

Integration of Genomics in Cancer Care

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Purpose

- * To introduce how genetics and genomics are integrated into cancer care from prevention to treatment**

Topics

- * **Etiology of Cancer**
- * **Cancer Risk Assessment**
- * **Tumor Profiling**
- * **Pharmacogenomics**
- * **Targeted Cancer Therapy**

Case Study

- * **Mr. J – 41 yrs of age, white, Northern European ancestry**
- * **Biopsy: right-sided colon cancer; plus two adenomatous polyps**
- * **No prior cancer history**
- * **Medical history otherwise unremarkable**

Case Study – Mr. J

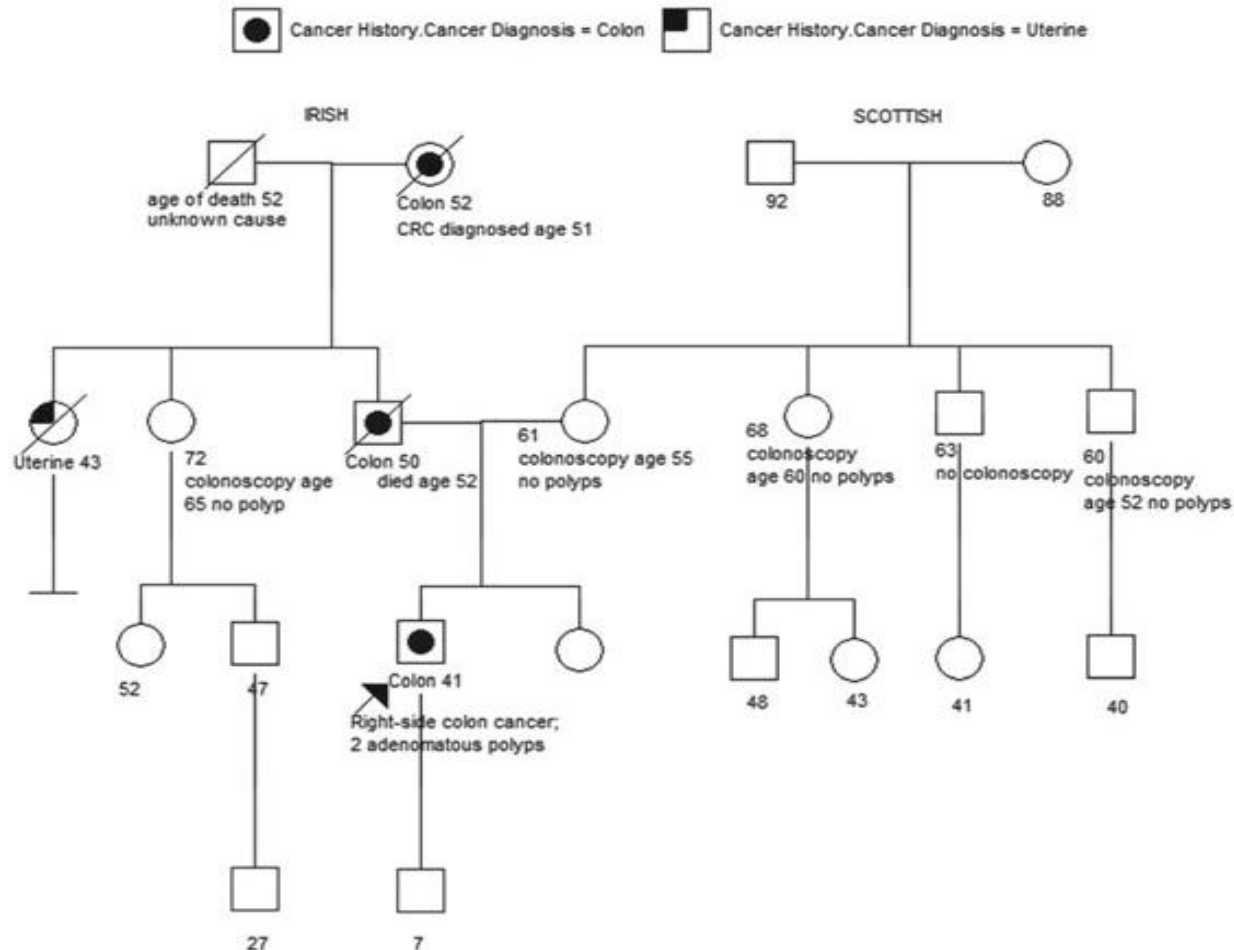
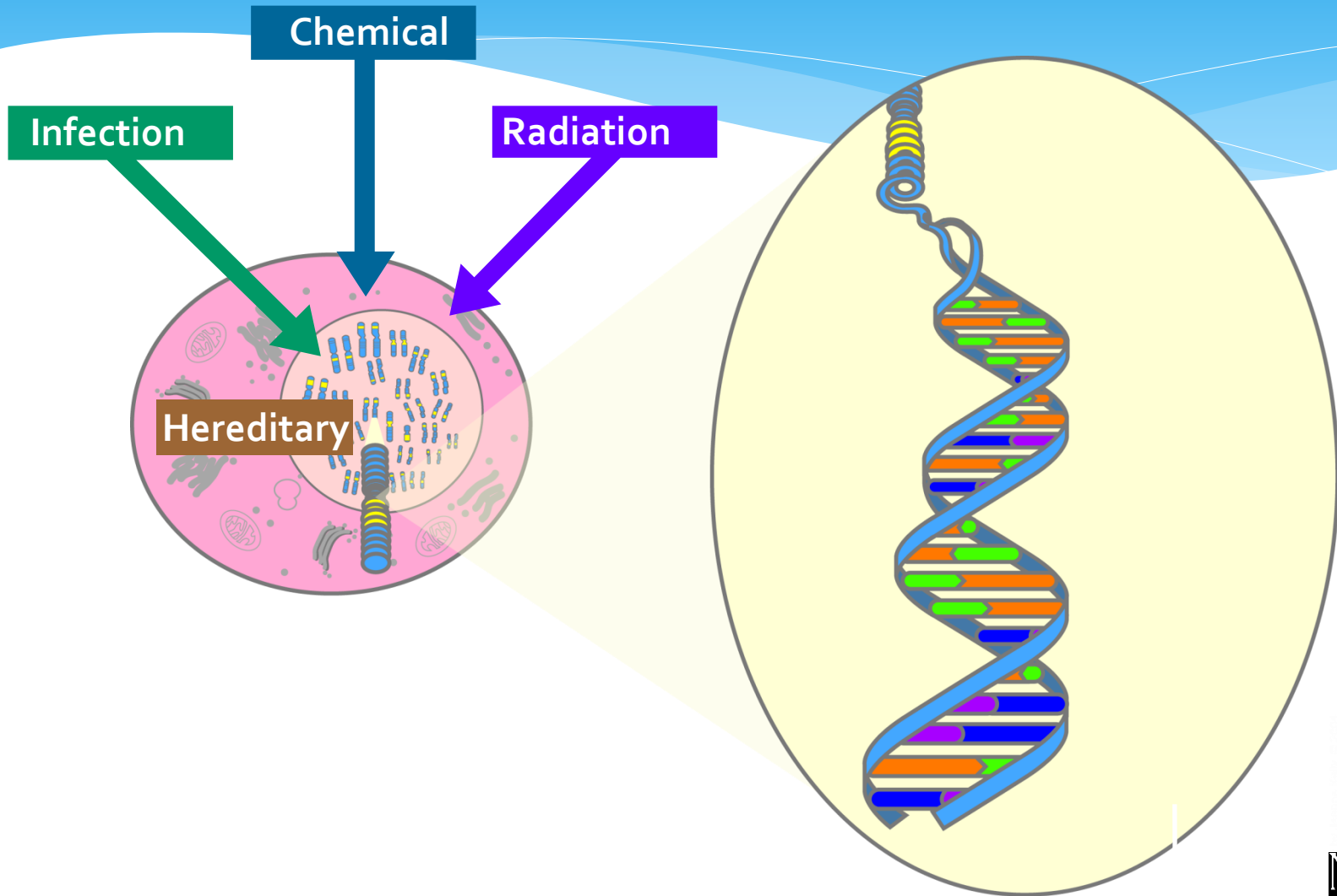
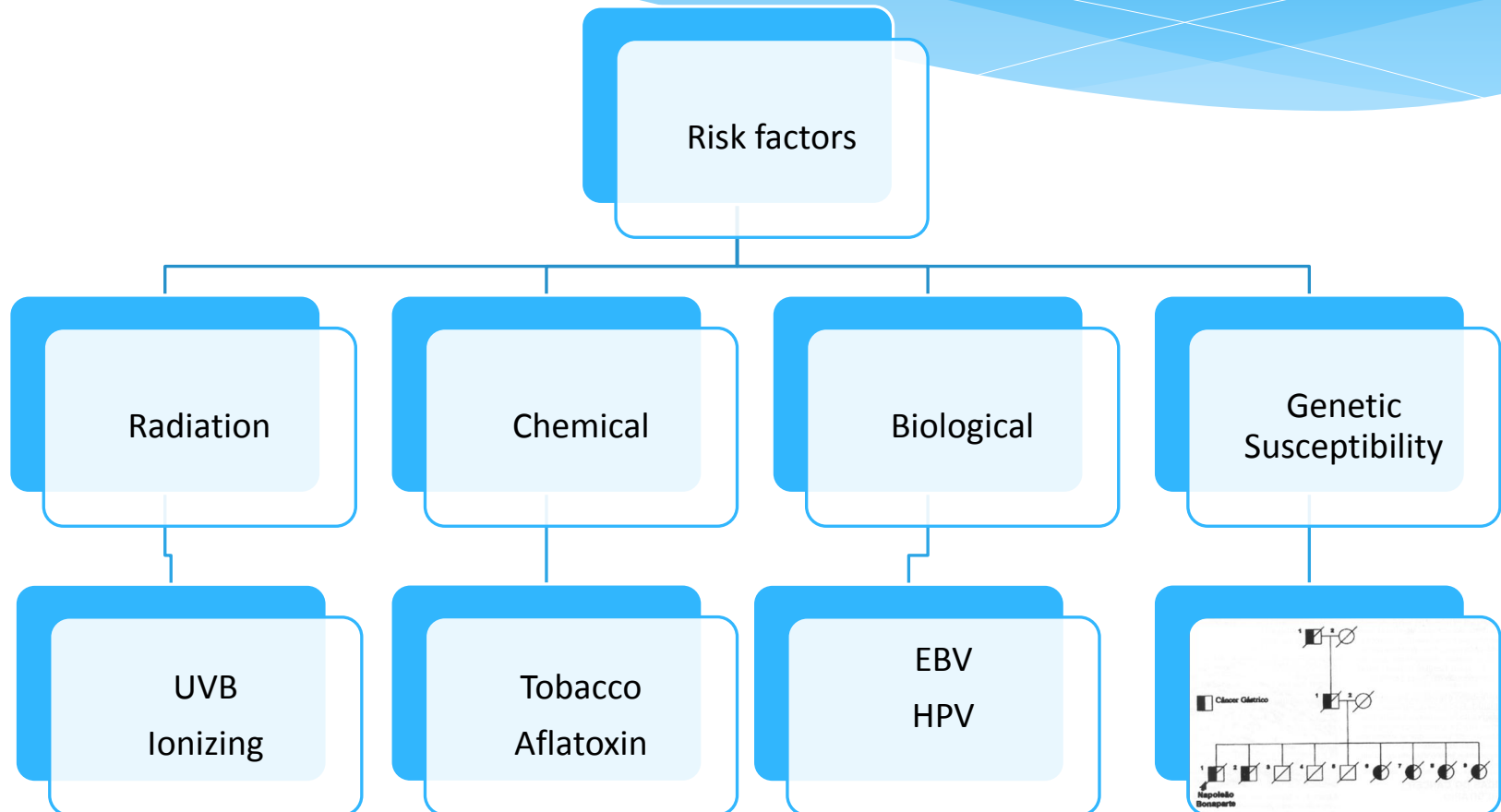


Figure 1. Four-generation pedigree with significant family history of colon and uterine cancers, in the paternal lineage; suspect for Lynch syndrome (fictitious case).

Etiology of Cancer

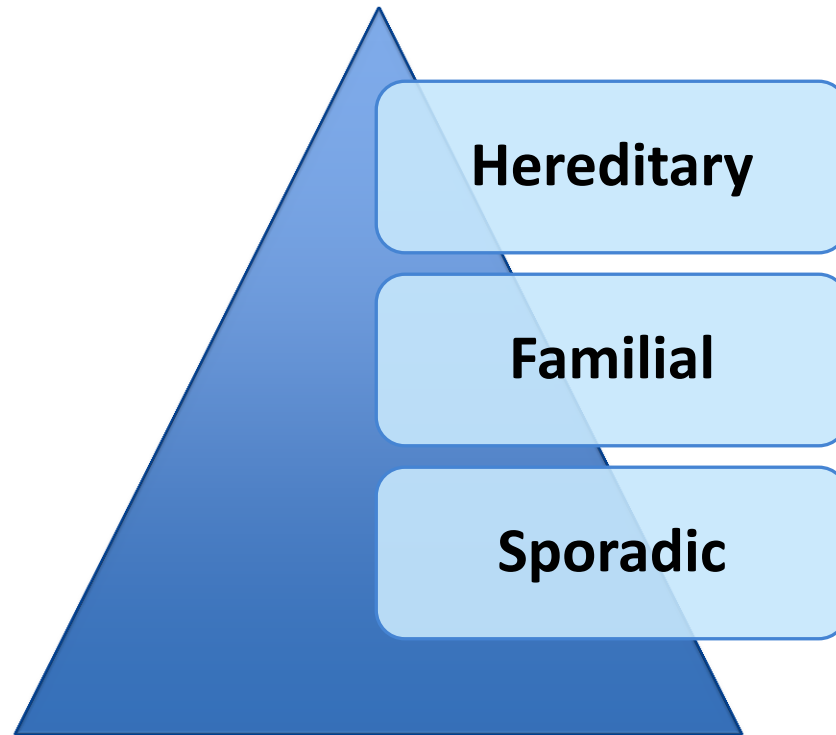


Etiology of Cancer



Etiology of Cancer

Classification of Tumors Due to Family History (FH)



Etiology of Cancer

Classification of Tumors Due to Family History



Sporadic

75% of all cancers
Age of onset typically that expected for the type of cancer
Somatic (acquired) mutations in a specific tissue (e.g., breast, colon)

Etiology of Cancer

Classification of Tumors Due to Family History



Familial

10%-15% of all cancers
Same cancer type occurring at
excepted age in more than
one close relative
Shared environmental +
genomic influences

Etiology of Cancer

Classification of Tumors Due to Family History



Hereditary

5%-10 of all cancers
Earlier age at onset than usual
May or may not have FH of same cancer or other cancers associated with a cancer syndrome
Single gene mutation in the germline (egg or sperm)

Etiology of Cancer

Somatic mutations

- Occur in non-germline tissues
- Are not heritable

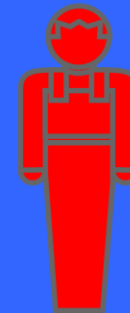


Non-heritable

Somatic mutation
(e.g., breast)

Germline mutations

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

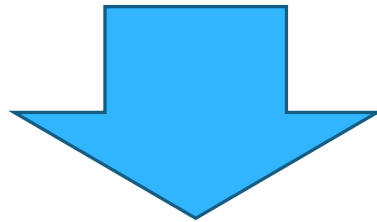


Mutation in
egg or sperm

All cells affected
in offspring

Etiology of Cancer

How important is to recognize the difference among acquired and heritable genetic mutations?



Key to appropriate referral for further evaluation

Cancer Risk Assessment (CRA)

Objectives
of CRA

Define
cancer risk

Identity
individuals
who may
benefit from
genetic
testing

Provide risk-
based cancer
screening
and risk
reduction
strategies

Assess
psychosocial
and cultural
implications of
risk
assessment

Provide
education,
counseling to
facilitate
informed
decision
making

Cancer Risk Assessment

How to recognize individuals for CRA?

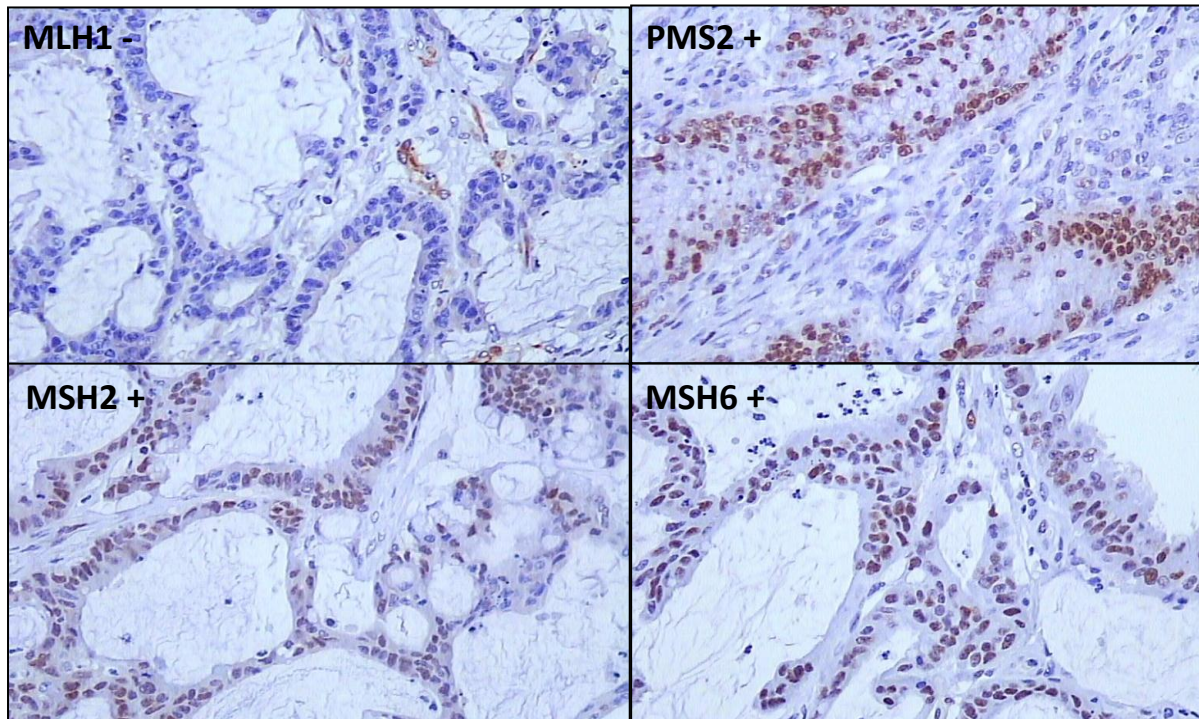
- ⌘ **Earlier age of cancer onset than expected**
- ⌘ **Same type of cancer in two or more close relatives**
- ⌘ **Two or more primary cancers in the same person**
- ⌘ **Constellation of cancers characteristic of a hereditary syndrome**
- ⌘ **Male breast cancer, ovarian cancer or medullary thyroid cancer, at any age**
- ⌘ **Previously identified cancer-associated mutation in the family**

Tumor Profiling

- * **Evaluation of genomic, proteomic and epigenomic expression factors for cancer diagnosis, prognosis and therapeutics**

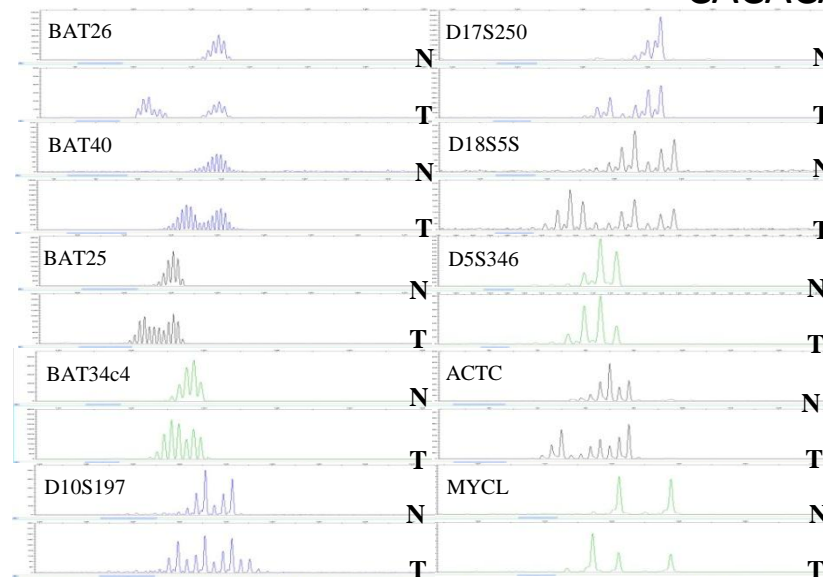
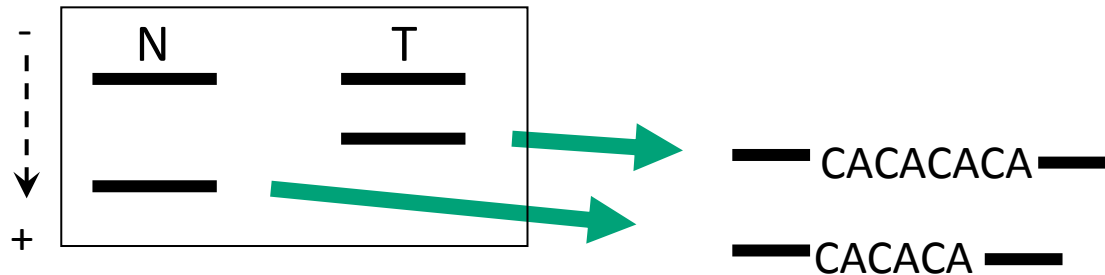
Case Study – Mr. J

Immunohistochemistry – test for protein expression of 4 genes associated with colorectal cancer
Result: absence of MLH1 expression



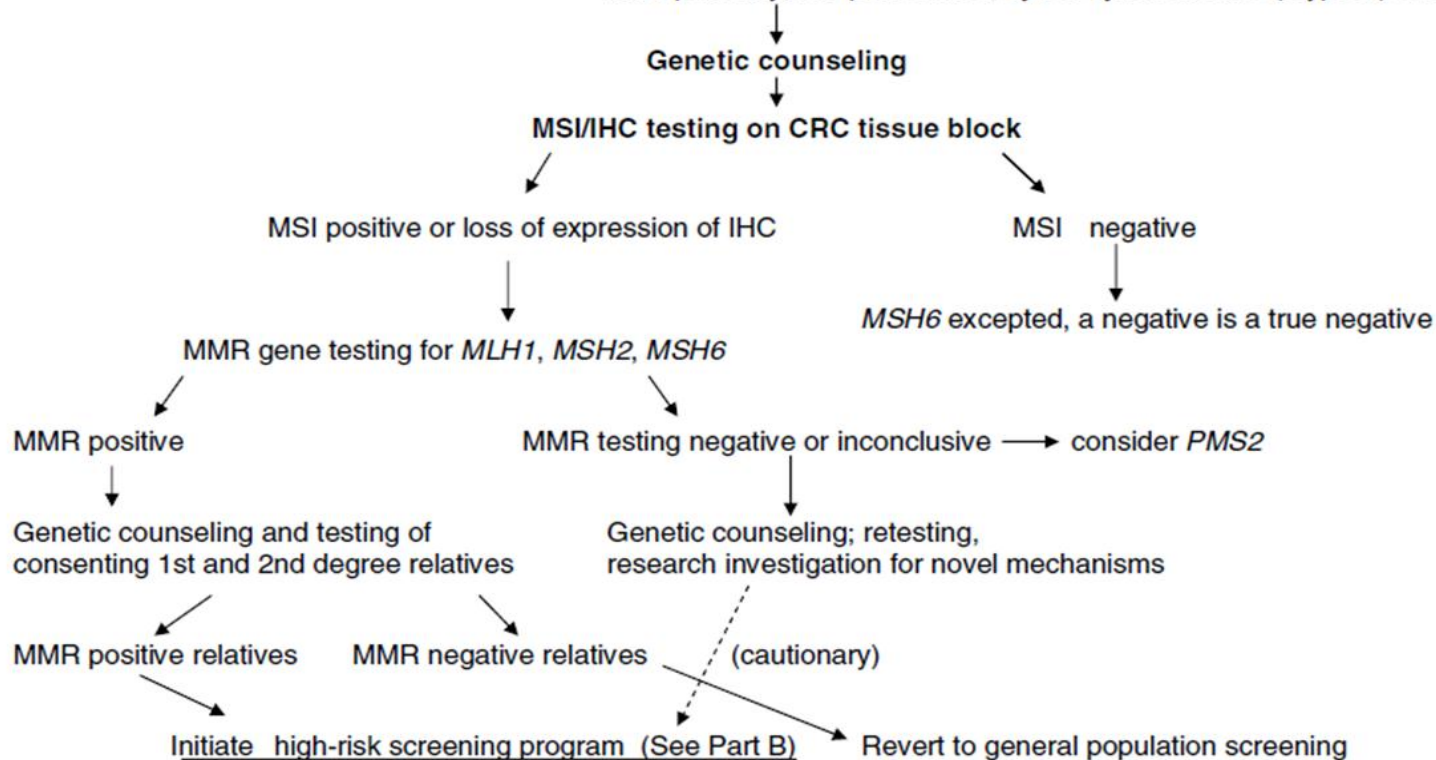
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Other evidence of germline mutation: MSI testing
Result: MSI-H (MSI-High)

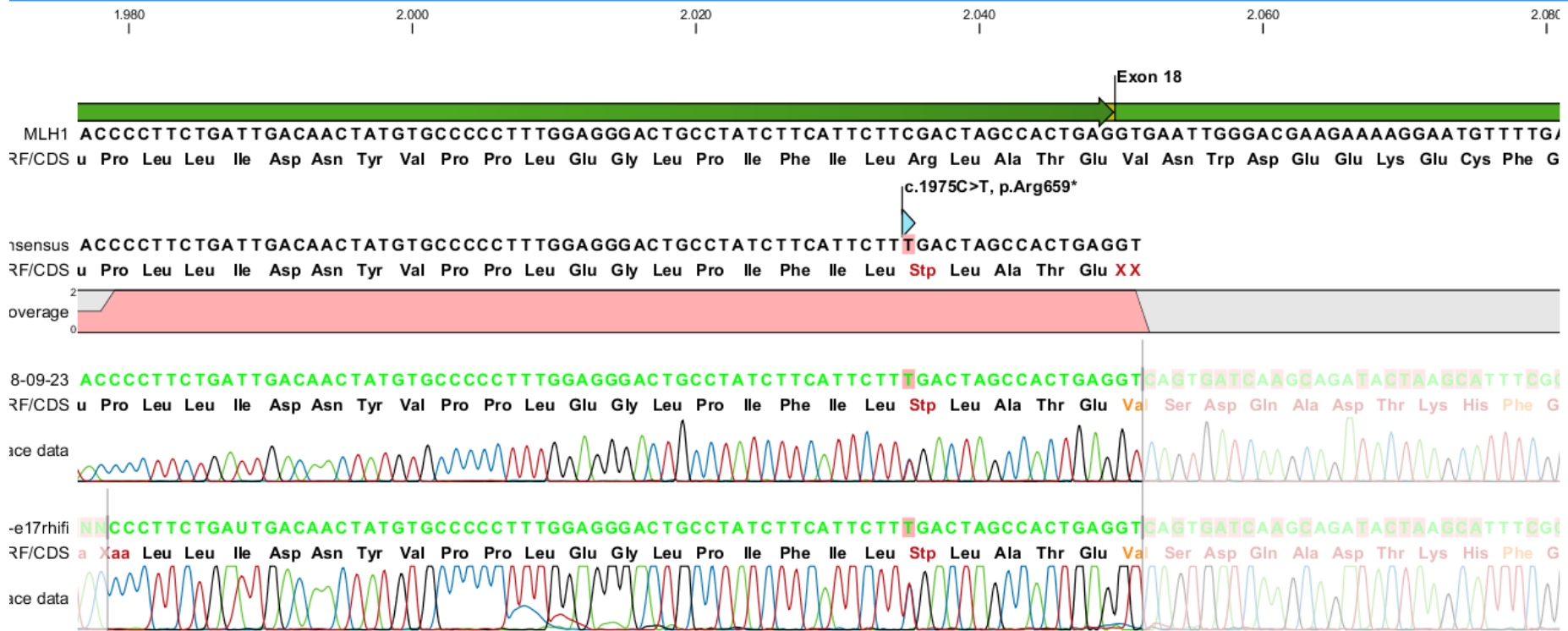


Case Study – Mr. J

Construct modified nuclear pedigree: Invoke Amsterdam I/II or Bethesda Criteria
Include all maternal and paternal 1st and 2nd degree relatives.
Record all cancer occurrences.
Invoke cardinal principles of Lynch syndrome.
Must consider adoption, incomplete FH, denial/poor cooperation,
false paternity, low penetrance, Lynch syndrome-like (atypical) family.

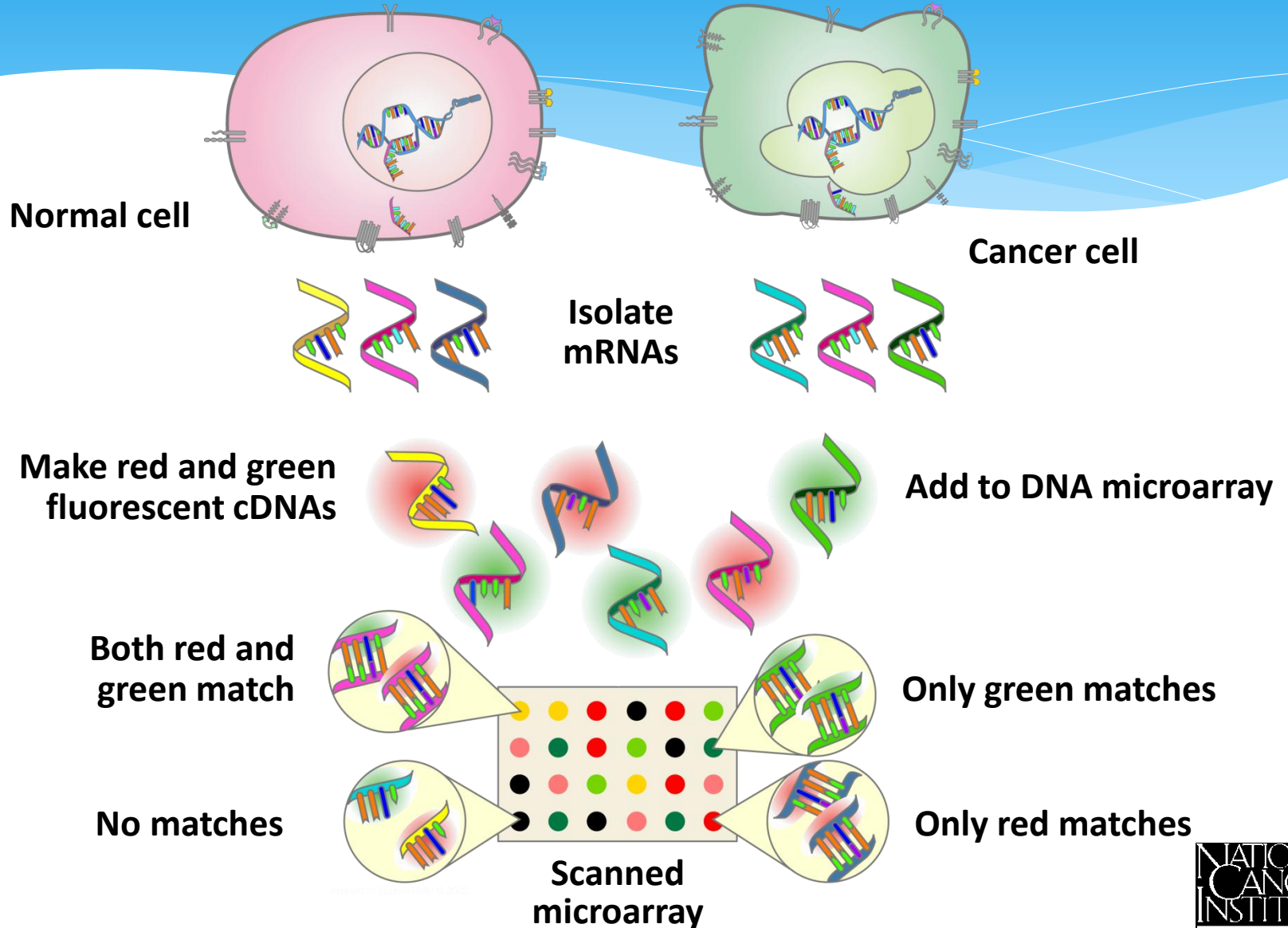


Case Study – Mr. J



Stop codon – exon 17 (c.1975C>T; p.Arg659*)

Tumor Profiling - Microarray



SNPs and Cancer Risk

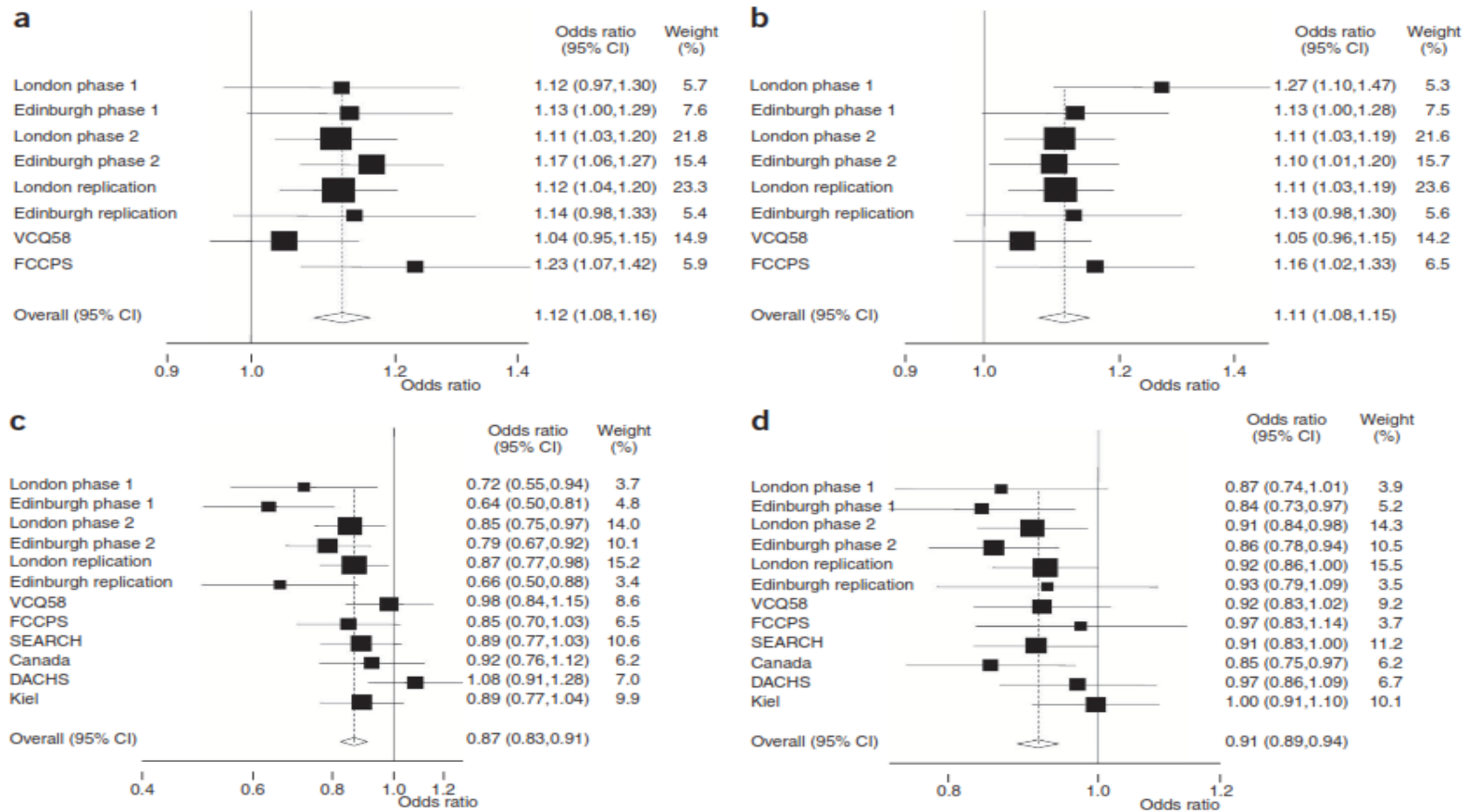
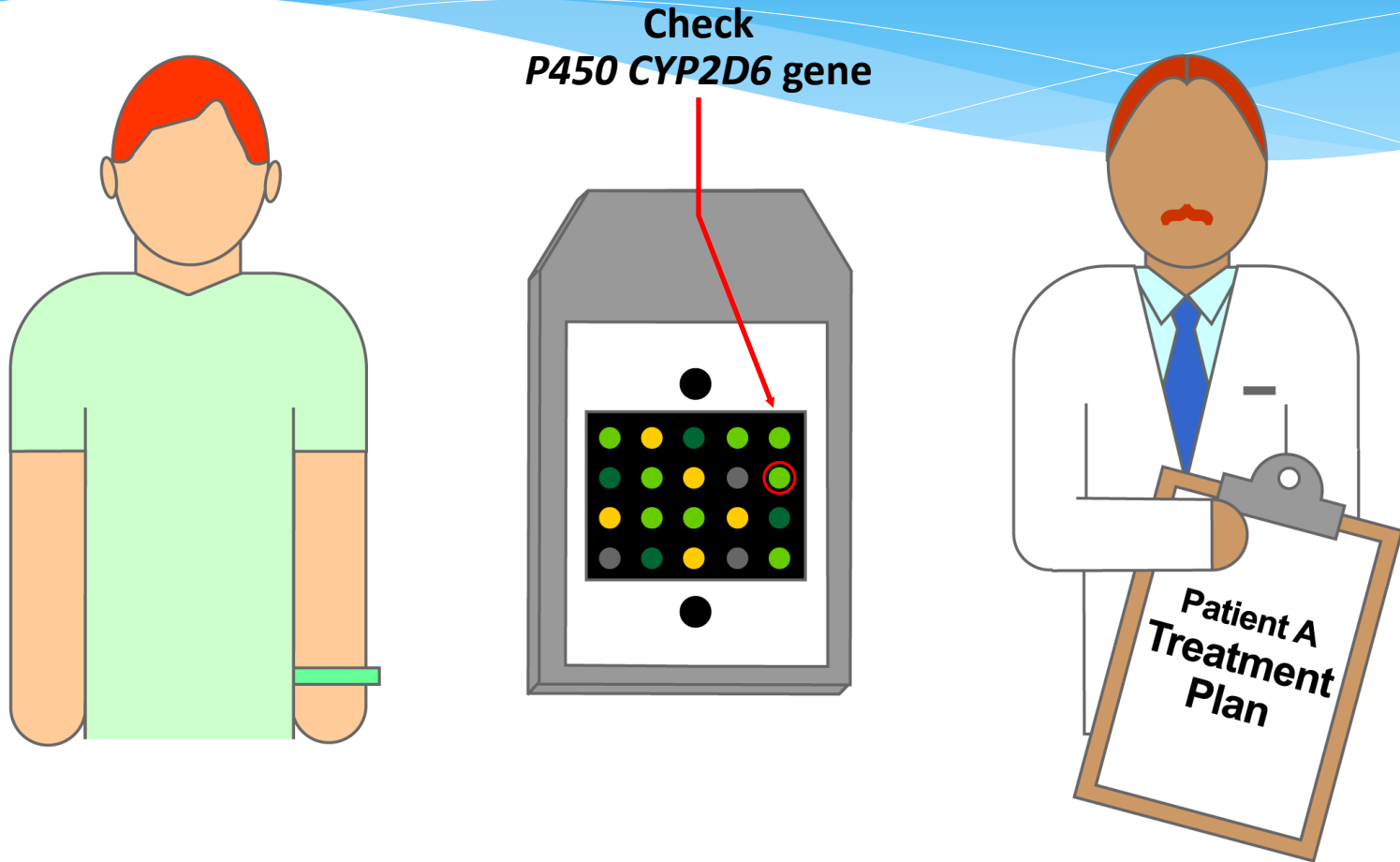
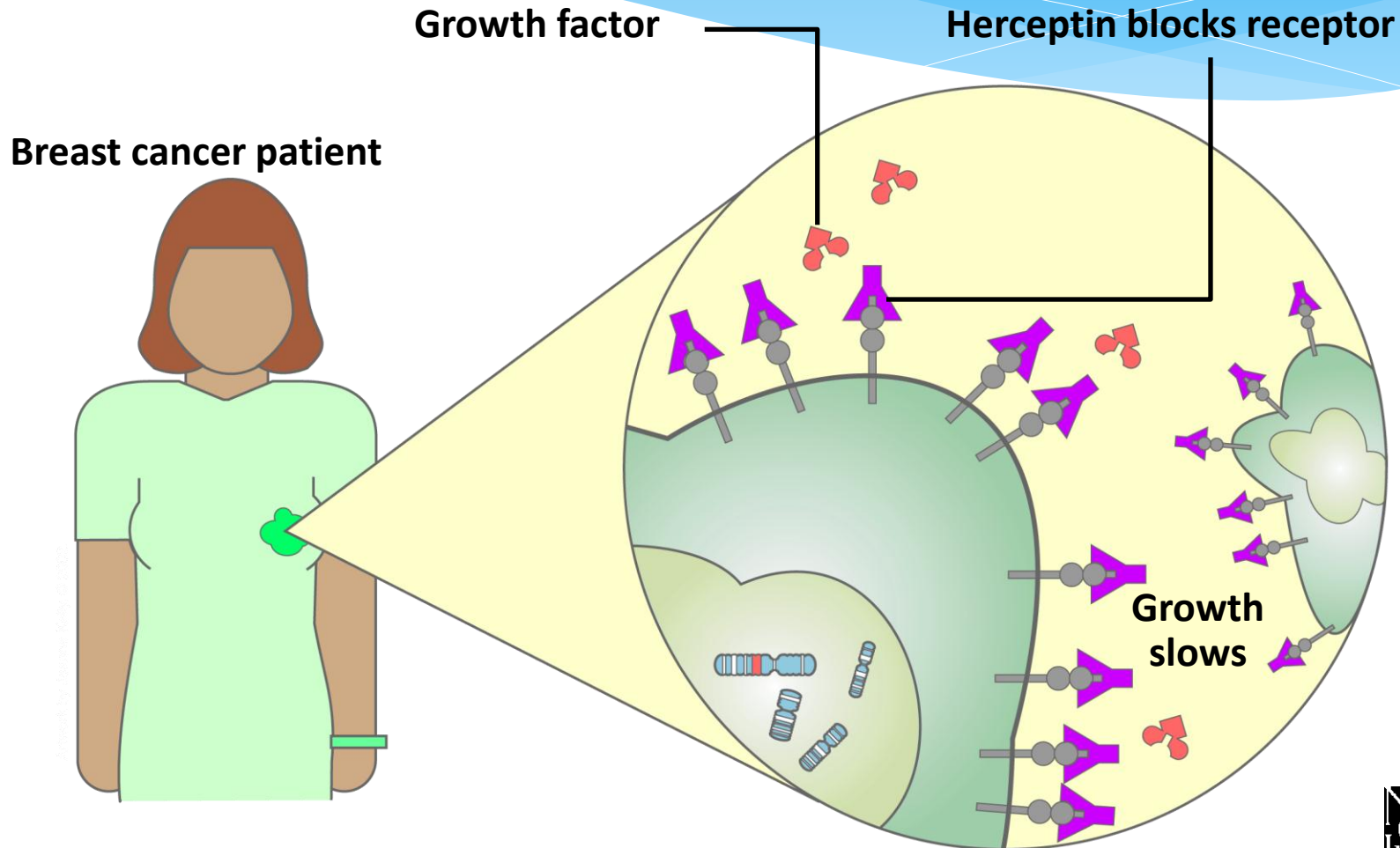


Figure 2 Forest plot of effect size and direction for the four SNPs associated with CRC. (a) rs961253. (b) rs4444235. (c) rs10411210. (d) rs9929218. Boxes denote allelic OR point estimates, their areas being proportional to the inverse variance weight of the estimate. Horizontal lines represent 95% CIs. The diamond (and broken line) represents the summary OR computed under a fixed-effects model, with the 95% CI given by its width. The unbroken vertical line is at the null value (OR = 1.0).

SNPs and Pharmacogenomics



Targeted Therapy - Trastuzumab



Targeted Therapy

Table 1. Selected Genetic Markers and Their Application in Cancer Treatment

Tumor	Genetic marker	Description-application	Drug-implication
Breast	<i>HER2</i> amplification	<i>HER2</i> -positive tumors indicates need for additional therapy.	Trastuzumab, lapatinib
Breast	OncotypeDx [®]	Microarray analysis of 21 genetic markers. Identifies if patients with early stage ER-positive, lymph node negative, <i>Her2</i> -negative tumors may benefit from adjuvant chemotherapy.	Chemotherapy evaluation
Colorectal cancer	OncotypeDx [®]	Microarray analysis of 12 genetic markers. Identifies if patients with stage II disease may benefit from adjuvant chemotherapy.	Chemotherapy evaluation
	<i>KRAS</i> mutation	Tumors with a <i>KRAS</i> mutation do not respond to treatment with EGFR monoclonal antibodies. <i>KRAS</i> status should be evaluated prior to treatment.	Cetuximab, panitumumab contraindicated
	<i>UGT1A1*28</i>	Patients with a germline <i>UGT1A1</i> variant are at risk for higher toxicity (especially neutropenia, diarrhea).	Irinotecan; consider dosage adjustment or alternate drug
Leukemia	<i>BCR-ABL</i>	Ph + CML; Ph + ALL. Presence of a <i>BCR-ABL</i> gene mutation indicates response to tyrosine kinase inhibitor therapy.	Imatinib, dasatinib, nilotinib
Non-small-cell lung cancer	<i>EGFR</i> mutation	<i>EGFR</i> mutation is associated with a better response to an <i>EGFR</i> -tyrosine-kinase inhibitor.	Erlotinib, gefitinib
Breast, ovarian	<i>BRCA1/BRCA2</i> mutation	Patients with a germline <i>BRCA</i> gene mutation who have disease progression following initial therapy may respond to treatment with PARP inhibitors.	Olaparib, for example
Melanoma	<i>BRAF</i> V600E mutation	Tumors with this <i>BRAF</i> mutation are sensitive to a kinase inhibitor	Vemurafenib indicated

Note. ER = estrogen receptor; EGFR = epidermal growth factor receptor; Ph = Philadelphia chromosome; CML = chronic myelogenous leukemia; ALL = acute lymphoblastic leukemia; PARP = poly ADP ribose polymerase.

Case Study – Mr. J

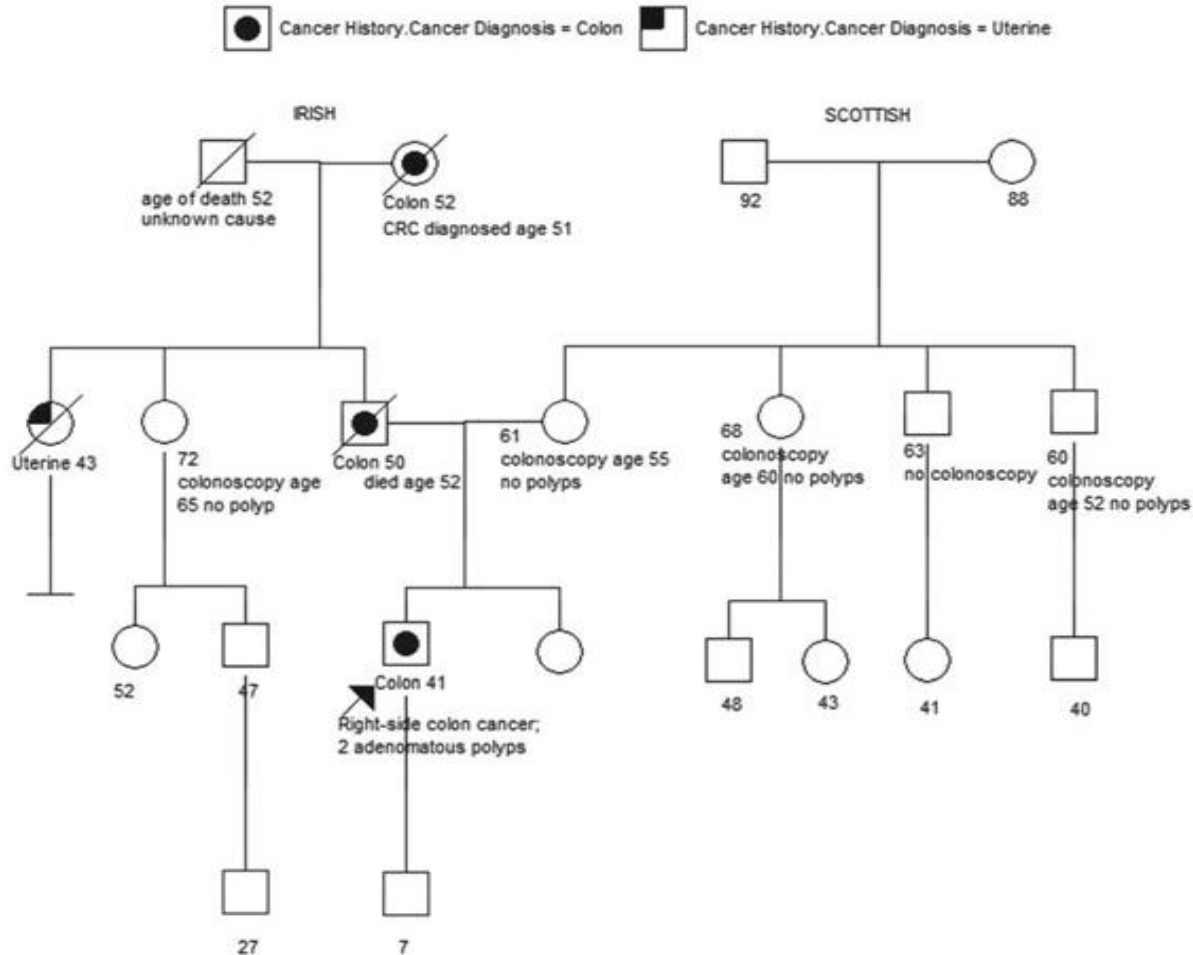


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Case Study – Mr. J

- * **MSI – Important to guiding treatment decision-making in early stage colon cancer**
- * **IHC – Important to guiding genetic testing strategy**
- * **Mutation detection – Important to guiding genetic counseling/testing for at-risk family members**

Closing Remarks

- * **Genomic care is now central to the care of patients with cancer**
- * **Nurses must be aware of developments in genomics and its impact in the cancer care continuum to help educate patients and support informed decision-making**