genetic! True negative vs. uninformative negative Includes Benign and Likely Benign. Will not be reported on. Follow based on family history Variant of Uncertain Significance Vast majority are benign differences Lab needs more time Follow based on family history

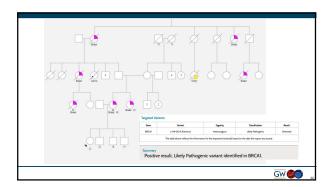


VUS Rates Falling Introduction of RNA testing is helping to reclassify VUSes, especially splice site variants RNA testing also helping to identify pathogenic intronic variants not typically captured by DNA alone Functional modeling harnesses machine learning to combine multiple lines of evidence for better predictions If a mutation is found or reclassified, implications are the same. Just another way to identify an underlying familial cancer syndrome Reclassification from pathogenic to VUS or Benign is VERY rare

Where we are now Testing is performed the same day Saliva or Blood Majority of patients are getting multi-gene panel testing Insurance coverage is good, self-pay prices reasonable (\$250) Results available in 2-3 weeks 7-10 days for some genes Cancer genetics is about prevention and targeted therapies





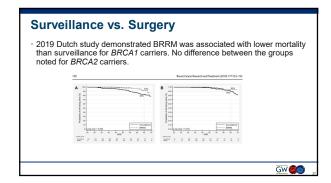


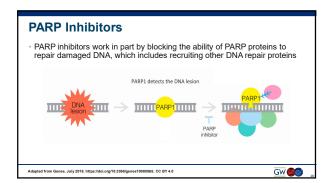


Cancer Sites [6-8,12,62,169]	BRCA1			BRCA2	
	Strength of Evidence	Magnitude of Absolute Risk	Strength of Evidence	Magnitude of Absolute Risk	
Breast (female)	***	High	***	High	
Ovary, falloplan tube, peritoneum	***	High	***	Moderate	
Breast (male)	+	Undefined	***	Low	
Pancreas	**	Very Low	***	Low	
Prostate ^a		Undefined	***	High	

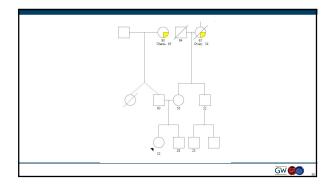


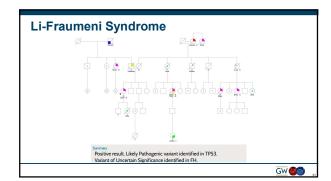






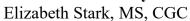
PARPi in Breast Cancer	
FDA Approved Metastatic breast cancer BRCA1 or BRCA2 positive- germline only HER2-negative	
Clinical Trials Includes other genes (PALB2, ATM, CHEK2, RAD51, BRIP1, NBN) Includes somatic BRCA mutations	
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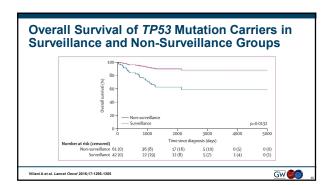


TP53 Associated (Cancers	
Primary Cancers	Also seen	
 Breast cancer 	 Lung 	
• ACC	 Melanoma 	
 Osteosarcoma 	 Thyroid cancer 	
 Brain/CNS 	 GI Tumors 	
 Leukemia 	 Kidney 	
	Gonadal germ cell	
Risk is appx 50% by age 30, 90% by a	ge 60. Higher for women than men.	
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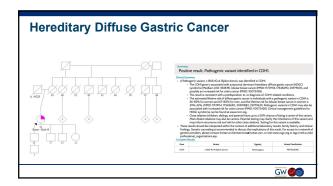
Confirmation of Result Becoming standard practice to confirm germline presence of *TP53* through confirmatory testing of a second cell line Can be confirmed on punch biopsy or eyebrow pluck Allele fraction can give a clue (looking for about 50%) Especially important if the case/fam hx doesn't look typical for *TP53*High de novo rate for *TP53*Other possibilities include CHIP, chemo effect, or mosaicism LiFT UP Study through DFCI and COH Important implications for care of patient and family members



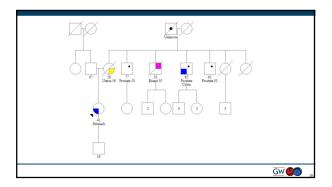




LFS Considerations Testing children is appropriate Breast cancers are more likely to be triple positive Whole body MRI preferred, can do breast, brain separately otherwise Radiation sensitivity With breast cancer, radiation therapy should be avoided if possible. Treat the cancer in front of you, even if there is an increased risk of developing a second malignancy



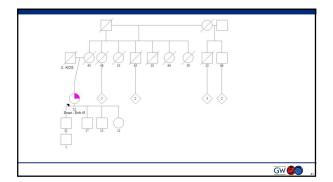


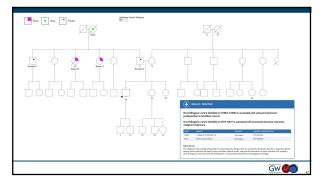


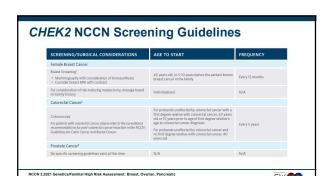
CDH1 Considerations

- Testing children is not recommended unless there is a family history of DGC at 25 or younger
- Cleft lip and palate can also be seen
- Screening studies are ongoing, but have not yet been shown to an effective means of detecting early stage

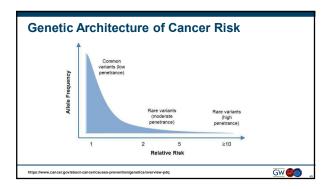
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Transgender Care	
General recommendations are to start screening mammogram at age 50, and 5-10 years after start of feminizing hormones Consider personalization based on strength of family history and personal risk factors Absence of data	
* www.transcare.ucsf.edu/guidelines	
UCSI Transporter Care, Department of Earshy and Community Medicine, University of Celfornia San Francisca. Guidelines for the Rinnary and Gender-Affirming Care of Transporder and Gender Nombionry Prospic 2nd edition. Deviatch Mill, ed. June 2016.	_

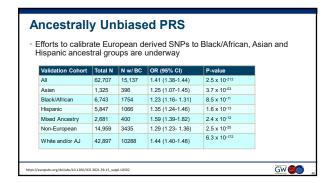


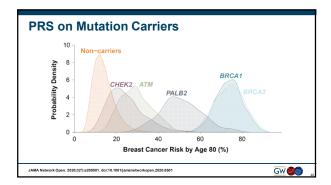
Polygenic Risk Scores

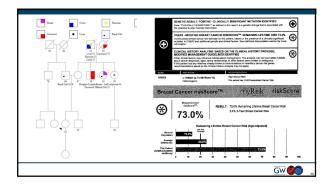
- Use data from GWAS (genome-wide association studies), large scale genetic analyses which focus on uncovering common DNA differences that influence disease and physical traits
- · Each variant has modest effect on risk
- · Combinations of variants can pose more substantial risk
- · Predicts risk in unaffected individuals
- Currently clinically available for breast cancer risk and for CHEK2 risk stratification for European and Ashkenazi Jewish ancestry



GWAS Genome Wide Associate Studies scan the entire genome looking for differences between the control group and the experimental group Currently 80% of participants are of European (non Ashkenazi) ancestry Beatis for many new advances Need information on underrepresented populations: Black, Hispanic, Asian to be truly informative and generalizable All of Us Study from NIH hoping to help address this issue







Fanconi Anemia Gene	es and Breast Cancer Risk
High-Risk Genes BRCA1 (FANCS) BRCA2 (FANCD1) PALB2 (FANCN)	16 genes associated with FA 6 have breast cancer risk implications Each of these accounts for 3% or less of all pathogenic variants
Moderate-Risk Genes BRIP1 (FANCJ) FANCD2 RAD51C (FANCO)	,
https://www.cancer.gov/types/breast/hp/breast-ovarian-genetics-pdq	GW ◎

Ovarian Cancer One of the most heritable cancers, 18% due to a mutation (Norquist, 2016) BRCA1, BRCA2, BRIP1, Lynch syndrome, RAD51C, RAD51D PALB2 under investigation

- BRCA carriers receive greatest benefit from PARPi Don't forget somatic mutation carriers!
- All epithelial ovarian cancer patients are candidates for genetic testing regardless of age or family history
- WISP trial available for carriers



Gene	Ovarian Cancer Risk	RRSO Rec Age	Other Cancer Risk
ATM	<3%	Manage based on fam hx	Breast, Pancreas
BRIP1	>10%	45-50	Breast (TNBC)
RAD51C	>10%	45-50	Breast (TNBC)
RAD51D	>10%	45-50	
BRCA1	39-58%	35-40	Breast
BRCA2	13-29%	40-45	Breast
MLH1/MSH2	>10%	Individualized	Colon, Uterine
MSH6	>10%	Insufficient data	Colon, Uterine
PMS2	<3%	Insufficient data	Colon, Uterine
PALB2	3-5%	Manage based on fam hx	Breast, Pancreas
STK11	>10% (non epithelial)	Not recommended	Colon, Pancreas, Stomach, Breast

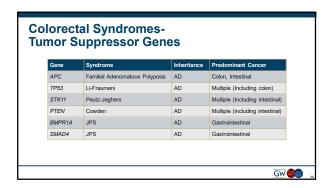
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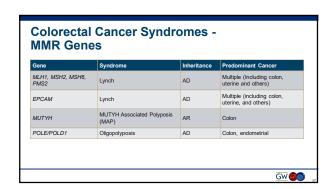
Negative Germline?

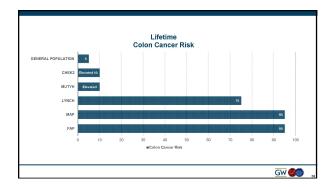
- It's not over yet...
 - 3% of breast cancers harbor BRCA somatic mutations
 - 7-8% of ovarian cancers harbor BRCA somatic mutations

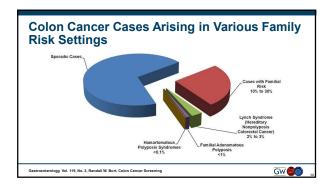
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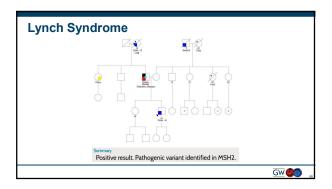
PARPi in Ovarian Cancer • Stage II-IV • Approved as first line maintenance for women with a complete or partial response to platinum chemotherapy • Stronger responses in those who are BRCA+

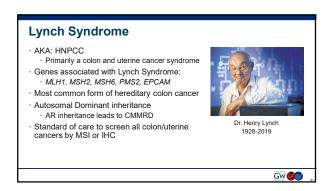


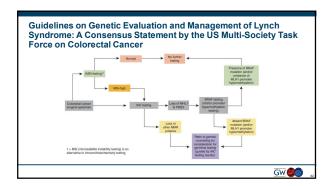


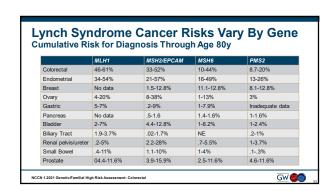


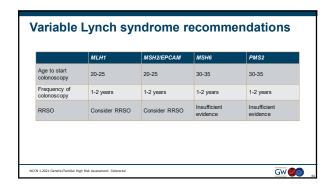






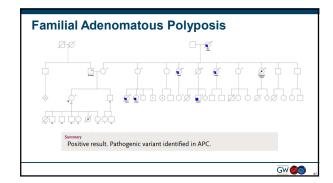




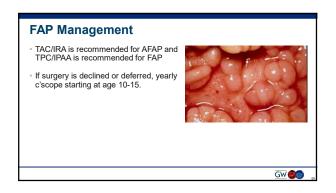


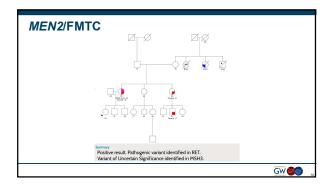
CMMRD	
Constitutional Mismatch Repair Deficiency When an individual has biallelic mutations in an MMR gene (AR Inheritance)	е)
Childhood cancer syndrome 16 fold increased risk in developing cancer	
• Heme, CNS, Colon cancers with risks starting at age 1	
Derm features overlapping with NF	
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Polyposis Syndromes Familial Adenomatous Polyposis (FAP): APC • AD inheritance • Classic: 100s-1000s of polyps • Attenuated (AFAP): 10-100 polyps (Avg 30) • 30% de novo rate for classic • Upper GI, CHRPE, desmoids, osteomas, thyroid, brain • APC I1307K = low risk mutation (AJ) MUTYH Associated Polyposis (MAP): MUTYH • AR inheritance • A single mutation slightly increases risk POLE/POLD1 • AD Inheritance

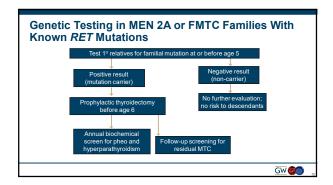


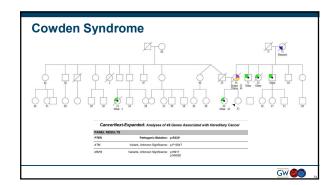
Type of Malignancy	Lifetime Risk
Small bowel: duodenum or periampulla	4-12%
Thyroid: Papillary thyroid cancer	1-2% (higher in women)
CNS: Medulloblastoma	<1%
Liver: Hepatoblastoma	1-2%
Pancreatic Adenocarcinoma	1%
Bile Duct Adenocarcinoma	<1%
Gastric Adenocarcinoma	<1% in Western cultures





MEN2 Clinical Features Medullary thyroid cancer MEN2A Pheochromocytoma, PC (50%) Parathyroid hyperplasia, HPT (15-30%) MEN 2B Pheochromocytoma (50%) Developmental Abnormalities: Ganglioneuromatosis, Mucosal Neuromas, Marfanoid Phenotype

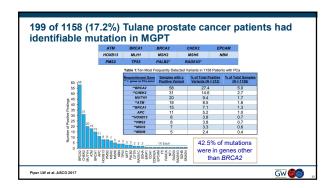


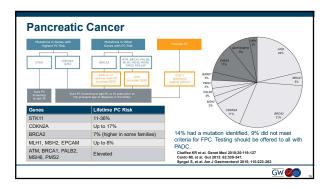






Prostate Cancer Genetic mutation carriers tend to have: Earlier age of diagnosis More aggressive disease Higher likelihood to metastasize FDA approval for use of PARPi ATM, BRCA1, BRCA2, BARD1, BRIP1, CDK12, CHEK2, FANCL, PALB2, RAD51B, RAD51C, RAD51D, RAD54 metastatic castration-resistant prostate cancer Phase III PROfound trial showed an OS benefit TRITON2 trial showed nearly 45% of men with BRCA2 mutations showed a tumor response In more than half of these men the response lasted at least 6 months





NCCN Update for Screening

- MRCP and/or EUS can be considered for:
 - Carriers of ATM, BRCA1, BRCA2, CDKN2A, Lynch syndrome, PALB2, STK11, TP53 with a family history of pancreatic cancer
 - 2 or more first degree relatives with pancreatic cancer, even in absence of a mutation
 - 3 or more first or second degree relatives with pancreatic cancer, even in absence of a mutation



PARPi and Pancreatic Cancer

- FDA Approval:
- First line maintenance therapy for patients whose disease has not progressed on at leat 16 weeks of platinum based chemotherapy
 BRCA1 or BRCA2 germline mutation



Gene	Effect on Pigment	Effect on Nevi	Penetrance	Other cancer risks
CDKN2A	No	Yes	High	Panc, Breast, cervical, GI, lymphoma, lung, Wilm's
CDK4	No	Yes	High	Similar to CDKN2A
TERT	No	Yes	High	Renal, bladder, AML, myeloproliferative neoplasms
POT1	No	No	High	Glioma, brain, breast, lung, CLL, endometrial
ACD	No	No	High	Breast, brain, lung, ovarian, cervical, colorectal, prostate, myeloproliferative neoplasms
TERF2IP	No	No	High	Similar to ACD
BAP1	No	Yes	High	Uveal melanoma, mesothelioma, renal, meningioma, paraganglioma
PTEN	Yes	Yes	High	Breast, thyroid, endometrium, colorectal, kidney
MC1R	Yes	No	Medium	None reported
BRCA2	No	No	Medium	Breast, ovarian, prostate, pancreas
MITE	Yes	Yes	Medium	Pancreatic, Renal

Genetic '	Testing	Referral	Criteria:
Affected	Patient	·e	

- All ovarian
- All pancreatic
- Colon or uterine under age 50
- All medullary thyroid
- All pheochromocytoma/ paraganglioma
- All adrenal cortical carcinoma
- Anyone Tested prior to 2014
- · Metastatic prostate cancer
- PREMM, BRCAPro, Penn II or other validated model showing a 10% risk for having a mutation

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Genetic Testing Referral Criteria: Breast Cancer Patients

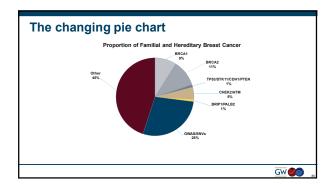
- Breast cancer under age 45
- Breast cancer and AJ heritage
- · Male breast cancer
- Triple neg breast cancer under age 60
- Bilateral breast cancer under age 50
- Metastatic, Her2- breast cancer
- Anyone Tested prior to 2014
- 2 or more family members with breast cancer at any age
- Family history of ovarian, pancreatic, or metastatic prostate cancer
- Dx <50 and limited or male dominated family structure



Genetic Testing Referral Criteria: Unaffected

- First or second degree relative that meets criteria
 - Per NCCN
 - · Not necessarily per insurance!
- Anyone Tested prior to 2014





Thank You!	
Feel free to email me with any questions or comments.	
Elizabeth Stark, MS, LCGC estark@mfa.gwu.edu	
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