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Immunotherapy and Targeted Therapy in Clinical Oncology: Commercial Perspective

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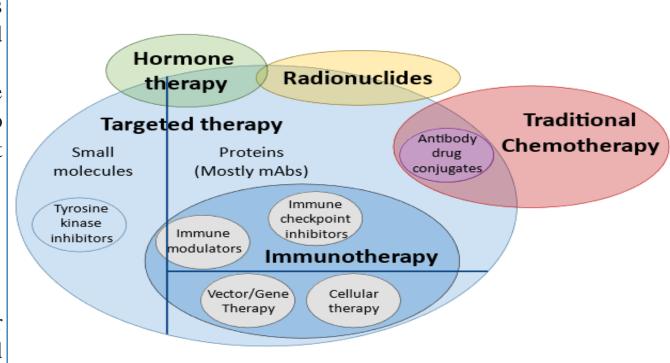


Immunotherapy and Targeted Therapy



Targeted Therapy

- Targeted therapy- uses drugs to target specific genes and proteins
- Patients with cancer will have a target for a certain drug, so they can be treated with that drug.
- 3 main types: 1. small molecule medicines 2. monoclonal antibodies
 3. immunotoxins
- cetuximab (Erbitux) –for advanced bowel cancer and HNC
- trastuzumab (Herceptin) for breast cancer and stomach cancer.

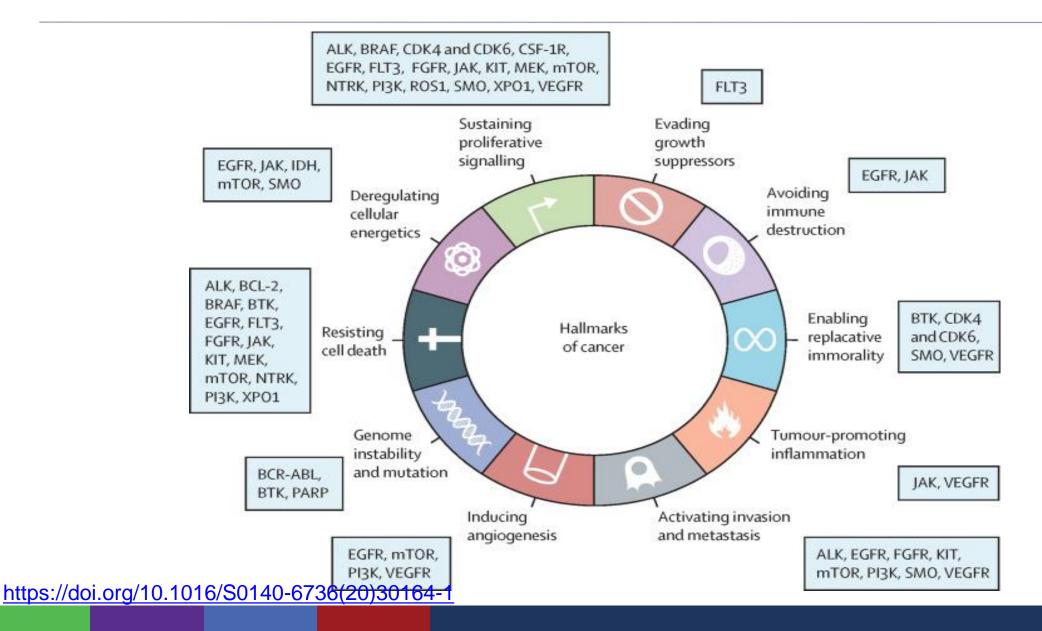


Immuno-Therapy

- Some monoclonal antibodies are also immunotherapy because they help turn the immune system against cancer.
- Example:
 Checkpoint
 inhibitors block
 proteins that stop the
 immune system
 attacking cancer
 cells.
 pembrolizumab
 (Keytruda)

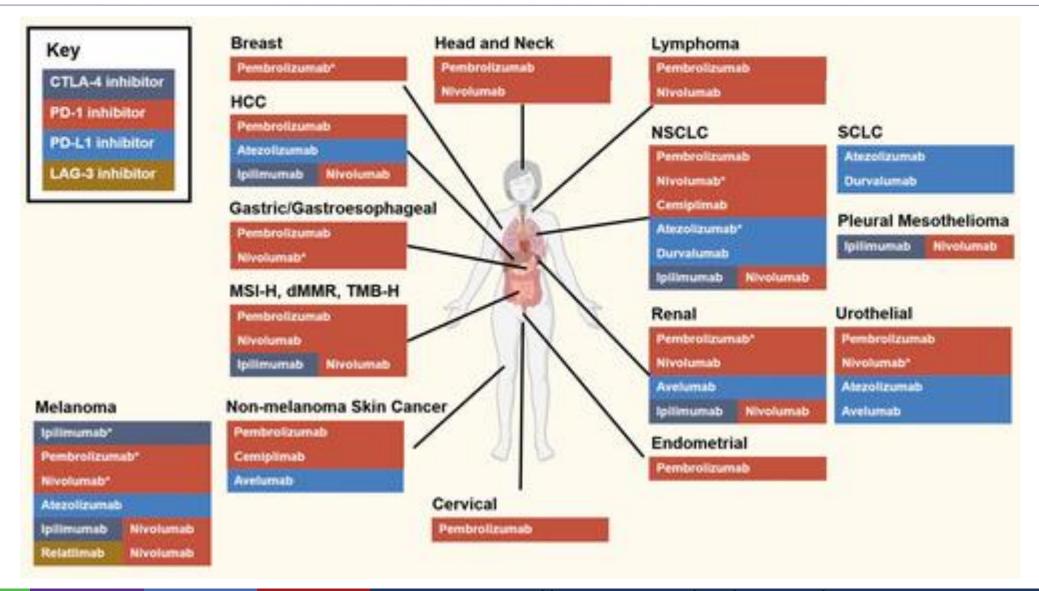
Small Molecules- Targeted Therapy





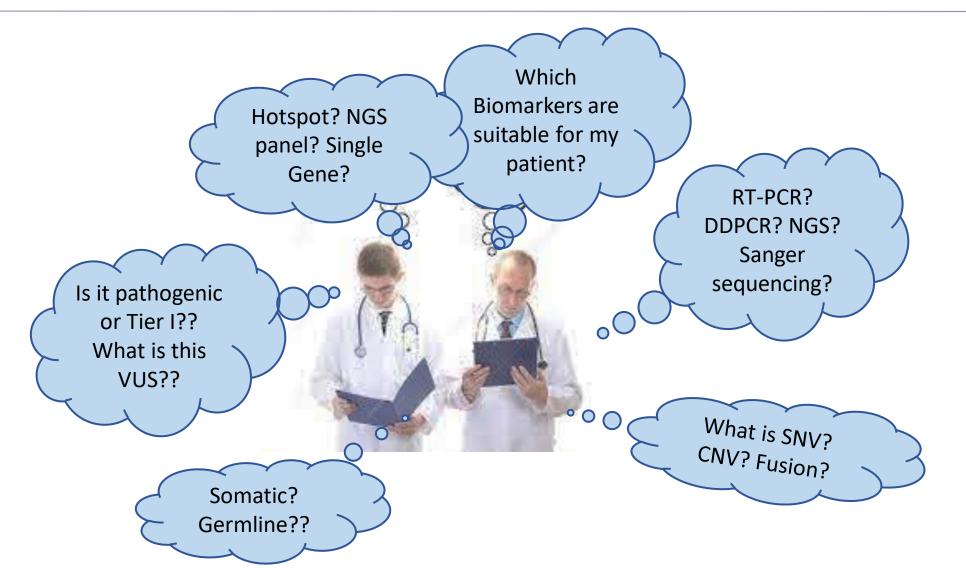
Immunotherapy: Immune checkpoint inhibitors





Detection of Biomarkers- Physician's Conundrum





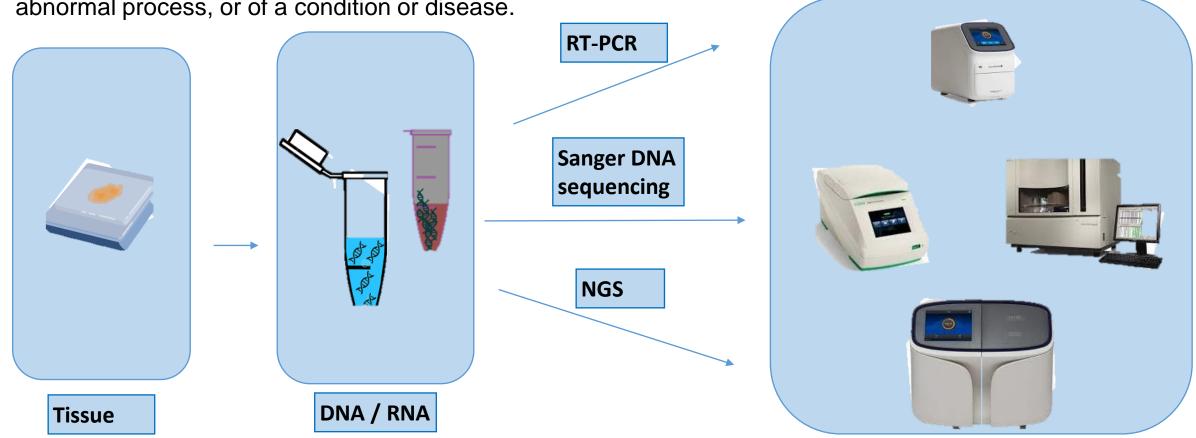
Molecular Diagnostics



Molecular Diagnostics: The process of identifying a disease by studying molecules, such as proteins, DNA, and RNA, in a tissue or fluid.

Bio-Marker: A biological molecule found in blood, other body fluids, or tissues that is a sign of a normal or

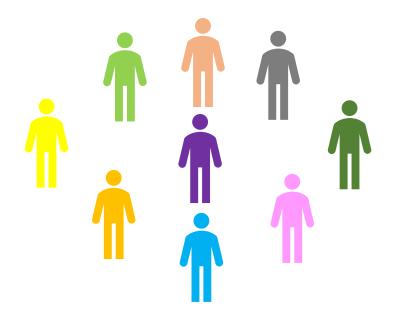
abnormal process, or of a condition or disease.



Why Target Testing

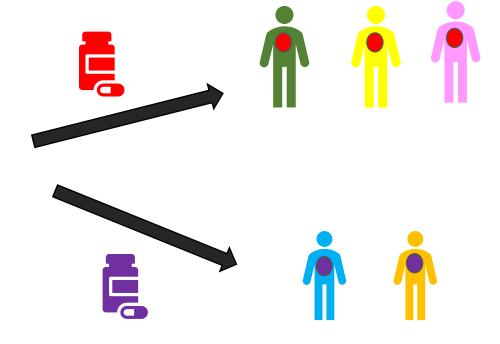


- Identify target in the patient to decide therapy options
- Enrich Patient pool which can benefit from target therapy

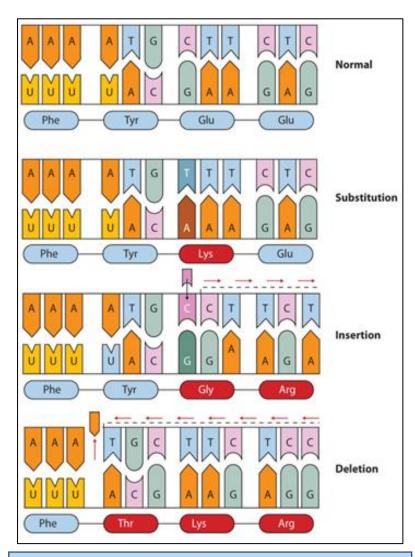


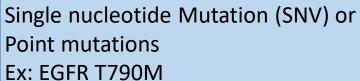


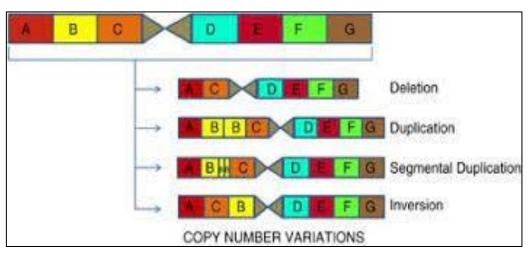
Biomarker/ Target Identification



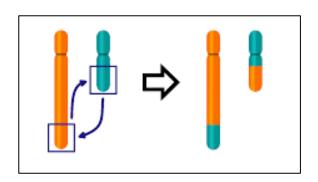
What are these "Targets"? What are their types?



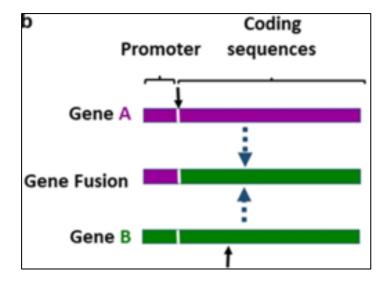




CNV –Copy Number Variants Ex: MET amplification



Translocations
Ex: BCR-ABL (Ph chromosome)



Gene Fusion Ex: ALK/ROS

Targets/ Biomarkers for Immunotherapy



Biomarkers in Use

PD-L1 Expression

- · Useful only in certain tumors
- Predictive for anti-PD-1/PD-L1

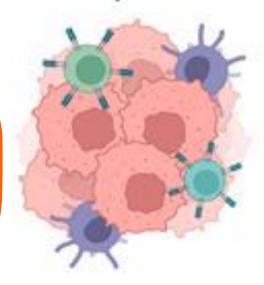
Tumor Mutational Burden

- Increases neoantigen expression
- Predictive for anti-CTLA-4 and anti-PD-1/PD-L1

Microsatellite Instability

 Reflects hypermutability
 Predictive for anti-CTLA-4 and acti-PD-1/PD-L1

Predictive Biomarkers of ICI Response



Biomarkers Under Investigation

Tumor Microenvironment

 Immune and angiogenesis profiles around tumor cells

Genomic Mutational Profile

 Defects in oncogenes, tumor suppressors and signalling

Gut Microbiome

 Better response with increased diversity, anabolic pathways and specific bacteria

Neutrophil/Lymphocyte Ratio

- Obtained from peripheral blood
- Imperfect and lacks standardized threshold

1. Tumor Mutation Burden- TMB



Tumor mutation burden and immune response

Tumor mutation burden (TMB) refers to the number of genetic changes (mutations) in a cancer cell. The immune system can identify cancer cells and activate an immune response by detecting these mutations.

Cancers with low mutation burden (TMB-low) have fewer mutations, decreasing the chance that one will activate the immune system.

Immune cells may not identify cancer cells for many reasons, including lack of appropriate markers on the cell surface.

Cold Tumor

Low TMB

Hot Tumor

High TMB – Better ICI response

Cancers with high tumor mutation burden (TMB-high) have more mutations, increasing the chance that at least one will activate an immune response.

TMB-HIGH
CANCERCELL

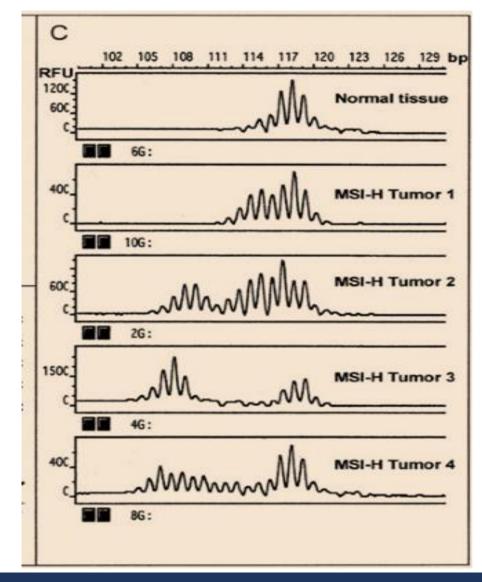
Immune cells can potentially identify cancer cells from specific markers that may be present on the cell surface due to cancer-related mutations.

TMB can be measured in a lab using detailed molecular testing of tumor biopsy tissue. Studies are currently evaluating the ability to also measure TMB from tumor genetic material in the blood, making testing easier and more accessible.

2. Microsatellite Instability (MSI)



- Microsatellites are regions of repeated DNA that change in length (show instability) when mismatch repair is not working properly.
- MSI High / low/ stable
- MSI-H: Predictive for ICI therapy
- MSI-H: eligibility to Pembrolizumab, ipilimumab, nivolumab treatment of colorectal cancer patients
- MSI-H: Diagnostic- Lynch Syndrome
- MSI-H = mismatch repair deficient (dMMR)
- dMMR is defined as at least 1 protein (MSH 2, MSH 6, PMS 2 and MLH 1) showing loss of expression IHC



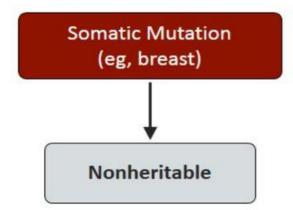
Mutations Types based on Origin



Somatic vs Germline Mutations

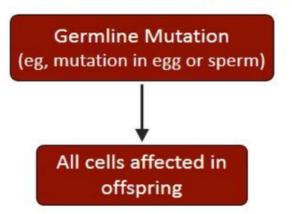
Somatic mutations

- Occur in non-germline tissues
- Are nonheritable



Germline mutations

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



Somatic Testing = Tumor testing. Samples: FFPE block, FFPE slides, Liquid biopsy, Whole blood in streck/paxgene tube for ctDNA analysis

Germline Testing: Testing for heredity. Sample: Blood in EDTA tube

How are these Targets detected: Molecular Techniques



Technique

Application

RT-PCR

Detection/ Quantification of a marker eg. BCR/ABL IS %

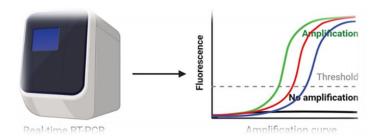
DD-PCR

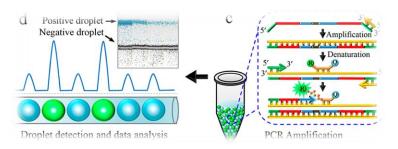
Highly sensitive detection/ Quantification of a marker eg. EGFR T790M mutation

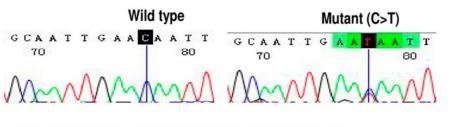
Sanger Sequencing Detection multiple mutations on a gene eg. JAK2 Exon 12, KRAS mutations

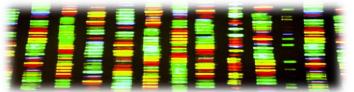
Next Generation Sequencing

Detection multiple mutations on multiple genes eg. All genes involved in lung cancer



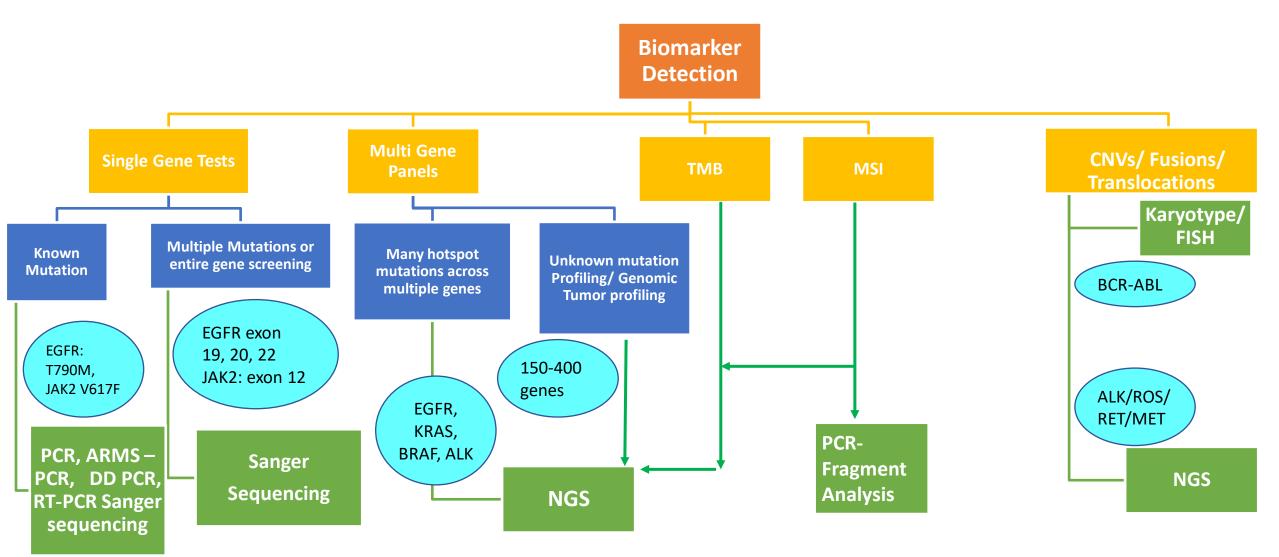






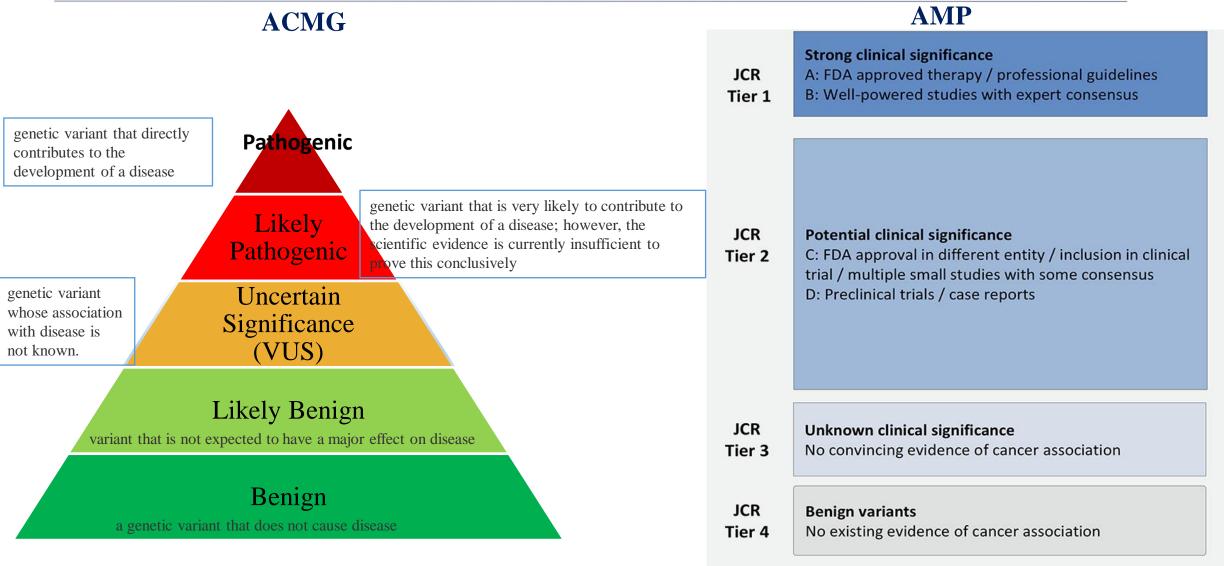
Target- Technique correlation: How to choose?





Reporting: Variant Categories





Applying Biomarker/ Target testing in Cancers



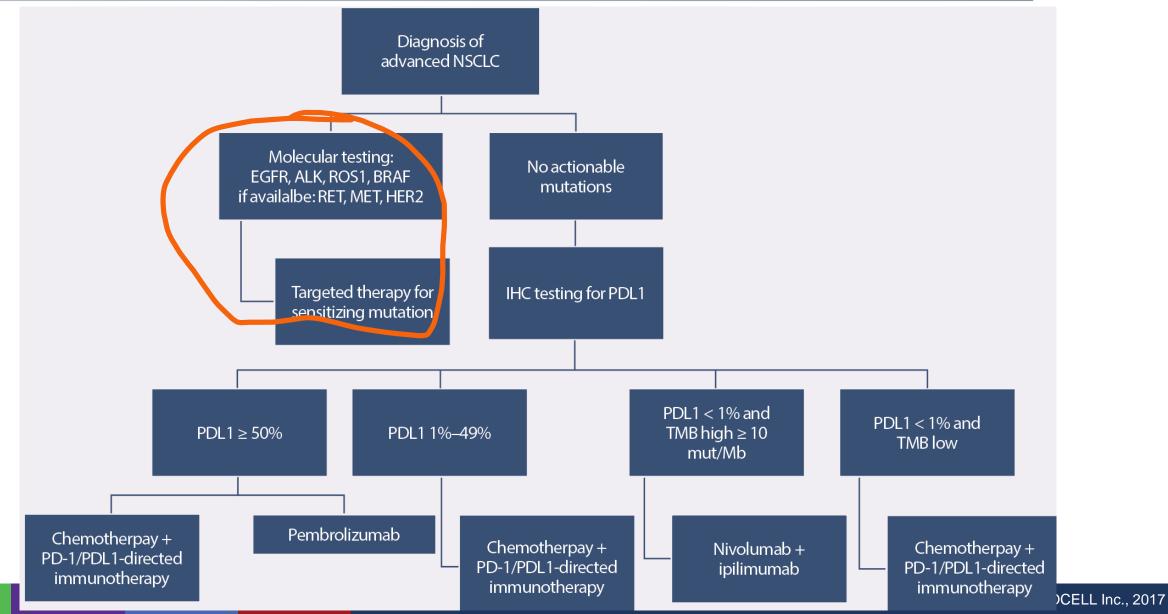


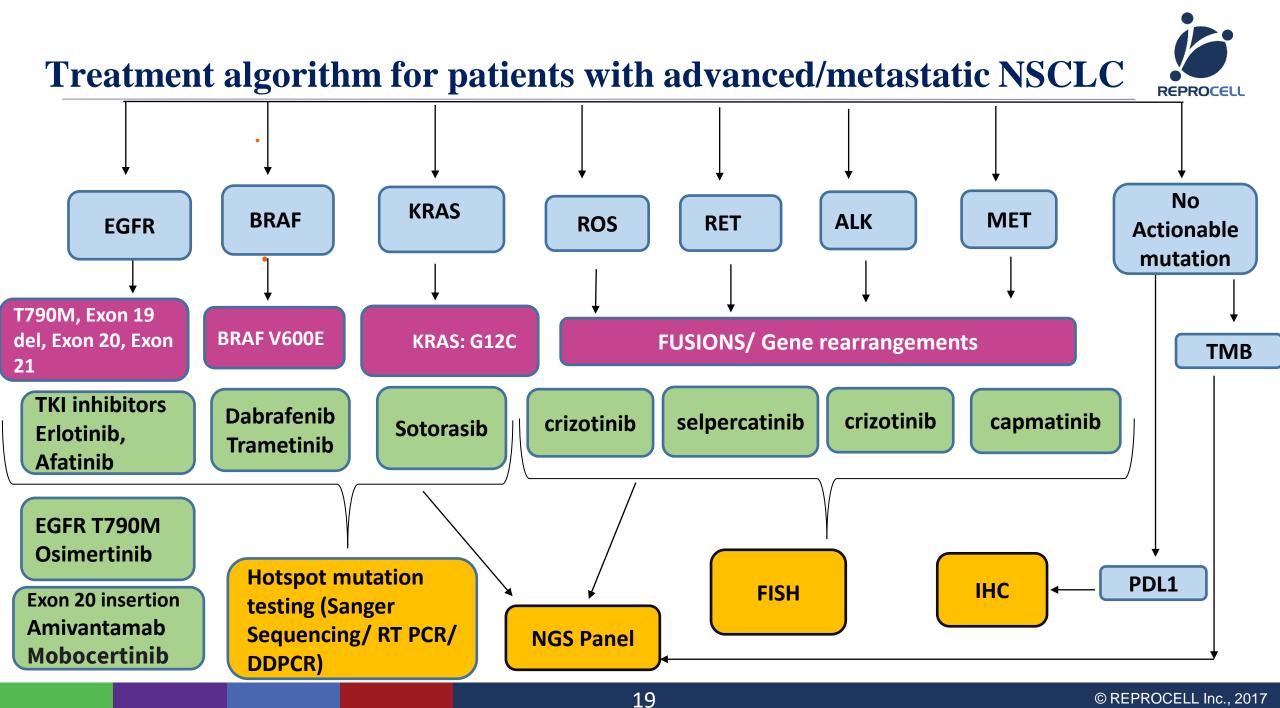
- Lung Cancer
- Breast Cancer
- Colorectal Cancer



Treatment Algorithm for Non-Small Cell Lung Cancer









BIOMARKERS FOR SYSTEMIC THERAPY IN METASTATIC BREAST CANCER

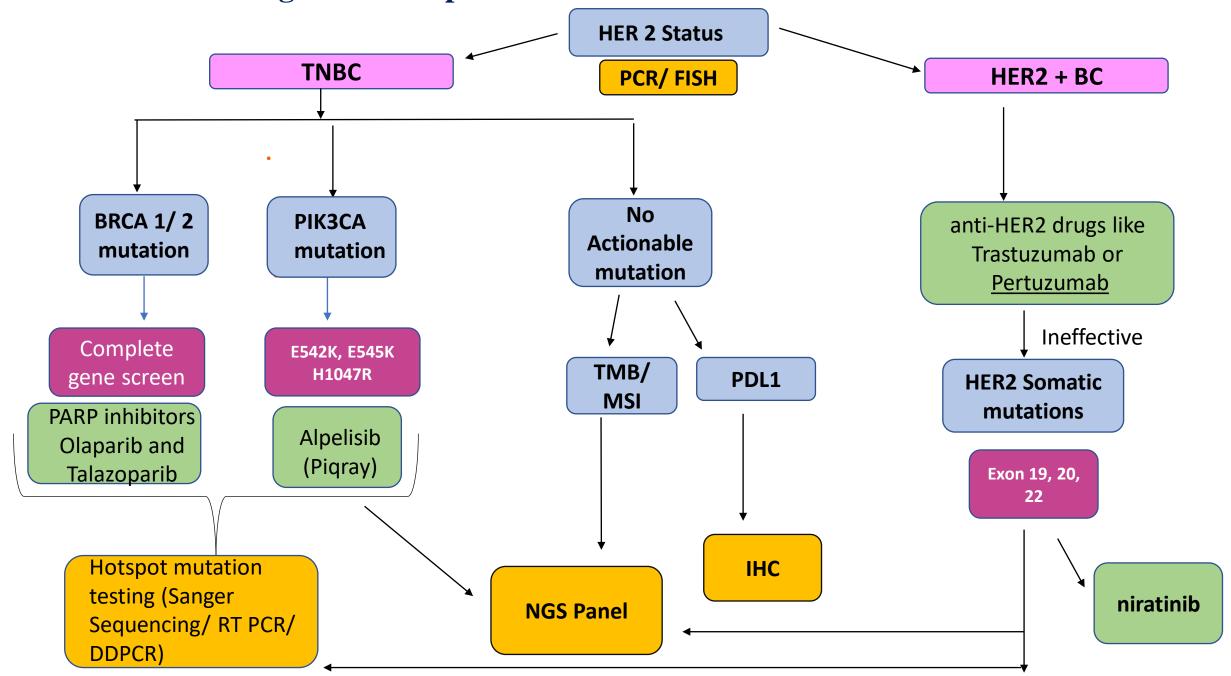
GUIDELINE UPDATE AT-A-GLANCE RECOMMENDATIONS

Test	Type of Recommendation	Quality of Evidence	Strength of Recommendation
Biomarker Tests Recommended	by the ASCO Expert Panel		
• PIK3CA	Evidence-based	• High	Strong
 Germline BRCA1 and BRCA2 	 Evidence-based 	High	Strong
• PD-L1	 Evidence-based 	 Intermediate 	Strong
 dMMR/MSI-H 	 Informal consensus-based 	• Low	Moderate
• TMB	 Informal consensus-based 	• Low	Moderate
 NTRK Fusions 	 Informal consensus-based 	• Low	Moderate
Biomarker Tests Not Recommen	ded by the ASCO Expert Panel		
• ESR1	Evidence-based	Insufficient	Moderate
• PALB2	Evidence-based	• Low	Moderate
• HRD	 Informal consensus-based 	• Low	Moderate
TROP2 expression	 Informal consensus-based 	• Low	Moderate
• ctDNA	 Informal consensus-based 	• Low	Moderate
• CTCs	 Informal consensus-based 	• Low	Moderate

Henry et al J Clin Oncol 2022 asco.org/breast-cancer-guidelines

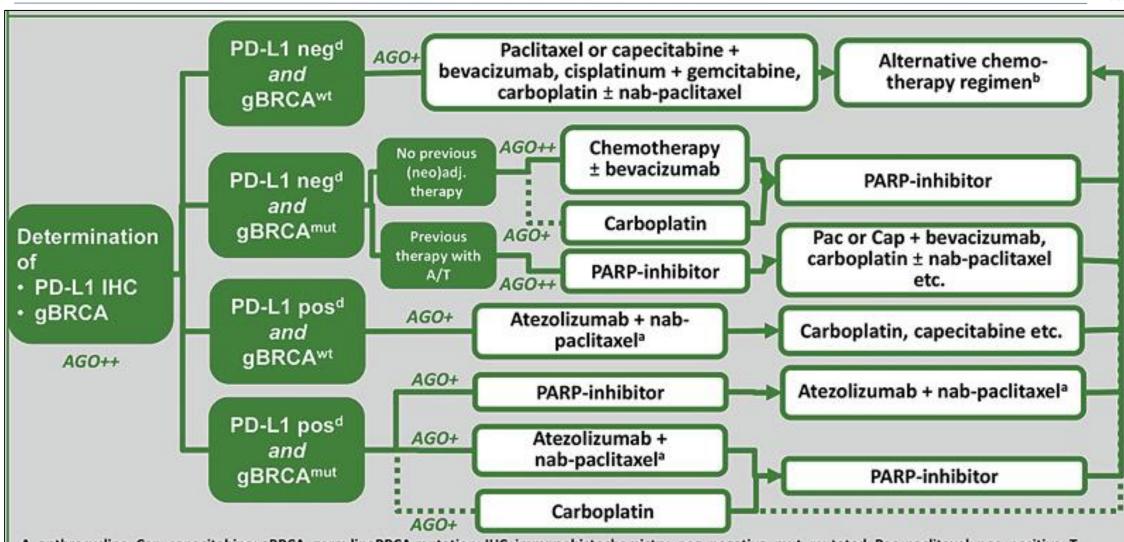
Abbreviations. CTCs, circulating tumor cells; ctDNA, circulating tumor DNA; dMMR, deficient mismatch repair, HRD, homologous recombination deficiency; MSI-H, microsatellite instability-high; NTRK, neutotrophic tyrosine receptor kinase; PD-L1, programmed cell death ligand-1; TMB, tumor mutational burden, TROP2, trophobiast cell-surface antigen 2

Treatment algorithm for patients with advanced/metastatic Breast cancer



Treatment Algorithm for Triple Negative Breast Cancer





https://www.karger.com/Article/Fulltext/511925

BRCA: Who Should get Tested?



PREDISPOSITION TEST

- A history of Breast/ Ovarian cancer in family
- A history of breast cancer at a young age in two or more blood relatives, such as your parents, siblings or children
- A close male relative with breast cancer
- A relative with a known BRCA1 or BRCA2 mutation

PATIENT TEST

- •A personal history of Breast / Ovarian cancer diagnosed before age 45
- •A personal history of triple negative breast cancer diagnosed at age 60 or younger
- •A personal history of two or more types of cancer
- •A personal history of breast cancer and one or more relatives with breast cancer diagnosed before age 50

ESMO consensus guidelines CRC patient Management



Recommendation 4: RAS testing

- RAS mutational status is a negative predictive biomarker for therapeutic choices involving EGFR antibody therapies in the metastatic disease setting [I, A]
 RAS testing should be carried out on all patients at the time of diagnosis of mCRC [I, A]
- · RAS testing is mandatory before treatment with the EGFR-targeted monoclonal antibodies cetuximab and panitumumab [I, A]
- A network of arrangements should be established to ensure the rapid and robust transit of tissue samples from referral centres to testing laboratories, to
 minimise the turnaround time and avoid delays in having this information available for all patients with mCRC
- Primary or metastatic colorectal tumour tissue can be used for RAS testing (see also Recommendation 3)
- RAS analysis should include at least KRAS exons 2, 3 and 4 (codons 12, 13, 59, 61, 117 and 146) and NRAS exons 2, 3 and 4 (codons 12, 13, 59, 61 and 117)

Recommendation 5: BRAF testing

Tumour BRAF mutation status should be assessed alongside the assessment of tumour RAS mutational status for prognostic assessment (and/or potential selection for clinical trials) [I, B]

Recommendation 6: Microsatellite instability testing

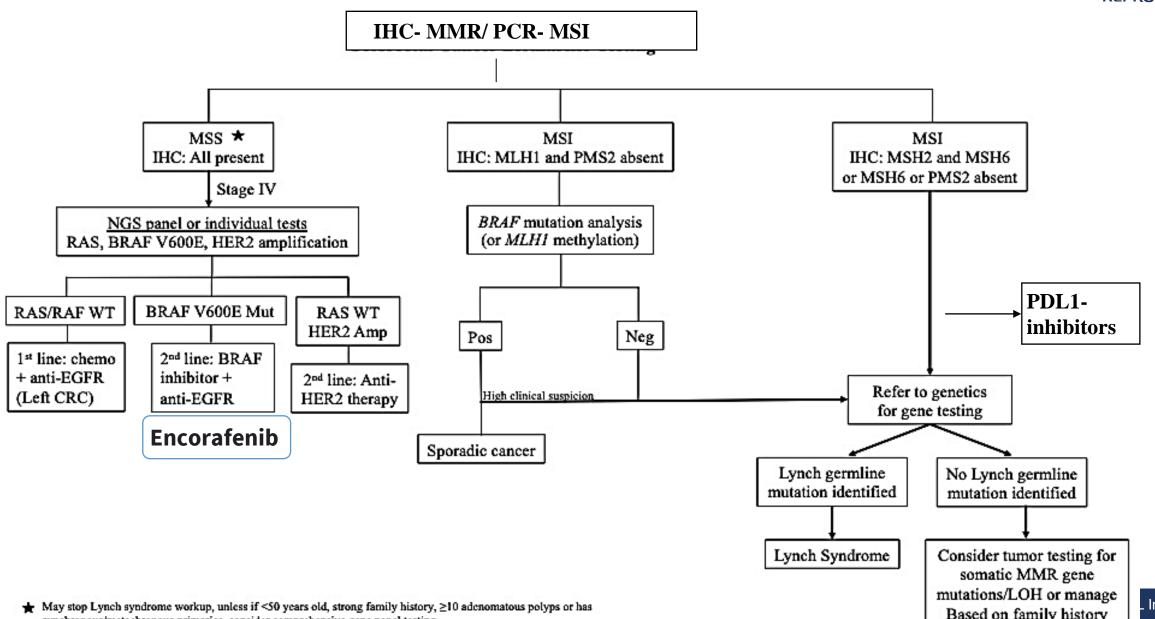
- MSI testing in the metastatic disease setting can assist clinicians in genetic counselling [II, B]
- . MSI testing has strong predictive value for the use of immune check-point inhibitors in the treatment of patients with mCRC. [II, B]

Recommendation 7: Biomarkers of chemotherapy sensitivity and toxicity

- DPD testing before 5-FU administration remains an option but is not routinely recommended [II, D]
- UGT1A1 phenotyping remains an option and should be carried out in patients with a suspicion of UGT1A1 deficiency as reflected by low conjugated bilirubin and in patients where an irinotecan dose of >180 mg/m² per administration is planned [95] [III, C]

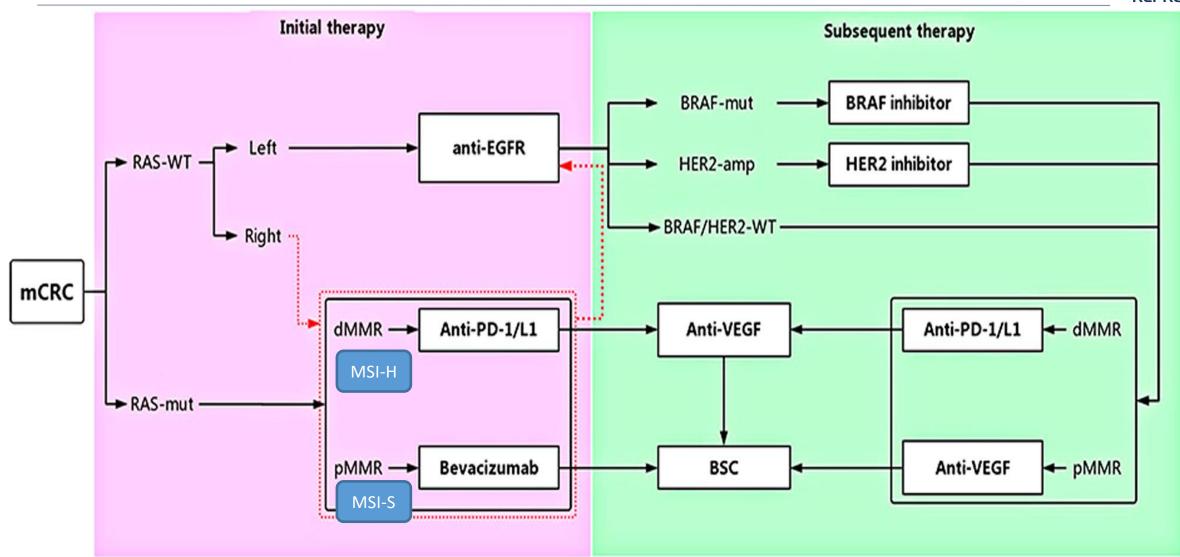
Treatment Algorithm in m-CRC





NCCN-recommended strategy for m-Colorectal cancer targeted therapy

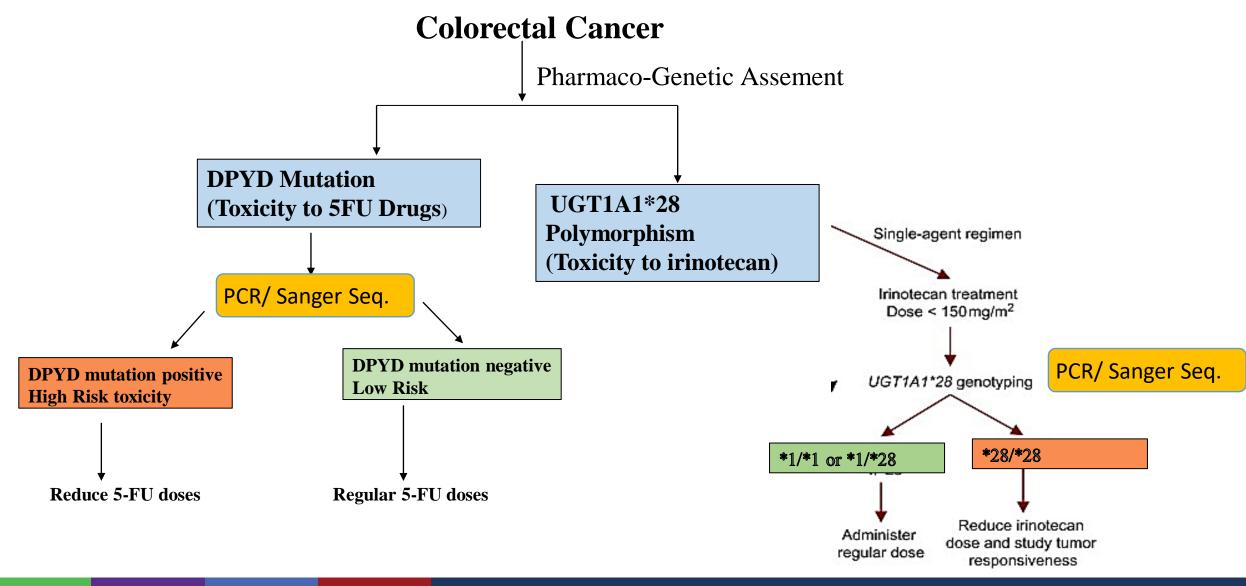




https://www.nature.com/articles/s41392-020-0116-z/figures/4

Biomarkers for CT sensitivity/ Toxicity- CRC





Bioserve: Biomarker Testing



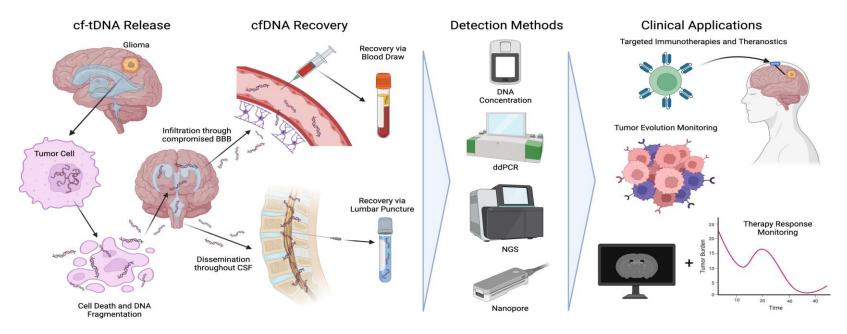
Marker	Cancer	Technique
EGFR, KRAS,NRAS, BRAF	LUNG, CRC, MELANOMA	Sanger sequencing, RT-PCR
Met Exon 14 skipping	Lung	PCR
RET/ ROS/ ALK/NTRK FUSIONS	LUNG, CRC	FISH, NGS
BRCA 1/ 2 MUTATIONS	BREAST	NGS
PIK3CA MUTAIONS	BREAST	Sanger Sequencing
IDH1/ 2 MUTATIONS	GIOMA, AML	Sanger Sequencing
KIT MUTATIONS	GIST	Sanger Sequencing
BCR/ABL TRANSCRIPT/FUSION	CML	FISH, RT-PCR
JAK2 MUTAIONS	MPN	Sanger Sequencing
HRR GENE MUTATIONS	BREAST, PROSTATE	NGS
TUMOUR MUTATION BURDEN	SOLID TUMORS	NGS
MSI	CRC, LUNG, BREAST	PCR+ Fragment Analysis

Marker	Cancer	Technique
PAN-CANCER LIQUID BIOPSY PANEL	PAN CANCER	NGS
CEBPA, NPM1, FLT3 mutations	AML	Sanger sequencing, RT-PCR
TERT mutation	Melanoma	Sanger Sequencing
HER 2 MUTATIONS	BREAST	Sanger Sequencing
IMANITIB RESISTANCE	CML	Sanger Sequencing
HER 2 AMPLIFICATION	BREAST, CRC	FISH, RT-PCR
MGMT METHYLATION ASSAY	GLIOMA	RT-PCR
CALR, MPL MUTATIONS	MPN	Sanger Sequencing
Comprehensive genomic profiling	Solid tumors	NGS

Future Directions



Liquid Biopsy for Biomarker Detection and residual Disease monitoring

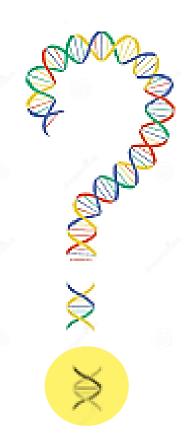


Gut microbiome- predict immune therapy response

- gut microbiota influences anti-tumor immunity
- thereby impacting the clinical responses and outcomes of the patients receiving cancer immunotherapy
- FMT in combination with checkpoint inhibitors are able to reprogram the tumor microenvironment and activate host immunity with favorable changes in immune cell infiltrates in patients with prostate cancer, melanoma, gastrointestinal and prostate cancer.
- NCT04758507, NCT03353402, NCT04130763, NCT05094167

Thank You!





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