

Molecular Tumor Board – How Can We Learn Together?

Patrick C. Ma, M.D., M.Sc.

Associate Professor

Eminent Scholar in Lung Cancer Research

Allen Comprehensive Lung Cancer Program

WVU Cancer Institute, West Virginia University

WV Clinical and Translational Science Institute

Morgantown, West Virginia, USA

October 6, 2017 (Friday)

What is a Tumor Board?

Cancer.Net (ASCO)



A “**Tumor Board**” is a group of doctors and other health care providers with different specialties that meets regularly at the hospital to discuss cancer cases and share knowledge. The board’s goal is to determine the best possible cancer treatment and care plan for an individual patient. Having fresh perspectives from other doctors makes it much easier to come up with that plan.

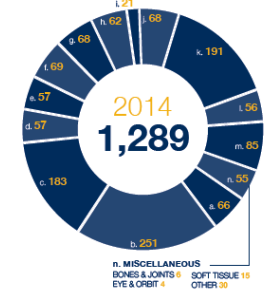
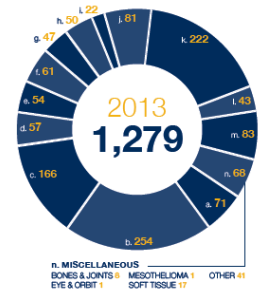
What is a Tumor Board?



WV CANCER REGISTRY DATA

The Mary Babb Randolph Cancer Center, flagship location for the WVU Cancer Institute, provided 41,888 patient visits in 2016, seeing patients for close to 20 different cancer types. West Virginia Cancer Registry data from the previous four years illustrates the broad range of cancer services provided by the WVU Cancer Institute. The Comprehensive Breast Cancer Program, accredited by the National Accreditation Program for Breast Centers, sees more than 200 analytic (new) cases each year.*

- a. BRAIN & OTHER NERVOUS SYSTEM
- f. BREAST
- g. DIGESTIVE SYSTEM
- d. ENDOCRINE SYSTEM
- e. FEMALE GENITAL SYSTEM
- i. LEUKEMIA
- g. LYMPHOMA
- h. MALE GENITAL SYSTEM
- l. MYELOMA
- j. ORAL CAVITY & PHARYNX
- k. RESPIRATORY SYSTEM
- f. SKIN (Melanoma)
- m. URINARY SYSTEM
- n. MISCELLANEOUS

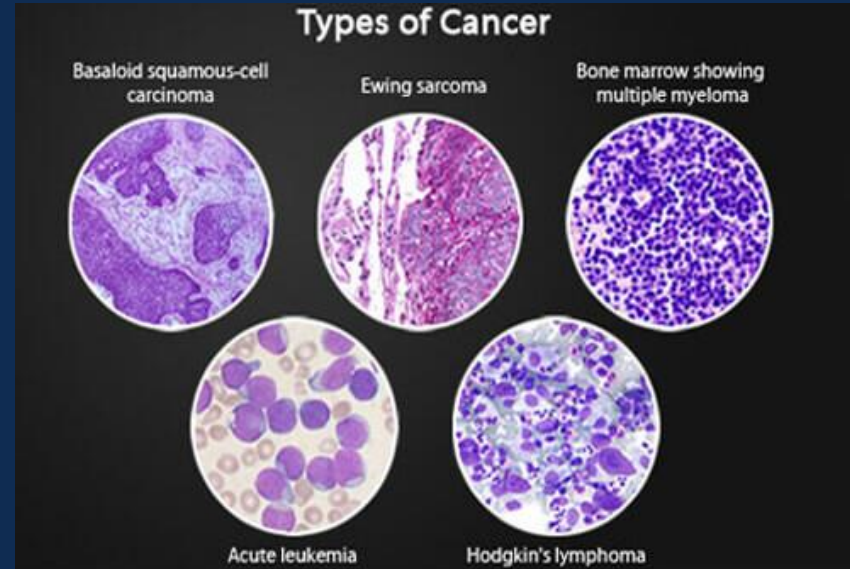
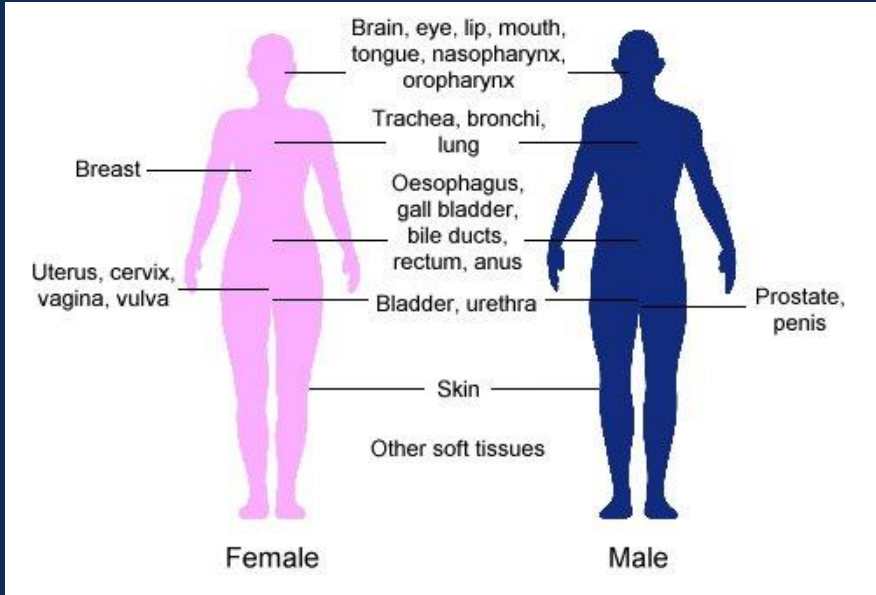


What is a Tumor Board?

MULTIDISCIPLINARY TUMOR BOARDS

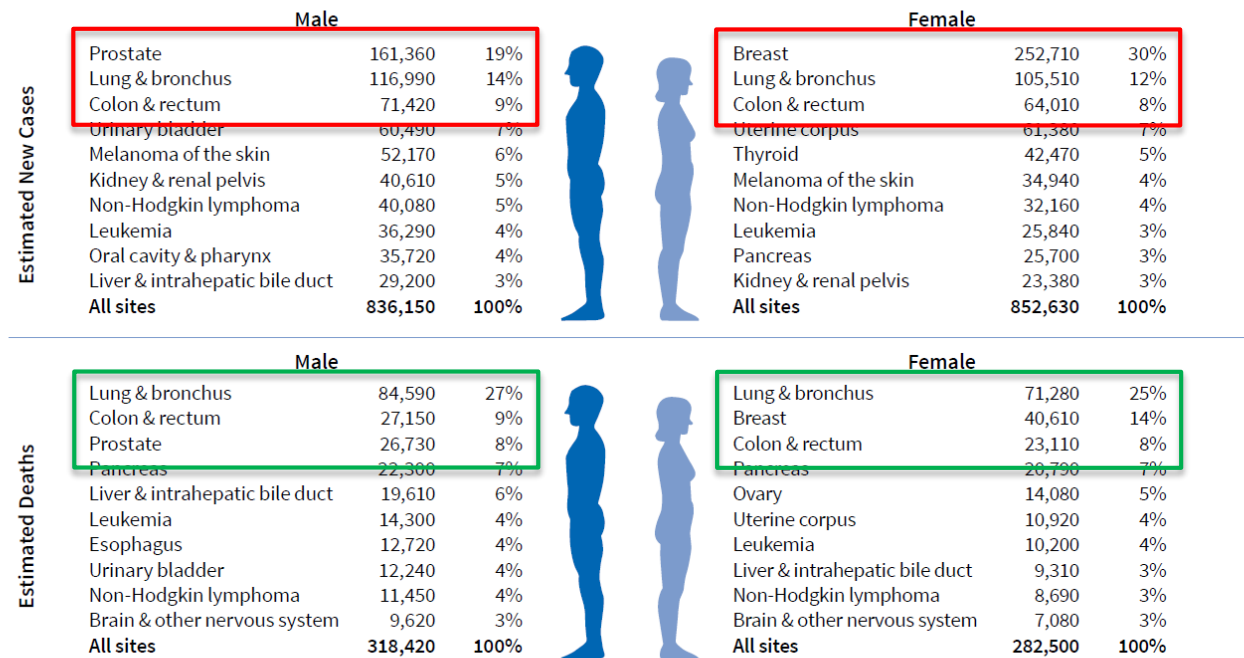
MONDAY	WEDNESDAY	THURSDAY	FRIDAY
BREAST Weekly, 12 pm Dr. Hannah Hazard Dr. M. Salkari	SARCOMA 1st and 3rd Wed, 7:30 am Dr. Brock Lindsey	BMT Weekly, 8 am Dr. Michael Craig	BRAIN Weekly, 7 am Dr. Javier Gonzalez
TUESDAY THORACIC Weekly, 4 pm Dr. Patrick Ma	HEAD AND NECK Weekly, 12 pm Dr. Tariya Fahey Dr. Mohammed Alimubarak	GYNECOLOGICAL ONCOLOGY 1st Thurs, 11 am Dr. Mohammed Adnan	MDS (BLOOD CANCERS) 1st, 3rd – 5th Fri, 12 pm Dr. Abdullah Kanate
	UROLOGY 2nd and 4th Wed, 4 pm Dr. William	GASTROINTESTINAL Weekly, 12 pm Dr. Alan Thomas	CUTANEOUS MALIGNANCIES 2nd Fri, 12 pm Dr. Alan Thomas

A Winning Team
 WVU EXPERTS POOL THEIR KNOWLEDGE FOR GREATER GOOD OF PATIENTS



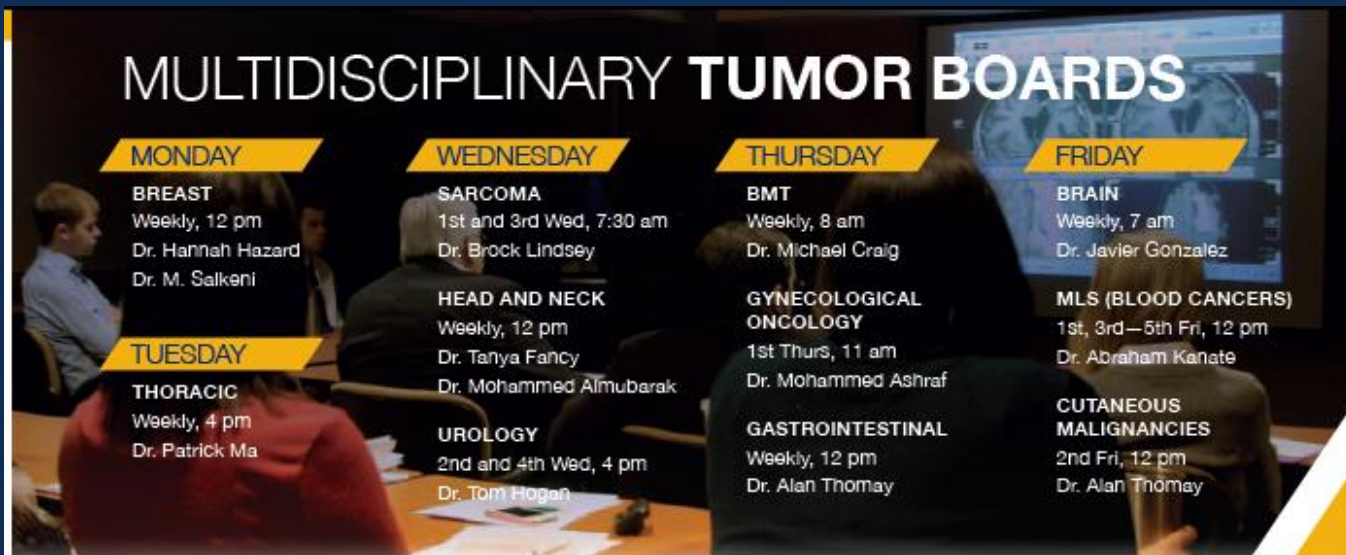
Cancer Sites – New Cases and Deaths

Figure 3. Leading Sites of New Cancer Cases and Deaths – 2017 Estimates



Estimates are rounded to the nearest 10, and cases exclude basal cell and squamous cell skin cancers and in situ carcinoma except urinary bladder.

What is a Tumor Board?



MULTIDISCIPLINARY TUMOR BOARDS

MONDAY	WEDNESDAY	THURSDAY	FRIDAY
BREAST Weekly, 12 pm Dr. Hannah Hazard Dr. M. Salkeni	SARCOMA 1st and 3rd Wed, 7:30 am Dr. Brock Lindsey	BMT Weekly, 8 am Dr. Michael Craig	BRAIN Weekly, 7 am Dr. Javier Gonzalez
TUESDAY	HEAD AND NECK Weekly, 12 pm Dr. Tahya Fancy Dr. Mohammed Almubarak	GYNECOLOGICAL ONCOLOGY 1st Thurs, 11 am Dr. Mohammed Ashraf	MLS (BLOOD CANCERS) 1st, 3rd—5th Fri, 12 pm Dr. Abraham Kanate
THORACIC Weekly, 4 pm Dr. Patrick Ma	UROLOGY 2nd and 4th Wed, 4 pm Dr. Tom Hogan	GASTROINTESTINAL Weekly, 12 pm Dr. Alan Thomay	CUTANEOUS MALIGNANCIES 2nd Fri, 12 pm Dr. Alan Thomay

A Winning Team

WVU EXPERTS POOL THEIR KNOWLEDGE FOR GREATER GOOD OF PATIENTS

The Cancer Genome Atlas

COMMENTARY

OPEN

The Cancer Genome Atlas Pan-Cancer analysis project

The Cancer Genome Atlas Research Network¹, John N Weinstein^{2,3}, Eric A Collisson⁴, Gordon B Mills³, Kenna R Mills Shaw^{5,6}, Brad A Ozenberger⁷, Kyle Ellrott^{8,9}, Ilya Shmulevich¹⁰, Chris Sander¹¹ & Joshua M Stuart^{8,9}

The Cancer Genome Atlas (TCGA) Research Network has profiled and analyzed large numbers of human tumors to discover molecular aberrations at the DNA, RNA, protein and epigenetic levels. The resulting rich data provide a major opportunity to develop an integrated picture of commonalities, differences and emergent themes across tumor lineages. The Pan-Cancer initiative compares the first 12 tumor types profiled by TCGA. Analysis of the molecular aberrations and their functional roles across tumor types will teach us how to extend therapies effective in one cancer type to others with a similar genomic profile.

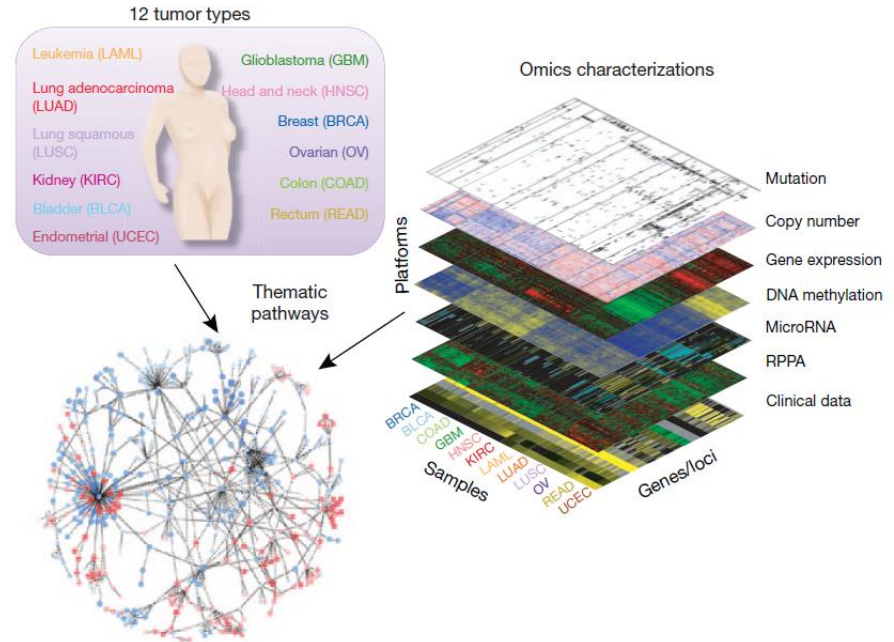


Figure 1 Integrated data set for comparing and contrasting multiple tumor types. The TCGA Pan-Cancer project assembled data from thousands of patients with primary tumors occurring in different sites of the body, covering 12 tumor types (top left) including glioblastoma multiformae (GBM), lymphoblastic acute myeloid leukemia (LAML), head and neck squamous carcinoma (HNSC), lung adenocarcinoma (LUAD), lung squamous carcinoma (LUSC), breast carcinoma (BRCA), kidney renal clear-cell carcinoma (KIRC), ovarian carcinoma (OV), bladder carcinoma (BLCA), colon adenocarcinoma (COAD), uterine cervical and endometrial carcinoma (UCEC) and rectal adenocarcinoma (READ).

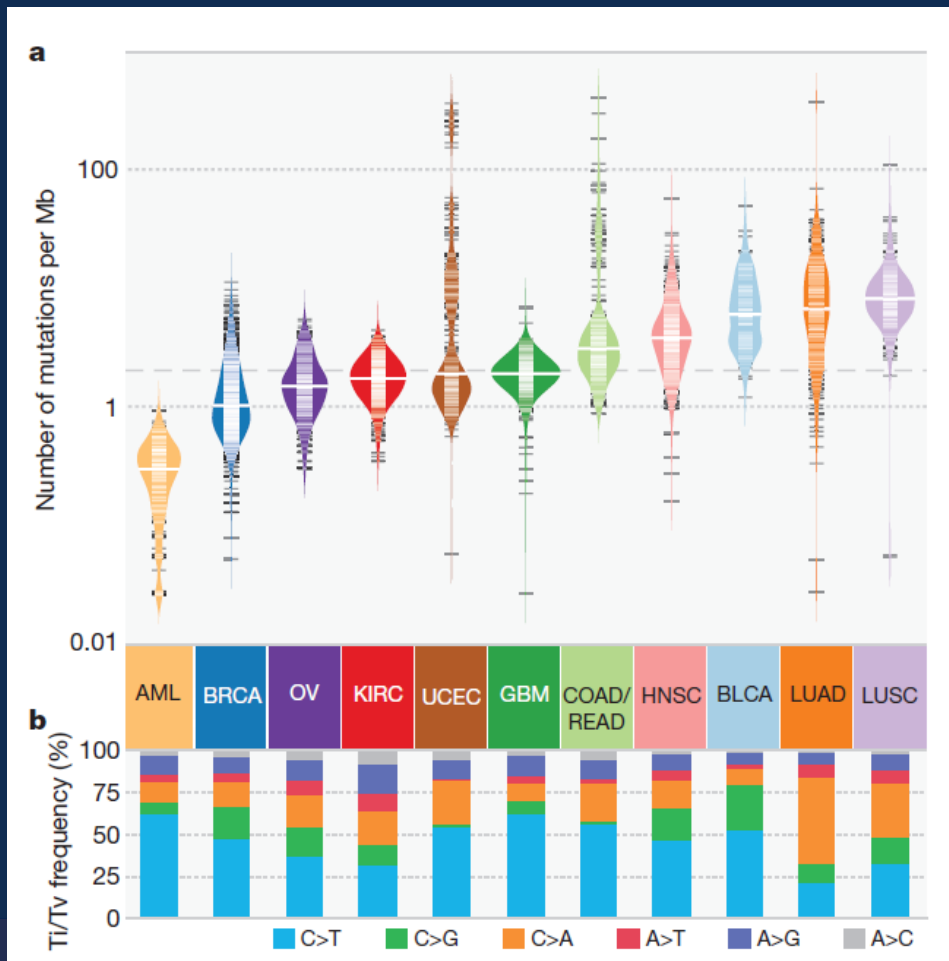
Cancer Genome

ARTICLE

OPEN
doi:10.1038/nature12634

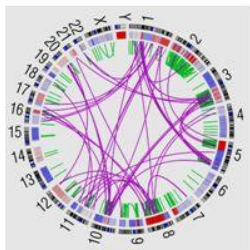
Mutational landscape and significance across 12 major cancer types

Cyriac Kandoth^{1*}, Michael D. McLellan^{1*}, Fabio Vandin², Kai Ye^{3,3}, Beifang Niu¹, Charles Lu¹, Mingchao Xie¹, Qunyuan Zhang^{1,3}, Joshua F. McMichael¹, Matthew A. Wyczalkowski¹, Mark D. M. Leiserson², Christopher A. Miller¹, John S. Welch^{4,5}, Matthew J. Walter^{4,5}, Michael C. Wendl^{1,3,6}, Timothy J. Ley^{1,3,4,5}, Richard K. Wilson^{1,3,5}, Benjamin J. Raphael² & Li Ding^{1,3,4,5}

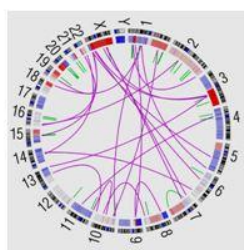


Cancer Genome Complexity and Heterogeneity

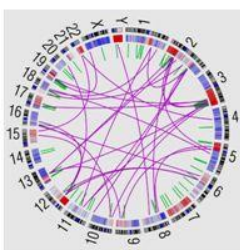
LUSC-66-2756



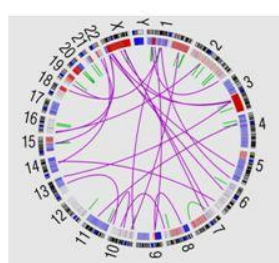
LUSC-34-2600



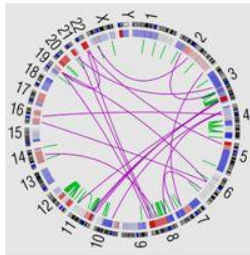
LUSC-43-3394



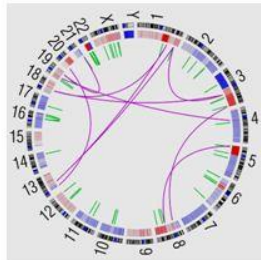
LUSC-34-2609



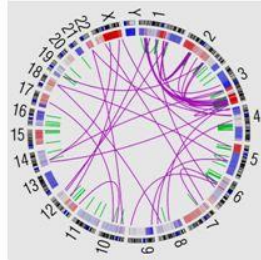
LUSC-56-1622



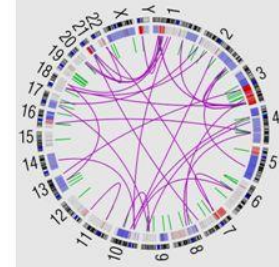
LUSC-60-2695



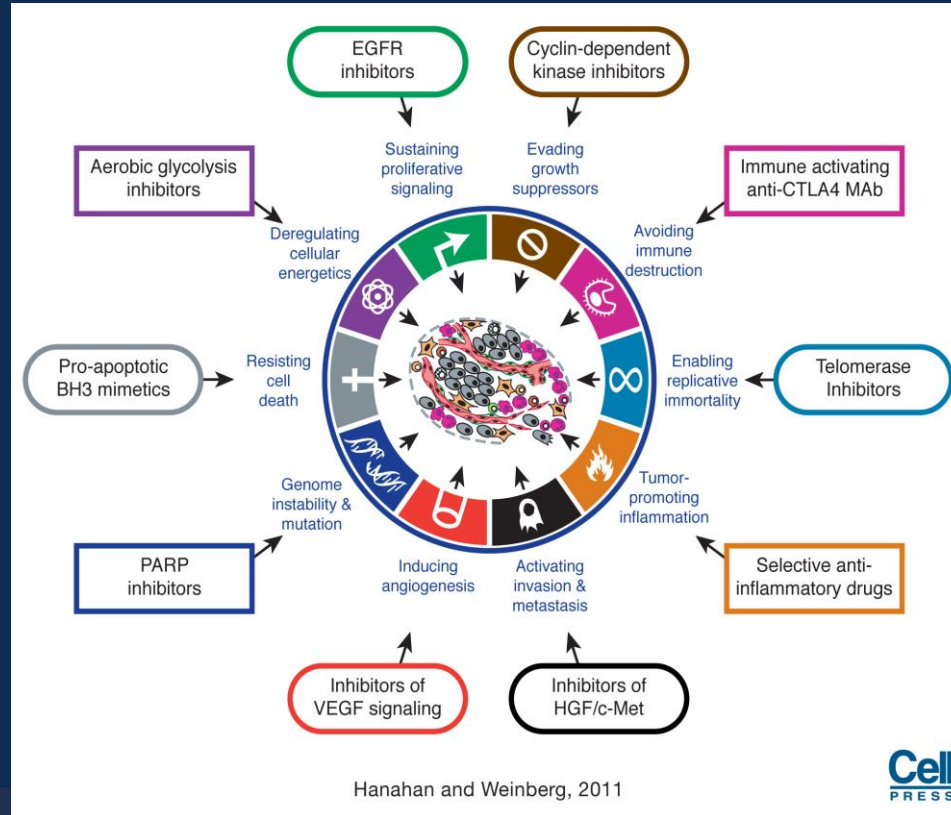
LUSC-60-2711



LUSC-60-2713

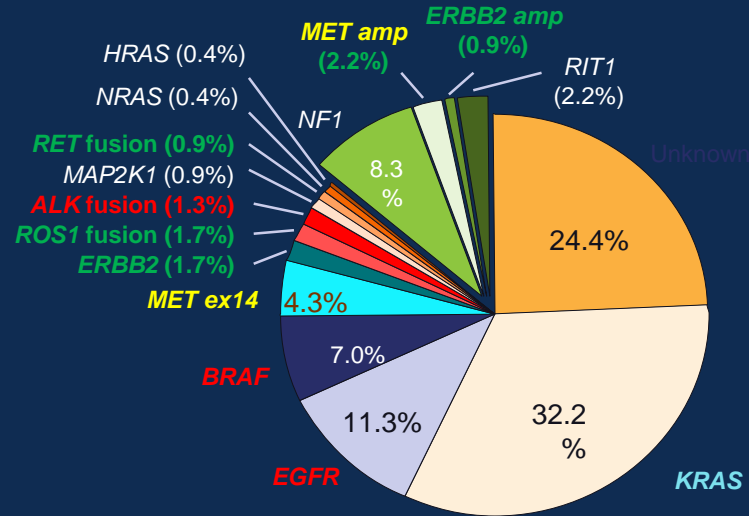


Hallmarks of Cancer - v.2.0



Lung Cancer Precision Therapy 2017

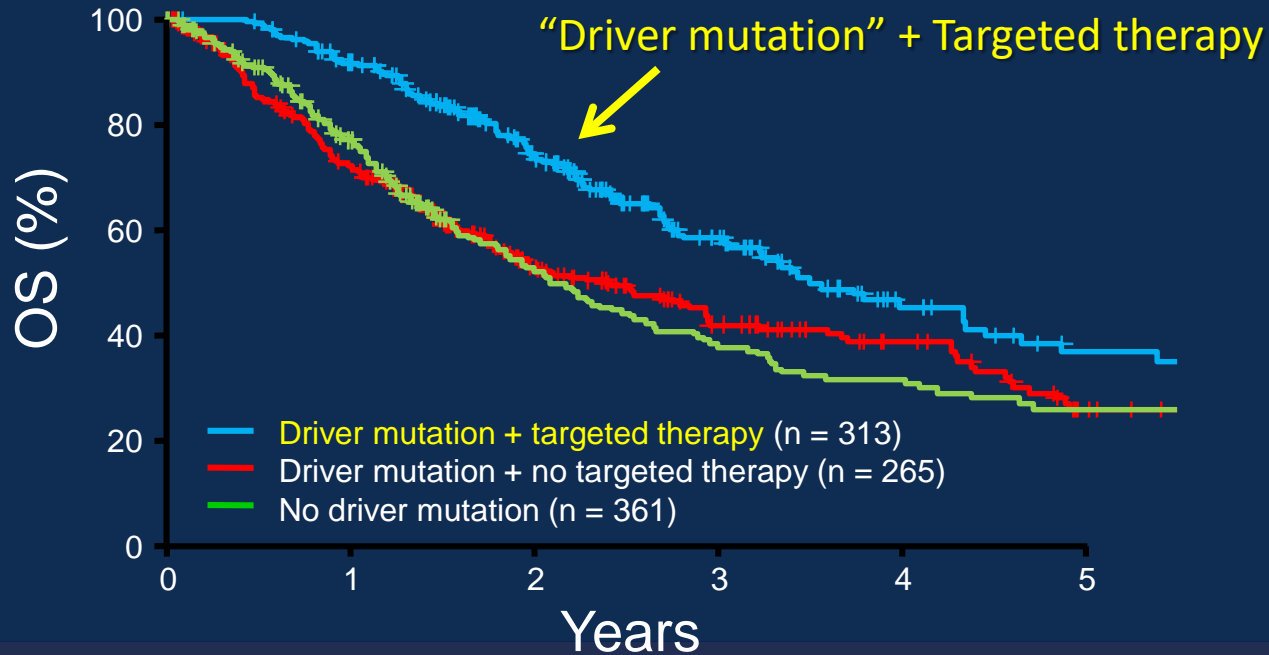
- Lung Cancer - composed of many distinct “driver” genomic mutations, often intrinsically actionable.



Genomic Classification - Lung Adenocarcinoma:
“Actionable Mutations”

Lung Cancer Mutation Consortium

LCMC data supported the OS being the best when pts with “Driver mutations” were treated with matching “Targeted therapy”



Molecular Tumor Board

WVU Cancer Institute – Molecular Tumor Board

- Launched September 2017
- Multidisciplinary Team Approach
- Objectives: Case Management and Educational
- Case review (clinical and molecular features) and literature review
- Attendee: Clinical – Surgical / Radiation / Medical Oncologists,
Radiologist
Pathology – Molecular Pathologist
Basic Scientists (Ph.D.'s)
Fellows, Graduate Students
Nurses, Mid-Level Practitioners, Social Workers,
Pharmacists, Cancer Prevention and Control

CASE 1:

- 46 y.o. Caucasian female, “never-smoker”, otherwise healthy
 - Presents with progressive forgetfulness, then acute confusion
 - ED: brain MRI: At least 6 metastatic foci, with the largest identified within the inferior left frontal lobe (~1.5 cm) associated with extensive vasogenic edema and mass effect with subfalcine herniation.

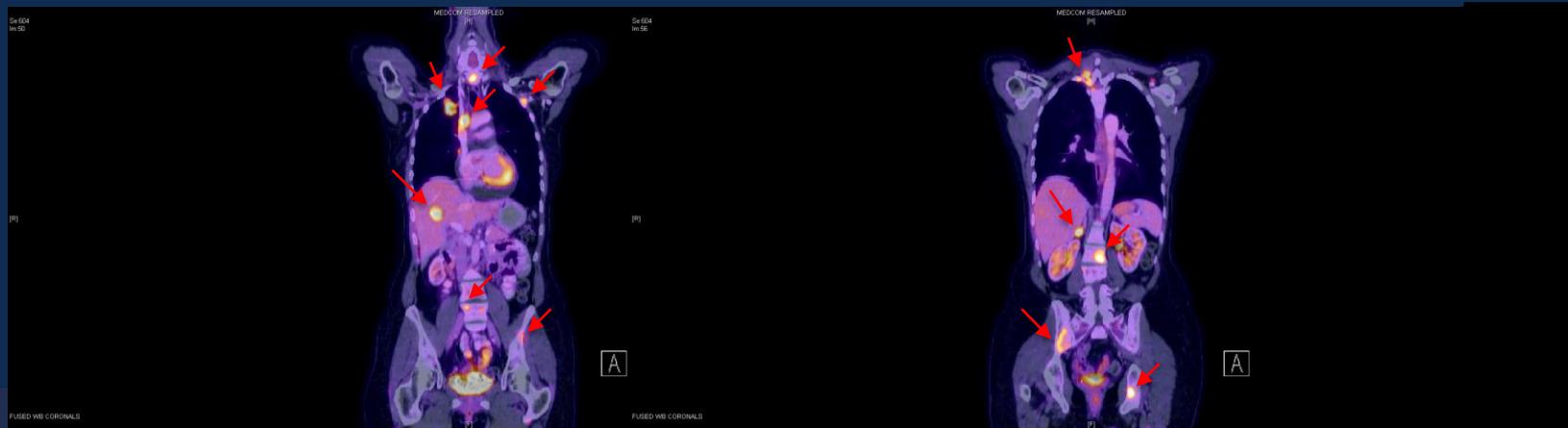


Craniotomy –
metastatic adenocarcinoma

CASE 1: 46 y.o. never-smoker female

- **PET/CT scan:**

- Hypermetabolic **lung mass** (RUL) (3.1 x 2.8 x 2 cm) with metastatic disease involving the **brain**, **skeletal system**, mediastinal **lymph nodes**, right **adrenal gland**, and **liver**.
- Small pericardial effusion.
- Postoperative changes from resection of left frontal lobe metastatic brain lesion. Multiple known brain metastasis are partially demonstrated
- Hypermetabolic enhancing lesion in the **left infraspinatus muscle** (2.1 x 1.5 cm) c/w necrotic metastatic lesion.

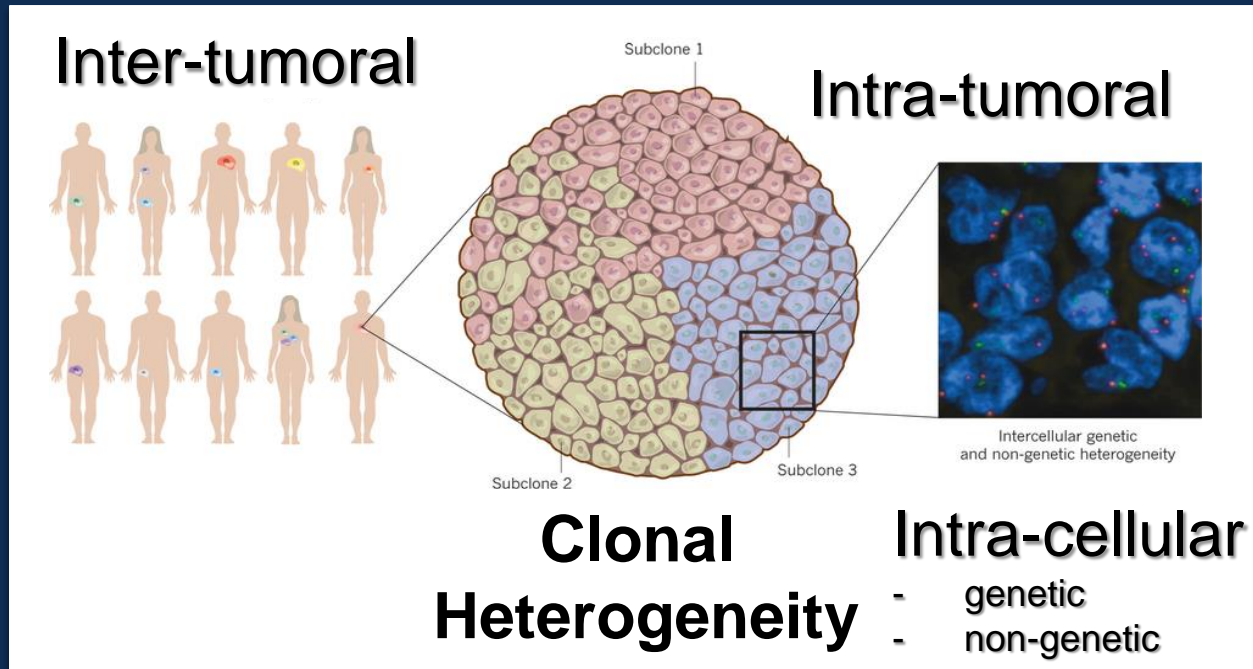


CASE 1: 46 y.o. never-smoker female

- Comprehensive Tumor Profiling (Caris CMI): **TP53 Exon 5 | C176F**
- **EGFR - Exon 19 | E746_A750del** → Started on first-line Erlotinib (Tarceva) 150 mg once daily by mouth (9/2015)
- Essential complete response in 4 months



Tumor Heterogeneity



CASE 1: 46 y.o. never-smoker female

- Guardant360 Liquid Biopsy:
- 9/2016 (1-yr post-erlotinib: several bone mets progression):
EGFR Exon 19 | E746_A750del



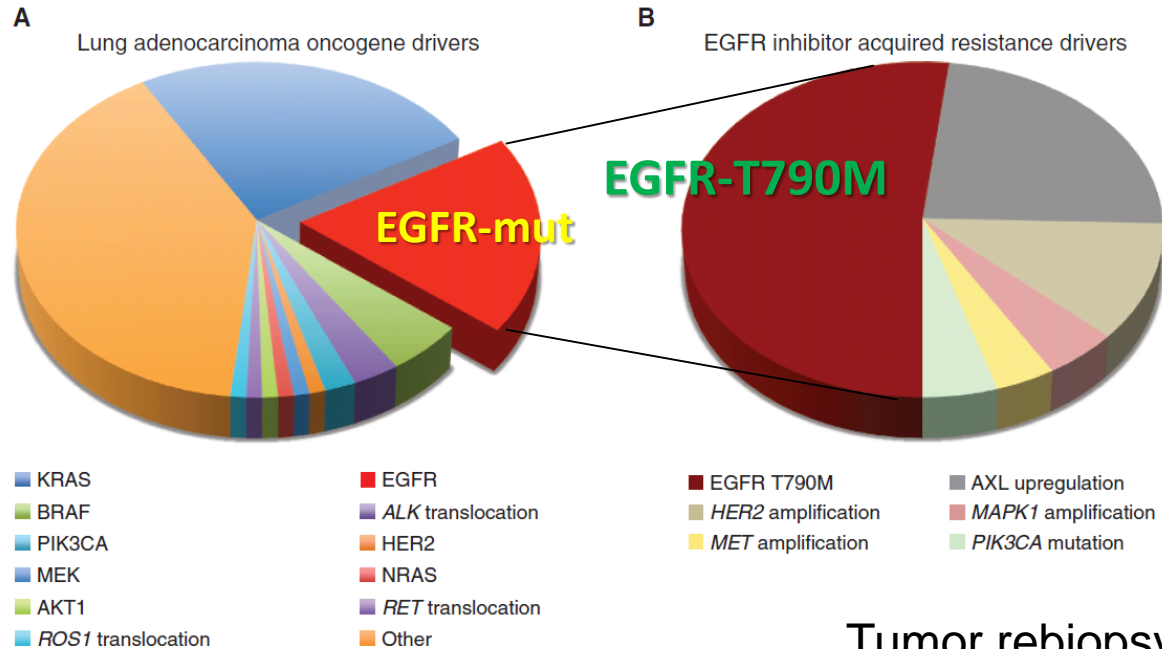
Sept 2016



Dec 2016

12/2016 (left ilium bone mets progression):
TP53 Exon 5 | C176F;
EGFR Exon 19 | E746_A750del
EGFR Exon 20 | T790M

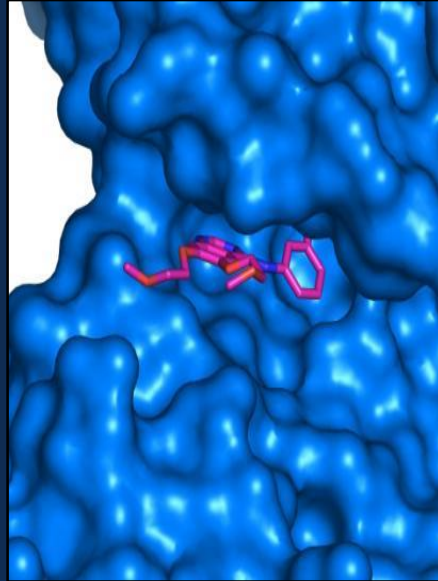
Targeting the New Achilles Heel in Drug Resistance



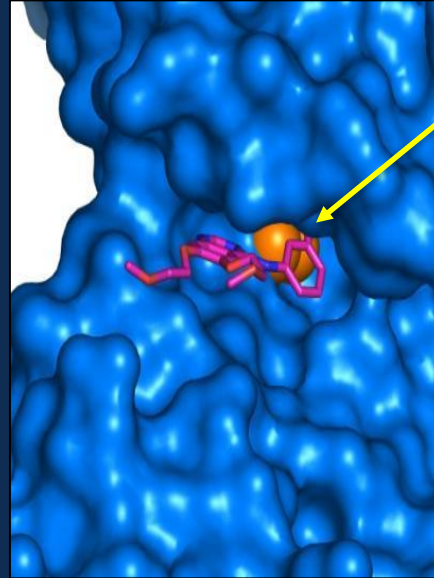
Tumor rebiopsy studies

Targeting the New Targets:

Overcoming Acquired Tumor Resistance to (EGFR) Targeted Therapy



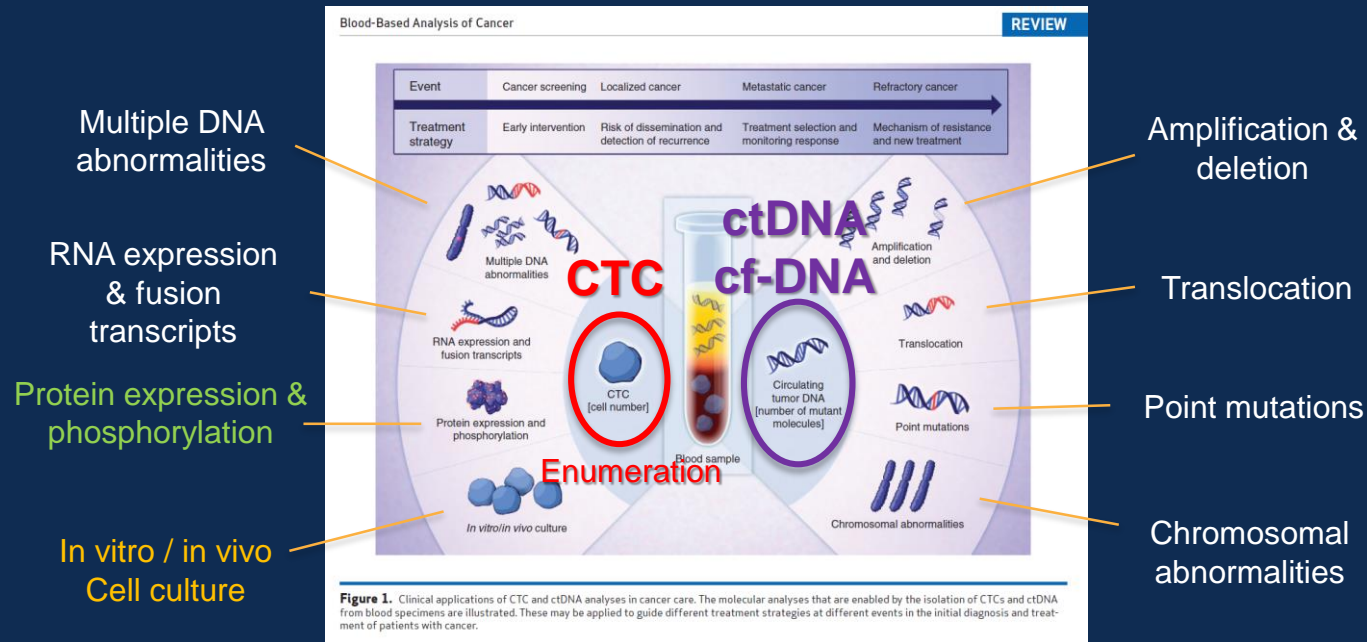
Wild-type receptor: erlotinib (chemical structure) snugly fits into the ATP-binding pocket of EGFR blocking its function



T790M

T790M mutant receptor: methionine (M) 790 (orange) protrudes into the ATP-binding pocket, leads to steric hindrance disallowing erlotinib to bind

“Liquid Biopsy” in Cancer Diagnosis and Therapy (2017)



Cobas EGFR Mutation v2 Plasma Test – FDA Approved June 1st, 2016
Liquid Biopsy – Molecular Gneomic Profiling

Targeting the New Targets:

Overcoming Acquired Tumor Resistance to (EGFR) Targeted Therapy

Guardant360 Tumor Response Map

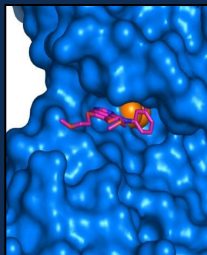
The Guardant360 Tumor Response Map illustrates the mutant allele percentage (% cfDNA) of observed somatic variants at each sample submission time point. The "Somatic Alteration Burden" value below refers to the maximum % cfDNA detected at each time point. Amplifications are not plotted, and only the first and last four test dates are plotted. Please see the Physician Portal (<https://portal.guardianhealth.com>) for the Tumor Response Map with all test dates.



Summary of Somatic Alterations & Associated Treatment Options

The percentage of altered cell-free DNA (% cfDNA) circulating in blood is related to the unique tumor biology of each patient. Factors that may affect the % cfDNA of detected somatic alterations include tumor growth, turn-over, size, heterogeneity, vascularization, disease progression, and treatment.

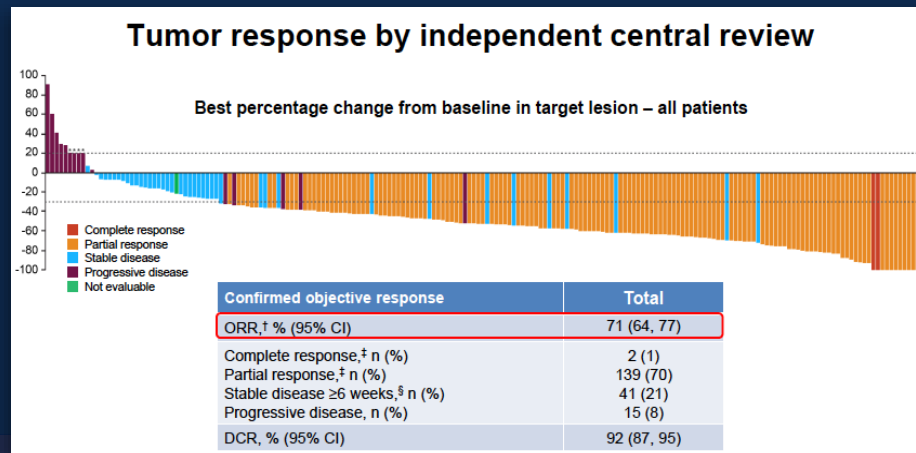
Alteration	Mutation Trend	% cfDNA or Amplification	FDA Approved in Indication <i>see page 3</i>	Available for Use in Other Indications <i>see page 3</i>	Clinical Drug Trials <i>see page 6</i>
EGFR	Exon 19 Deletion	6.7	Afatinib	None	Trials Available
	T790M	5.4	Osimertinib Lack of Response: Erlotinib, Gefitinib	Afatinib, Cetuximab	Trials Available
TP53	C176F	0.5	None	None	Trials Available



Third-Generation T790M-Targeting TKIs

- These novel and highly promising drugs largely spare EGFR WT signaling and preferentially block mutant/T790M signaling, leading to potentially wider therapeutic indices
- AZD9291 (osimertinib) – 56% response rate in T790M+ patients, well-tolerated, some rash, pneumonitis (3%), now FDA-approved

AURA-2: Osimertinib



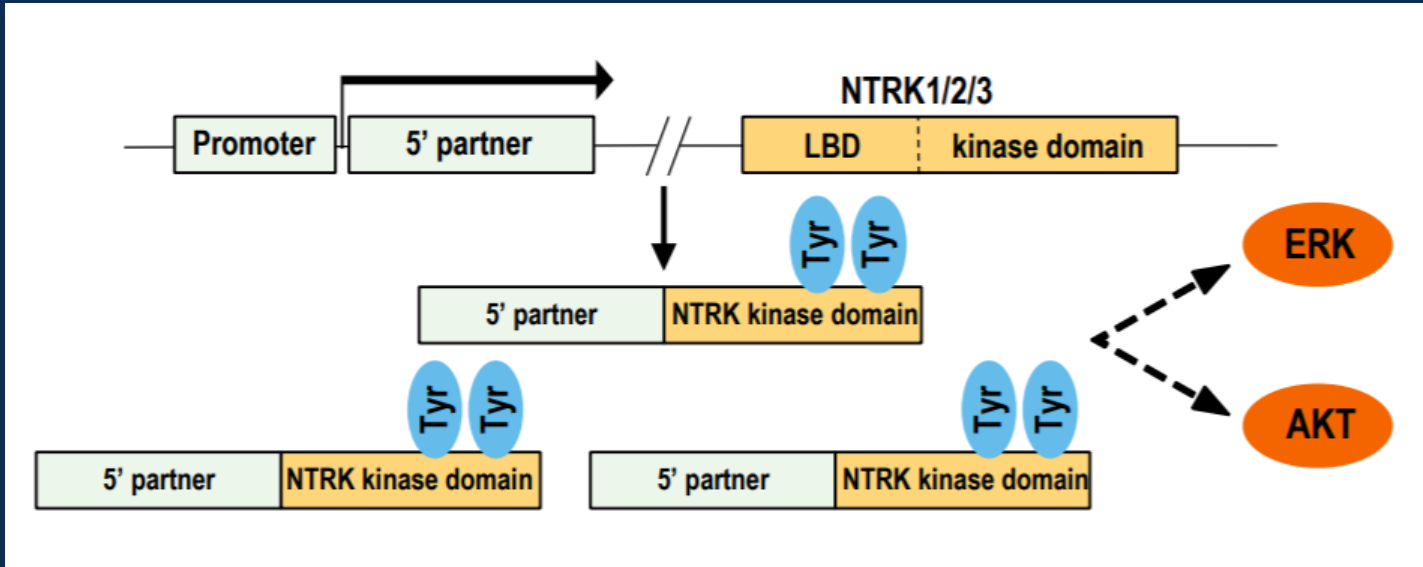
Case 2

- 76 year old Caucasian male, never-smoker, PMHx: Diabetes, Hypothyroidism, Eczema, CAD, HTN, HLD seen as a referral from Cleveland Clinic Genomic Medicine Institute for Radioactive-Iodine Refractory **Papillary Thyroid Cancer**.
- Dec 2008 - Total thyroidectomy with pathology showing papillary thyroid carcinoma, 1/5 regional LN +. Radio-iodine study followed by RAI-ablation completed
- October 2015 - Recurrence of disease noted being RAI resistance, Lenvatinib was started.
- December 2015 - Significant proteinuria thought to be related to lenvatinib, initially dose reduced but eventually stopped Feb 2016.

Case 2

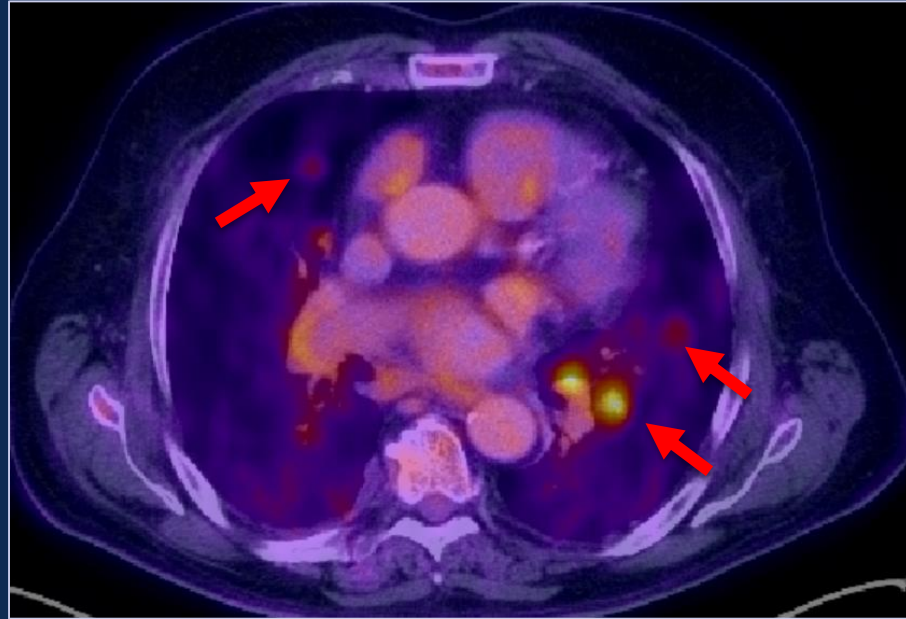
- March 2016. Started on Sorafenib 200 mg daily, complicated by uncontrolled HTN and hence stopped after few months
- May 2016. All cancer directed therapies stopped. CT scan revealed B/L pulmonary nodules increased in size from 2015. Increased right hilar LN, small effusions
- **September 2016**. Molecular genomic tumor profiling reported presence of *ETV6-NTRK3* fusion and TERT promoter
- Enrolled in NTRK inhibitor clinical trial study

NTRK pathway



Case 2: Metastatic lung lesions

PET/CT



Case 2

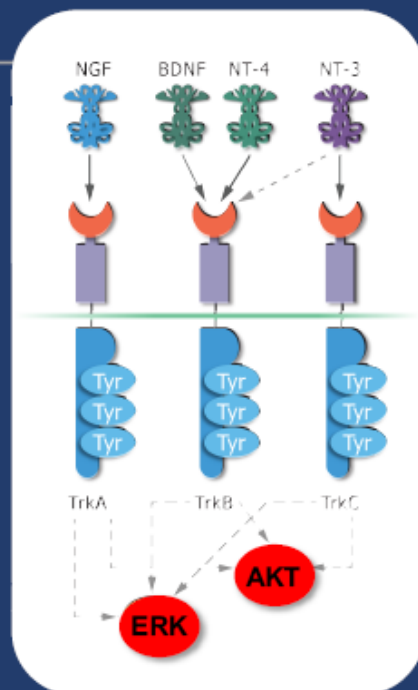
- April 4, 2017 - Completed 5 cycles of NTRK inhibitor. Restaging CT Chest showed Partial Response going in to stable disease
- June 2017 - Had an embolic stroke from atrial fibrillation, with resultant right-sided hemiparesis/hemiplegia, aphasia; undergoing rehab. Drug held for 2 weeks
- July 2017 - Restarted drug on trial protocol.
- Completed 12 cycles by September 2017.
- August 2017 - Restaging CT scan → ongoing partial response

TRK Fusions are Oncogenic and Signal Through Canonical Downstream Pathways

Normal TRK Proteins

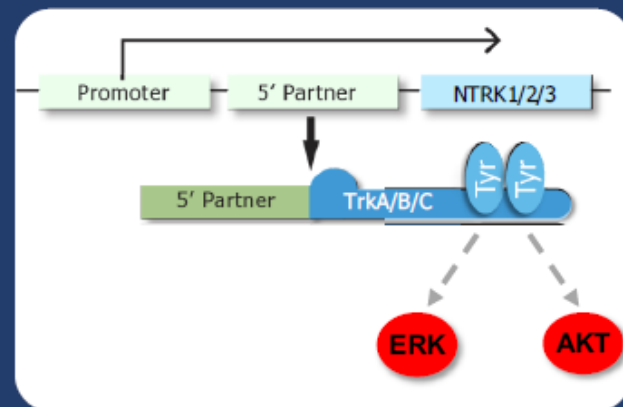
Family of neurotrophin receptors

- TrkA (*NTRK1*) → Pain, thermoregulation
- TrkB (*NTRK2*) → Movement, memory, mood, appetite, body weight
- TrkC (*NTRK3*) → Proprioception



TRK Fusions

- Ligand binding domain replaced by 5' fusion partner; highly expressed by promoter of 5' fusion gene
- Ligand-independent activation



TRK Fusions Found in Diverse Cancer Histologies

TRK FUSION FREQUENCY

<5%

CNS

- ✓ Astrocytoma
- ✓ Brain low-grade glioma
- ✓ Glioblastoma

GI

- ✓ Colorectal cancer
- ✓ Cholangiocarcinoma
- ✓ GIST
- ✓ Pancreatic cancer

Head and neck

- ✓ Squamous cell carcinoma

Lung

- ✓ Adenocarcinoma
- ✓ Large cell neuroendocrine

Other

- ✓ Acute myeloid leukemia
- ✓ Breast invasive carcinoma
- ✓ Melanoma
- ✓ Sarcoma

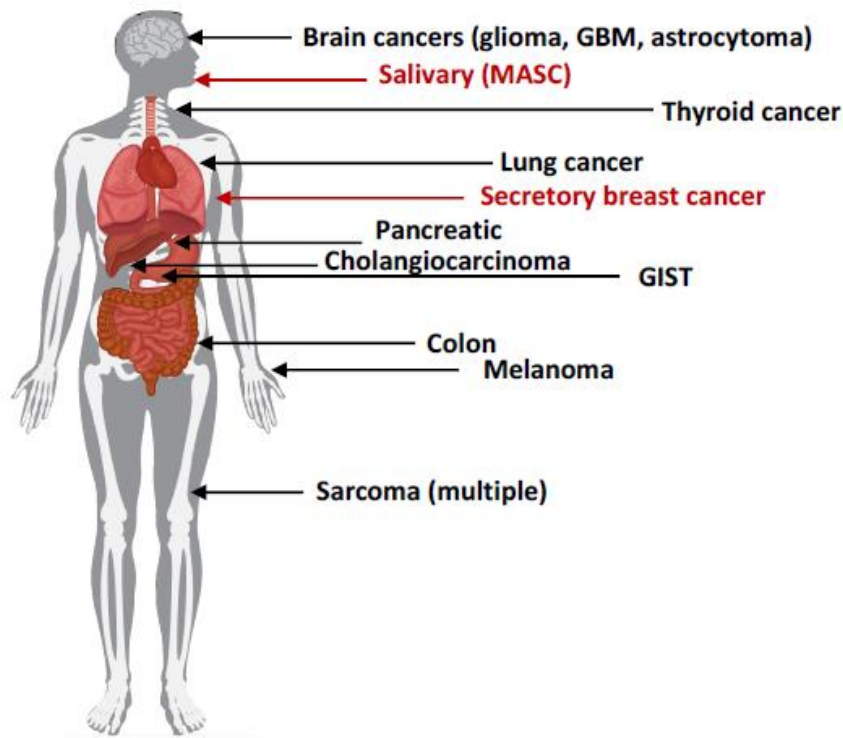
5–25%

- ✓ Congenital mesoblastic nephroma
- ✓ Papillary thyroid cancer
- ✓ Pontine glioma
- ✓ Spitz tumors

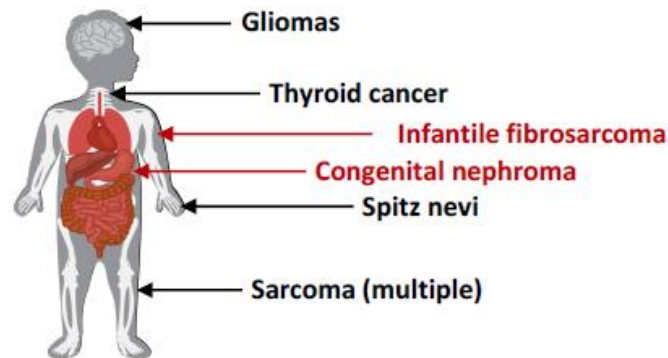
>75%

- ✓ Mammary analogue secretory carcinoma (MASC) of the salivary glands
- ✓ Secretory breast carcinoma
- ✓ Infantile fibrosarcoma

TRK fusions found in diverse cancer histologies

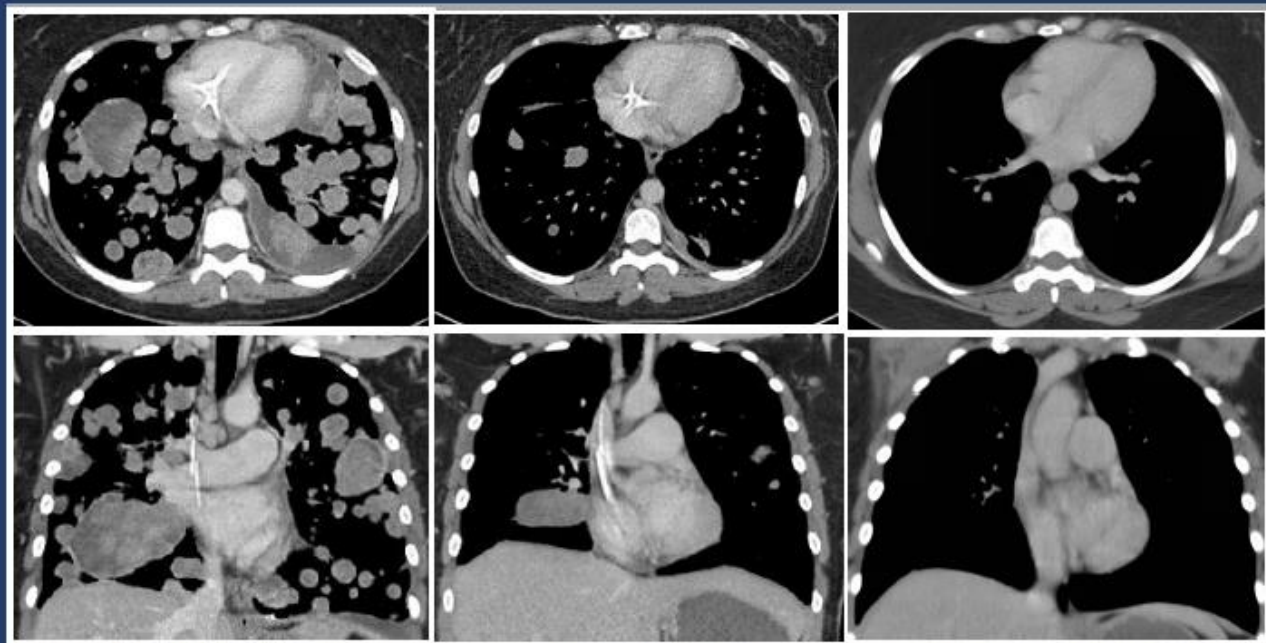


- Common cancer with low TRK fusion frequency
- Rare cancer with high TRK fusion frequency



Patient #1: LMNA-NTRK1 fusion soft tissue sarcoma

- 42 yo female with undifferentiated sarcoma progressed through epirubicin, ifosfamide, sorafenib, and doxorubicin
- 100mg BID
- Rapid resolution of dyspnea and hypoxemia
- Confirmed partial response
- Currently on study in cycle 14



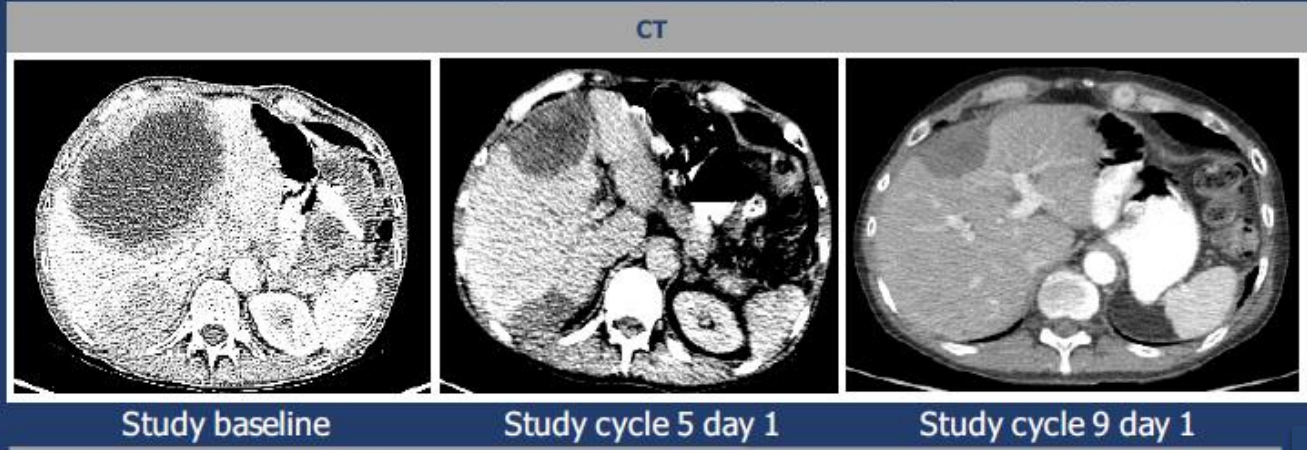
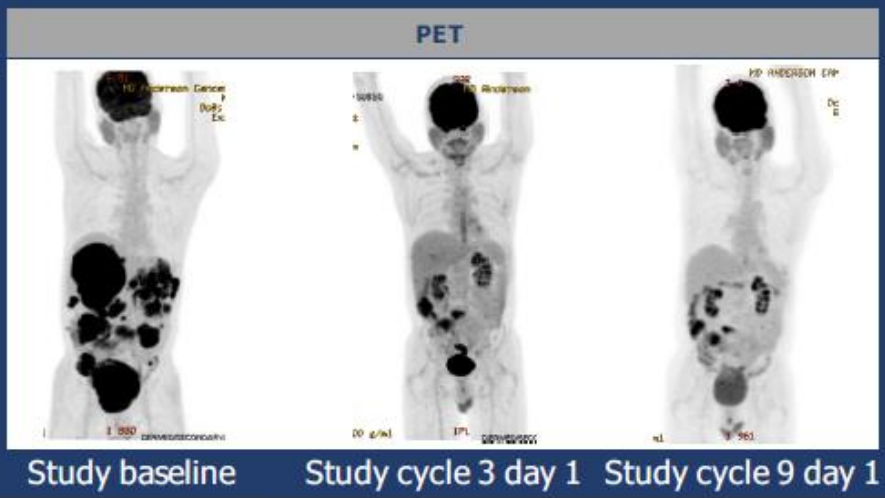
Study baseline

Study cycle 3 day 1

Study cycle 13 day 1

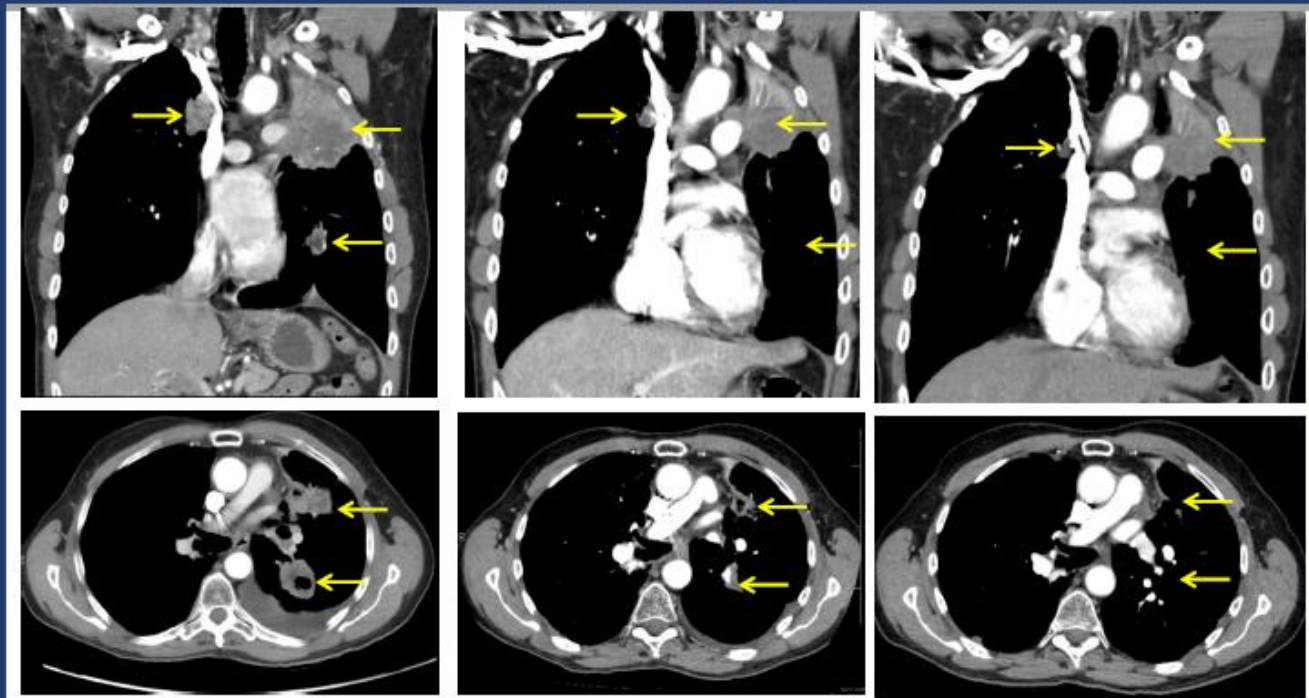
Patient #2: ETV6-NTRK3 fusion GIST

- 55 yo male with GIST progressed through imatinib, sunitinib, sorafenib, nilotinib, and regorafenib
- 150mg BID
- Confirmed partial response
- Currently on study in cycle 10



Patient #4: ETV6-NTRK3 fusion mammary analogue secretory carcinoma of the salivary gland (MASC)

- 66 yo male progressed through radiotherapy, dasatinib, GDC-0941+ erlotinib, and ABBV-399
- 100mg QD*
- Confirmed partial response
- Currently on study in cycle 7



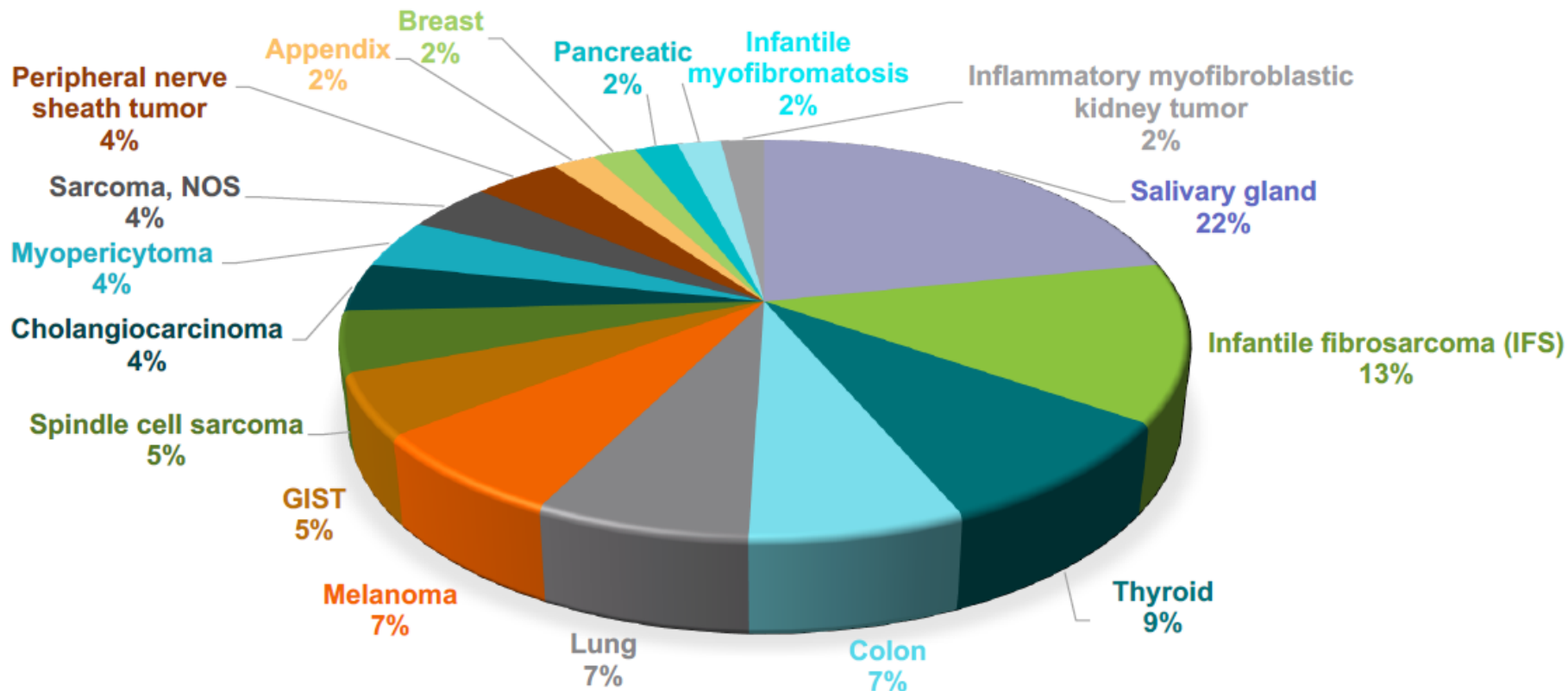
Study baseline

Study cycle 3 day 1

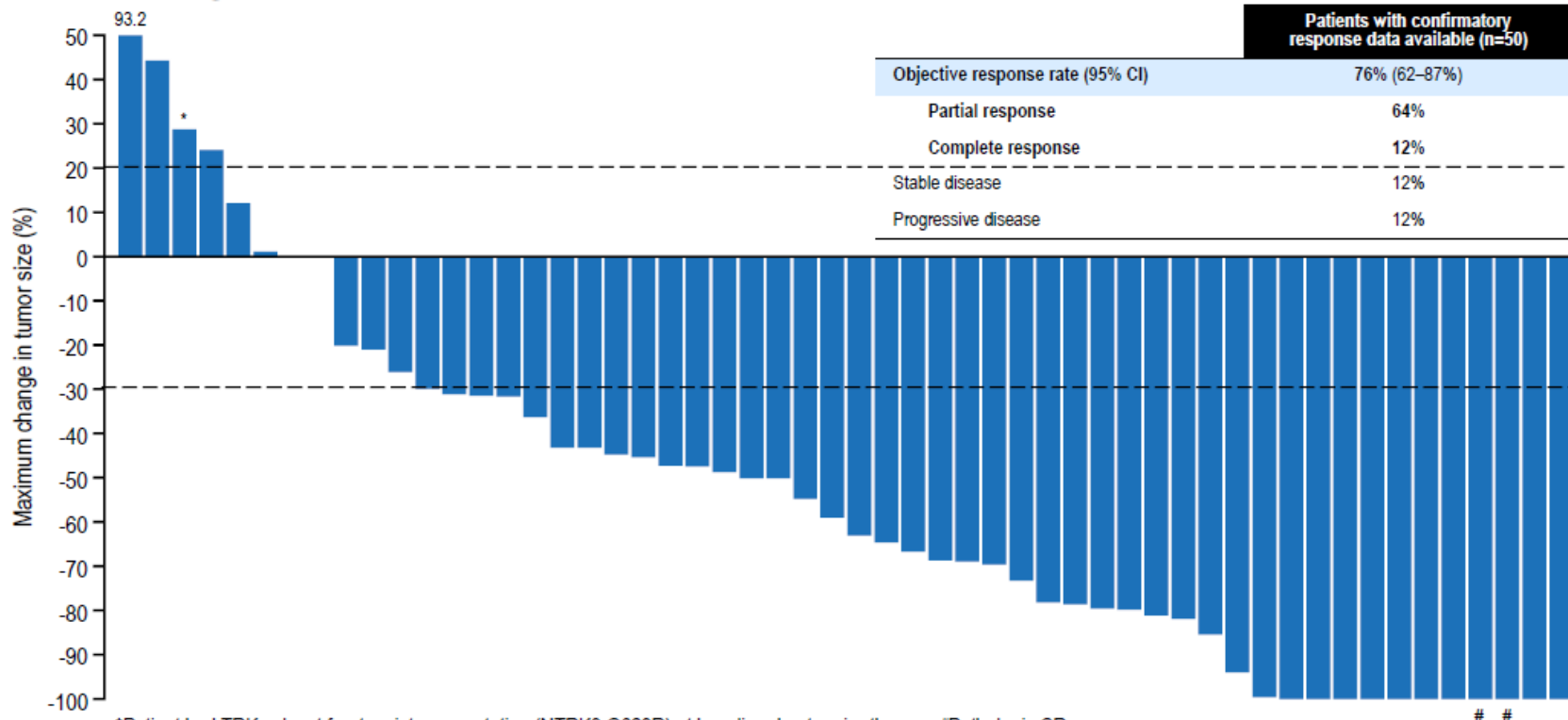
Study cycle 7 day 1

* Patient enrolled at 100mg BID and dose reduced to 100mg QD on C1D2 due to transient dizziness possibly related to drug

Diversity of cancer types treated (n=17)



Efficacy of larotrectinib in TRK fusion cancers



*Patient had TRK solvent front resistance mutation (NTRK3 G623R) at baseline due to prior therapy; #Pathologic CR

Note: One patient not shown here. Patient experienced clinical progression and no post-baseline tumor measurements were recorded.

ETV6-NTRK3 secretory breast cancer patient



Baseline



Day 6



Day 20

**14F, prior therapy: 4 lines of chemotherapy and repeated resections
Treated with larotrectinib under expanded access**

Case 3

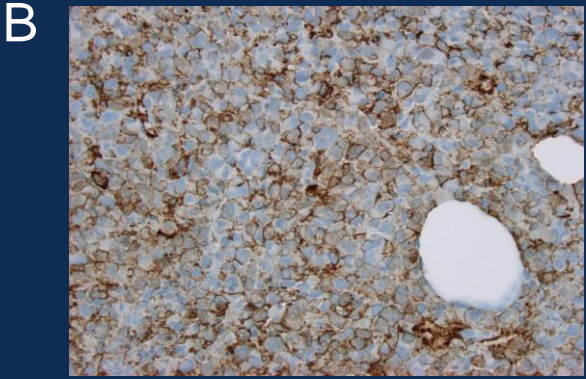
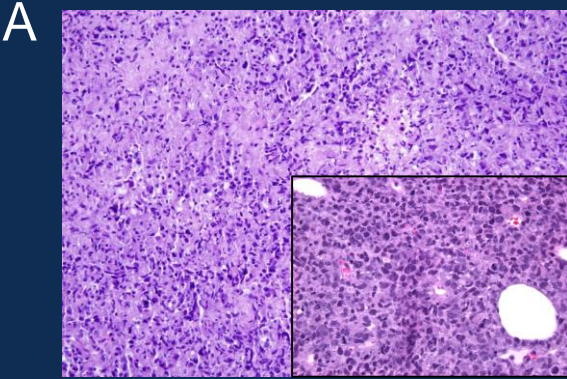
- 50-year-old Caucasian male with a 30 pack-year smoking history, who was healthy except hypertension, presented with a sore lump over the right lower chest wall.
- CT of the chest, abdomen, and pelvis in January 2015 confirmed an enlarging underlying soft tissue mass arising within the chest wall soft tissue/pleural tissue space, centered over the intercostal space, measuring 3.0 x 5.9 x 5.0 cm. An enlarged right mid-hilar lymph node (2.7 x 3.1 x 2.7 cm) was also seen.
- Ultrasound-guided needle biopsy on the chest wall mass → pathology review at the Mayo Clinic, revealed an extensively necrotic, poorly differentiated malignant neoplasm possibly representing a **malignant SFTP (Solitary Fibrous Tumor of the Pleura)**

Case 3

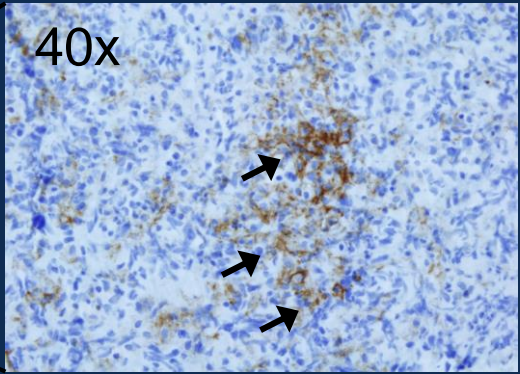
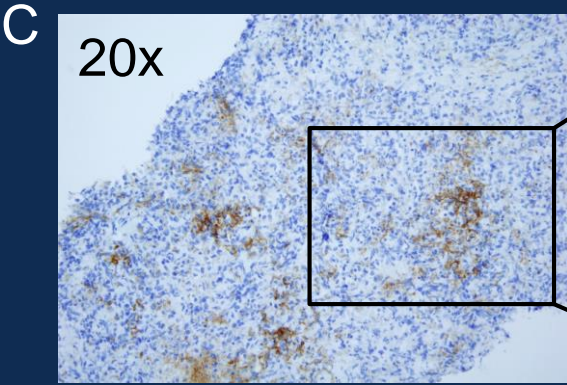
- Diffusely positive for CD34 and focally cytokeratin-positive. INI-1 expression was retained. WT-1, calretinin, ER, CD31, ERG protein, FLI-1 and high MW cytokeratin were negative.
- PET/CT imaging in February 2015 showed no evidence of metastatic lesions in the abdomen, pelvis, or bones, and a bone scan was also negative for any distant metastases.
- The right lower chest wall mass was found intensely hypermetabolic with SUV 23, with the right hilar mass having SUV 20.

Case 3 - Pathology

H&E



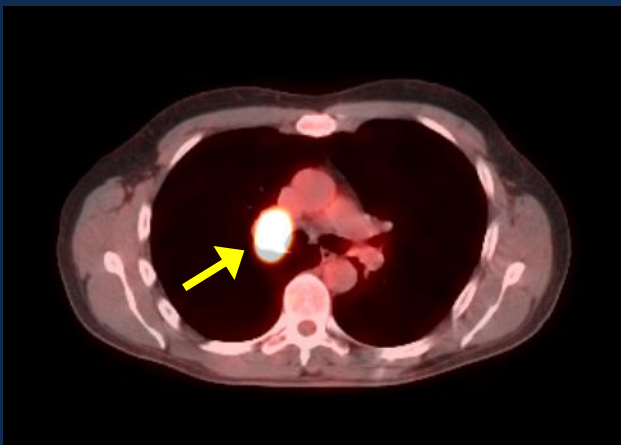
CD34



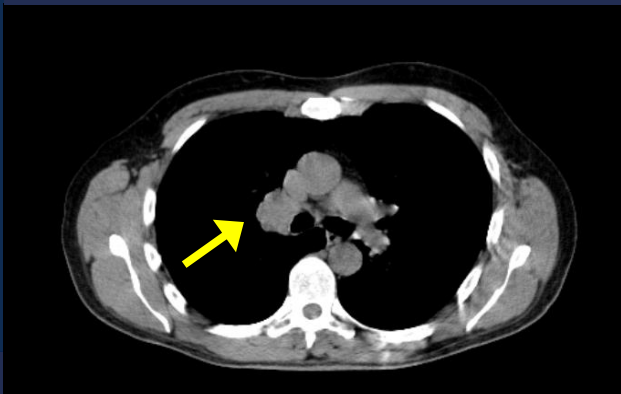
PD-L1

Case 3 - Radiology

A



B



C



D



Case 3

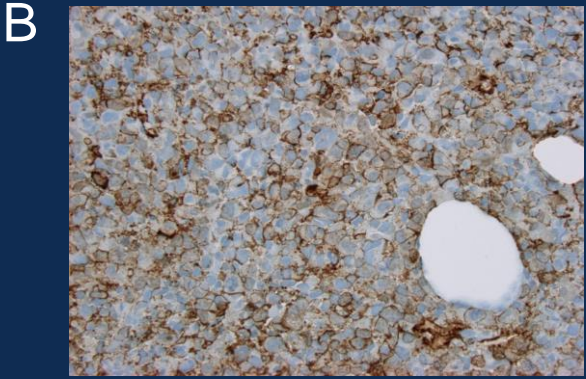
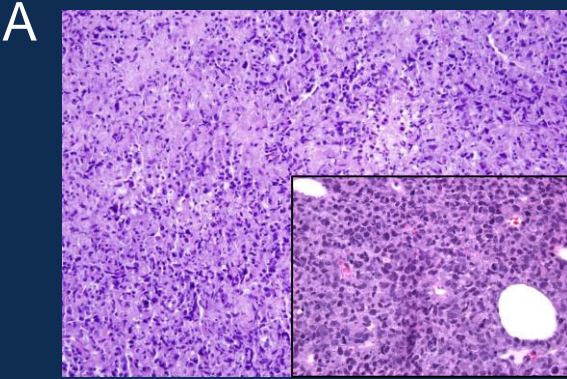
- 1st line concurrent chemoRT: Restaging by PET/CT – shrinkage of the primary chest wall mass to 2.8 x 1.4 cm and persistence of the right hilar nodal mass.
- However, new PET-avid soft tissue foci seen in the deep subcutaneous tissue of the left posterior abdominal wall.
- Punch biopsy of the new subcutaneous mass revealed pathological characteristics similar to the previous biopsy, confirming metastatic dissemination.
- 2nd line - palliative gemcitabine/docetaxel → severe life-threatening treatment-related toxicities: neutropenic fever, pneumonia, severe fatigue, mouth sores, oral candidiasis, nausea/vomiting, decreased oral intake, and prerenal acute renal failure.

Case 3

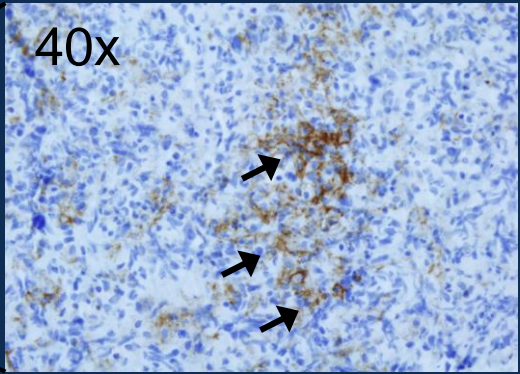
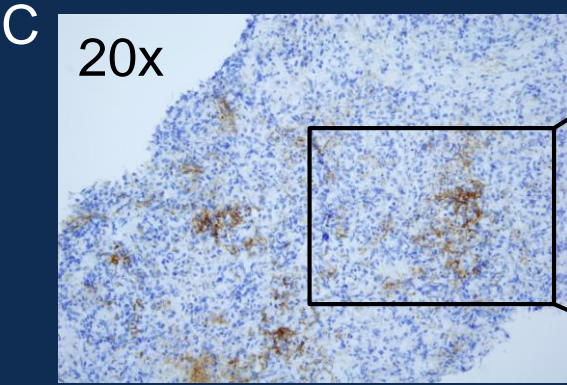
- MD Anderson Rx for malignant SFTP:
Temozolomide and Bevacizumab (temozolomide 150 mg/m² orally on days 1-7 and days 15-21, plus Bevacizumab 5 mg/kg IV on days 8 and 22, repeat every 28 days cycle) → further progression of the disease
- Comprehensive multi-platform molecular tumor profiling (CMI) while patient was undergoing temozolomide/ bevacizumab therapy. CMI-X tumor profiling confirmed positive **PD-1** and **PD-L1** (2+, 5%) in IHC.
- Temozolomide/bevacizumab tx was eventually found ineffective.
- Key genomic alterations: **TP53-V157F** (56%), **CDKN2A-R112P** (62%), and **MLH1-E234Q** (52%).

Case 3 - Pathology

H&E



CD34

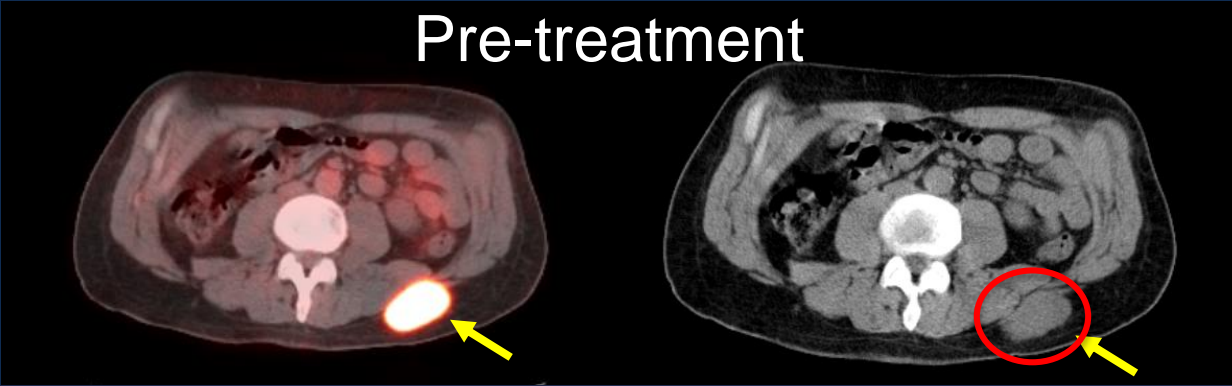


PD-L1

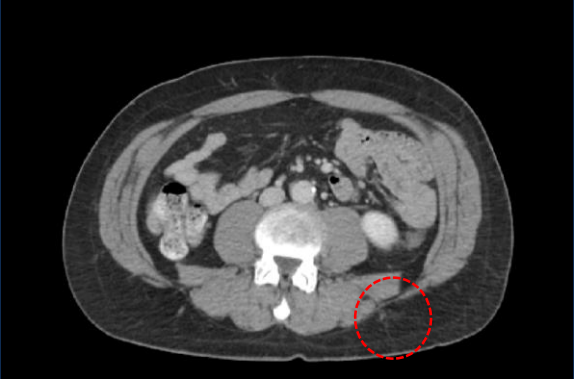
Case 3

- initiated treatment with an anti-PD-1 immune checkpoint monoclonal antibody in October 2015; **Pembrolizumab** was selected for compassionate off-label use, using the FDA-approved dose regimen of 2 mg/kg i.v every 3 weeks.
- December 2015 after 2 cycles, excellent tolerance and remarkable prompt shrinkage response of the left abdominal wall mass, confirmed by CT scan.
- Restaging CT imaging s/p 5th cycle of pembrolizumab confirmed a near-CR in February 2016.
- Restaging via CT scan imaging in July 2016 confirmed an ongoing persistent near-complete response with no new disease recurrence.
- Durable near-complete response after up to a **total of 26 cycles** of pembrolizumab infusions, and he remains well with excellent tolerance without significant adverse effects except for mild dry skin.

Case 3 - Radiology

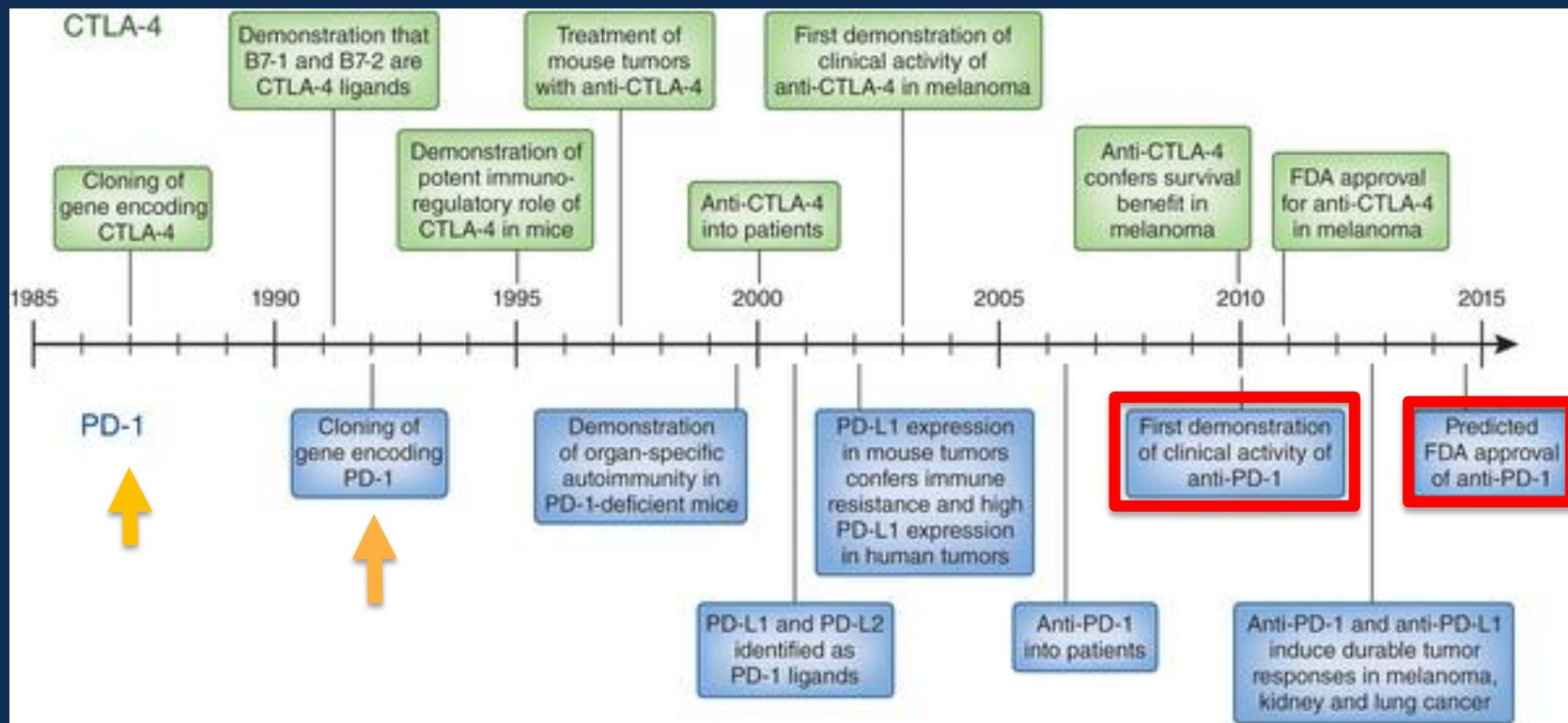


s/p C2



s/p C12 - Pembrolizumab

Progress of Immuno-Oncology



New Era - Cancer Immunotherapy: Targeting Immune Checkpoints

The NEW ENGLAND JOURNAL of MEDICINE

ORIGINAL ARTICLE

Nivolumab versus Docetaxel in Advanced Nonsquamous Non-Small-Cell Lung Cancer

H. Borghaei, L. Paz-Ares, L. Horn, D.R. Spigel, M. Steins, N.E. Ready, L.Q. Chow, E.E. Vokes, E. Felip, E. Holgado, F. Barlesi, M. Kohlhäufi, O. Arrieta, M.A. Burgio, J. Fayette, H. Lena, E. Poddubskaya, D.E. Gerber, S.N. Gettinger, C.M. Rudin, N. Rizvi, L. Crinò, G.R. Blumenschein, Jr., S.J. Antonia, C. Dorange, C.T. Harbison, F. Graf Finckenstein, and J.R. Brahmer

ABSTRACT

BACKGROUND

Nivolumab, a fully human IgG4 programmed death 1 (PD-1) immune-checkpoint inhibitor antibody, disrupts PD-1-mediated signaling and may restore antitumor immunity.

METHODS

In this randomized, open-label, international phase 3 study, we assigned patients with nonsquamous non-small-cell lung cancer (NSCLC) that had progressed during or after platinum-based doublet chemotherapy to receive nivolumab at a dose of 75 mg or 3 mg per kilogram of body weight every 2 weeks or docetaxel at a dose of 75 mg per square meter of body-surface area every 3 weeks. The primary end point was overall survival.

RESULTS

Overall survival was longer with nivolumab than with docetaxel. The median overall survival was 12.2 months (95% confidence interval [CI], 9.7 to 15.0) among patients in the nivolumab group and 9.4 months (95% CI, 8.1 to 10.7) among patients in the docetaxel group (hazard ratio for death, 0.73; 96% CI, 0.59 to 0.92; $P < .00001$). The median progression-free survival was 5.1 months (95% CI, 4.5 to 5.6) among patients in the nivolumab group and 4.2 months (95% CI, 3.7 to 4.7) among patients in the docetaxel group (hazard ratio for progression, 0.79; 96% CI, 0.64 to 0.97; $P < .00001$).

Pembrolizumab versus docetaxel for previously treated, PD-L1-positive, advanced non-small-cell lung cancer (KEYNOTE-010): a randomised controlled trial

Roy S Herbst, Paul Baas, Dong-Wan Kim, Enriqueta Felip, José L Pérez-García, Ji-Youn Han, Julian Molina, Joo-Hang Kim, Catherine Duran, Myung-Ju Ahn, Margarita Mojem, Mary J Fidlar, Gilberto de Castro Jr, Marcelo Garrido, Gregory M Lubiniecki, Yue Shentu, Ellie Im, Marisa Dolid-Filhard, Edward B Garon

Summary

Background Despite recent advances in the treatment of advanced non-small-cell lung cancer, there remain few effective treatments for progressive disease. We assessed the efficacy of pembrolizumab for previously treated, PD-L1-positive, advanced non-small-cell lung cancer.

Methods We did this randomised, open-label, phase 2/3 study at 202 academic medical centres in 24 countries with previously treated non-small-cell lung cancer with PD-L1 expression on at least 1% of tumour cells (1:1:1) in blocks of six per stratum with an interactive voice-response system to receive pembrolizumab 2 mg/kg, pembrolizumab 10 mg/kg, or docetaxel 75 mg/m² every 3 weeks. The primary endpoints were overall survival and progression-free survival both in the total population and in patients with PD-L1 expression on at least 1% of tumour cells. We used a threshold for significance of $p < 0.00825$ (one-sided) for the analysis of overall survival and progression-free survival. This trial is registered at ClinicalTrials.gov, number NCT01287585.

Findings Between Aug 28, 2013, and Feb 27, 2015, we enrolled 1034 patients: 345 allocated to pembrolizumab 2 mg/kg, 346 allocated to pembrolizumab 10 mg/kg, and 343 allocated to docetaxel. By Sept 30, 2015, 521 patients in the total population, median overall survival was 10.4 months with pembrolizumab 2 mg/kg, 12.2 months with pembrolizumab 10 mg/kg, and 8.5 months with docetaxel. Overall survival was significantly longer for pembrolizumab 2 mg/kg versus docetaxel (hazard ratio [HR] 0.71, 95% CI 0.58–0.88; $p = 0.0008$) and for pembrolizumab 10 mg/kg versus docetaxel (0.61, 0.49–0.75; $p < 0.0001$). Median progression-free survival was 3.9 months with pembrolizumab 2 mg/kg, 4.0 months with pembrolizumab 10 mg/kg, and 4.0 months with docetaxel, with no significant difference between pembrolizumab 2 mg/kg and docetaxel (0.88, 0.74–1.05; $p = 0.07$) or for pembrolizumab 10 mg/kg versus docetaxel (0.79, 95% CI 0.66–0.94; $p = 0.004$). Among patients with at least 50% of tumour cells with PD-L1, overall survival was significantly longer with pembrolizumab 2 mg/kg than with docetaxel (17.3 months vs 8.2 months; HR 0.54, 95% CI 0.38–0.77; $p < 0.0002$) and with pembrolizumab 10 mg/kg versus docetaxel (17.3 months vs 8.2 months; 0.50, 0.36–0.70; $p < 0.0001$). Likewise, for this patient population, progression-free survival was significantly longer with pembrolizumab 2 mg/kg than with docetaxel (median 4.1 months; HR 0.59, 95% CI 0.44–0.78; $p < 0.0001$) and with pembrolizumab 10 mg/kg versus docetaxel (5.2 months vs 4.1 months; 0.59, 0.45–0.78; $p < 0.0001$). Grade 3–5 treatment-related adverse events were common with pembrolizumab than with docetaxel (43 [13%] of 339 patients given 2 mg/kg, 43 [13%] of 339 patients given 10 mg/kg, and 109 [35%] of 309 given docetaxel).

Atezolizumab versus docetaxel in patients with previously treated non-small-cell lung cancer (OAK): a phase 3, open-label, multicentre randomised controlled trial

Achim Rittmeyer, Fabrice Barlesi, Daniel Waterkamp, Keunchil Park, Fortunato Ciardiello, Joachim von Pawel, Shitish M Gadgeel, Toyooki Hida, Dariusz M Kowalski, Manuel Cabo Doks, Diego L Cortinovis, Joseph Leach, Jonat han Polkoff, Carlos Barrios, Fairouz Kabbinavar, Osvaldo Arén Fronteira, Filigno De Marinis, Diego L Cortinovis, Joseph Leach, Jonat han Polkoff, Carlos Barrios, Fairouz Kabbinavar, David R Gandara, for the OAK Study Group*

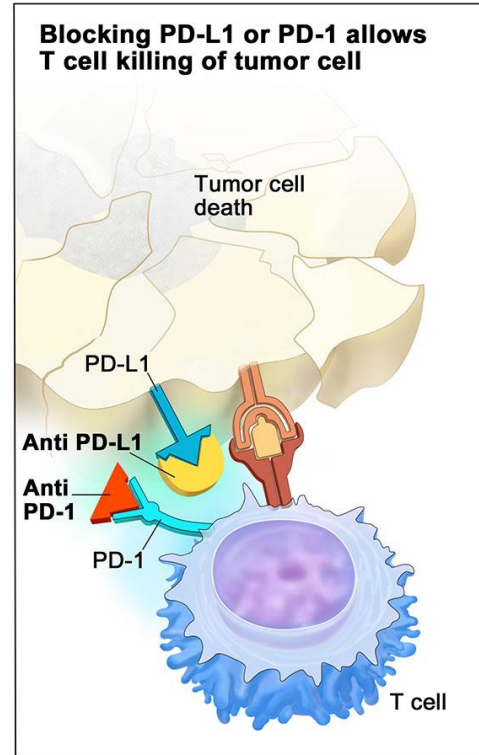
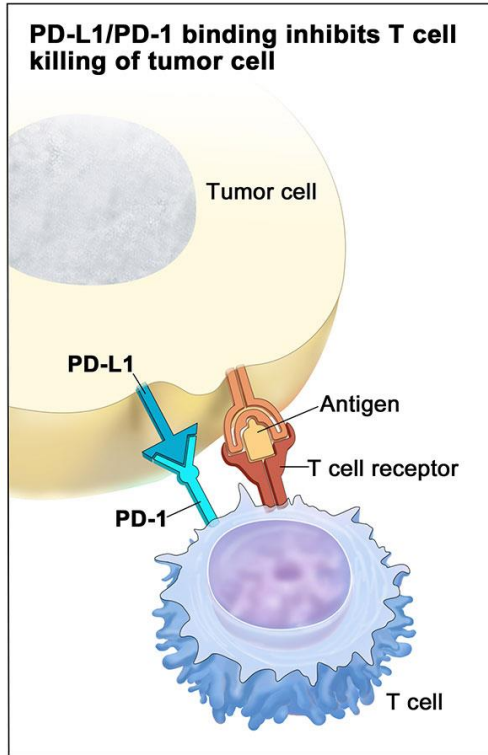
Summary

Background Atezolizumab is a humanised antiprogrammed death-ligand 1 (PD-L1) monoclonal antibody that inhibits PD-L1 and programmed death-1 (PD-1) and PD-L1 and B7-1 interactions, reinvigorating antitumor immunity. We assessed its efficacy and safety versus docetaxel in previously treated patients with non-small-cell lung cancer.

Methods We did a randomised, open-label, phase 3 trial (OAK) in 194 academic or community oncology centres in 31 countries. We enrolled patients who had squamous or non-squamous non-small-cell lung cancer, were 18 years or older, had measurable disease per Response Evaluation Criteria in Solid Tumors, and had an Eastern Cooperative Oncology Group performance status of 0 or 1. Patients had received one to two previous cytotoxic chemotherapy regimens (one or more platinum based combination therapies) for stage IIIB or IV non-small-cell lung cancer. Patients with a history of autoimmune disease and those who had received previous treatments with docetaxel, CD137 agonists, anti-CTLA4, or therapies targeting the PD-L1 and PD-1 pathway were excluded. CD137 agonists were assigned (1:1) to intravenously receive either atezolizumab 1200 mg or docetaxel 75 mg/m² every 3 weeks by permuted block randomisation (block size of eight) via an interactive voice or web response system. Coprimary endpoints were overall survival in the intention-to-treat (ITT) and PD-L1-expression population TC1/2/3 or IC1/2/3 (±1% PD-L1 on tumour cells or tumour-infiltrating immune cells). The primary efficacy analysis was done in the first 850 of 1225 enrolled patients. This study is registered with ClinicalTrials.gov, number NCT02008227.

Findings Between March 11, 2014, and April 29, 2015, 1225 patients were recruited. In the primary population, 425 patients were randomly assigned to receive atezolizumab and 425 patients were assigned to receive docetaxel. Overall survival was significantly longer with atezolizumab in the ITT and PD-L1-expression populations. In the ITT population, overall survival was improved with atezolizumab compared with docetaxel (median overall survival was 13.8 months [95% CI 11.8–15.7] vs 9.6 months [8.6–11.2]; hazard ratio [HR] 0.73 [95% CI 0.62–0.87]; $p = 0.0003$). Overall survival in the TC1/2/3 or IC1/2/3 population was improved with atezolizumab compared with docetaxel ($n = 222$; median overall survival was 15.7 months [95% CI 12.6–18.0] with atezolizumab [$n = 241$] compared with docetaxel [HR 0.74 [95% CI 0.58–0.93]; $p = 0.0102$). Patients in the PD-L1-expression population (TC1/2/3 or IC1/2/3) also had improved survival with atezolizumab compared with docetaxel (HR 0.75 [95% CI 0.59–0.96]; $p = 0.0003$). In the PD-L1-expression population, progression-free survival was significantly longer with atezolizumab compared with docetaxel (median 4.1 months [95% CI 3.7–4.5] vs 3.9 months [3.5–4.3]; HR 0.79 [95% CI 0.64–0.97]; $p = 0.0001$).

Targeting Immune PD-L1/PD-1 Checkpoint Pathway



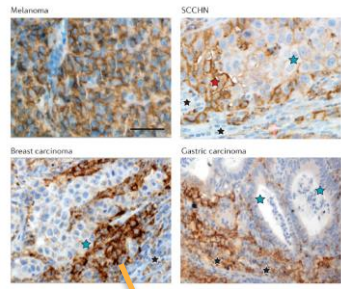
© 2015 Terese Winslow LLC
U.S. Govt. has certain rights

Targeting PD-L1/PD-1 Checkpoint Pathway

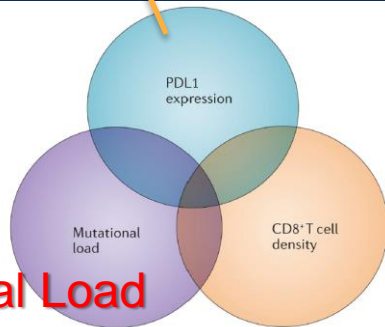
Nat Rev Cancer. 2016 May ; 16(5): 275–287. doi:10.1038/nrc.2016.36.

Mechanism-driven biomarkers to guide immune checkpoint blockade in cancer therapy

Suzanne L. Topalian¹, Janis M. Taube^{2,3,4}, Robert A. Anders⁴, and Drew M. Pardoll³



PD-L1 expression



Total Mutational Load

Figure 4. Multifactorial biomarkers of clinical response to PD1 pathway blockade

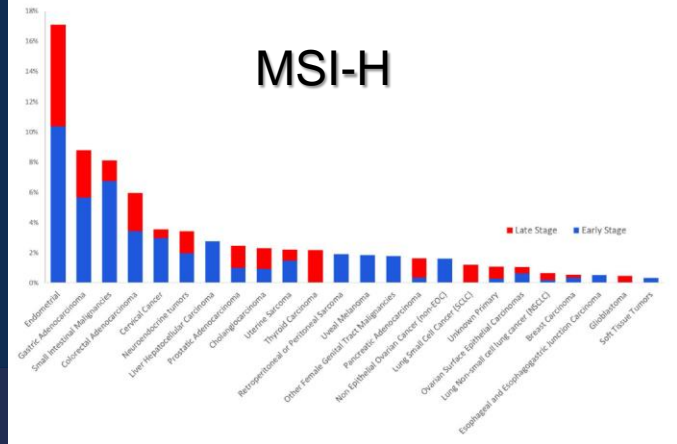
Science

REPORTS

Cite as: D. T. Le *et al.*, *Science* 10.1126/science.aan6733 (2017).

Mismatch-repair deficiency predicts response of solid tumors to PD-1 blockade

Dung T. Le,^{1,2,3} Jennifer N. Durham,^{1,2,3*} Kellie N. Smith,^{1,3*} Hao Wang,^{3*} Bjarne R. Bartlett,^{2,4*} Laveet K. Aulakh,^{2,4} Steve Lu,^{2,4} Holly Kemberling,³ Cara Wilt,³ Brandon S. Luber,³ Fay Wong,^{2,4} Nilofer S. Azad,^{1,3} Agnieszka A. Rucki,^{1,3} Dan Laheru,³ Ross Donehower,³ Atif Zaheer,⁵ George A. Fisher,⁶ Todd S. Crocenzi,⁷ James J. Lee,⁸ Tim F. Greten,⁹ Austin G. Duffy,⁹ Kristen K. Ciombor,¹⁰ Aleksandra D. Eyring,¹¹ Bao H. Lam,¹¹ Andrew Joe,¹¹ S. Peter Kang,¹¹ Matthias Heldhoff,³ Ludmila Danilova,^{1,3} Leslie Cope,^{1,3} Christian Meyer,³ Shilin Zhou,^{1,3,4} Richard M. Goldberg,¹² Deborah K. Armstrong,³ Katherine M. Bever,³ Amanda N. Fader,¹³ Janis Taube,^{1,3} Franck Housseau,^{1,3} David Spetzler,¹⁴ Nianqing Xiao,¹⁴ Drew M. Pardoll,^{1,3} Nickolas Papadopoulos,^{3,4} Kenneth W. Kinzler,^{3,4} James R. Eshleman,¹⁵ Bert Vogelstein,^{1,3,4} Robert A. Anders,^{1,3,15} Luis A. Diaz Jr.^{1,2,3,†}



Caris Precision Oncology Alliance (POA) - WVU Cancer Institute

Caris Precision Medicine Network Expands



"A comprehensive understanding of the molecular drivers of cancer has become critically important in recent years to develop innovative and better ways to treat our patients under the paradigm of personalized cancer care."

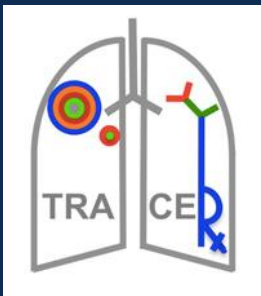
Dr. Patrick Ma

Associate Professor of Medicine and
Co-leader of the Sara Crile Allen &
James Frederick Allen Lung Cancer
Program, WVU Cancer Institute



CARIS CENTERS
OF EXCELLENCE

FOR PRECISION MEDICINE NETWORK™



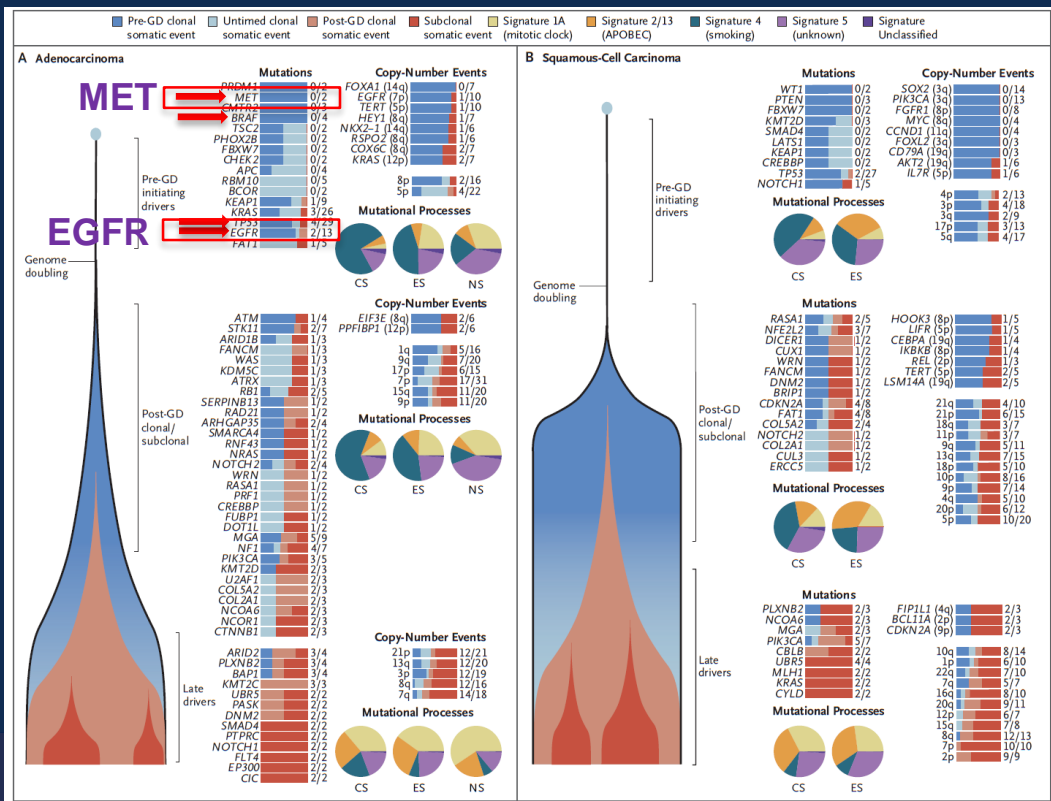
TRACERx NSCLC Evolution through Therapy

N Engl J Med 2017; 376:2109-2121, June 1, 2017

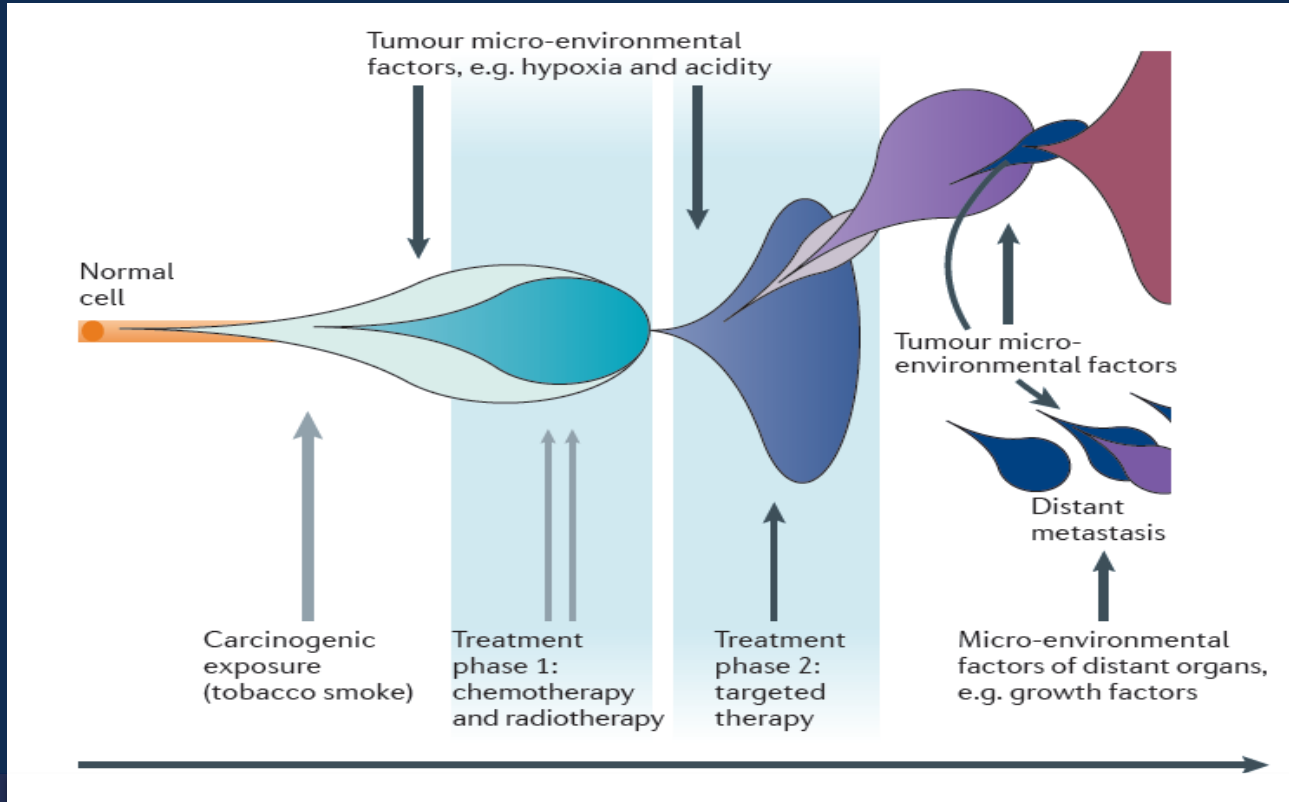
“Intratumor Heterogeneity”

(CNV, mutations) is widespread and asso. with decreased disease-free survival and is caused by chromosomal instability.

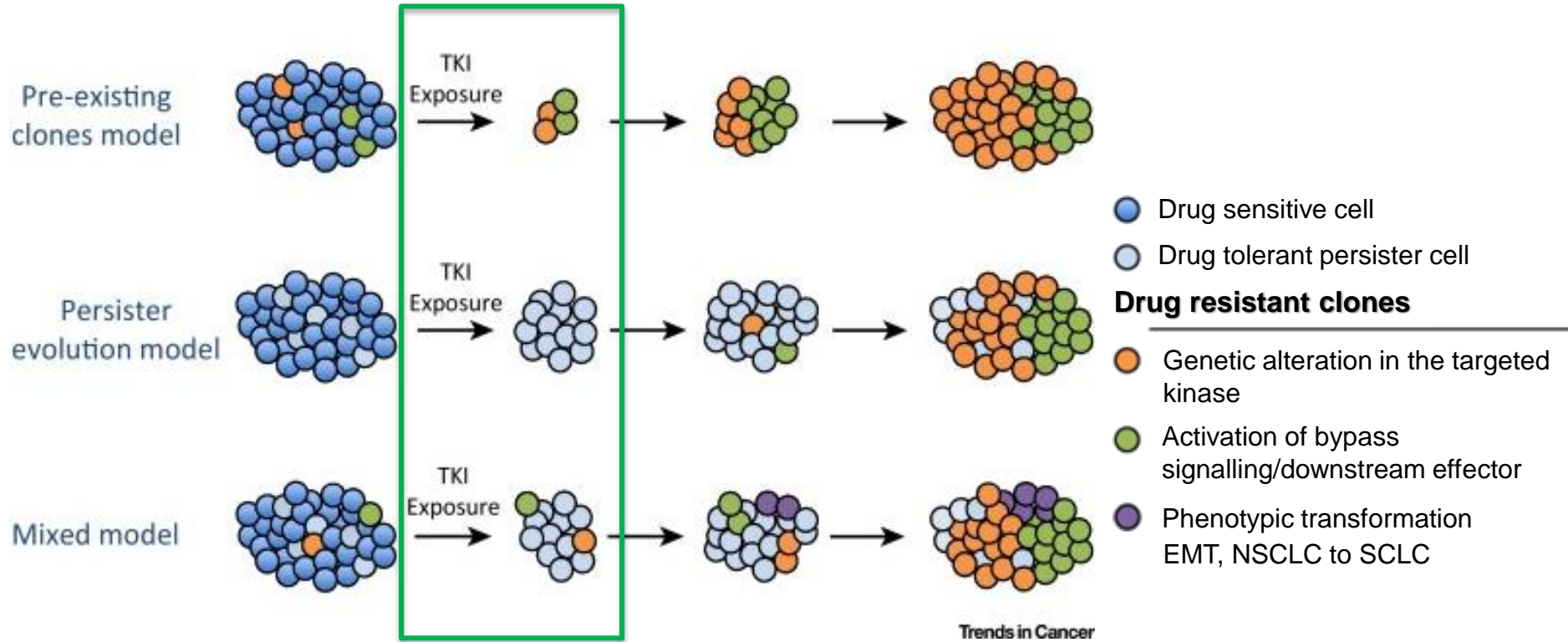
Key driver mutations, and almost always clonal (TP53, BRAF, **EGFR, MET**)



Combating Resistance: Cancer Genome Evolutionary Adaptation, Tumor Heterogeneity and Drug Resistance



Mechanisms of Targeted Therapy Drug Resistance



Combating Resistance: Cancer Genome Evolutionary Adaptation, Tumor Heterogeneity and Drug Resistance

OnCore Protocol No.: WVU011117
IRB No.: 1704546158

Title: Serial ProspECtive bIopsy for Appalachian Lung Cancer Molecular Profiling (**SPECIAL**) Study

PI: Ma, Patrick

Protocol Status: OPEN TO ACCRUAL

Translational Research in Elucidating Molecular Mechanism of Tumor Resistance in Lung Cancer Therapies

WVU011117 - Serial ProspECtive biopsy for Appalachian Lung Cancer Molecular Profiling (SPECIAL) Study

- Serial tumor biopsies-rebiopsies and blood collection on treatments
- Genomics sequencing and profiling
- Proteomics-Metabolomics profiling
- Predictive biomarkers discovery for treatment response-resistance
- Novel evolutionary treatment strategies and drug discovery