

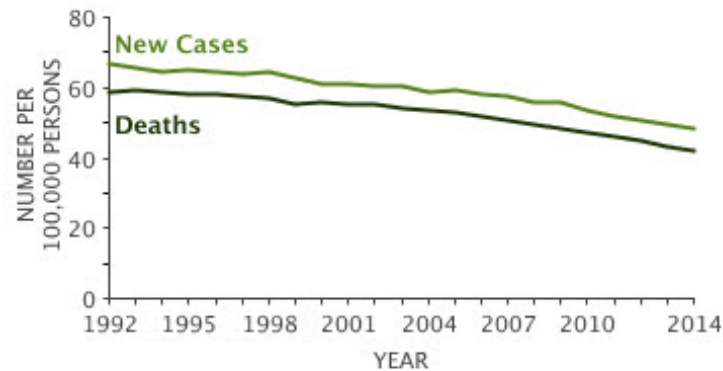
# Optimizing EBUS-TBNA in an Era of Personalized Medicine



**Nancy P. Caraway, M. D.**

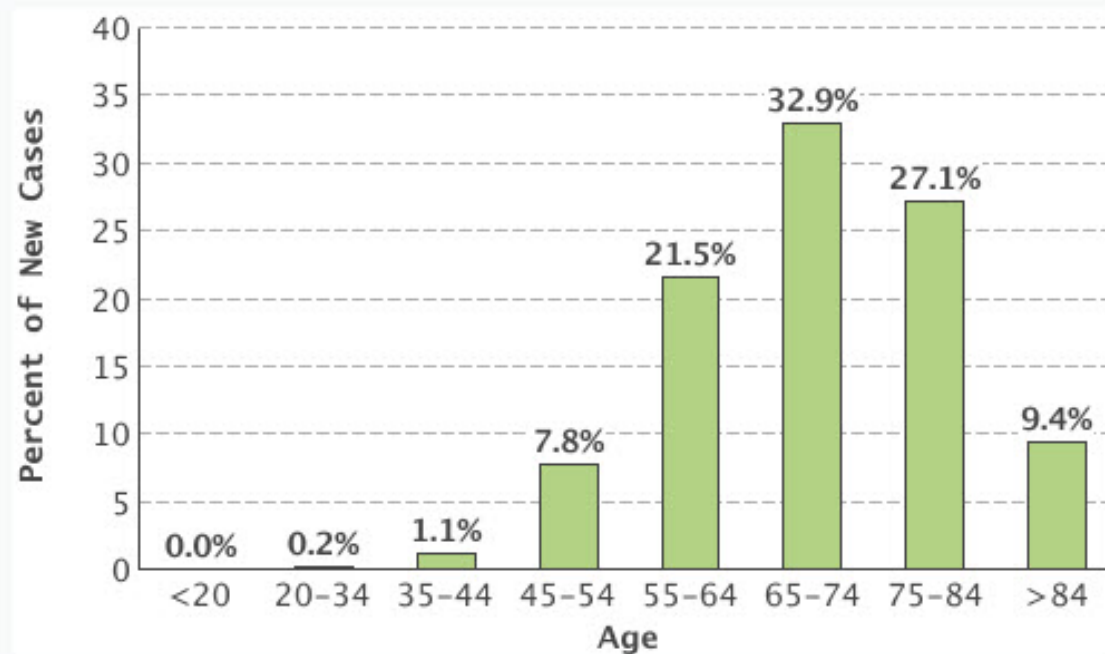
# Lung Cancer Incidence

Estimated New Cases in 2017	222,500
% of All New Cancer Cases	13.2%
Estimated Deaths in 2017	155,870
% of All Cancer Deaths	25.9%



Percent Surviving 5 Years
<b>18.1%</b>
2007-2013

Percent of New Cases by Age Group: Lung and Bronchus Cancer



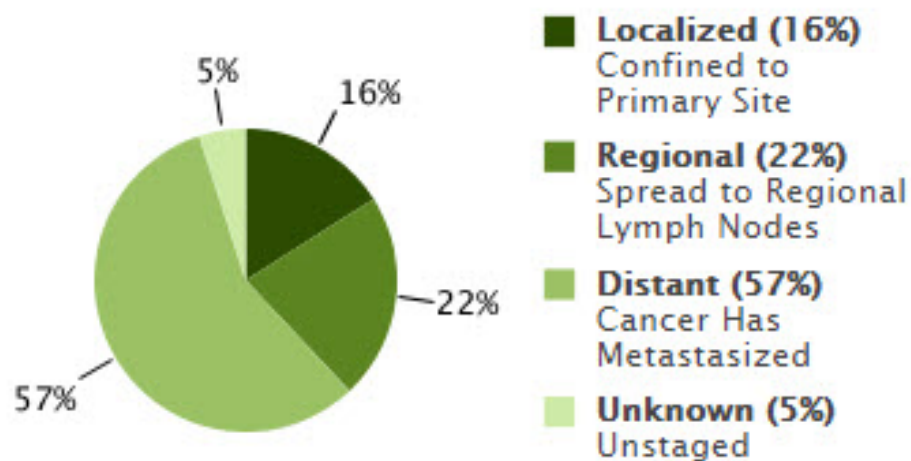
Lung and bronchus cancer is most frequently diagnosed among people aged 65-74.

Median Age At Diagnosis
<b>70</b>

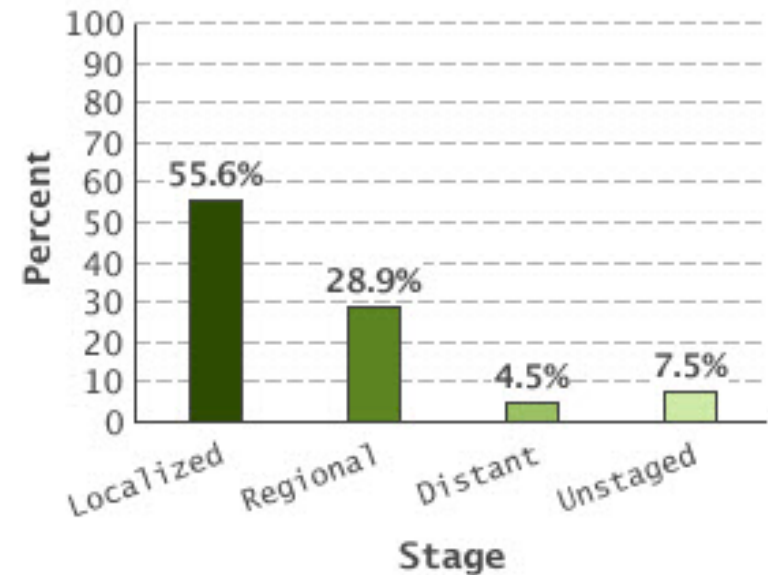
# Survival by Stage at DX

Percent of Cases & 5-Year Relative Survival by Stage at Diagnosis: Lung and Bronchus Cancer

Percent of Cases by Stage



5-Year Relative Survival



SEER 18 2007-2013, All Races, Both Sexes by SEER Summary Stage 2000

# AJCC on Lung Cancer Staging

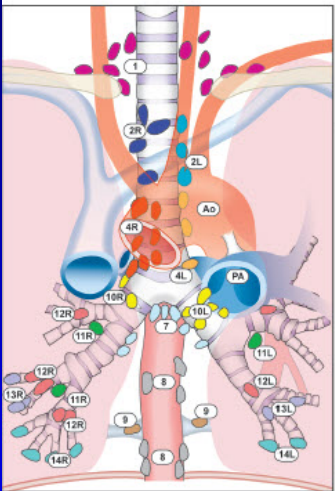
## 7<sup>th</sup> Edition

## 8<sup>th</sup> Edition

American Joint Committee on Cancer

### Lung Cancer Staging

7<sup>th</sup> EDITION



**Supraclavicular zone**

- 1 Low cervical, supraclavicular, and sternal notch nodes

**Superior Mediastinal Nodes**

*Upper zone*

- 2R Upper Paratracheal (right)
- 2L Upper Paratracheal (left)
- 3a Pre-vascular
- 3p Retrotracheal
- 4R Lower Paratracheal (right)
- 4L Lower Paratracheal (left)

**Aortic Nodes**

*AP zone*

- 5 Subaortic
- 6 Para-aortic (ascending aorta or phrenic)

**Inferior Mediastinal Nodes**

*Subcarinal zone*

- 7 Subcarinal

*Lower zone*

- 8 Paraesophageal (below carina)
- 9 Pulmonary ligament

**N<sub>1</sub> Nodes**

*Hilar/Interlobar zone*

- 10 Hilar
- 11 Interlobar

*Peripheral zone*

- 12 Lobar
- 13 Segmental
- 14 Subsegmental

**Regional Lymph Nodes (N)**

**NX** Regional lymph nodes cannot be assessed

**N0** No regional lymph node metastases

**N1** Metastasis in ipsilateral peribronchial and/or ipsilateral hilar lymph nodes and intrapulmonary nodes, including involvement by direct extension

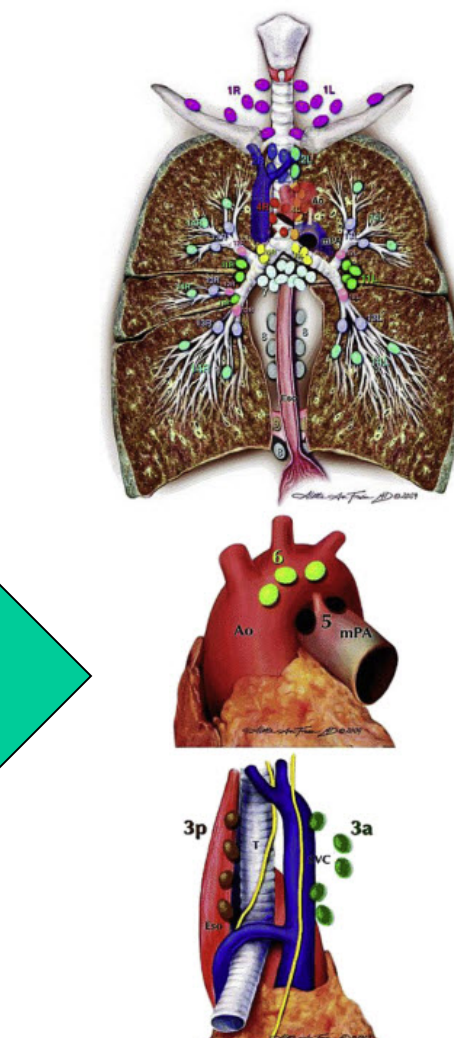
**N2** Metastasis in ipsilateral mediastinal and/or subcarinal lymph node(s)

**N3** Metastasis in contralateral mediastinal, contralateral hilar, ipsilateral or contralateral scalene, or supraclavicular lymph node(s)

**ILLUSTRATION**

The IASLC lymph node map shown with the proposed amalgamation of lymph into zones.

(© Memorial Sloan-Kettering Cancer Center, 2009.)



**Supraclavicular zone**

- 1 Low cervical, supraclavicular, and sternal notch nodes

**SUPERIOR MEDIASTINAL NODES**

*Upper zone*

- 2R Upper Paratracheal (right)
- 2L Upper Paratracheal (left)
- 3a Prevascular
- 3p Retrotracheal
- 4R Lower Paratracheal (right)
- 4L Lower Paratracheal (left)

**AORTIC NODES**

*AP zone*

- 5 Subaortic
- 6 Para-aortic (ascending aorta or phrenic)

**INFERIOR MEDIASTINAL NODES**

*Subcarinal zone*

- 7 Subcarinal

*Lower zone*

- 8 Paraesophageal (below carina)
- 9 Pulmonary ligament

**N1 NODES**

*Hilar/Interlobar zone*

- 10 Hilar
- 11 Interlobar

*Peripheral zone*

- 12 Lobar
- 13 Segmental
- 14 Subsegmental

# N Subclassification & Distant Mets

## N (Regional Lymph Nodes)

N0	No regional node metastasis
N1	Metastasis in ipsilateral pulmonary or hilar nodes
N2	Metastasis in ipsilateral mediastinal/subcarinal nodes
N3	Metastasis in contralateral mediastinal/hilar, or supraclavicular nodes

## M (Distant Metastasis)

M0	No distant metastasis
M1a	Malignant pleural/pericardial effusion <sup>c</sup> or pleural /pericardial nodules or separate tumor nodule(s) in a contralateral lobe;
M1b	Single extrathoracic metastasis
M1c	Multiple extrathoracic metastases (1 or >1 organ)

M1a *Pl. Dissem*

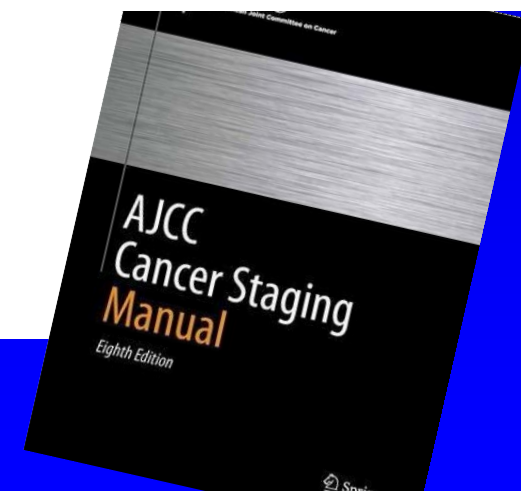
M1a *Contr. Nod*

M1b *Single*

M1c *Multi*

TABLE 4 ] N Subclassification

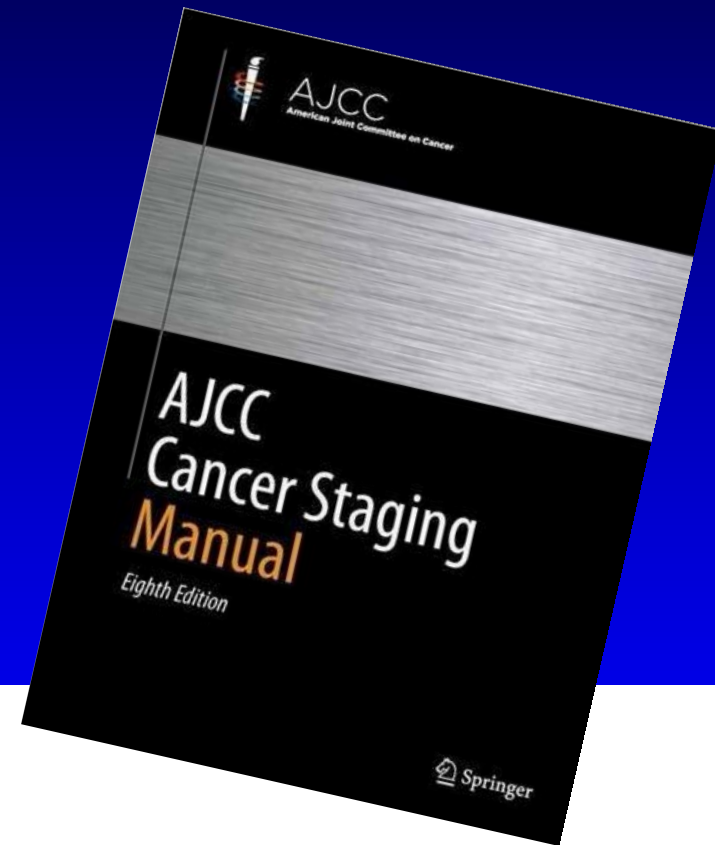
Category	Subclass	Description
Nx		Regional lymph nodes cannot be assessed
N0		No regional lymph node involvement
N1	N1a	Single-station N1 involvement
	N1b	Multiple-station N1 involvement
N2	N2a1	Single-station N2 without N1 involvement (skip)
	N2a2	Single-station N2 with N1 involvement
	N2b	Multiple-station N2 involvement
N3		N3 lymph node involvement



# Lung Cancer Grouping & Survival

TABLE 5 ] Lung Cancer Stage Grouping (Eighth Edition)

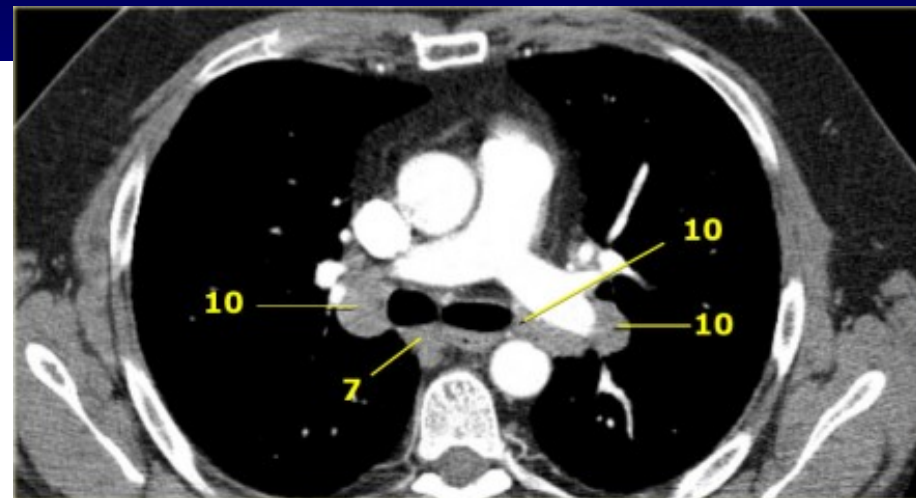
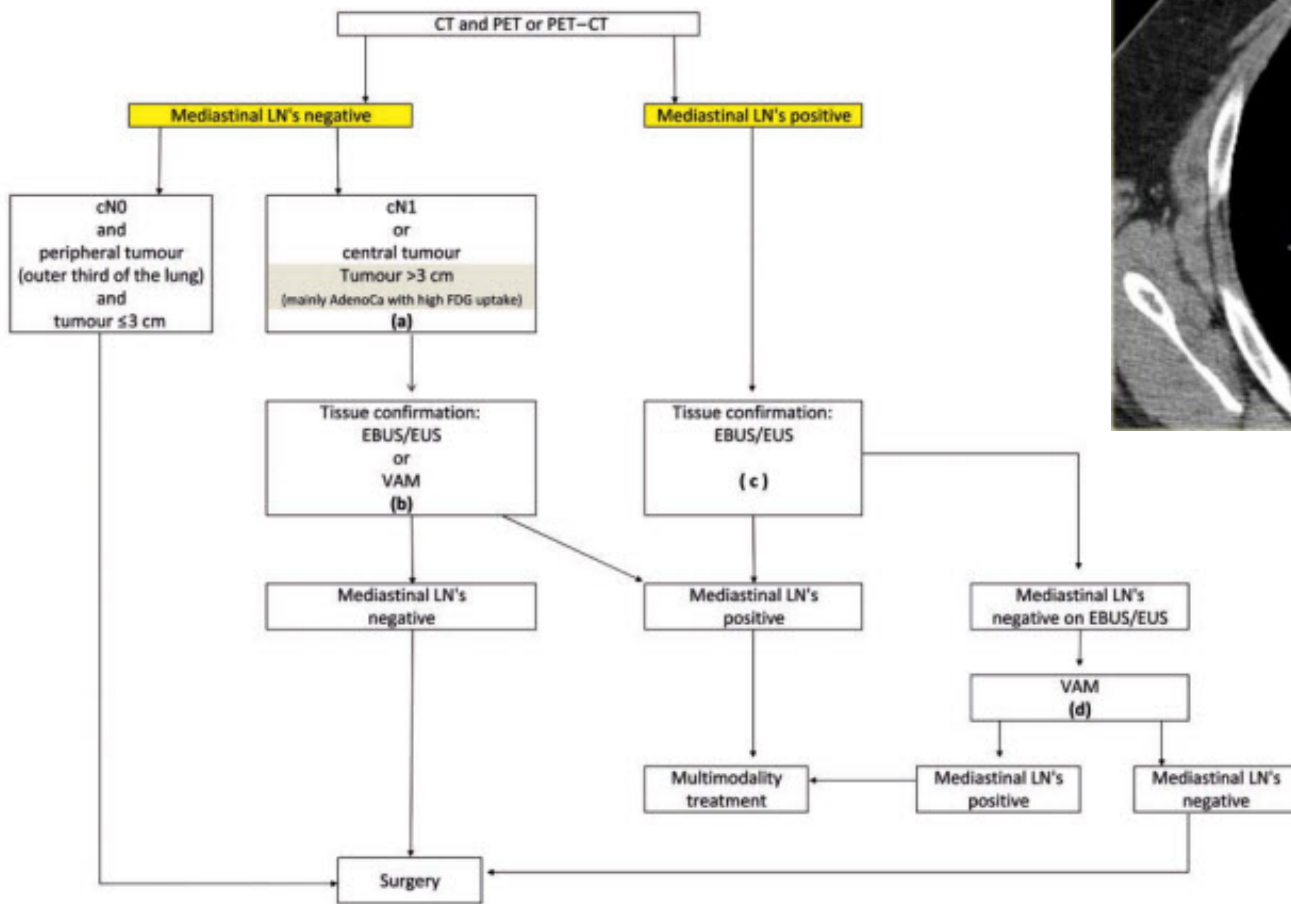
T/M	Label	N0	N1	N2	N3
T1	T1a $\leq 1$	IA1	IIB	IIIA	IIIB
	T1b >1-2	IA2	IIB	IIIA	IIIB
	T1c >2-3	IA3	IIB	IIIA	IIIB
T2	T2a <i>Cent, No PI</i>	IB	IIB	IIIA	IIIB
	T2a >3-4	IB	IIB	IIIA	IIIB
	T2b >4-5	IIA	IIB	IIIA	IIIB
T3	T3 >5-7	IIB	IIIA	IIIB	IIIC
	T3 <i>Inv</i>	IIB	IIIA	IIIB	IIIC
	T3 <i>Satell</i>	IIB	IIIA	IIIB	IIIC
T4	T4 >7	IIIA	IIIA	IIIB	IIIC
	T4 <i>Inv</i>	IIIA	IIIA	IIIB	IIIC
	T4 <i>Ipsi Nod</i>	IIIA	IIIA	IIIB	IIIC
M1	M1a <i>Contr Nod</i>	IVA	IVA	IVA	IVA
	M1a <i>PI Dissem</i>	IVA	IVA	IVA	IVA
	M1b <i>Single</i>	IVA	IVA	IVA	IVA
	M1c <i>Mult</i>	IVB	IVB	IVB	IVB



5-Year Survival (%)

Type	IA1	IA2	IA3	IB	IIA	IIB	IIIA	IIIB	IIIC	IVA	IVB
Clinical	92	83	77	68	60	53	36	26	13	10	0
Pathologic	90	85	80	73	65	56	41	24	12	-	-

# CT & PET or PET-CT in Staging Lung Cancer



<http://www.radiologyassistant.nl/en/p4646f1278c26f/mediastinum-lymph-node-map.html>

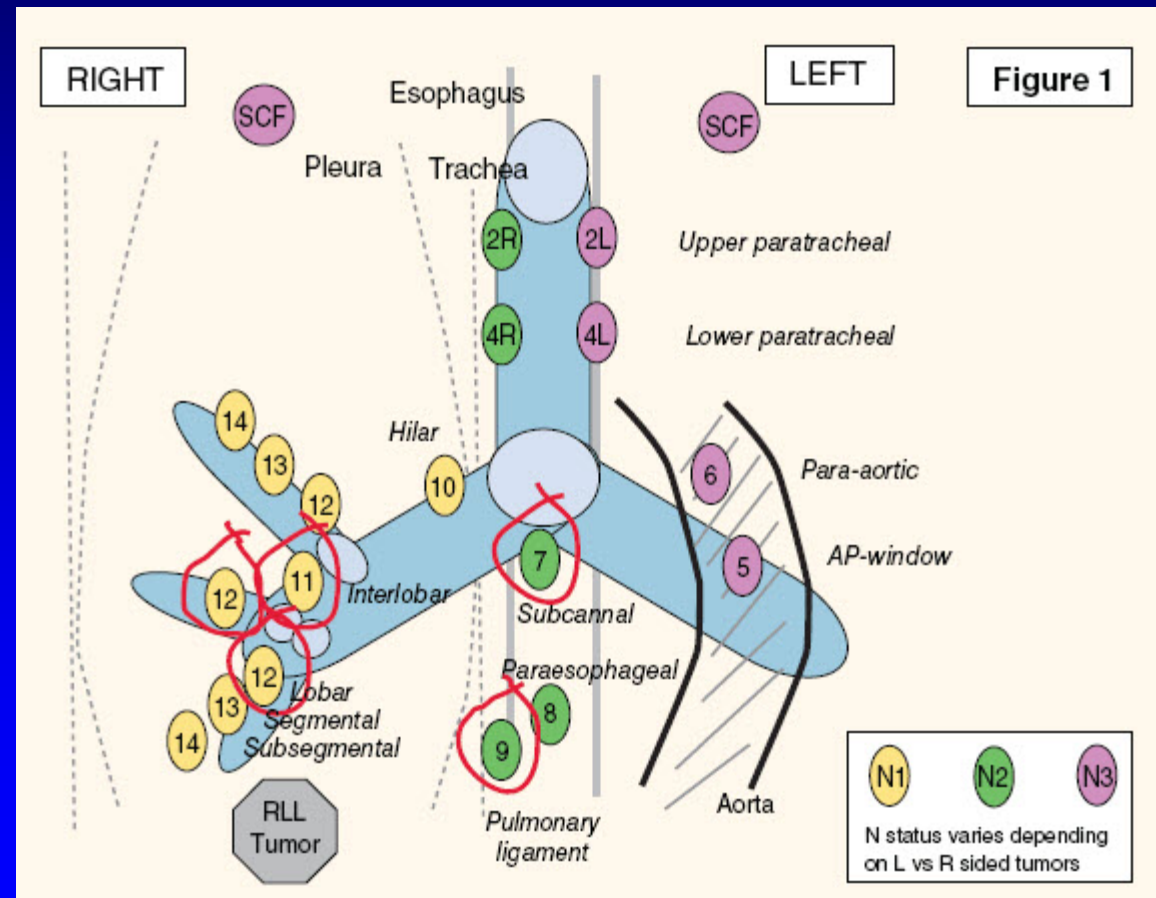
# EBUS-TBNA & EUS

- **EBUS-TBNA: LN sample**

- 2R and 2L
- 4R and 4L
- 7
- 10R and 10L
- 11R and 11L
- Sometimes 12

- **EUS: LN sample**

- 8
- 9





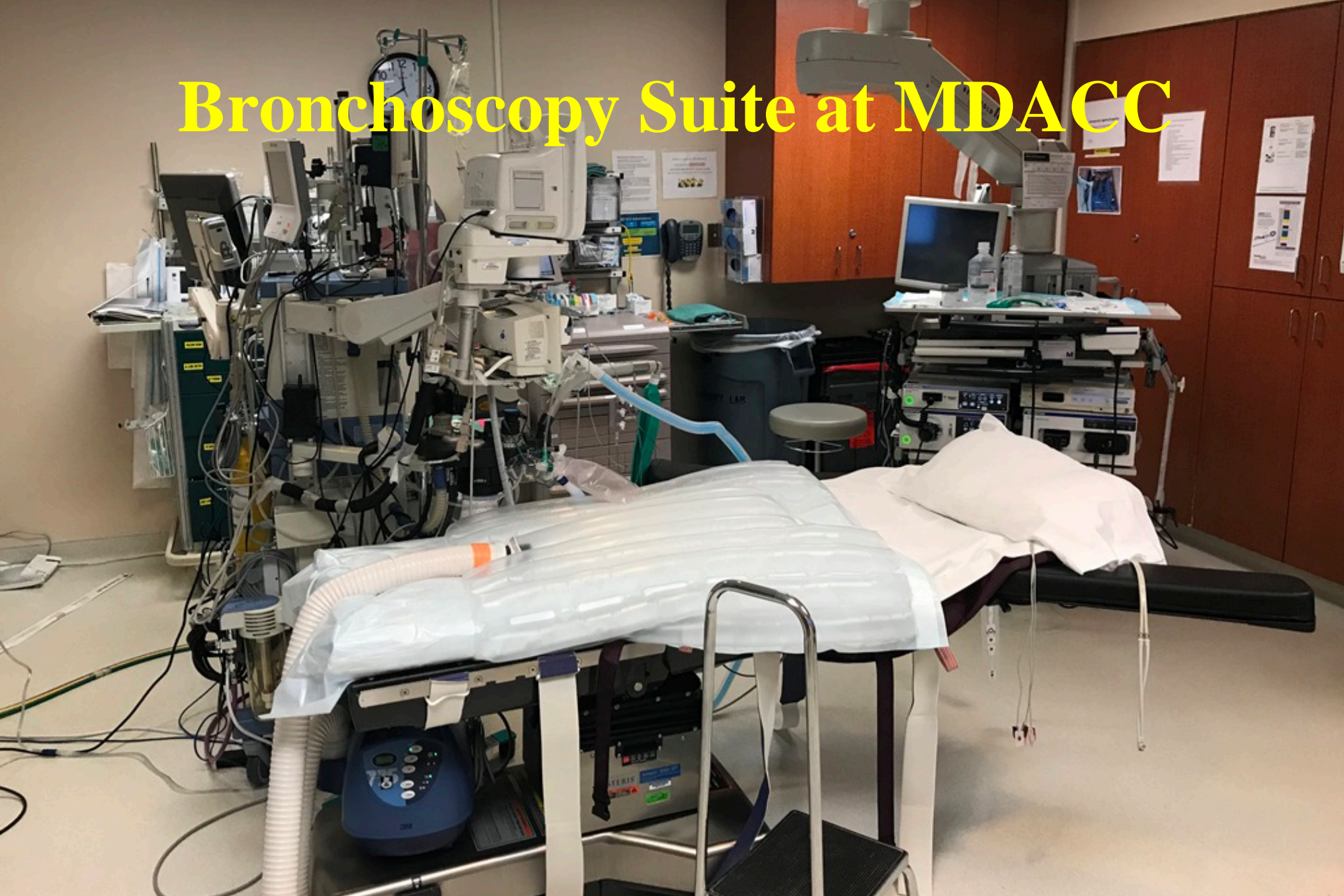
# Indications for EBUS-TBNA

- Initial staging & restaging of lung cancer
- Evaluating mediastinal adenopathy with no prior malignancy
- Evaluating mediastinal adenopathy with non-lung primary
- Evaluating mediastinal adenopathy with known non-lung carcinoma and now new lung mass
- Obtain material for mutational analysis

# **The EBUS-TBNA ROSE Advantage**

- **Real time evaluation of procured material**
- **Improves diagnostic yield**
- **Sampling stopped when adequate material obtained**
- **Specimen triaged for ancillary studies**
- **Communication between the pulmonologist and pathology faculty & staff**

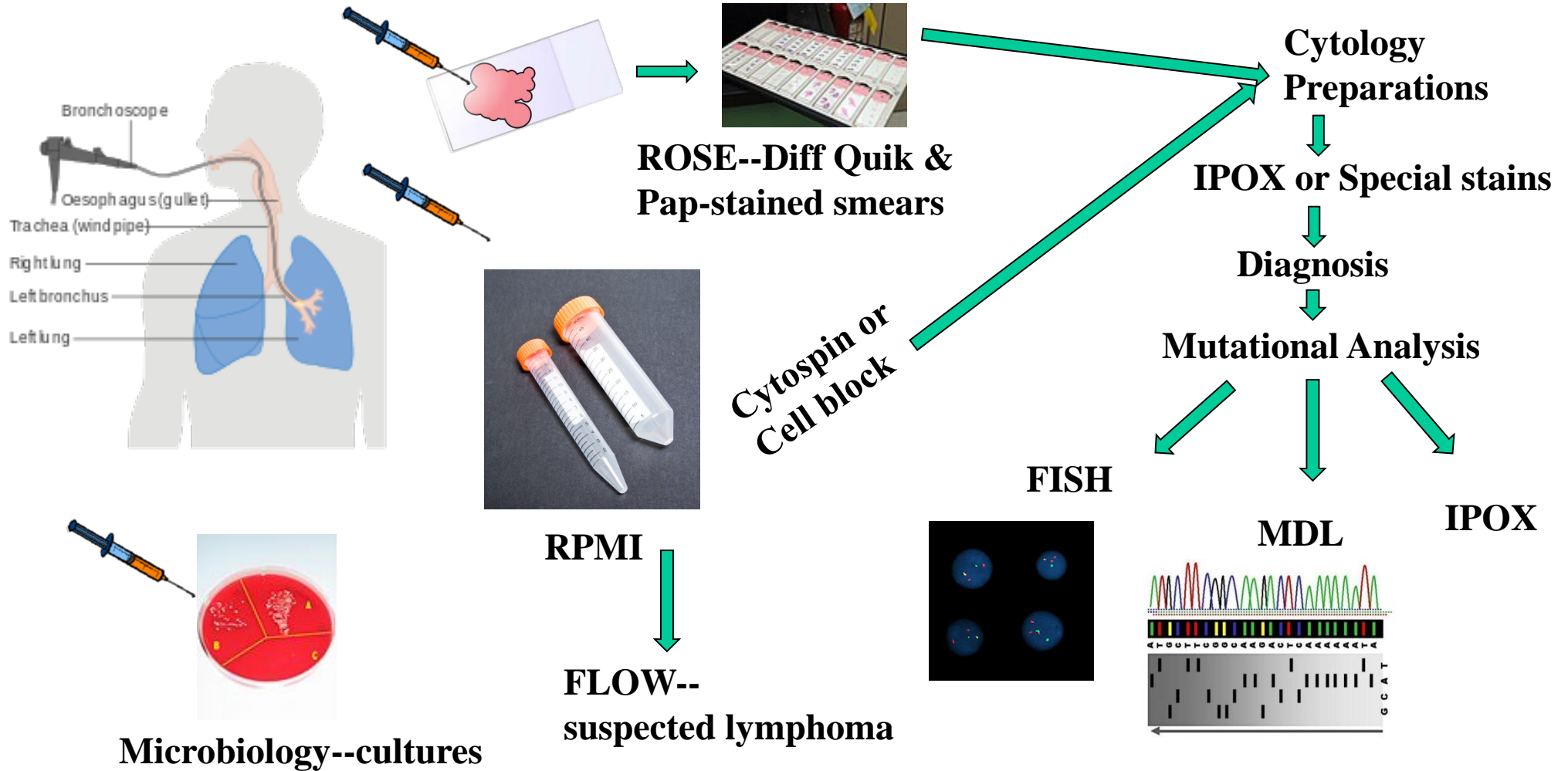
# Bronchoscopy Suite at MDACC



# Goals for Tissue Procurement

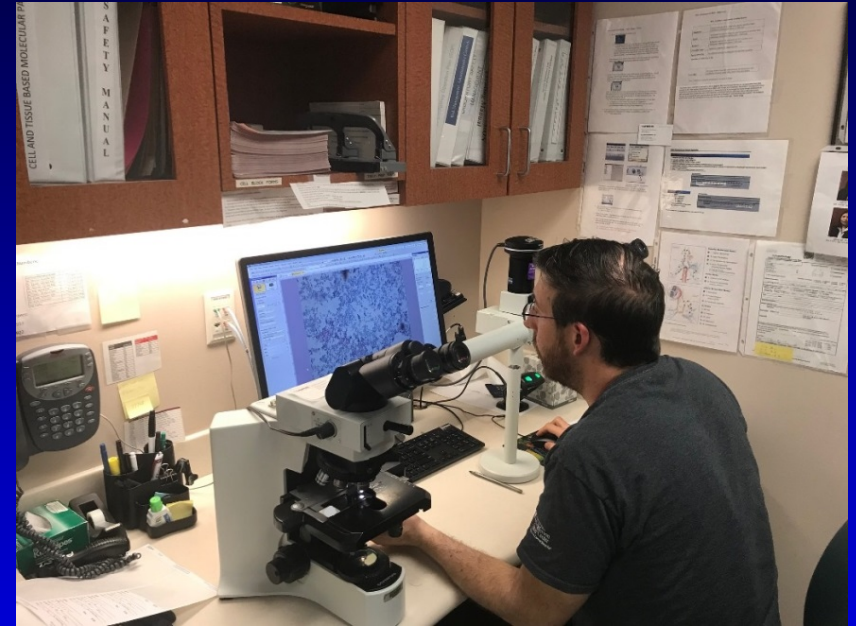
- Adequate sampling of lymph node
- Benign vs. Malignant
- Metastatic lung vs. non-lung carcinoma vs. other
- Histologic subtype
- Molecular testing

# EBUS TBNA with ROSE for Dx & Mutational Analysis



# ROSE for EBUS-TBNA

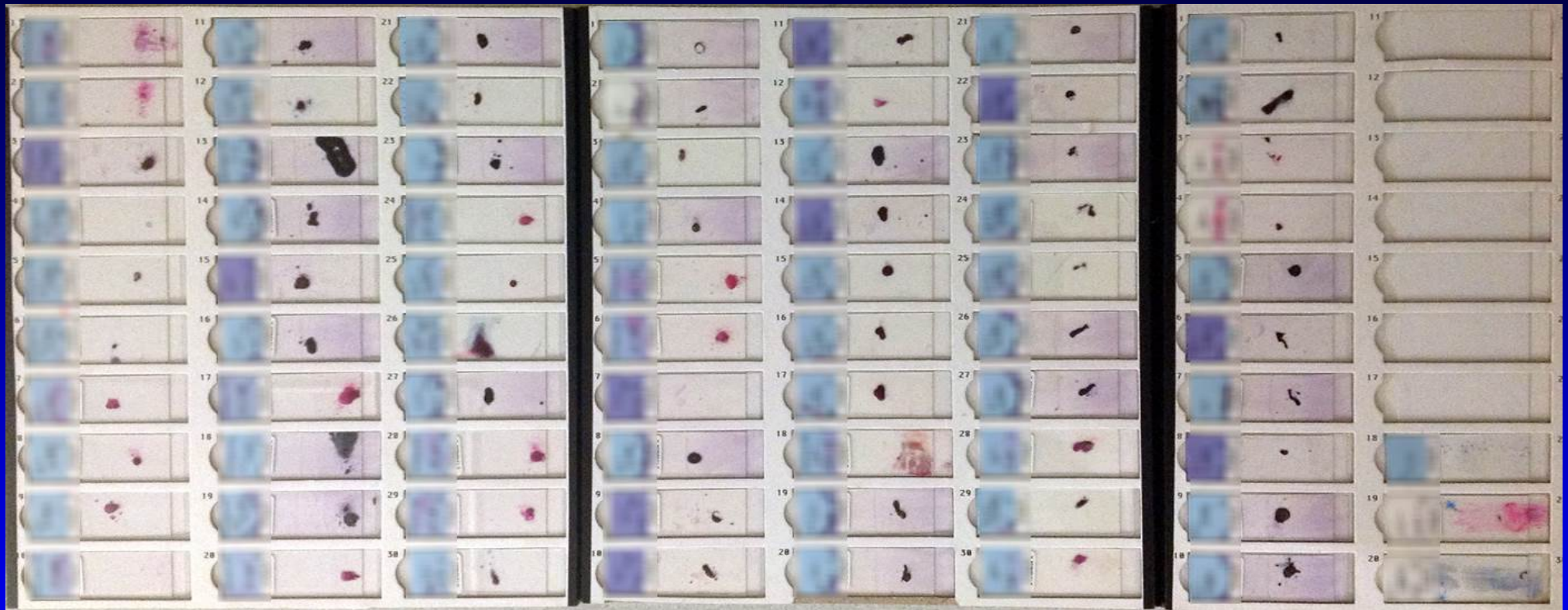
- Bronchoscopy suite contains a lab prep area & microscope
- ~3 min from main cyto lab
- Cytotech performs ROSE
  - Adequate sampling of LN
  - Malignant → ancillary studies
- Telepathology capability; not used on every case



# Performance ROSE

- **Senior Cytotechnologists**
- **Cytopathologists**
  - Direct viewing of slides
  - Telepathology
- **Trainees (fellows and residents)**

# What EBUS Can Look Like Without ROSE



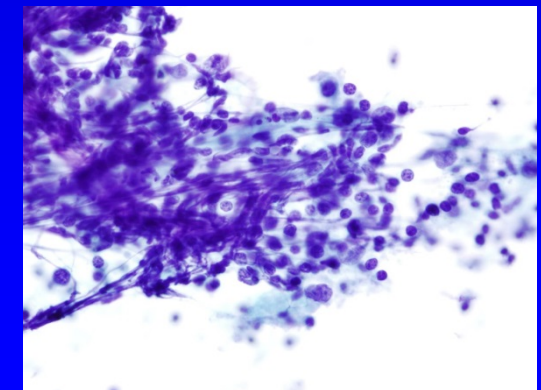
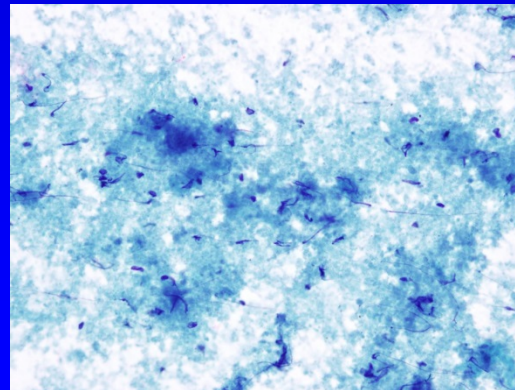
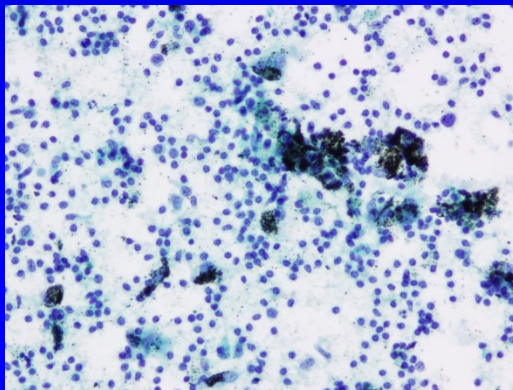
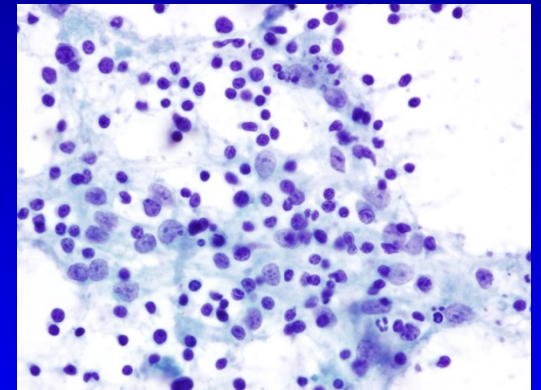
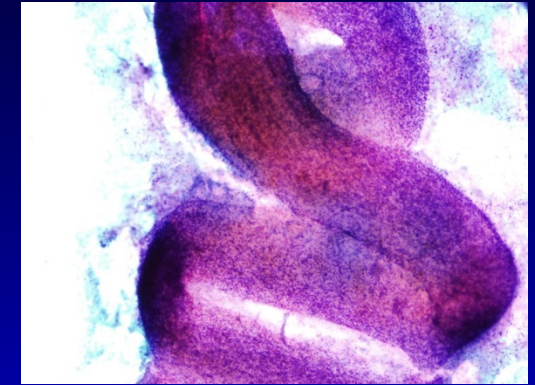
## EBUS, station 7, FNA

Received 72 direct smears and 1 cell block prepared from 6 needle passes. Passes #1-5 non-diagnostic. Pass #6 has some lymphocytes and two slides with few cells of carcinoma.



# Specimen Adequacy

- Number of lymphocytes/hpf
- Germinal center cells
- Anthracotic pigment-laden macrophages
- Extensive necrosis



# **Positive Cases on ROSE for Non-Small Cell Lung Carcinoma**

- **10 smears (5 DQ & 5 Pap)**
- **Extra Pass for CB**
- **1<sup>st</sup> & last H&E slide**
- **10 USS up front**
- **Not uncommon need more  
USS for MDL**

# ROSE Reimbursement

- Utilization Review and reimbursement of cytology services in EBUS-TBNA

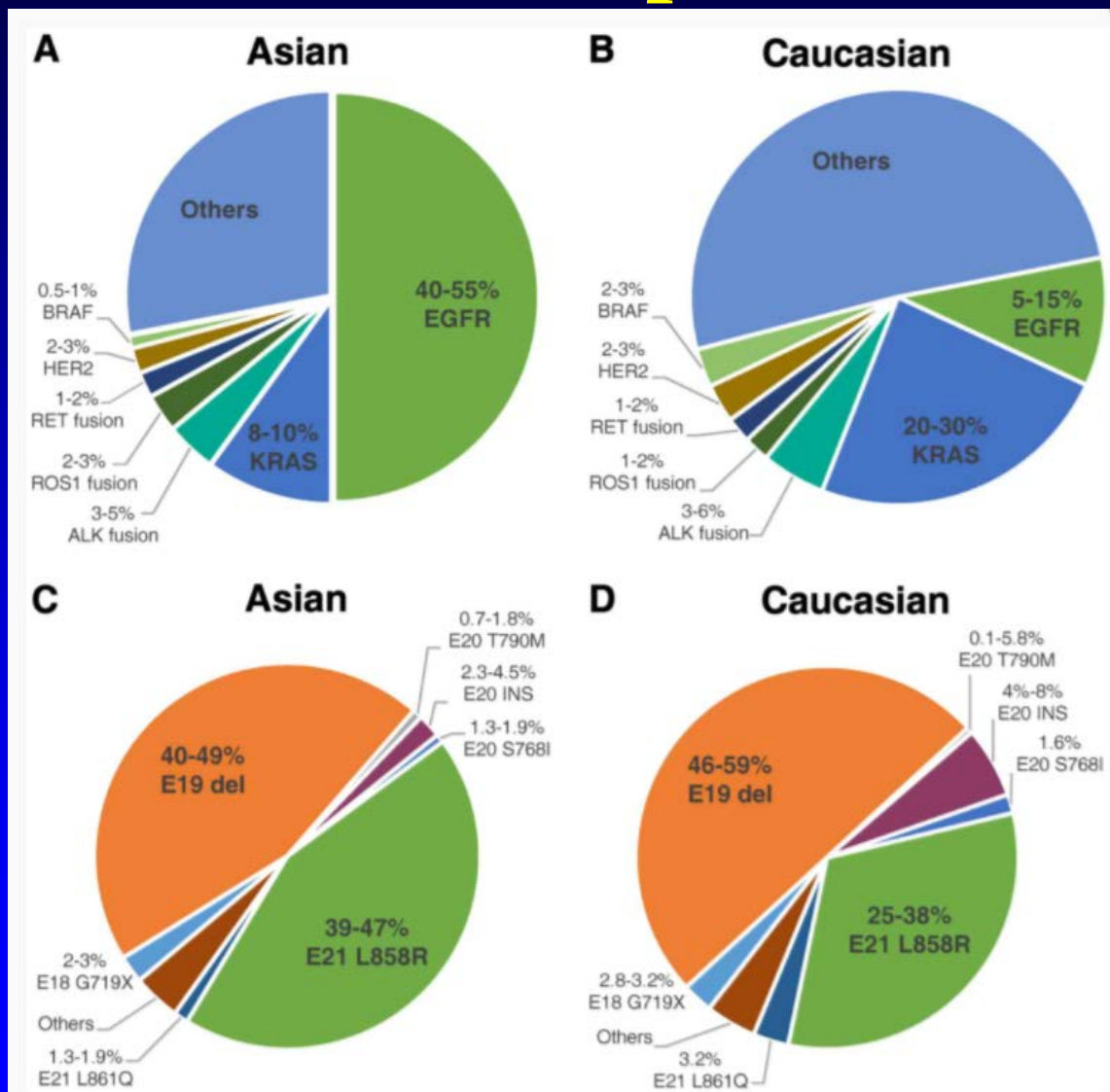
**Table 4** Cost calculation.

Financial considerations	Technical component	Professional component	Global component	Net gain /loss (range)
Reimbursement	\$46.73	\$113.97	\$160.70	
Cost (using telecytology)	\$109.00	\$55.71	\$164.71	-\$4.01
Cost (non-telecytology)	\$109.00	\$101.45	\$210.45	-\$49.75
Cost (overall average)	\$109.00	\$85.60	\$194.60	-\$33.90

# Triaging Specimens for Non-Small Cell Lung Carcinoma

- **Diagnosis**
  - **Adenocarcinoma vs. Squamous cell carcinoma**  
**Limited immunopanel (TTF-1 and p40); save material for mutational analysis (MDL)**
  - **1-2% of NSCLC cannot be classified as adenocarcinoma or squamous carcinoma; MDL may be helpful**

# Mutation Frequencies in Asian and Caucasian Populations



# Targeted Therapy

<b>DRUG</b>	<b>TARGET</b>	<b>INDICATION</b>
<b>Erbtinib (Tarceva) Afatinib (Gilotrif) Gefitinib (Iressa)</b>	<b>EGFR</b>	<b>NSCLC (EGFR exon 19 del or 21 substitution)</b>
<b>Osimertinib (Tagrisso)</b>	<b>EGFR T790M</b>	<b>NSCLC with T790M mutation</b>
<b>Necitumumab (Portazza)</b>	<b>EGFR</b>	<b>Squamous cell ca</b>
<b>Ceritinib (Zykadia) Alectinib (Alecensa) Brigatinib (Alunbrig)</b>	<b>ALK</b>	<b>NSCLC with ALK fusion</b>
<b>Crizotinib (Xalkori)</b>	<b>ALK, ROSI</b>	<b>NSCLC with ALK fusion or ROSI gene alteration</b>
<b>Debrafenib (Tafinlar)</b>	<b>BRAF</b>	<b>NSCLC with BRAF V600E mutation</b>
<b>Tranmetinib (Mekinist)</b>	<b>MEK</b>	<b>NSCLC with BRAF V600E mutation</b>
<b>Pembrolizumab (Keytruda)</b>	<b>PD-L1</b>	<b>NSCLC</b>

# PD-L1

- An immuno-peroxidase stain for PD-L1, 22C3 AB, performed on tumor cells within the cell block preparation, yielded the following result: membranous staining in \_\_\_\_\_% of cells (\_\_\_\_+ staining intensity).



Clinical Trials:  Newly diagnosed  Previously treated

### LIVING LONGER IS POSSIBLE

If your tumor has high levels of **PD-L1**, KEYTRUDA is proven to help patients live longer compared to chemotherapy and could be your first treatment option

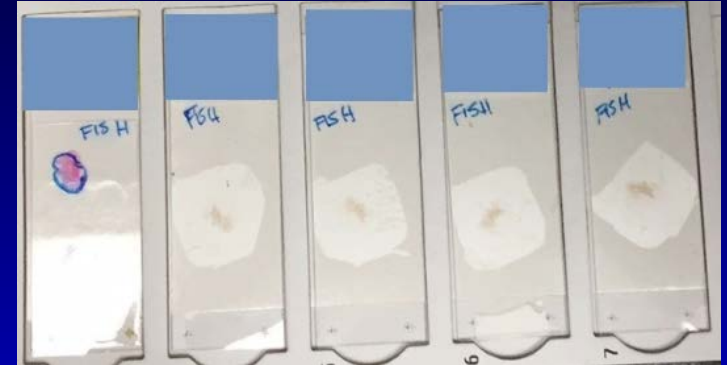
A clinical trial compared patients with advanced non–small cell lung cancer who received KEYTRUDA with those who received chemotherapy containing platinum. All patients in the trial had no previous drug treatment, tested positive for biomarker PD-L1 at a level of 50% or more, and tested negative for an abnormal EGFR or ALK gene. 154 patients received 200 mg of KEYTRUDA every 3 weeks, and 151 patients received chemotherapy.

### HELPED PATIENTS LIVE LONGER

More patients treated with KEYTRUDA were alive at the time of follow-up compared to patients treated with chemotherapy. 29% (44 of 154 patients) treated with KEYTRUDA were not alive at follow-up compared to 42% (64 of 151 patients) treated with chemotherapy.

# Triaging FNA for FISH Analysis

- **Most frequent FISH request**
  - ALK, RET, MET, ROSI
- **Need monolayer smear with tumor cells in the center of the slide**
- **Slide preparations for cytogenetics**
  - 4 USS & 1 H&E (tumor circled)
  - or DQ smears scored and marked (1-2 probes can be applied per slide)





# EX: Mutational Analysis Request at MDACC for Lung Adenocarcinoma

- **FISH Tests**

- ROS1
- RET
- CMET
- ALK

- **Immuno Tests**

- PD-L1
- MSI

- **MDL Tests**

- BRAF V600E
- EGFR Mutation
- KRAS Mutation
- RET Mutation
- EML4/ALK FUSIONS
- ROS1 FUSIONS

# What Tissue is Suitable for MDL?

## 2017 Statement

Recommendation: Pathologists may use either cell blocks or other cytologic preparations as suitable specimens for lung cancer biomarker molecular testing.

Expert consensus opinion: Laboratories should use, or have available at an external reference laboratory, clinical lung cancer biomarker molecular testing assays that are able to detect molecular alterations in specimens with as little as 20% cancer cells.

Strong recommendation: Laboratories should not use total EGFR expression by IHC testing to select patients for EGFR-targeted TKI therapy.

(Arch Pathol Lab Med. doi: 10.5858/arpa.2017-0388-CP)

Table 1 Tumor fraction and DNA yield from concurrently acquired fine needle aspiration and core needle biopsy samples

	<i>FNA smears</i> (n = 24)	<i>CNB</i> (n = 24)	<i>P-value</i>
<i>Tumor fraction</i>			
Mean	54%	39%	<i>P</i> = 0.003
Median	60%	30%	
Range	25–90%	20–70%	
<i>DNA yield</i>			
Mean	6.6 ng/μl	17.5 ng/μl	<i>P</i> = 0.01
Median	3.6 ng/μl	12.9 ng/μl	
Range	0.36–21 ng/μl	0.27–55 ng/μl	

Abbreviations: CNB, core needle biopsy; FNA, fine needle aspiration.

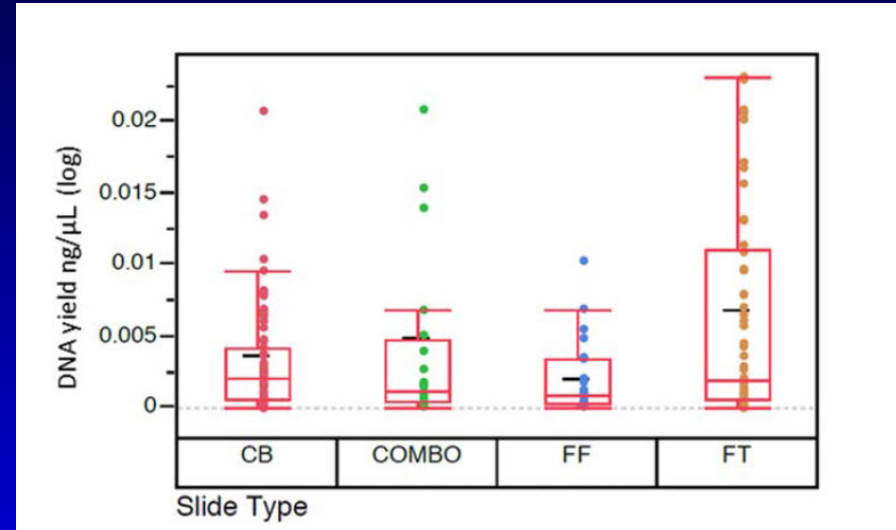
Roy et al.

# **Optimal Cytology Preparations for Mutational Analysis**

- **Validation studies needed for laboratory**
- **Types of preparations**
  - **Smears vs. liquid-base preparations**
  - **Papanicolaou vs. Diff-Quik smears**
  - **Role of cell block preparations**
  - **Combining different preparations**
  - **Other techniques (cell transfer/lift)**
  - **ProCore Bx??? (Acta Cytologica 2106;60:254-259)**

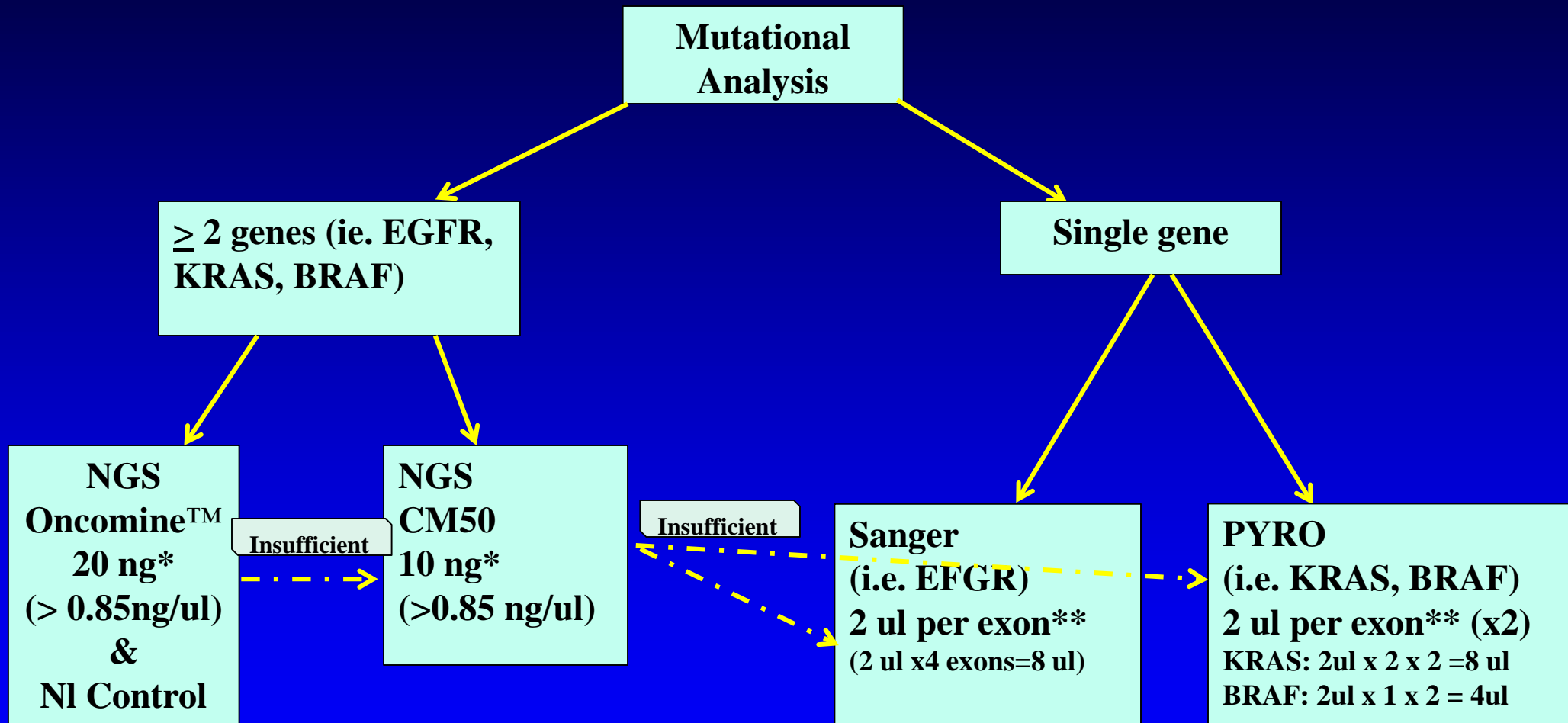
# Cytology Preparations vs. DNA Yield

- **Frosted tip slides had highest DNA yield over cell blocks, frosted slides and combination**
- **No experience with liquid base specimens**



Roy et al

# Mutation Testing Algorithm (MDL)



\*Recommended

\*\*Recommended (1000ng/ul) Almost never have that

Courtesy of Dr. Sinchita Roy-Chowdhuri

# How Much Material Do I Send for MDL???

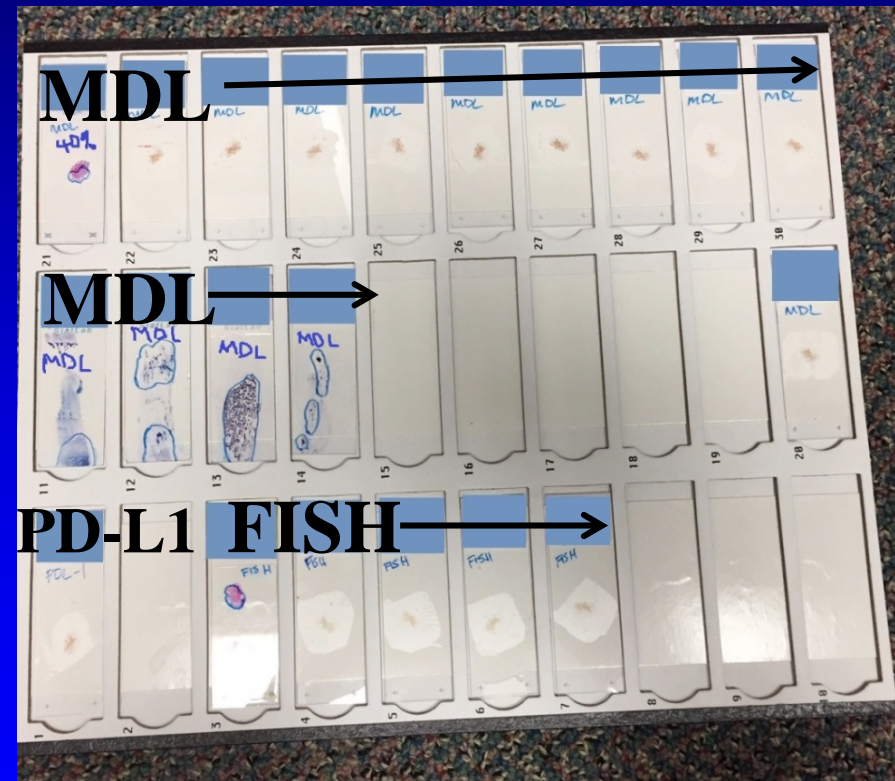
“As much as  
You can!!!”



- 1 cell ~ 6-7 pg of DNA
- For every 1 ng of DNA need ~143-166 intact cells
- NGS (Ion Torrent PGM) needs 10 ng, so rounding up ~ 2000 intact tumor cells
- Oncomine needs 20 ng, so 2X (~4000)
- Future ?????

# Triaging FNA for Mutational Analysis

- **MDL**
  - 10 USS & 1 H&E (tumor circled; % of tumor) plus 2 DQ & 2 PAP smears (tumor circled, estimate % of tumor present) submitted
- **FISH**
  - ALK, RET, MET, ROSI
  - 4 USS & 1 H&E (tumor circled) submitted to cytogenetics
- **Immunostains**
  - PD-L1
  - MSI (4 slides +)



# Triaging FNA for Mutational Analysis

- MDL requested on limited material
- Circle tumor; give tumor %
- Write a note to document slides sent
- Request slide to be returned with remaining tumor
- Designated areas circled scrapped (DQ & Pap-stained combined)

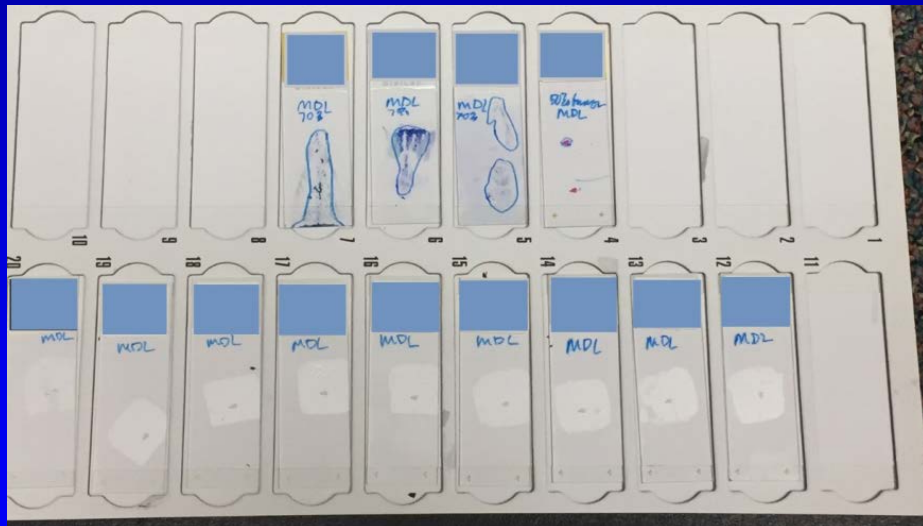
Diamond pen used to etch back of slide; marker to highlight the etched area after coverslip removed





# Mutational Analysis with CM50

- Smears and cell block from the same case can be combined for MDL



## Solid Tumor Genomics Assay v1

Clinical test requisition for mutation studies on the following genes was received: *BRAF*, *EGFR*, *KRAS*

A next generation sequencing (NGS)-based analysis for the detection of somatic mutations in the coding sequence of 128 genes and selected copy number variations (amplifications) in 49 genes (overlap: 134 genes total) was performed on the DNA extracted from the sample in our CLIA-certified molecular diagnostics laboratory. Interpretative findings are reported in the gene summary table(s) below followed by specific details of detected amplification and/or mutations.

### Interpretation Key:

- Circled/Bold: Mutation or amplification detected
- Underlined: Mutation testing requested (ordered gene)
- Asterisk: Additional confirmation studies in progress

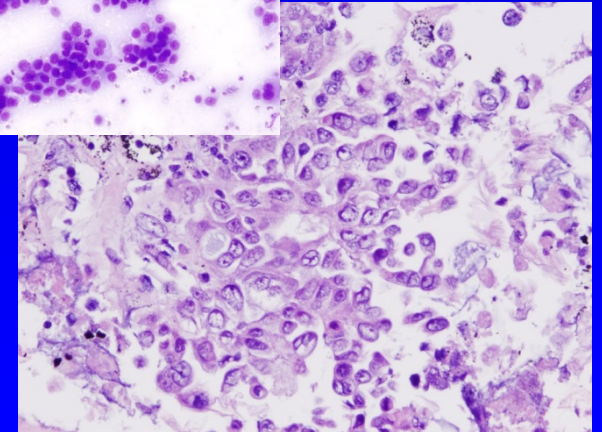
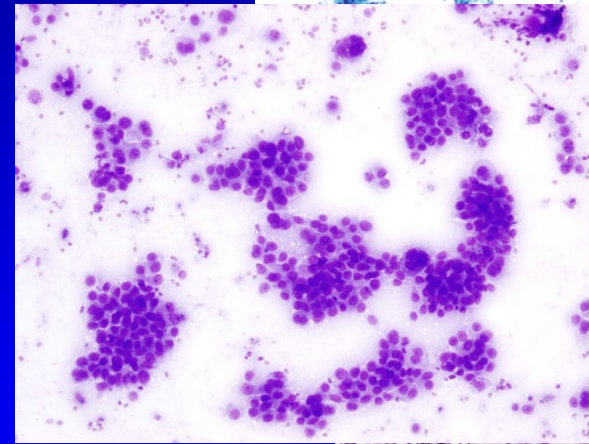
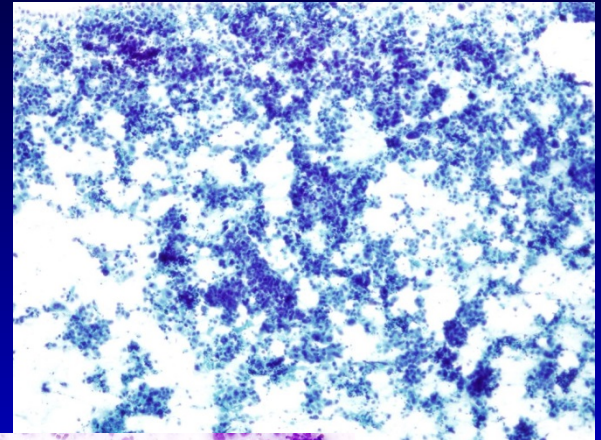
### GENE SUMMARY:

#### Molecular Diagnostics

<i>ABL1</i>	<i>CSF1R</i>	<i>FGFR2</i>	<i>IDH1</i>	<i>MLH1</i>	<i>PTPN11</i>	<b>TP53</b>
<i>AKT1</i>	<i>CTNNB1</i>	<i>FGFR3</i>	<i>IDH2</i>	<i>MPL</i>	<i>RB1</i>	<i>VHL</i>
<i>ALK</i>	<u><i>EGFR</i></u>	<i>FLT3</i>	<i>JAK2</i>	<i>NOTCH1</i>	<i>RET</i>	
<i>APC</i>	<i>ERBB2</i>	<i>GNA11</i>	<i>JAK3</i>	<i>NPM1</i>	<i>SMAD4</i>	
<i>ATM</i>	<i>ERBB4</i>	<i>GNAQ</i>	<i>KDR</i>	<i>NRAS</i>	<i>SMARCB1</i>	
<u><i>BRAF</i></u>	<i>EZH2</i>	<i>GNAS</i>	<i>KIT</i>	<i>PDGFRA</i>	<i>SMO</i>	
<i>CDH1</i>	<i>FBXW7</i>	<i>HNF1A</i>	<u><i>KRAS</i></u>	<i>PIK3CA</i>	<i>SRC</i>	
<b>CDKN2A</b>	<i>FGFR1</i>	<i>HRAS</i>	<i>MET</i>	<i>PTEN</i>	<i>STK11</i>	

# Mutational Analysis with Oncomine

- Request: BRAF, EGFR, KRAS
- Submit Tumor & Normal control



# Mutational Analysis Performed on FNA specimen with Oncomine

## Solid Tumor Genomics Assay v1

Clinical test requisition for mutation studies on the following genes was received: *BRAF*, *EGFR*, *KRAS*

A next generation sequencing (NGS)-based analysis for the detection of somatic mutations in the coding sequence of 128 genes and selected copy number variations (amplifications) in 49 genes (overlap: 134 genes total) was performed on the DNA extracted from the sample in our CLIA-certified molecular diagnostics laboratory. Interpretative findings are reported in the gene summary table(s) below followed by specific details of detected amplification and/or mutations.

### Interpretation Key:

- Circled/Bold: Mutation or amplification detected
- Underlined: Mutation testing requested (ordered gene)
- Asterisk: Additional confirmation studies in progress

### GENE SUMMARY:

## Molecular Diagnostics

<i>ABL1</i>	<i>CCND1</i>	<i>EZH2</i>	<i>IGF1R</i>	<i>MPL</i>	<i>PIK3R1</i>	<i>SRC</i>
<i>ACVRL1</i>	<i>CCNE1</i>	<i>FBXW7</i>	<i>IL6</i>	<i>MSH2</i>	<i>PNP</i>	<i>STAT3</i>
<i>AKT1</i>	<i>CD274</i>	<i>FGFR1</i>	<i>JAK1</i>	<i>MTOR</i>	<i>PPARG</i>	<i>STK11</i>
<i>ALK</i>	<i>CD44</i>	<i>FGFR2</i>	<i>JAK2</i>	<i>MYC</i>	<b><i>PPP2R1A</i></b>	<i>TERT</i>
<i>APC</i>	<i>CDH1</i>	<i>FGFR3</i>	<i>JAK3</i>	<i>MYCL1</i>	<i>PTCH1</i>	<i>TET2</i>
<i>APEX1</i>	<i>CDK4</i>	<i>FGFR4</i>	<i>KDR</i>	<i>MYCN</i>	<i>PTEN</i>	<i>TIAF1</i>
<i>AR</i>	<i>CDK6</i>	<i>FLT3</i>	<i>KIT</i>	<i>MYD88</i>	<i>PTPN11</i>	<b><i>TP53</i></b>
<i>ARAF</i>	<i>CDKN2A</i>	<i>FOXL2</i>	<i>KNSTRN</i>	<i>MYO18A</i>	<i>RAC1</i>	<i>TSC1</i>
<i>ATM</i>	<i>CHEK2</i>	<i>GAS6</i>	<b><i>KRAS</i></b>	<i>NF1</i>	<i>RAF1</i>	<i>TSC2</i>
<i>ATP11B</i>	<i>CSF1R</i>	<i>GATA2</i>	<i>MAGOH</i>	<i>NF2</i>	<i>RB1</i>	<i>U2AF1</i>
<i>BAP1</i>	<i>CSNK2A1</i>	<i>GATA3</i>	<i>MAP2K1</i>	<i>NFE2L2</i>	<i>RET</i>	<i>VHL</i>
<i>BCL2L1</i>	<i>CTNNB1</i>	<i>GNA11</i>	<i>MAP2K2</i>	<i>NKX2-1</i>	<i>RHEB</i>	<i>WT1</i>
<i>BCL9</i>	<i>DCUN1D1</i>	<i>GNAQ</i>	<i>MAPK1</i>	<i>NKX2-8</i>	<i>RHOA</i>	<i>XPO1</i>
<i>BIRC2</i>	<b><i>DDR2</i></b>	<i>GNAS</i>	<i>MAX</i>	<i>NOTCH1</i>	<i>RPS6KB1</i>	<i>ZNF217</i>
<i>BIRC3</i>	<i>DNMT3A</i>	<i>HNF1A</i>	<i>MCL1</i>	<i>NPM1</i>	<i>SF3B1</i>	
<u><i>BRAF</i></u>	<u><i>EGFR</i></u>	<i>HRAS</i>	<i>MDM2</i>	<i>NRAS</i>	<i>SMAD4</i>	
<i>BRCA1</i>	<i>ERBB2</i>	<i>IDH1</i>	<i>MDM4</i>	<i>PAX5</i>	<i>SMARCB1</i>	
<i>BRCA2</i>	<i>ERBB3</i>	<i>IDH2</i>	<i>MED12</i>	<i>PDCD1LG2</i>	<i>SMO</i>	
<i>BTK</i>	<i>ERBB4</i>	<i>IFITM1</i>	<i>MET</i>	<i>PDGFRA</i>	<i>SOX2</i>	
<i>CBL</i>	<i>ESR1</i>	<i>IFITM3</i>	<i>MLH1</i>	<i>PIK3CA</i>	<i>SPOP</i>	

# Summary

- **EBUS-TBNA aids in staging patients with lung cancer**
- **EBUS-TBNA with ROSE can help ensure adequate material for diagnosis and molecular testing**
- **Triaging and optimizing material prior to mutational analysis is an important step**
- **Mutational analysis can be performed on cytology material**

**Thank You**

