

Practical Use of Genetic Testing in Clinical Practice

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The UCSF logo is located in the bottom right corner of the slide. It consists of the letters "UCSF" in a white, sans-serif font, centered within a dark blue square background.

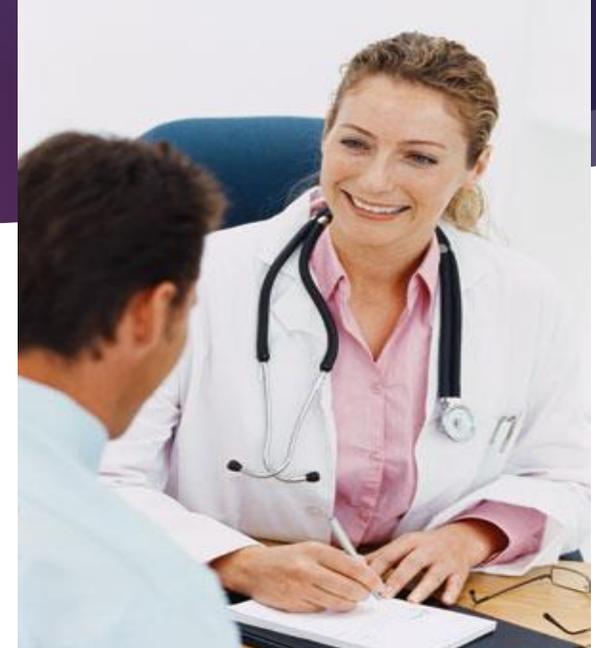
UCSF

Disclosures

None

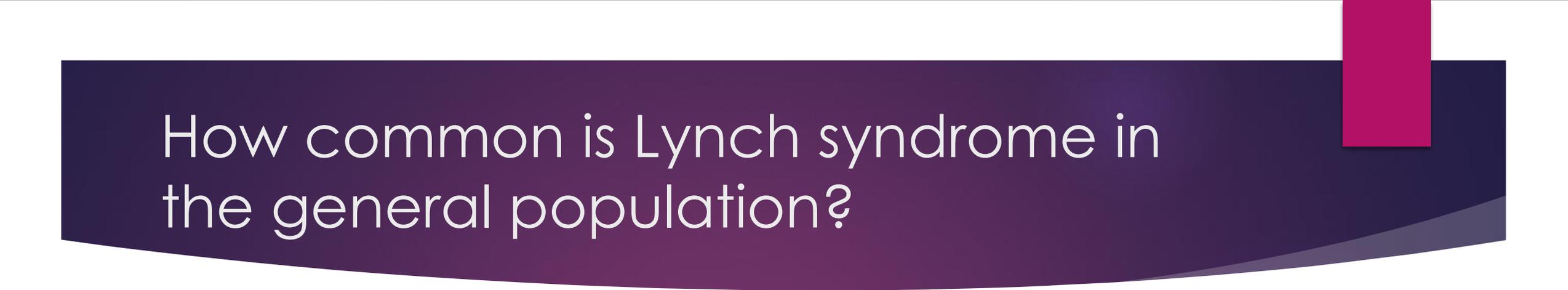
Patient's story...

- ▶ “Until my diagnosis, there were no less than **thirteen doctors** who could have taken detailed family history from the members of my family and referred them for genetic testing.
- ▶ It never occurred prior to the time, **I was diagnosed with a cancer.**”



Objectives

- ▶ Primer to genetic testing and counseling
- ▶ Universal tumor testing
- ▶ Criteria for referral to genetic testing



How common is Lynch syndrome in the general population?

A) 1:300

B) 1:1000

C) 1:5000

D) 1:10,000

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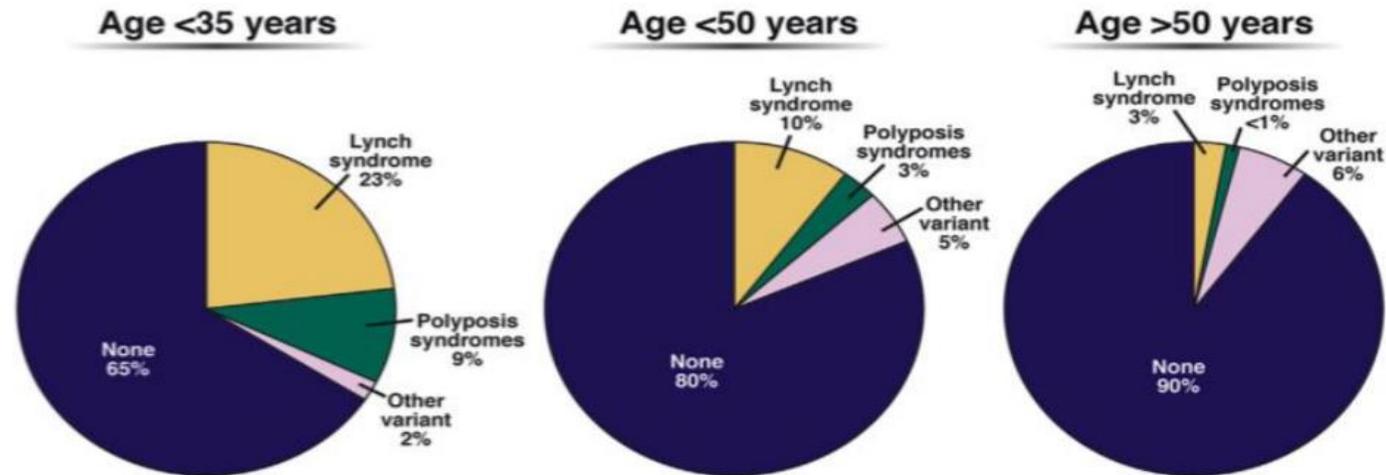
C) 1:5000

D) 1:10,000



1:300

Genes associated with CRC



Lynch syndrome	Polyposis syndromes	Other pathogenic variants	
		High penetrance	Moderate/low penetrance
<i>MLH1</i>	<i>APC</i>	<i>BRCA1</i>	<i>CHEK2</i>
<i>MSH2</i>	<i>MUTYH</i>	<i>BRCA2</i>	<i>ATM</i>
<i>MSH6</i>	<i>SMAD4</i>	<i>TP53</i>	<i>NBN</i>
<i>PMS2</i>	<i>BMPR1A</i>	<i>PALB2</i>	<i>BARD1</i>
	<i>PTEN</i>	<i>CDKN2A</i>	<i>BRIP1</i>
	<i>POLE</i>		

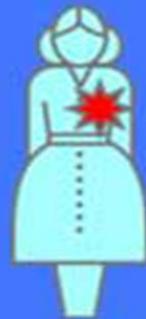
Lynch syndrome

Gene	CRC risk by age 80	Age to start CRC surveillance	Frequency
MLH1	40-61%	20-25 Y	1-2 years
MSH2(EPCAM)	33-52%		
MSH6	10-44%	30-35 Y	
PMS2	8-20%		

Mutations: Somatic and Germline

Somatic mutations

- Occur in nongermline tissues
- Are nonheritable



⇒ Nonheritable

Somatic mutation
(e.g., breast)

Germline mutations

- Present in egg or sperm
- Are heritable
- Cause cancer family syndrome



Mutation in
egg or sperm

All cells
affected in
offspring

Adapted by Joanne Kelly, © 2004.

Universal Tumor Testing for CRC

- ▶ Microsatellite instability (MSI)/ Immunohistochemistry (IHC) tumor testing to all CRC
- ▶ Does not require written consent

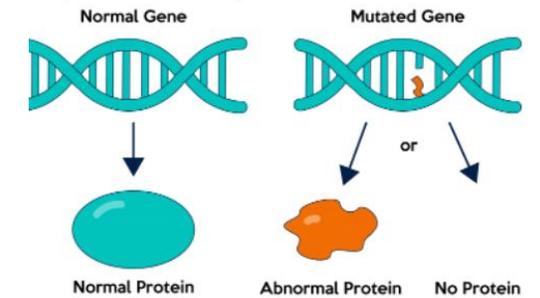
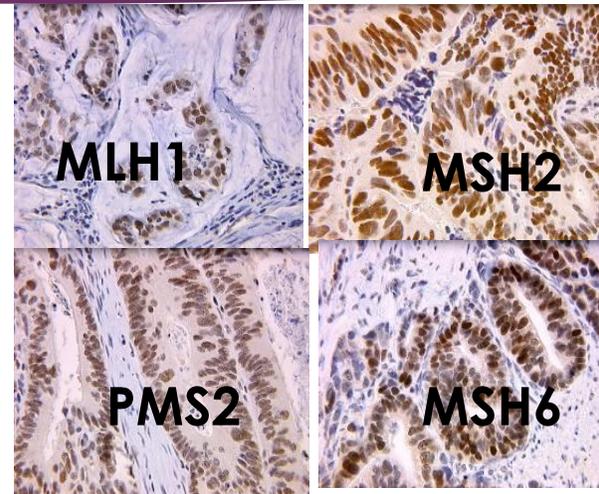
Microsatellite Instability Testing (MSI)

- ▶ Microsatellite
 - ▶ Stretches of DNA with repetitive sequences of nucleotide
 - ▶ (eg. AAAAA or CGCGCGCG)
- ▶ Susceptible to errors when MMR genes are mutated



Immunohistochemistry (IHC)

- Easily available tumor test
- Antibodies used to stain for MMR proteins



Mismatch
repair gene
mutated



Abnormal
Protein



Lack of
staining on
IHC

Immunohistochemistry tumor tissue

IHC Normal
MMR Proficient

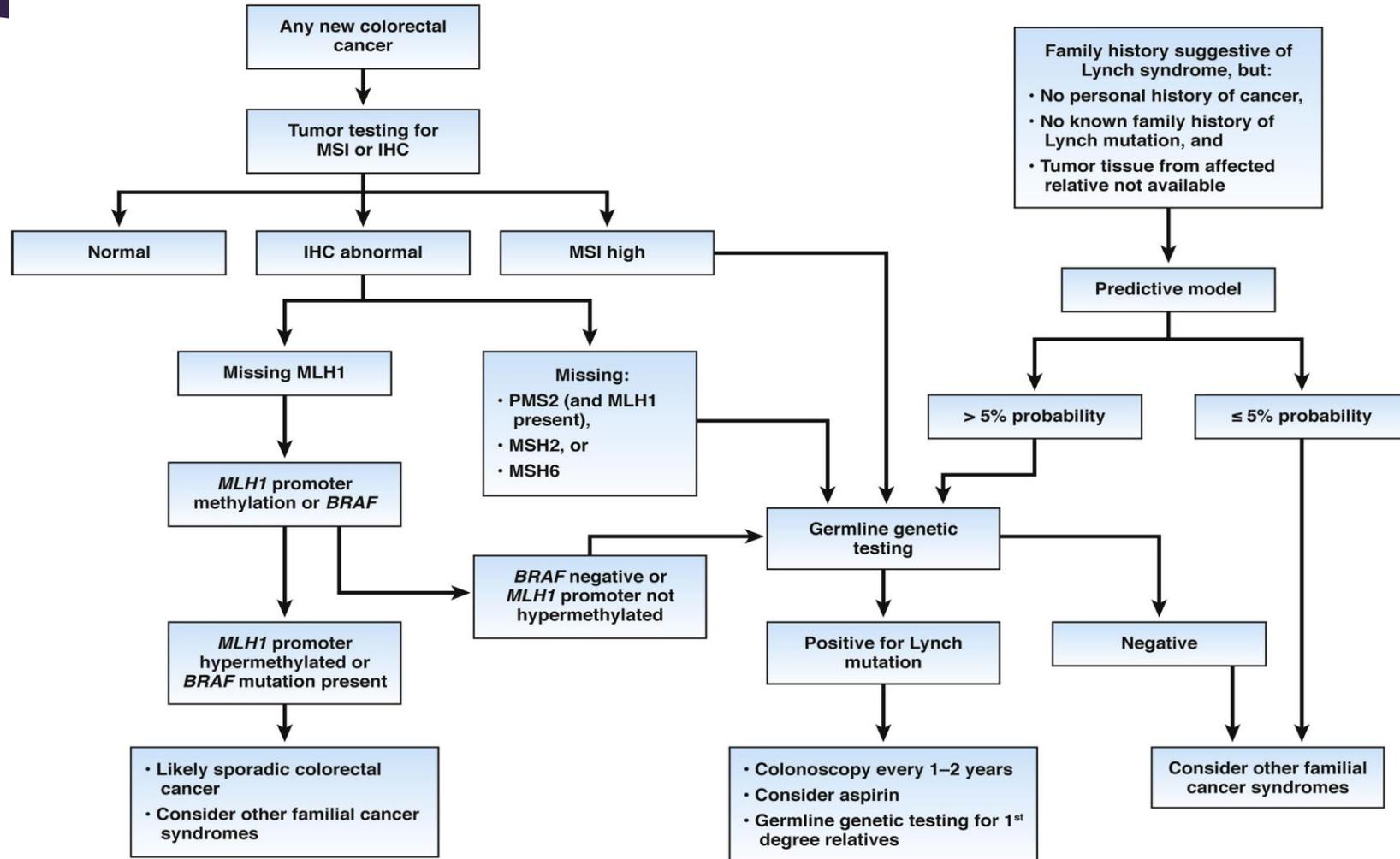
MLH1 expression: **Present**
PMS2 expression: **Present**
MSH2 expression: **Present**
MSH6 expression: **Present**

IHC Abnormal
dMMR
MMR Deficient

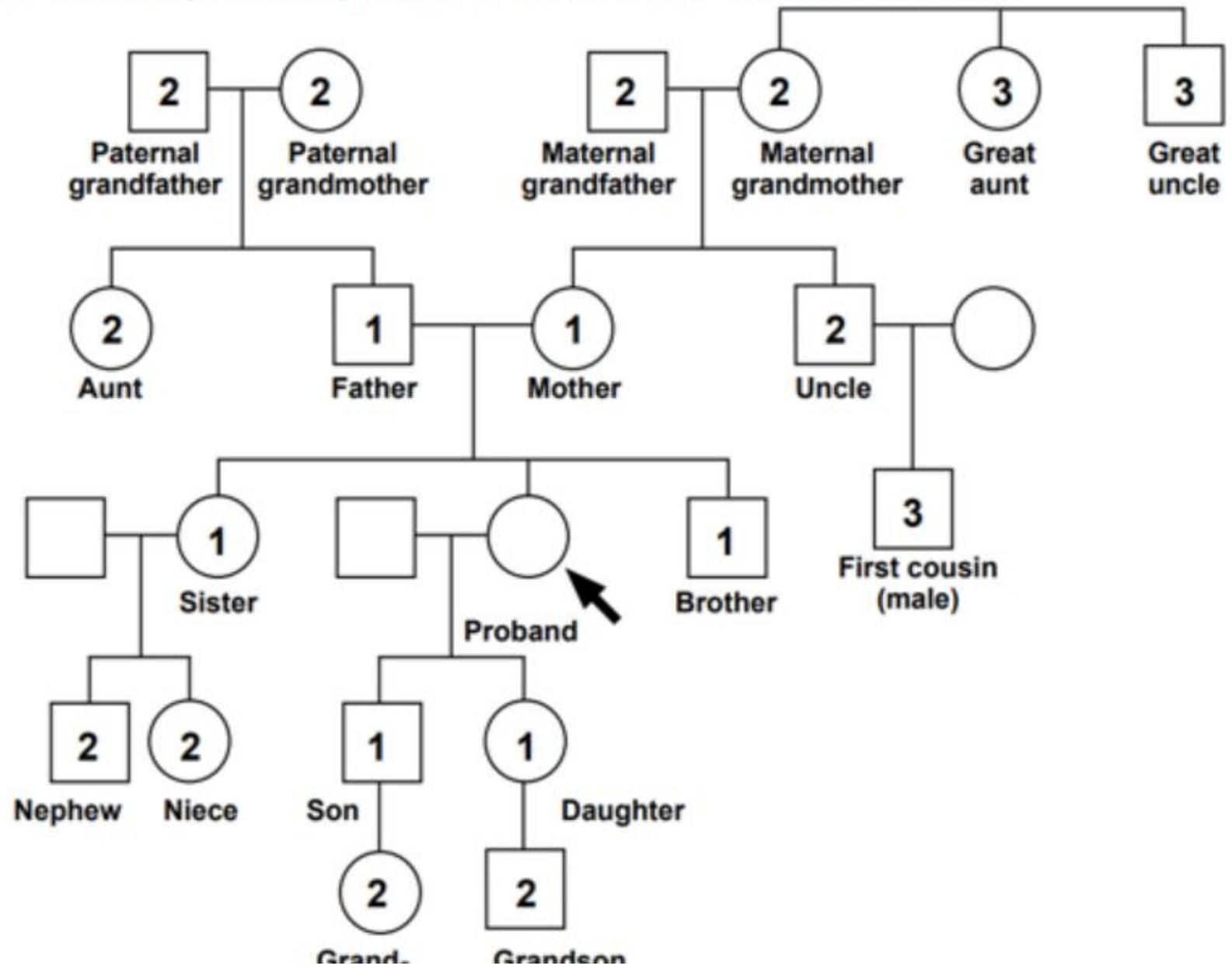
MLH1 expression: **Absent**
PMS2 expression: **Absent**
MSH2 expression: Present
MSH6 expression: Present

Diagnosis and Management of Lynch Syndrome

Clinical Decision Support Tool



PEDIGREE: FIRST-, SECOND-, AND THIRD-DEGREE RELATIVES OF PROBAND¹⁶



When to refer for genetic testing

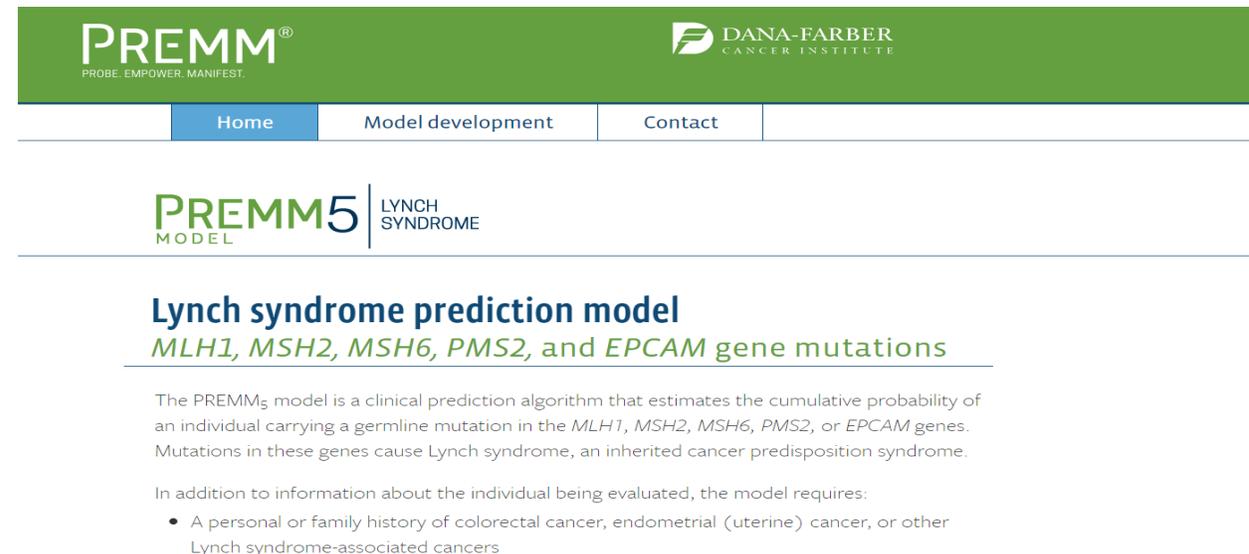
- ▶ ≥ 10 adenomatous polyps
- ▶ ≥ 2 hamartomatous polyps

- ▶ Tumor testing with MMR deficiency
- ▶ Family history of known genetic syndrome

- ▶ Meets NCCN criteria for Lynch testing
- ▶ Meets PREMM5 model cutoff ($>2.5\%$)

Lynch Syndrome Prediction Model

- ▶ **PREMM 5** (<https://premm.dfci.harvard.edu/>)
 - ▶ Sex /Age
 - ▶ History of colorectal cancer and other Lynch related cancer
 - ▶ Number of first and second degree relatives with CRC/
Endometrial cancer



The screenshot shows the top portion of the PREMM5 website. At the top is a green header bar with the PREMM logo on the left and the Dana-Farber Cancer Institute logo on the right. Below the header is a navigation menu with three items: 'Home', 'Model development', and 'Contact'. The 'Home' item is highlighted in blue. Below the navigation menu is a section titled 'PREMM5 LYNCH SYNDROME MODEL'. Underneath this title is a sub-heading 'Lynch syndrome prediction model' followed by 'MLH1, MSH2, MSH6, PMS2, and EPCAM gene mutations'. The main content area contains a paragraph describing the model and a list of requirements for evaluation.

PREMM[®]
PROBE. EMPOWER. MANIFEST.

DANA-FARBER
CANCER INSTITUTE

Home | Model development | Contact

PREMM5 | LYNCH SYNDROME
MODEL

Lynch syndrome prediction model

MLH1, MSH2, MSH6, PMS2, and EPCAM gene mutations

The PREMM₅ model is a clinical prediction algorithm that estimates the cumulative probability of an individual carrying a germline mutation in the *MLH1*, *MSH2*, *MSH6*, *PMS2*, or *EPCAM* genes. Mutations in these genes cause Lynch syndrome, an inherited cancer predisposition syndrome.

In addition to information about the individual being evaluated, the model requires:

- A personal or family history of colorectal cancer, endometrial (uterine) cancer, or other Lynch syndrome-associated cancers

NCCN Criteria for Lynch syndrome

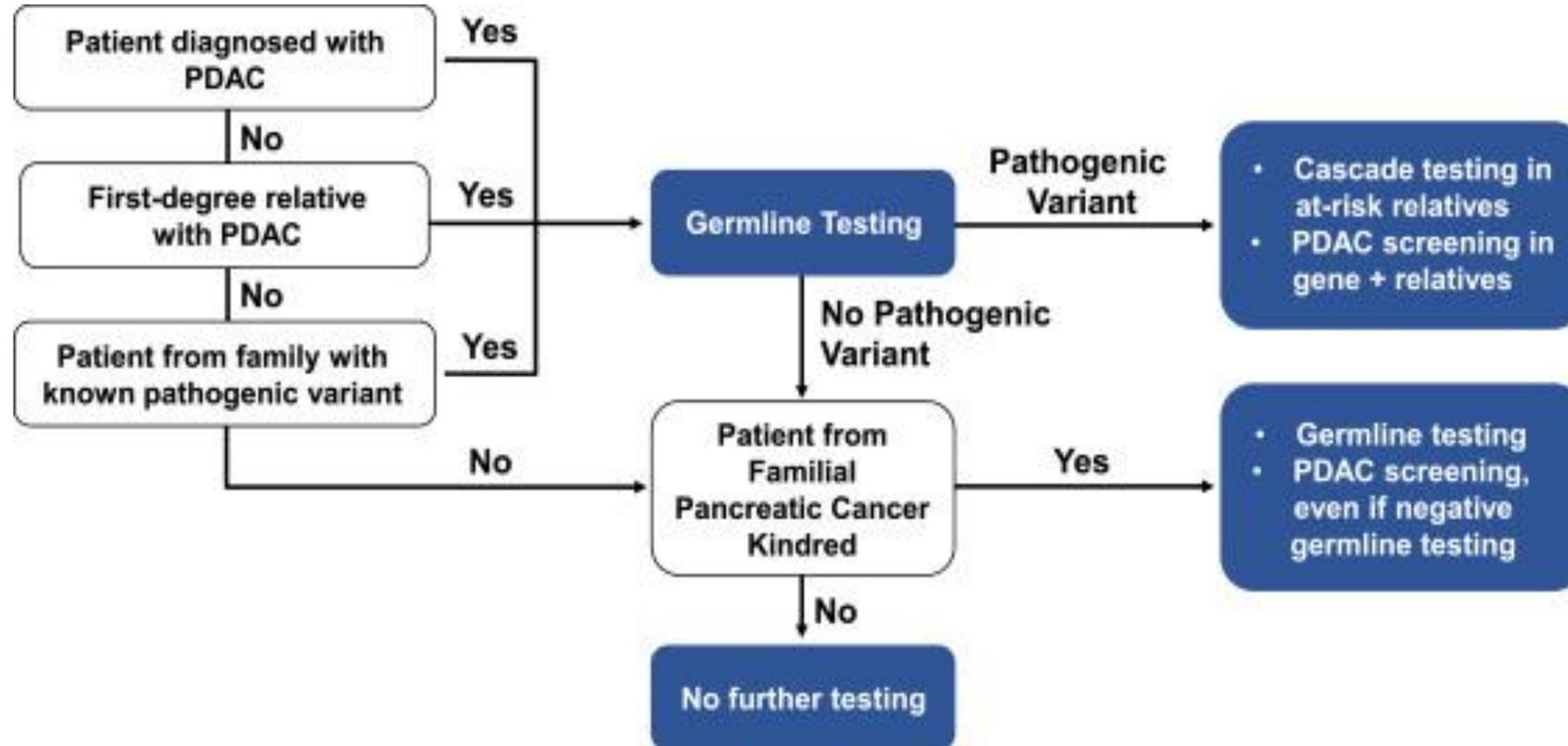
- ▶ ≥ 1 FDR with a CRC or endometrial cancer < 50 y
- ▶ ≥ 1 FDR with a CRC or endometrial cancer and a synchronous or metachronous **LS-related cancer*** regardless of age
- ▶ ≥ 2 FDR or SDR with **LS-related cancers**, including ≥ 1 diagnosed < 50 y
- ▶ ≥ 3 FDR or SDR with **LS-related cancers** regardless of age

*glioblastoma, small bowel, stomach, pancreas, bile duct, gallbladder, kidney, ureter, bladder, prostate, sebaceous skin tumors

Pancreatic Cancer

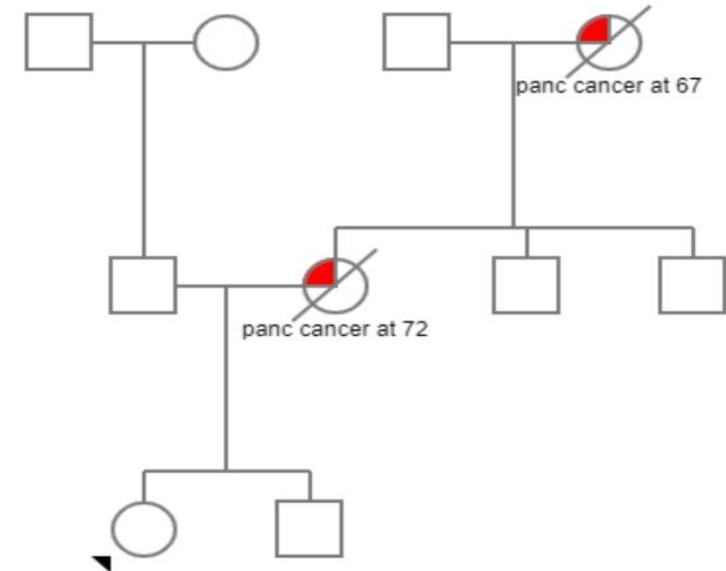
- ▶ 10% of pancreatic cancers are due to genetic causes
 - ▶ (ATM, BRCA1/2, CDKN2A, Lynch Genes, PALB2, STK11, TP53)

How to approach a patient with family history of pancreas cancer



Familial Pancreatic Cancer (FPC) Kindred

- ▶ A family with two or more individuals with pancreatic cancer and two of these have a first degree relationship to each other (parent, child, sibling)
- ▶ Absence of known hereditary syndrome



Genetic counseling



President George W. Bush signs H.R. 493, the Genetic Information Nondiscrimination Act of 2008, Wednesday, May 21, 2008, in the Oval Office. White House photo by Eric Draper.

- ▶ Limitation of testing review
 - ▶ Not all genes are known
 - ▶ Finding on VUS, genes with not well defined penetrance or management plan

- ▶ GINA (Genetic Information Non-discrimination Act)-2008
 - ▶ Excluded- Life insurance, disability, long term care insurance

Genetic Testing sample

- ▶ **Saliva sample/ Buccal swab**
- ▶ Blood test
- ▶ Skin biopsy fibroblast culture
(bone marrow transplant recipient, active/recent hematologic malignancy)

- ▶ Turnaround time: 2-3 weeks
- ▶ Out of pocket cost- \$250



Genetic Testing Results

Positive Mutation
Pathogenic/Likely Pathogenic
Deleterious/ Suspected Deleterious

- Positive Results

Negative

- Manage according to family history
- Family testing not needed

VUS
(Variant of Unknown Significance)

- Manage according to family history
- Family testing not needed

Timely genetic testing is critical

- ▶ Timely genetic testing can alter:
 - ▶ Type of surgery
 - ▶ Hysterectomy/oophorectomy at the time of colectomy
 - ▶ Extended colectomy
 - ▶ Therapeutics -
 - ▶ Genomic instability may lead to neo-antigen expression and susceptibility to immunotherapy and better outcome
 - ▶ Lynch syndrome vaccine trials underway

Resources

GI Cancer Genetics

- ▶ www.findageneticcounselor.org (Find a genetic counselor)
- ▶ www.cgaigc.com/find-a-gi-genetics-clinic(GI cancer genetics clinic)
- ▶ www.plsd.eu (Prospective Lynch Syndrome Database (PLSD) - gene, organ specific cancer risk)
- ▶ www.nccn.org
- ▶ www.Ask2me.org (Calculate the risk of cancers associated with genes)

Patient information resources

- ▶ www.Kintalk.org
- ▶ www.facingourrisk.org
- ▶ www.alivenkickn.org
- ▶ www.nostomachforcancer.org

Conclusion

- Technological advances have made genetic testing more affordable
- Timely diagnosis of genetic cancer syndromes can alter therapeutics and outcomes for the patients and their families
- Family history and referral to genetic testing should be a routine practice for all

Thank you!
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UCSF

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San Francisco