

Current Status of Targeted Therapy for NSCLC

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Potential Conflicts of Interest

- Served on advisory board for:
 - Genentech, Pfizer and BMS -- targeted therapy agents
 - Oncimmune--biomarkers lung cancer: Research grant pending approval; iSense: Research grant pending

IASLC/ATS/ERS International Classification of Lung Adenocarcinoma

www.JTO.Org

**Travis et al J Thorac Oncology
2011; 6:244-283**

Specimen Handling for Molecular Testing

- **Molecular Testing of NSCLC is Hampered by Small Diagnostic Specimen**
- **Limited Tissue is Further Reduced by Diagnostic IHC Testing and Aggressive Tissue Block Facing**
- **“Molecular Only” Testing Offers a High Success Rate at University of Colorado**

Suitability of EBUS-TBNA Samples

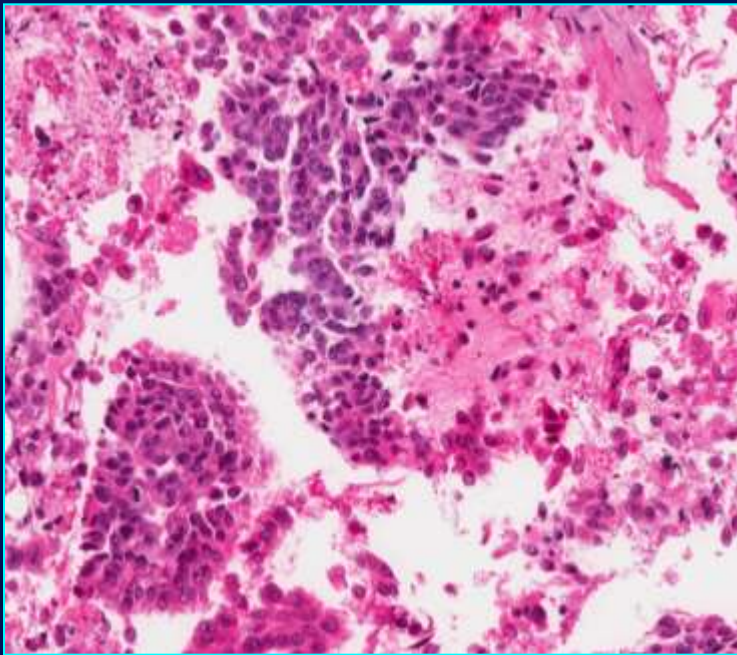
- Final Diagnosis by EBUS in Whom Subtype Was Classified 77% (CI 73-80%)
 - Rate of NOS Was Significantly Reduced When IHC Performed (OR 0.50)
- EGFR Mutational Analysis Was Possible in 90% of the Patients (Adequate Samples)
 - These Were Experienced EBUS Centers

Transthoracic Core Needle Biopsy BATTLE Lung Trial

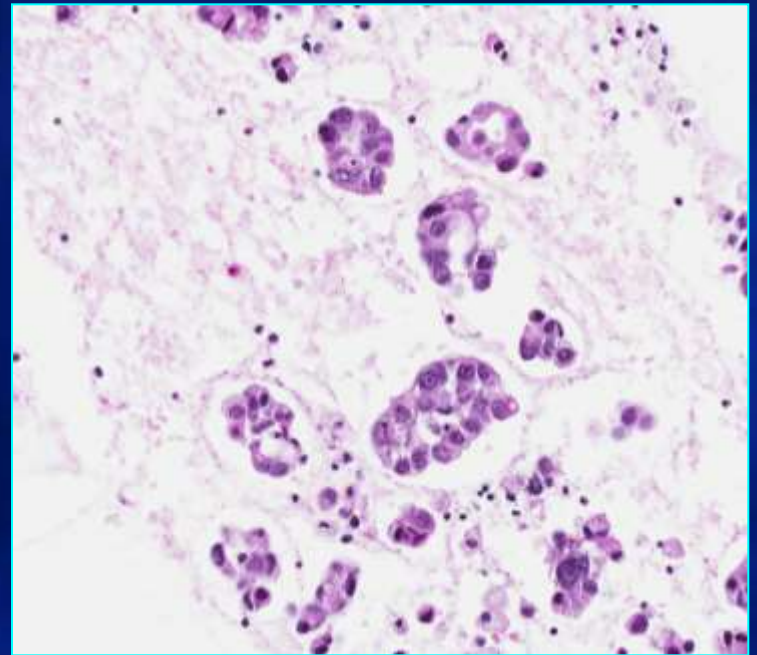
- **20G needle Bx in 151 NSCLC patients screened for trial**
- **83% specimens were adequate for biomarker testing (11 markers)**
- **Metastatic lesions were 5X more likely to yield diagnostic tissue**
- **Pneumothorax in 15%; chest tube 9%**

Use of Cytological Material for Molecular Diagnosis of Lung Cancer

Cell Block - Histology Sections



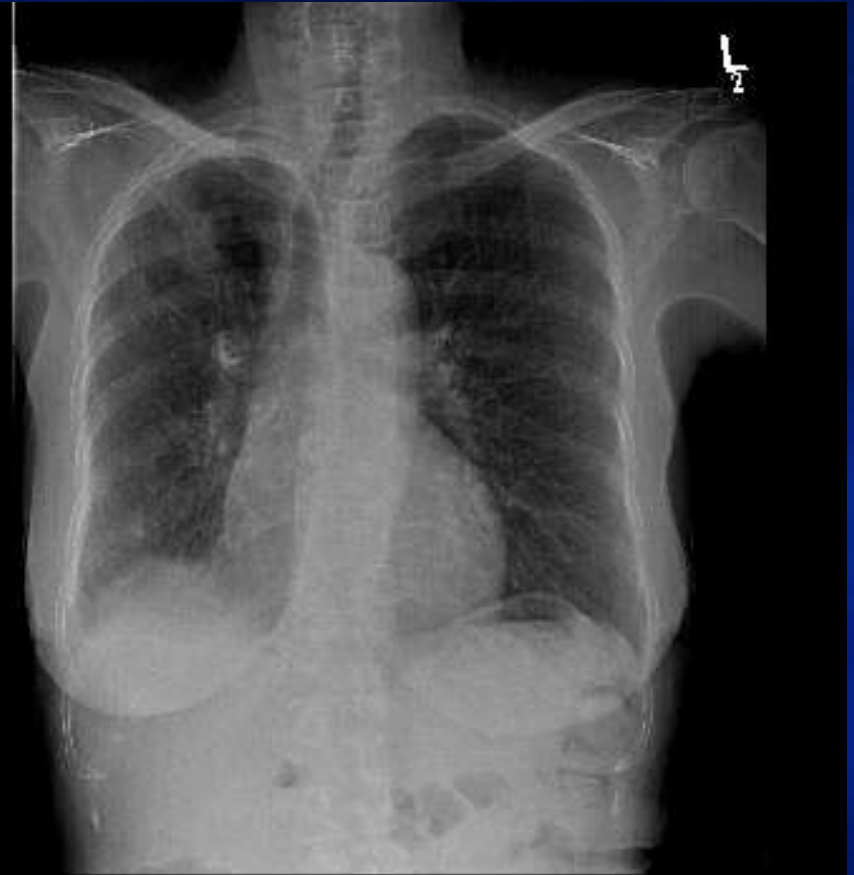
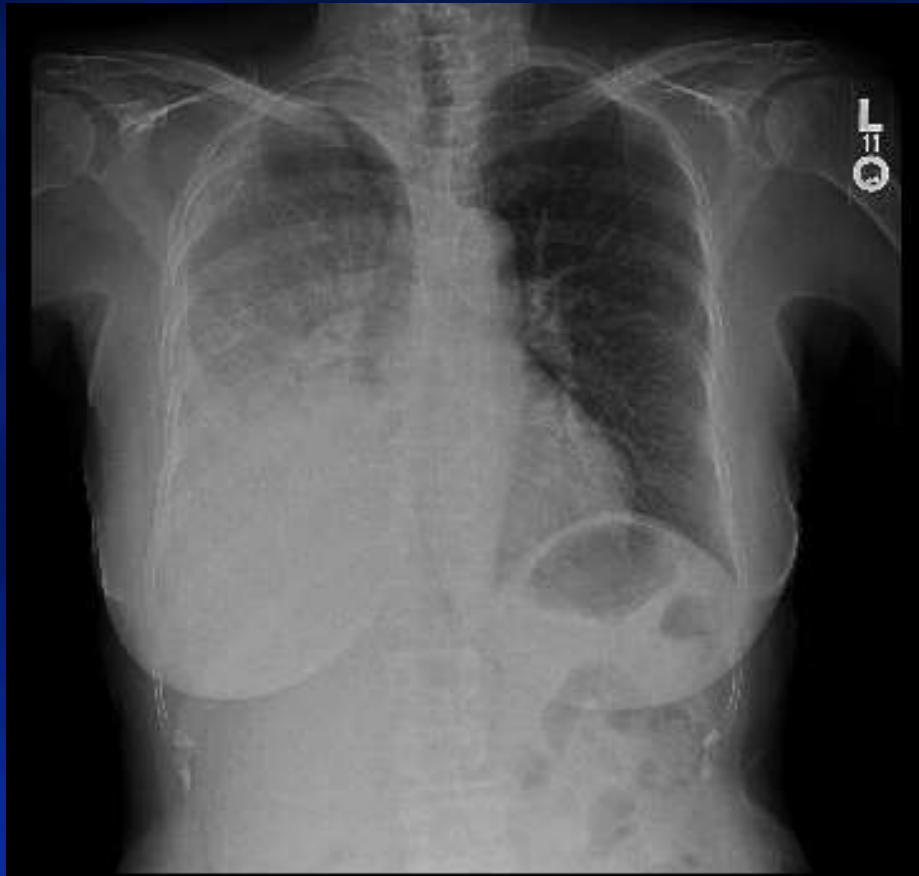
Fine Needle Aspiration



Pleural Fluid

EGFR Mutations in Tyrosine Kinase Domain

- 90% of lung cancer EGFR mutations
 - Exon 19 deletion LREA sequence (del E746-A750)
 - * Exon 21 mutation; L858R (leucine to arginine substitution)
- Resistant mutation exon 20
 - T790M
 - Exon 20 insertion

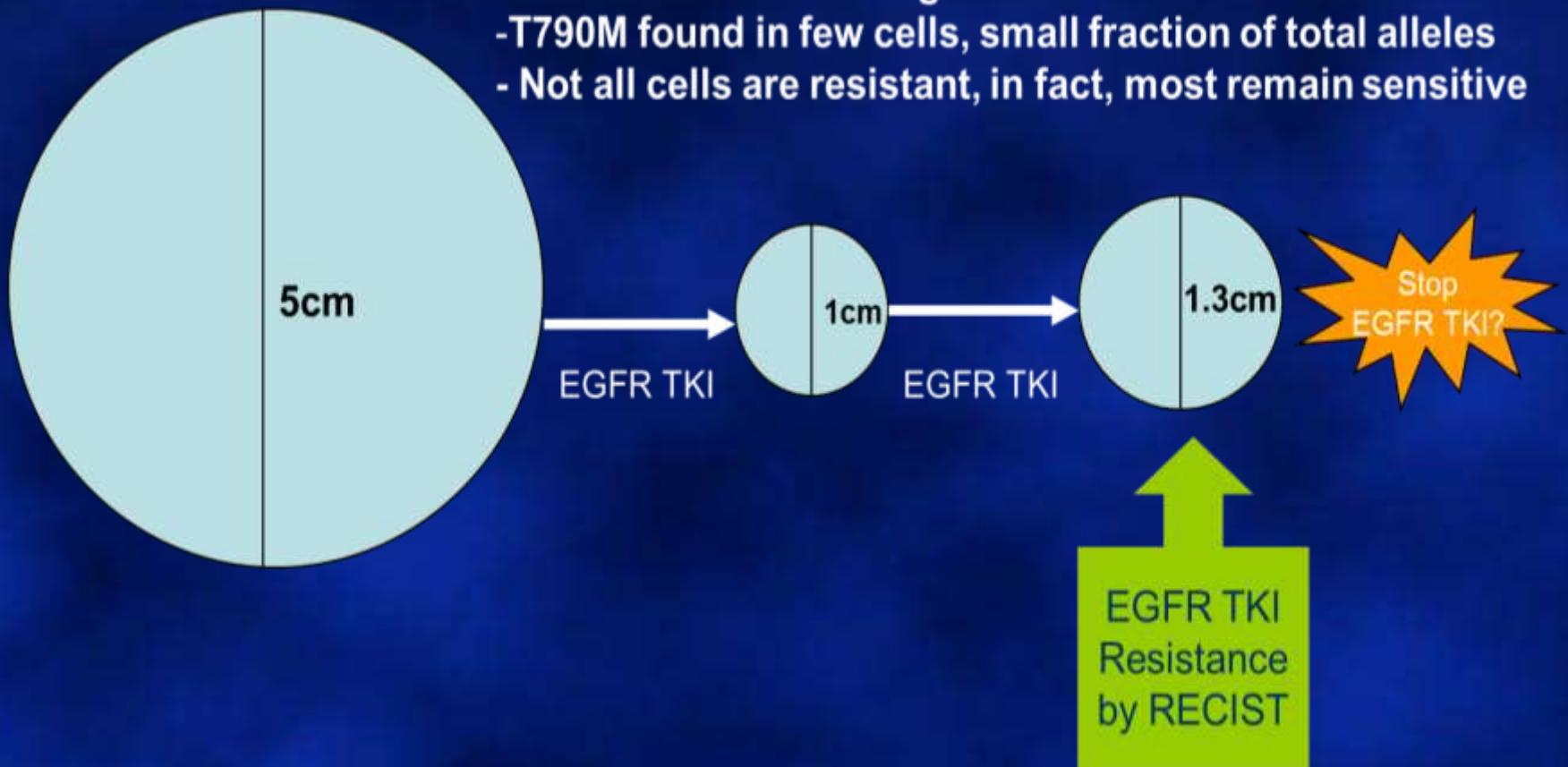


EGFR mutation: Exon 19 deletion

RECIST Criteria for Progression A Signal to Stop the EGFR TKI?

At the development of acquired resistance:

- All Cells Remain Oncogene – Addicted
- T790M found in few cells, small fraction of total alleles
- Not all cells are resistant, in fact, most remain sensitive



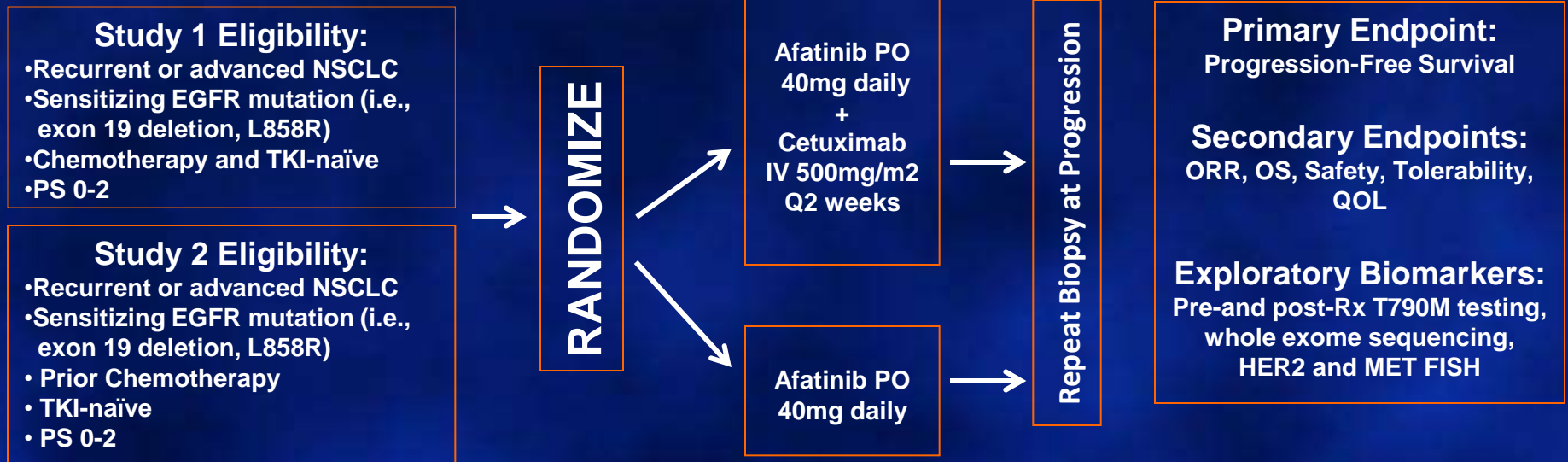
Continuation of TKI + Local Rx for TKI PD on Erlotinib or Crizotinib

Study	N pts	PFS1	PFS2
Colorado	25	10	6.2
MSKCC	18	19	10

Weickhardt A, et al, Proc ASCO 2012 # 7526

Yu A, et al, Proc ASCO 2012 # 7527

Randomized trials of afatinib plus cetuximab versus afatinib alone in treatment-naïve and previously treated patients and with advanced, *EGFR* mut+ NSCLC



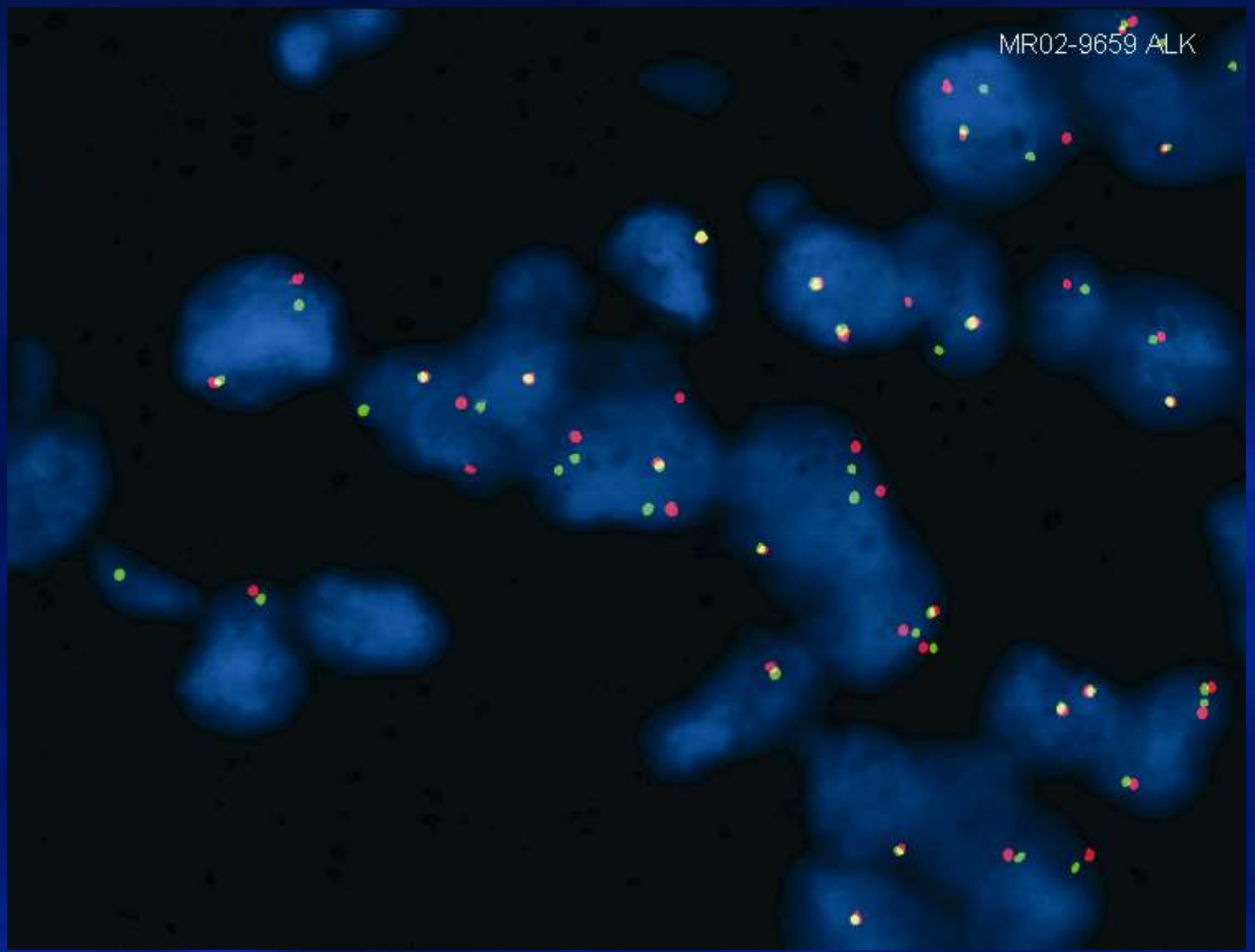
Initial Evaluation:
PET-CT
Brain CT or MRI
ECG, Echo/MUGA
Tumor Molecular
Analysis

CT Scans q8 Wks

Third Generation EGFR TKIs

- Irreversible Binding of Activating and T790M Mutations But Not Wildtype
 - WZ4002 (DFCI) - tool
 - CO-1686 (Clovis) OR=4/6 at highest dose (900mg BID-reformulation ongoing)
 - AP26113 (Ariad) – ALK, too OR 0/3 in T790M
 - TAS-2913 (Taiho)
 - AZD9291 (AZ) Phase 1 ongoing
 - Z650 (HEC Pharma)

MR02-9659 ALK



EML4-ALK in NSCLC

- Screened 141 NSCLC based on ≥ 2 of following: female, Asian, NS/light smoker, and adenocarcinoma
- 19 of 141 (13%) were ALK mutations
 - 31 (22%) were EGFR mutations
 - 91 (65%) were wild type for both

Shaw et al. J Clin Oncol 2009; 27:4247.

Preselection Screening for ALK In German Lung Cancer Patients

- Patients were negative for EGFR mutations
- 61 patients; mean age 56; 90% ACA
 - 41% female
- 16% were positive for EML4-ALK
 - 60% female, smoked less and were younger

Tufman et al J Thorac Oncol 2014; 9:109

FDA Approves Crizotinib (Xalkori): August 26, 2011

- For advanced or metastatic NSCLC that is ALK transformation positive
- FDA approved Vysis ALK Break apart FISH probe kit (Abbott Molecular, Inc)
- 255 patients; 96% adenoca; 95% metastatic; 94% had prior treatment; median age 52 yrs
- ORR was 50% and 61% in 2 studies with MDR of 42 and 48 weeks; only 1% CR
- Mild visual disturbances; G3/4 SAE in 4%
Fatal pneumonitis in 1.6% (1st 2 mos)

Parallel FISH and IHC Tests of ALK Reveal Major Discordances

- FISH by Abbott BA probe (Vysis) and Dako split probe in Rennes and Bordeaux cohorts respectively (2 probes equivalent in 100)
- IHC assayed in both centers using Abcam (clone 5A4)
 - In Rennes used Ventana automation immunostainer
 - In Bordeaux used Bond-maX automation immunostainer

Parallel FISH and IHC Tests of ALK Reveal Major Discordances

- 3244 consecutive NSCLC cases analyzed at two French Centers
- FISH and or IHC was + in 150 (4.6%)
 - Only 80 of 150 were + by both
 - FISH pos ; IHC neg in 36
 - FISH neg; IHC pos in 19
 - FISH noncontributory; IHC + in 15

Ongoing Randomized Trials of Crizotinib in ALK + NSCLC

PROFILE 1007 (N=318)

- ALK-FISH positive
- 1 prior chemotherapy (platinum-based)

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Crizotinib 250 mg BID (n=159)
[continuous]

pemetrexed 500 mg/m² or
docetaxel 75 mg/m² (n=159)
infused on day 1 of a 21-day cycle

PROFILE 1014 (N=334)

- ALK-FISH positive, non-squamous NSCLC
- No prior treatment for advanced disease

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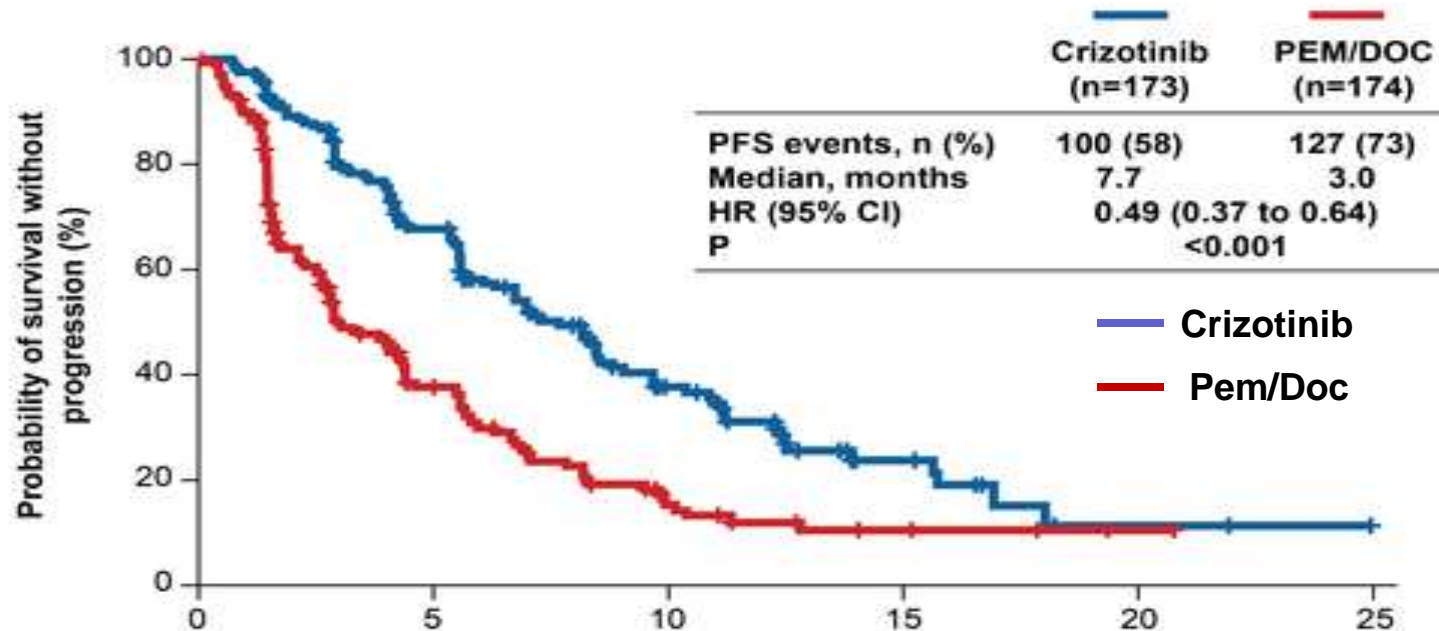
Crizotinib 250 mg BID (n=167)
[continuous]

Crossover on PD

pemetrexed/cisplatin or
pemetrexed/carboplatin (n=167)
infused on day 1 of a 21-day cycle

Crizotinib vs Chemotherapy

**PROFILE 1007 Primary Endpoint:
PFS by Independent Radiologic Review**



No. of Patients at Risk

	0	5	10	15	20	25
Crizotinib	173	93	38	11	2	0
Pem/Doc	174	49	15	4	1	0

Pem/Doc = pemetrexed/docetaxel

Shaw A, et al.^[24]

Phase 1/2 Trials of 2nd Generation ALK Inhibitors

Study	N pts	OR (N)	OR
LDK378 All*	114	66*	75%*
LDK CRZ pre*	79	45*	78%*
AP26113 pre**	24	15	73
CH5424802^	46	39	83

*ASCO 8010 OR=CR+PR+uPR; **ASCO 8031; ^ASCO 8033 all 2013

Is there space for second generation inhibitors of EGFR and ALK?

ROS1 Receptor Tyrosine Kinase Rearrangement & Crizotinib Rx

- FISH break apart assay: 250mg BID
- 25 treated and evaluable
 - Median age 51; 79% NS; 97% ACA
 - Prior Rx median 1 (range 0-7)
- ORR was 56%; 2CR; 12 PR; 8 stable
 - Median Rx duration 24 weeks
 - 6 months PFS probability 71%

Ou et al J Clin Oncol 2013; 31: abst 8032

RET Fusions

- About 1% of NSCLC
- KIF5B-RET or
 - CCDC6-RET
- Mutually Exclusive of EGFR, KRAS, and ALK Mutations

Strategies for Identifying ROS1 and RET mutations

- Occur in 2,000 to 3,000 ACA per year
- Screened 51 ACA that were negative for EGFR, ALK, KRAS, ROS1
 - 8 patients (15%) had RET fusions
- Screened 35 mutation neg (LCMC), never smokers with ACA
 - RET in 15% and ROS1 in 15%

J Clin Oncol 2013; 31 abstr 8024 (U of CO) and 8067 (MSKCC)

Vandetanib

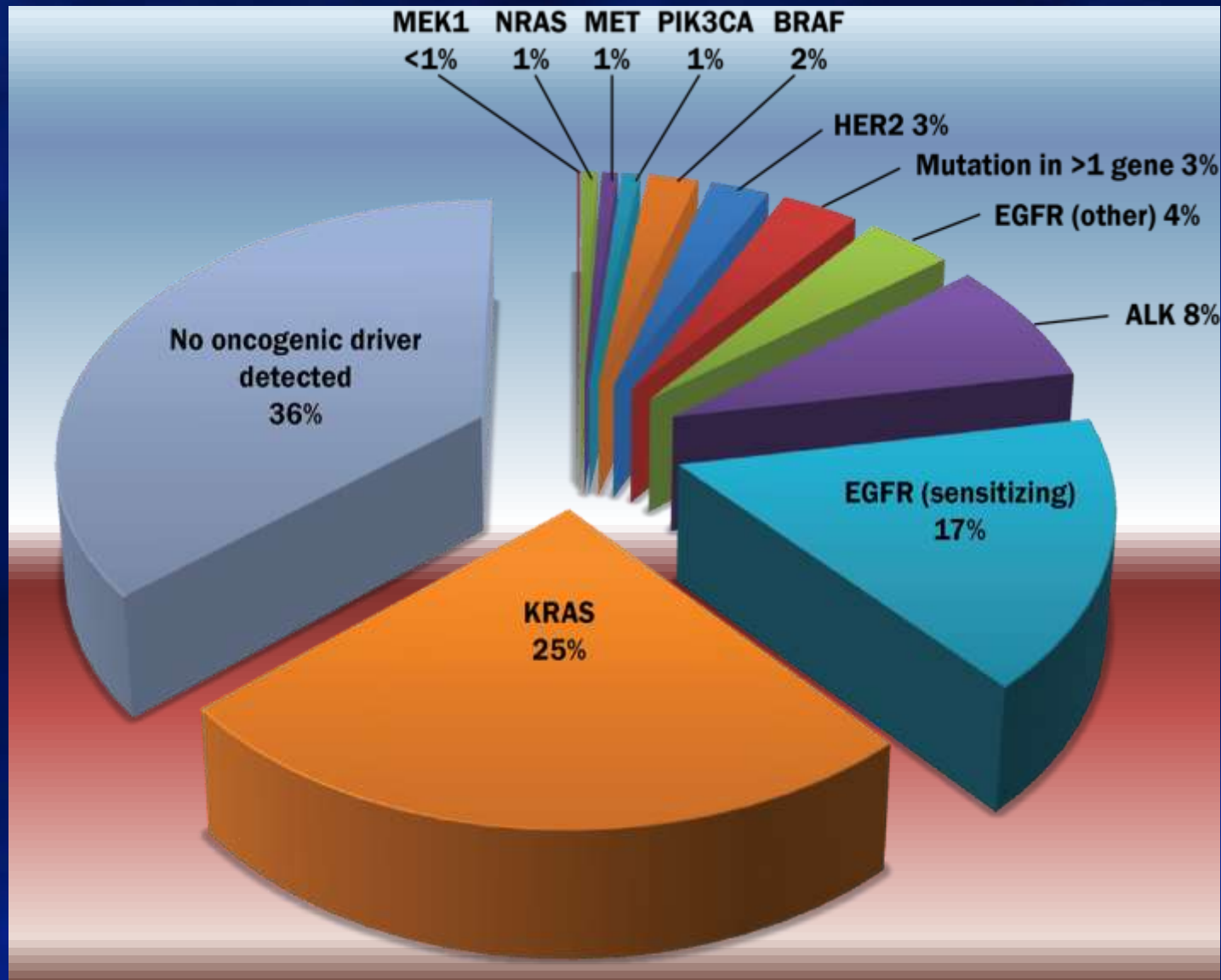
- **MultiKinase Inhibitor of RET**
 - VEGFR-2
 - EGFR
 - **Carbozantinib**: Blocks multiple RTK including RET
 - Approved for medullary thyroid cancer, and some preliminary activity in Lung ACA with RET

Lung Cancer Mutation Consortium in Lung Adenocarcinoma

- 1007 patients tested for mutation
 - 733 tested for all 10 genes
 - 60% female; median age 63
 - 34% NS; 58% former smokers
- Driver mutation found in 62%
 - Two mutations in 4%

J Clin Oncol 2013; 31: abst 8019

Lung Cancer Mutation Consortium: Incidence of Drive Mutations



Lung Cancer Mutation Consortium in Lung Adenocarcinoma

- 938 had follow-up data
 - 264 with driver mutation and Rx with a targeted agent: **MST 3.5 yrs**
 - 313 driver mutation and no Rx with targeted agent: **MST 2.4 yrs**
 - 361 with no driver mutation had a MST of **2.1 years**

French Nationwide Screening for Six Driver Mutations

- EGFR, ALK, HER2, KRAS, BRAF, PI3KCA
- First 10,000 reported after one year
 - 63% males; 83% (ex)smokers
 - 76% adenoca
- 27% KRAS; 9% EGFR; 4% ALK
 - 1% HER2; 1.6% BRAF
- **Feasibility of nationwide mutational testing**

Dabrafenib Receives Breakthrough Status by FDA for BRAF Mutated LC

- Has clinical activity in BRAF V600E melanoma patients
 - **Nearly 3000 LC patients per year USA**
- 17 patients with this mutation treated 150 mg PO BID
 - Median age 69; 12/17 males; all ACA and 13 were former smokers
- 7 PRs of 13 (54%) evaluable: DuR was 6+, 24+, 29 and 49 weeks

HER2 Mutations in Lung Cancer

- **HER2 insertions in exon 20 identified in 65 (1.7%) of 3,800 tested**
 - **Median age 60; 69% women; 52% NS; all ACA**
- **16 patients Rx; 11 PRs**
 - **DCR of 93% (n=15) with trastuzumab based Rx**
 - **DCR of 100% in 3 Rx with afatinib**

Genomic Characterization of Squamous Cell Lung Cancer

- **Comprehensive Genomic Characterization of Squamous Cell Lung Cancer: TCGA**

Nature 2012; 489:519-525

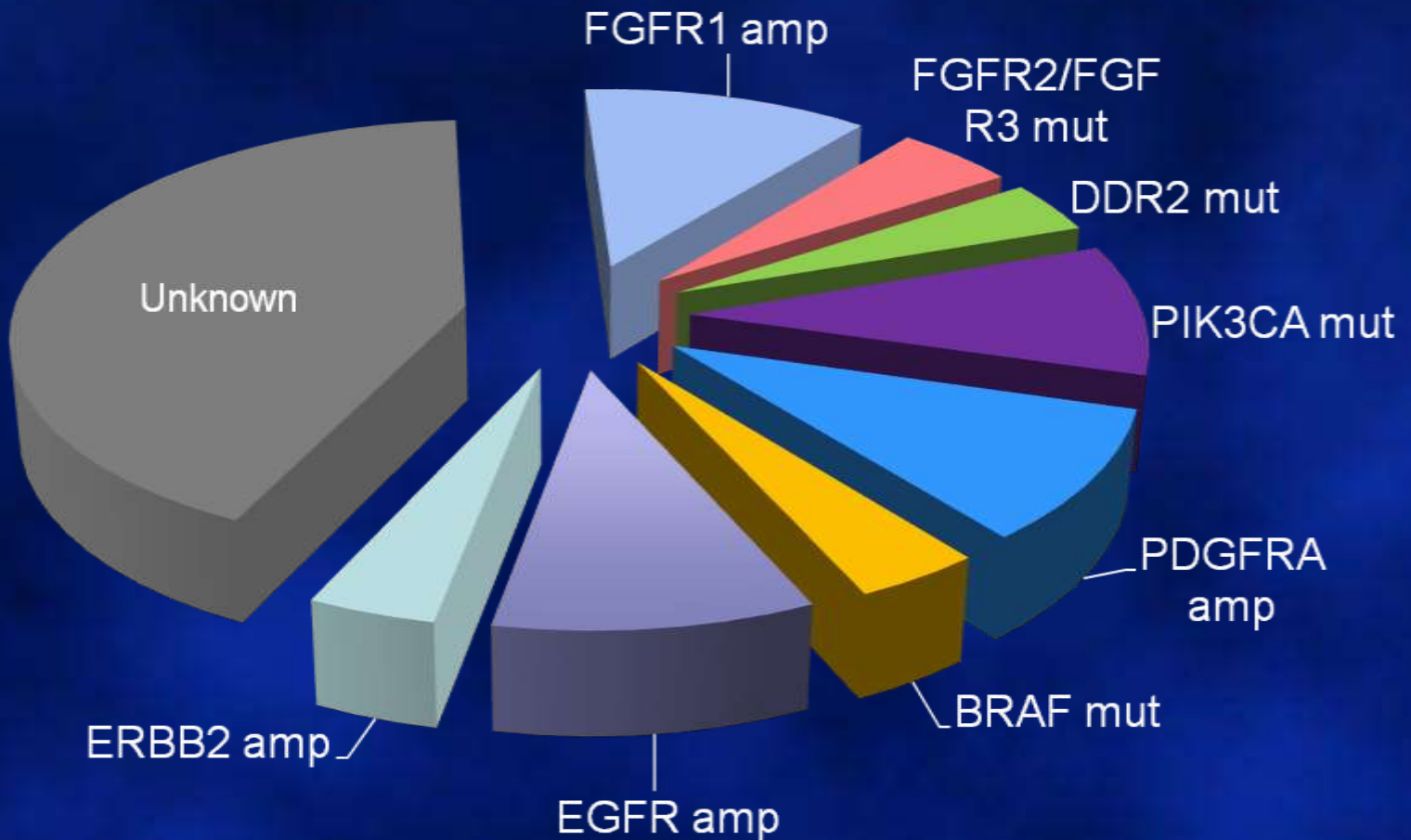
- **Integrative and Comparative Genomic Analysis of Lung Squamous Carcinomas in East Asian Patients**

Kim et al. J Clin Oncol 2013; 32:121-128

Significantly Mutated Genes in Squamous Cell Lung Cancer

TP53	81%
CCKN2A	15%
PIK3CA	16%
MLL2	20%
NFE2L2	15%
KEAP1	12%
PTEN	8%
NOTCH1	8%
RB1	7%

Genomic Characterization of Squamous Cell Lung Cancers: Potentially Targetable/Actionable Alterations



TCGA: Squamous Cell

- Previously Unreported Mutation were Loss-of-Function in HLA-A Class I Major Histocompatibility Gene
- Significantly Altered Pathways*
 - NFE2L2 and KEAP1 in 34% (oxidative stress response)
 - SOX2, TP63, NOTCH1 and 2 (squamous differentiation)

(Somatic mutations, copy number alterations, or up or down regulation)
Nature 2012;489:519

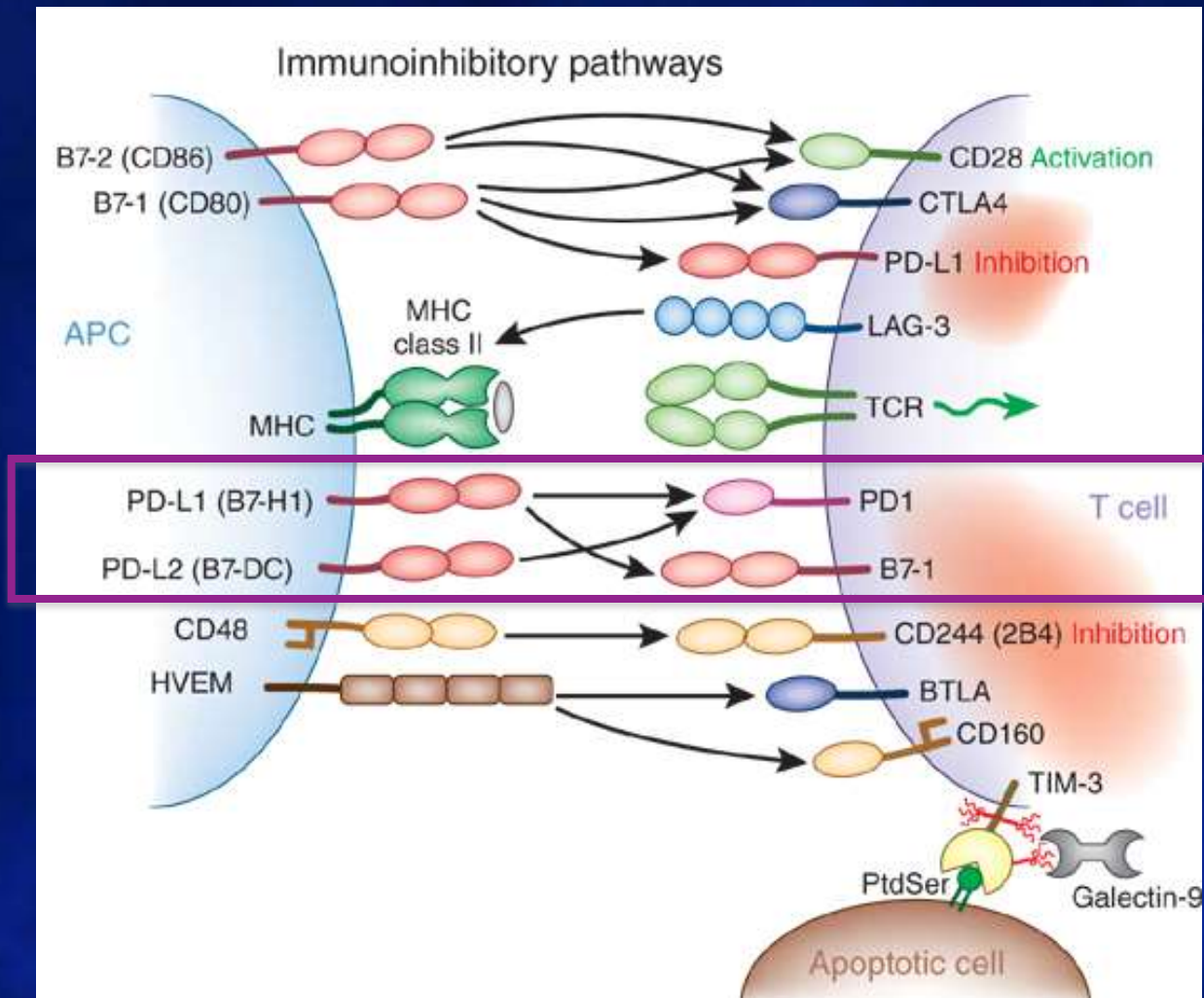
Squamous Cell Carcinoma: Korean Study

- Seven Most Common Genes Mutated
 - TP53, RB1, PTEN, NFE2L2, KEAP1, MLL2, and PIK3CA
- Mutations Were Similar to TCGA
- Identified Recurrent Occurrence of FGFR3-TACC3 Fusion
 - Potentially actionable

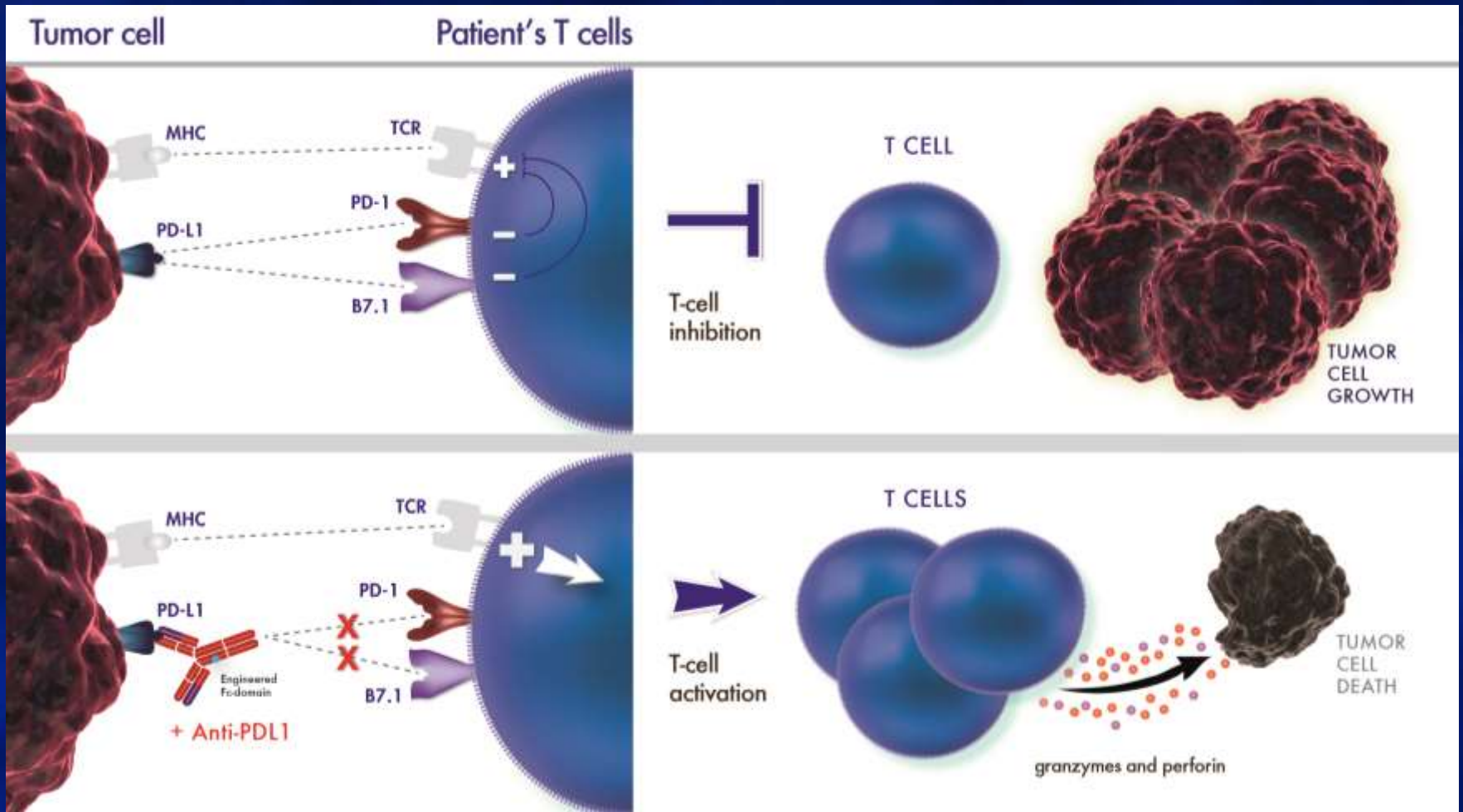
FGFR Gene Fusions in Diverse Cancers

- Whole exome sequencing (DNA) and transcriptome sequencing (RNA)
- Various cancers tested for FGFR fusions
 - Present in many cancers including **Squamous cell LC and HNSCC**
 - Six lung cancer with fusions of FGFR1 (1), FGFR2 (1), FGFR3 (4)
- **FGFR fusions are active kinases**

How Do PD-L1 Blockade and PD-1 Blockade Differ?



MPDL3280A (Anti-PDL1) Inhibits the Binding of PD-L1 to PD-1 and B7.1



Blocking PD-L1 restores T-cell activity, resulting in tumor regression in preclinical model. Binding to PD-L1 leaves PD-1/PD-L2 interaction intact and may enhance efficacy and safety.

Preliminary Efficacy Data in NSCLC

Target	Agent	N	ORR%	SD>24 wks	24 wk PFSR
PD-1	Nivolumab- BMS-936558 1-10 mg/kg	129	17.1%	10.1%	--
	Nivolumab 5,10 mg/kg + Platinum Doublet	56	~45%	--	36-71%
	Lambrolizumab MK-3475	5	1/5	--	--
PD-L1	BMS-936559 1-10 mg/kg	49	10%	12%	31%
	MPDL-3280A	41	22%	12%	46%

www.MYCANCERGENOME.ORG

- Select a disease
- Select a gene
- Select a mutation

Vanderbilt-Ingram Cancer Center

NCI Supported Genomic Clinical Trials

- **SWOG 1400**
 - Biomarker Driven master Protocol for Second Line Treatment of Lung cancer (4-6 arms)
- **ALCHEMIST**
 - Adjuvant Lung Cancer Enrichment Marker Identification and Sequencing Trial

ALCHEMIST TRIAL

- Resected non-squamous Stage IB,II,III
EGFR and ALK in CLIA Lab
 - Rest of tissue sent for testing in The Cancer Genome Atlas study
 - **EGFR mutations** enrolled in Alliance A081105 adjuvant erlotinib (n=410)
 - **ALK fusions** enrolled in E4512 adjuvant crizotinib (n=360)

SWOG 1400

- **Second line squamous cell mutational testing platform**
- **PI3K (PIK3CA mut): GDC0032 or chemo**
- **CDK4/6: PD0332991 or chemo**
 - **Mut CCND1,CDK4/6 amp,CDKN2 mut**
- **FGFR (amp,fus,mut): AZD4547 or chemo**
- **HGF (mut MET or amp):AMG102+erlotinib or chemo**
- **If no match then consider antiPDL1 vs CT**

Suggested Mutations Approach in 2014

EGFR

ALK

KRAS

Additional FISH Testing 2014

ROS

RET

MET amp



WARNING:

SMOKING CAUSES
92% OF ORAL CANCERS
QUIT: 1 800-438-2000



Marlboro