

Precision Medicine in Metastatic NSCLC

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Professor and Chair

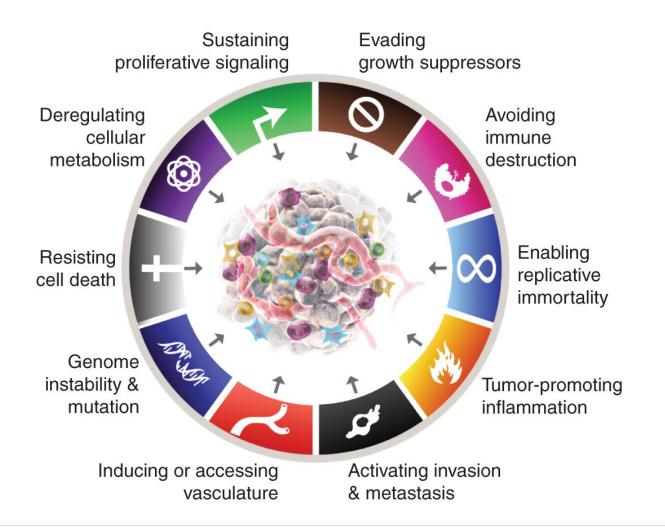
Department of Medical Oncology and Therapeutics Research

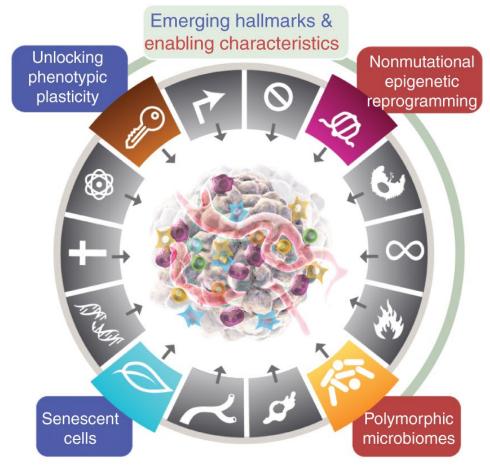


Objectives

- Precision Medicine at City of Hope
- Lung Cancer Overview
- Precision Medicine Overview
- Therapeutic Strategies
- Germline Testing and Strategies
- Artificial Intelligence for Precision Medicine

Hallmarks of Cancer: New Dimensions

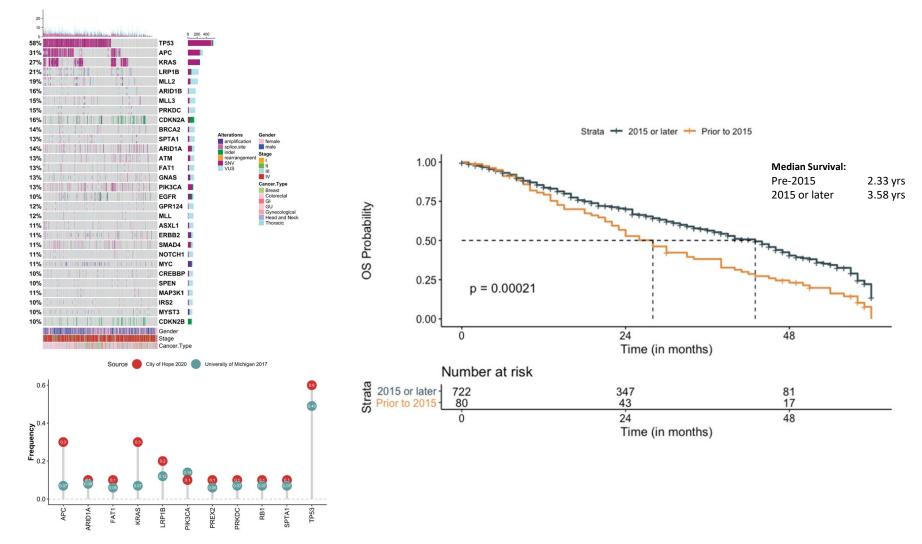




Hanahan, Cancer Discov, 2022

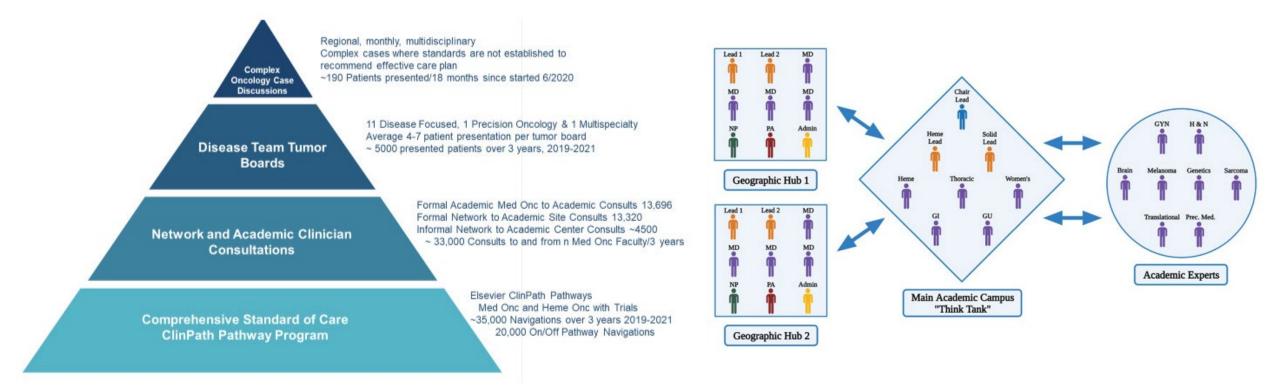


Precision Medicine in Solid Tumors at COH



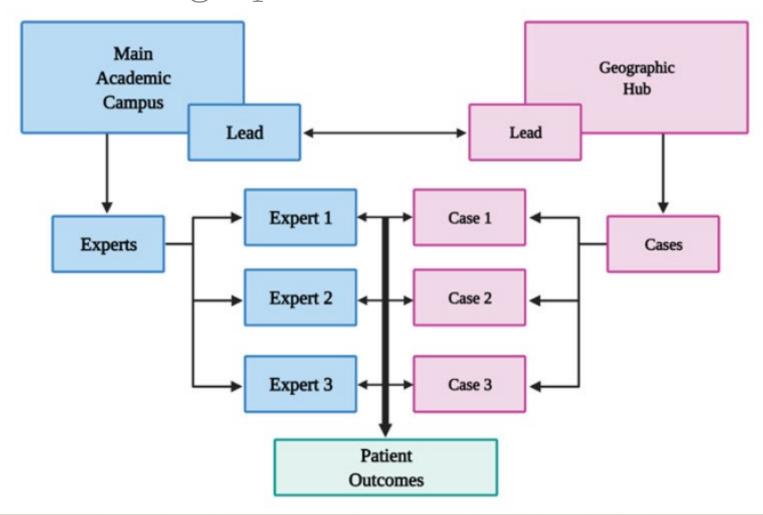
Roosan et int. Salgia, Cancers, 2021

COH Enterprise: Decision Support Network



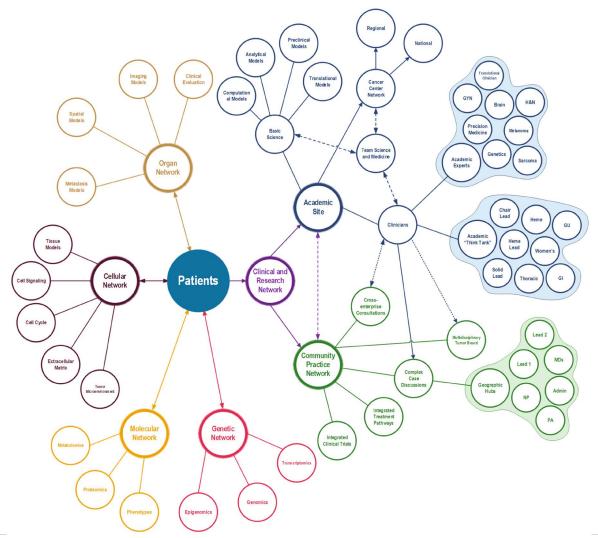
Bosserman et int. Salgia, JCM, 2022

Complex Oncology Case Discussion Algorithm-Academic and Geographic Network

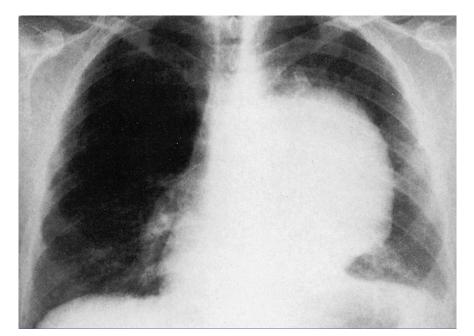


Bosserman et int. Salgia, JCM, 2022

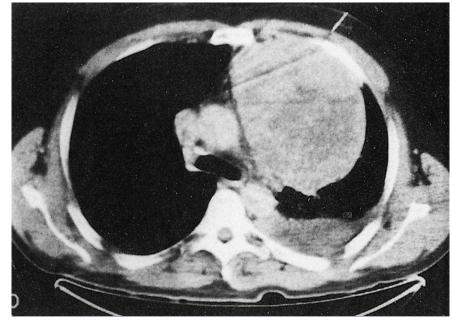
Enhancing the Patient Network

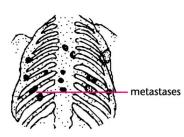


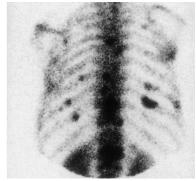
Lung Cancer













Salgia et al., 2010

Lung Cancer Incidence and Mortality

Ма	ale	Fema	Female				
Prostate	299,010	29%	Breast	310,720	32%		
Lung & bronchus	116,310	11%	Lung & bronchus	118,270	12%		
Colon & rectum	81,540	8%	Colon & rectum	71,270	7%		
Urinary bladder	63,070	6%	Uterine corpus	67,880	7%		
Melanoma of the skin	59,170	6%	Melanoma of the skin	41,470	4%		
Kidney & renal pelvis	52,380	5%	Non-Hodgkin lymphoma	36,030	4%		
Non-Hodgkin lymphoma	44,590	4%	Pancreas	31,910	3%		
Oral cavity & pharynx	41,510	4%	Thyroid	31,520	3%		
Leukemia	36,450	4%	Kidney & renal pelvis	29,230	3%		
Pancreas	34,530	3%	Leukemia	26,320	3%		
All sites	1,029,080		All sites	972,060			

Lung & bronchus	65,790	20%
Prostate	35,250	11%
Colon & rectum	28,700	9%
Pancreas	27,270	8%
Liver & intrahepatic bile duct	19,120	6%
Leukemia	13,640	4%
Esophagus	12,880	4%
Urinary bladder	12,290	4%
Non-Hodgkin lymphoma	11,780	4%
Brain & other nervous system	10,690	3%
All sites	322,800	

Male

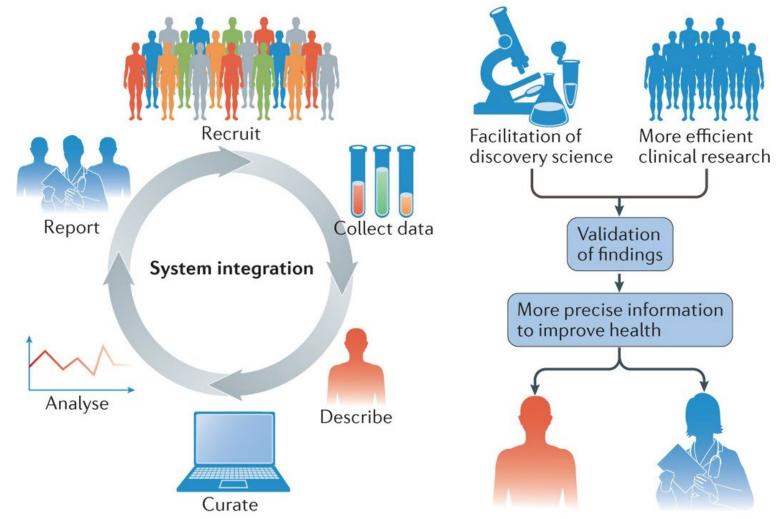
59,280	21%
42,250	15%
24,480	8%
24,310	8%
13,250	5%
12,740	4%
10,720	4%
10,030	3%
8,360	3%
8,070	3%
288,920	
	42,250 24,480 24,310 13,250 12,740 10,720 10,030 8,360 8,070

Female

Siegel et al., CA Cancer J. Clin., 2024

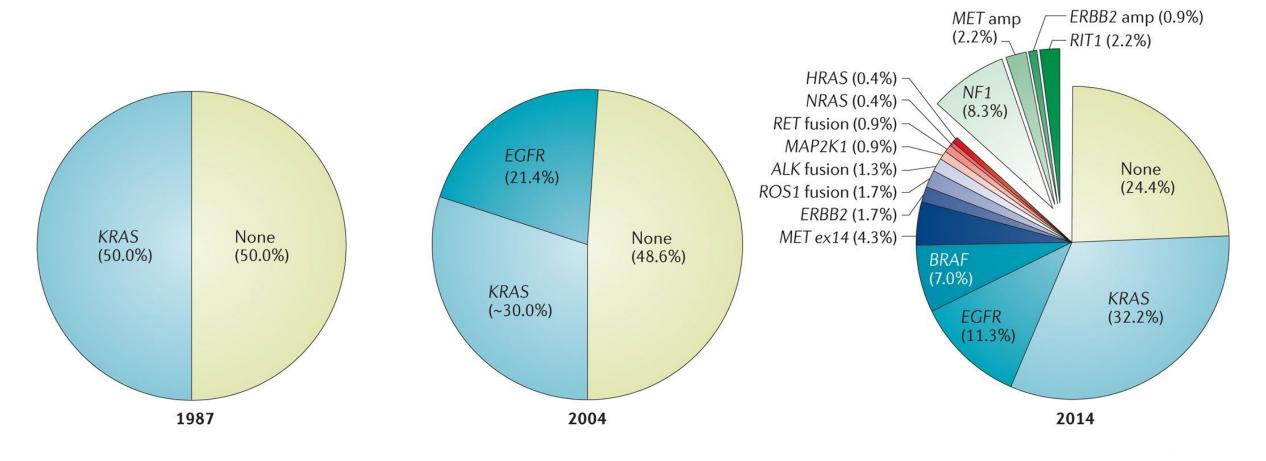
Estimated Deaths

Precision Medicine System and Goals

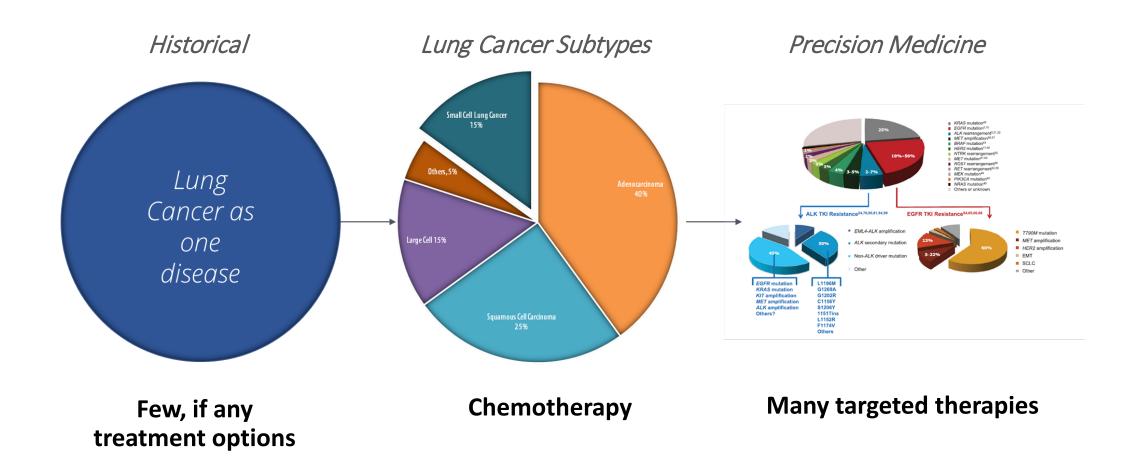


Antman and Loscalzo, Nature Reviews Cardiology, 2016

Lung Cancer Genomic Knowledge Has Evolved

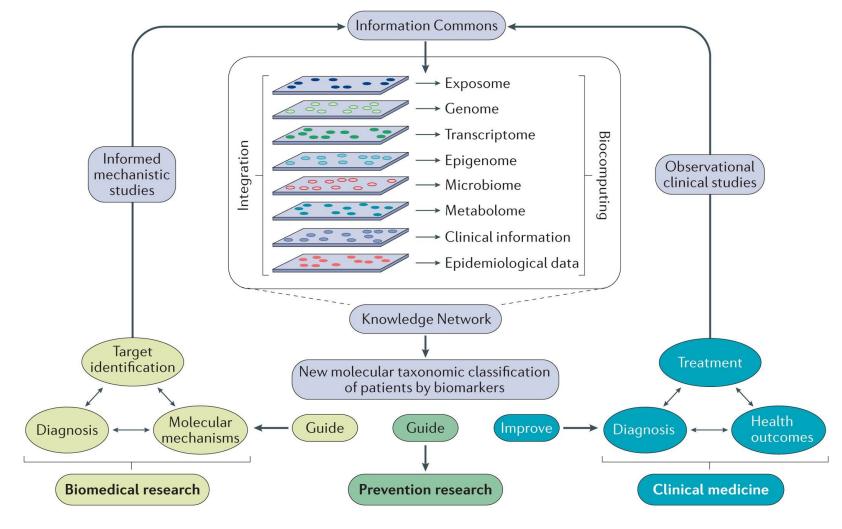


Lung Cancer- Therapeutic Evolution



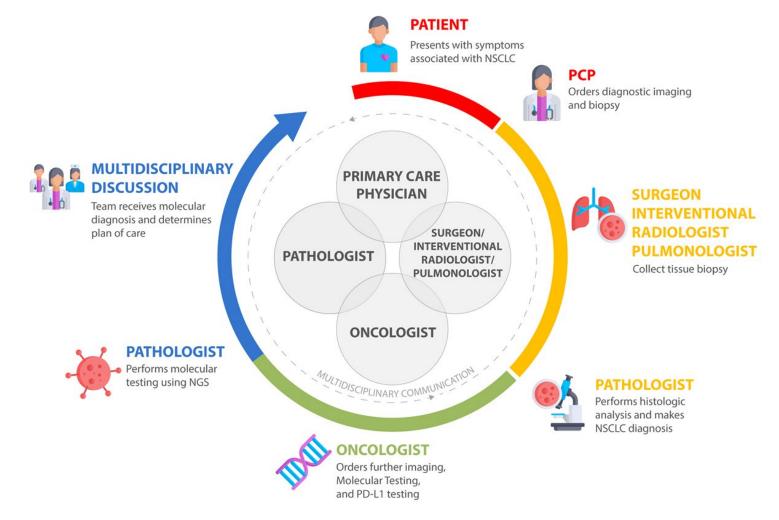
Salgia, 2019

Lung Cancer-Precision Medicine Strategy



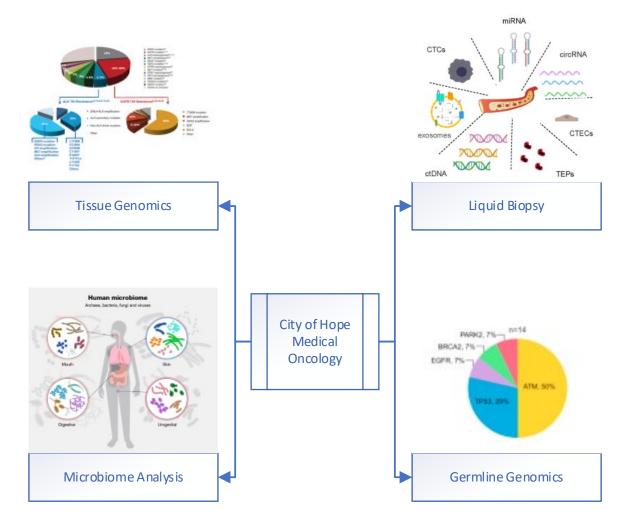
Vargas and Harris, Nature Reviews Cancer, 2016

Precision Medicine Multidisciplinary Care Model



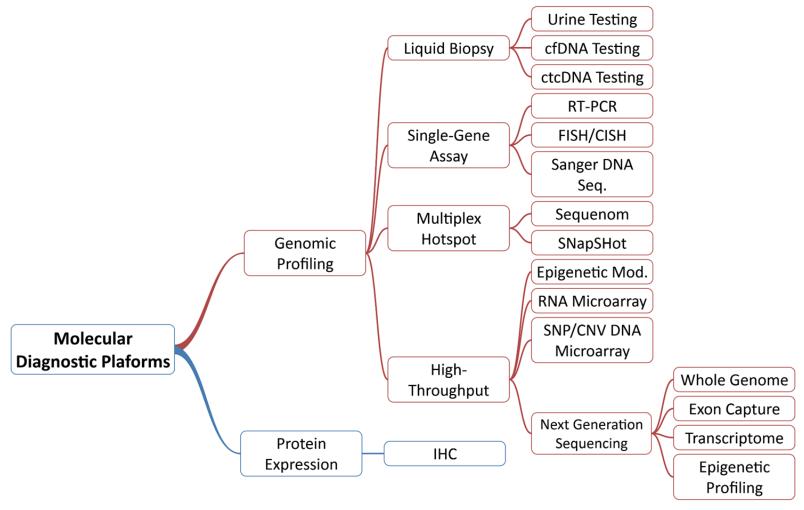
Salgia and Mambetsariev, J. Clin. Med., 2020

Precision Medicine Tools



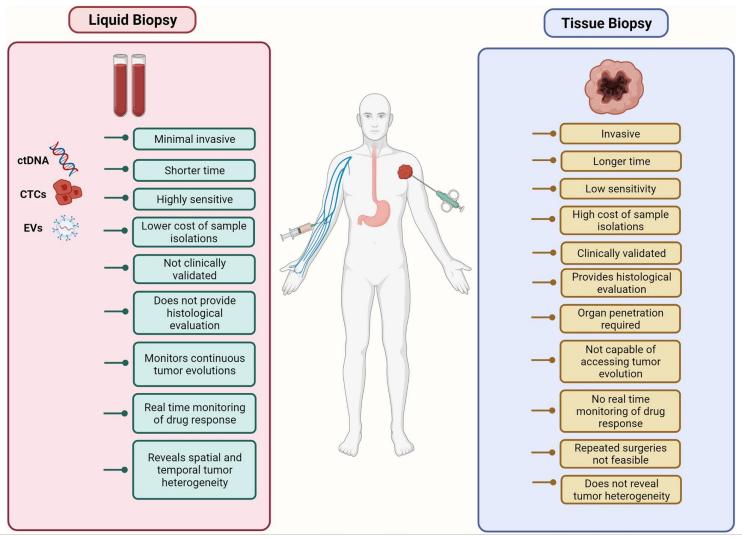
Salgia, 2024

Precision Medicine- Diagnostic Platform Options



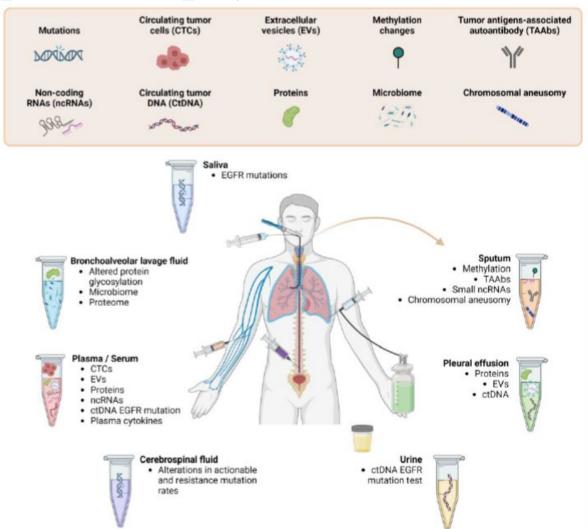
Hensing, Mambetsariev, Salgia; 2017; in Pass et al. IASLC Thoracic Oncology 2nd Ed.

Liquid Biopsy Versus Tissue Biopsy



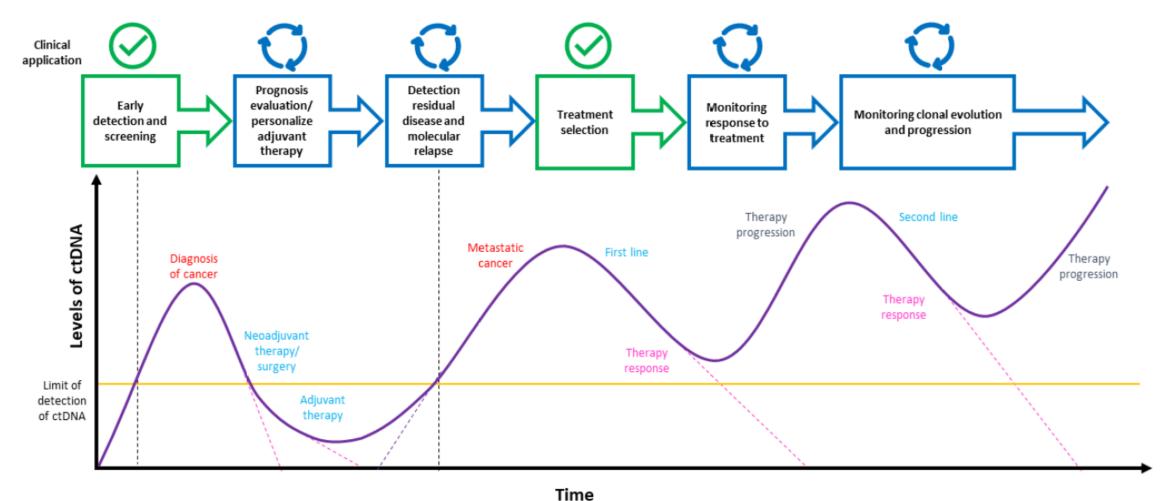
Lone et al., Mol Cancer, 2022

Utility of Liquid Biopsy in Cancer



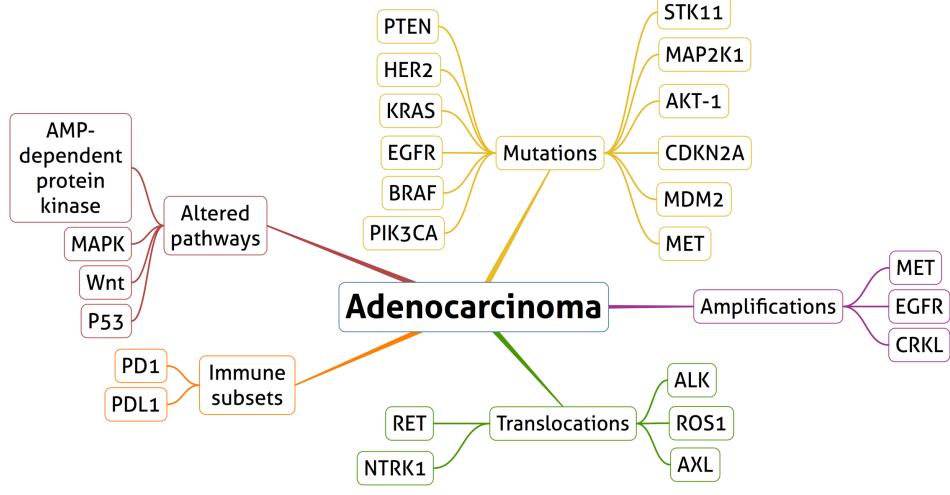
Brockley et al., Cancers, 2023

Clinical Applications of Liquid Biopsies



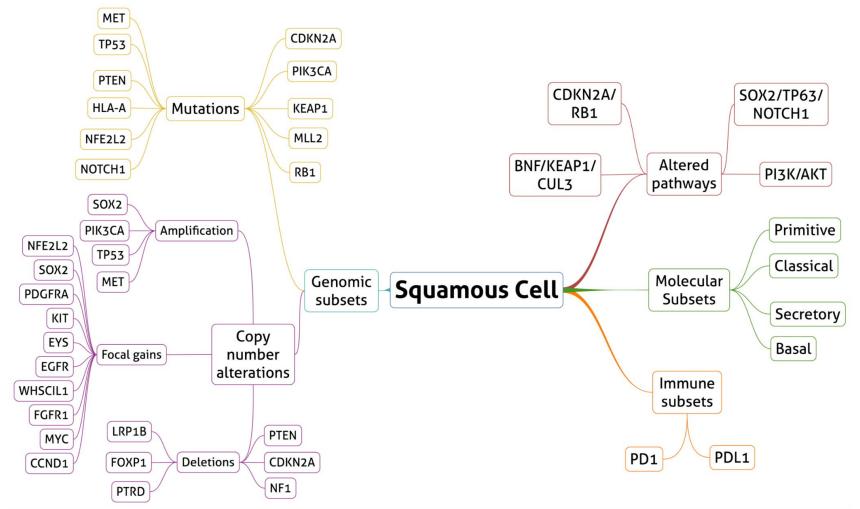
Caputo et al., Explor Target AntitumorTher., 2022

Lung Cancer- Adenocarcinoma Heterogeneity



Hensing, Mambetsariev, Salgia; 2017; in Pass et al. IASLC Thoracic Oncology 2nd Ed.

Lung Cancer- Squamous Heterogeneity



Hensing, Mambetsariev, Salgia; 2017; in Pass et al. IASLC Thoracic Oncology 2nd Ed.

Lung Cancer- Heterogeneity



Mambetsariev et int. Salgia, PlosOne, 2020

When to consider genomic tumor testing for "NSCLC"

Initial Presentation

- o Is there enough tissue for molecular testing?
- o Is a biopsy required?
- o Is there pleural fluid for potential biopsy?

Features of a particular patient/scenario to consider:

- Histology (Squamous cell carcinoma vs Non-Squamous)
 - If squamous, does it have adenocarcinoma features?
- Stage
 - Stage I-III vs Stage IV
- o Age
 - Is the patient younger than 50 or older than 65
- Prior Testing/Treatment
 - Previous history of other cancer types
 - Previous molecular testing results (should you re-biopsy?)
 - Previous treatment for early-stage vs recurrence

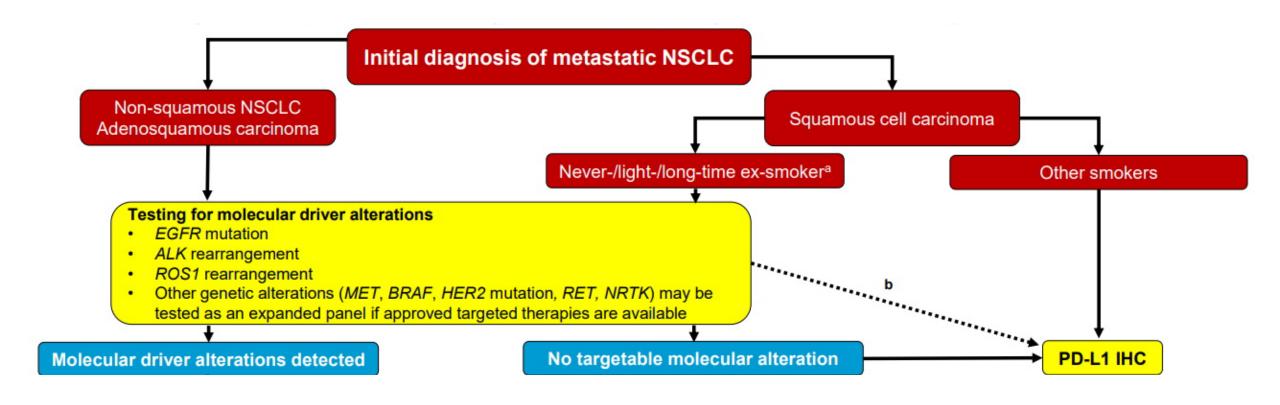
NSCLC Biomarker Testing Rates

Nonsquamous NSCLC	Decisions in Academic Setting (n = 51)	Decisions in Community Setting (n = 253)	Total # of Decisions (N = 304)	
ALK Test Status				
Awaiting test results	6	74	80	
Did not order test	0	19	19	
Negative	42	157	199	
Positive	3	3	6	
ALK testing rate, n (%)	51 (100)	234 (92)	285 (94)	
EGFR Test Status				
Awaiting test results	6	76	82	
Did not order test	0	16	16	
Negative/wild type	19	100	119	
Non-sensitizing	7	27	34	
Sensitizing	19	34	53	
EGFR testing rate, n (%)	51 (100)	237 (94)	288 (95)	
ROS1 Test Status				
Awaiting test results	6	72	78	
Did not order test	0	37	37	
Negative	43	142	185	
Positive	2	2	4	
ROS1 testing rate, n (%)	51 (100) ^a	216 (85) ^a	267 (88)	
Nonsquamous and Squamous NSCLC	Decisions in Academic Setting (n = 45)	Decisions in Community Setting (n = 282)	Total # of Decisions (N = 327) ^b	
PD-L1 Test Status				
Negative	15	57	72	
Positive (≥ 50% TPS)	8	55	63	
Positive (1%-49% TPS)	5	47	52	
Did not order test	17	123	140	
PD-L1 testing rate, n (%)	28 (62)	159 (56)	187 (57)	

Reported Study	EGFR	ALK	ROS1	MET	RET	NTRK	BRAF	KRAS	PD-L1 Expression
Inal et al. [66]	62%	23%	N/A	N/A	N/A	N/A	N/A	43%	N/A
Gutierrez et al. [67]	69%	65%	25%	15%	14%	N/A	18%	34%	N/A
Gierman et al. [68]	54%	51%	43%	N/A	N/A	N/A	29%	N/A	N/A
Presley et al. [69]	100%	95%	~15%	~15%	~15%	~15%	~15%	~15%	~15%
Illei et al. [94]	N/A	53.1%	N/A						
Hussein et al. [95]	~60%	~50%	N/A						
Mason et al. [29]	94%	92%	85%	N/A	N/A	N/A	N/A	N/A	56%
Audibert et al. [105]	68%	67%	32%	6%	8%	0%	12%	0%	N/A
Khozin et al. [142]	64%	61%	N/A	N/A	N/A	N/A	N/A	N/A	8.3%
Nadler et al. 2018 [143]	37%	35%	N/A	N/A	N/A	N/A	N/A	N/A	1.2%
Nadler et al. 2019 [106]	35.5%	32.9%	5.7%	N/A	N/A	N/A	0.1%	N/A	5.7%

Mason et al., J Clin Pathw, 2018 Rajurkar et int. Salgia, JCM, 2020

Stage IV NSCLC Molecular Testing Algorithm by Histology



Chong-Kin et al., Respirology, 2020

Genomic Tumor Testing in Lung Cancer

What to consider when choosing test type for these patients:

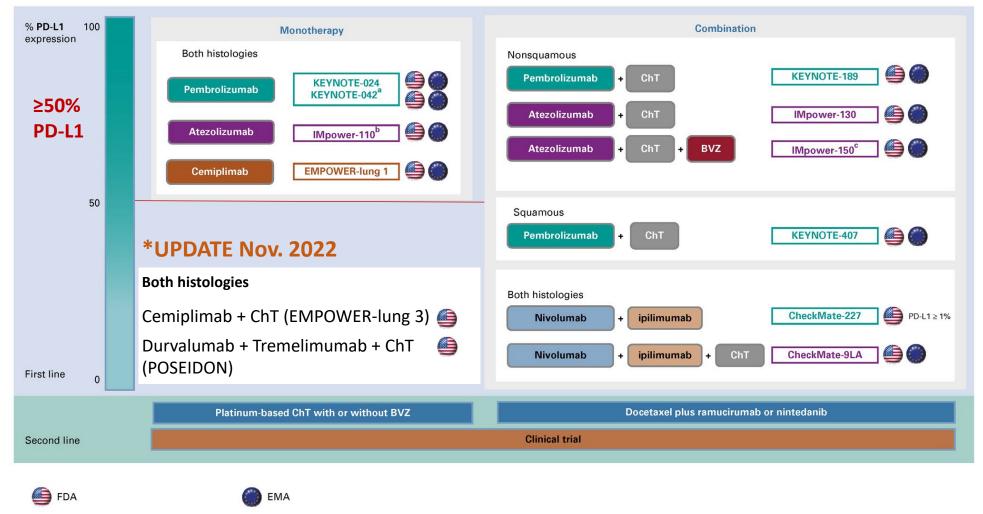
- Comprehensive panels tissue biopsy
 - HopeSeq
 - FoundationOne CDX
 - o Tempus
 - CARIS
 - Neogenomics
- Liquid Biopsy
 - o Blood
 - Guardant 360
 - FoundationOne Liquid CDX
 - Tempus
 - Neogenomics
 - Cerebrospinal Fluid
 - Biocept
- Specific markers (e.g. PDL-1)
 - Full NGS panel (EGFR, ALK, KRAS, MET, ROS1, BRAF, NTRK, RET, HER-2, and more with up to >450 genes)
 - o IHC (PD-L1)
 - 22C3 (Pembrolizumab, Cemiplimab)
 - SP142 (Atezolizumab)
 - 28-8 (Nivolumab)
 - SP263 (Durvalumab)

Interpreting Biomarker Testing in Lung Cancer

- What to consider when a patient has an actionable mutation?
 - o EGFR, ALK, KRAS G12C, MET, BRAF V600E, RET, ROS1, NTRK fusion, HER2
 - o Is there a rare subtype such as EGFR exon 20, exon 18, or MET fusion?
 - Are there any co-mutations such as KRAS-KEAP1-STK11 or EGFR-RB1-TP53?
- PD-L1 results
 - 0% PD-L1 negative

 - ≥50% PD-L1 positive (high expression)
 - Are there any mutations that may cause hyper-progression? (KRAS-KEAP1-STK11 or actionable mutations such as EGFR)
- Do any of the genomic results dispute the histologic diagnosis?
- Germline results:
 - o BRCA-1 or BRCA-2 becoming more common in lung cancer
 - ATM, PARK2, TP53, EGFR and others

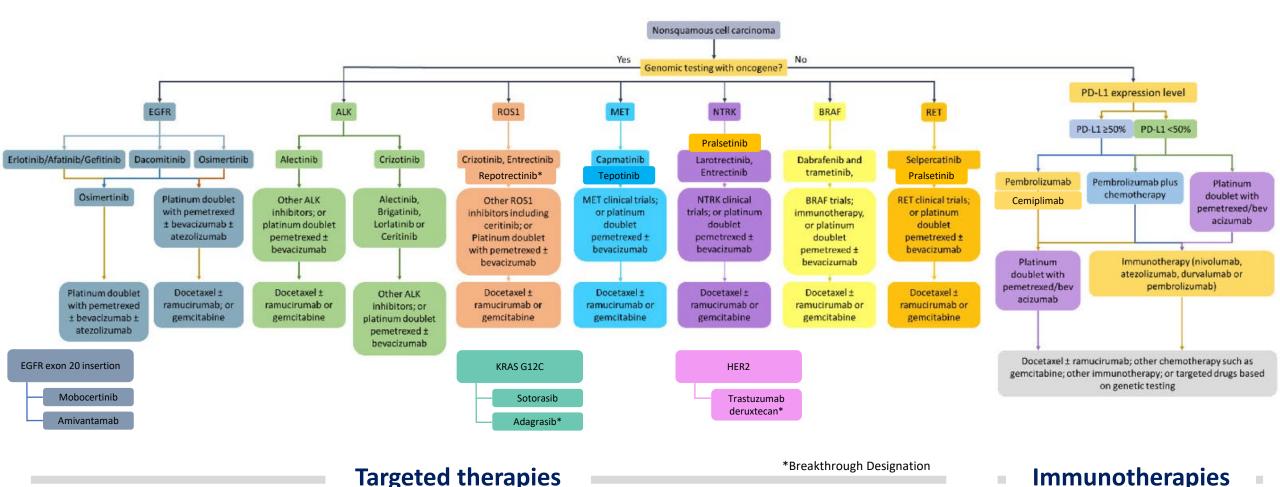
Immunotherapy: Stage IV NSCLC



Reck, Remon, and Hellmann; J. Clin. Oncol., 2022

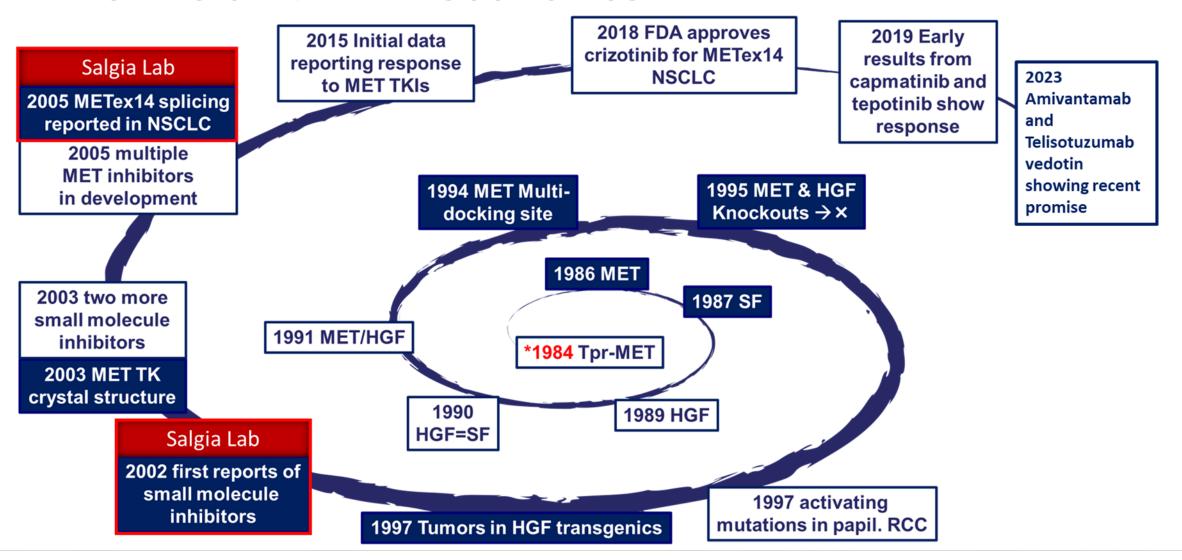
CITY OF HOPE

Lung Cancer Targeted Therapeutic Strategies



Rajurkar et int. Salgia, JCM, 2020 (Updated 2022)

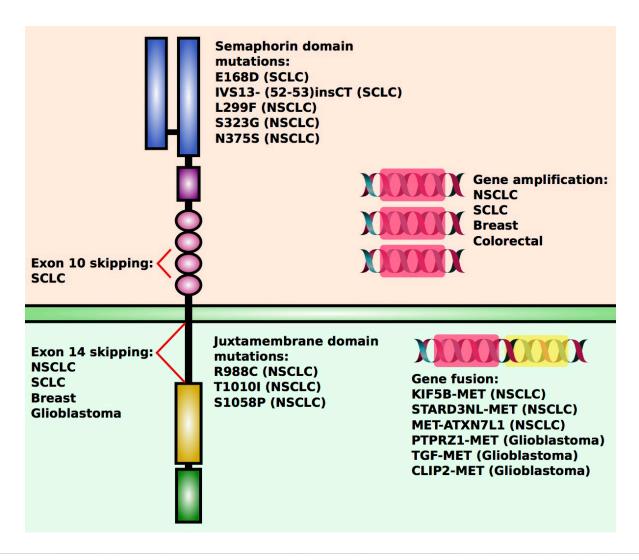
Timeline of MET Discoveries



CITY OF HOPE

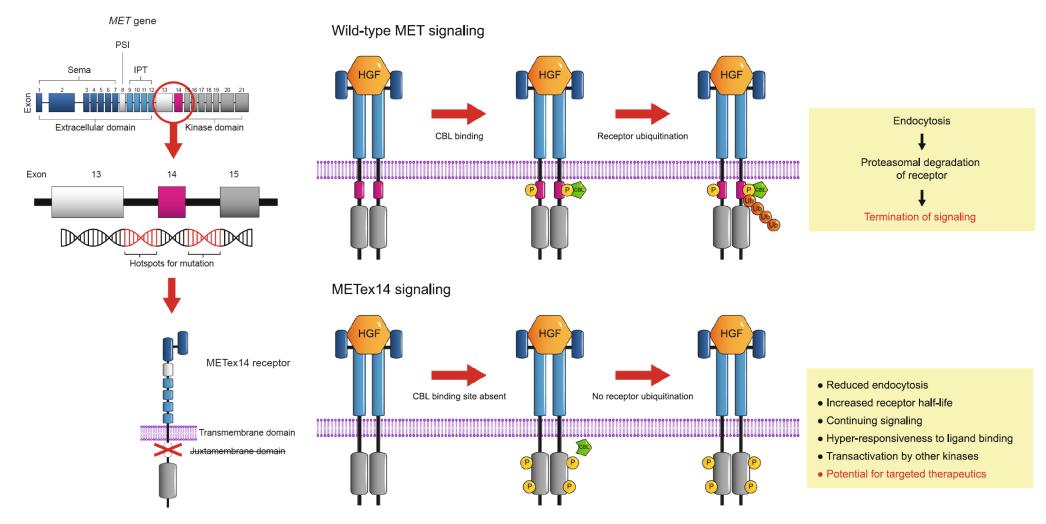
31

Genetic Alterations of MET Found in Solid Tumors



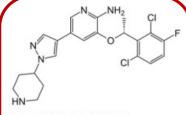
Malik, et int. Salgia, ACR 2020

MET Exon 14 Mutation and Downstream Signaling



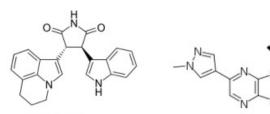
Salgia R., Sattler M., et al. Cancer Treatment Reviews, 2020

New Generation of Small-molecule MET Inhibitors



Crizotinib (PF-02341066)

- FDA-approved as an ALK inhibitor
- Inhibit MET, ALK, ROS1 and RON
- IC₅₀ for MET: 0.0002 μM



Tivantinib (ARQ197)

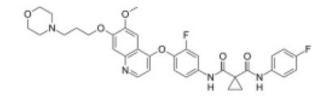
- Phase II combined with erlotinib in NSCLC
- Inhibit MET
- IC50 for MET: N/A

Savolitinib (HMPL-504; AZD6094)

- Phase II with NSCLC
- Inhibit MET
- IC₅₀ for MET: 0.003 μM

Cabozantinib (XL184, BMS907351)

- FDA-approved for renal and thyroid cancers
 Inhibit MET, VEGFR2, AXL and RET
- IC₅₀ for MET: 0.0013 μM
- Phase II with NSCLC

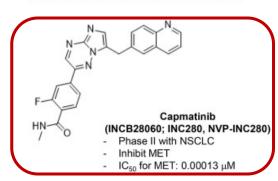


Foretinib (XL880; EXEL-2880; GSK1363089; GSK089)

- Inhibit MET and VEGFR2
- IC₅₀ for MET: 0.0004 μM
- Phase I in combination with erlotinib in NSCLC

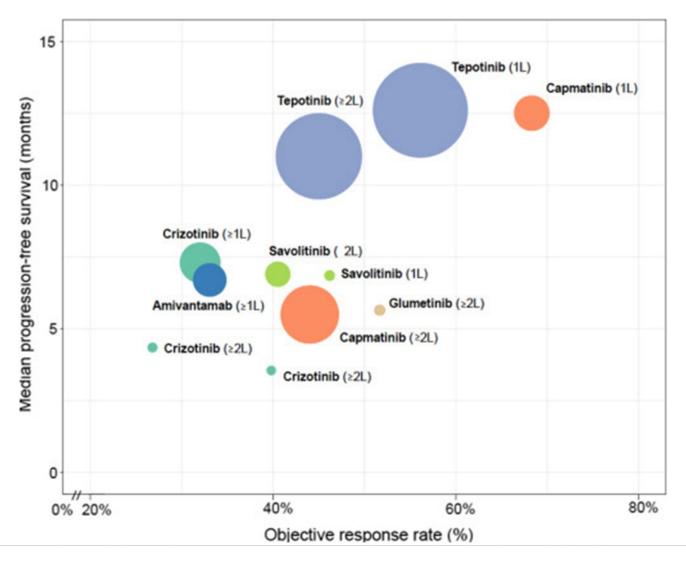
Tepotinib (EMD 214063)

- Phase II with MET-altered NSCLC
- Inhibit MET
- IC₅₀ for MET: 0.001 μM



Wang et al. J Hematol Oncol, 2019

Efficacy of MET Inhibitors



Remon et al., JCO, 2023

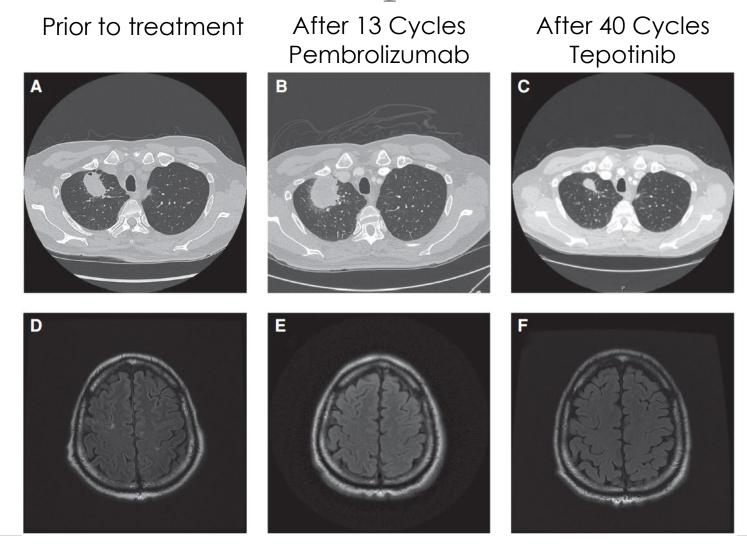
Case #1: MET

- 66-year-old male, never smoker, initially presented with dyspnea on exertion, and a cardiac MRI revealed biventricular masses consistent with metastatic disease.
- Additionally, CT showed a 5.7 cm RUL mass, subcentimeter pulmonary nodules, mediastinal and hilar lymphadenopathy, hepatic masses, bilateral adrenal nodules, and lytic lesions.
- Brain MRI showed multiple small sub-centimeter lesions.
- Liver biopsy confirmed stage IV lung adenocarcinoma (T3N3M1c) with PD-L1 70%, but as insufficient for molecular testing.

Case #1: MET Cont.

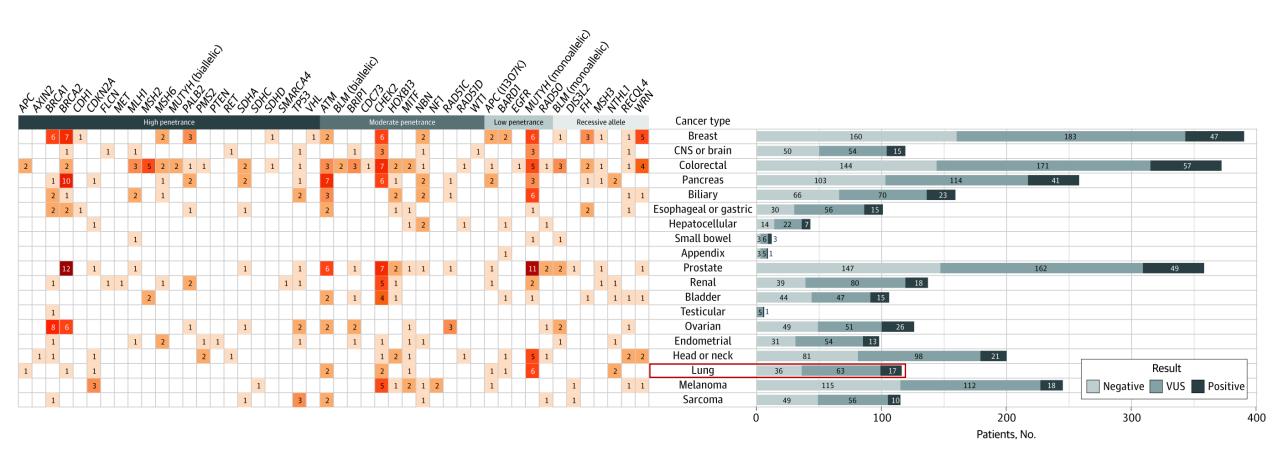
- He started Carboplatin/Paclitaxel for two cycles with Bevacizumab for the second cycle and developed G3 fatigue and G3 neuropathy; therapy was switched to Pembrolizumab.
- NGS was performed on a LN EBUS and revealed MET exon 14 splice site mutation.
- He developed PD in the chest and enrolled in a clinical trial with Tepotinib.
- The time course of the disease in both the chest and brain can be seen in (**Figure 1**).
- Patient continues treatment with Tepotinib (currently on cycles 83-84).

Case #1: Treatment Response

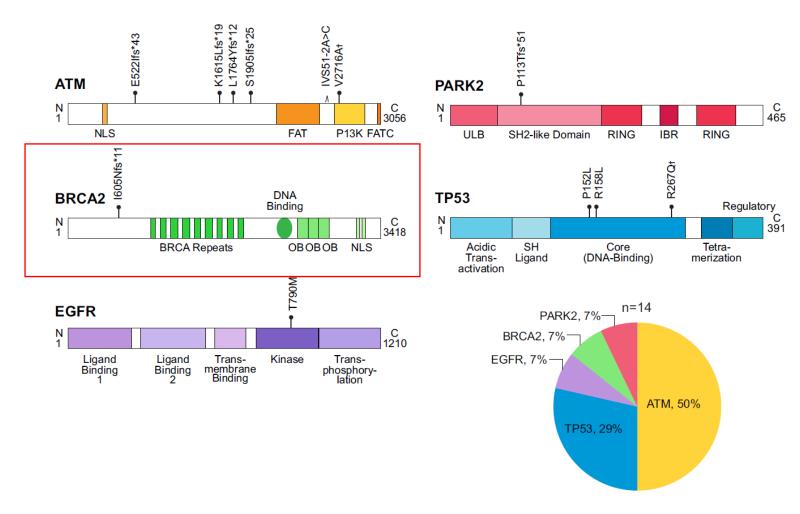


Roth et int. Salgia, Cold Spring Harb. Mol. Case Stud. 2020

Lung Cancer- Germline Mutations Prevalent



Germline Variants in Lung Adenocarcinoma Cases



Parry et al., Journal of Thoracic Oncology, 2017

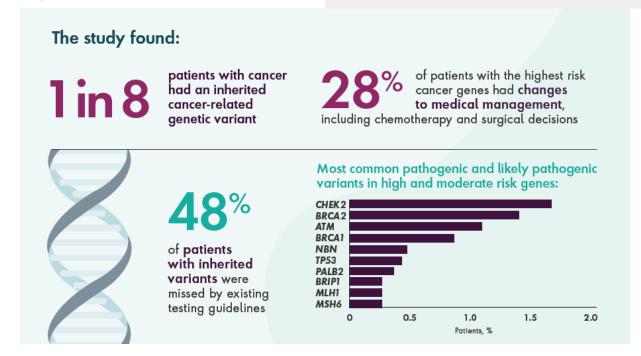
COH-Universal Germline Testing on all patients

Groundbreaking study supports germline genetic testing for <u>all</u> cancer patients

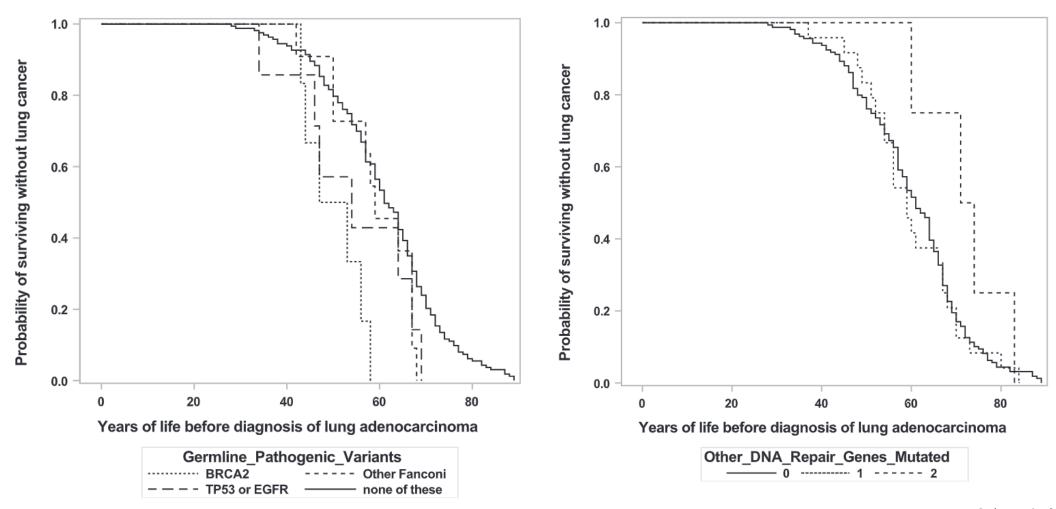
In a study published in JAMA Oncology, Mayo Clinic and Invitae researchers provided genetic testing and counseling to patients as part of their standard cancer care.

The INTERCEPT study

- The largest known multicenter study of universal testing of patients with cancer
- Published in JAMA Oncology
- Includes 2,984 patients with a new or active cancer diagnosis, across a broad mix of solid tumor cancer stages and types



Lung Cancer- Germline Mutations Affect Survival



Reckamp et int. Salgia, Weitzel, Cancer, 2021

Case #2: Lung Cancer BRCA2

- 26-year-old female, no tobacco exposure or significant past medical history, initially presented with dyspnea.
- CTA of the chest demonstrated a RUL mass with mediastinal involvement creating superior vena cava syndrome along with associated pulmonary embolisms and a large right sided pleural effusion secondary to complete right bronchial tree collapse (Figure 1).
- Pleural effusion and biopsy revealed poorly differentiated lung adenocarcinoma.
- MRI brain demonstrated greater than 20 supratentorial and infratentorial lesions with the largest being 1.1 cm.
- PETCT demonstrated innumerable bilateral pulmonary micronodules, multiple osseous lesions including the right seventh rib, spine, sacrum, and left iliac wing (Figure 3).
- Clinically staged as IVB (T3N2M1c) primary lung adenocarcinoma.

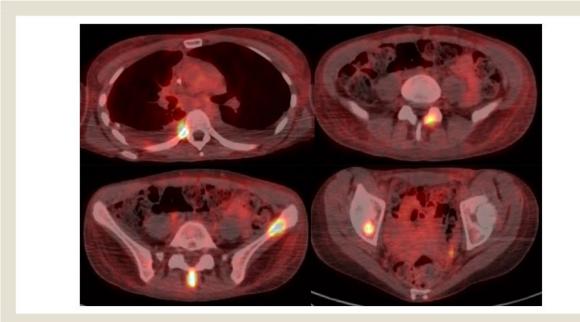
Waddington et int. Salgia, Clinical Lung Cancer, 2021

Case #2: Work Up Imaging

Figure 1 Computed tomography (CT) angiography of the chest demonstrates a right upper lobe mass with mediastinal involvement creating superior vena cava (SVC) syndrome along with associated pulmonary embolisms and a large right sided pleural effusion secondary to complete right bronchial tree collapse.



Figure 3 PET-CT demonstrates innumerable bilateral pulmonary micronodules, multiple osseous lesions including the right seventh rib, spine, sacrum, and left iliac wing.



Case #2: Molecular Findings

- NeoGenomics genetic testing was negative for EGFR, ALK, and ROS1 while PD-L1 (22C3) testing demonstrated high expression with a tumor proportion score of 90%.
- Ashion GEM ExTra with DNA and RNA sequencing was notable for BRCA2 S497*, TMB: Low at 4 muts/Mb and MSI: Stable.
- Liquid biopsy utilizing Guardant360 detected BRCA2 S497* with 51.6% of cfDNA.
- Germline BRCA1/2 analysis with CustomNext-Cancer was positive for the pathogenic mutation BRCA2 S497* (Table 1).
 - o Patient's Mother (aged 47) and maternal half-brother (aged 21) tested positive for the BRCA2 mutation.

Table 1	The Patient's Germline Mutation in the BRCA2 Gene								
Gene	Chromosome	HGVS DNA Reference	HGVS Protein Reference	Variant Type	Predicted Effect	dbSNP/ dbVar ID	Genotype (Heterogous or Homozygous	ClinVarID	
BRCA2	13	NM_000059.3 (BRCA2): c.1490C>G (p.Ser497Ter)	NP_000050.2: p.Ser497Ter	Nonsense, stop-gain	Premature truncation	nsv4449760	Heterozygous	RCV000774812.3	

Waddington et int. Salgia, Clinical Lung Cancer, 2021

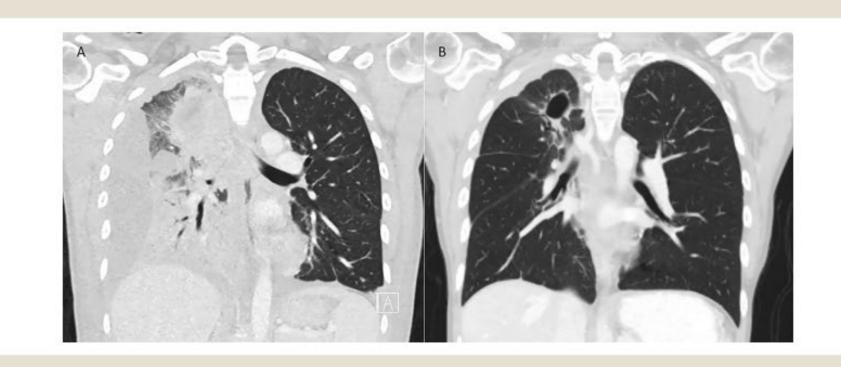
Case #2: Treatment

- Received one cycle of inpatient Carboplatin/Pemetrexed.
- Palliative radiation to her right lung and mediastinum with a total dose of 30 Gy delivered in 10 fractions along with WBRT reaching a total dose of 30 Gy over 10 fractions.
- Initiated combination Carboplatin (AUC 5)/Pemetrexed (500 mg/m2)/Pembrolizumab (200 mg) every 3 weeks for 5 cycles.
- Olaparib (150 mg, BID) was initiated with Pembrolizumab (200 mg), now every 6 weeks, for maintenance therapy (Figure 2).
- Patient continues therapy and is clinically and radiologically stable.

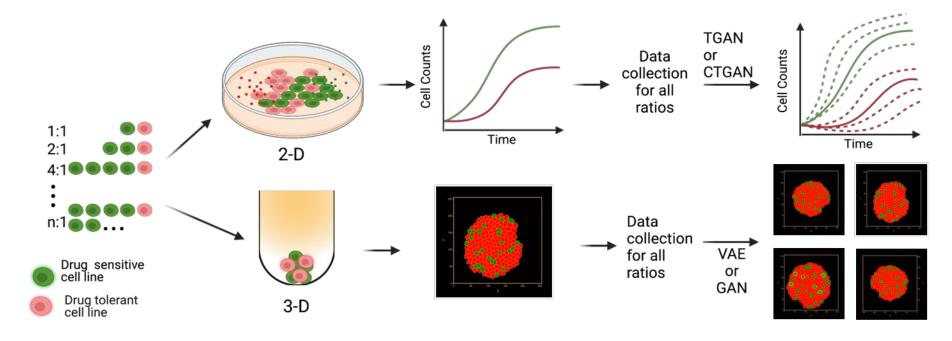
Case #2: Treatment Response

Figure 2

High resolution CT of the chest notes significant mediastinal adenopathy, bilateral pleural effusions and diffuse pulmonary nodules with the largest being in the posterior subsegment of the right upper lobe before (A) and after (B) treatment with chemotherapy/immunotherapy and olaparib.



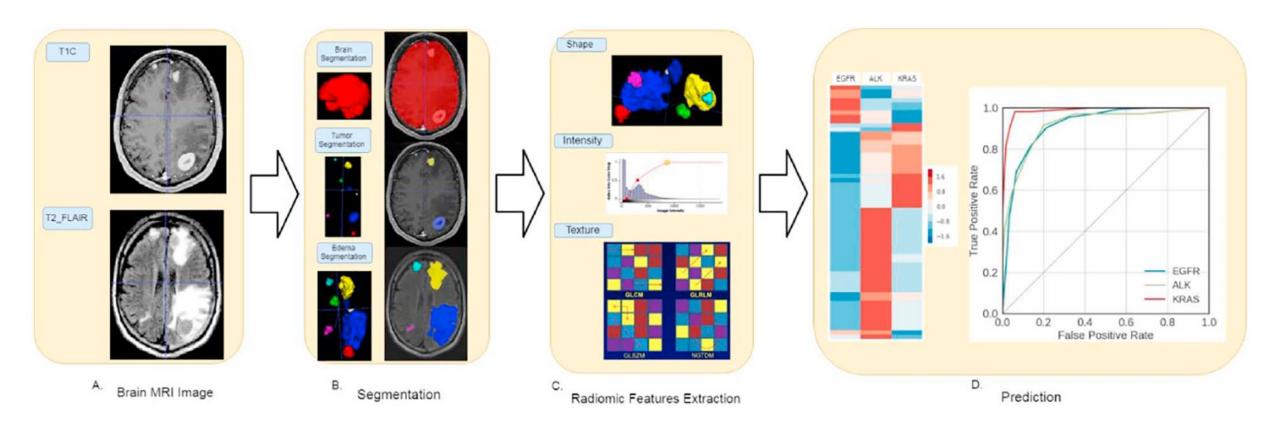
AI in Pre-clinical Research: Synthetic Tabular and Image Data Generated Using Deep Learning Algorithms (TGAN/CTGAN, VAE/GAN) Data



TGAN – Tabular Generative Adversarial Networks
CTGAN – Conditional Tabular Generative Adversarial Networks
VAE - Variational Autoencoders
GAN - Generative Adversarial Networks

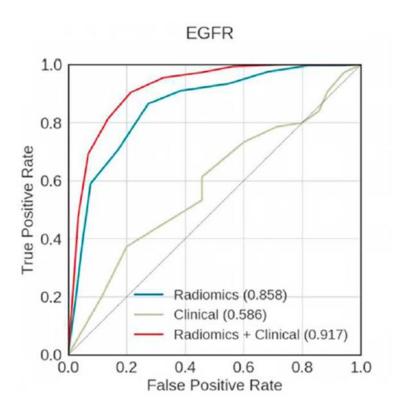
Achuthan et int. Salgia, J. Biosci, 2022

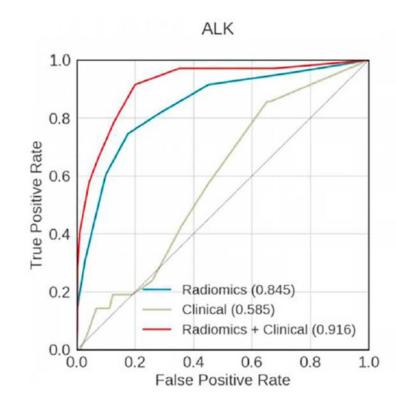
AI in Clinical Research: Predicting Mutational Status from MRI AI Models of Brain Metastases

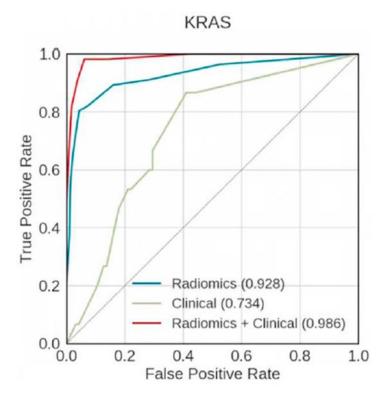


Chen et int. Salgia, Frontiers Oncology, 2021 Chen et int. Salgia, Magnetic Reason Imaging, 2020

AI in Clinical Research: Predicting Mutational Status from MRI AI Models of Brain Metastases

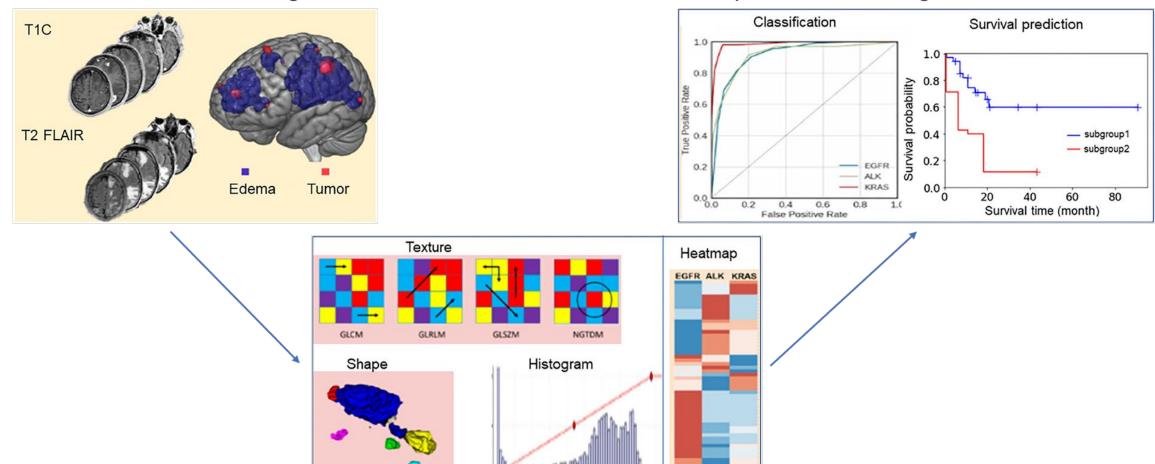






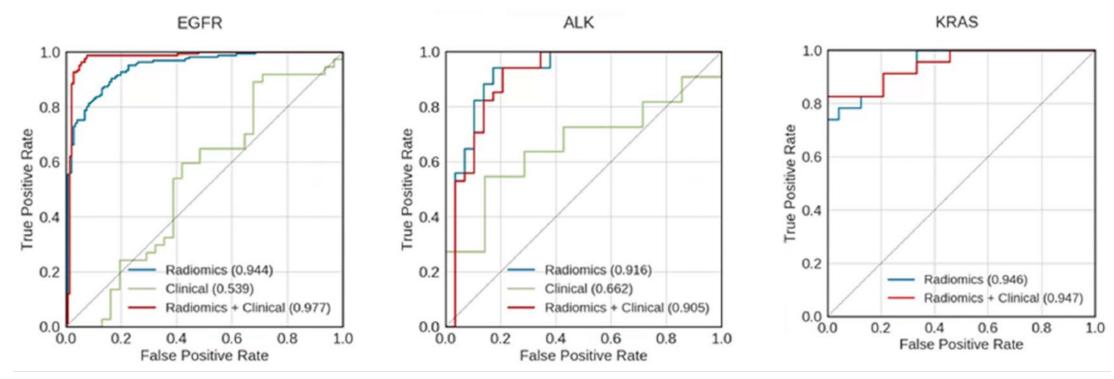
AI in Clinical Research: Predicting Survival from MRI AI Models of Brain Metastases

Schema for brain tumor segmentation, radiomic feature extraction, and predictive modeling



Chen et int. Salgia, Magnetic Resonance Imaging, 2021

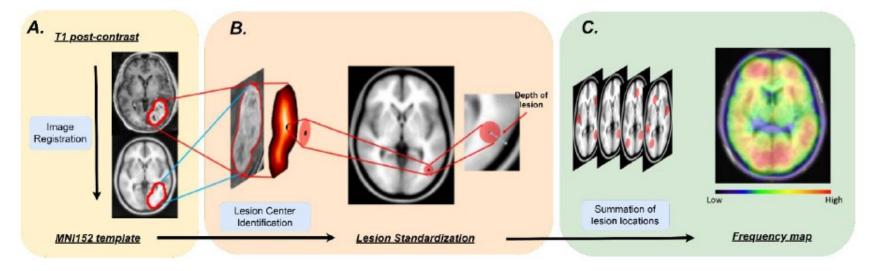
AI in Clinical Research: Predicting Survival from MRI AI Models of Brain Metastases

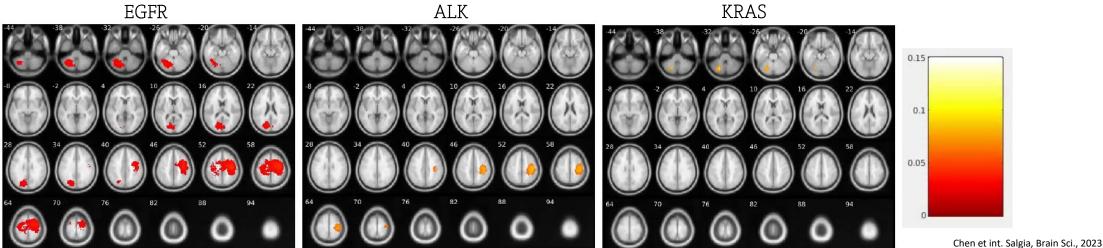


Mutation	Accuracy	AUC*	Sensitivity	Specificity
EGFR	94.90%	0.977	96.00%	94.00%
ALK	84.10%	0.905	88.00%	81.00%
KRAS	83.00%	0.947	83.00%	83.00%

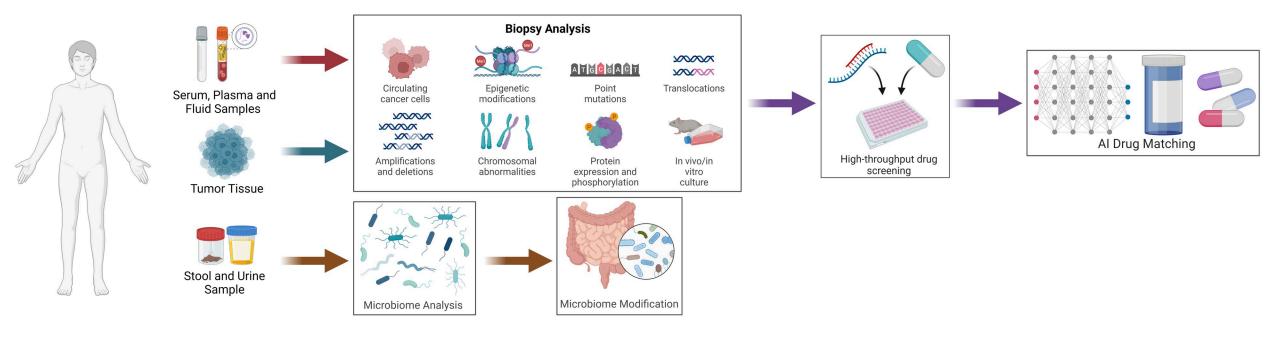
Chen et int. Salgia, Magnetic Resonance Imaging, 2021

AI in Clinical Research: Predicting Distribution of Brain Metastases Based on Mutation Status from MRI AI Models





Future of Precision Medicine



Summary

- Next-generation Sequencing is necessary in patient workup and follow up care
- Precision Medicine relies on timely access to NGS testing and appropriate therapeutic initiation if available
- Genomic and Germline alterations have uncovered therapeutic options for more and more patients
- Artificial intelligence can enhance our understanding of Precision Medicine and clinical decision making
- More therapeutic research and breakthroughs necessary to achieve Personalized Medicine

Acknowledgment

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