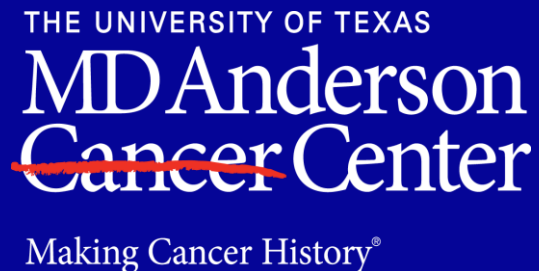


# Breast Cancer Session

W. Fraser Symmans, M.D.

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Director, Translational Research Program, the Alliance For Clinical Trials, NCI



# Disclosures

**Nuvera Biosciences, Inc.**  
**Delphi Diagnostics, Inc.**

**Co-founder and scientific advisor**  
**Co-founder and scientific advisor**

**IONS Pharmaceuticals**

**Stock**

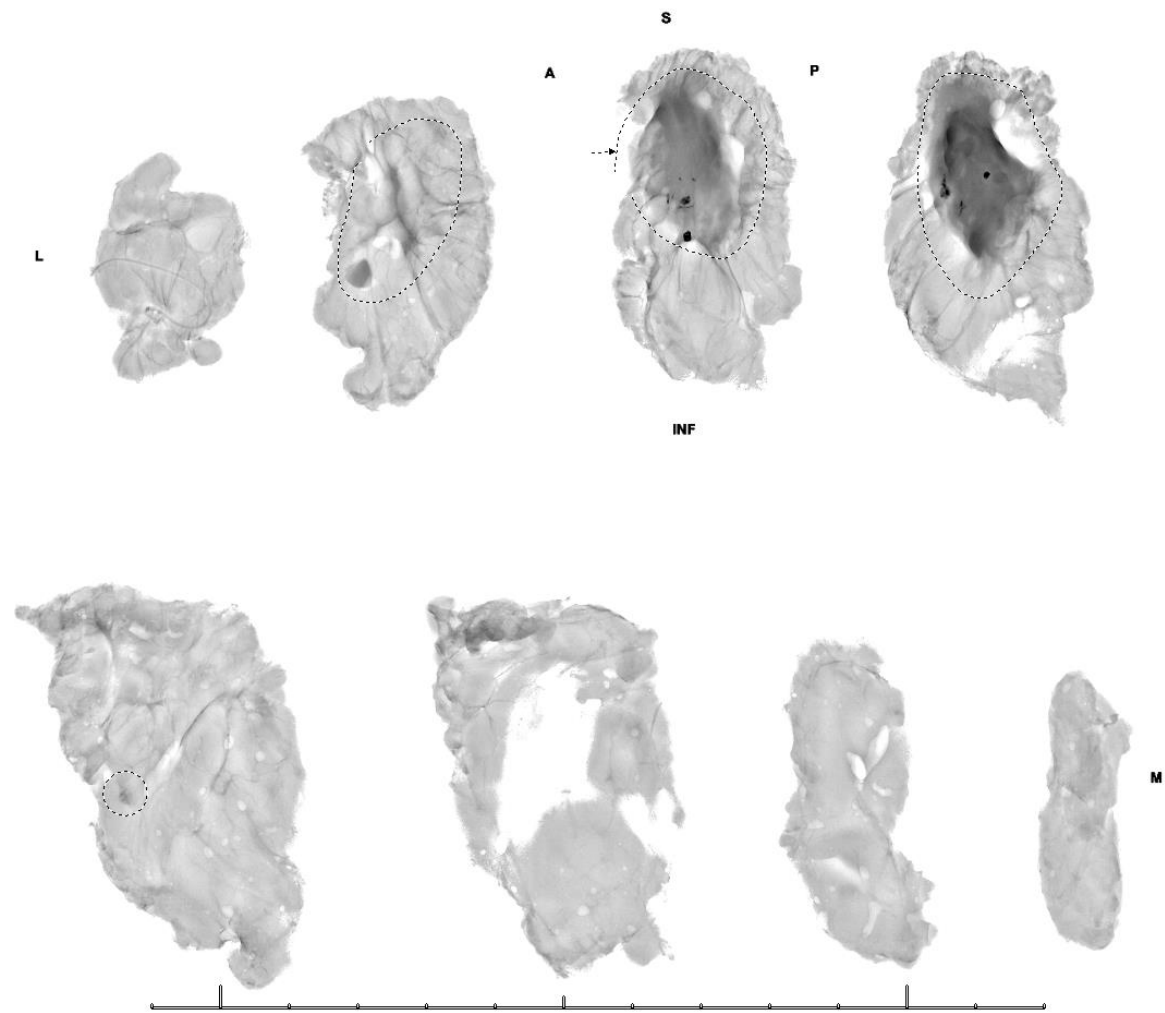
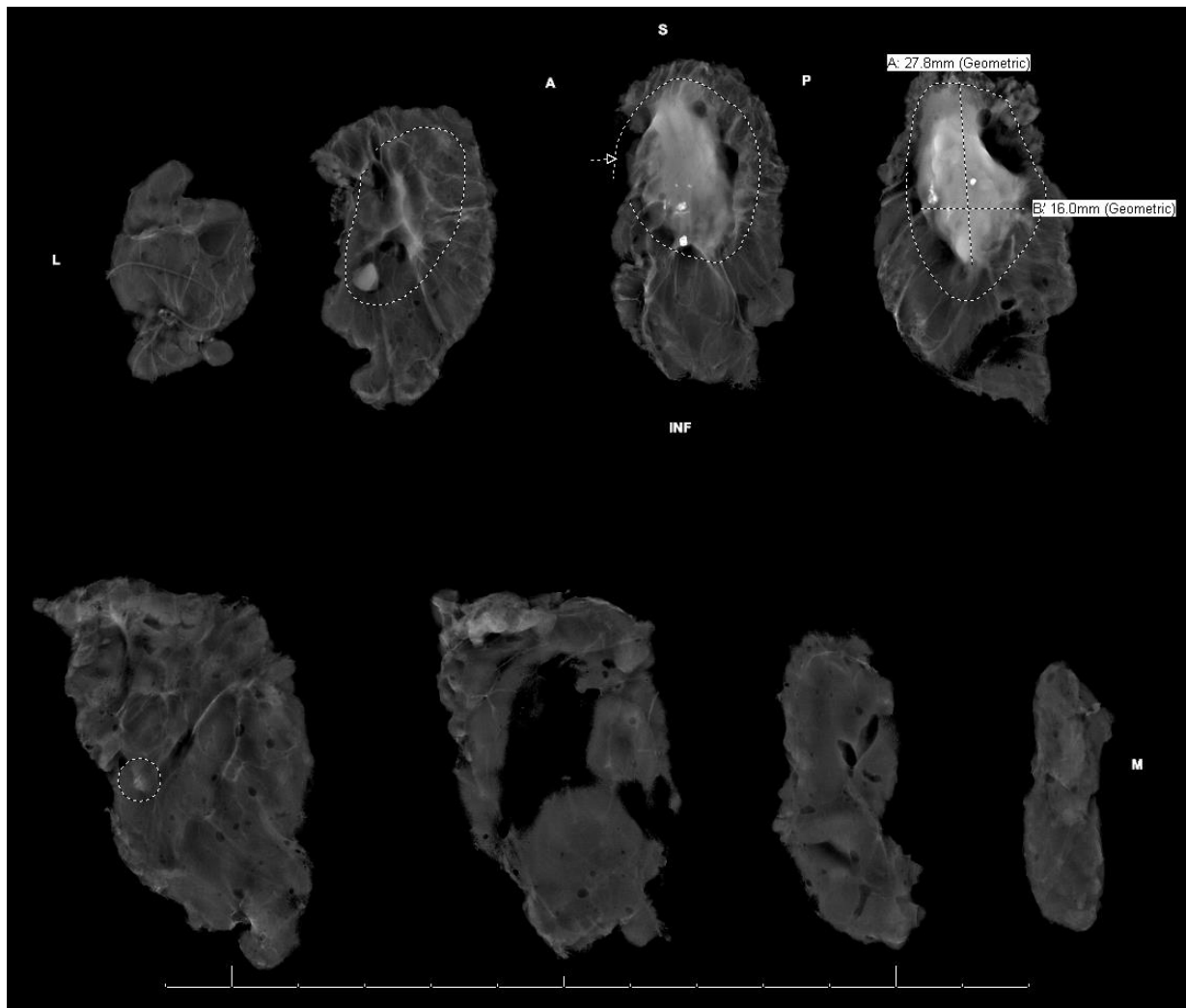
**Almac Diagnostics**  
**Merck**

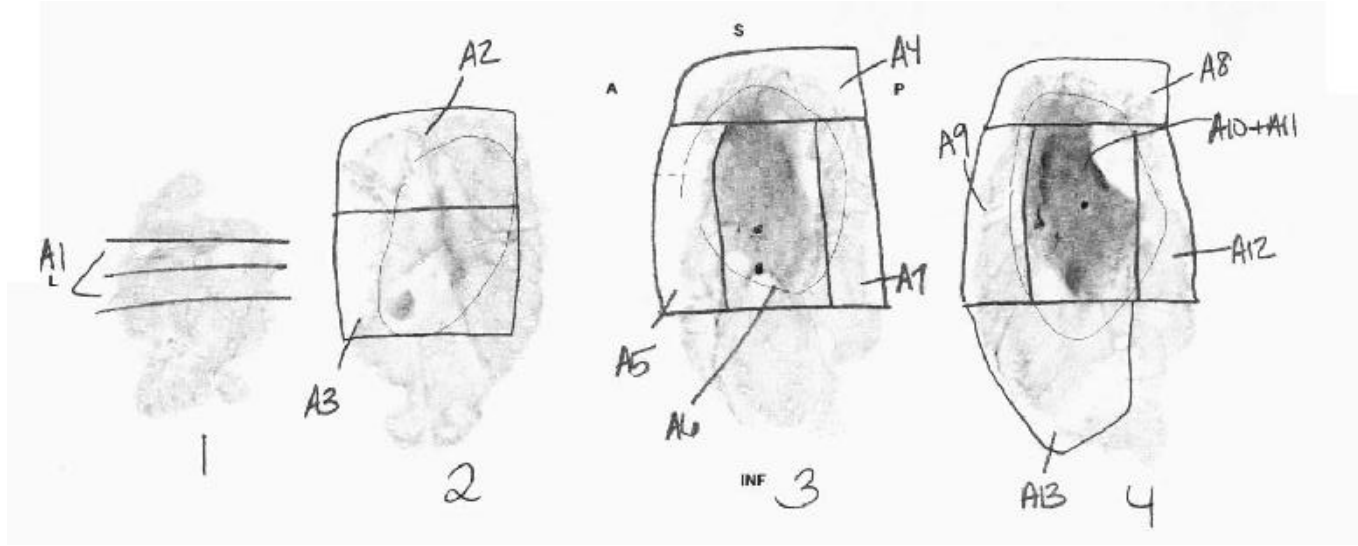
**Honorarium from advisory board**  
**Honorarium from advisory board**

**AbbVie**  
**Luminex**  
**Merck**

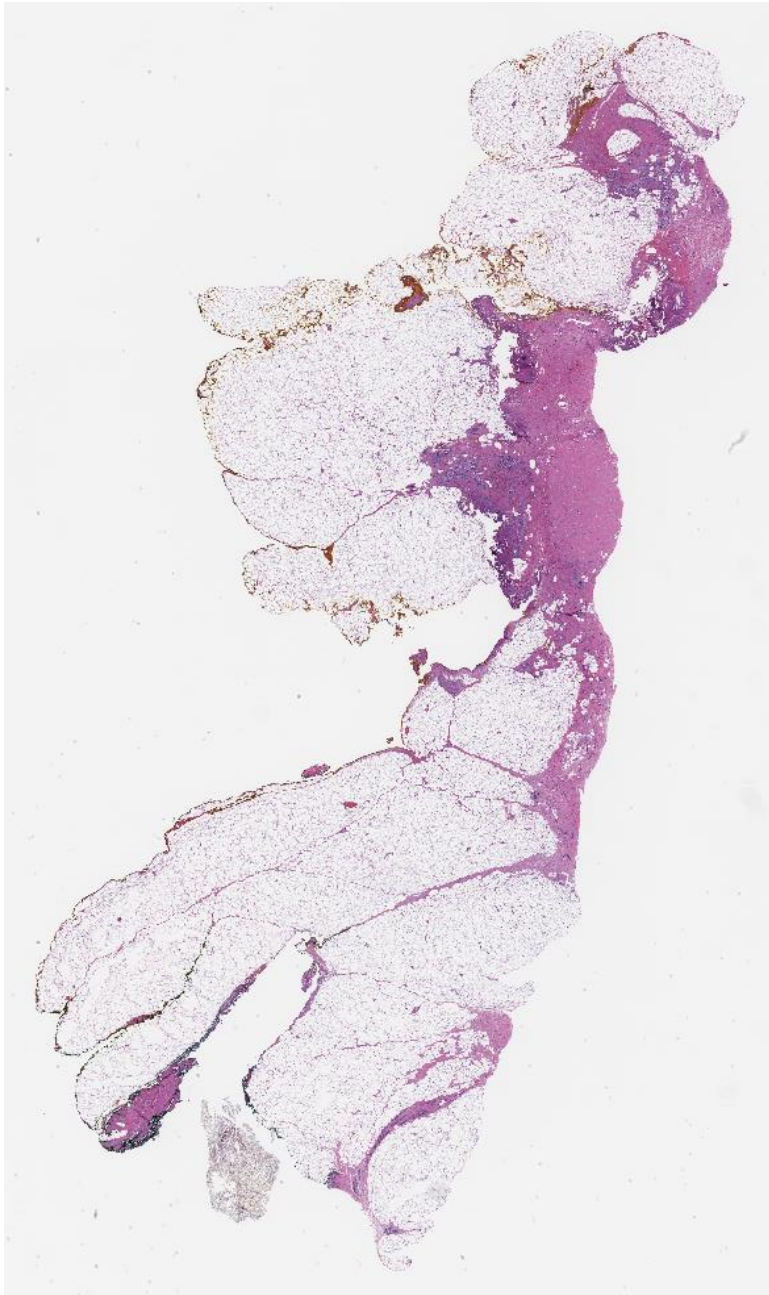
**Travel expenses to speak at meeting**  
**Travel expenses to speak at meeting**  
**Travel expenses to speak at meeting**

How we assess a lumpectomy breast resection specimen

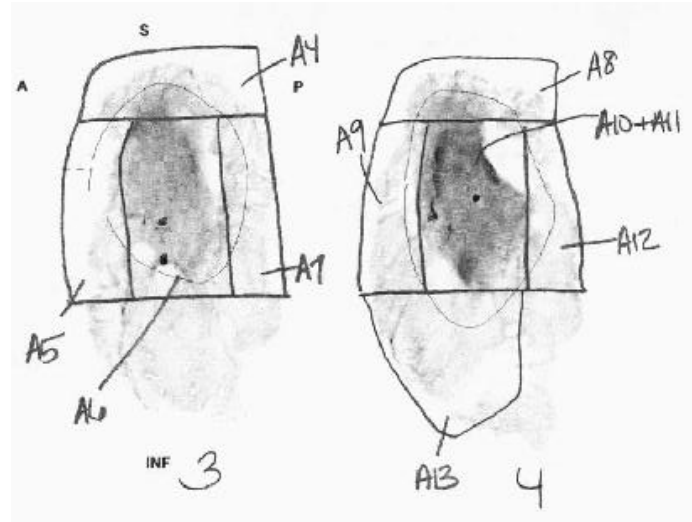
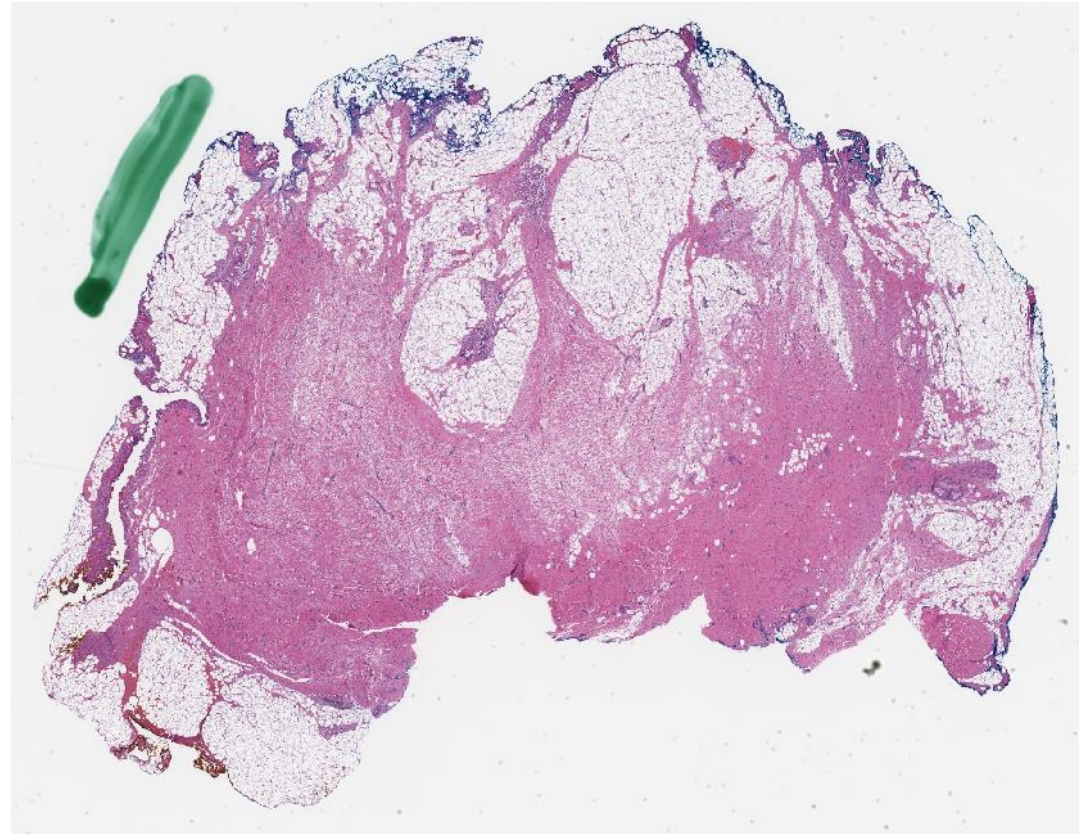




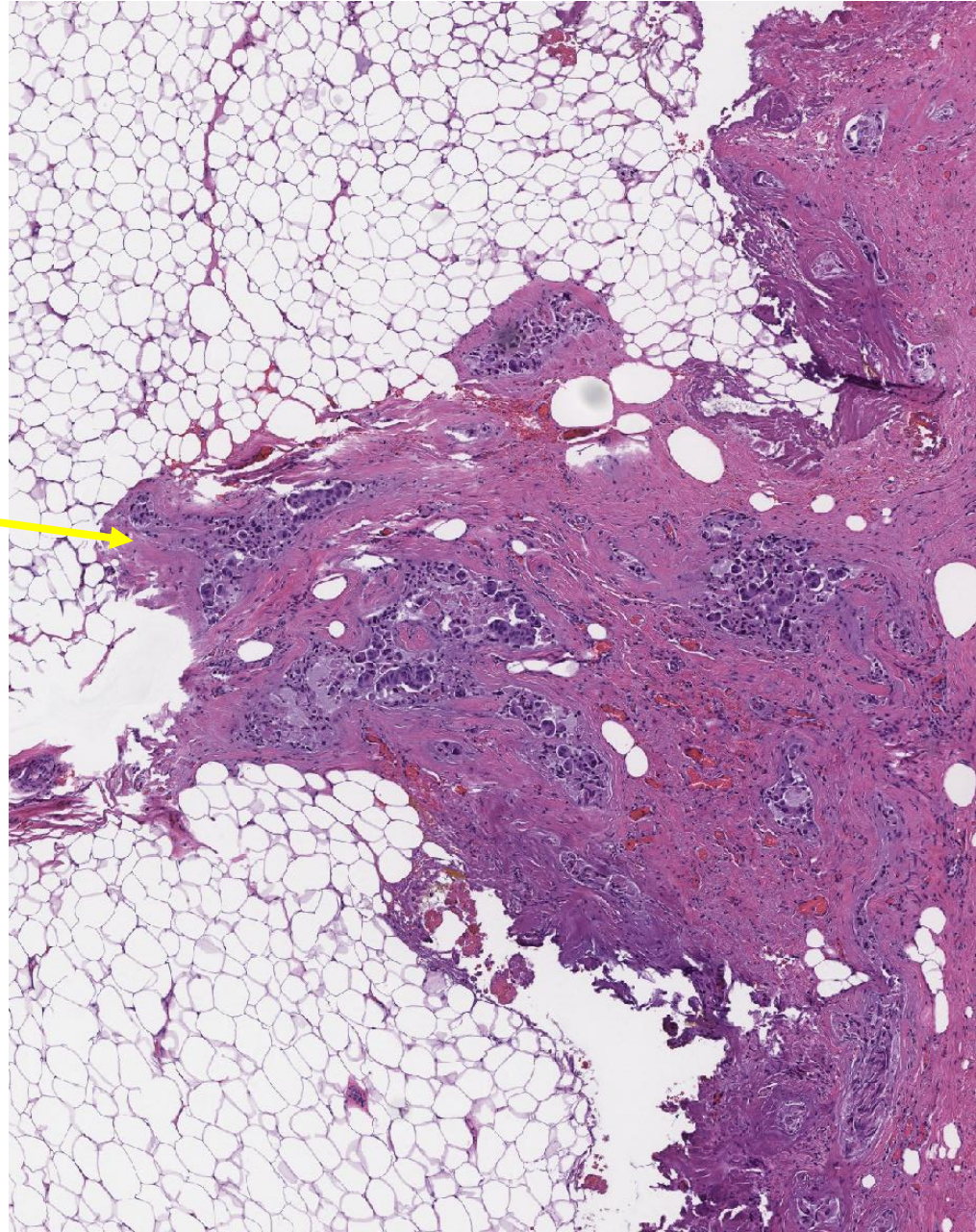
A5



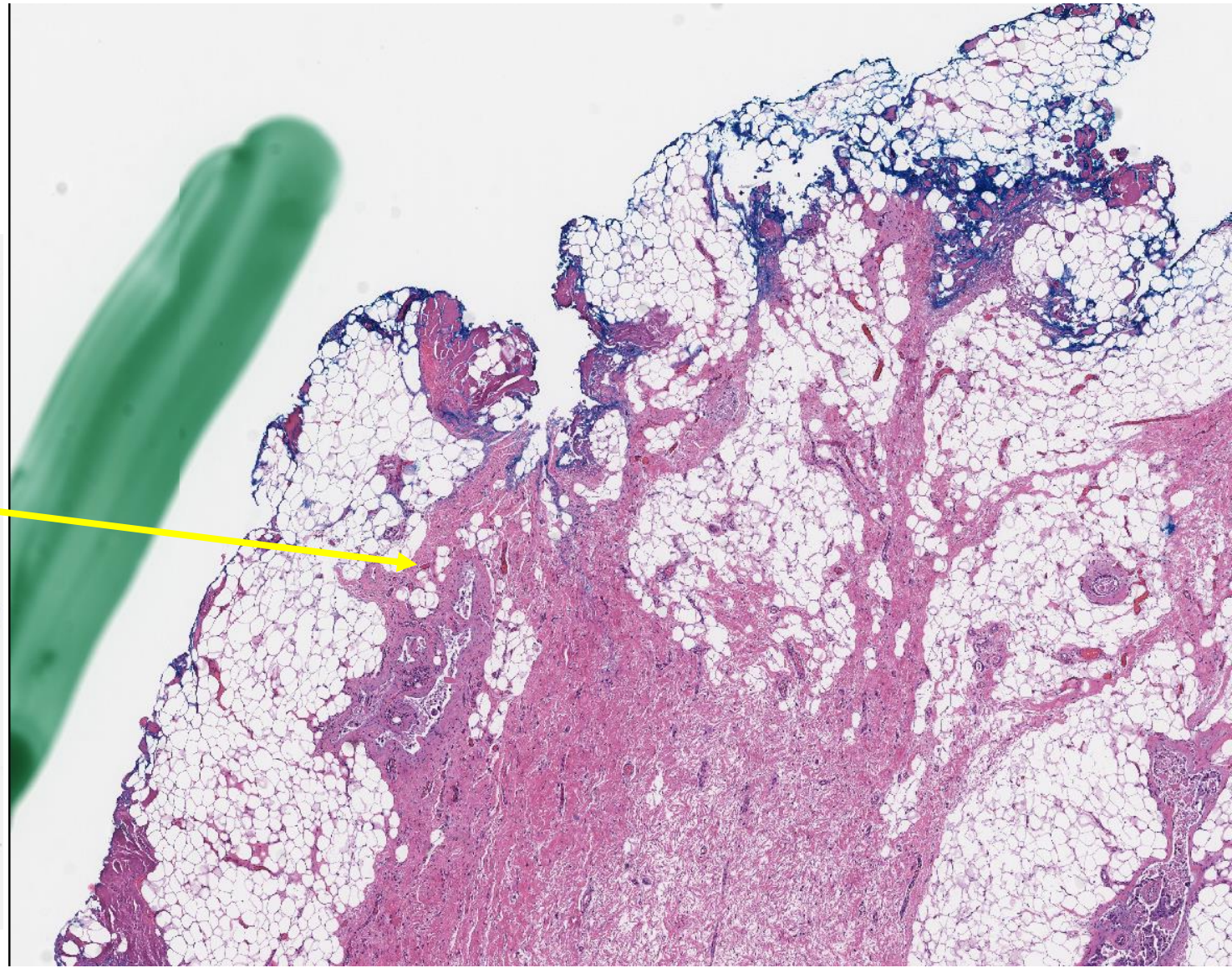
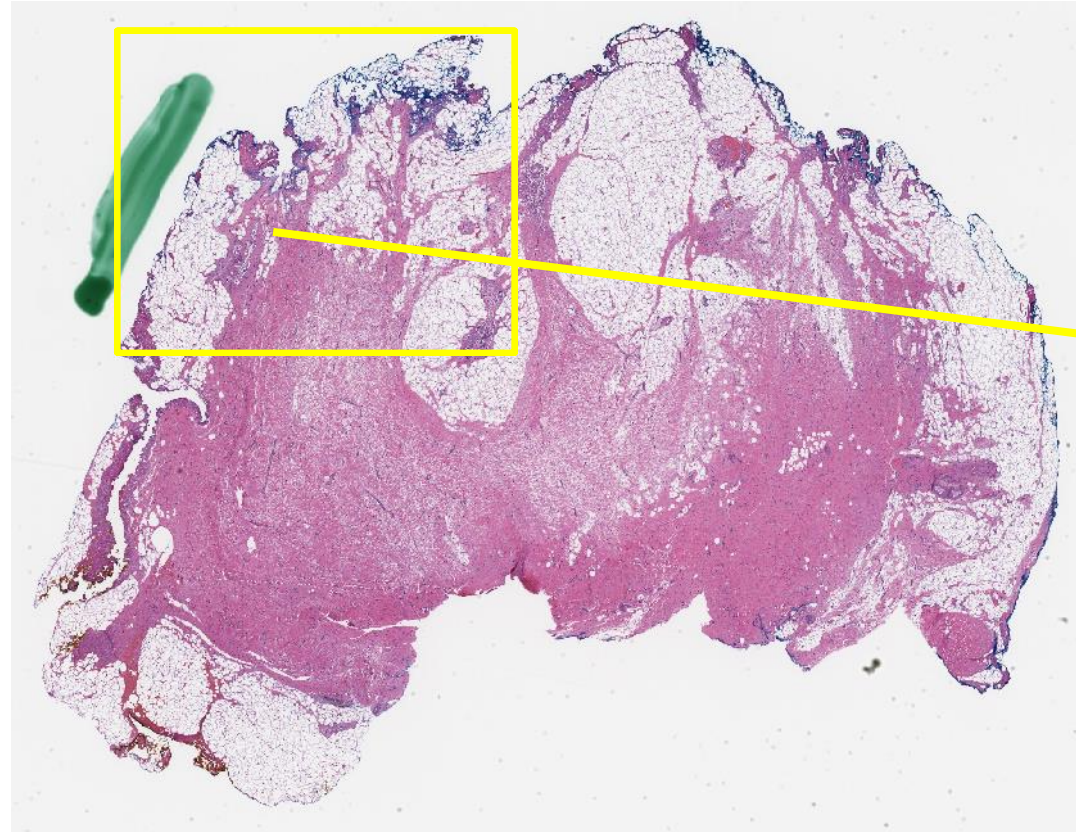
A8



A5



A8





How we assess a mastectomy specimen for a non-palpable disease

# Case: Pathologist's Preparatory Notes

49 year old with multifocal HER2+ left breast cancer (T2, N1)

## Imaging

- A) Tumor at 6 o'clock, 6 cm from nipple (biopsy + clip)
- B) Tumor at 5 o'clock (anterior), 1 cm from nipple (biopsy + clip)
- C) Intervening tumor 1.3 cm at 5 o'clock, 5 cm from nipple
- D) Tumor 0.6 cm at 5 o'clock, 1.7 cm lateral to B).

Ultrasound showed one 2 cm axillary node with enlarged cortex

## Pathology From Biopsy

IDC grade 3, HR+ / HER2- in tumors A) and B)

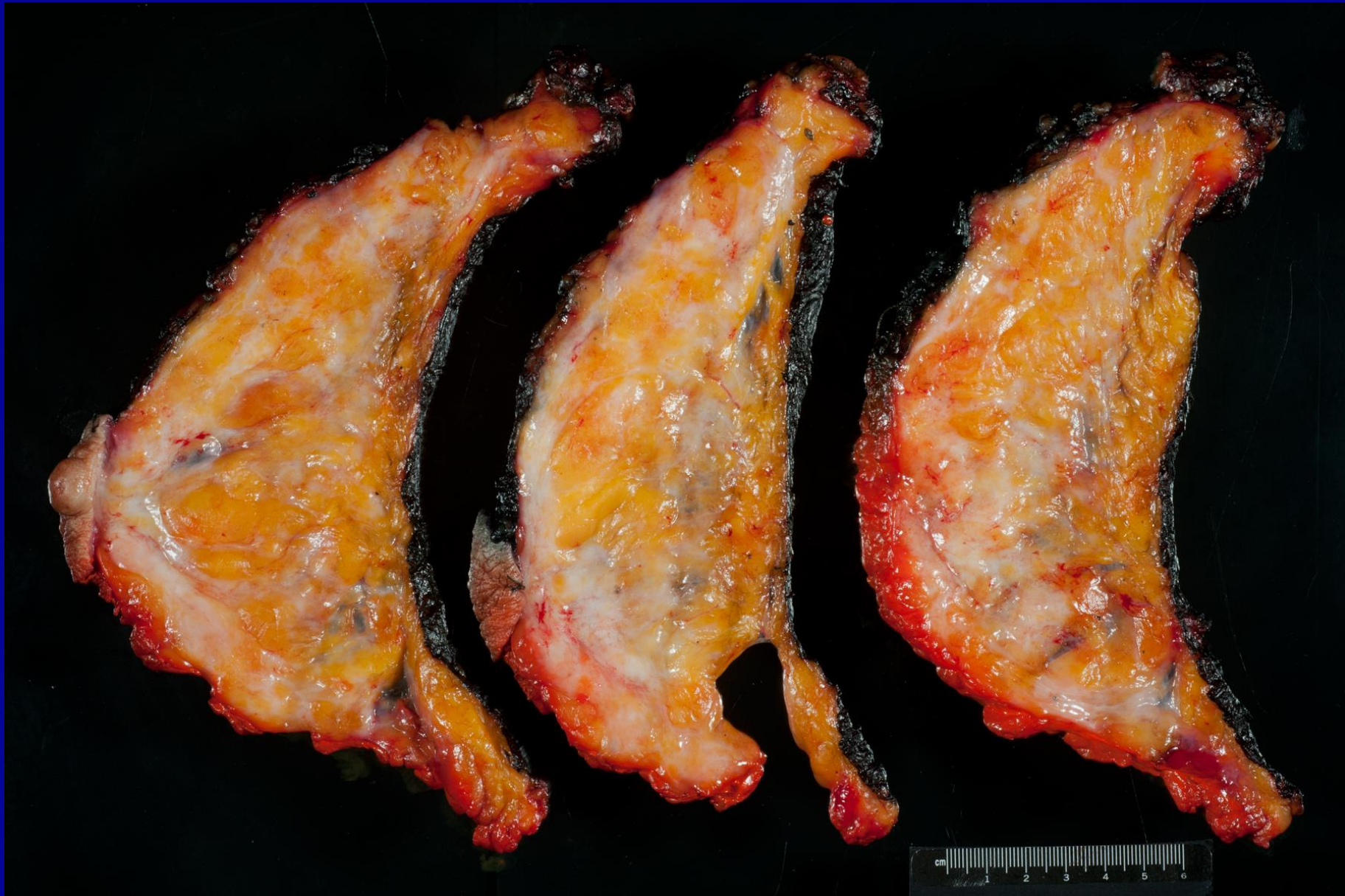
Metastatic breast cancer in LN, HR+, HER2+

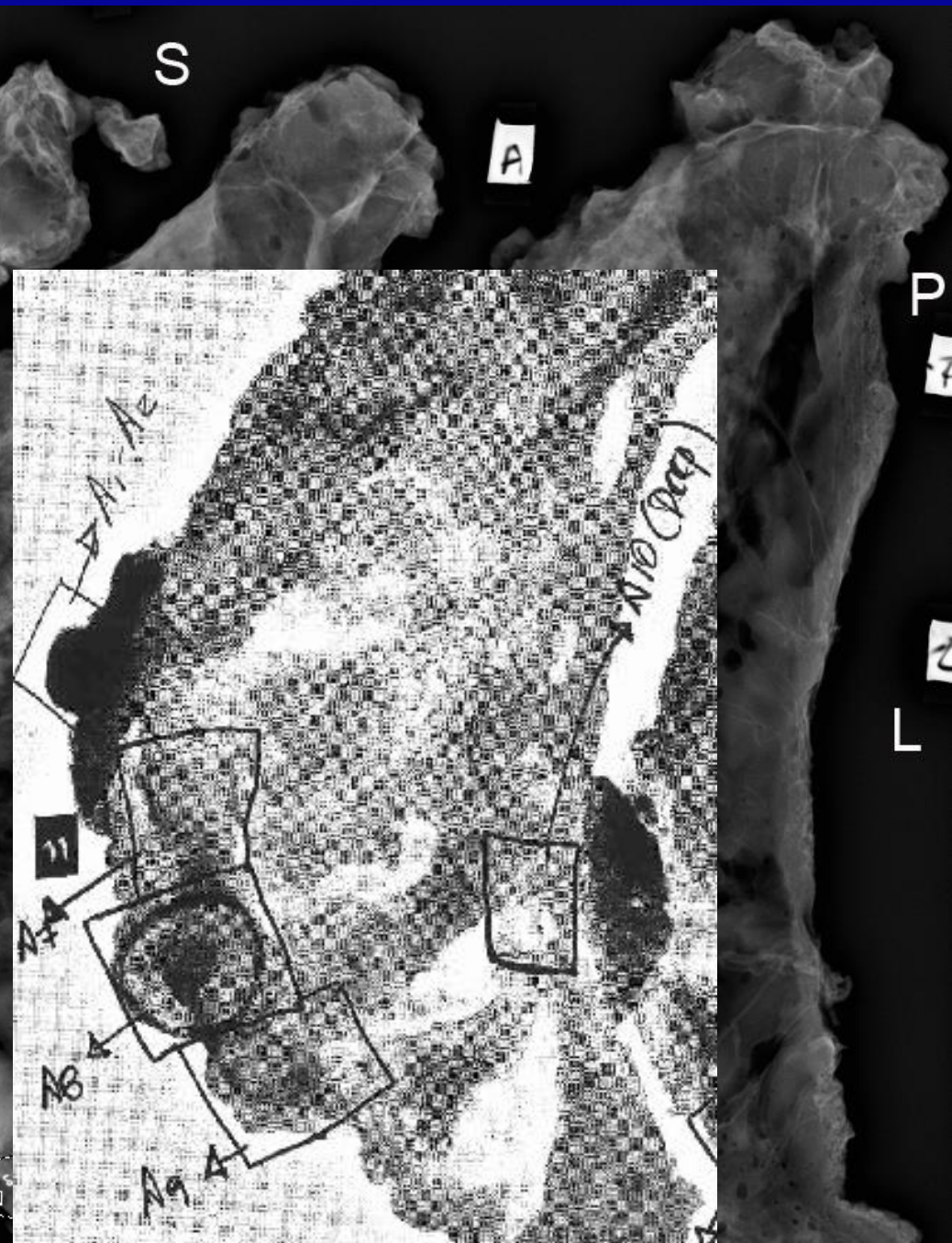
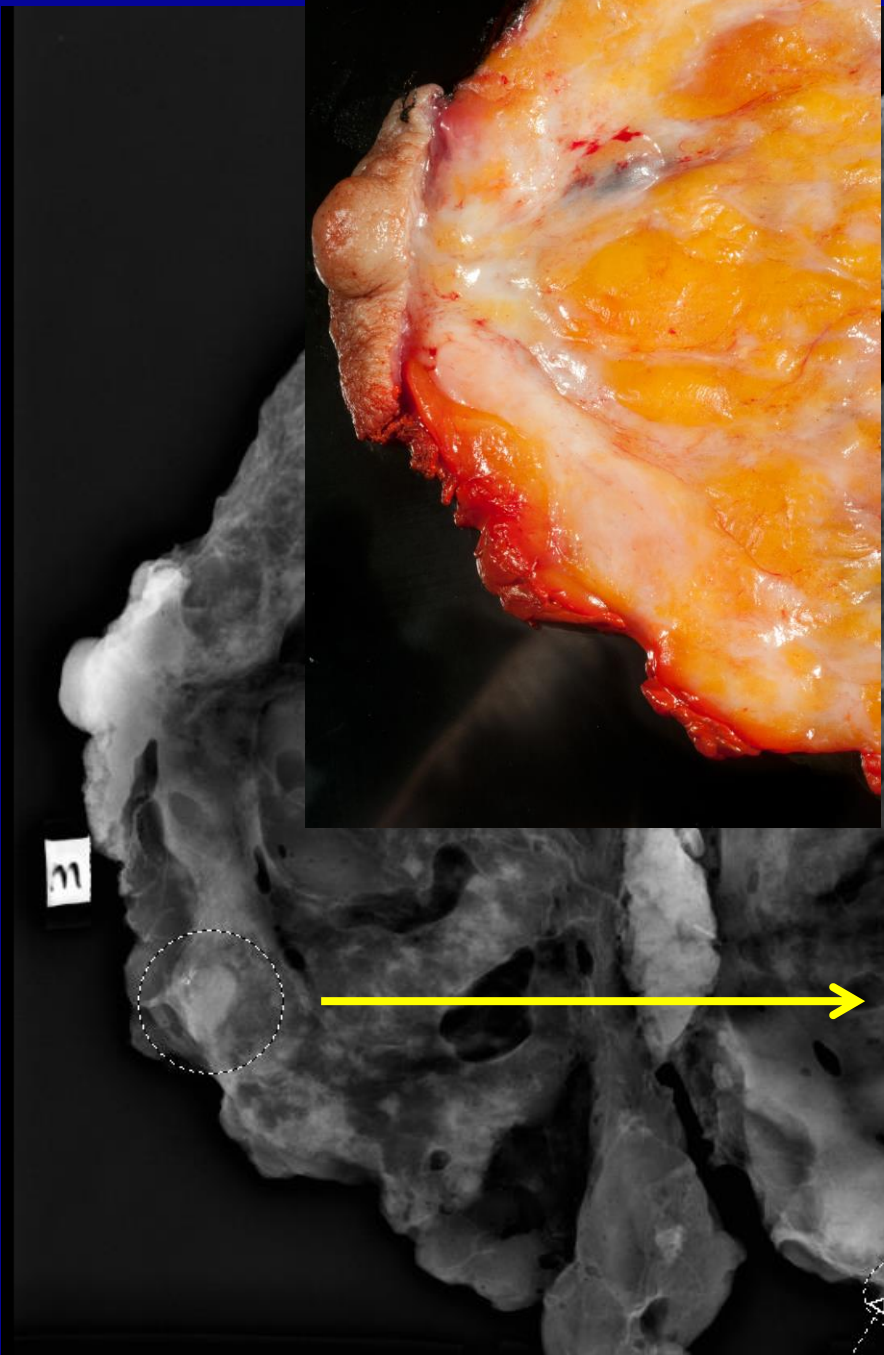
## Treatment & Response

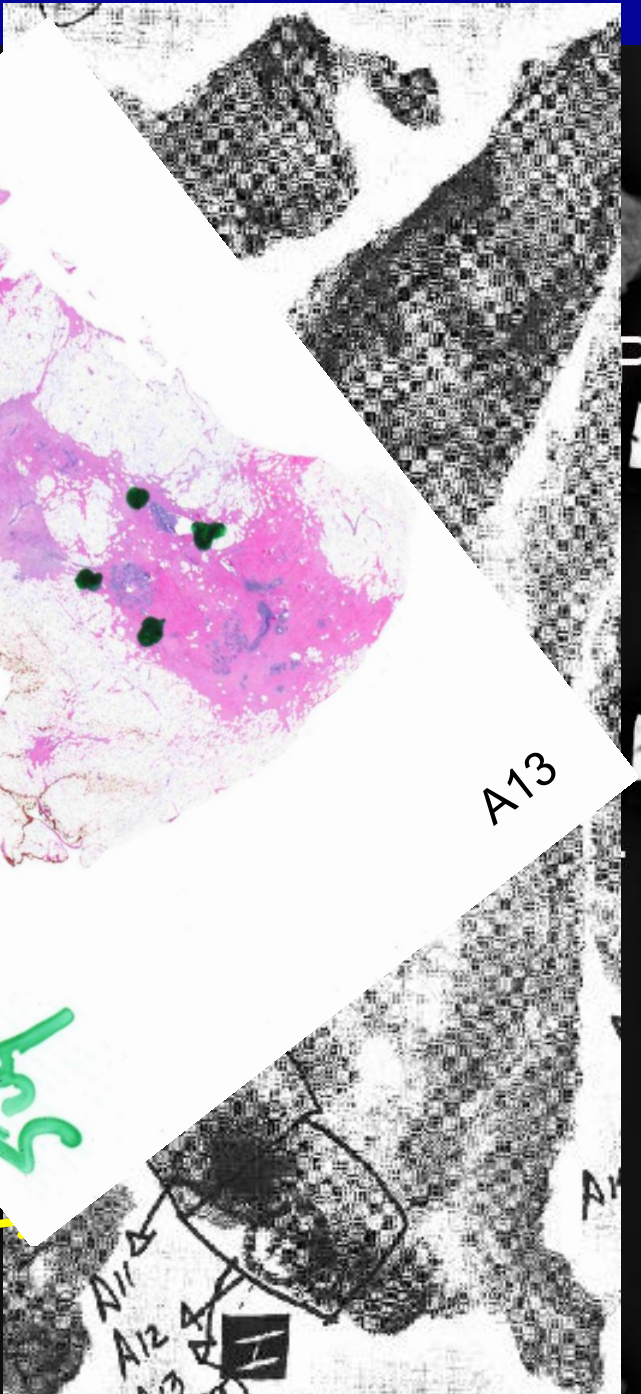
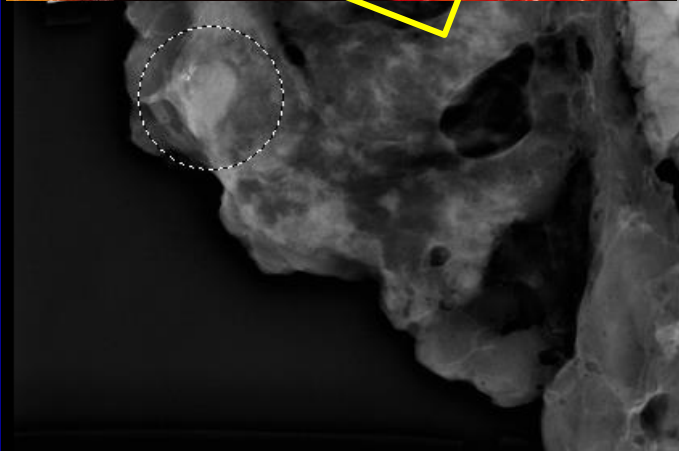
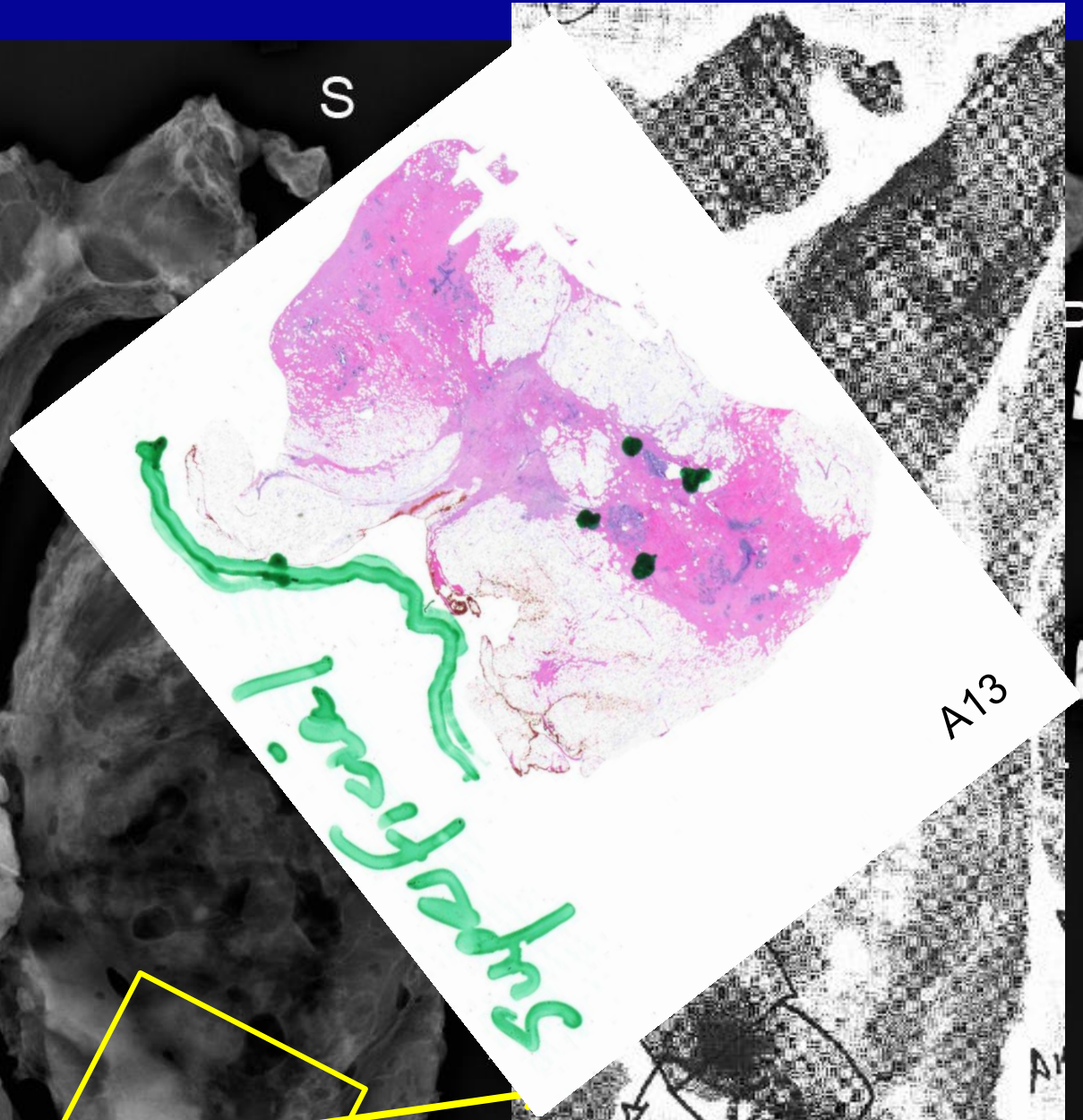
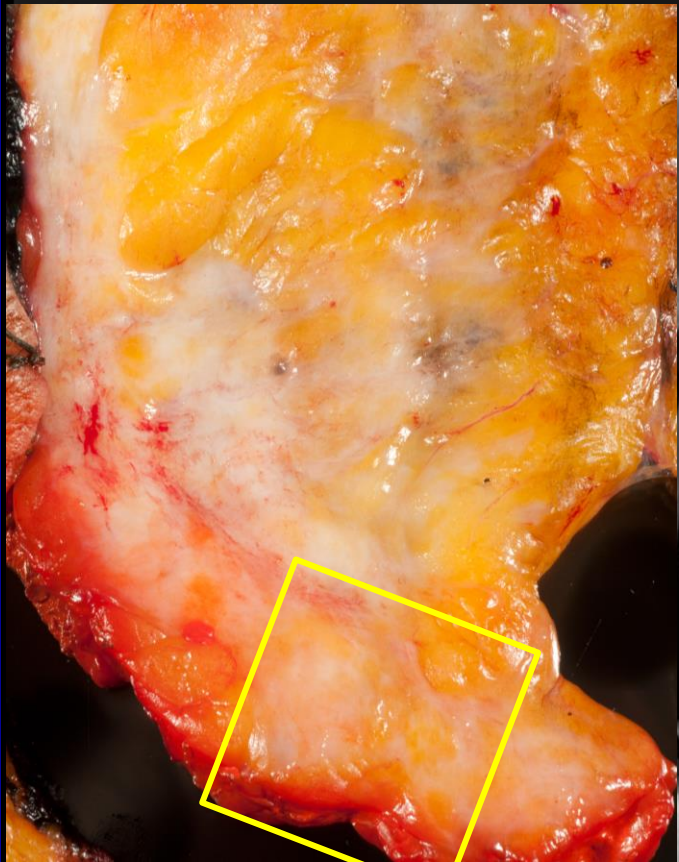
3-weekly FEC x 4 then weekly paclitaxel x 12 with concurrent trastuzumab throughout (FEC-H/T-H)

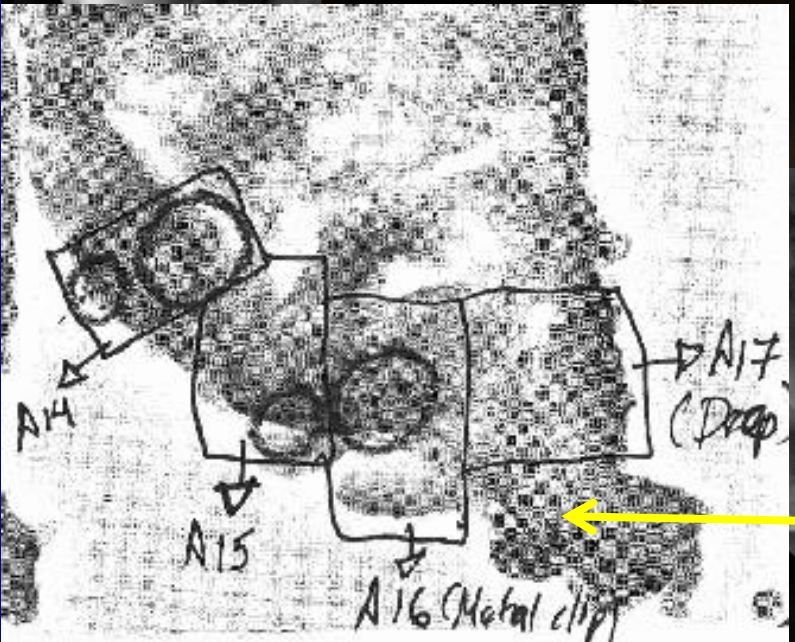
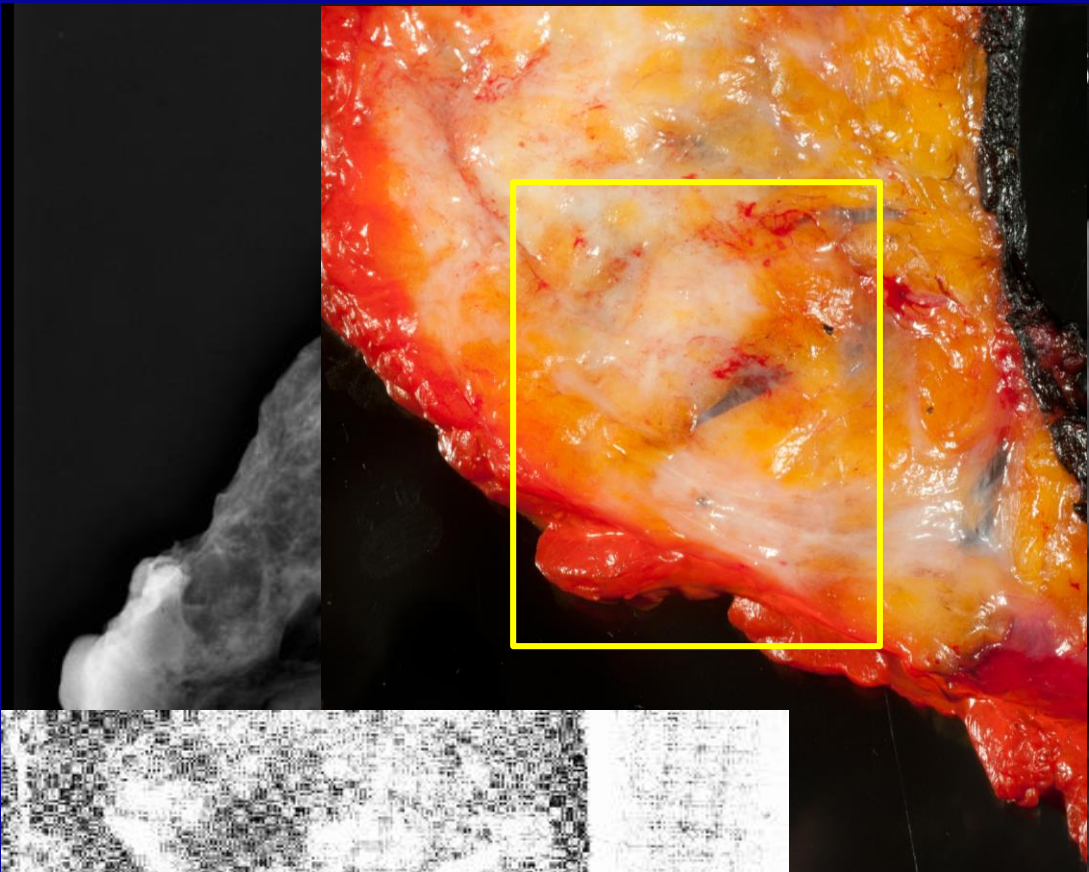
Pre-surgical MRI = complete response

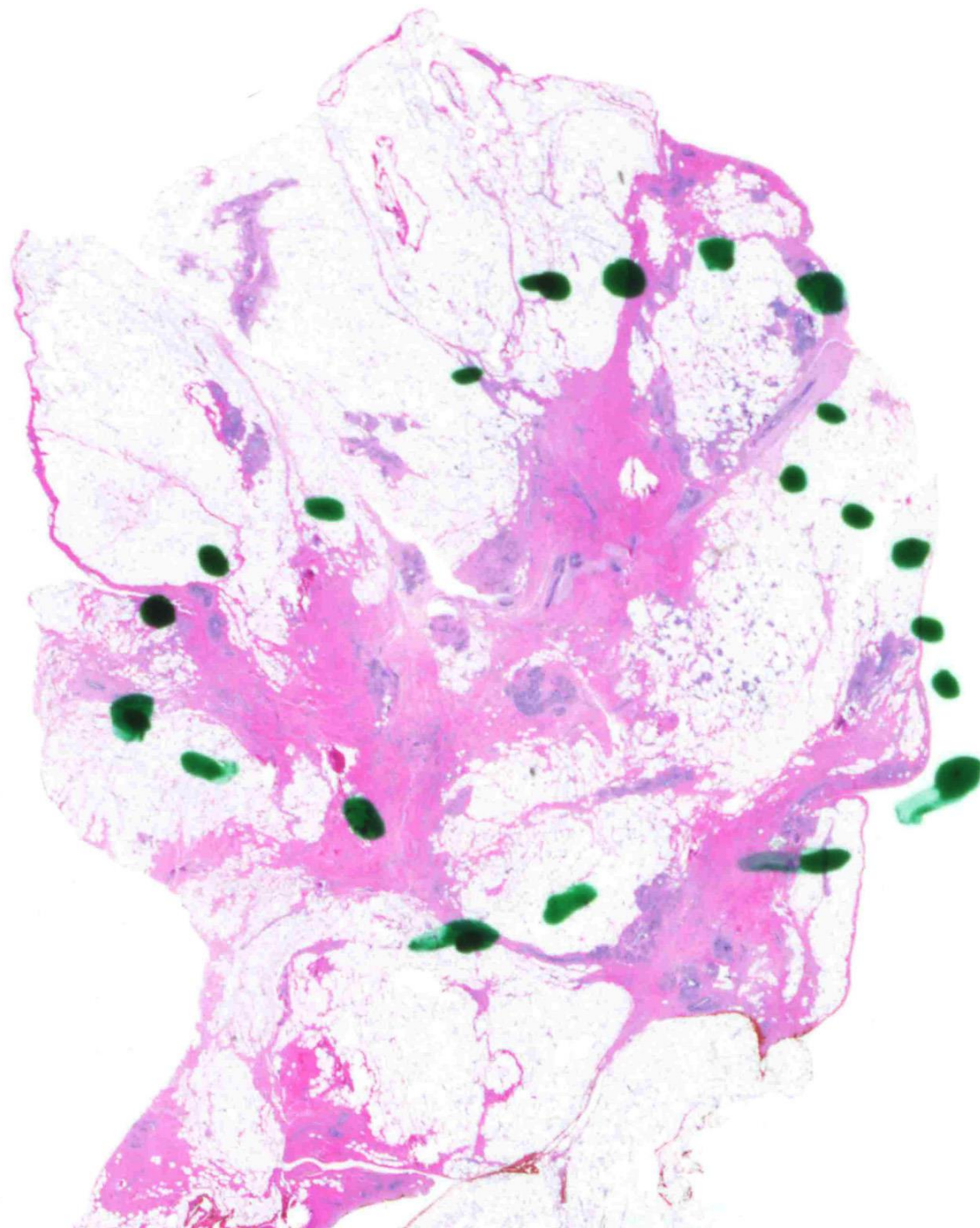
For skin-sparing mastectomy and axillary node dissection

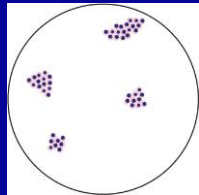
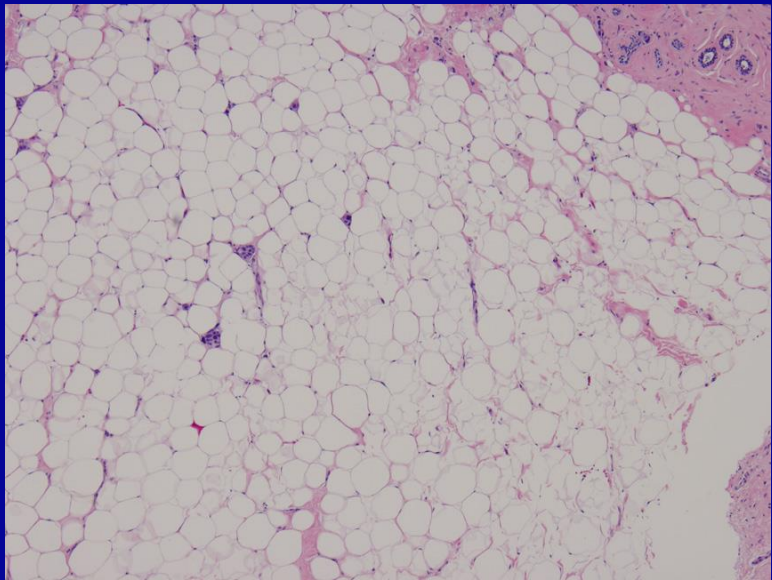
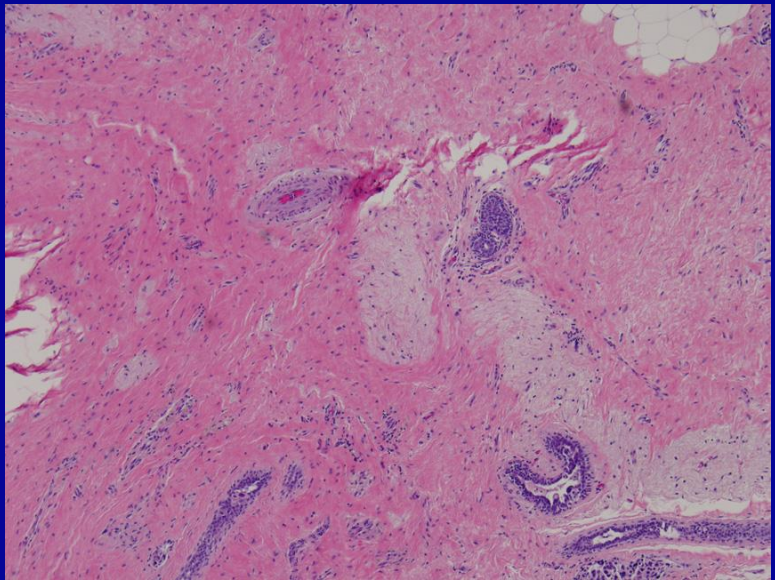




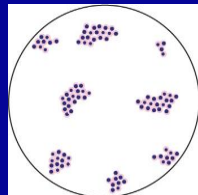
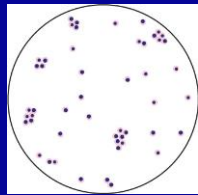




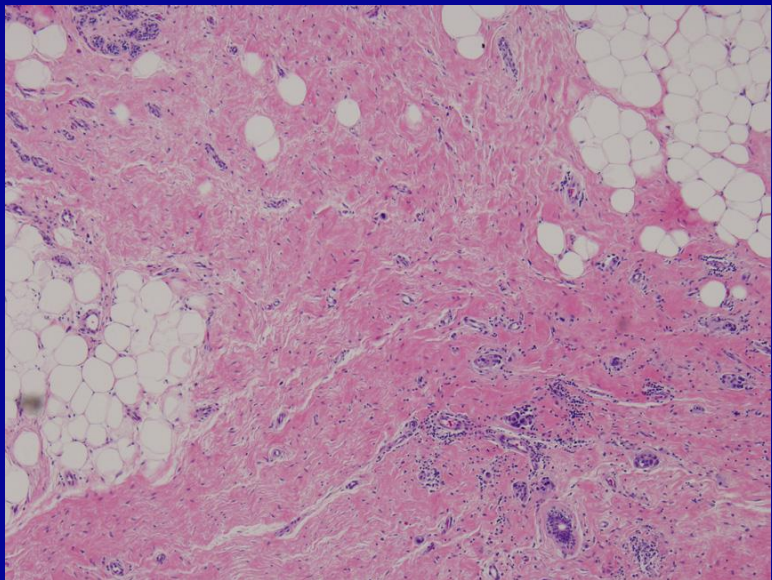
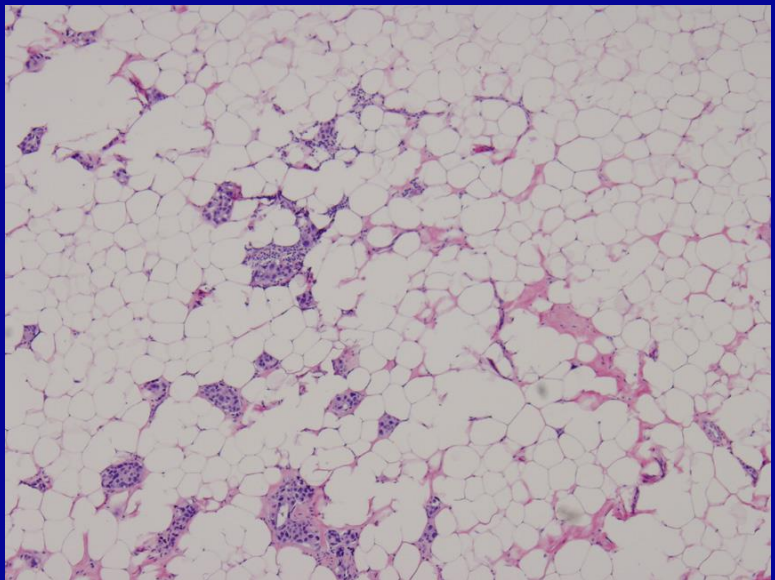
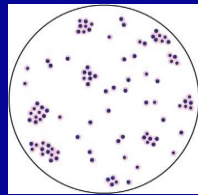




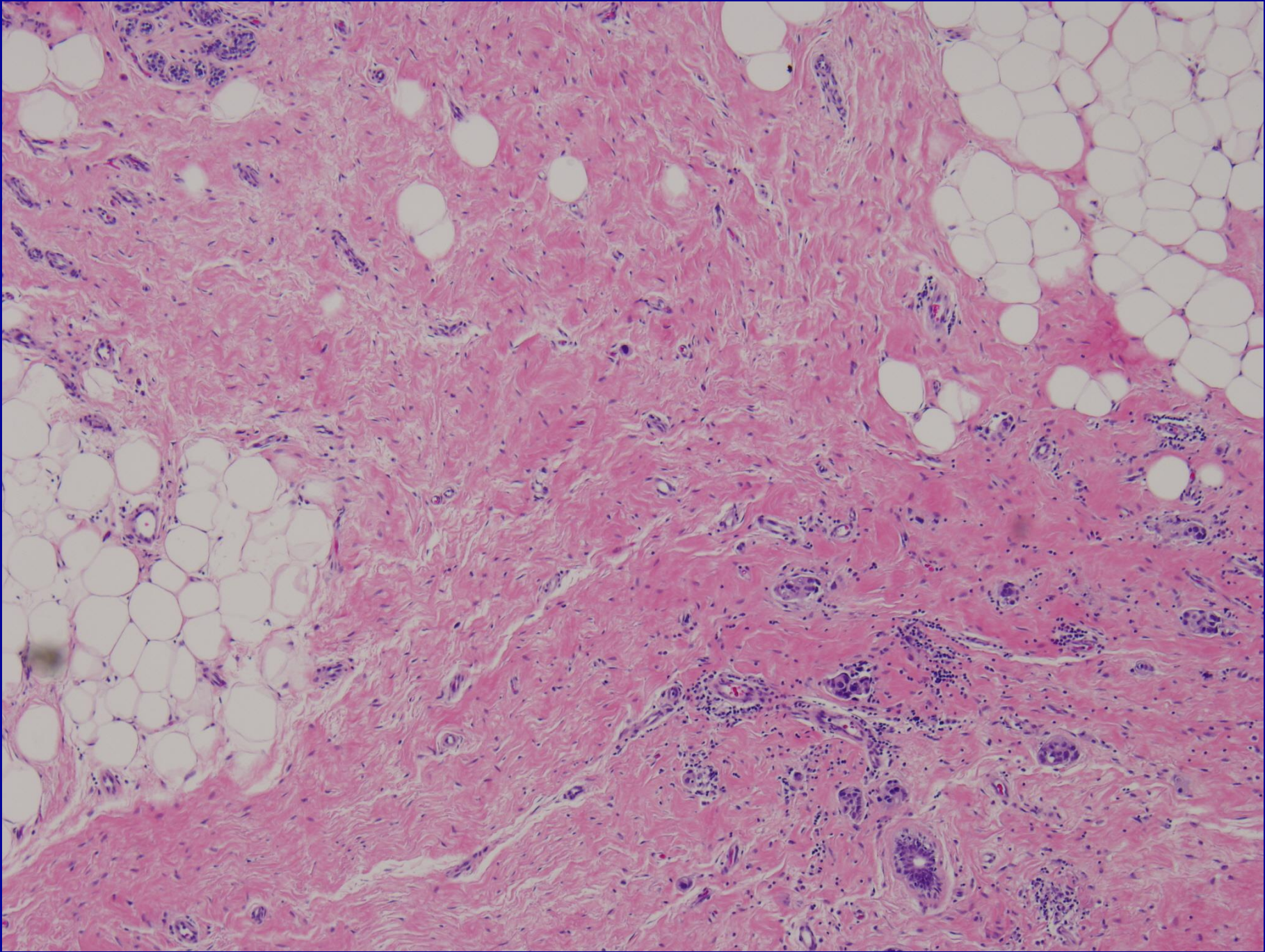
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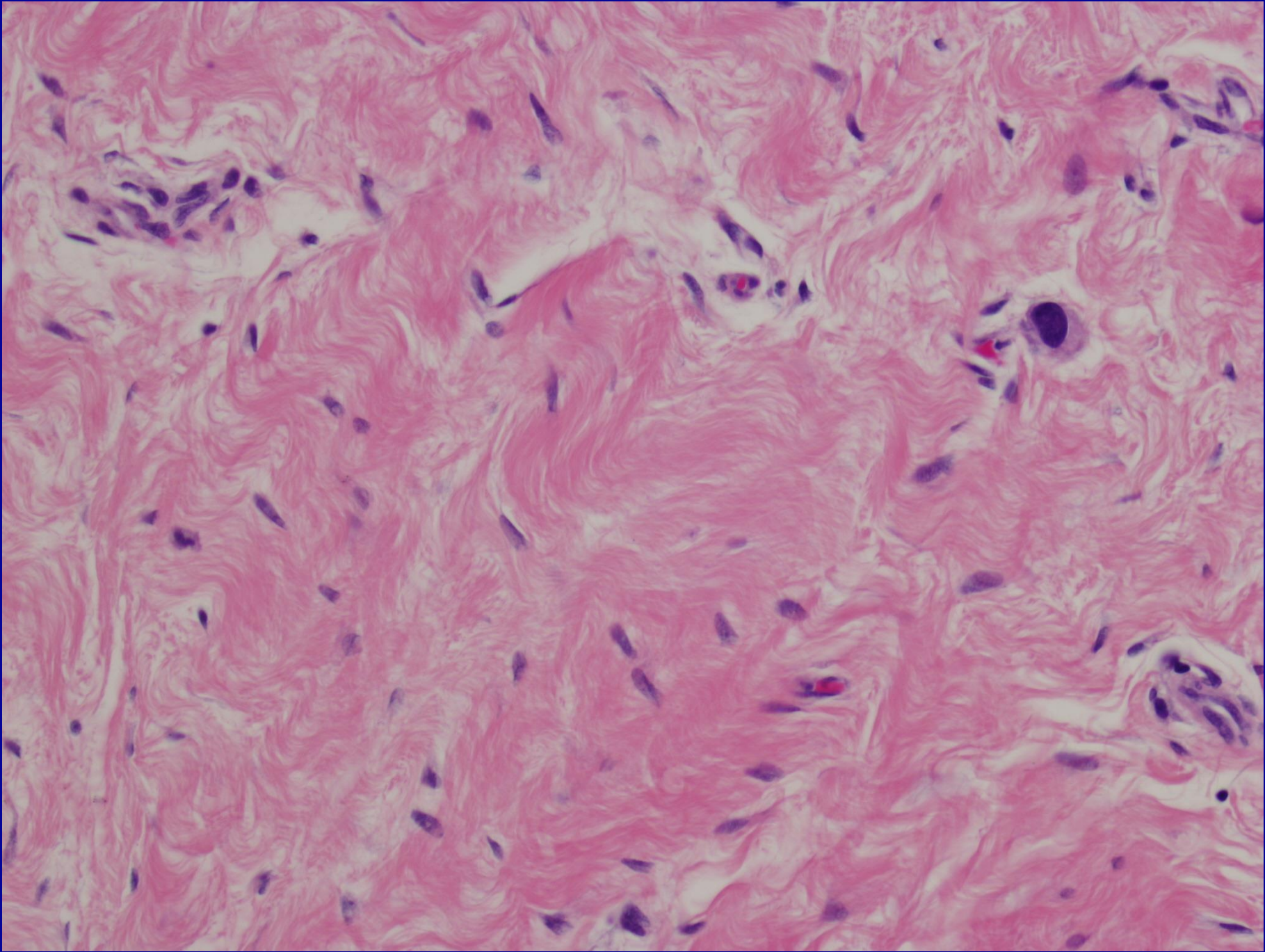


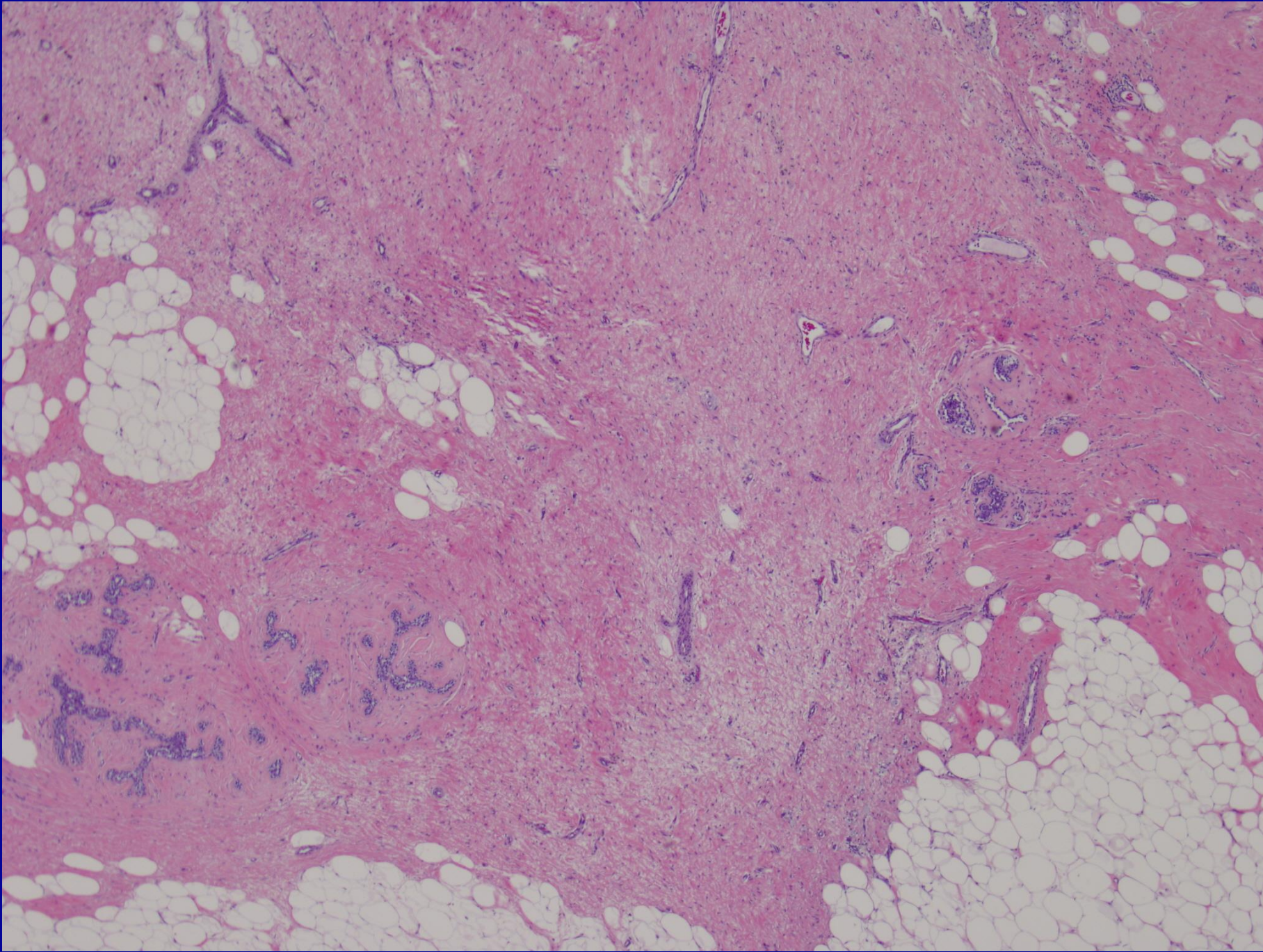
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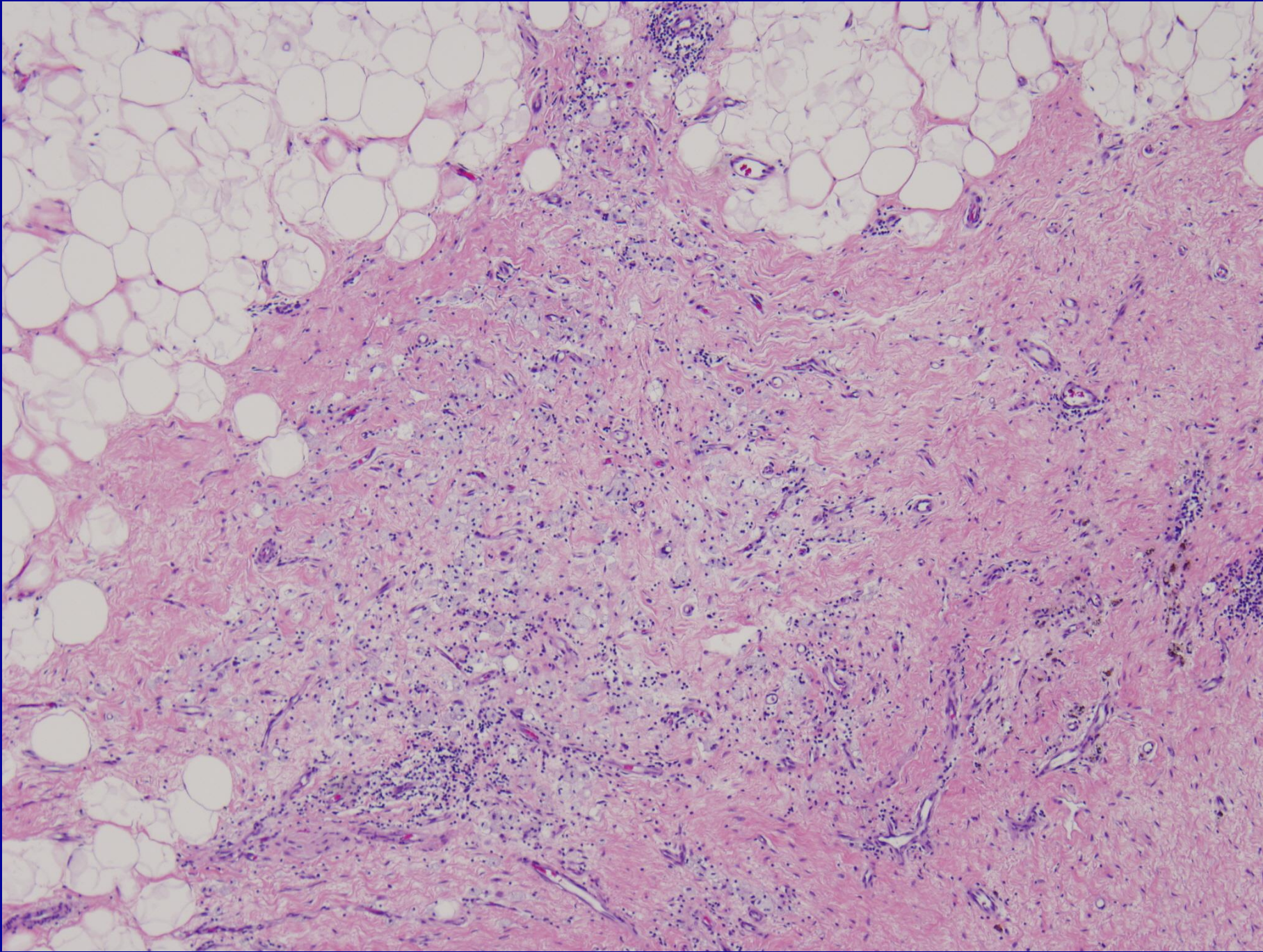


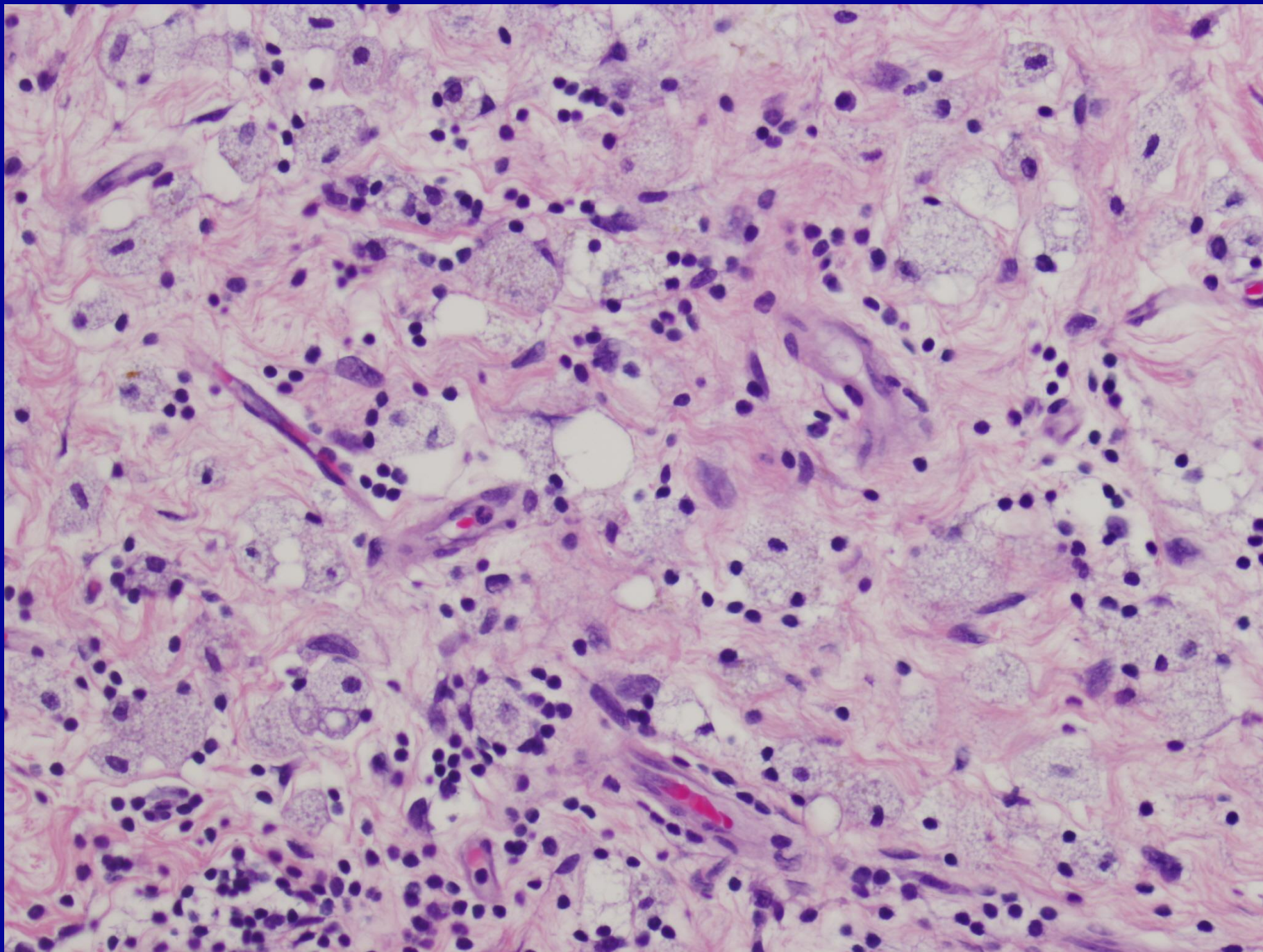












# Resections: Before We Start

- Clinical summary
  - Pathology requisition form
  - Direct communication with pathologist
  - Electronic medical record
- Knowledge of the pre-surgical pathology findings
  - Report
  - Digital image

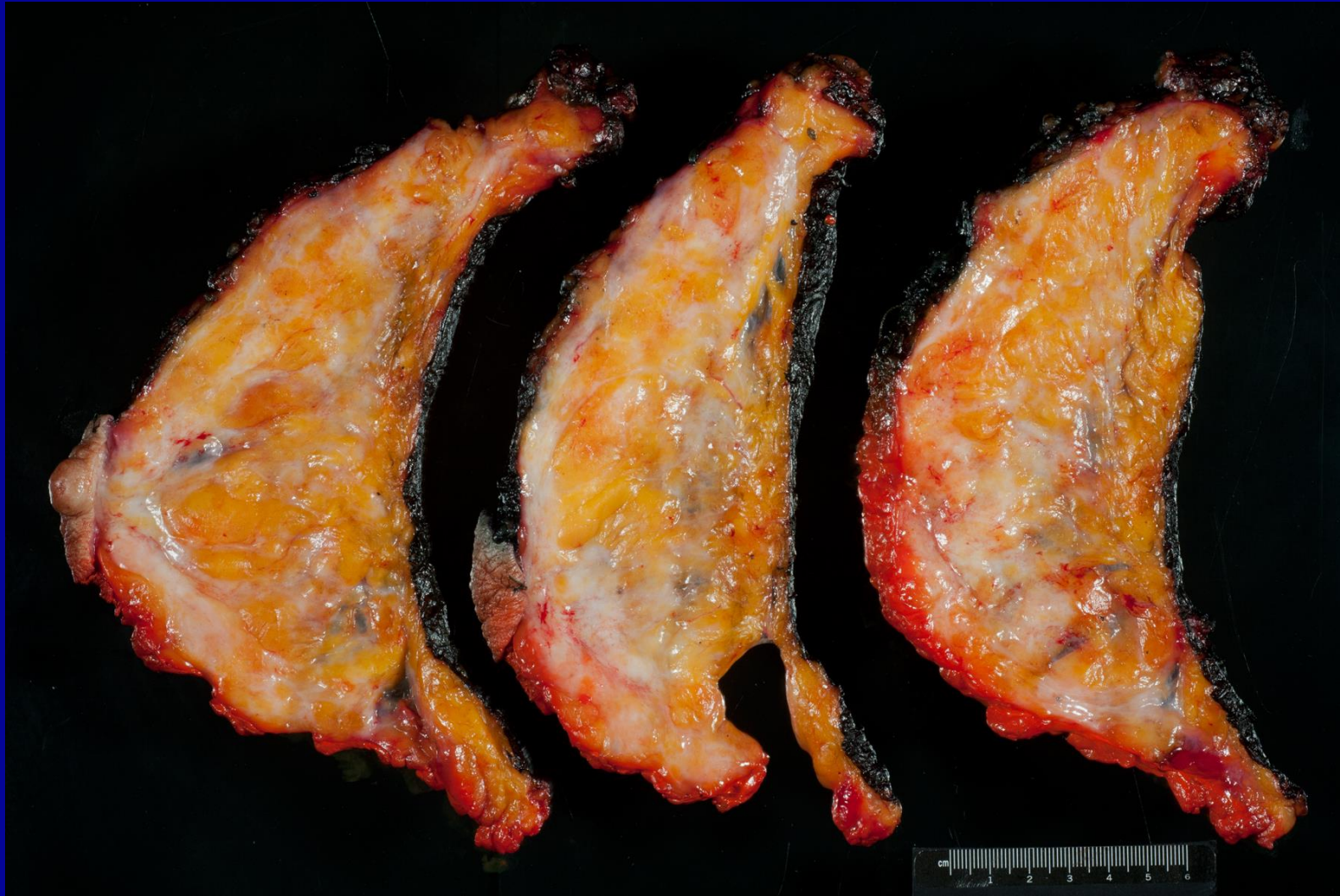
# Localization and Extent

- Medical records
- Clips, coils, seeds, wires, tatoos, ...
- Specimen radiography
- Sliced specimen radiography
- Section maps
- Digital measurement tools

# Response to Chemotherapy

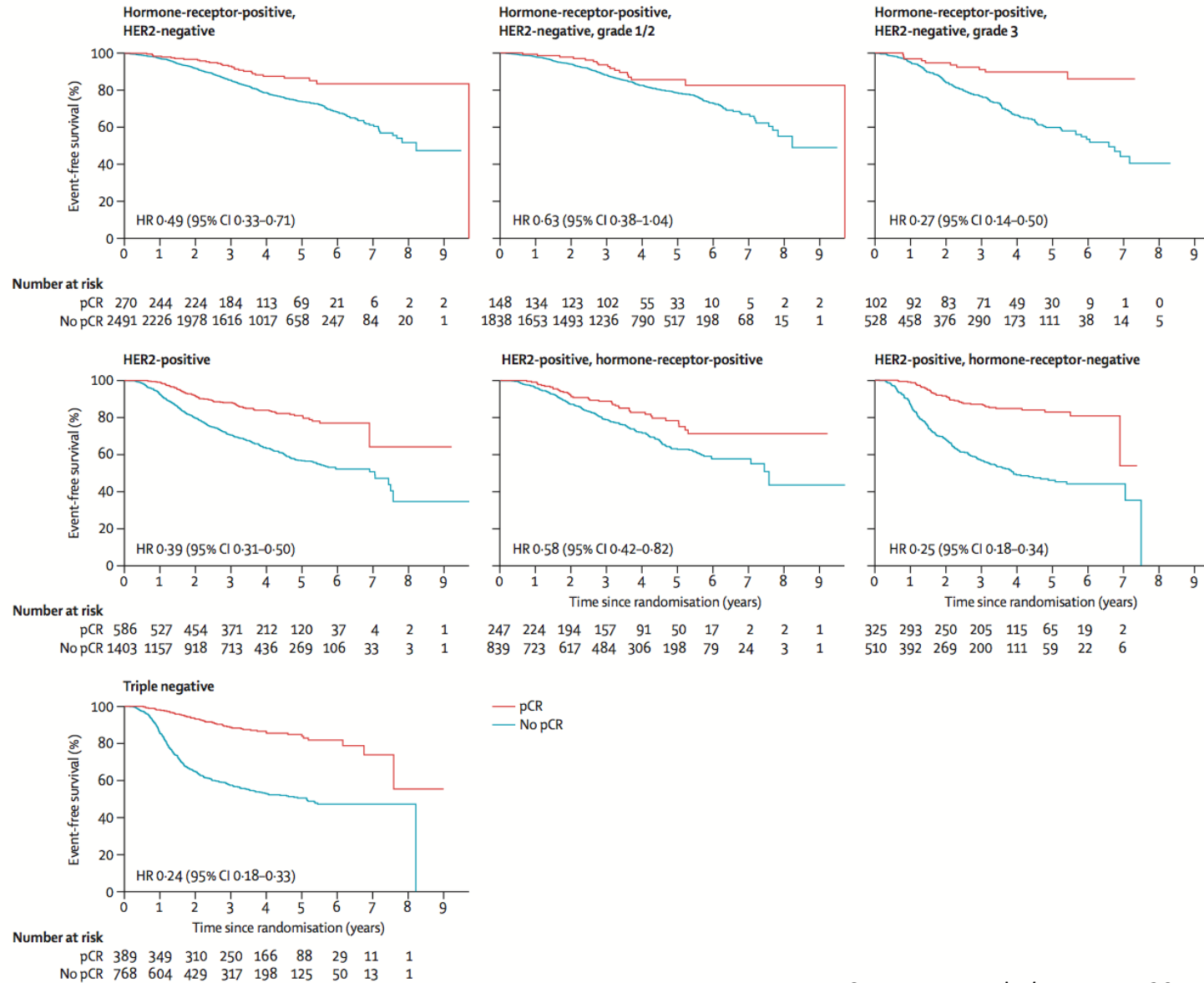


## The 3 Informative Slices Of The 13 Slices From The Mastectomy After Neoadjuvant Chemotherapy

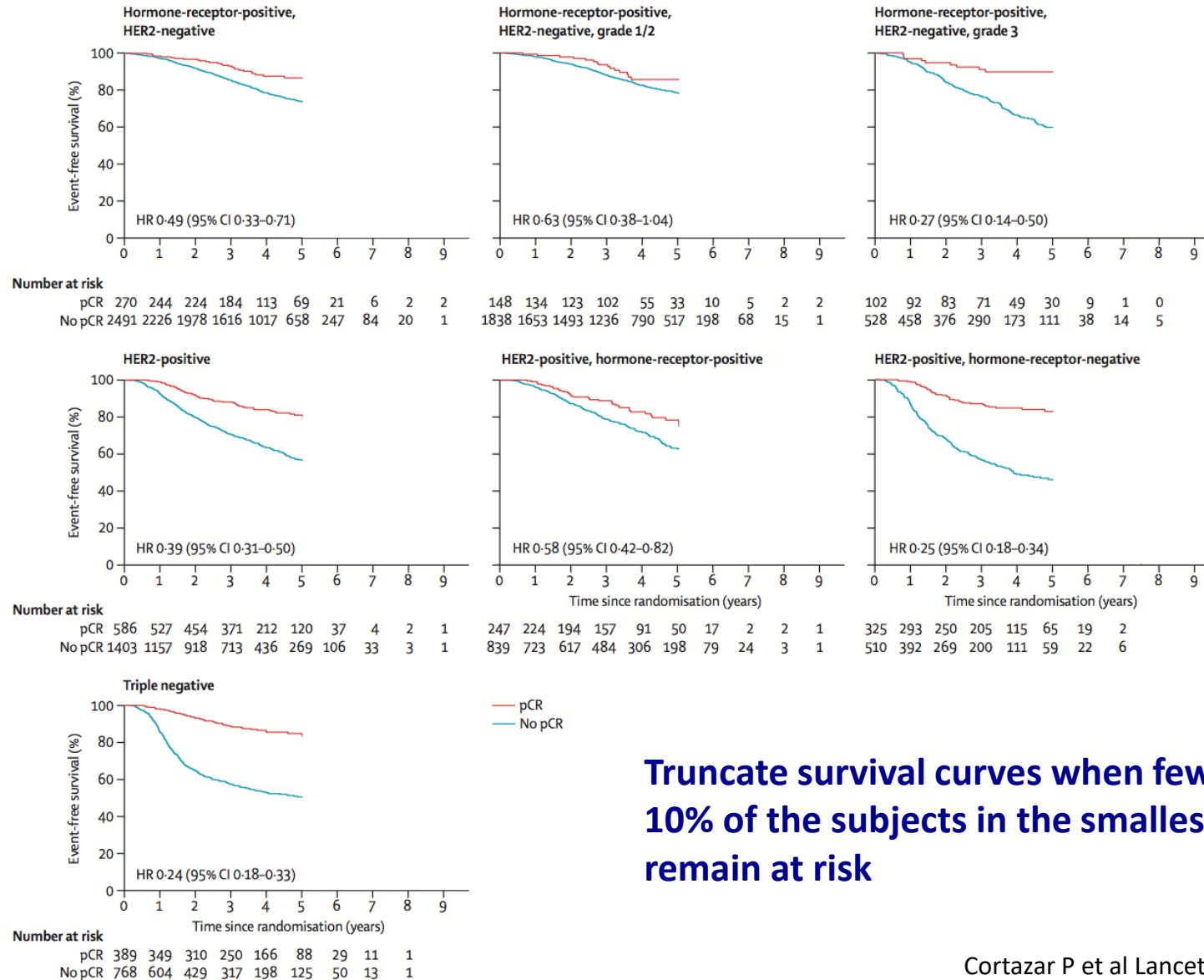


**Can you imagine a primary endpoint for clinical trials that is defined by absence of disease, but relies on preferences of local sites to identify and sample the correct area within each resection specimen?**

# Pathologic Complete Response (pCR) indicates good prognosis



# Pathologic Complete Response (pCR) indicates good prognosis



**Truncate survival curves when fewer than 10% of the subjects in the smallest group remain at risk**

Cortazar P et al Lancet 384:164-72, 2014

LEFT BREAST, SKIN-SPARING MASTECTOMY:

RESIDUAL INVASIVE DUCTAL CARCINOMA WITH TREATMENT EFFECT.

INVASIVE CARCINOMA MEASURES 1.3 X 1.3 CM AND CONTAINS APPROXIMATELY  
5% CANCER CELLULARITY BY AREA.

FOCAL LYMPHOVASCULAR INVASION IS PRESENT.

SCATTERED SMALL FOCI OF INTRADUCTAL CARCINOMA (DCIS) SURROUNDING  
INVASIVE CARCINOMA AND ADJACENT TO PRIOR BIOPSY SITES.

DCIS IS PRESENT 3 MM FROM INFERIOR SUPERFICIAL MARGIN.

INVASIVE CARCINOMA IS PRESENT AT LEAST 3 MM FROM INFERIOR SUPERFICIAL  
MARGIN.

MICROCALCIFICATIONS ASSOCIATED WITH DCIS AND BENIGN BREAST TISSUE.

LYMPH NODES, LEFT AXILLA, LEVELS I AND II, DISSECTION:

Fourteen lymph nodes, no carcinoma identified (0/14).

LEFT BREAST, NEW INFERIOR MARGIN, EXCISION:

No tumor present.

LEFT BREAST, NEW INFERIOR LATERAL MARGIN, EXCISION:

No tumor present.

COMMENTS: There was no residual invasive carcinoma at the sites of the clips. Pathologic  
AJCC Stage y-pT1c, y-pN0. Residual disease with pathologic findings c/w RCB-I.

## **Recommendations for standardized pathological characterization of residual disease for neoadjuvant clinical trials of breast cancer by the BIG-NABCG collaboration**

V. Bossuyt<sup>1\*</sup>, E. Provenzano<sup>2</sup>, W. F. Symmans<sup>3</sup>, J. C. Boughey<sup>4</sup>, C. Coles<sup>5</sup>, G. Curigliano<sup>6</sup>, J. M. Dixon<sup>7</sup>, L. J. Esserman<sup>8</sup>, G. Fastner<sup>9</sup>, T. Kuehn<sup>10</sup>, F. Peintinger<sup>11,12</sup>, G. von Minckwitz<sup>13</sup>, J. White<sup>14</sup>, W. Yang<sup>15</sup>, S. Badve<sup>16</sup>, C. Denkert<sup>17</sup>, G. MacGrogan<sup>18</sup>, F. Penault-Llorca<sup>19</sup>, G. Viale<sup>20</sup> & D. Cameron<sup>21</sup> of the Breast International Group-North American Breast Cancer Group (BIG-NABCG) collaboration

*Annals of Oncology 00: 1–12, 2015 doi:10.1093/annonc/mdv161*

## **Standardization of pathologic evaluation and reporting of postneoadjuvant specimens in clinical trials of breast cancer: recommendations from an international working group**

Elena Provenzano<sup>1</sup>, Veerle Bossuyt<sup>2</sup>, Giuseppe Viale<sup>3</sup>, David Cameron<sup>4</sup>, Sunil Badve<sup>5</sup>, Carsten Denkert<sup>6</sup>, Gaëtan MacGrogan<sup>7</sup>, Frédérique Penault-Llorca<sup>8</sup>, Judy Boughey<sup>9</sup>, Giuseppe Curigliano<sup>10</sup>, J Michael Dixon<sup>11</sup>, Laura Esserman<sup>12</sup>, Gerd Fastner<sup>13</sup>, Thorsten Kuehn<sup>14</sup>, Florentia Peintinger<sup>15,16</sup>, Gunter von Minckwitz<sup>17</sup>, Julia White<sup>18</sup>, Wei Yang<sup>19</sup> and W Fraser Symmans<sup>20</sup> on behalf of the Residual Disease Characterization Working Group of the Breast International Group-North American Breast Cancer Group (BIG-NABCG) collaboration

*Modern Pathol 00: 1–17, 2015 doi:10.1038/modpathol.2015.74*

# Summary of Recommendations

Mandate of this working group committee was limited to recommendations for clinical trials

Provide the following information:

1. pCR (ypT0 ypN0 and ypT0/is ypN0) *versus* residual disease,
2. ypT and ypN Stage using the current AJCC/UICC staging system, &
3. Residual cancer burden (RCB)

**A single standardized approach to macroscopic and microscopic pathologic examination makes it easy to reliably provide all 3 results!**

Post-treatment yp-Stage (AJCC)

# 8<sup>th</sup> Edition AJCC (2017) Summary of Changes

Categories of *yp*-Stage are the same as for *p*-Stage

**Primary Tumor:** The Expert Panel clarified that ....

- Measurement for *ypT* is based on the largest focus of *continuous* residual invasive carcinoma
- Treatment-related fibrosis adjacent to residual invasive carcinoma is not included in the *ypT* measurement
- When multiple foci are present the (*m*) modifier is included

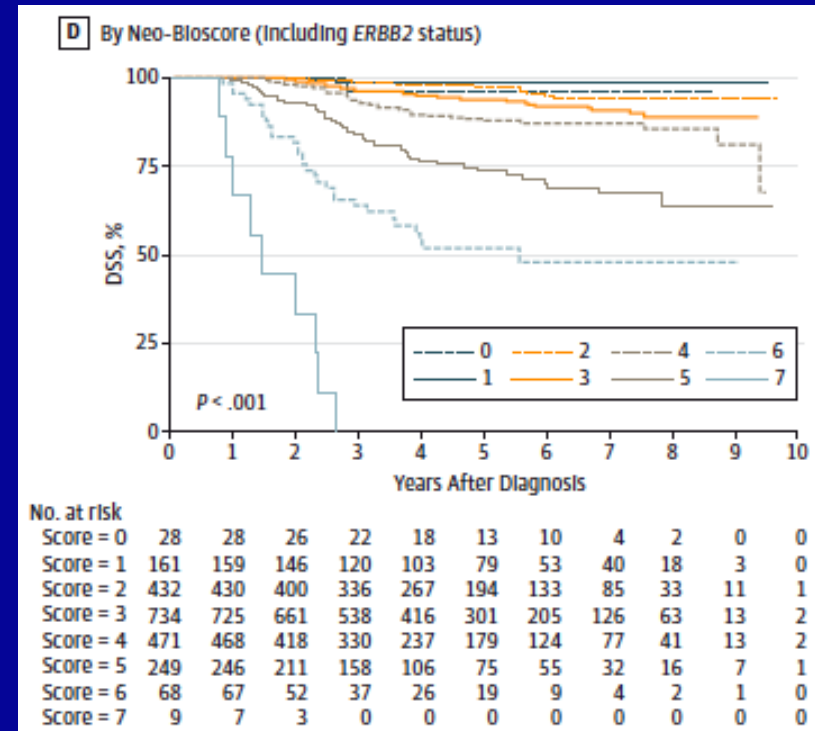
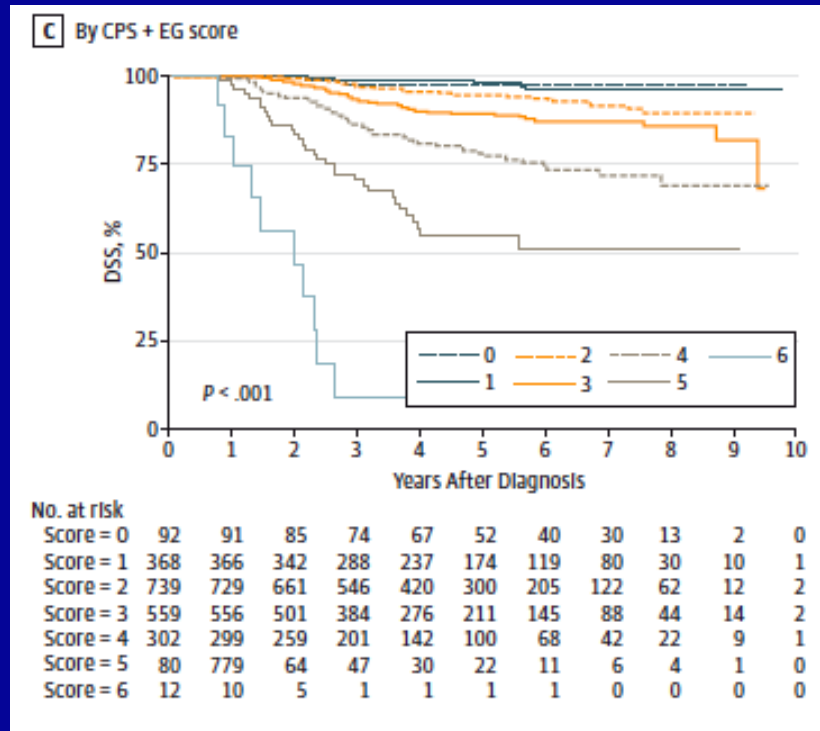
**Nodal Metastases:** The Expert Panel clarified that ....

- Measurement for *ypN* is based on the largest focus of *continuous* residual tumor
- Treatment-related fibrosis adjacent to residual invasive carcinoma is not included in the *ypT* measurement
- Isolated tumor cells *ypN0i* rules out pCR



# NeoBioscore Modification of CPS-EG

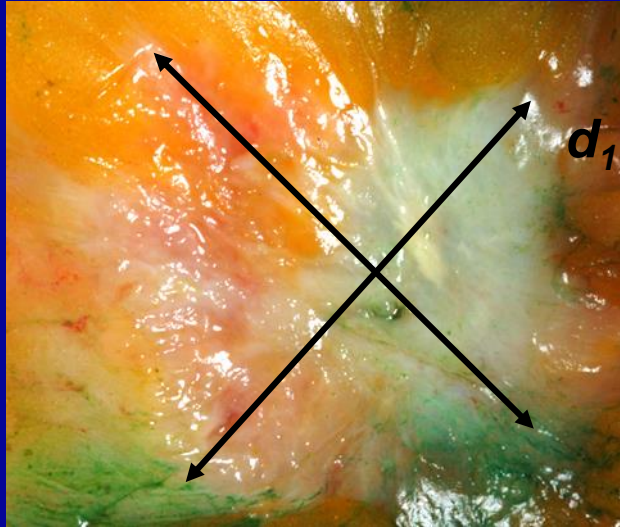
Pre-Rx Stage (c)		Pre-Rx Pathobiology						Post-Rx Stage (yp)	
c Stage	=	ER (1%)	=	N Grade	=	HER2	=	yp Stage	=
I - IIA	0	Positive	0	1 - 2	0	Positive	0	0 - I	0
IIB - IIIA	1	Negative	1	3	1	Negative	1	IIA - IIIB	1
IIIB - IIIC	2							IIIC	2



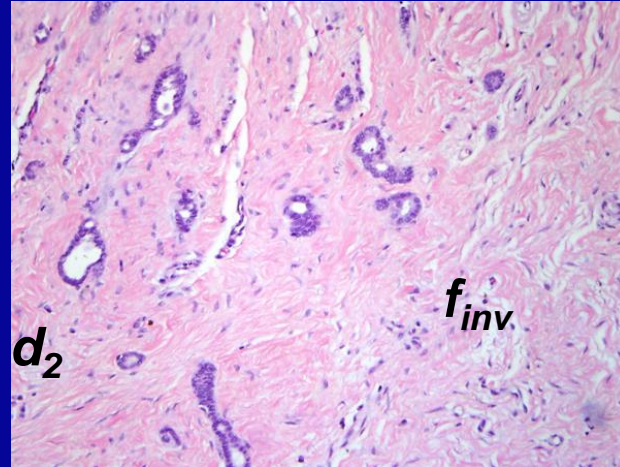
Residual Cancer Burden (RCB)

# Residual Cancer Burden (RCB)

Primary Tumor Bed

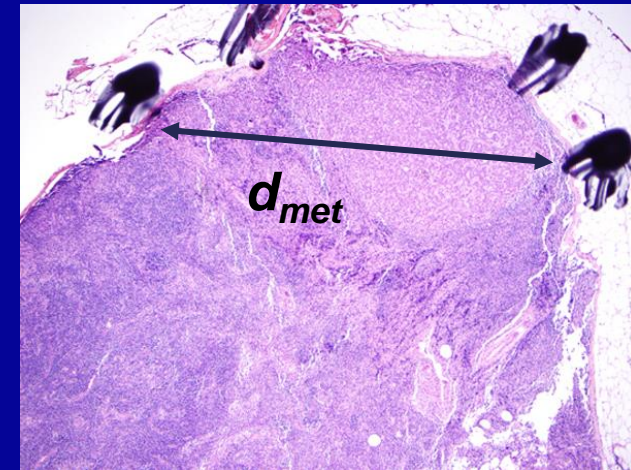


$$d_{prim} = \sqrt{d_1 d_2}$$



$$f_{inv} = \% \text{ area with invasive CA}$$

Lymph Nodes



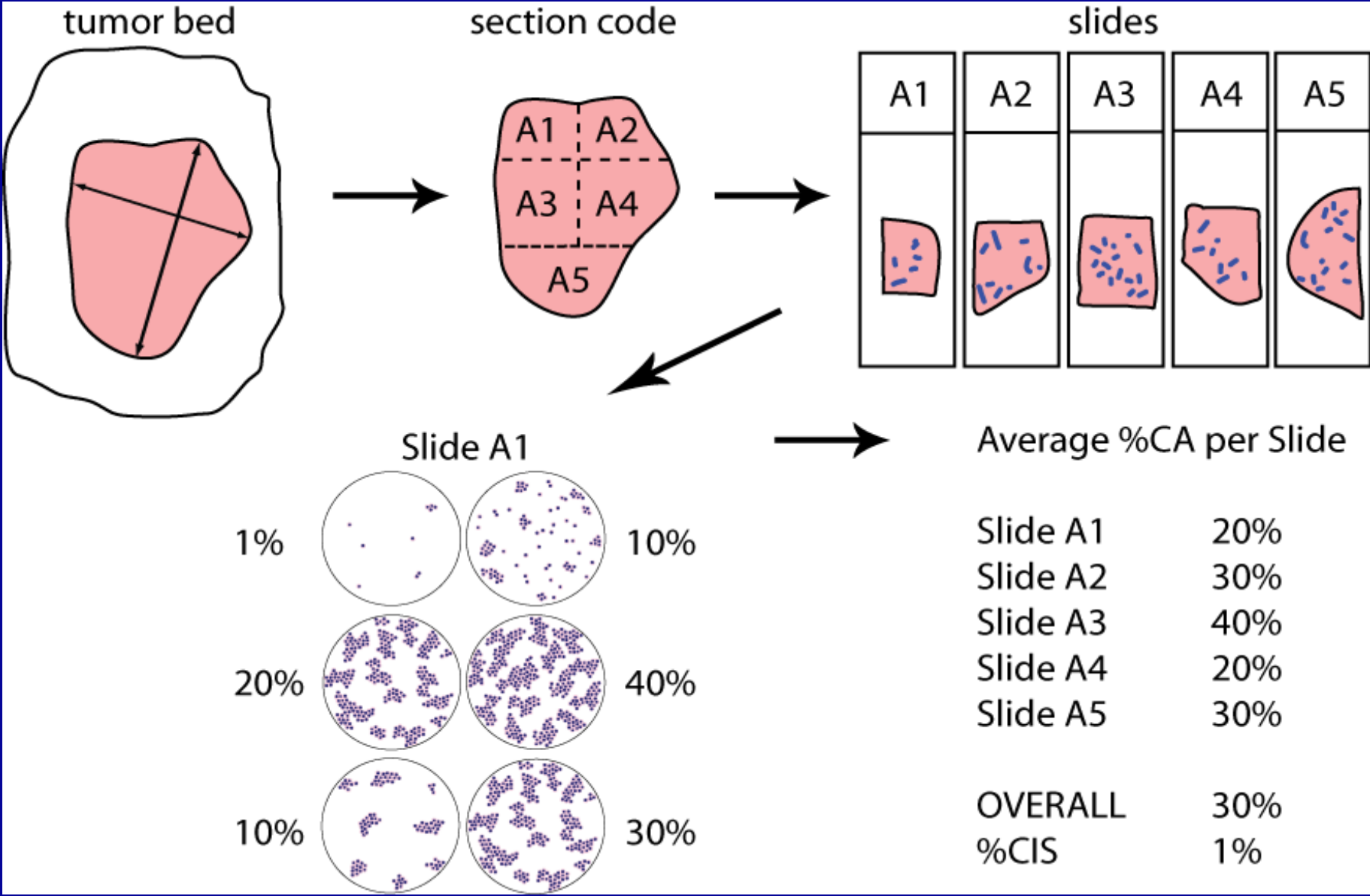
*LN = Number of Positive Nodes*

*d<sub>met</sub> = size largest metastasis*

## DRFS Following Neoadjuvant T/FAC Chemotherapy (N=241)

Variable	Hazard Ratio (95% CI)	P value
Primary tumor bed size ( $d_{prim}$ )	1.24 (1.04-1.48)	0.02
Fraction of invasive cancer ( $f_{inv}$ )	7.37 (2.16-25.1)	0.001
Number of positive lymph nodes ( $LN$ )	1.11 (1.04-1.19)	0.002
Size of largest metastasis ( $d_{met}$ )	1.17 (0.99-1.38)	0.06

# Pathologic Assessment Of The Primary Tumor Bed



# Example: Pathologist's Preparatory Notes

52 year old with triple-negative right breast cancer (T2, N0)

## Imaging

Tumor in right breast, 11-12 o'clock position, 8 cm from nipple

2.7 cm mass + minute satellites, overall 3.4 cm greatest dimension

Metal clip placed in tumor at time of diagnostic core biopsy

Ultrasound of regional nodal basins did not show any abnormal LNs

## Pathology From Biopsy

IDC grade 3, HR- / HER2- (TNBC)

## Treatment & Response

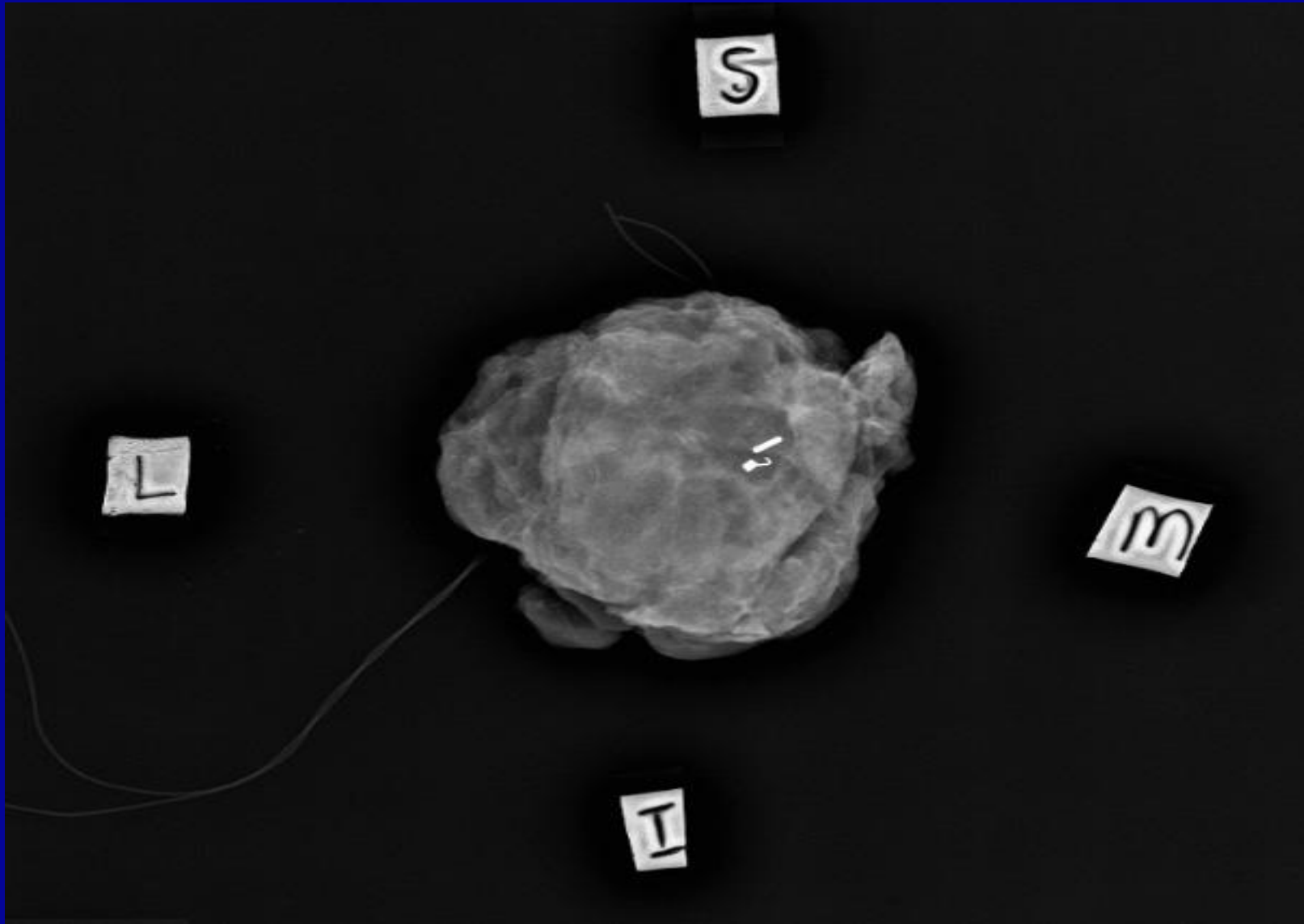
Weekly paclitaxel x 12 then 3-weekly FAC x 4 (T/FAC)

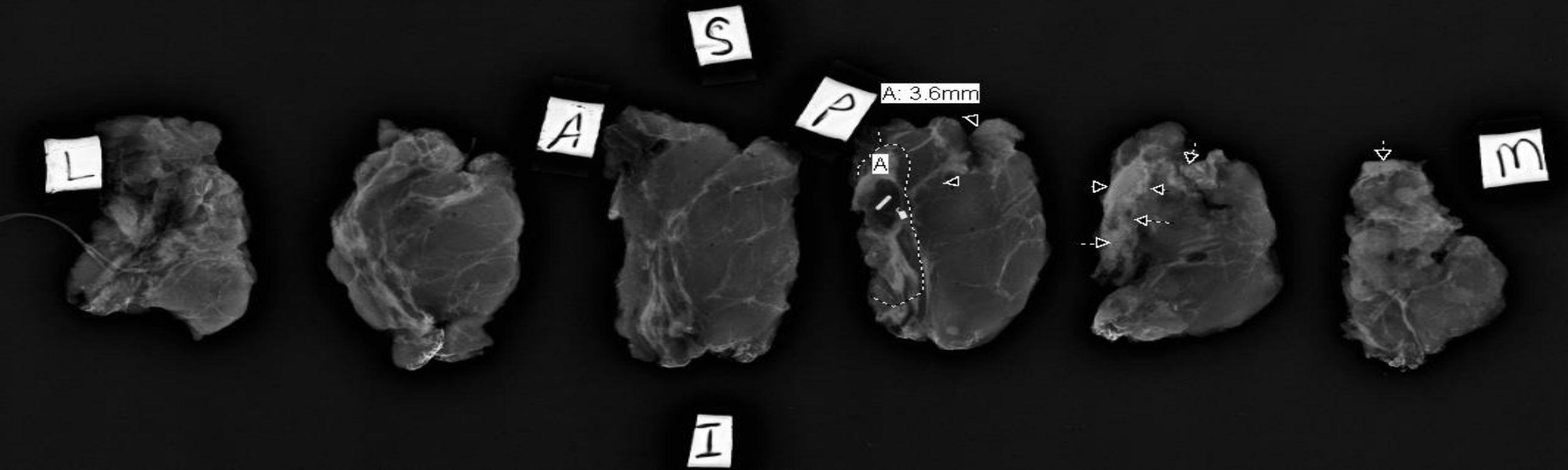
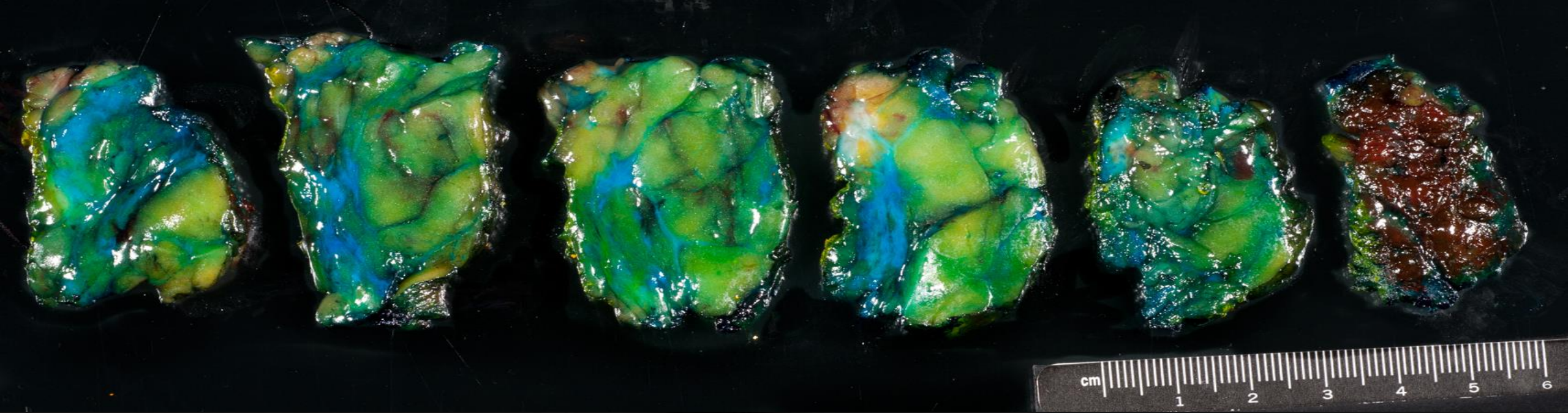
Residual architectural distortion, but no mass

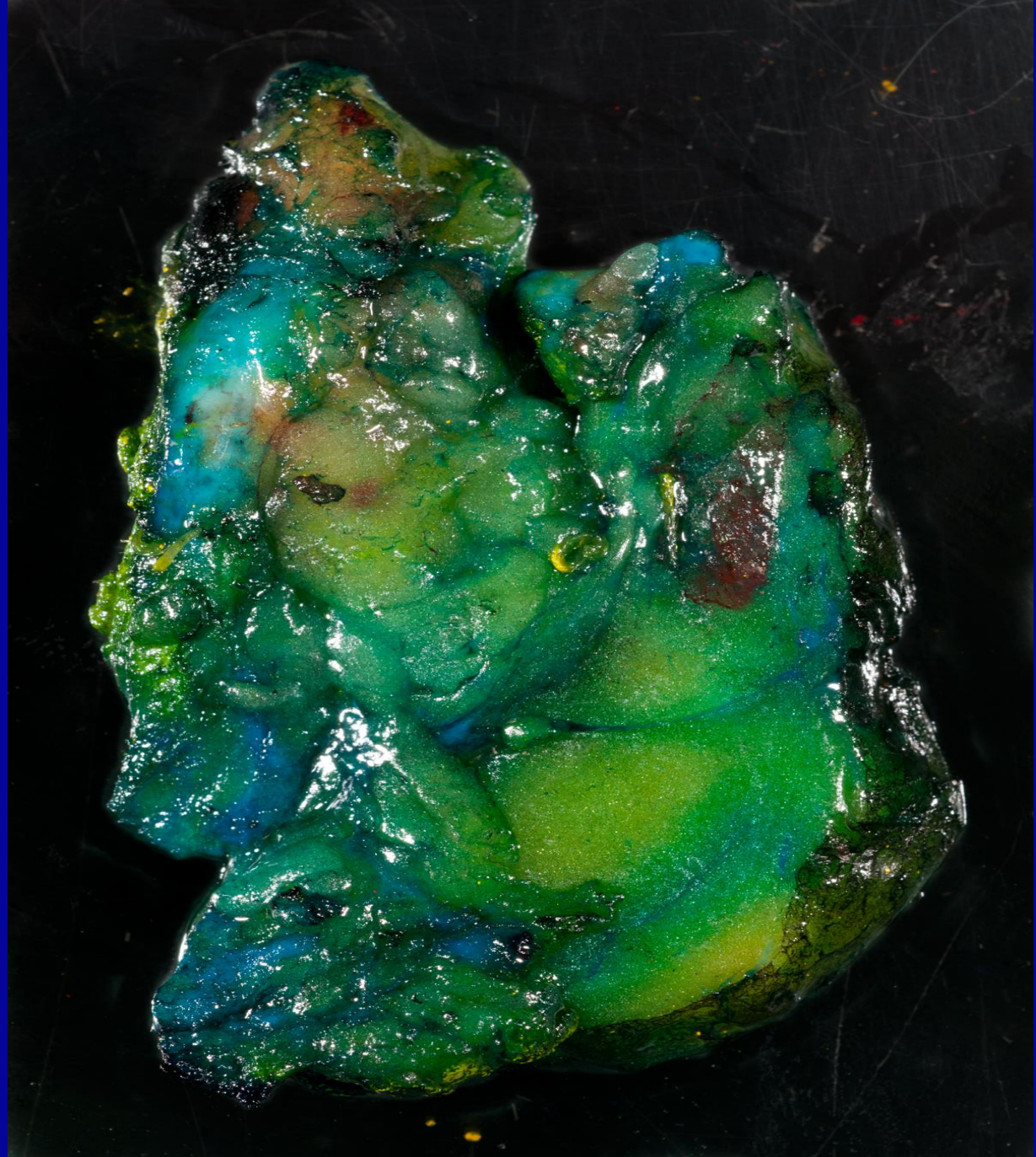
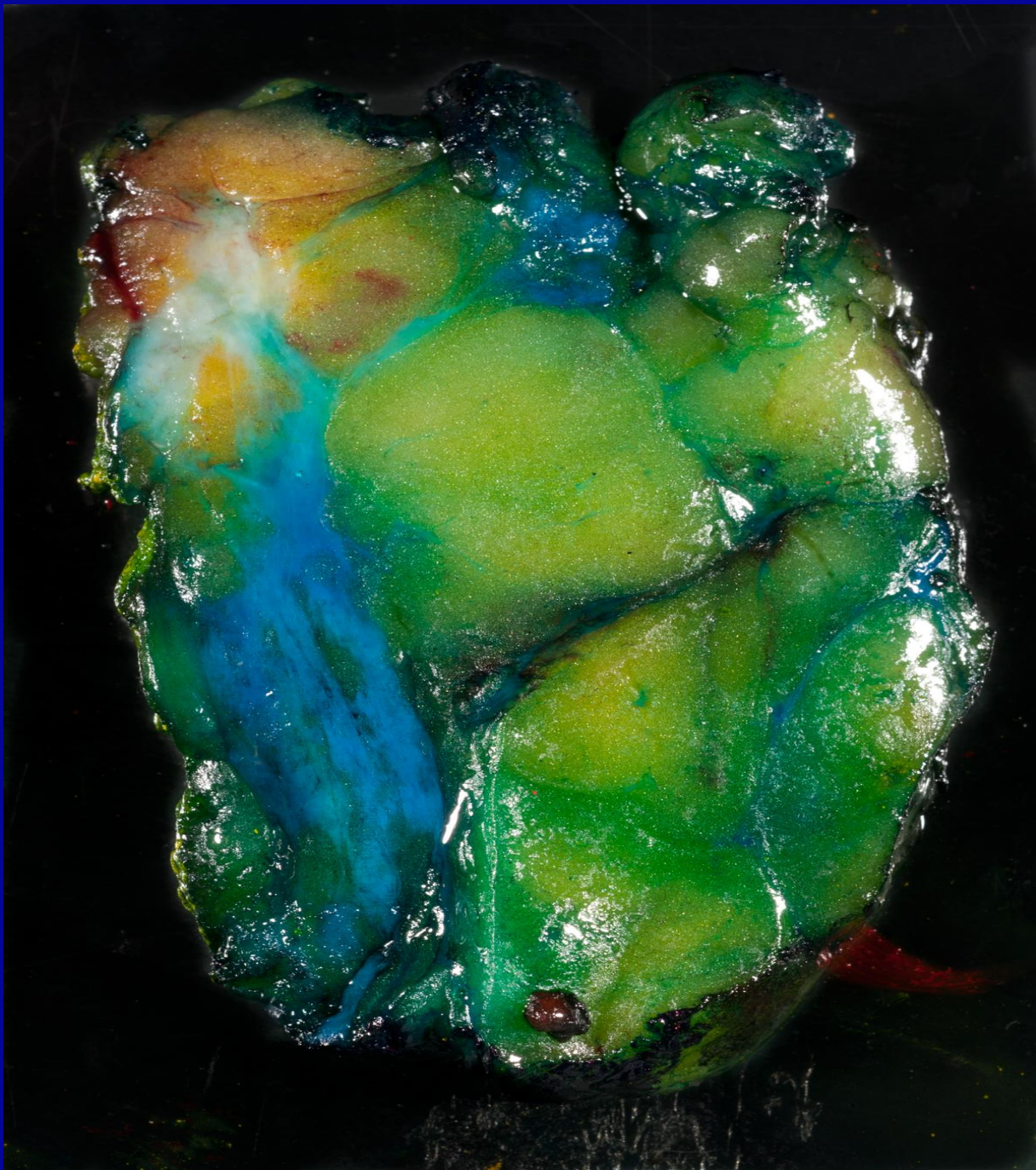
Radioactive seed placed in tumor on morning of surgery

For segmental mastectomy and sentinel node biopsy procedure

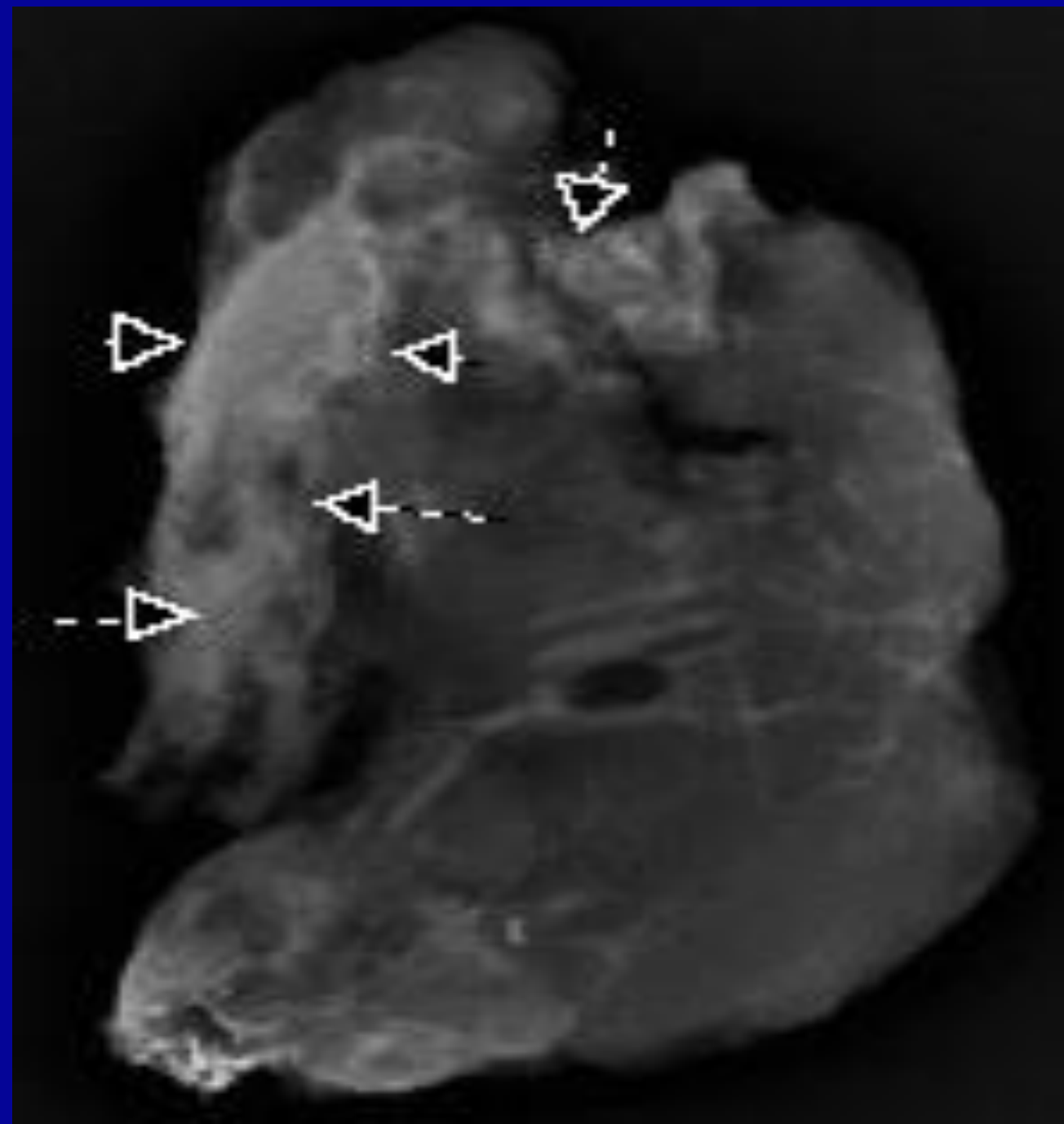
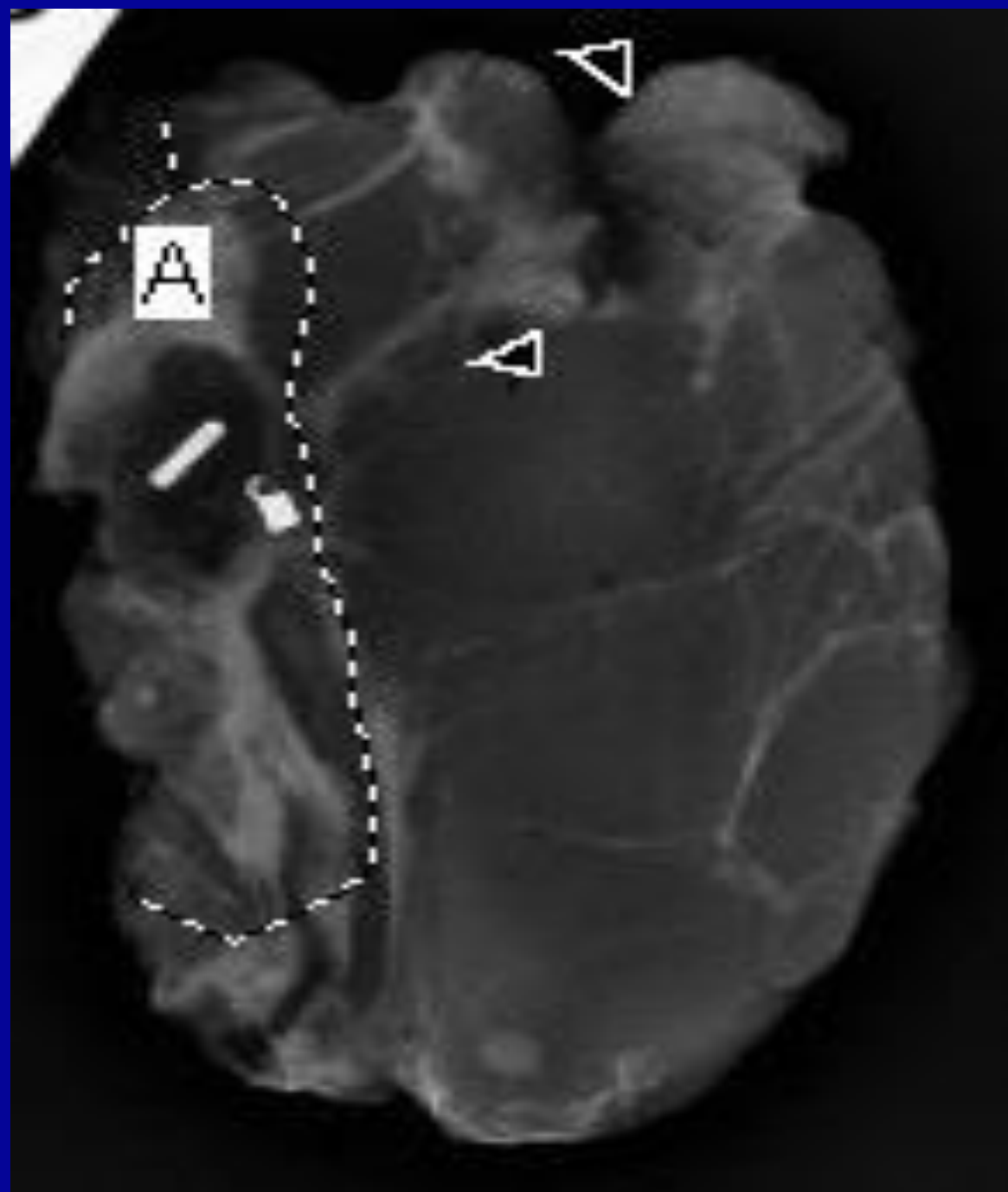
The clip and the seed are in the specimen

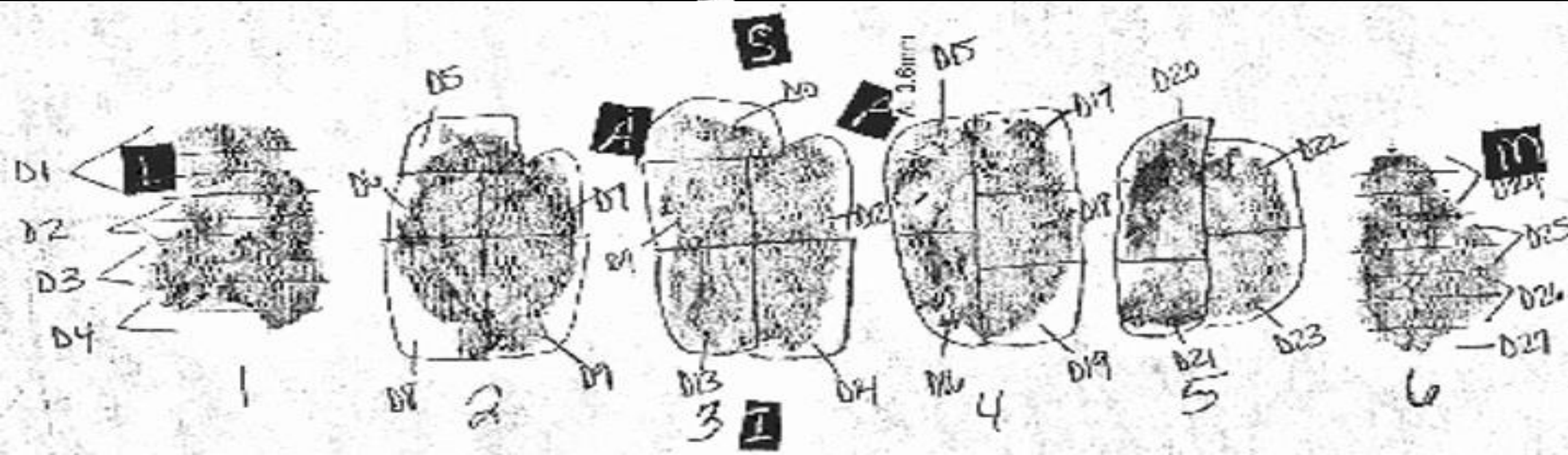
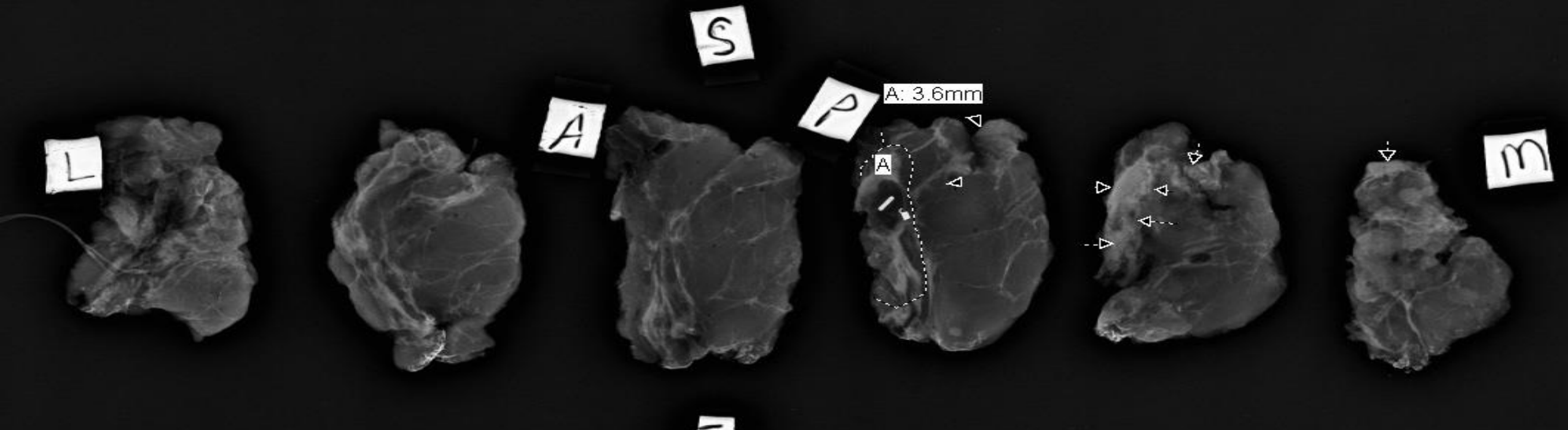




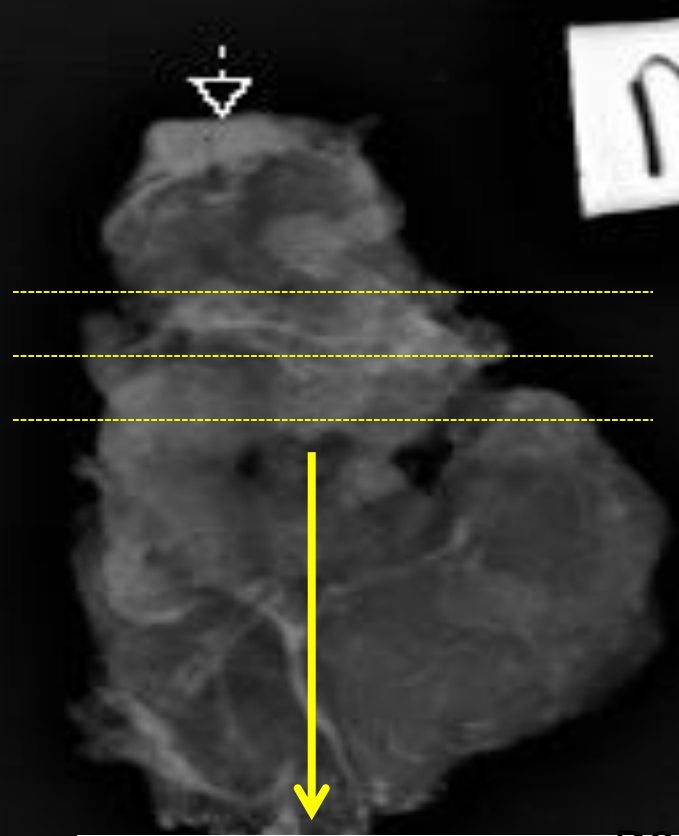
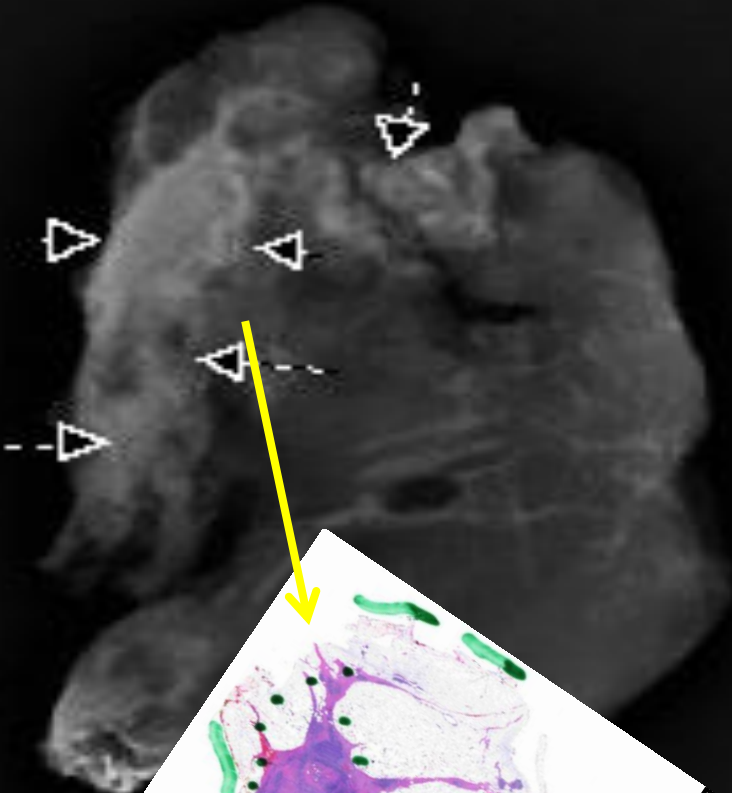




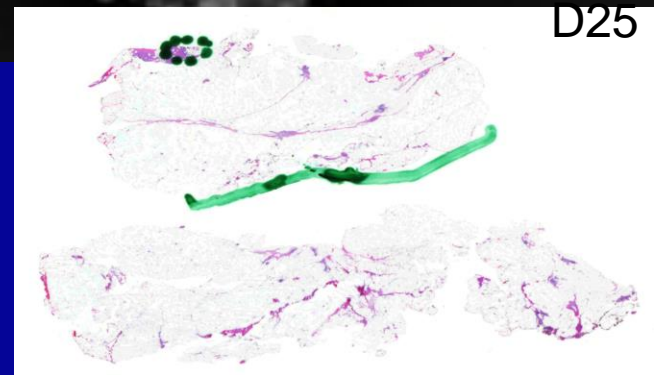
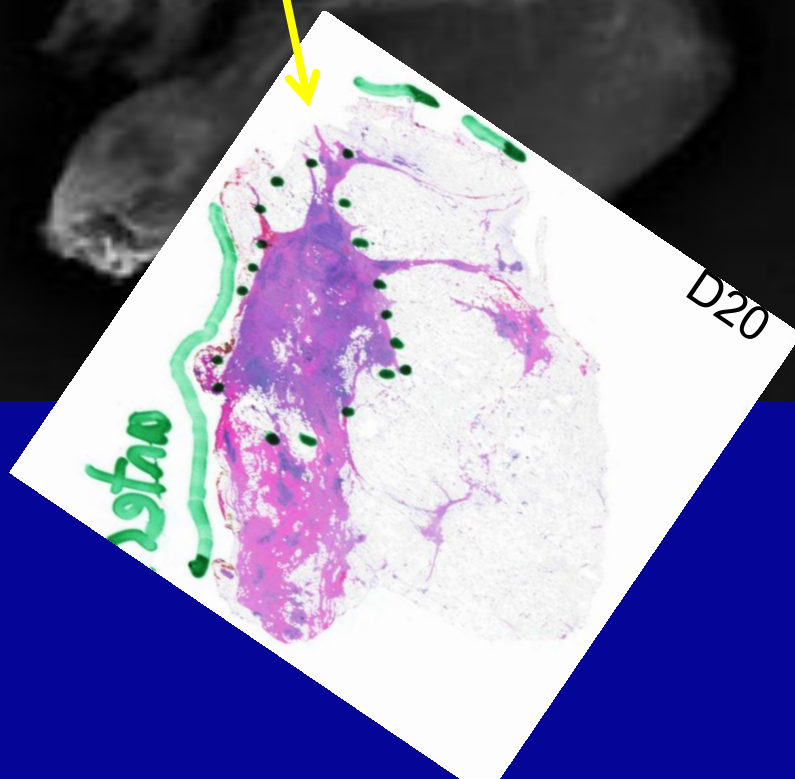
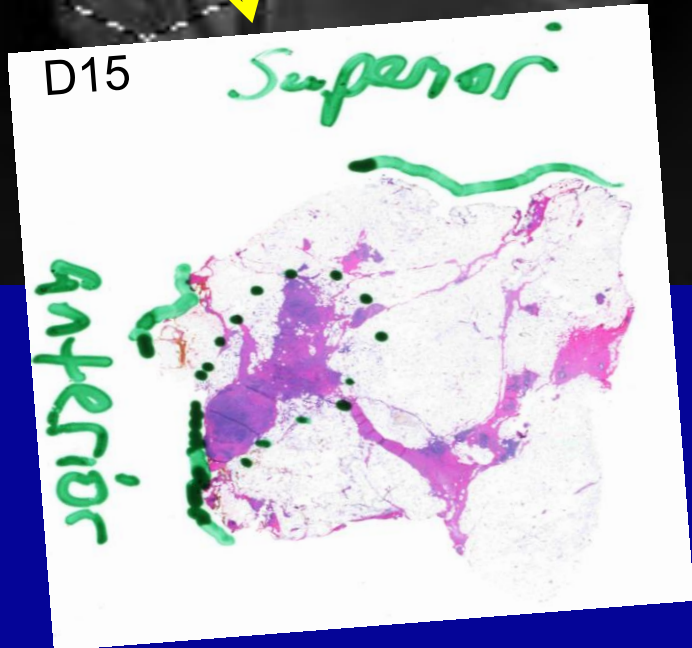


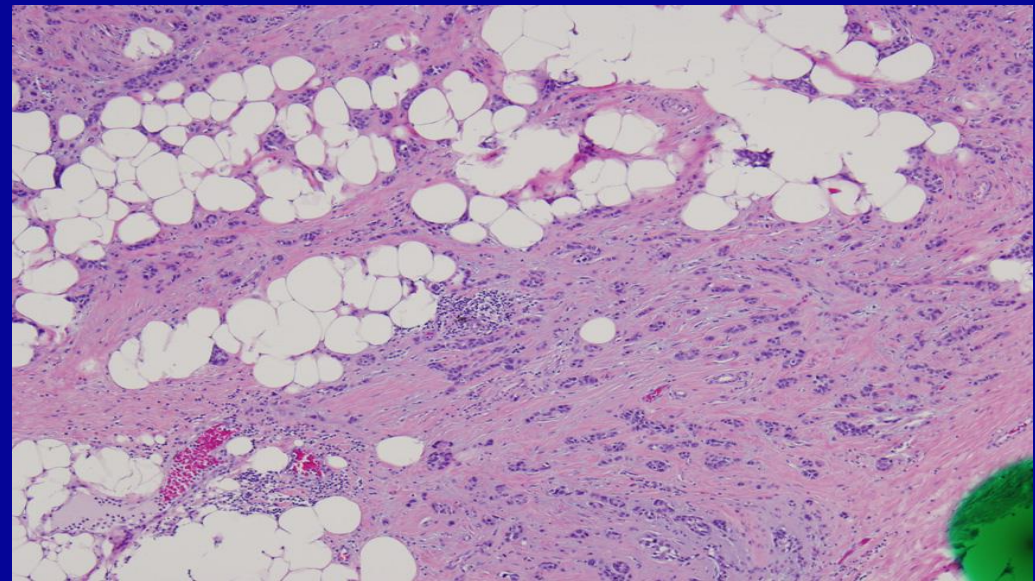
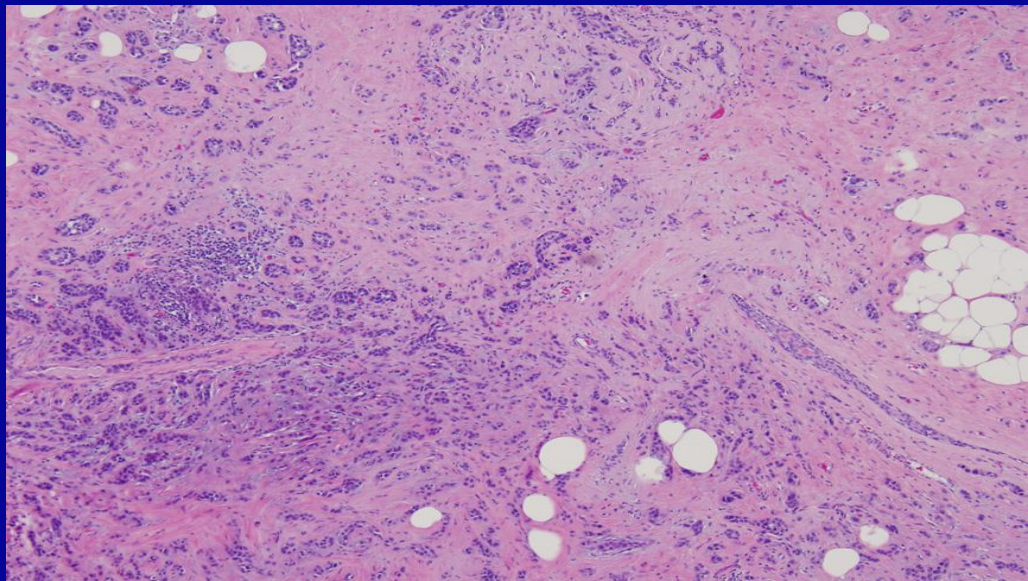
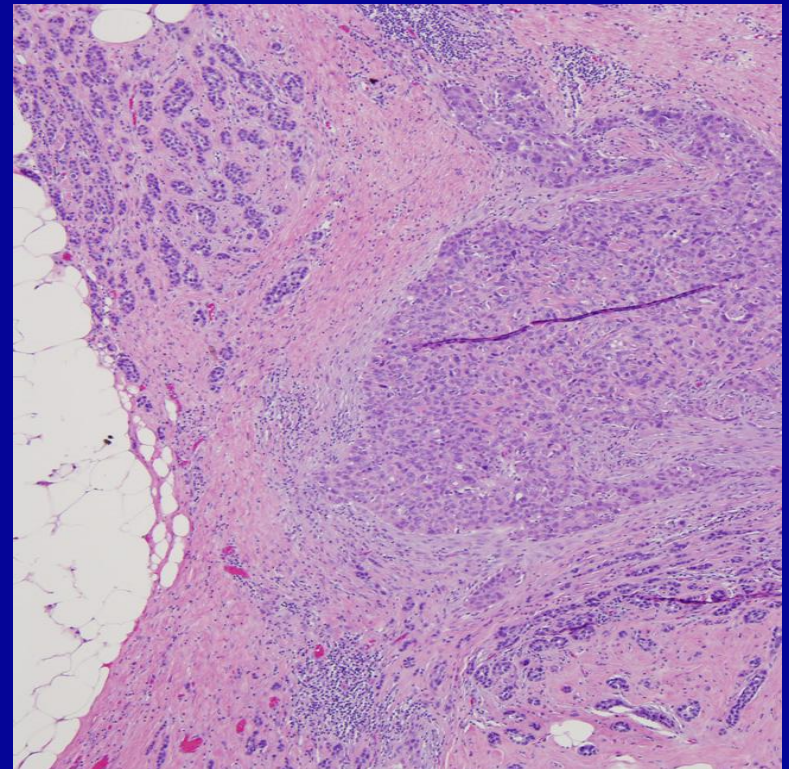
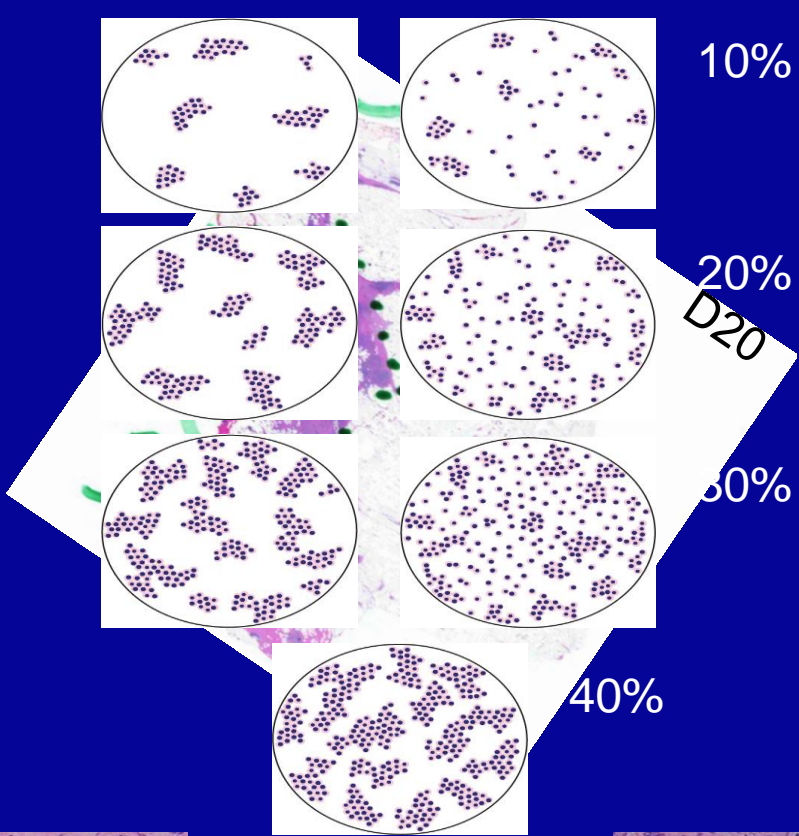
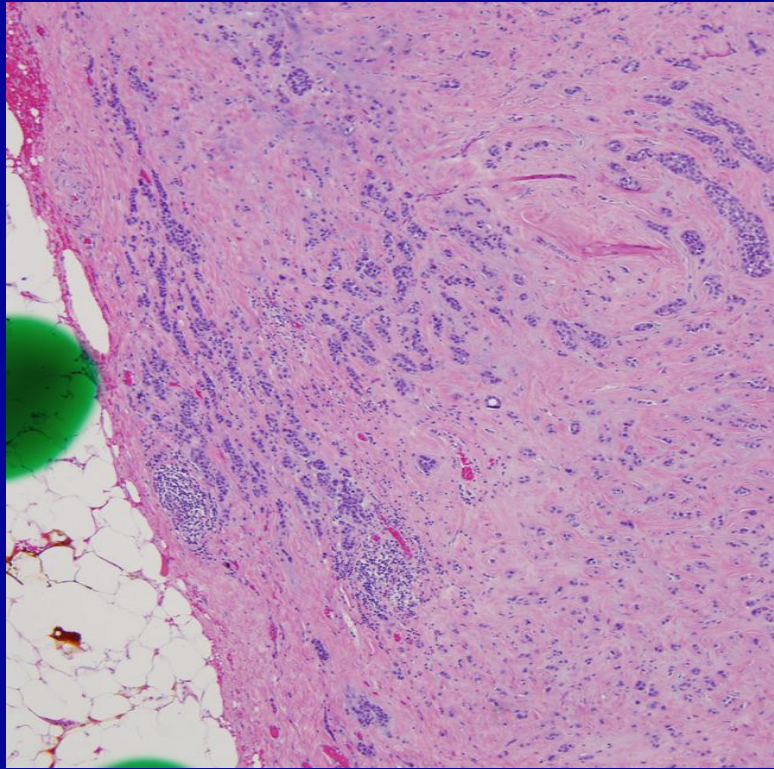


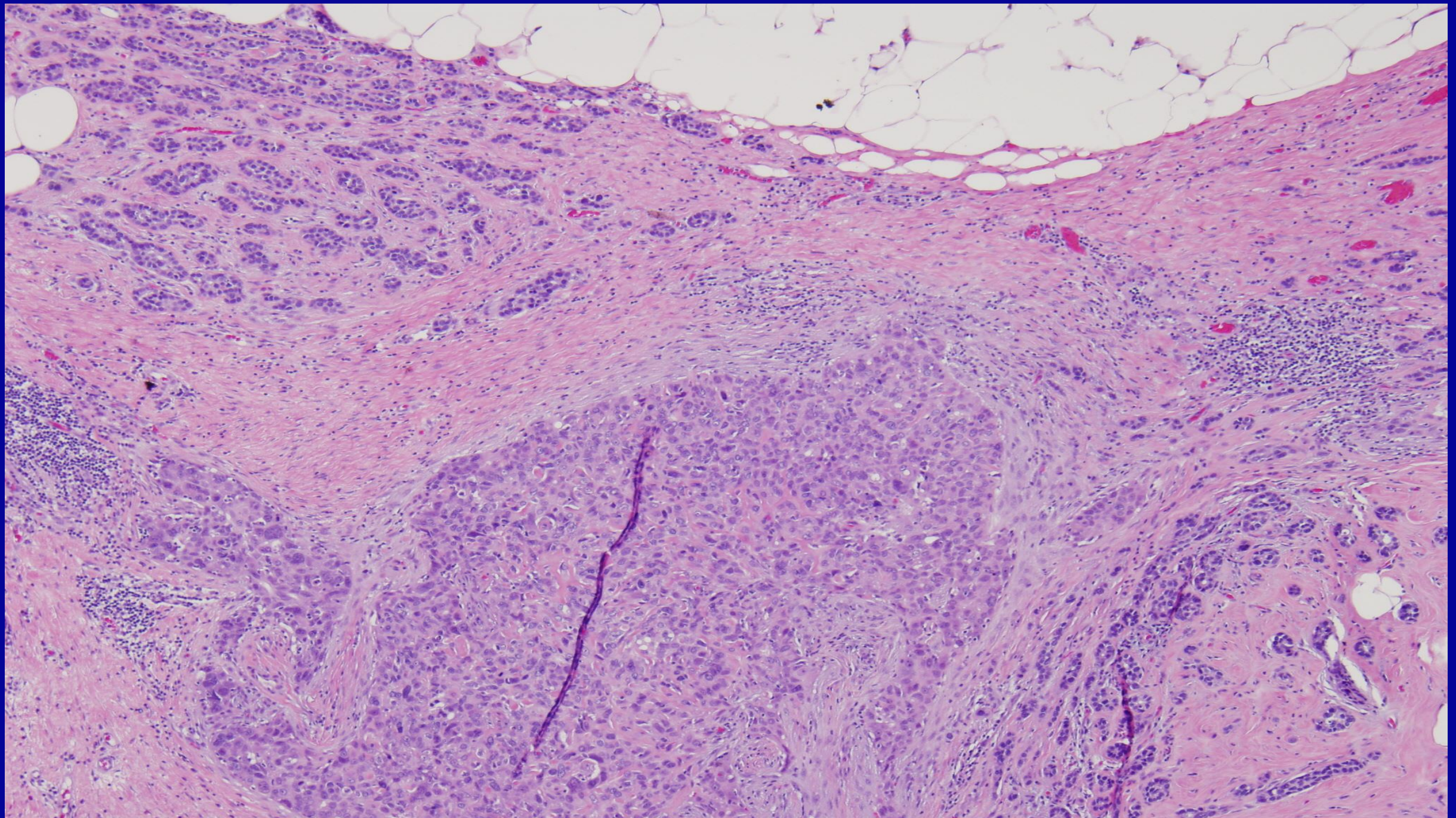
A: 3.6mm

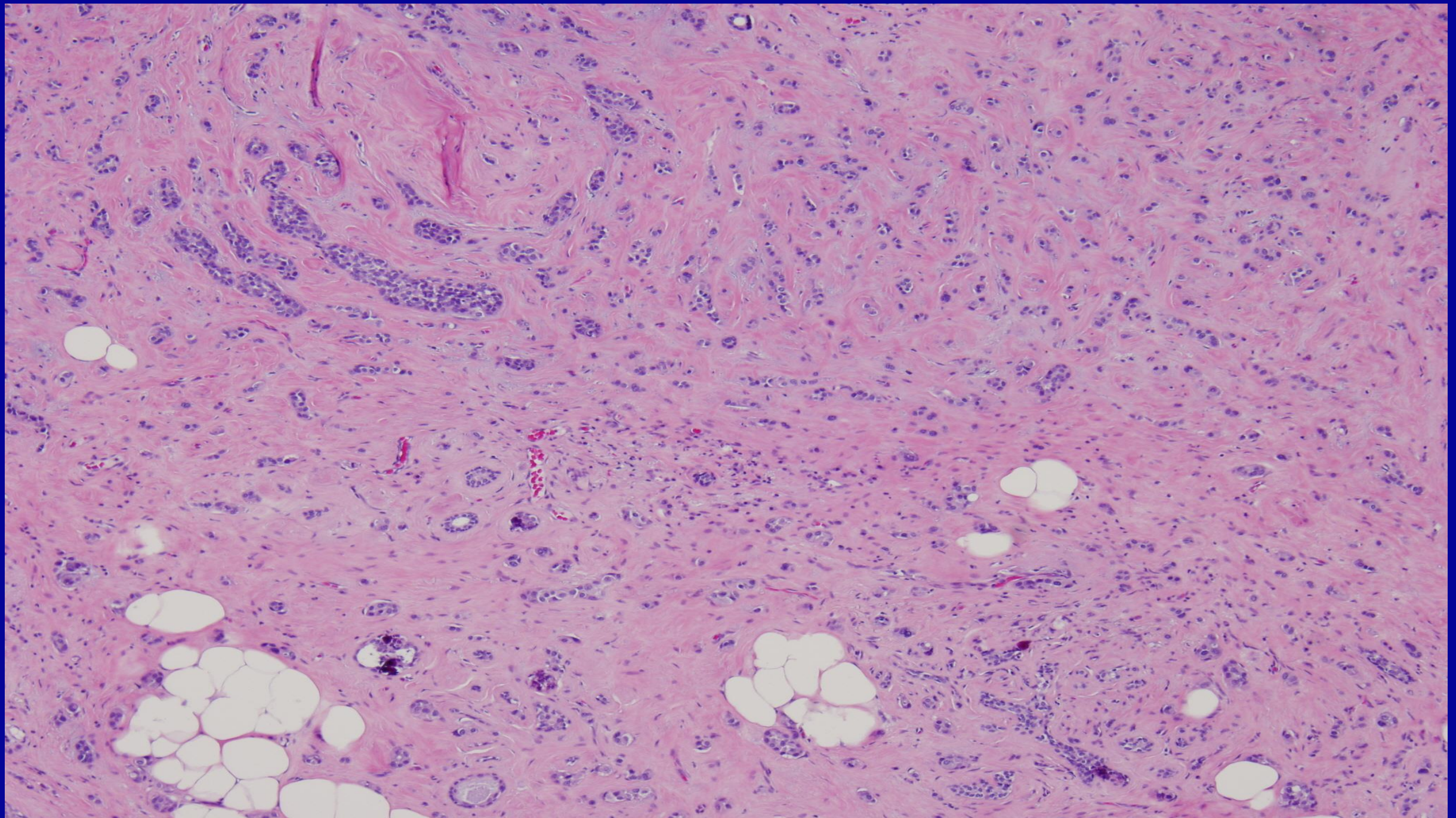


m









# Residual Cancer Burden Calculator

\*Values must be entered into all fields for the calculation results to be accurate.

## (1) *Primary Tumor Bed*

Primary Tumor Bed Area:  (mm) X  (mm)  
Overall Cancer Cellularity (as percentage of area):  (%)  
Percentage of Cancer That Is *in situ* Disease:  (%)

## (2) *Lymph Nodes*

Number of Positive Lymph Nodes:   
Diameter of Largest Metastasis:  (mm)

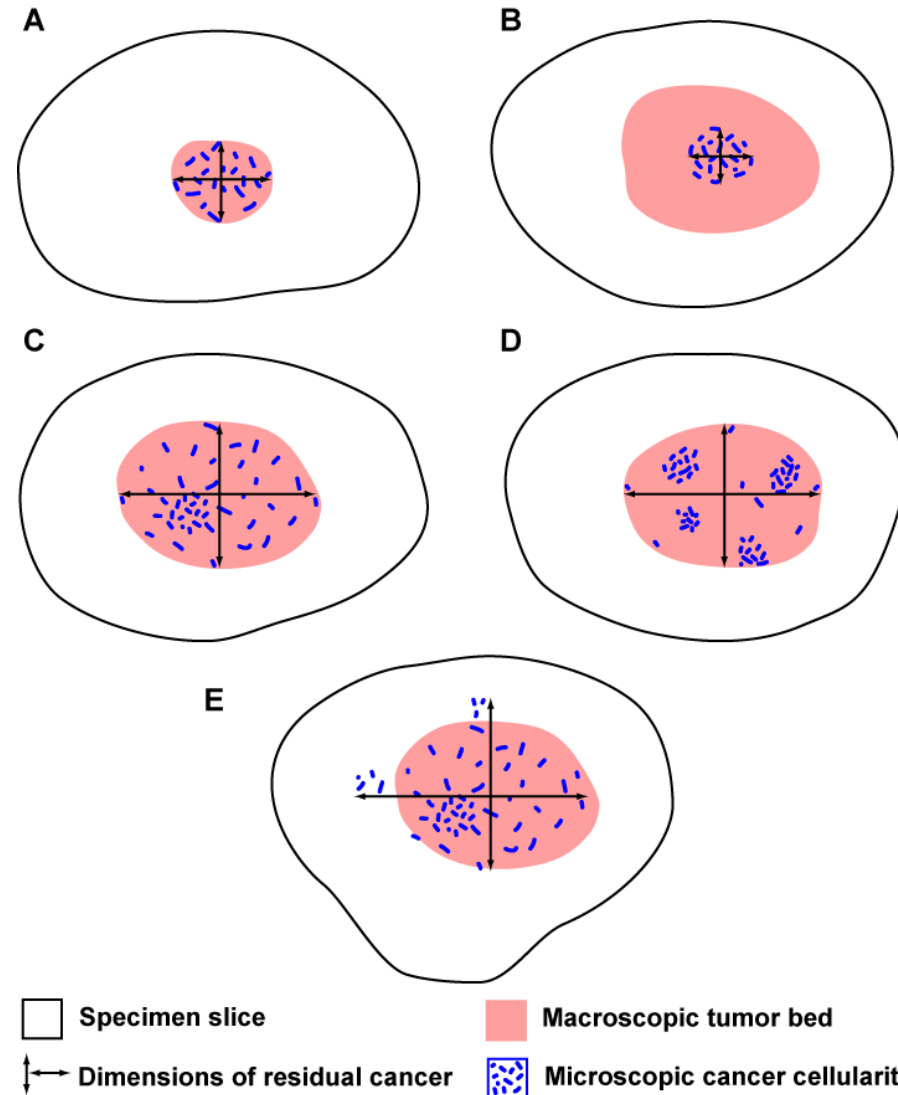
Reset

Calculate

Residual Cancer Burden:   
Residual Cancer Burden Class:

Google terms: residual cancer burden breast  
[www.mdanderson.org/breastcancer\\_RCB](http://www.mdanderson.org/breastcancer_RCB)

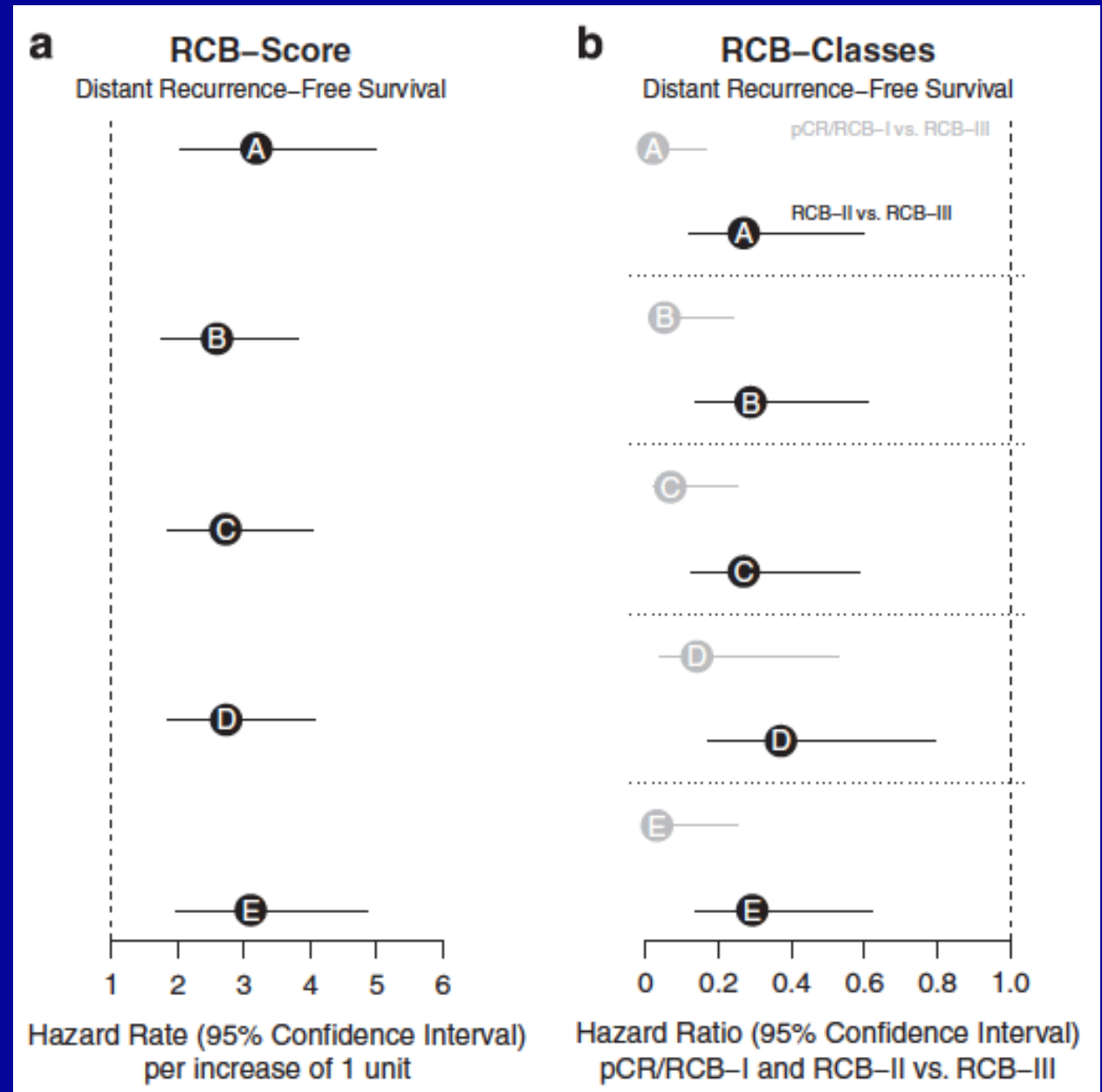
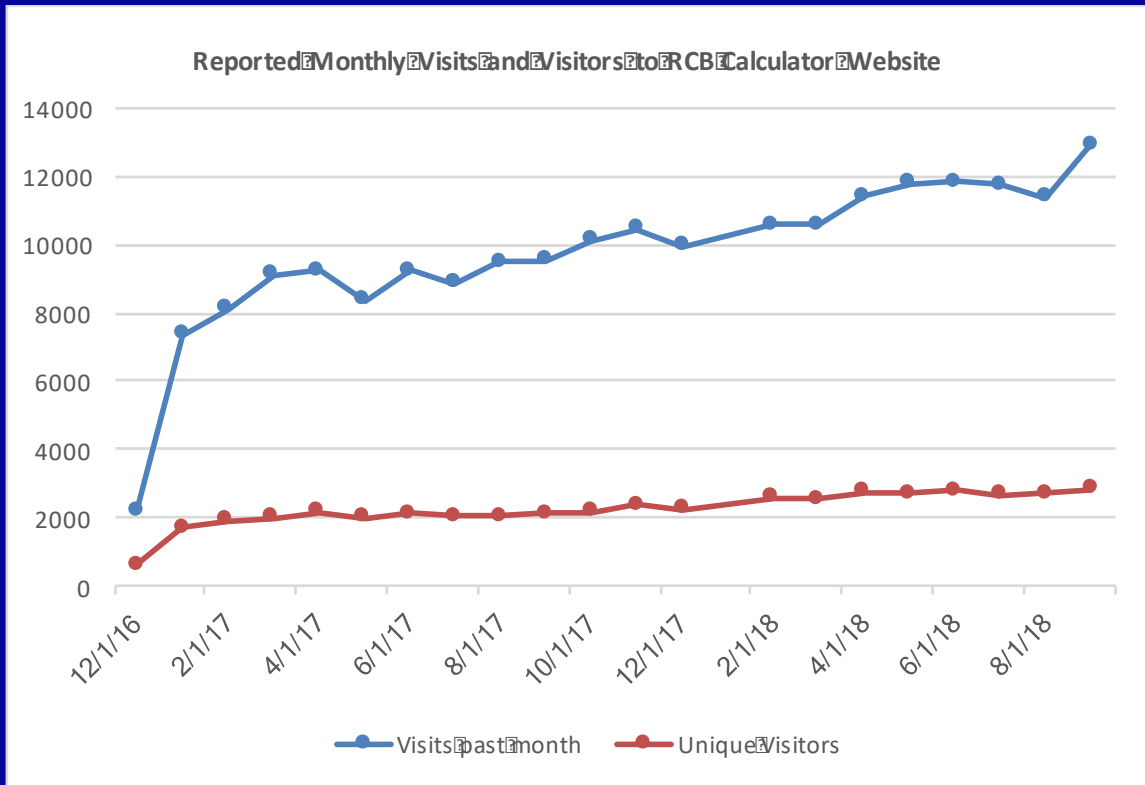
# Measure Residual Invasive Primary Tumor Bed: Macroscopic Findings Qualified By Microscopic Findings



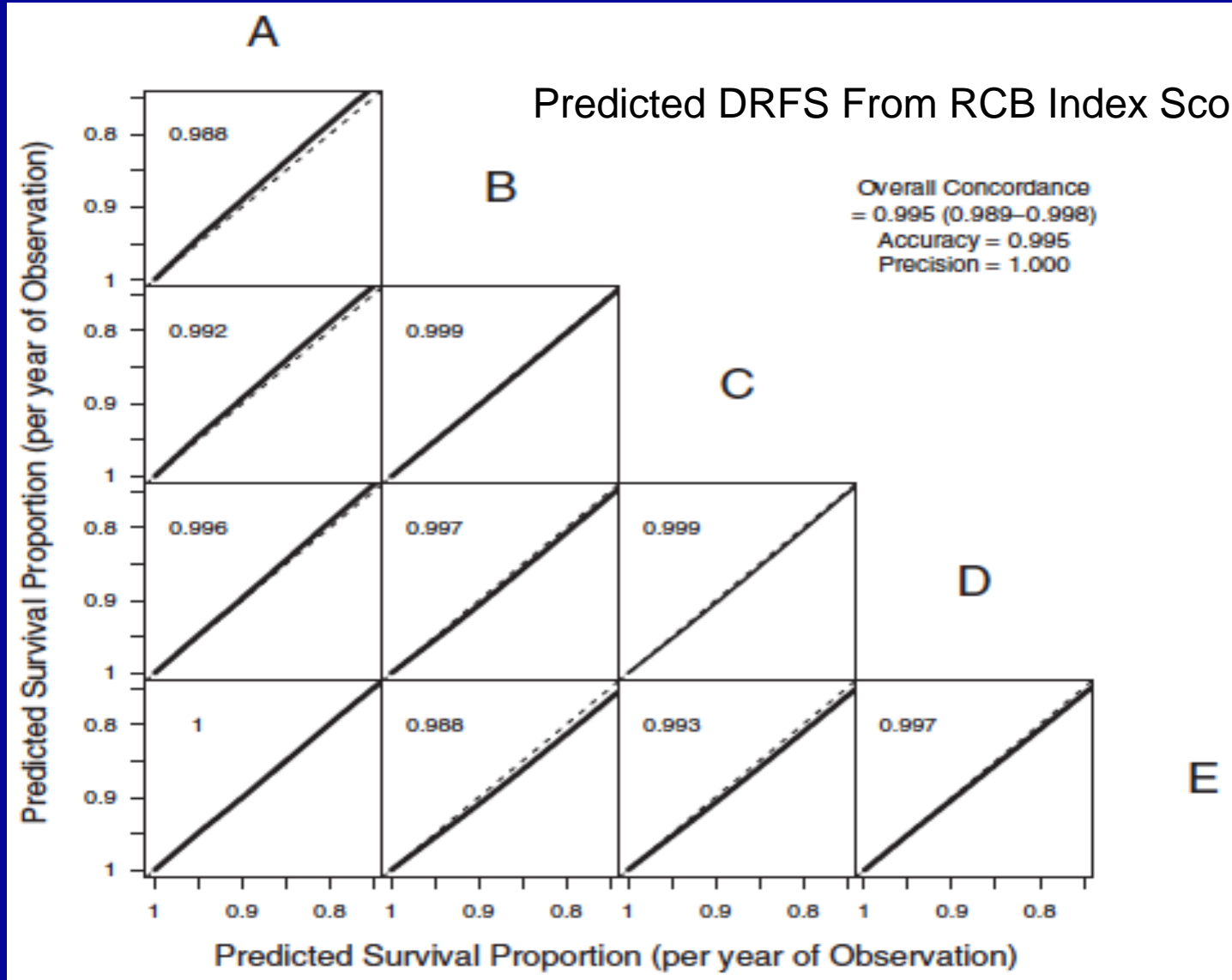
See downloadable protocol and illustrations at [www.mdanderson.org/breastcancer\\_RCB](http://www.mdanderson.org/breastcancer_RCB)



[www.mdanderson.org/breastcancer\\_RCB](http://www.mdanderson.org/breastcancer_RCB)



# Inter-Pathologist Concordance of Predicted DRFS



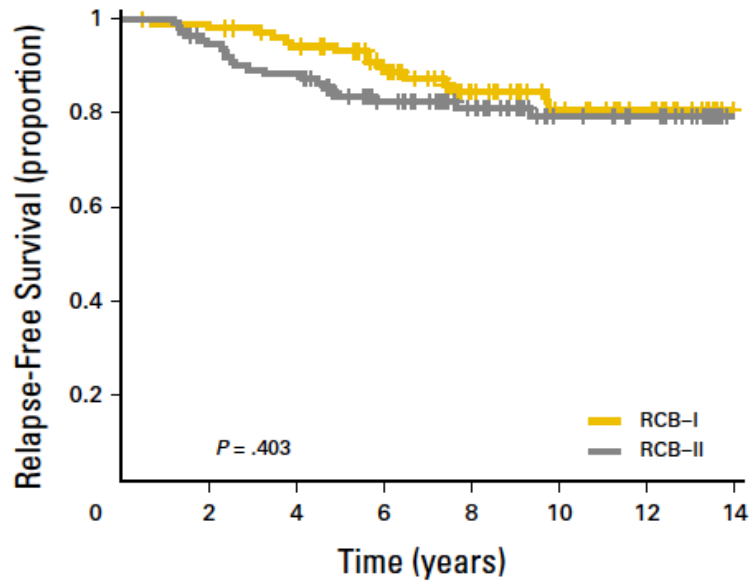
# RCB Classes Within *yp*-Stage Categories (RFS)

*yp*-Stage I

*yp*-Stage II

*yp*-Stage III

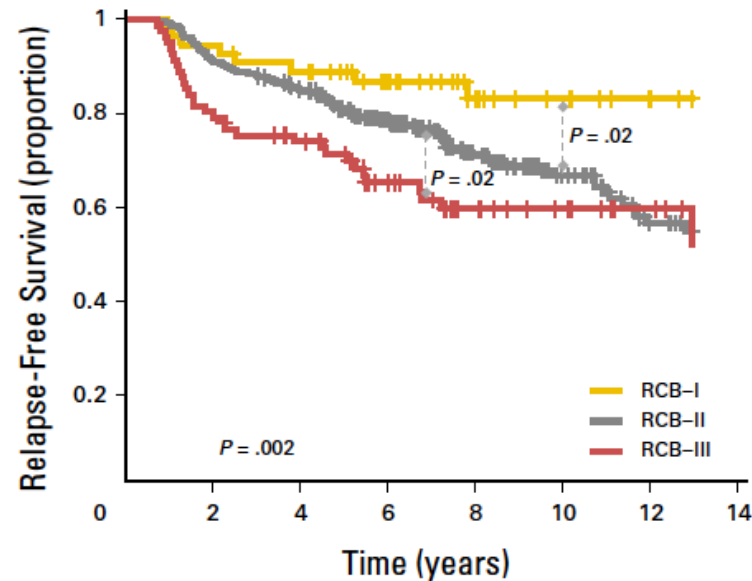
**A**



No. at risk:

RCB-I	106	103	96	76	53	40	30	13
RCB-II	112	104	97	76	58	39	34	16

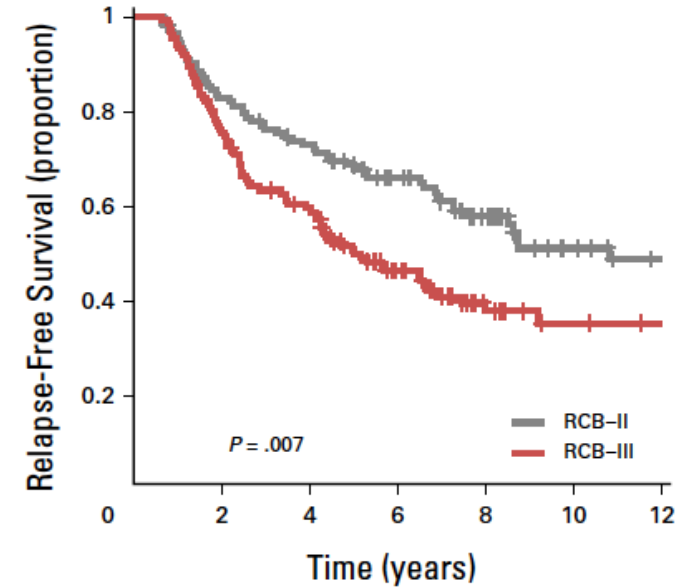
**B**



No. at risk:

RCB-I	55	51	47	36	21	16	11
RCB-II	282	257	233	180	105	62	40
RCB-III	81	64	56	39	39	19	11

**C**

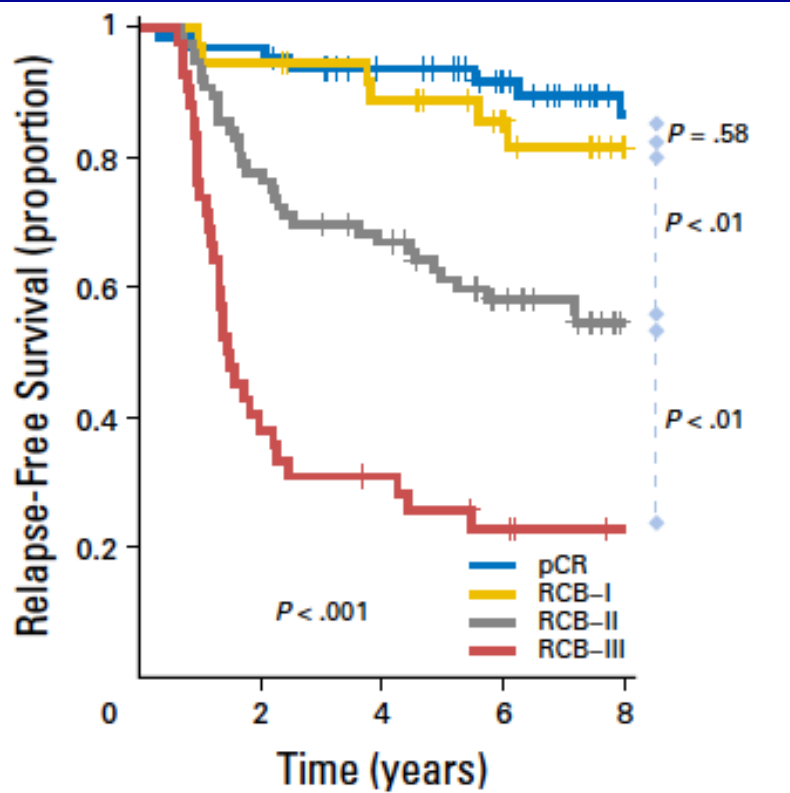


No. at risk:

RCB-II	124	101	87	69	49	26
RCB-III	135	101	77	46	22	13

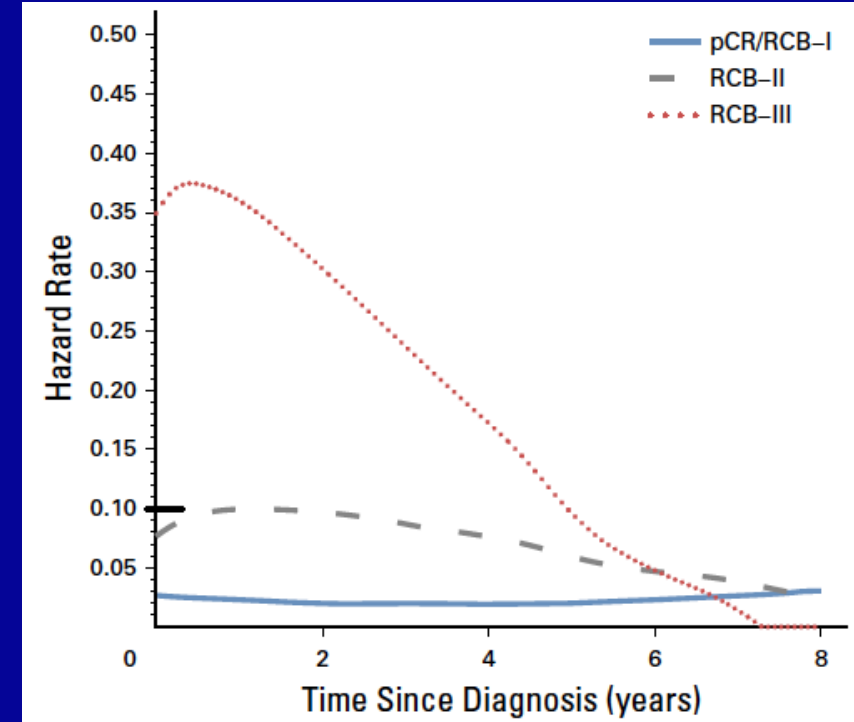
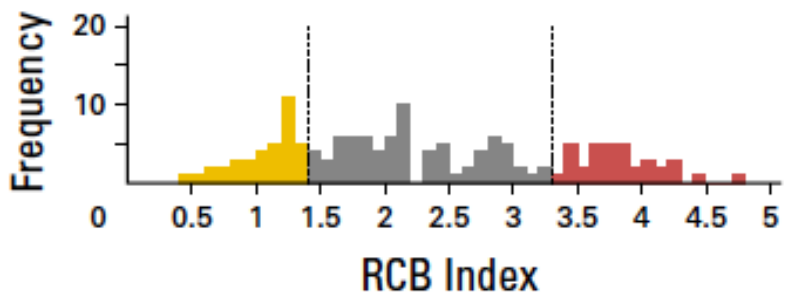
Figure 1. Symmans et al JCO 2017;35:1049-60

# TNBC



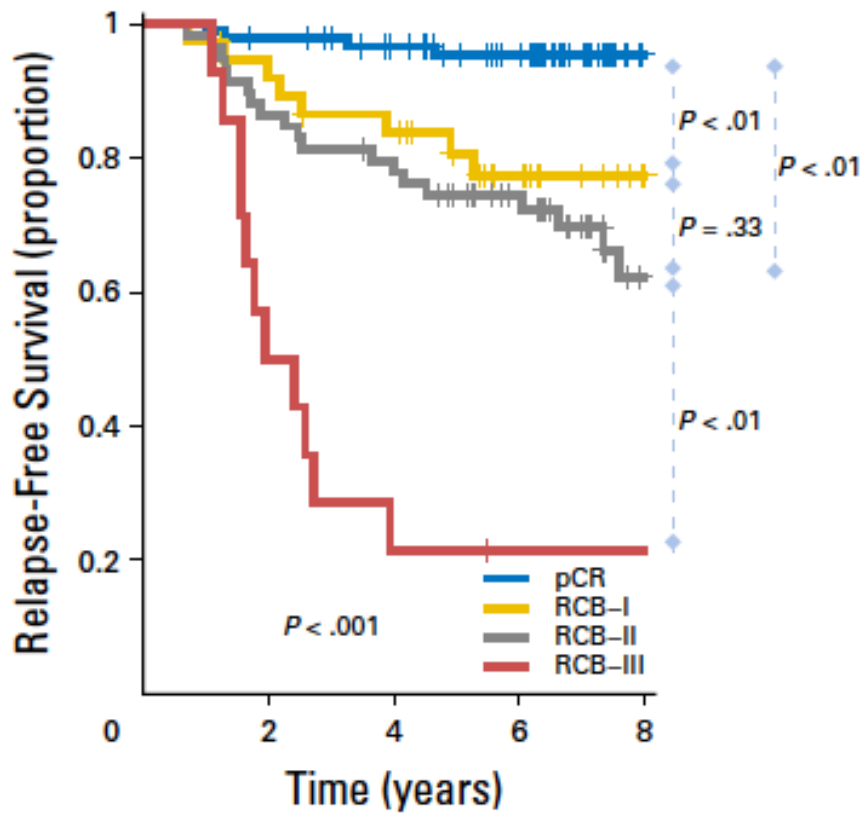
No. at risk:

	0	2	4	6	8
pCR	64	61	53	43	29
RCB-I	37	35	31	23	13
RCB-II	76	59	49	36	23
RCB-III	42	16	12	8	5



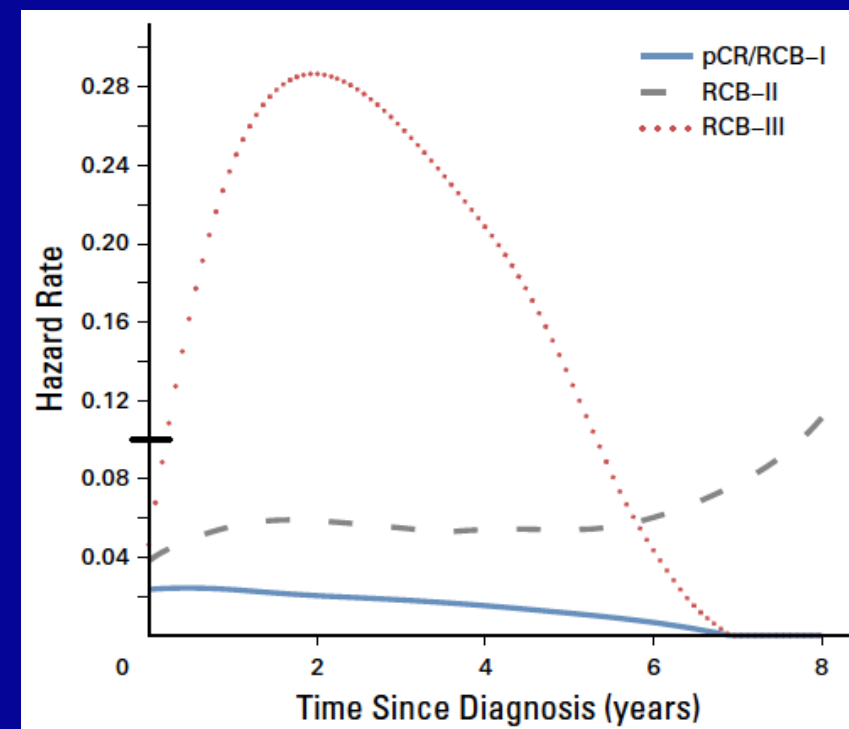
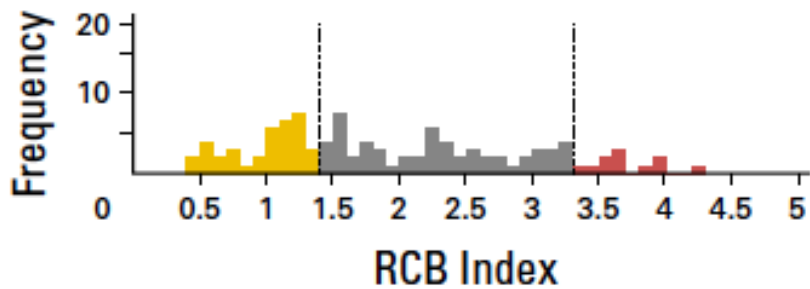
Multivariate Cox Regression				
TNBC Cases in MDACC RCB Study (n = 219)				
	HR	lower 95	upper 95	p =
Age	0.99	0.97	1.01	NS
Grade 3	0.96	0.54	1.69	NS
c-Stage III	1.19	0.73	1.96	NS
<b>RCB index</b>	<b>1.89</b>	<b>1.56</b>	<b>2.29</b>	<b>&lt;0.01</b>

# HER2+ (H+T/FEC)



No. at risk:

	0	2	4	6	8
pCR	92	89	82	70	37
RCB-I	38	34	30	19	6
RCB-II	59	51	45	34	14
RCB-III	14	7	3	2	2

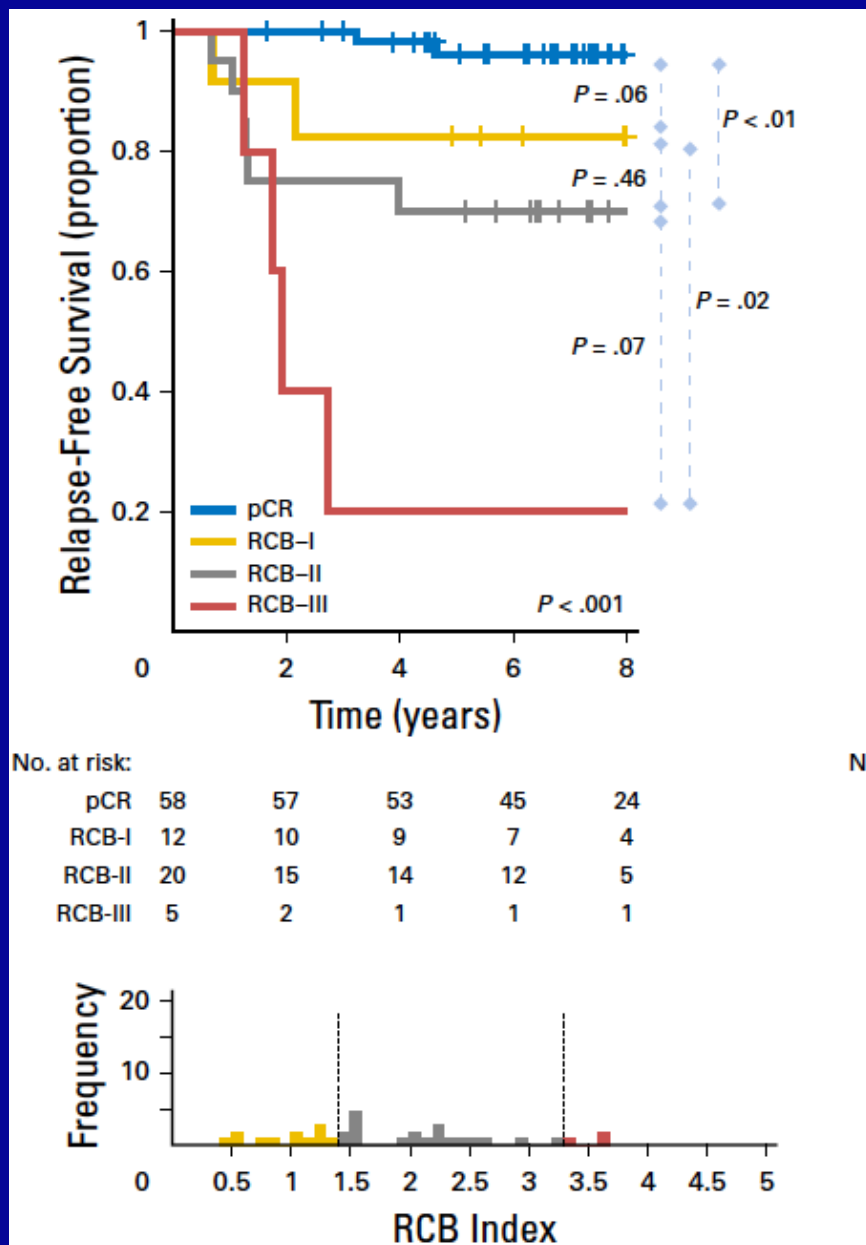


## Multivariate Cox Regression

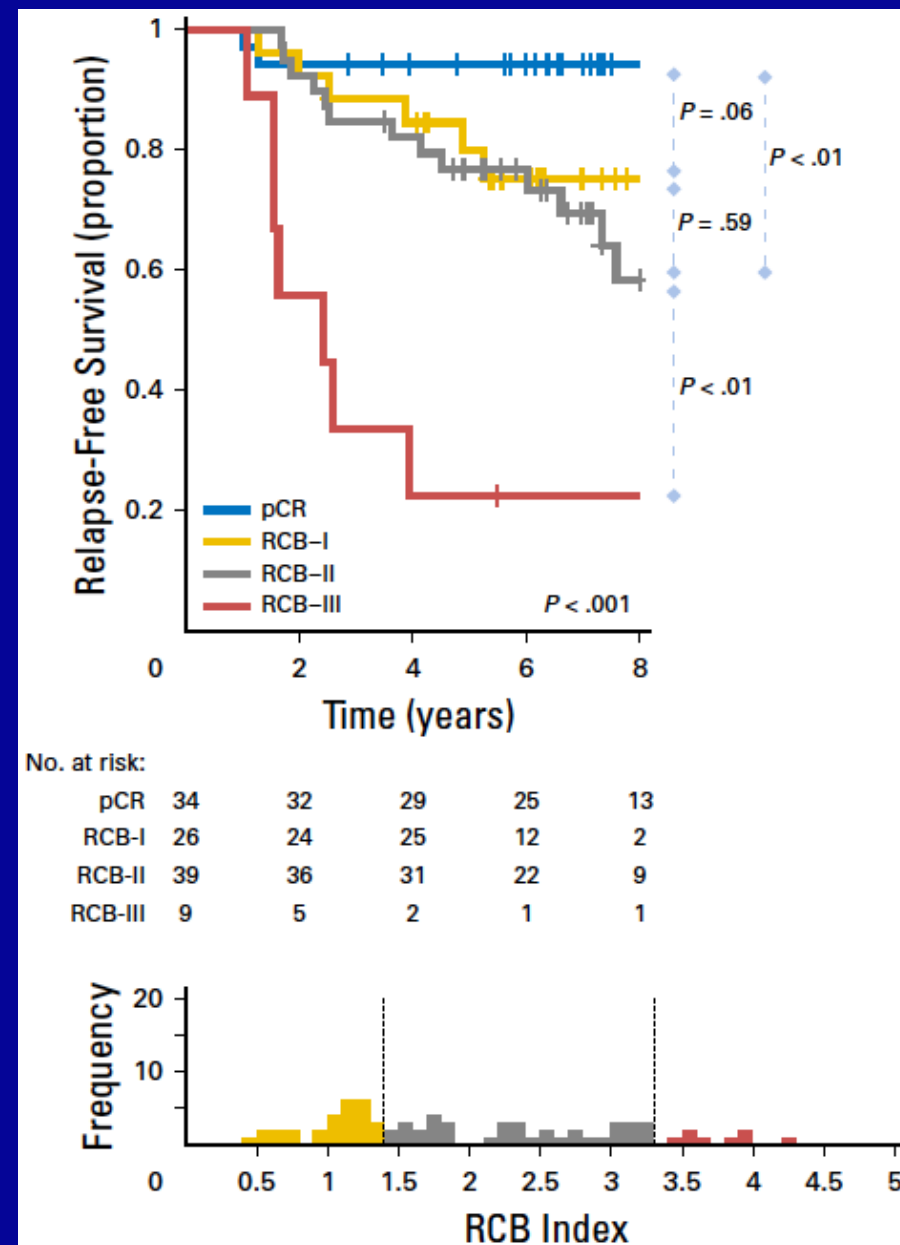
### HER2+ Cases in MDACC RCB Study (n = 203)

	HR	lower 95	upper 95	p =
Age	1.02	0.99	1.06	NS
Grade 3	0.56	0.29	1.10	NS
c-Stage III	1.33	0.70	2.55	NS
Multifocal	1.48	0.72	3.07	NS
pCR	0.34	0.08	1.35	NS
<b>RCB index</b>	<b>1.80</b>	<b>1.26</b>	<b>2.59</b>	<b>&lt;0.01</b>

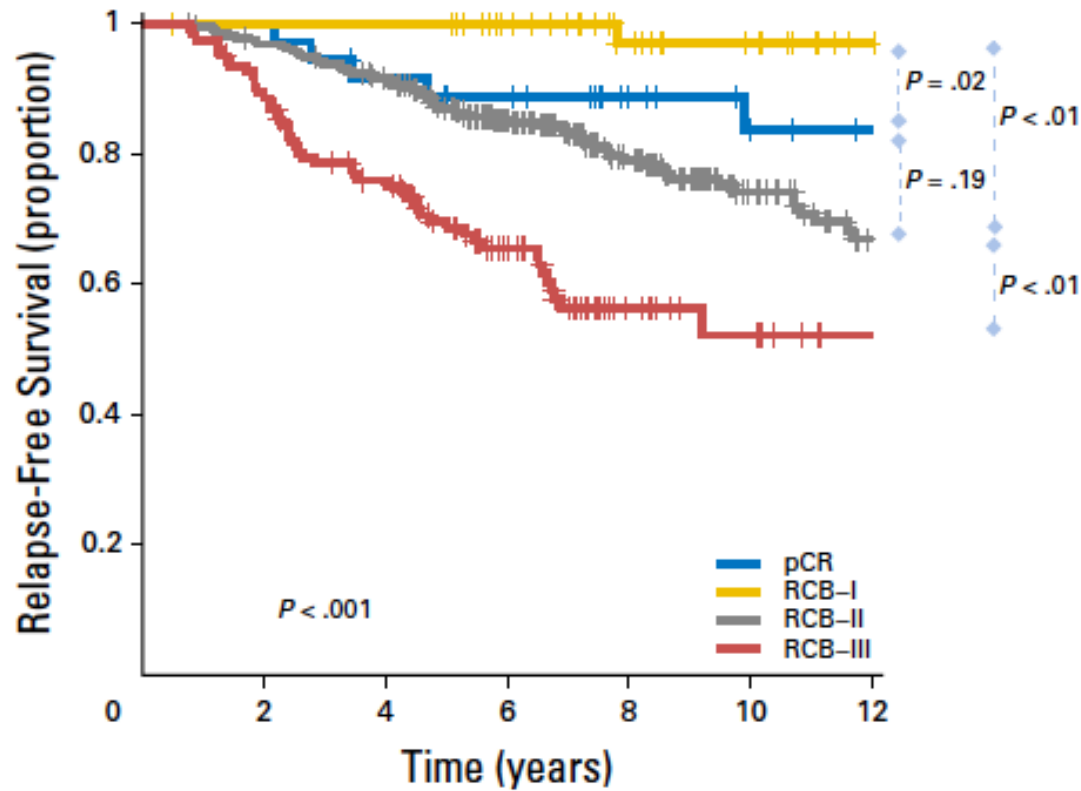
### HER2+/HR- (H+T/FEC)



### HER2+/HR+ (H+T/FEC)

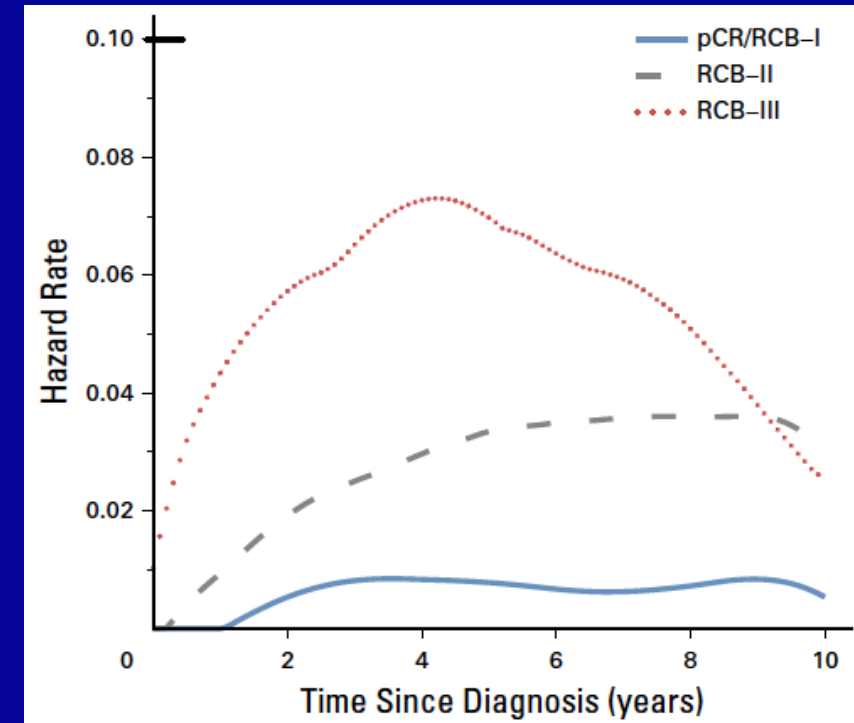
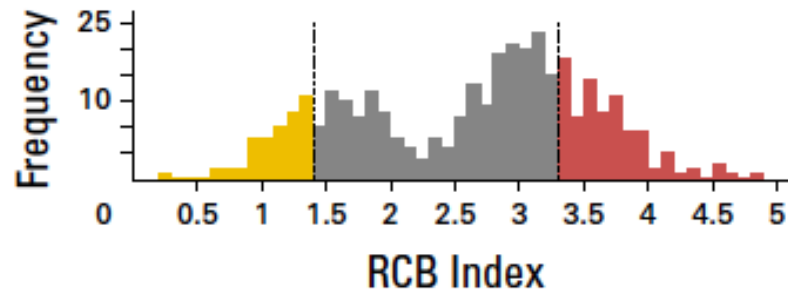


# HR+/HER2-



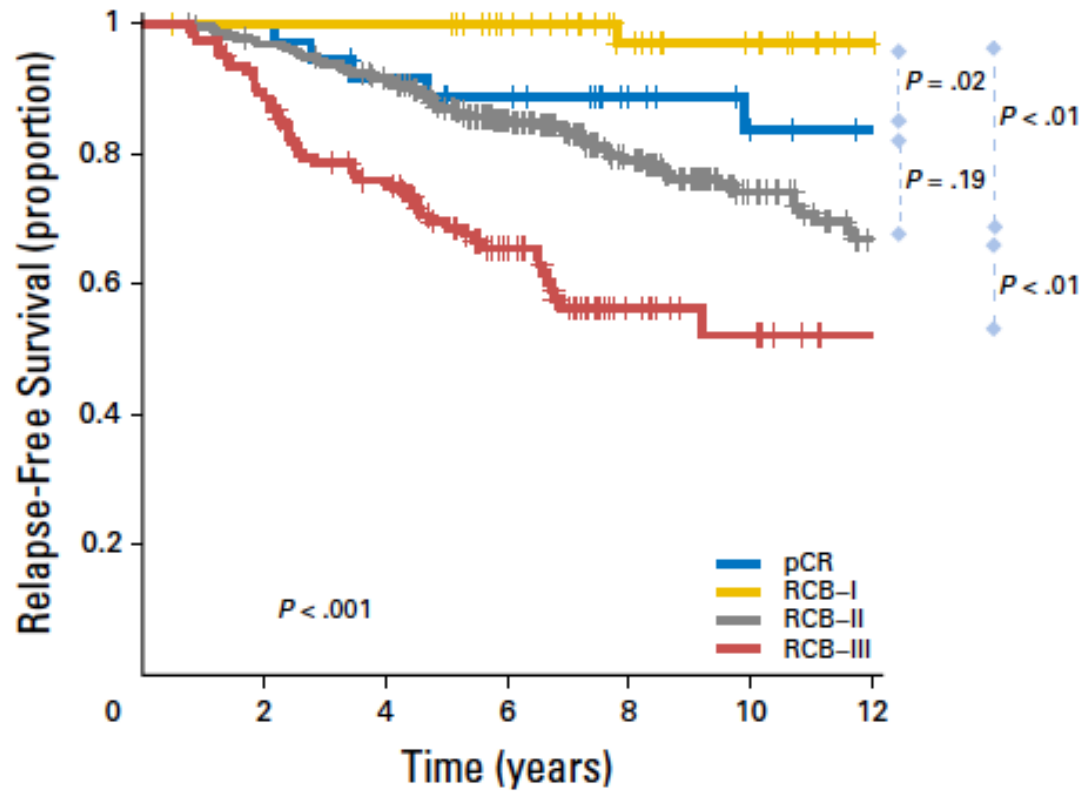
No. at risk:

	0	2	4	6	8	10	12
pCR	36	36	32	29	21	17	14
RCB-I	52	51	51	44	32	26	15
RCB-II	290	278	258	199	127	73	48
RCB-III	123	108	88	55	22	13	5



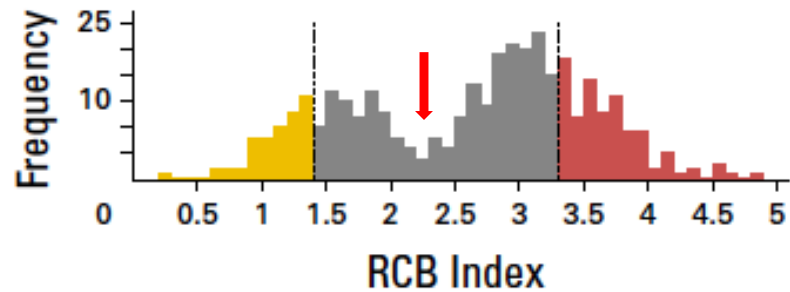
Multivariate Cox Regression				
HR+/HER2- Cases in MDACC RCB Study (n = 501)				
	HR	lower 95	upper 95	p =
Age	0.99	0.97	1.01	NS
Grade 3	1.23	0.82	1.84	NS
<b>c-Stage III</b>	<b>2.53</b>	<b>1.73</b>	<b>3.71</b>	<b>&lt;0.01</b>
<b>RCB index</b>	<b>1.95</b>	<b>1.57</b>	<b>2.41</b>	<b>&lt;0.01</b>

# HR+/HER2-

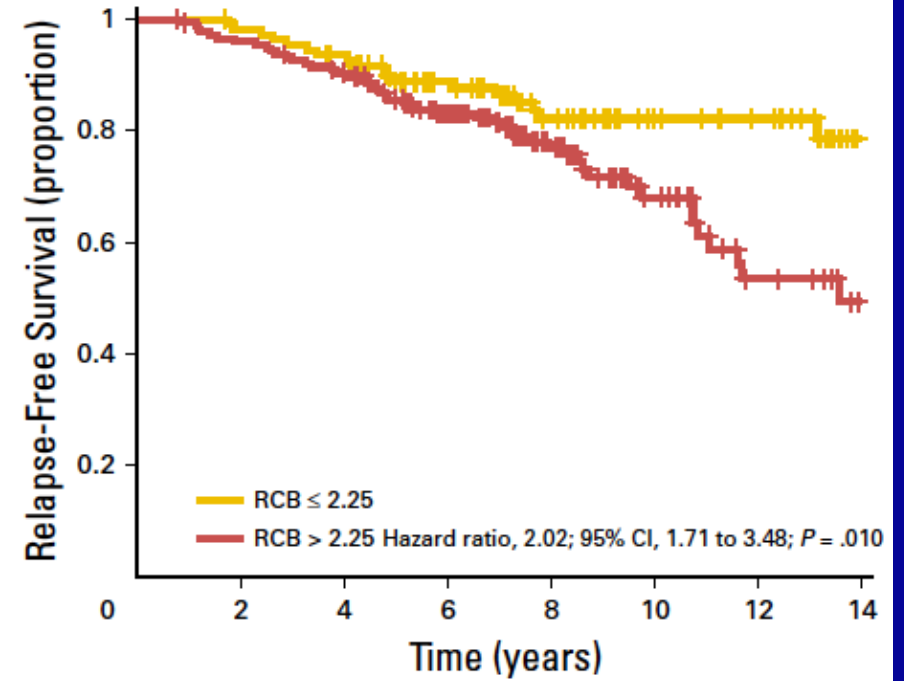


No. at risk:

	0	2	4	6	8	10	12
pCR	36	36	32	29	21	17	14
RCB-I	52	51	51	44	32	26	15
RCB-II	290	278	258	199	127	73	48
RCB-III	123	108	88	55	22	13	5



## RCB-II Only



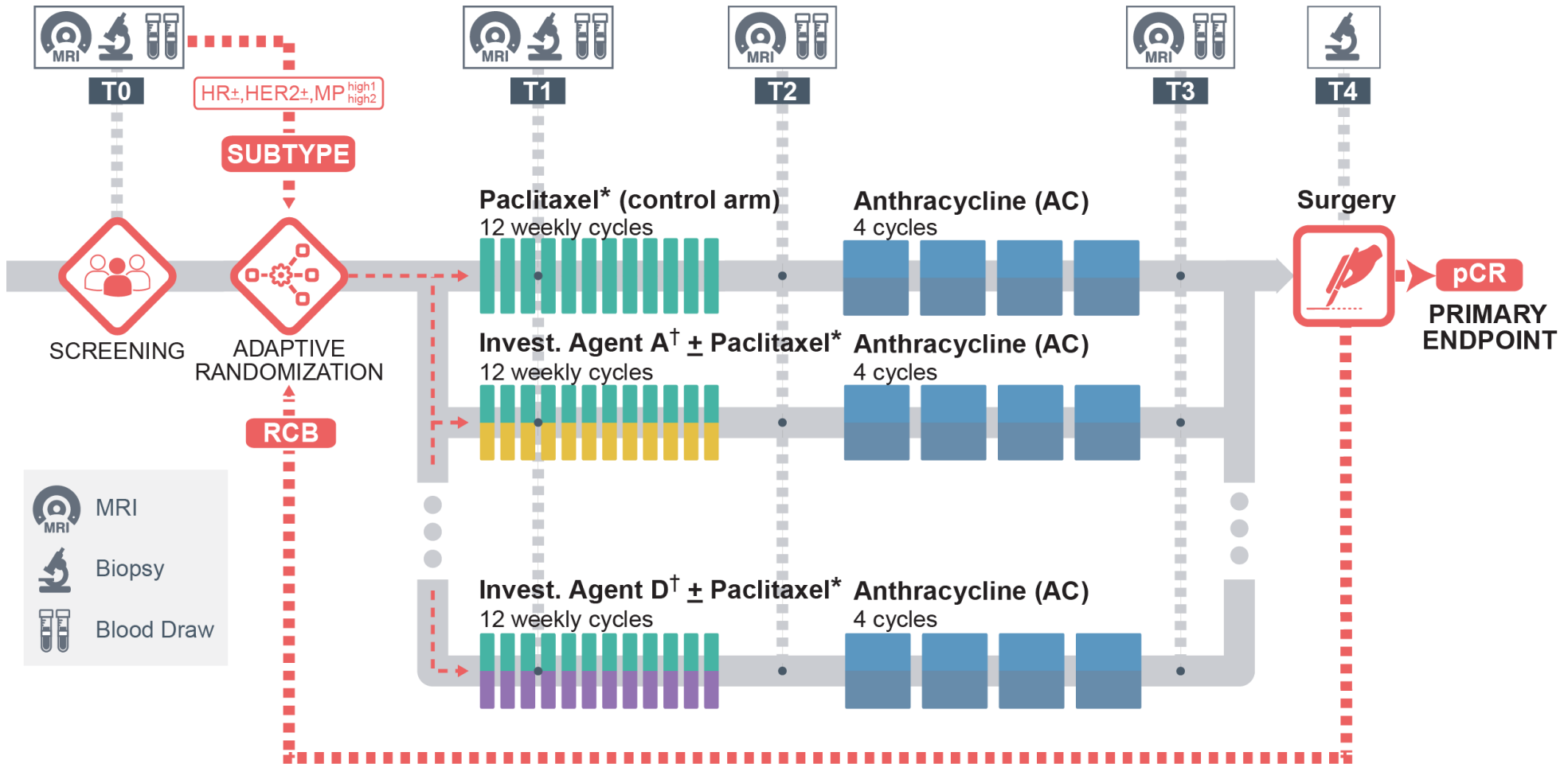
No. at risk:

	0	2	4	6	8	10	12	14
RCB $\leq$ 2.25	111	108	101	79	55	37	30	
RCB $>$ 2.25	179	170	170	120	72	36	18	



# Independent Validation in Prospective Clinical Trials

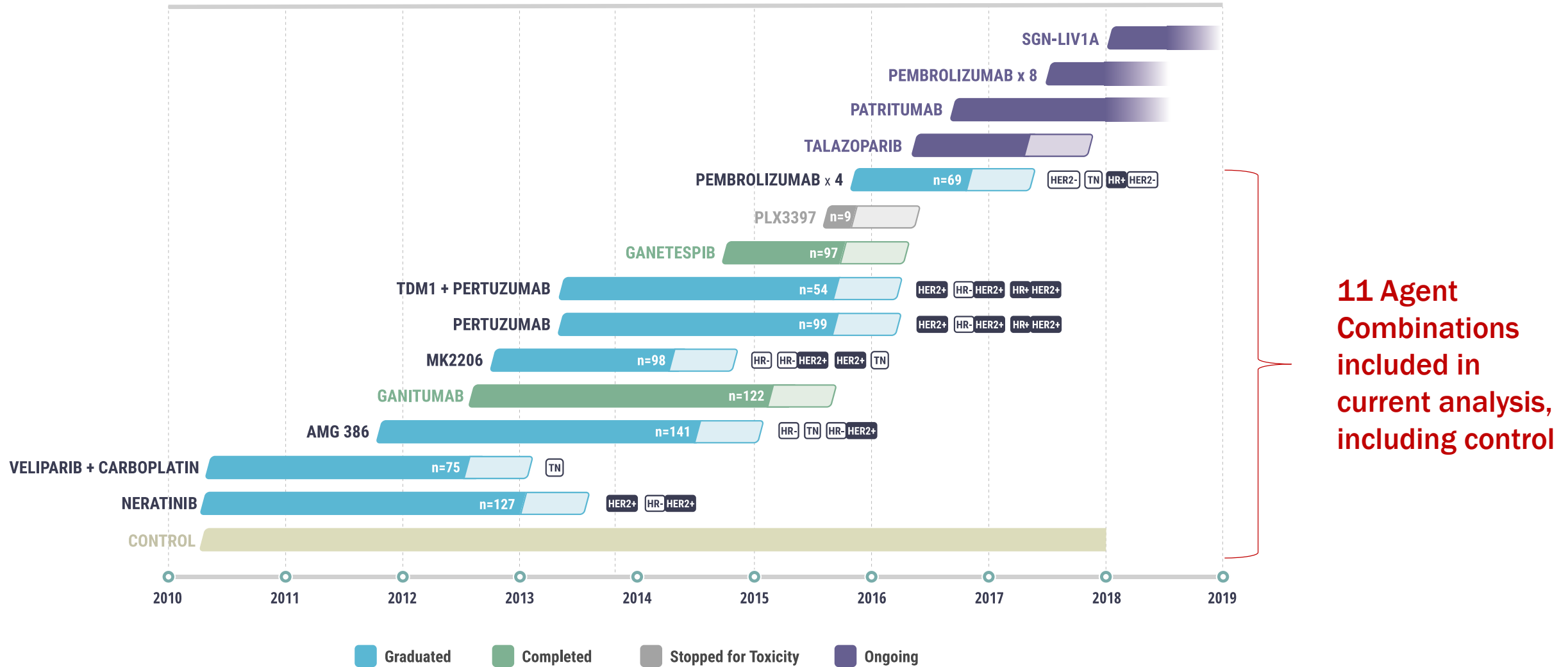
# I-SPY2 Trial: Adaptive Randomization



\*Patients who are HER2+ may also receive trastuzumab (Herceptin)

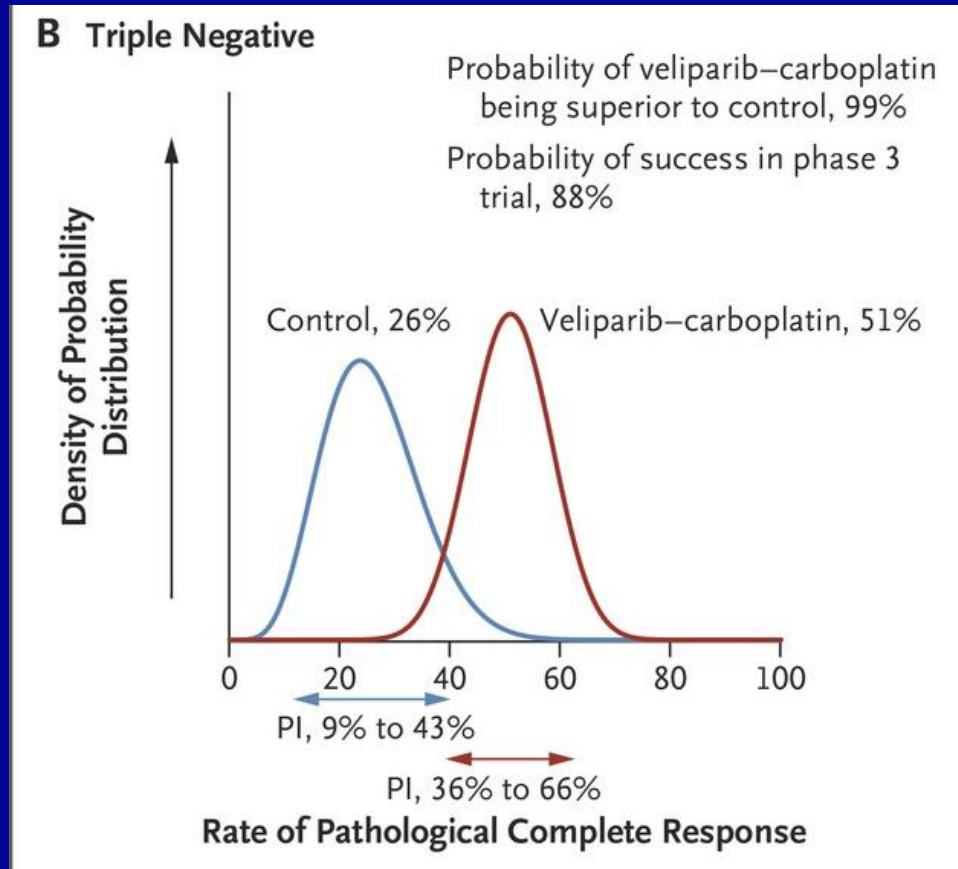
†An investigational combination of one or more agents may be used to replace all or some of the standard therapy

# I-SPY2 Trial: Agent Timeline



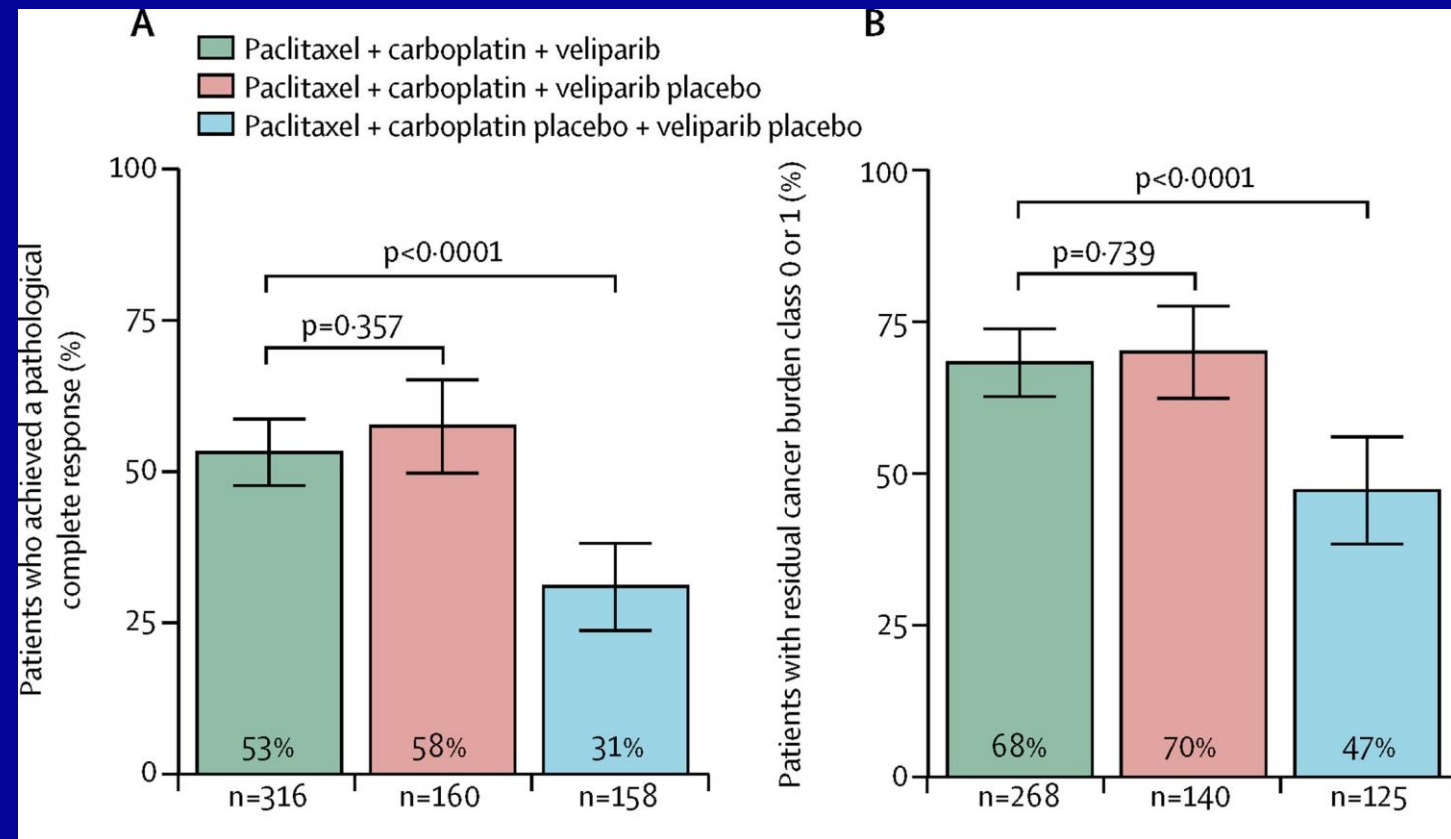
# Implementation In Randomized Clinical Trials

I-SPY2 : Adaptive Randomization Phase 2.  
Addition of Veliparib and Carboplatin to  
Weekly Paclitaxel



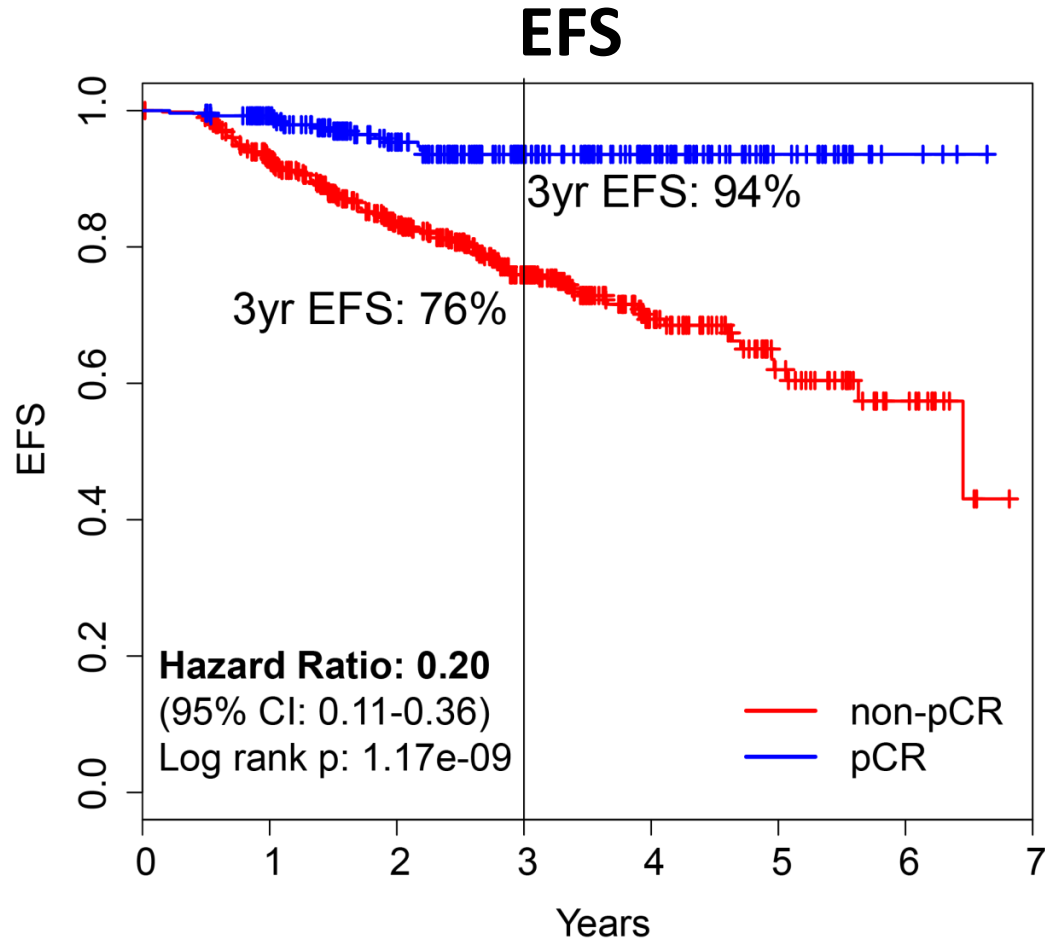
Rugo H, et al. *NEJM* 2016;375:23-34

Brightness Trial (AFT8): Randomized Phase 3.  
Addition of Veliparib and Carboplatin, or Carboplatin to Weekly  
Paclitaxel

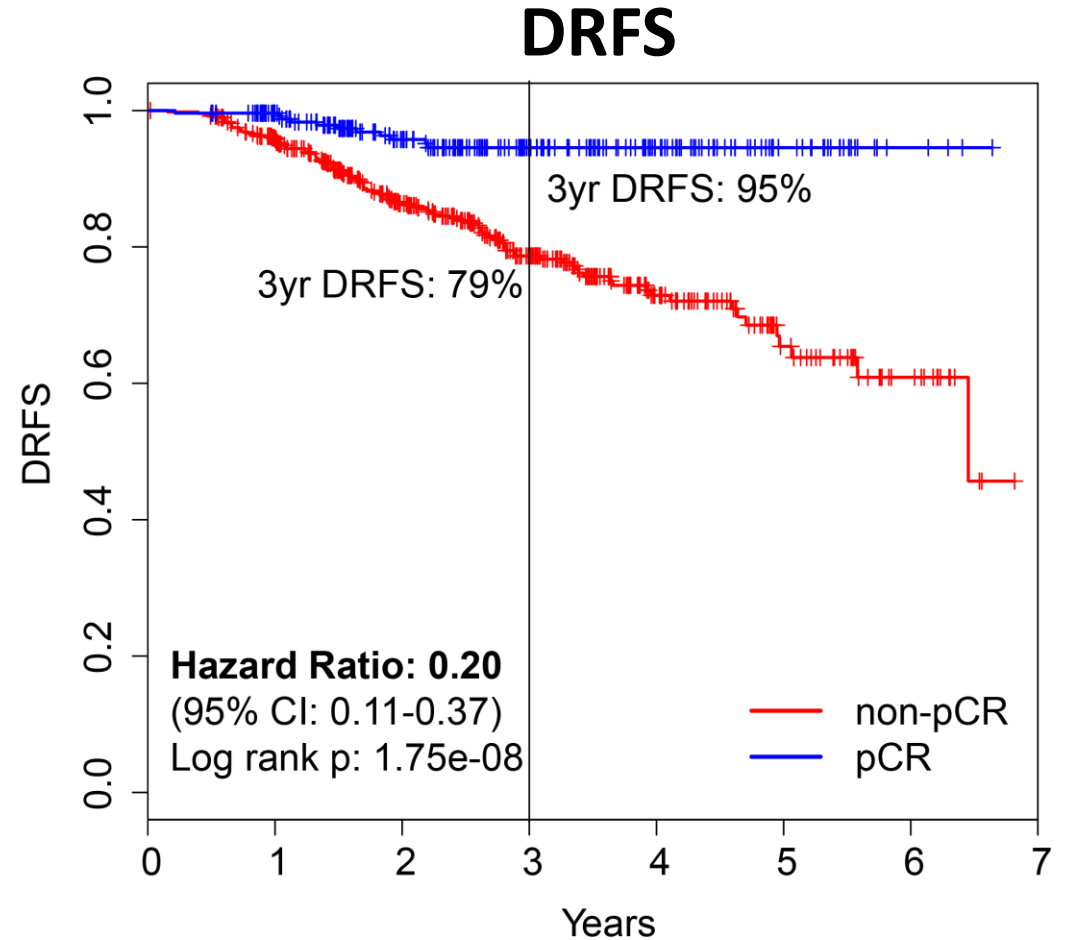


Loibl S, et al. *Lancet Oncol* 2018;19:497-509

# pCR is a highly significant predictor of EFS and DRFS

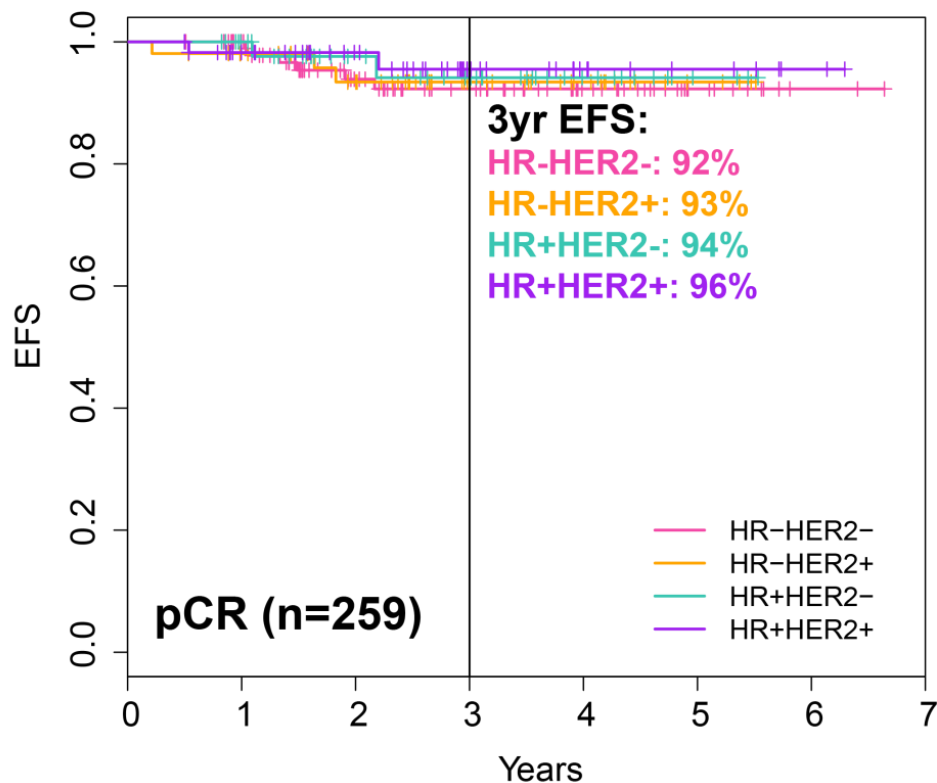


Number at Risk		0	1	2	3	4	5	6	7
non-pCR	487	418	288	186	89	40	13	0	0
pCR	259	232	166	109	59	23	4	0	0

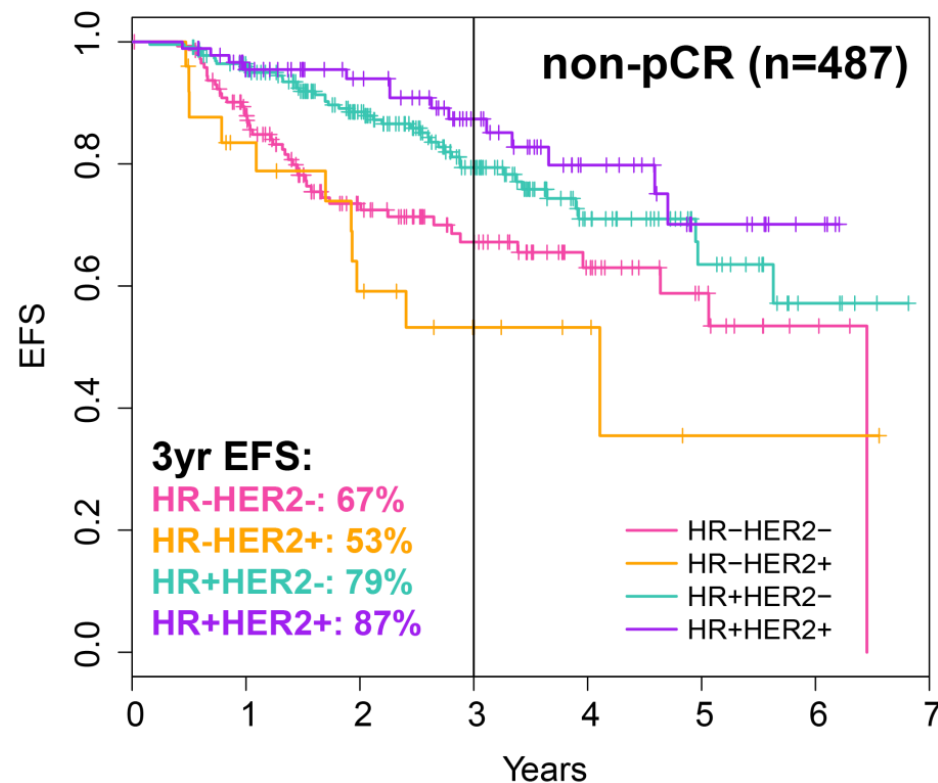


Number at Risk		0	1	2	3	4	5	6	7
non-pCR	487	430	295	193	95	41	14	0	0
pCR	259	233	167	110	60	24	4	0	0

# EFS by pCR & non-pCR, by Subtype

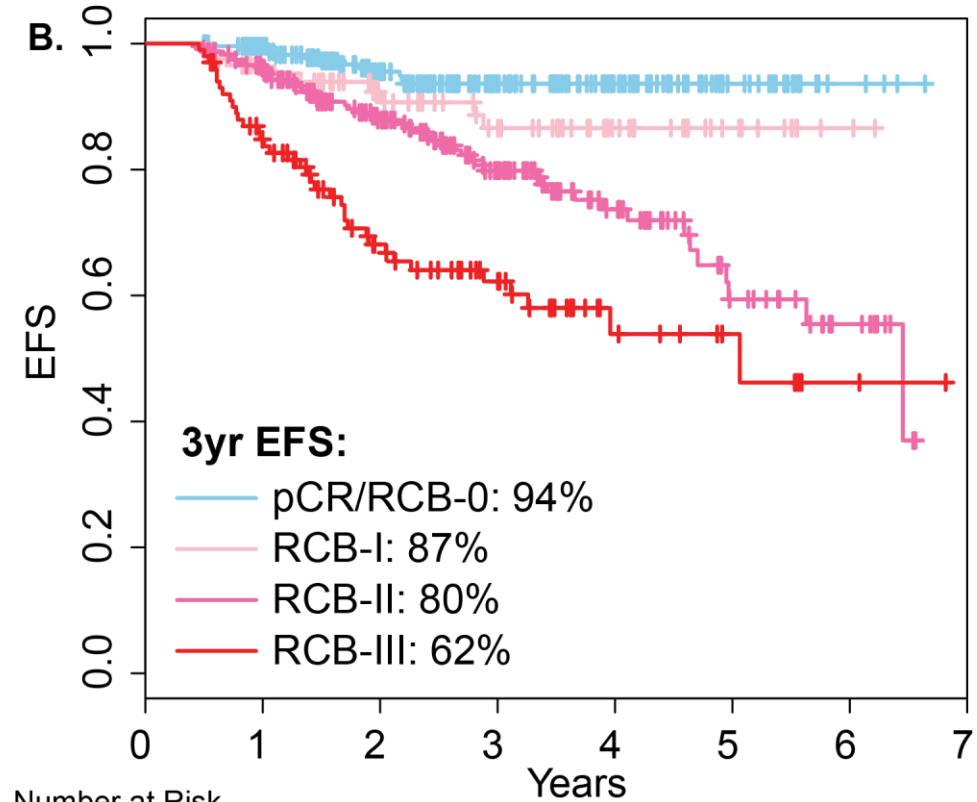
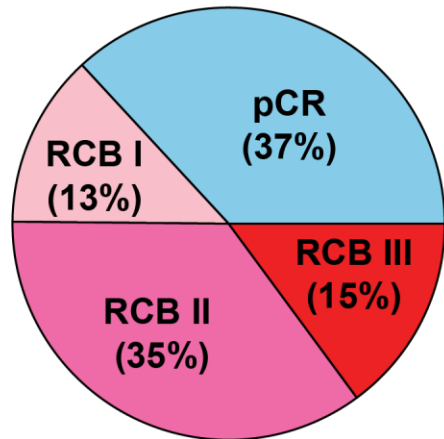
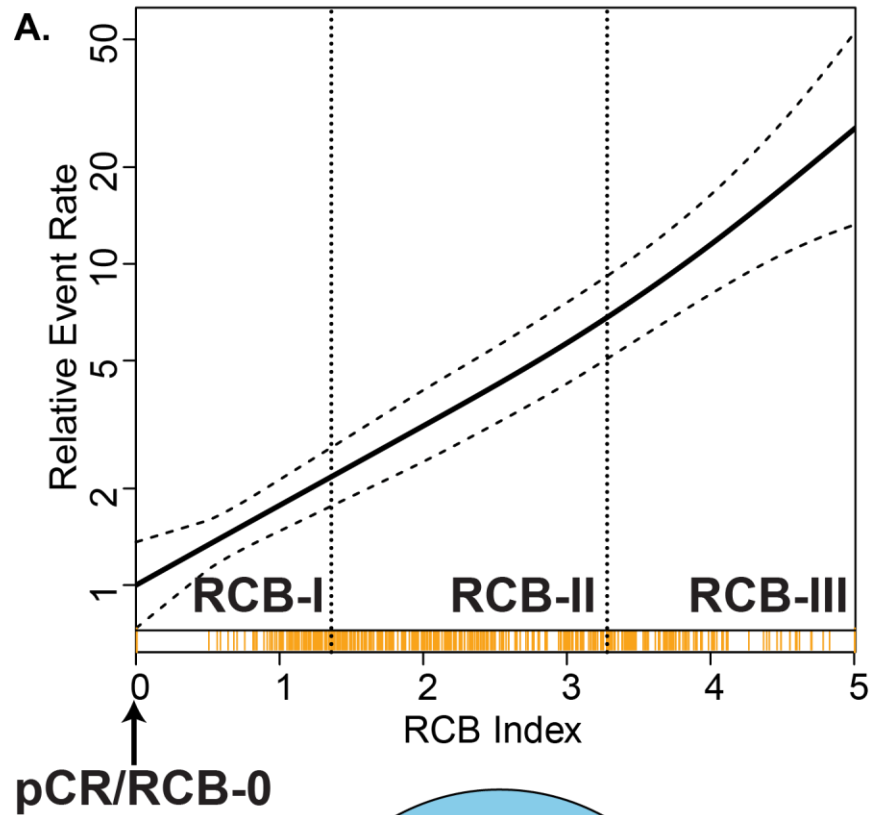


Number at Risk	0	1	2	3	4	5	6	7
HR-HER2-	100	92	61	44	25	10	2	0
HR-HER2+	52	47	39	23	13	4	0	0
HR+HER2-	49	44	29	22	11	3	0	0
HR+HER2+	58	49	37	20	10	6	2	0



Number at Risk	0	1	2	3	4	5	6	7
HR-HER2-	145	118	70	48	24	12	3	0
HR-HER2+	25	18	12	7	4	1	1	0
HR+HER2-	226	204	144	89	39	17	5	0
HR+HER2+	91	78	62	42	22	10	4	0

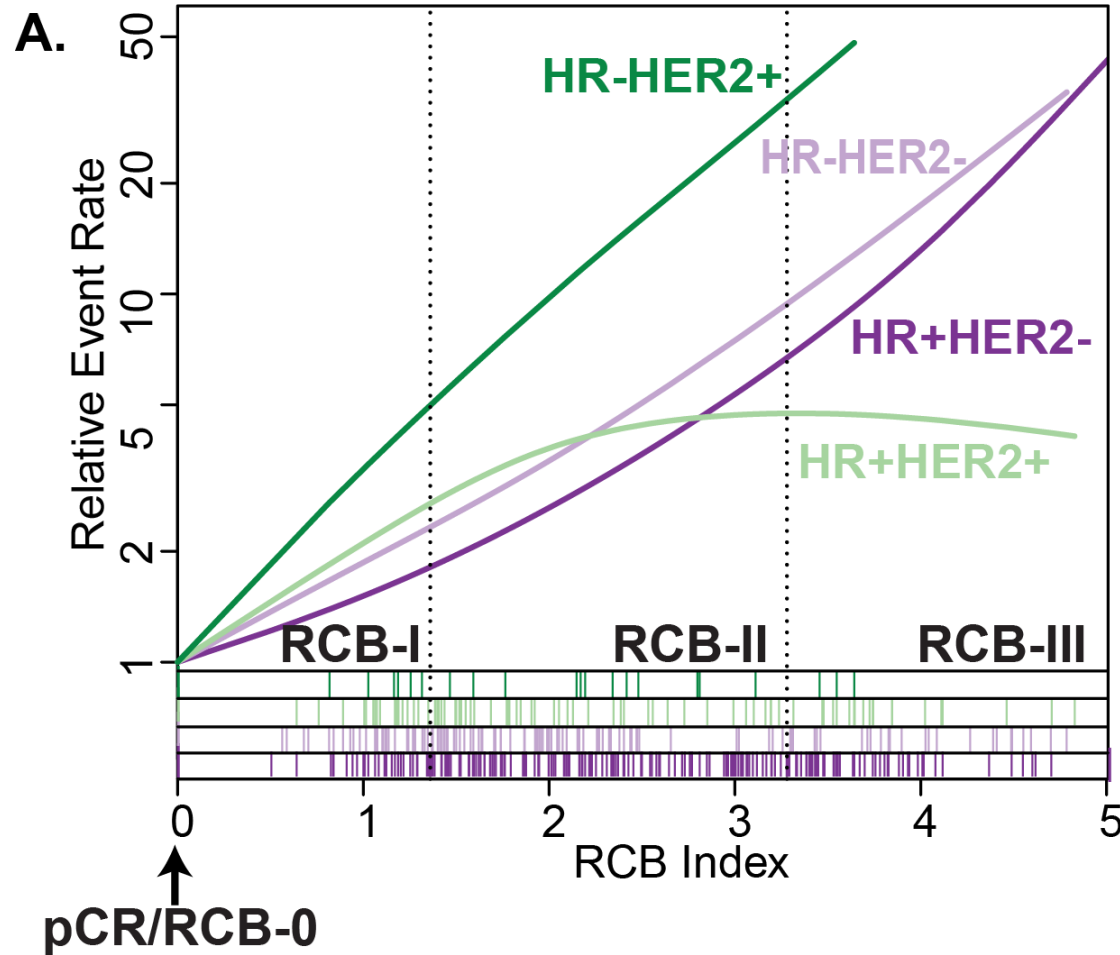
# Residual Cancer Burden in I-SPY2 (Site Pathologists' Reporting)



Number at Risk

	0	1	2	3	4	5	6	7
0	250	222	156	104	56	22	4	0
I	88	75	55	41	23	11	2	0
II	239	214	144	92	48	21	9	0
III	101	79	51	35	12	7	2	0

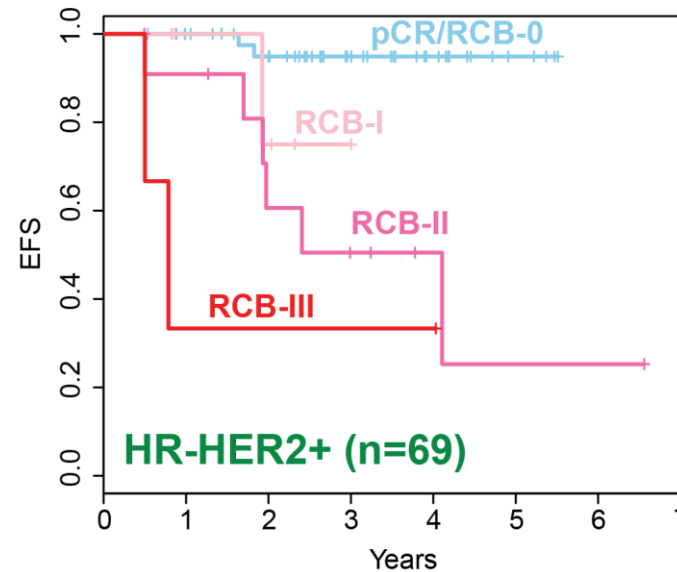
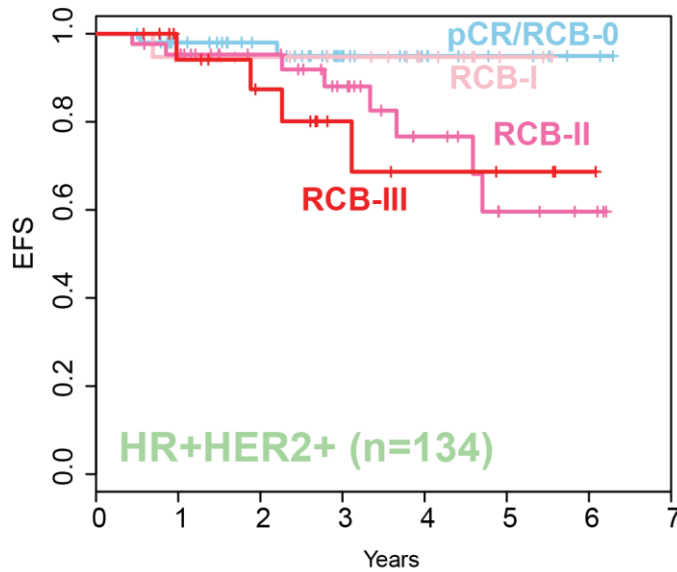
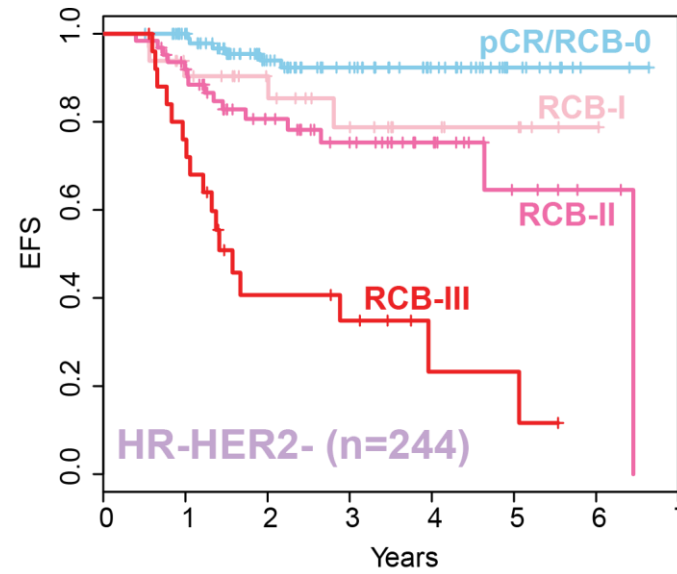
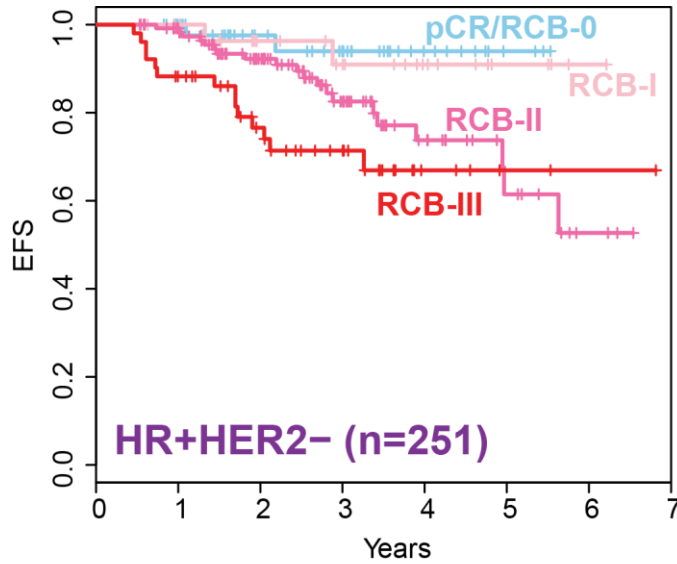
# Relative Risk According to RCB Index Within Subtypes



Relative risk of relapse within 3 years in breast cancer subtypes, according to RCB index.



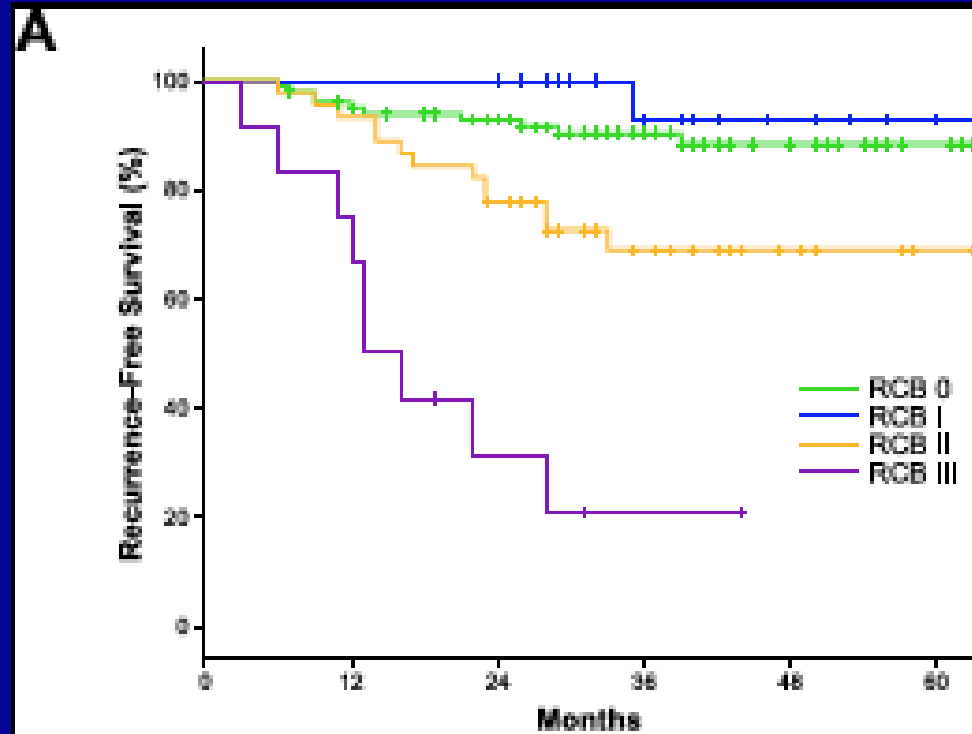
# Prognosis According to RCB Index Within Subtypes



# RCB in TNBC

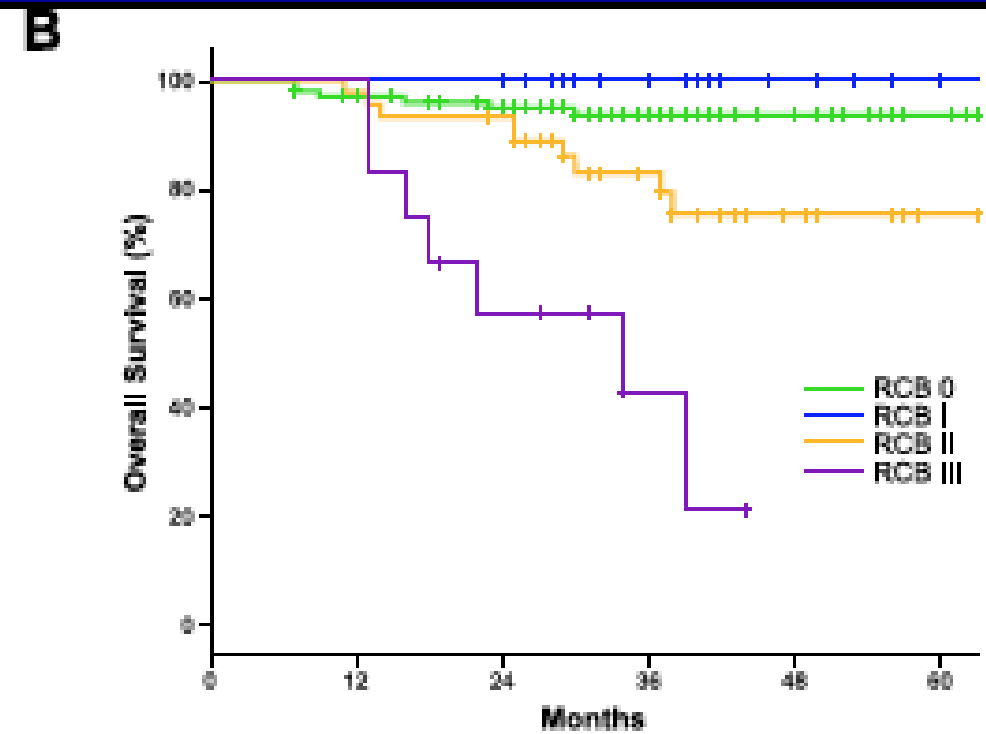
Treated with Carboplatin and Docetaxel (6 cycles)

## Recurrence Free Survival



	No. at risk <sup>a</sup>					
	0	12	24	36	48	60
RCB 0	100	91	80	51	29	19
RCB I	23	23	20	12	6	1
RCB II	45	42	34	18	8	4
RCB III	12	8	3	1	0	0

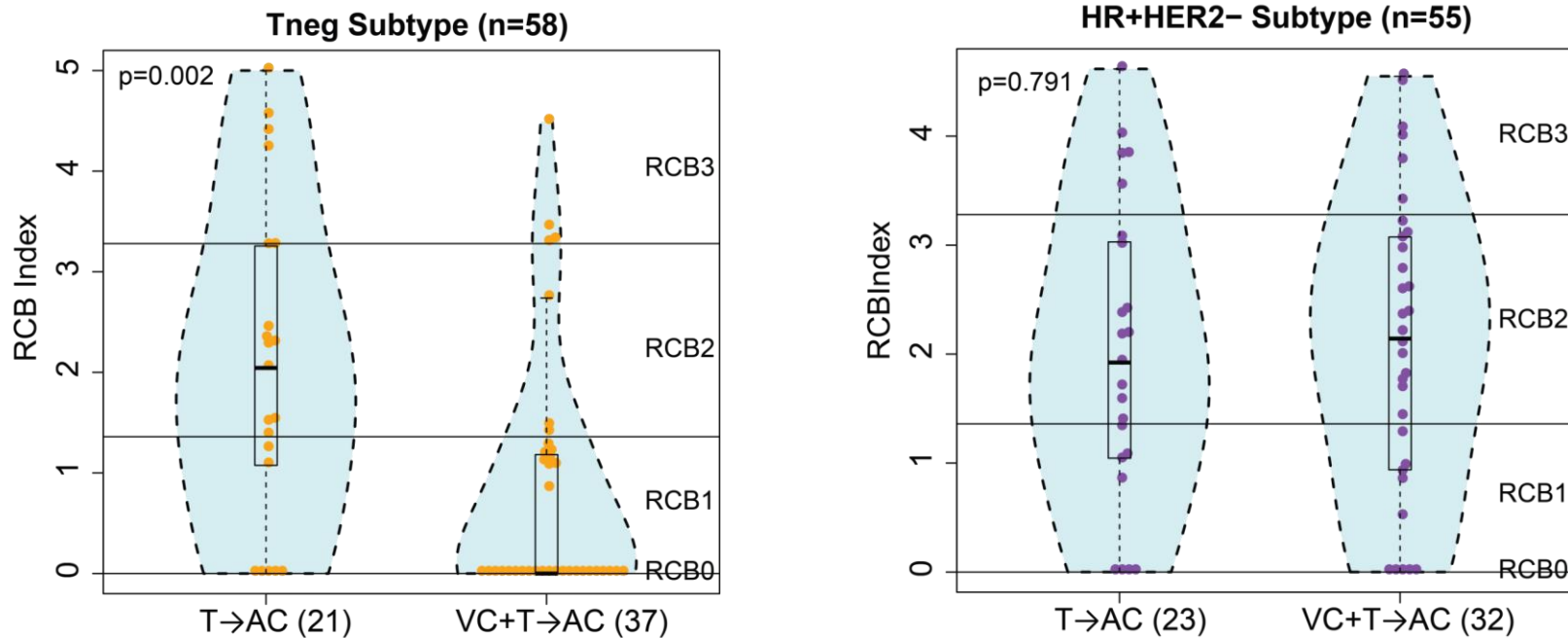
## Overall Survival



	No. at risk <sup>a</sup>					
	0	12	24	36	48	60
RCB 0	100	93	82	53	30	20
RCB I	23	23	20	13	6	1
RCB II	45	44	41	23	10	5
RCB III	12	12	6	2	0	0

# Comparison In A Phase II Randomized Trial

I-SPY2 Trial: Addition of Veliparib and Carboplatin to Weekly Paclitaxel



## Response Endpoint

pCR OR = 4.56, p = 0.013

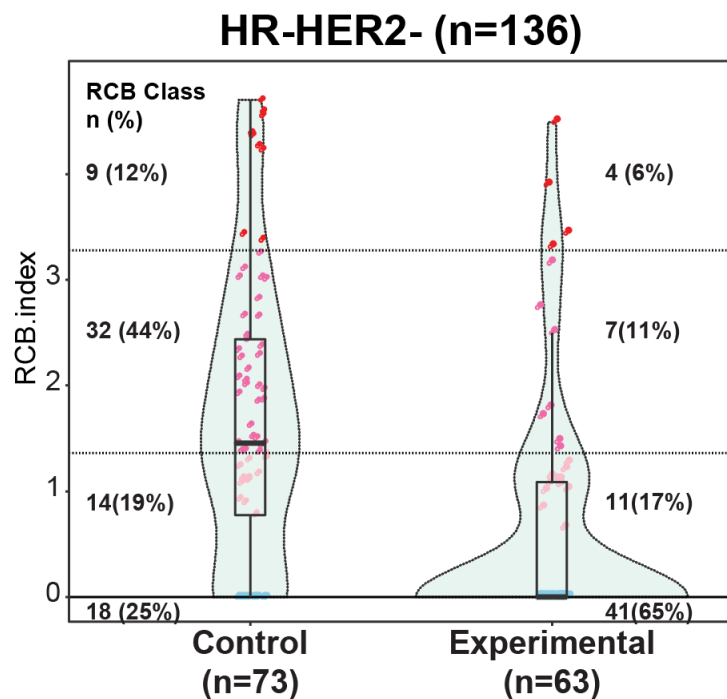
OR = 0.88, p = 1

pCR/RCB-I OR = 8.19, p = 0.0005

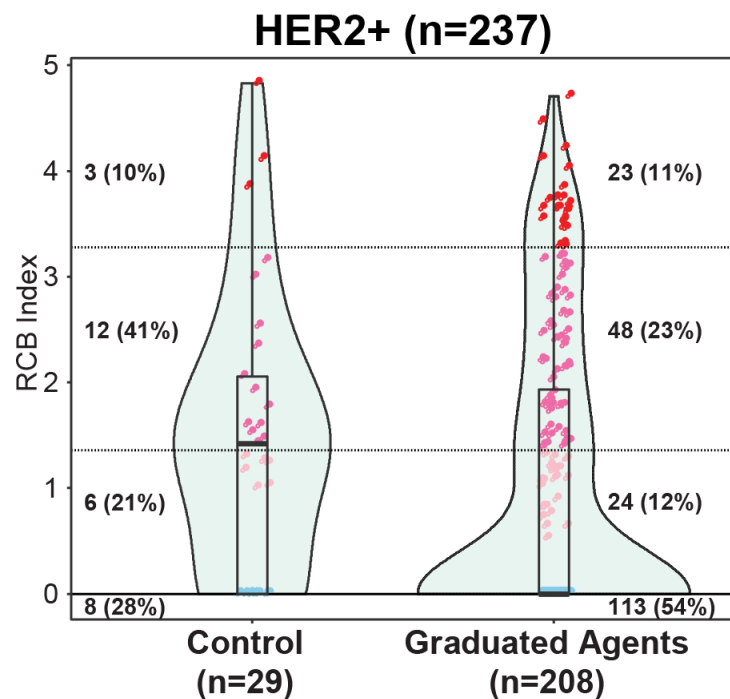
OR = 0.85, p = 1

# At A Glance: Responses In The Whole Trial

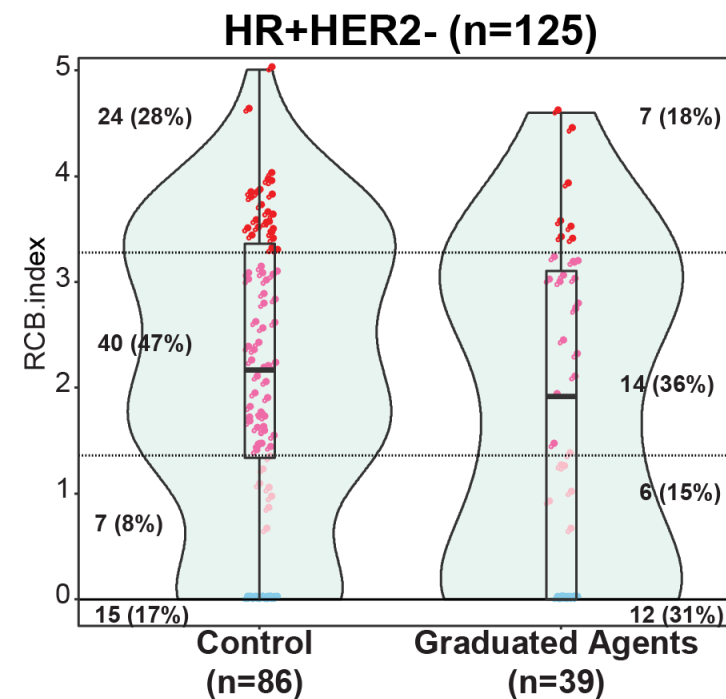
Comparison of RCB index distribution between graduated treatments and control treatment



**Odds Ratio for achieving pCR:**  
5.61 (2.55-12.83)



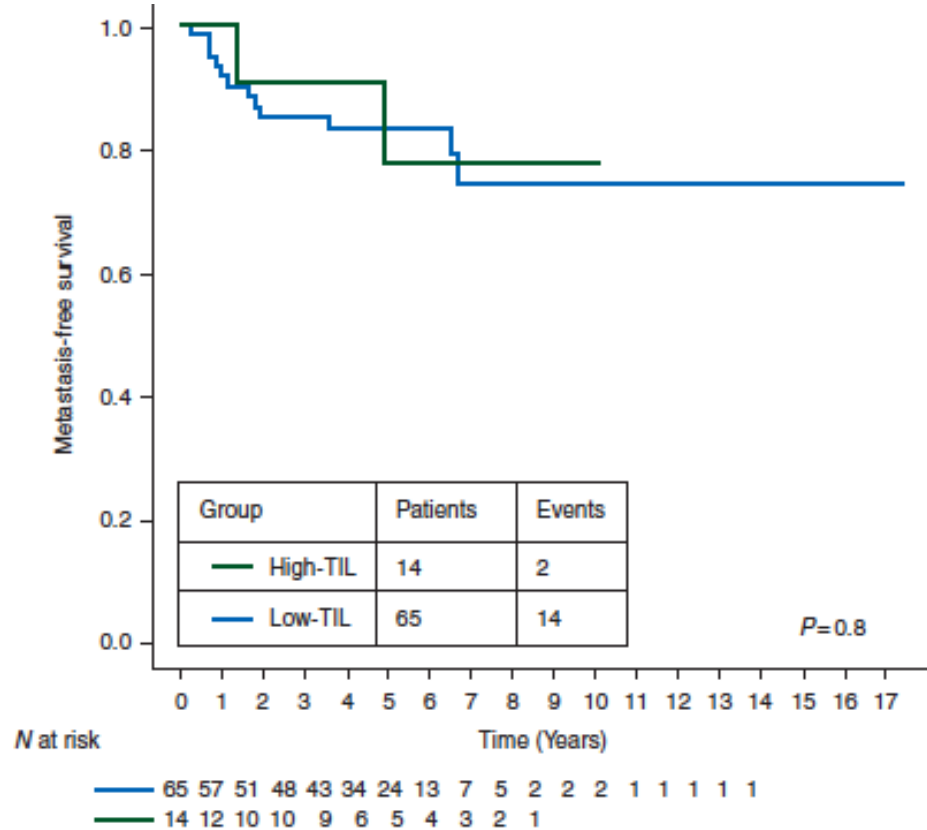
**Odds Ratio for achieving pCR:**  
3.11 (1.25-8.50)



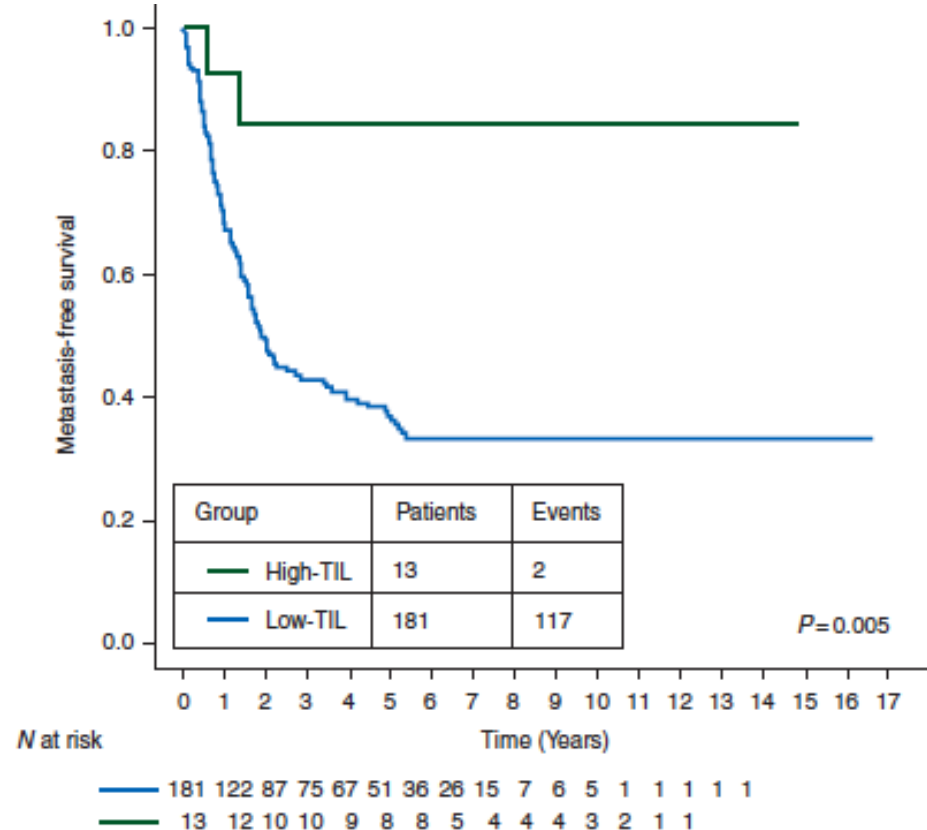
**Odds Ratio for achieving pCR:**  
2.09 (0.78-5.51)

# TILs in the Specimen After Neoadjuvant Chemotherapy

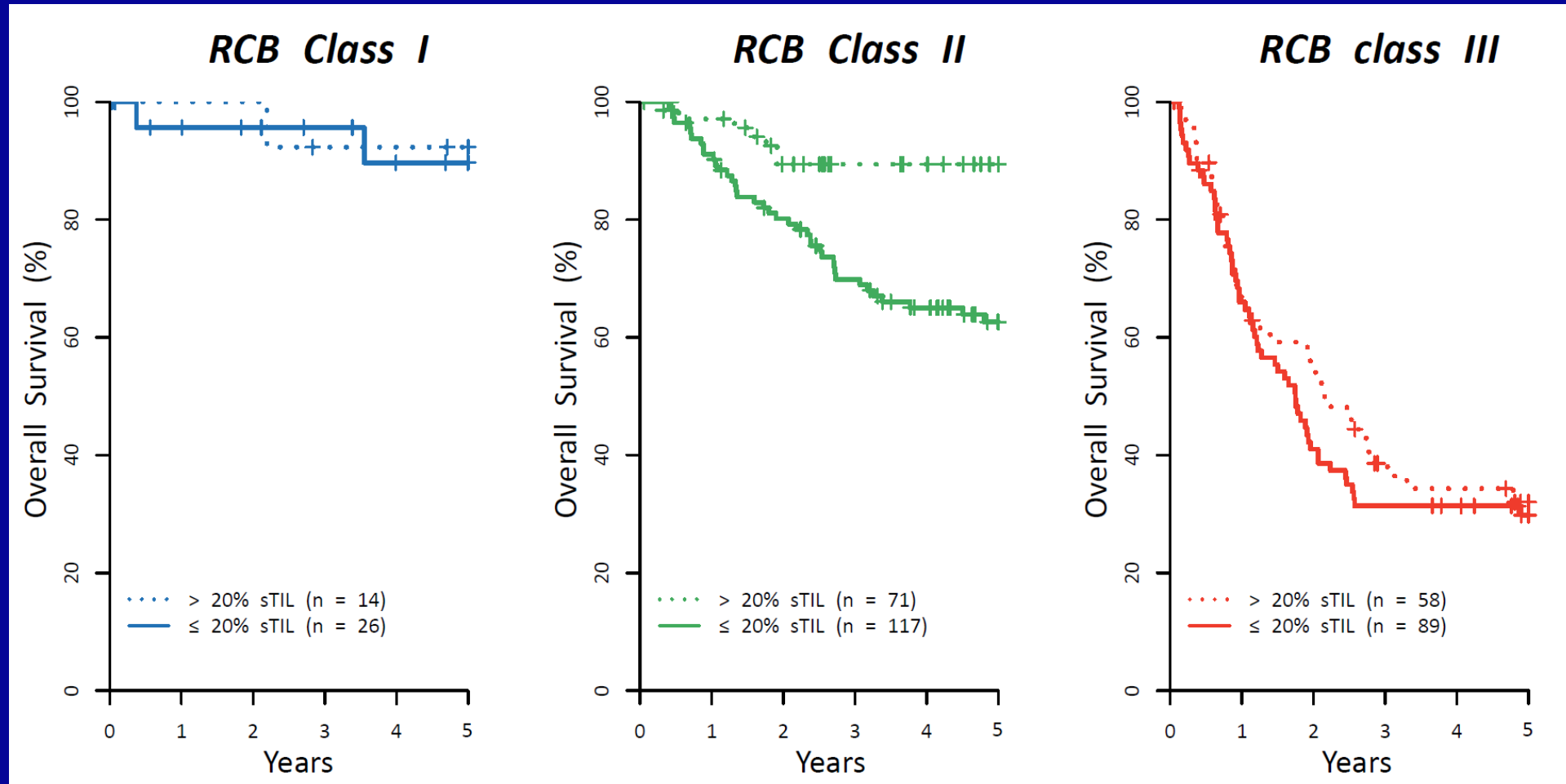
## ypT1 and ypN0



## yp >T1 and/or ypN >0



# TILs in Residual TNBC



# **Recommendations for standardized pathological characterization of residual disease for neoadjuvant clinical trials of breast cancer by the BIG-NABCG collaboration**

V. Bossuyt<sup>1\*</sup>, E. Provenzano<sup>2</sup>, W. F. Symmans<sup>3</sup>, J. C. Boughey<sup>4</sup>, C. Coles<sup>5</sup>, G. Curigliano<sup>6</sup>, J. M. Dixon<sup>7</sup>, L. J. Esserman<sup>8</sup>, G. Fastner<sup>9</sup>, T. Kuehn<sup>10</sup>, F. Peintinger<sup>11,12</sup>, G. von Minckwitz<sup>13</sup>, J. White<sup>14</sup>, W. Yang<sup>15</sup>, S. Badve<sup>16</sup>, C. Denkert<sup>17</sup>, G. MacGrogan<sup>18</sup>, F. Penault-Llorca<sup>19</sup>, G. Viale<sup>20</sup> & D. Cameron<sup>21</sup> of the Breast International Group-North American Breast Cancer Group (BIG-NABCG) collaboration

## **Standardization of pathologic evaluation and reporting of postneoadjuvant specimens in clinical trials of breast cancer: recommendations from an international working group**

Elena Provenzano<sup>1</sup>, Veerle Bossuyt<sup>2</sup>, Giuseppe Viale<sup>3</sup>, David Cameron<sup>4</sup>, Sunil Badve<sup>5</sup>, Carsten Denkert<sup>6</sup>, Gaëtan MacGrogan<sup>7</sup>, Frédérique Penault-Llorca<sup>8</sup>, Judy Boughey<sup>9</sup>, Giuseppe Curigliano<sup>10</sup>, J Michael Dixon<sup>11</sup>, Laura Esserman<sup>12</sup>, Gerd Fastner<sup>13</sup>, Thorsten Kuehn<sup>14</sup>, Florentia Peintinger<sup>15,16</sup>, Gunter von Minckwitz<sup>17</sup>, Julia White<sup>18</sup>, Wei Yang<sup>19</sup> and W Fraser Symmans<sup>20</sup> on behalf of the Residual Disease Characterization Working Group of the Breast International Group-North American Breast Cancer Group (BIG-NABCG) collaboration

# Recommendation

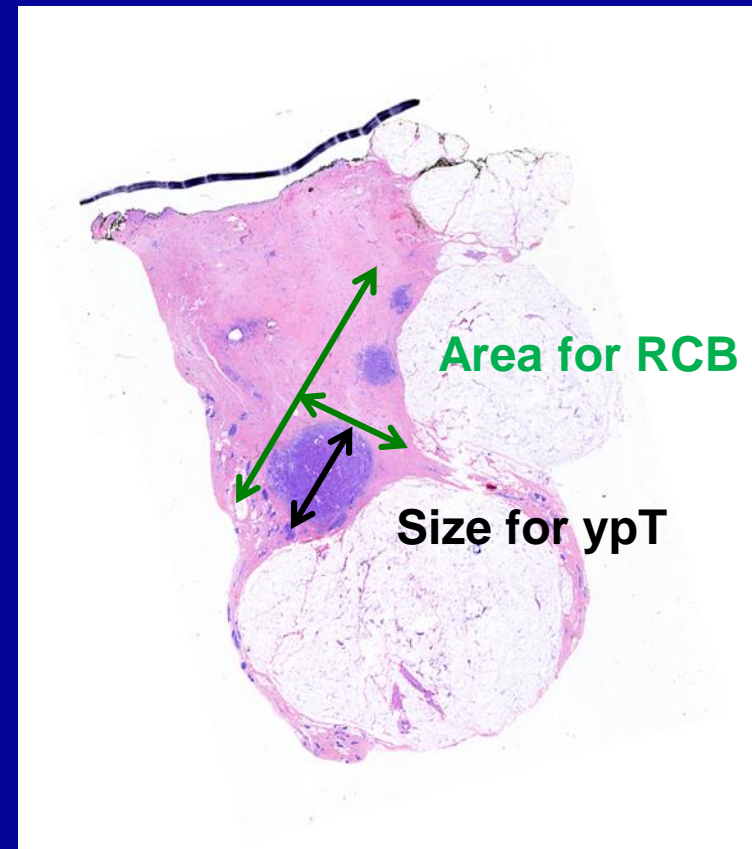
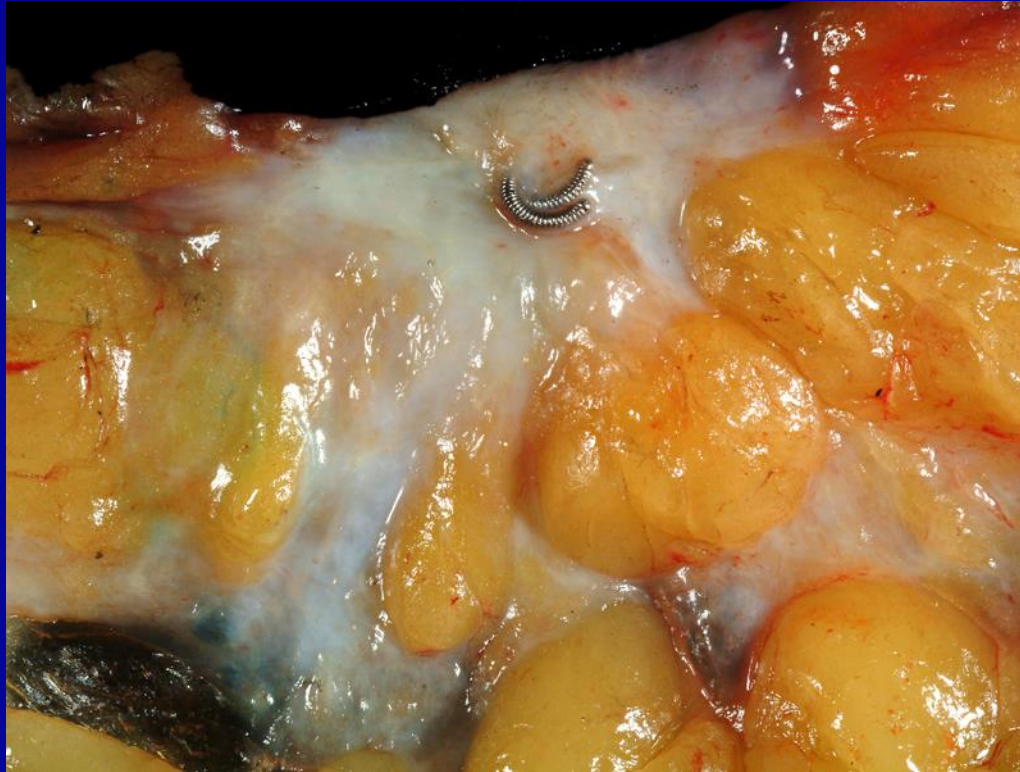
Provide the following information:

1. pCR (ypT0 ypN0 and ypT0/is ypN0) *versus* residual disease,
2. ypT and ypN Stage using the current AJCC/UICC staging system,
3. Residual cancer burden (RCB)

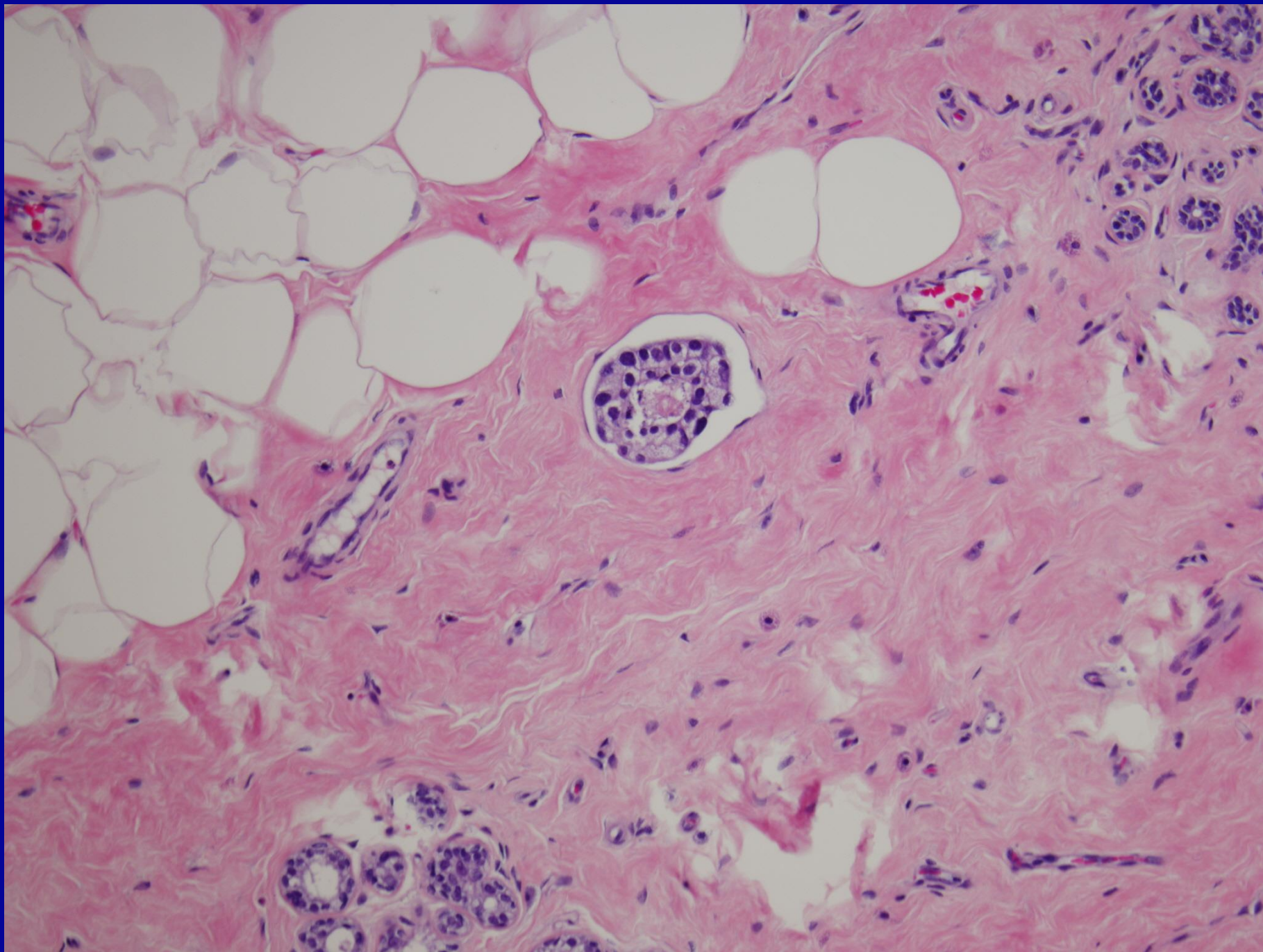
**A single standardized approach to macroscopic and microscopic pathologic examination makes it easy to reliably provide all 3 results!**



# Defining The Size Of The Residual Invasive Cancer



Same principle for measuring metastatic cancer in lymph nodes



# *yp*Stage and RCB Are Different Systems

## Main differences for interpreting residual disease in the breast

1. Primary tumor dimensions:
  - longest continuous invasive cancer is measured to interpret *ypT* for Stage
  - overall extent of invasive cancer (including gaps) is used to measure tumor area for RCB
2. Lymphovascular invasion (LVI):
  - not measured for *ypT* (so can be *ypT0* if only LVI) but cannot be called pCR
  - Interpreted as residual invasive cancer for RCB
3. Cellularity of residual invasive cancer:
  - not considered for Stage
  - **AVERAGE CELLULARITY** across the residual AREA of invasive disease for RCB

# *yp*Stage and RCB Are Different Systems

## Main differences for interpreting residual disease in the regional lymph nodes

### 1. Dimension of nodal metastases:

- longest continuous metastatic deposit is interpreted for *yp*N
- overall length of largest metastasis (inclusive of gaps) used for RCB and this would include extranodal extension

### 2. Isolated tumor cells < 0.2 mm (ITC):

- *yp*N0(i+), but cannot be called pCR
- interpreted as metastatic cancer for RCB, and measurement used if largest metastasis

# Excerpt From Our CAP Synoptic Report ...

## **Lymph Nodes**

**Total number of lymph nodes examined (sentinel and non-sentinel): 28**

**Number of sentinel lymph nodes examined: 0**

### **Lymph node involvement**

**Number of lymph nodes with macrometastases: 1**

**Number of lymph nodes with micrometastases: 0**

**Number of lymph nodes with isolated tumor cells: 0**

**Extranodal extension: 0**

## **RCB Index Input Variables**

**Dimensions of residual primary tumor area: 100 mm X 80 mm**

**Percent cancer cellularity of residual tumor area: 0.01%**

**Percent of residual cancer that is in situ: 100%**

**Total number of lymph nodes with residual tumor: 1**

**Size of largest metastasis: 4.5 mm**

**RCB index computed value (optional): 1.29**

**RCB class (optional): RCB-I**

## **Pathologic staging (pTNM)**

**Primary tumor (invasive carcinoma) (ypT): ypT0is**

**Regional lymph nodes (ypN): ypN1**

**Additional modifiers:**

**Distant metastasis (ypM): ypMX**

# Online Educational Resources

**Google Search Term: “RCB Breast” or “Residual Cancer Burden Breast”**

Educational Videos – Macroscopic and Microscopic Evaluation

<https://www.mdanderson.org/for-physicians/clinical-tools-resources/clinical-calculators/residual-cancer-burden.html>

Protocol for Pathologists – Detailed SOP Document

<https://www.mdanderson.org/education-and-research/resources-for-professionals/clinical-tools-and-resources/clinical-calculators/calculators-rcb-pathology-protocol2.pdf>

Calibration of Percent Cellularity by Area – Computer Generated Examples

<https://www.mdanderson.org/education-and-research/resources-for-professionals/clinical-tools-and-resources/clinical-calculators/calculators-cellularity-guide.pdf>

# Multigene Assays Foe Breast Cancer

# Survival Risk of Breast Cancer in the Adjuvant Setting

The probability that micrometastatic disease ever existed, survived the entirety of treatments, and was able to reawaken and flourish

Original burden of cancer prior to treatment

Biology of the cancer – natural history prognosis

Sensitivity to chemotherapy

- e.g. extent of residual cancer after neoadjuvant treatment

Sensitivity to radiation therapy

Sensitivity to adjuvant endocrine therapy

- Duration, type, bisphosphonates, adherence

Constitutional health that promotes dormancy

- Immunity, metabolism, prevention agents, other



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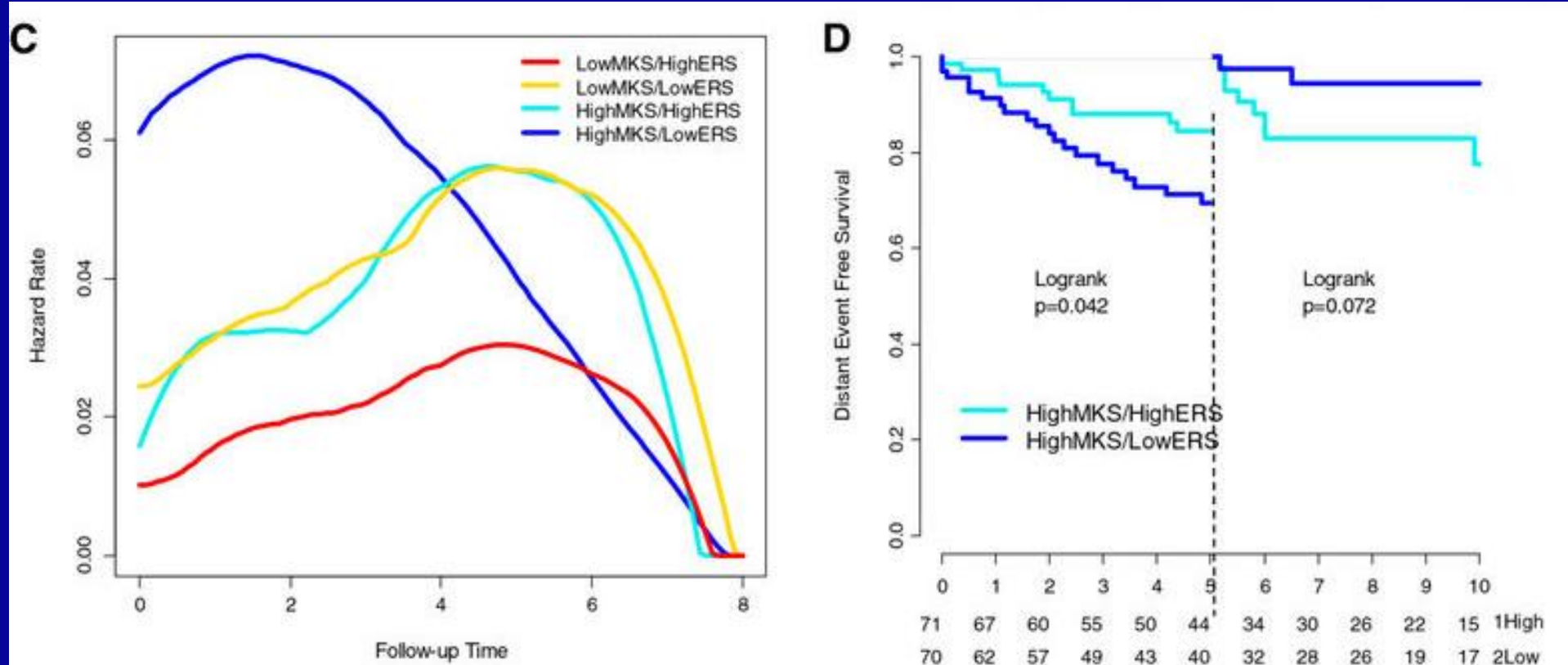
Sensitivity to adjuvant endocrine therapy

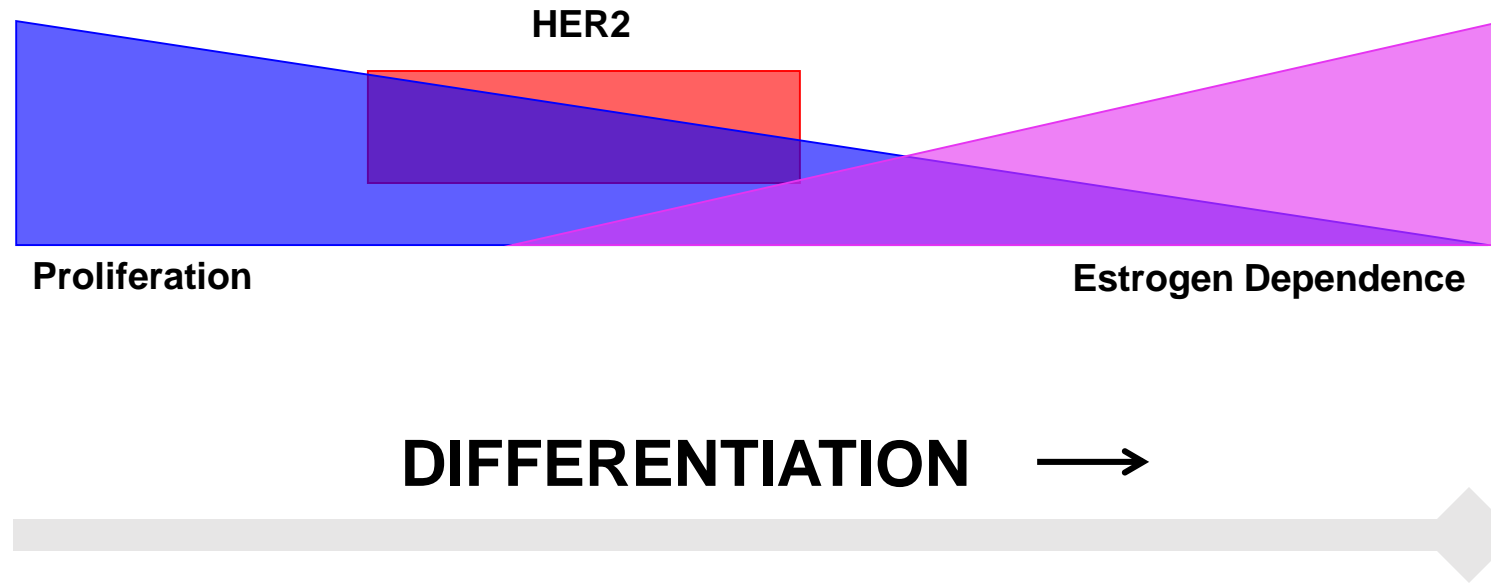
- Duration, type, bisphosphonates, adherence

Constitutional health that promotes dormancy

- Immunity, metabolism, prevention agents, other

# Time-Dependent Risk Related to Proliferation and to Endocrine Activity in HR+/HER2- Breast Cancers





Standard markers are generally sufficient for classification.

Multi-gene expression panels have favorable analytical characteristics that can:

- improve prognostic / theranostic performance.
- be useful when standard markers are indeterminate.

Different multi-gene expression panels have similar prognostic performance.

# Have New Tests Met Clinical Need?

Therapeutic Decision (implied action)	HR+ / HER2-				HER2+				TNBC			
	I	II	III	IV	I	II	III	IV	I	II	III	IV
<b>Natural History</b> Ultra-low Risk (no Rx)	✓ 1	Expression - - -			-	-	-	-	?	-	-	-
<b>Targeted Rx Alone</b> Ultra-low Residual Risk (no CT)	✓ 5	?	-	-	-	-	-	-	-	-	-	-
<b>CT + Targeted Rx</b> Lower Residual Risk (standard of care)	✓ 5	-	-	-	✓ 1	✓ 1	✓ 1	✓ 1	HER2 IHC/FISH			
<b>CT + Targeted Rx</b> High residual risk (clinical trial)	-	-	-	-	-	-	-	-	-	-	-	-
<b>Susceptibility</b> (prophylactic)	✓ 1				Germ line genetics ✓ 1				✓ 1			

Modified from ASCO 2013 Annual Meeting: ASCO/ESMO Presidents' Symposium

# PHENOTYPE

**Classification**  
PAM50 subtypes (& others)  
HR+/HER2-

**Prognosis: natural history**

70-gene Mammaprint

**Prognosis: endocrine therapy**

21-gene Recurrence Score

Breast Cancer Index

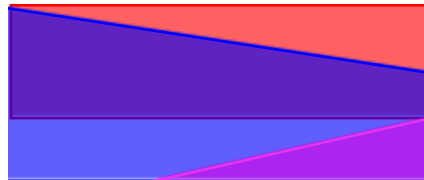
EndoPredict

Prosigna ROR score

**Receptor Activity**

SET index

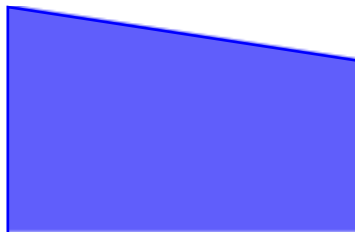
HER2+



**Classification**

*HER2-like* subtype of PAM50

TNBC



**Classification**

*Basal-like* subtype of PAM50

Vanderbilt subtypes

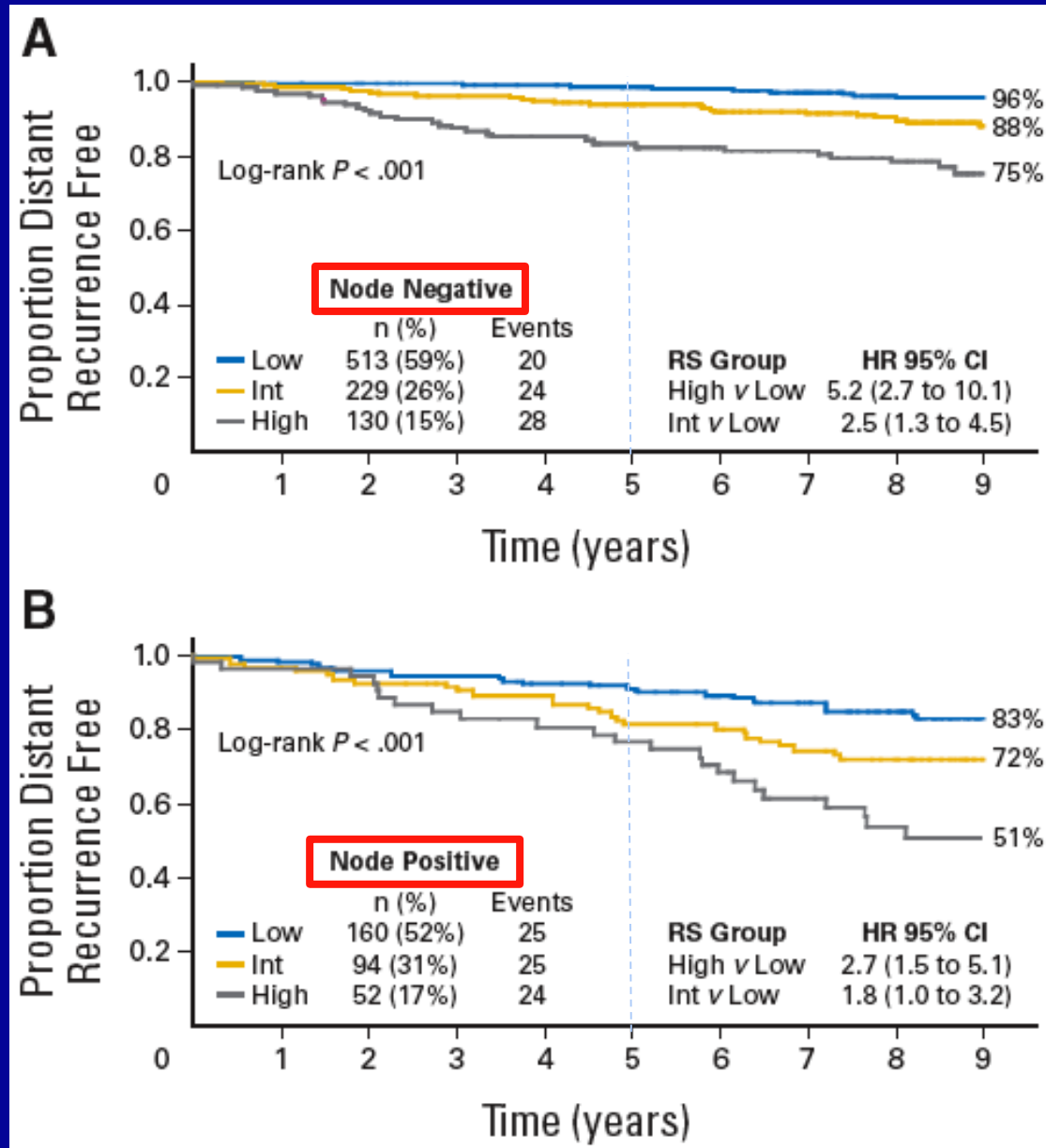
**Prognosis: chemotherapy**

Prognostic signature (immune)

**Response & Prognosis**

MDACC algorithm, Myriad HRD assay

# 21-Gene Recurrence Score: Tamoxifen or Anastrozole



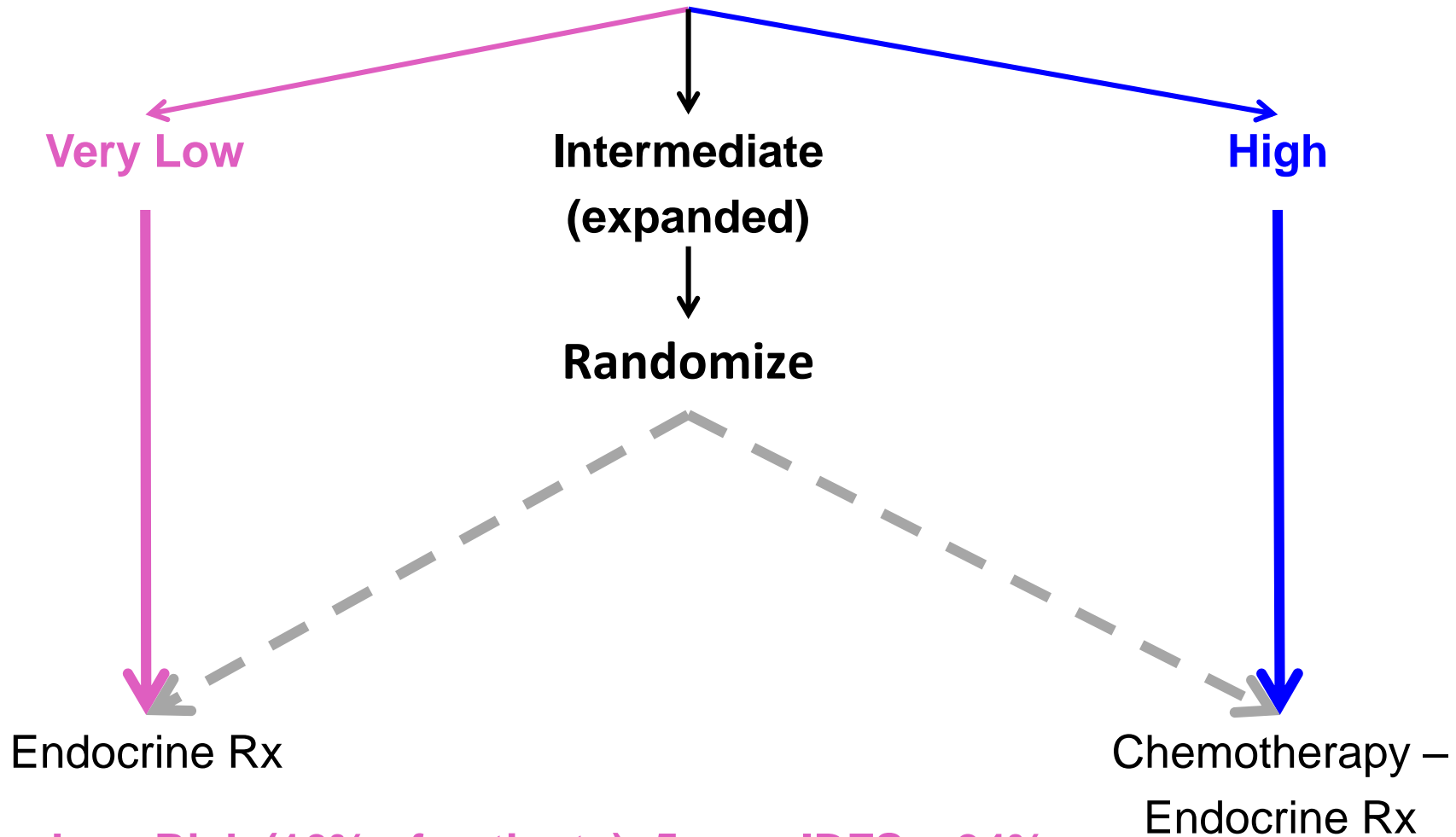
Postmenopausal women

Nodal Status Matters

Hazard ratio for RS group was adjusted for tumor size, grade, age, and treatment

# TAILORx Trial

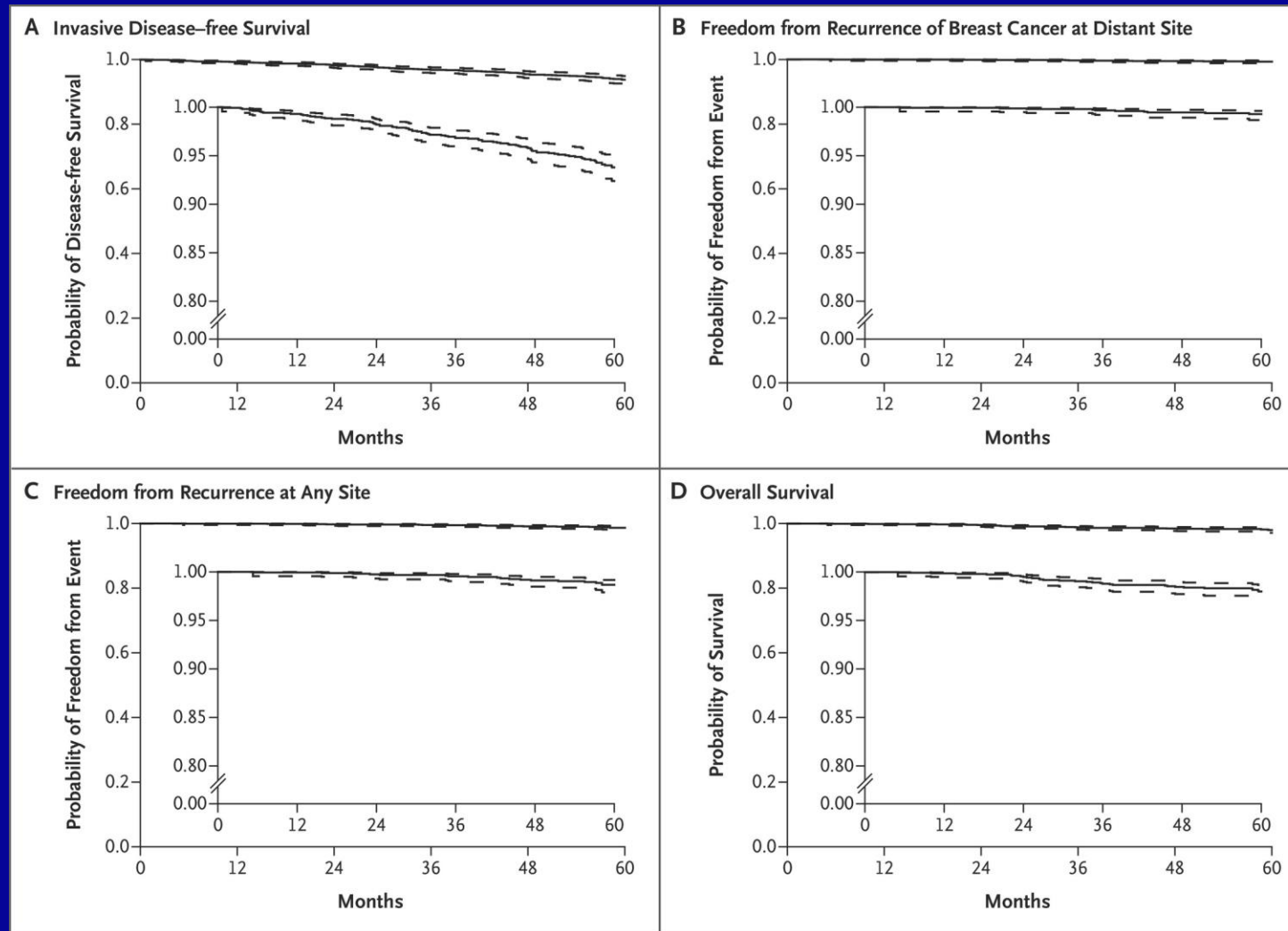
Recurrence Score in ER+, LN-



**Low Risk (16% of patients): 5-year IDFS = 94%**

Sparano et al NEJM 2015;373:2005-14.

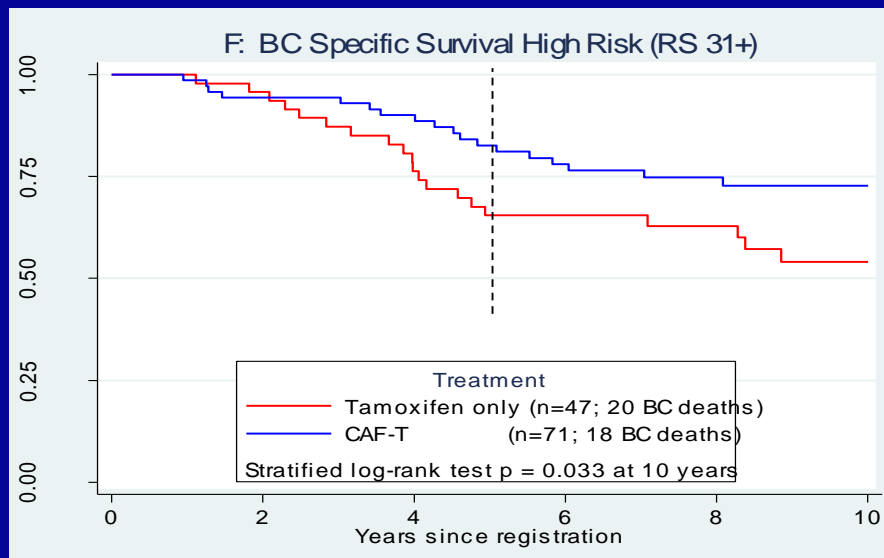
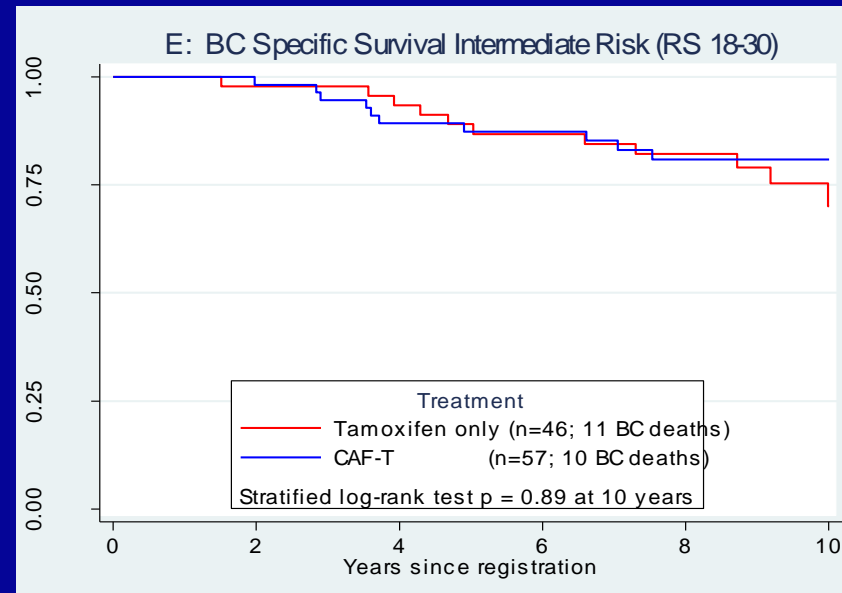
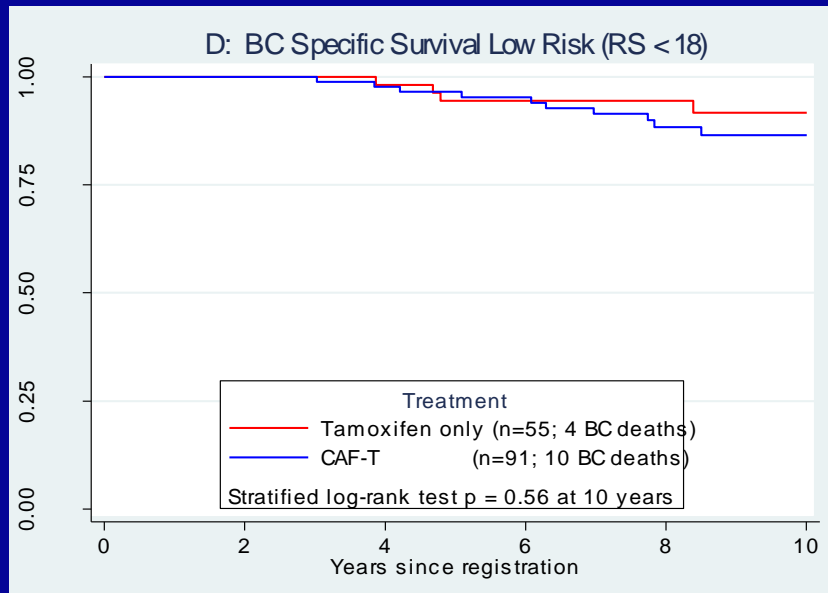
# Kaplan–Meier Estimates in the Analyses of Invasive Disease–free Survival, Freedom from Recurrence of Breast Cancer at a Distant Site, Freedom from Recurrence at Any Site, and Overall Survival.



Sparano JA et al. N Engl J Med 2015;373:2005-2014



# Sequential Chemo-endocrine Therapy, Node-positive: SWOG-8814 Results for Recurrence Score



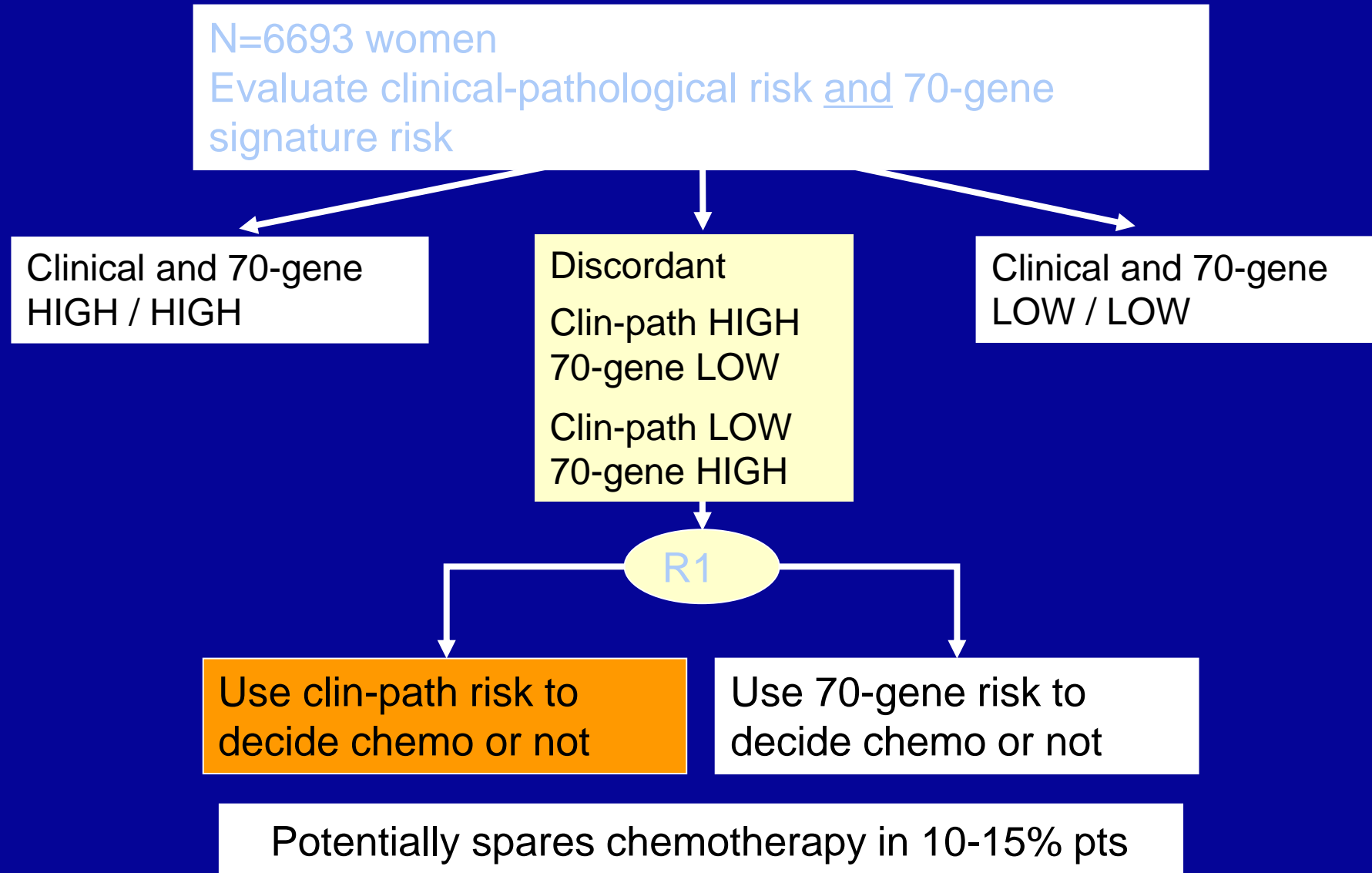
**Interaction p = 0.021**

**GHI unable to identify reliable multi-gene panel for chemoprediction using RNAseq from FFPE samples**

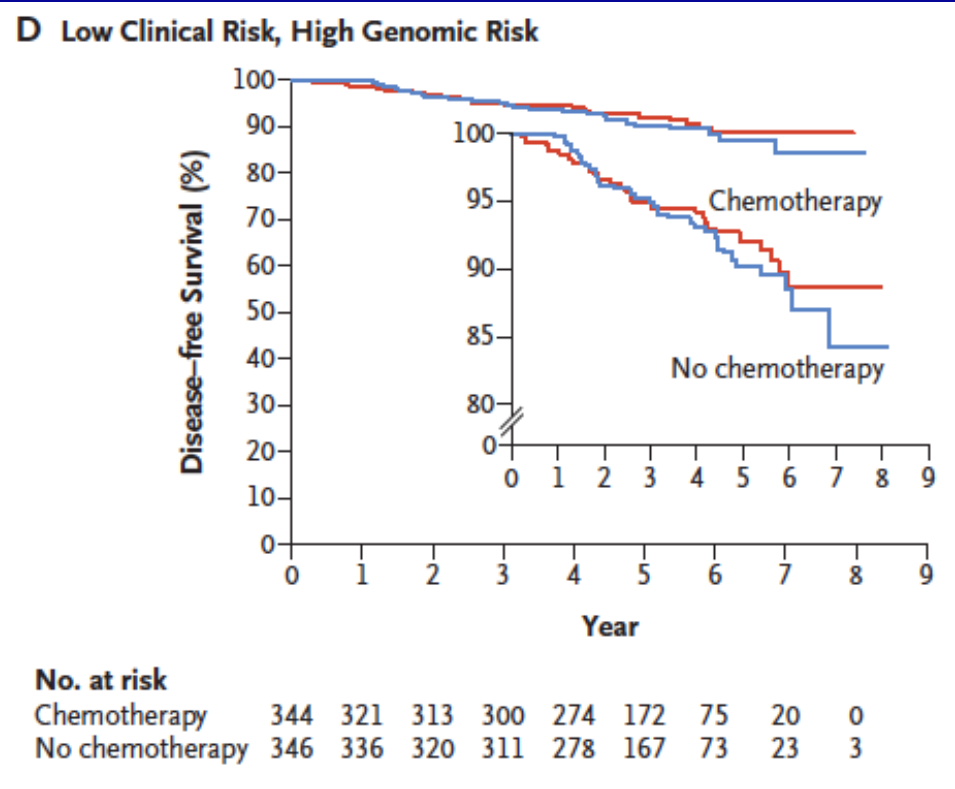
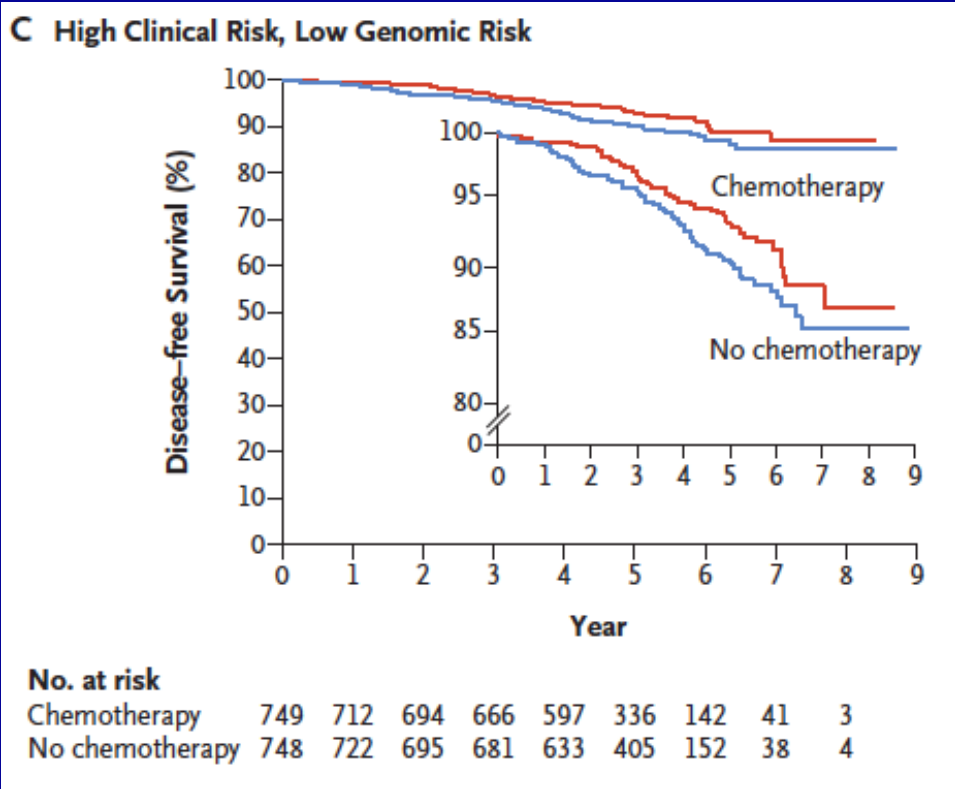
**Albain et al SABCS December, 2015**

**Albain et al. Lancet Oncol 2010, 11:55-65**

# EORTC-BIG MINDACT TRIAL

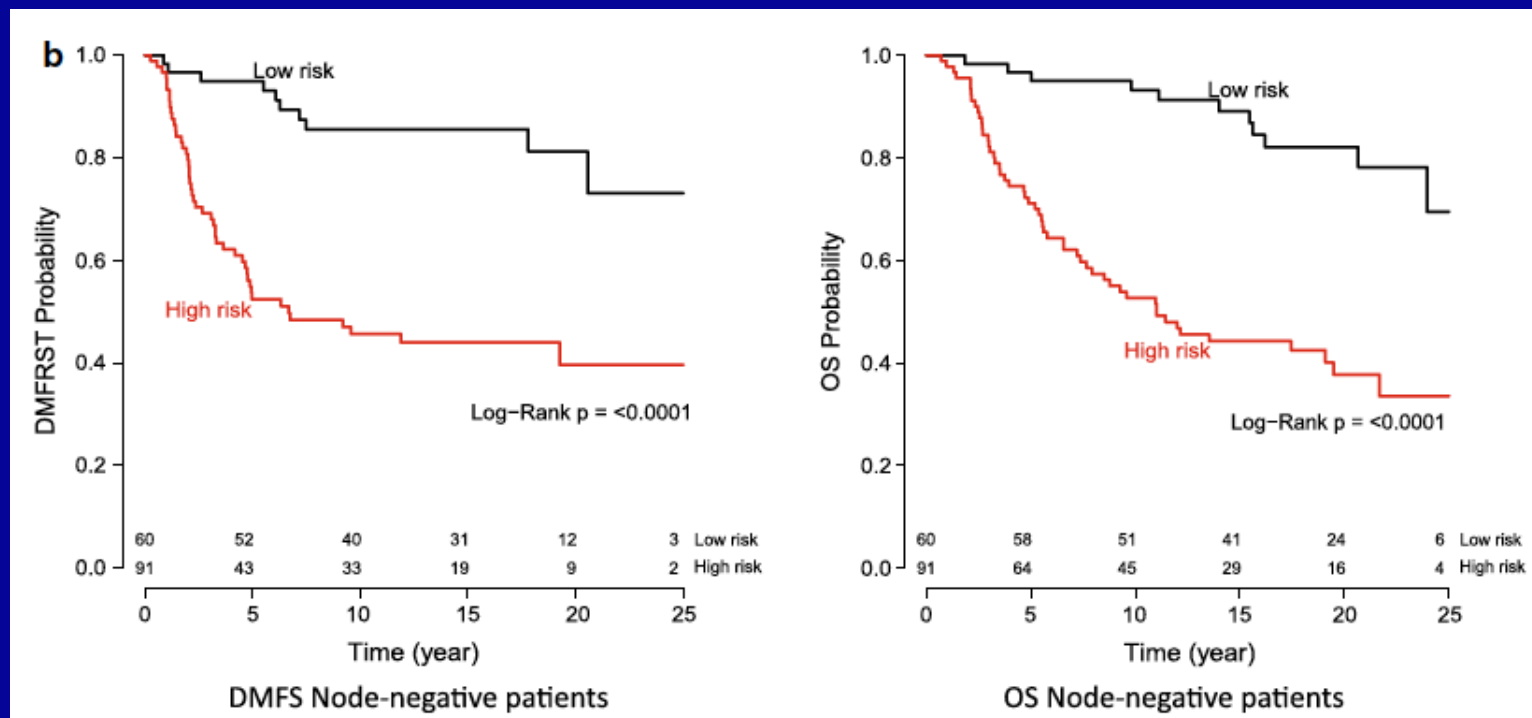


# MindACT Trial: 70-gene Mammaprint Assay



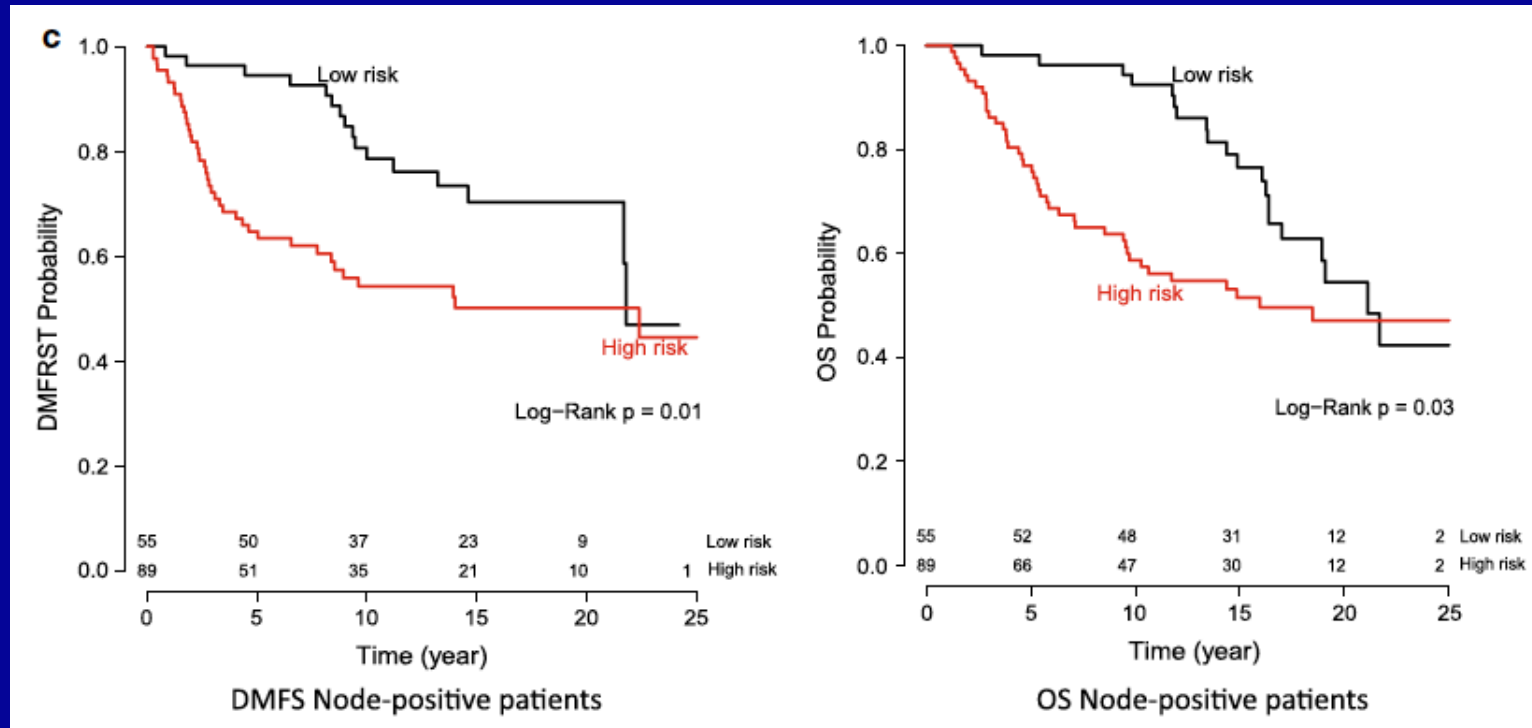
# 70-Gene Mammaprint Assay Very Long Term

Node-Negative



# 70-Gene Mammamprint Assay Very Long-Term

## Node-Positive



# Summary

## **Prognostic biology can be time-dependent**

- **Initial 5 years = Proliferation**
- **Years 5 to 10 = Endocrine**

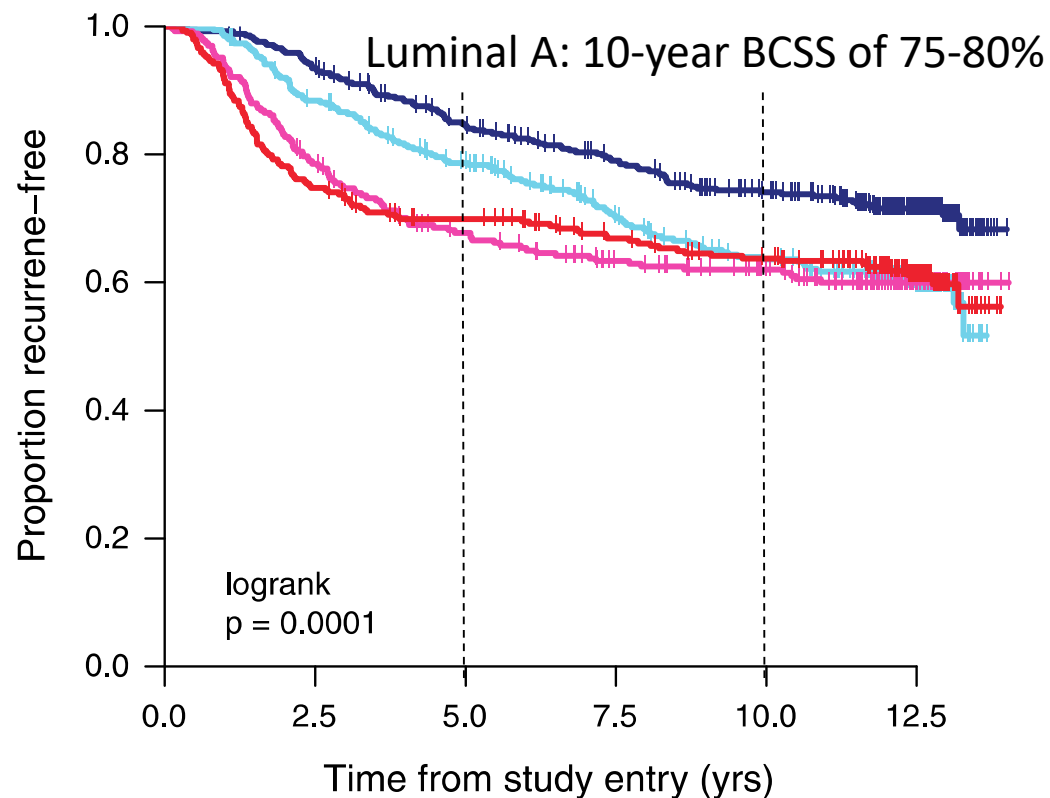
## **Performance of prognostic tests is context-dependent**

- **All the boats (large & small) rise on the incoming tide**
  - **Burden of disease retains importance**

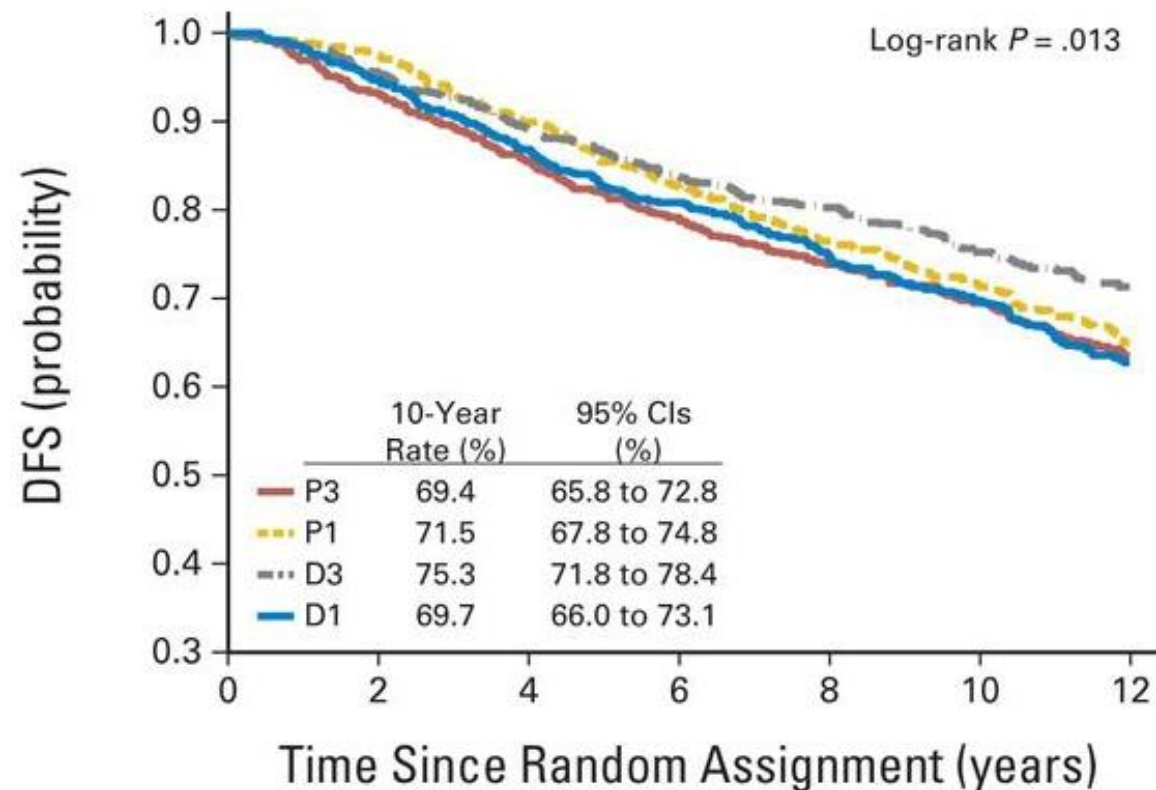
Residual Risk and Prediction: Chemotherapy +/- Endocrine Therapy

# Stage II-III Breast Cancer: Long-term Residual Risk

**CALGB-9741, AC/T chemo, all subtypes**



**ECOG E-1199, AC/T chemo, HR+/HER2-**

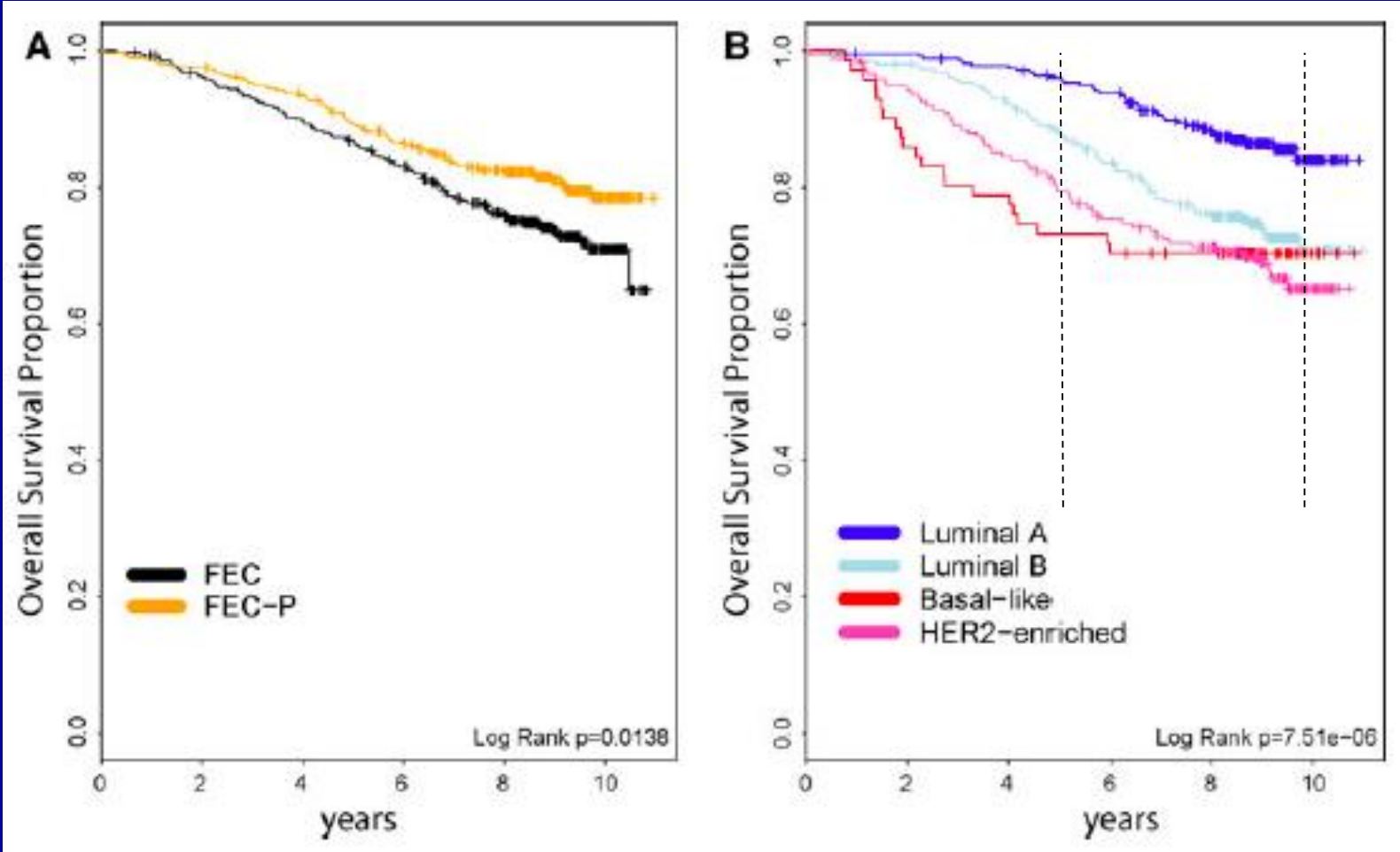


	Number at risk					
	0	2.5	5.0	7.5	10.0	12.5
LuminalA	414	385	335	293	256	125
LuminalB	338	295	253	203	172	81
Her2Enriched	266	207	171	146	126	57
BasalLike	293	219	195	173	159	69

	No. at risk						
	0	2	4	6	8	10	12
P3	715	653	572	505	452	377	197
P1	706	673	600	525	458	385	189
D3	731	683	621	562	513	436	223
D1	735	683	599	523	458	381	187

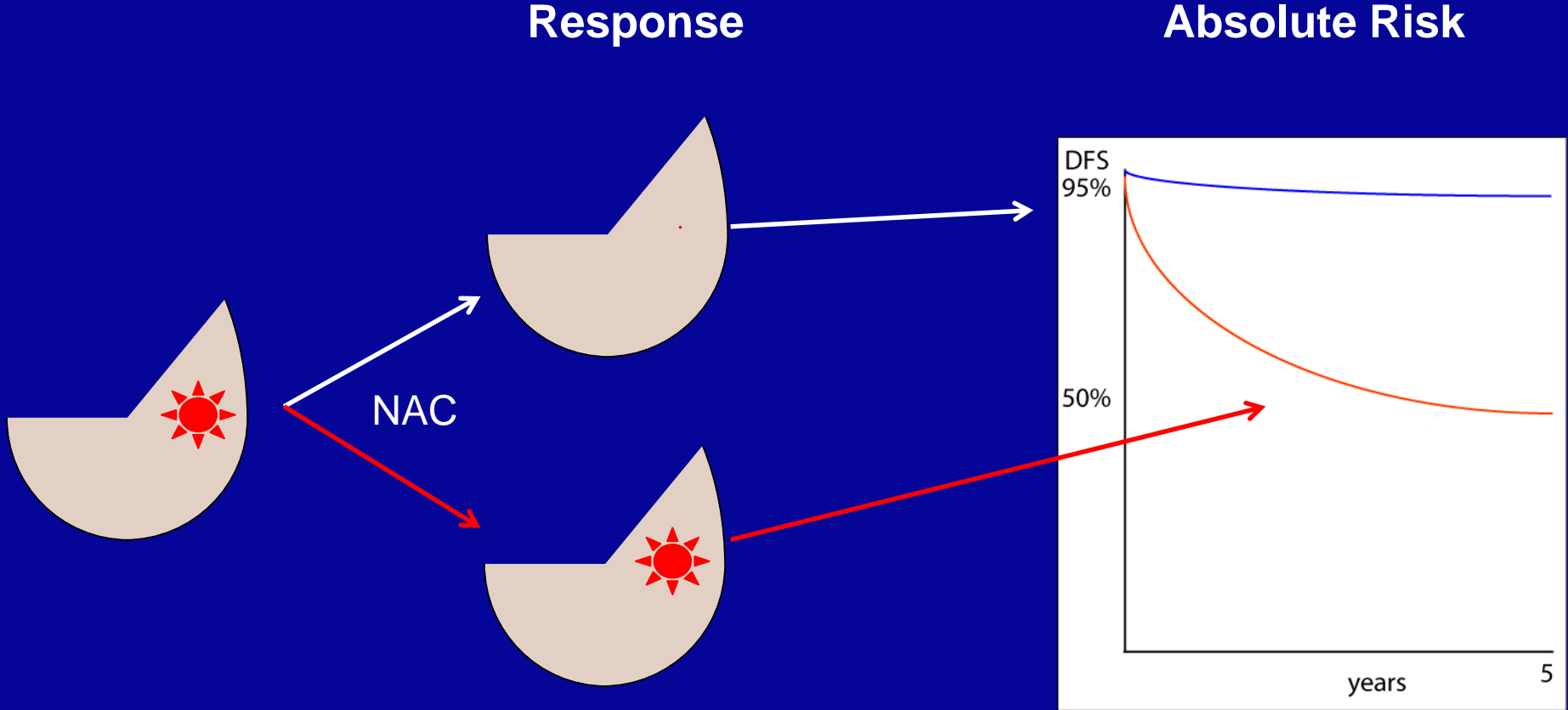


# GEICAM 9906: FEC +/- weekly Paclitaxel, Node-Positive Breast Cancer



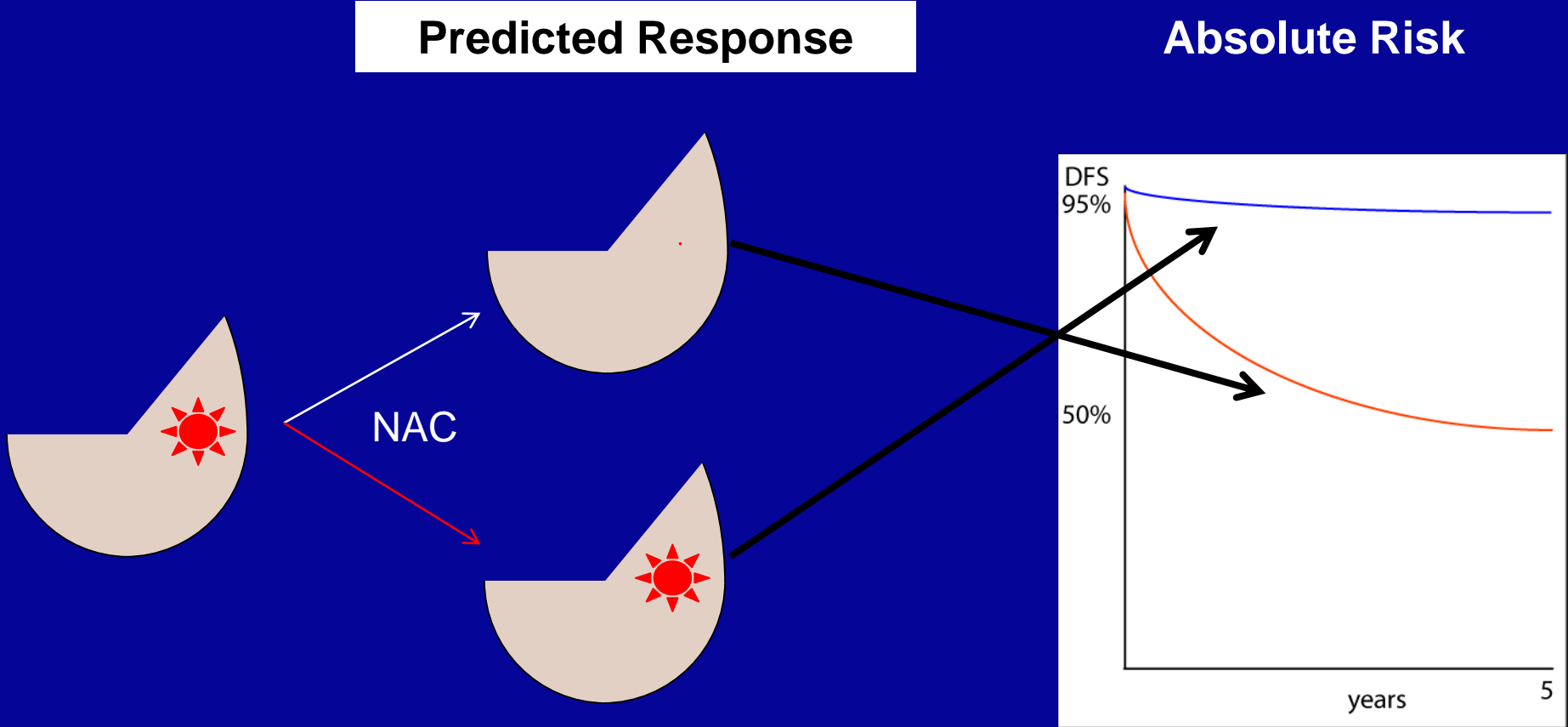
**PAM50 Intrinsic Subtypes did not predict benefit from paclitaxel**

# Prediction of Chemotherapy Response and Survival



Model depends on alignment of **BOTH** the predicted response outcome and the absolute survival probability

# Prognostic Biomarkers: Paradox

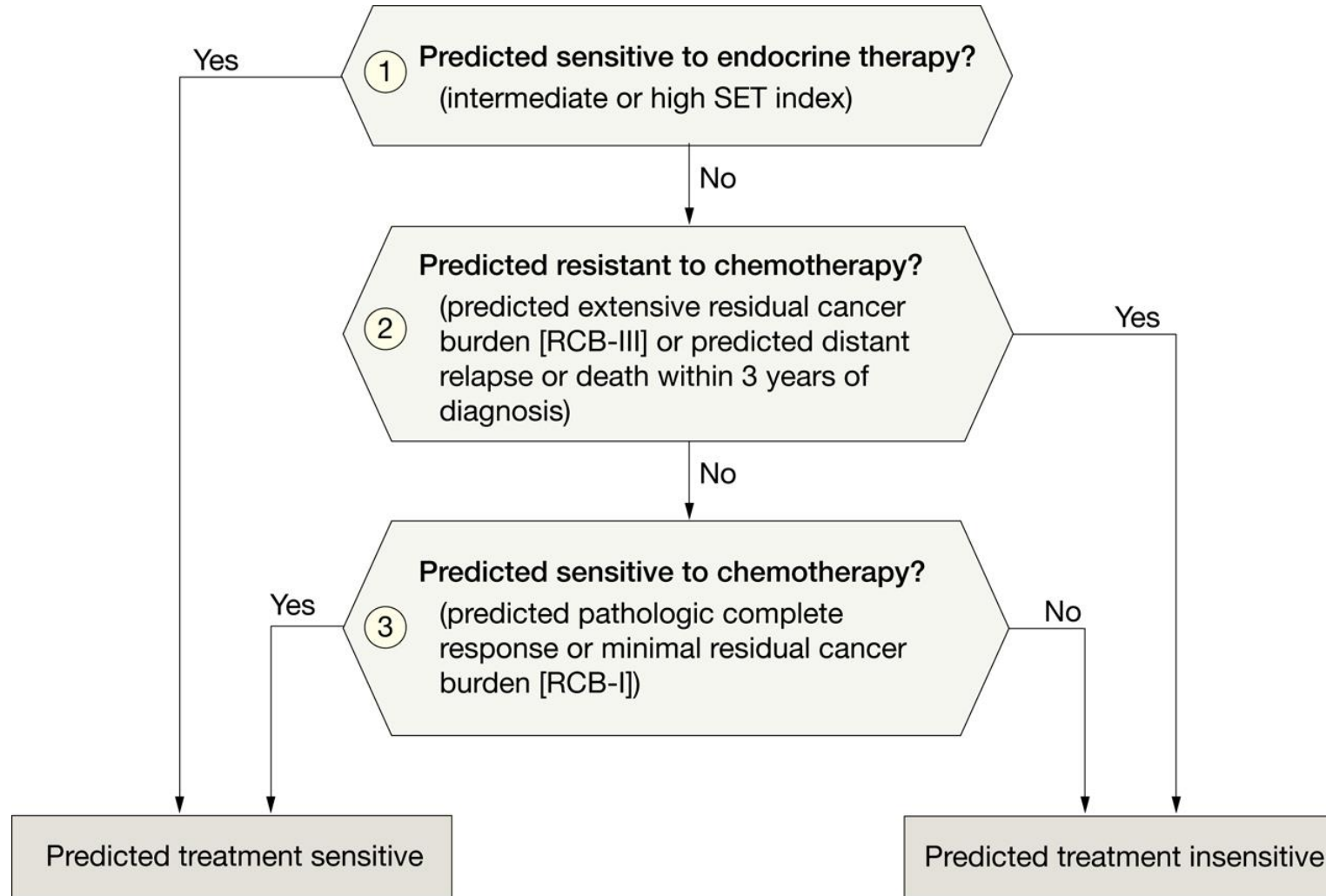


**Indolent tumor biology associated with poor response but good prognosis.**

# Summary of the Chemopredictive Properties of Molecular Classification/Prognostic Tests

Test	Subset	Chemosensitive Group	Response: PPV for pCR	Survival: relative
Grade	all	High grade	≈ 30%	worse
Ki67	all	High proliferation	≈ 30%	worse
Recurrence Score	HR+/HER2-	High	≈ 20-30%	similar
Mammaprint	HR+/HER2-	High Risk	≈ 20%	worse
PAM50	HR+/HER2-	Lum B	≈ 50%	worse
PAM50	TNBC	Basal-like	≈ 50%	similar
GGI	all	High grade	≈ 50%	worse
DLDA-30	all	Predicted pCR	≈ 50%	worse

**Proof of Concept Strategy For T → A Chemotherapy +/- Endocrine Therapy**  
**Developed Separately For ER+/HER2- and ER-/HER2- Cancers (N = 310)**



Hatzis, C. et al. JAMA 2011;305:1873-1881

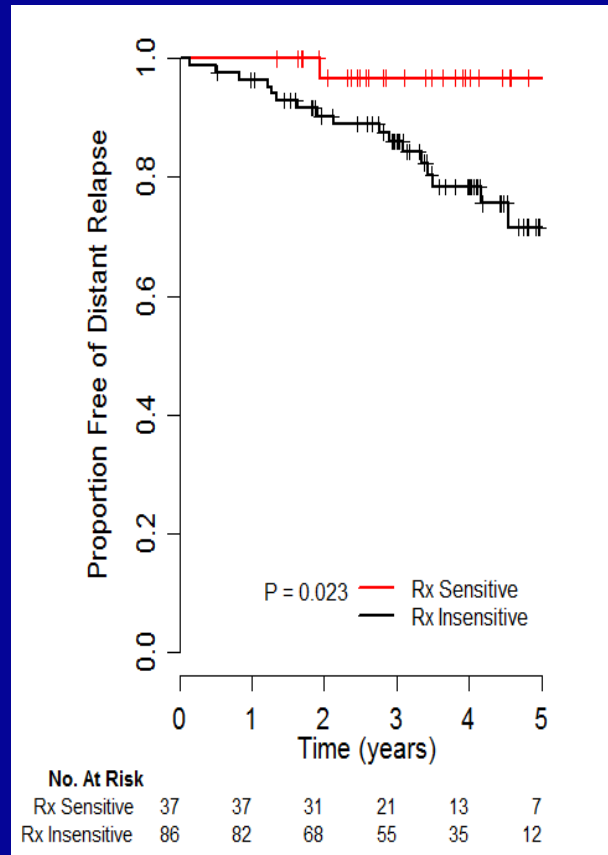
**JAMA**

# Response & Survival Outcomes in Validation Cohort

N = 198 Chemotherapy +/- Endocrine Therapy (99% Stage II-III)

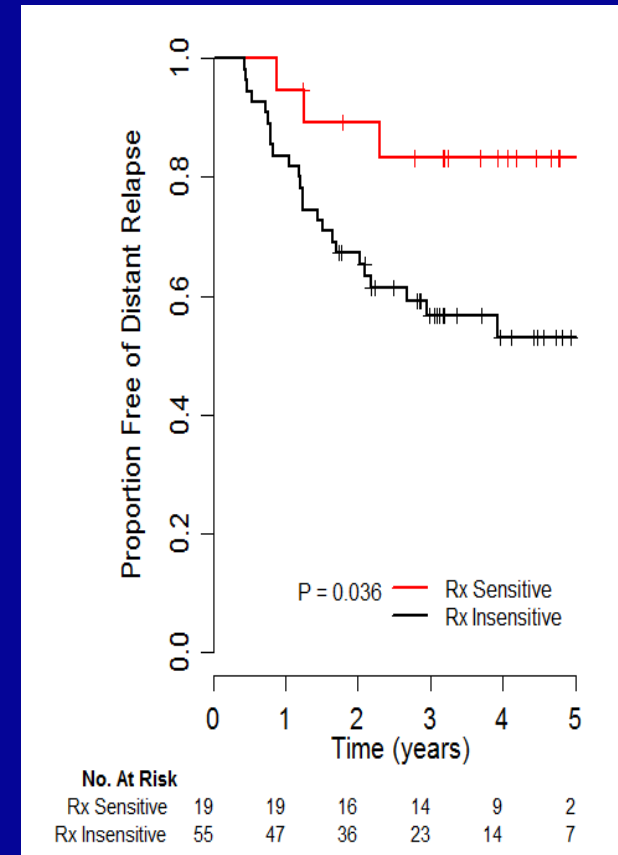
**HR+ / HER2-**

PPV for response 42% (95% CI 15-72)

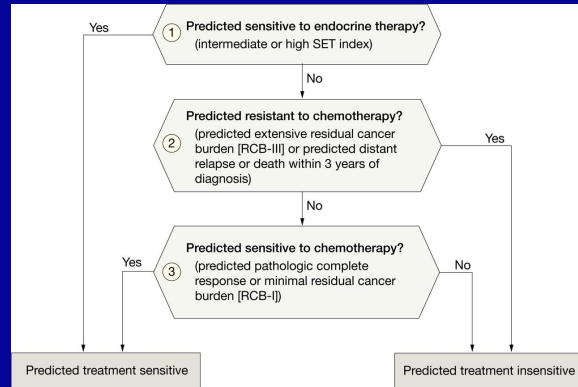


**HR- / HER2- (TNBC)**

PPV for response 83% (95% CI 36-100)



# TNBC: Logistic Algorithm vs. Random Forest Algorithm to Combine Gene Expression Signatures (GES)



cN-

GES: TILs

GES: RCB-III  
or  
RCB-II & relapse

GES: pCR/RCB-I  
&  
no relapse

Kappa values: 0.24 - 0.34

Triplicate Runs  
23 TNBCs

Kappa values: 0.63 - 0.91

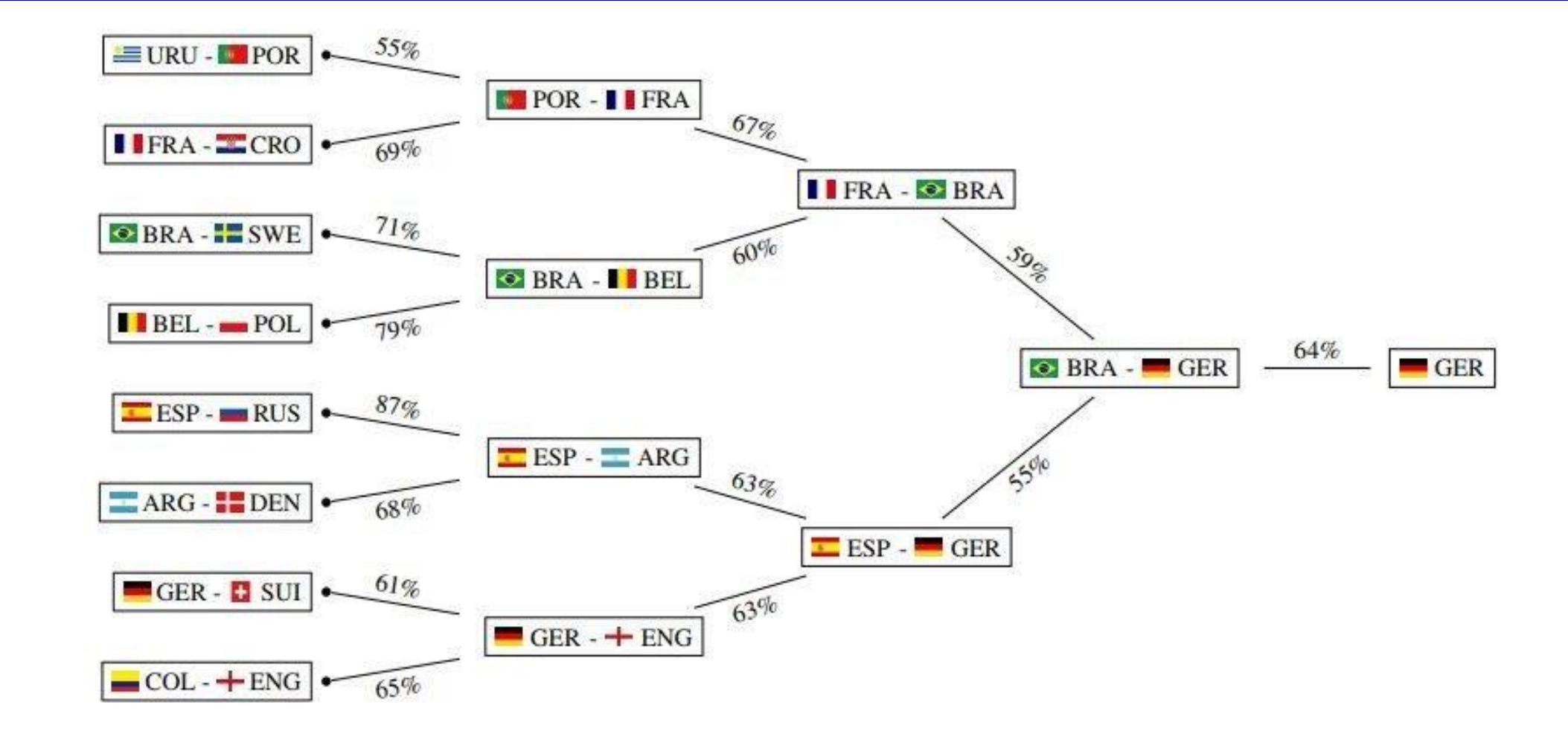
PPV: 0.60 - 0.65

Validation Cohort in TNBC  
Predicting pCR/RCB-I

PPV: 0.60 - 0.65

# Machine Learning Algorithms To Combine Predictions

Example: Random Forest Approach

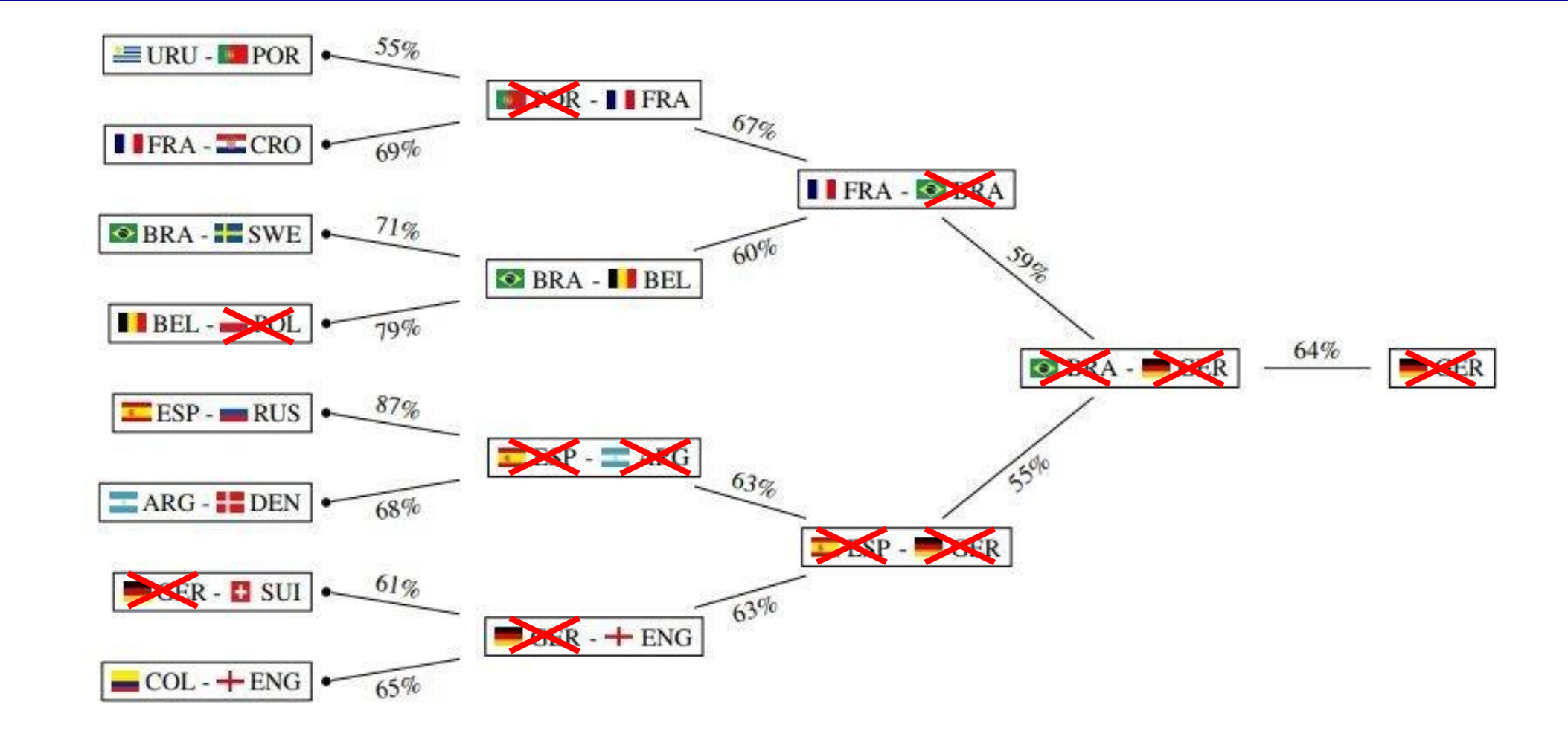


Groll et al. Technical University of Dortmund, MIT Technology Review, June 12, 2018



# Machine Learning Algorithms To Combine Predictions

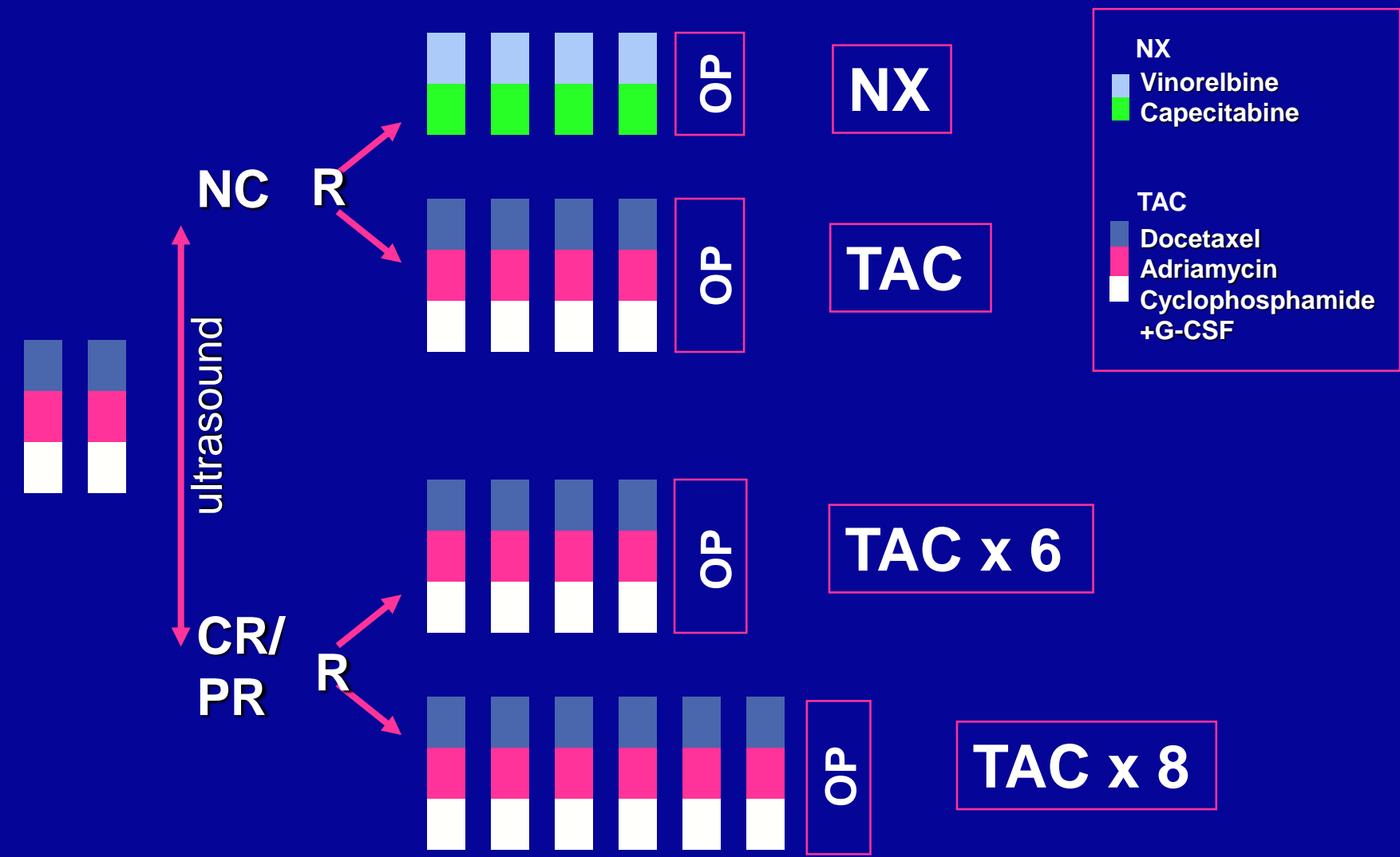
Example: Random Forest Approach



Groll et al. Technical University of Dortmund, MIT Technology Review, June 12, 2018

# Using Radiologic Response To Guide Neoadjuvant Chemotherapy: GEPAR-TRIO Trial

Core biopsy:  
uni/bilateral  
cT2-4  
cN0-3  
size  $\geq 2$  cm\*



\*excluding low risk (T2 + ER/PR pos. + cNO + G1/2 + > 35y.)

# Consequences of Response Guidance, by HR Status: GEPAR-TRIO Trial

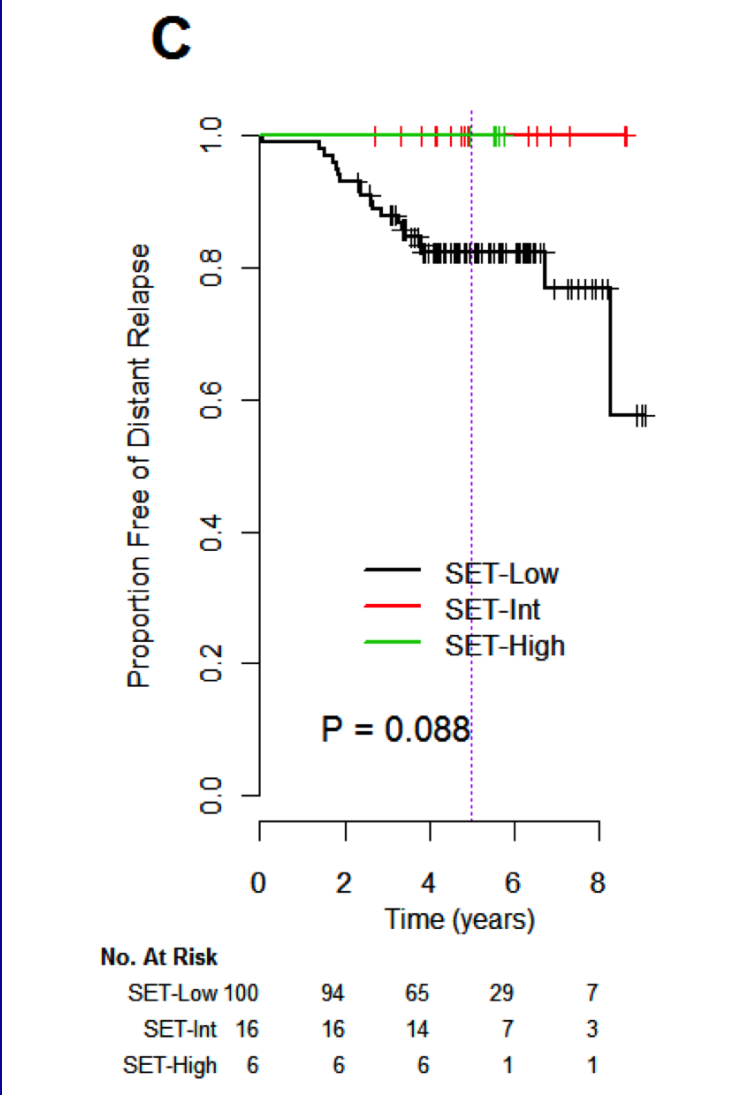
**Table 2.** ORs for pCR and HRs for DFS Between Response-Guided Compared With Conventional Chemotherapy and HRs for pCR According to Hormone Receptor Status and Phenotypes

Subgroups by HR and Phenotype*	No. of Patients	pCR Rate		pCR		DFS				
		No.	%	Conventional v Response Guided		pCR v No pCR		Conventional v Response Guided		
				OR	P	HR	P	HR	P	
HR										
Positive	1,295	105	8.2	1.34	.155	1.53	.122	0.56	< .001	
Negative	717	238	33.2	1.17	.330	4.89	< .001	0.94	.663	
Test for interaction							.004		.008	

# SET Index: ER-related Gene Expression w/o Proliferation

ER+, Stage II-III  
 66% clinically LN+  
 Neoadjuvant T/FAC chemotherapy  
 Surgery / XRT  
 Adjuvant Tam &/or AI  
 (N=122)

Association With Pathologic Response		
SET Class	pCR / RCB-I	Chi-square test
Low	35 / 100	NS
Intermediate / High	6 / 22	



# Response from Chemotherapy and Predicted SET

ER+, neoadjuvant T/FAC, adjuvant Tam &/or AI (N=122)

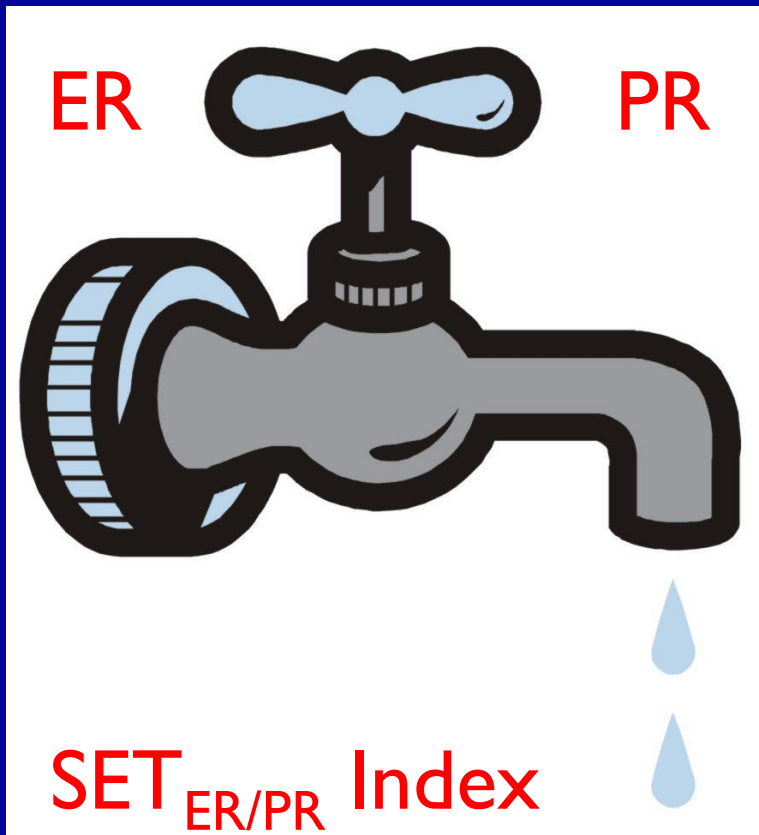
Cox Regression Model for DRFS			
Factor	HR	95% CI	P value
RCB Index	2.07	1.20 - 3.60	0.009
SET Index	0.19	0.05 - 0.69	0.011
RCB * SET	1.49	0.99 - 2.24	0.054

**RCB index = Residual Cancer Burden in breast and regional lymph nodes after completion of chemotherapy**

**SET index = genomic predicted Sensitivity to Endocrine Therapy**

# Sensitivity to Endocrine Therapy: $SET_{ER/PR}$ Index

## ER-Related Transcription in Primary Disease



Gene expression correlated with expression of ER & PR

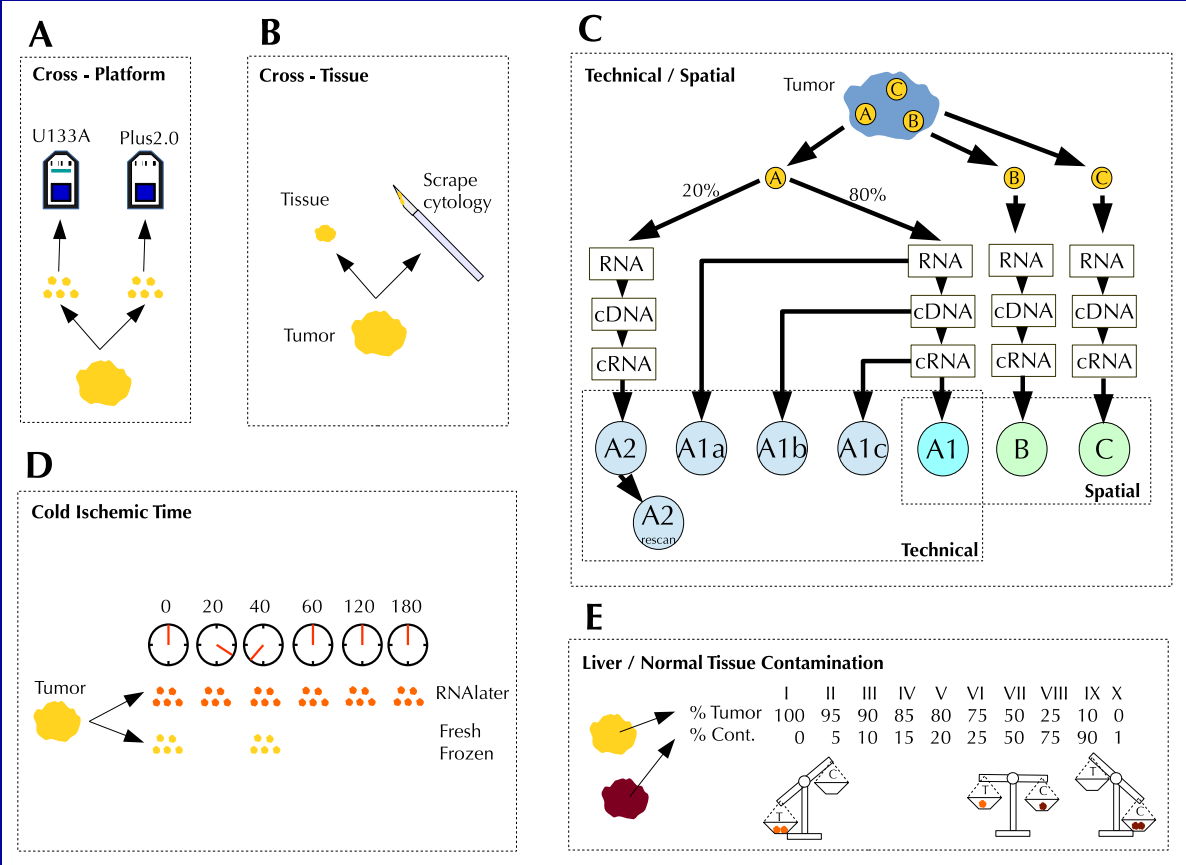
- Exclude proliferation-related genes
- Select if robust to pre-analytical and analytical conditions
- Measured relative to reference genes

165 → 18

Translation to assay for use with FFPE samples

- Simple workflow
- Affordable technology
- Robust technical measurements

# Biospecimen Studies To Filter Genes for the SET2,3 Test



## SET<sub>ER/PR</sub> Genes

ABAT, ADCY1, AZGP1, CA12, CD2, CD3D, DNAJC12, ESR1, KCNE4, MAPT, MRPS30, NAT1, NPY1R, PDZK1, QDPR, SCUBE2, SLC39A6, STC2

## Reference Genes

AK2, APPBB2, ATP5J2, DARS, LDHA, TRIM2, UBE2Z, UGP2, VDAC2, WIPF2

## RNA4

ESR1, ERBB2, PGR, AURKA

pT pN

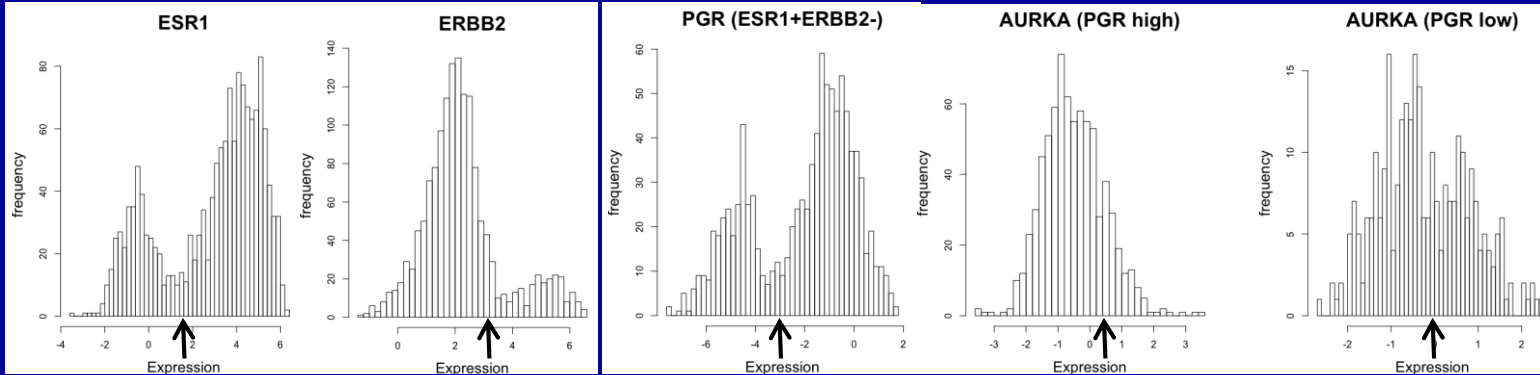
SET<sub>ER/PR</sub> Index

Prognostic Risk

# SET<sub>2,3</sub> Test: Risk-Adjusted Cutpoints for the SET<sub>ER/PR</sub> Index

## RNA4

## Risk Score

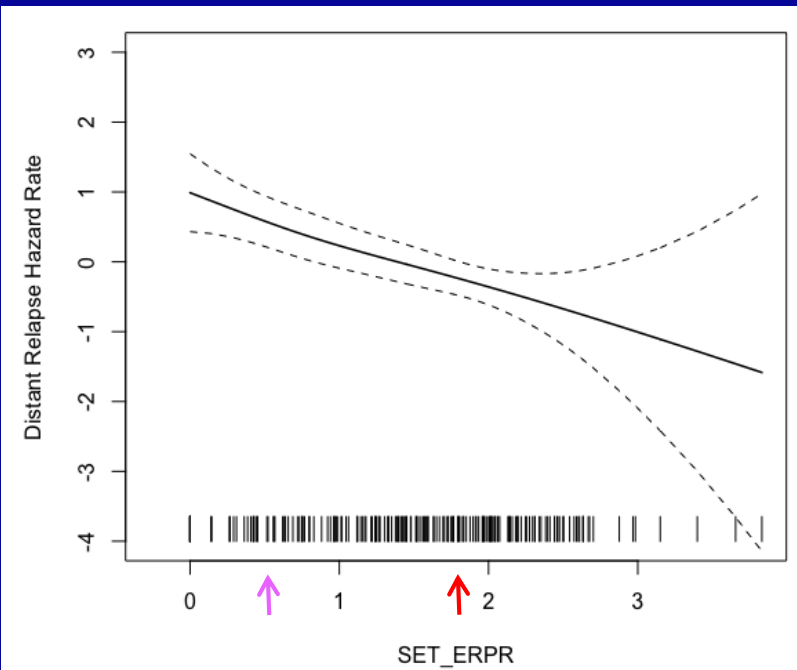


cN Stage: cN- (0 votes), cN+ (2 votes)

cT Stage: cT1-2 (0 votes), cT3-4 (1 vote)

RNA4: low risk (0 votes), not low risk (1 vote)

Low Risk if  $\leq 1$  vote, High Risk if  $\geq 2$  votes



SET<sub>ER/PR</sub> Index Cutpoint = 0.5 if Low Risk

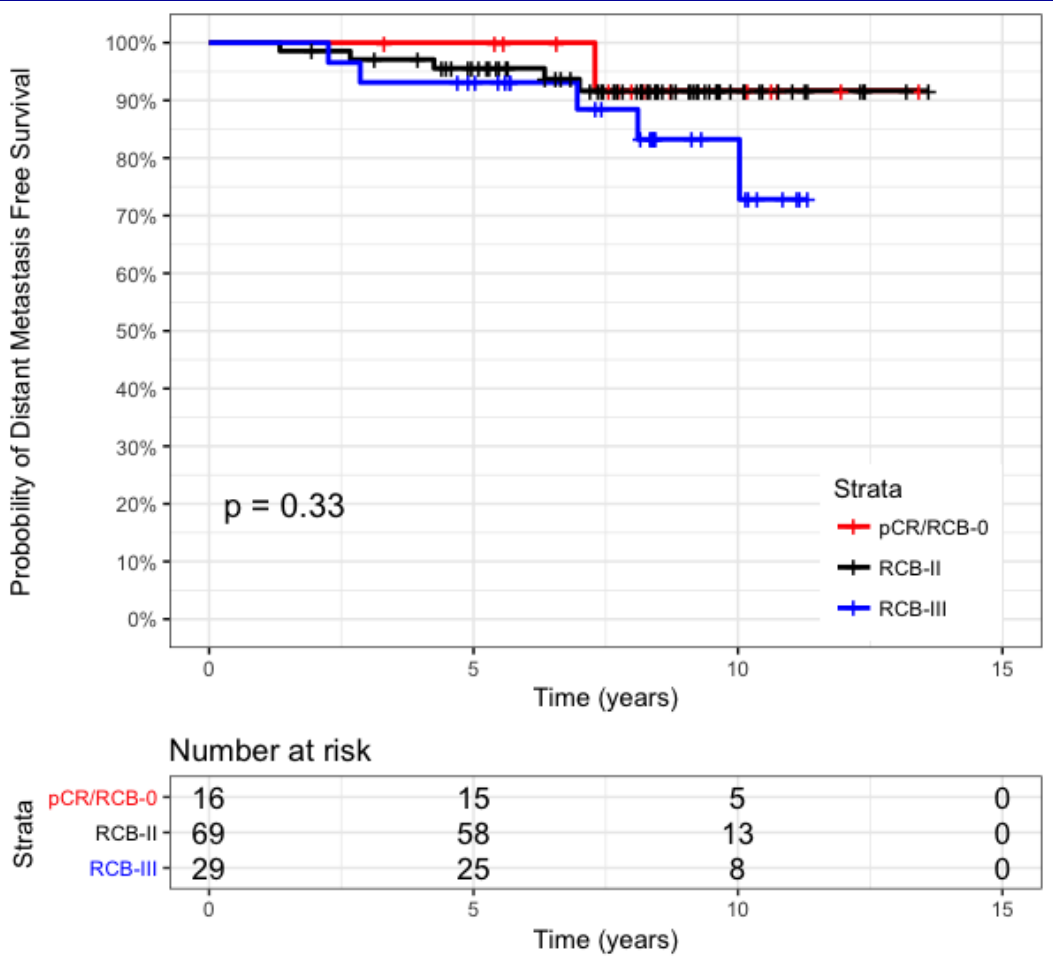
SET<sub>ER/PR</sub> Index Cutpoint = 1.8 if High Risk



# SET2,3 Classes: Prognostic Influence of Response To NAC

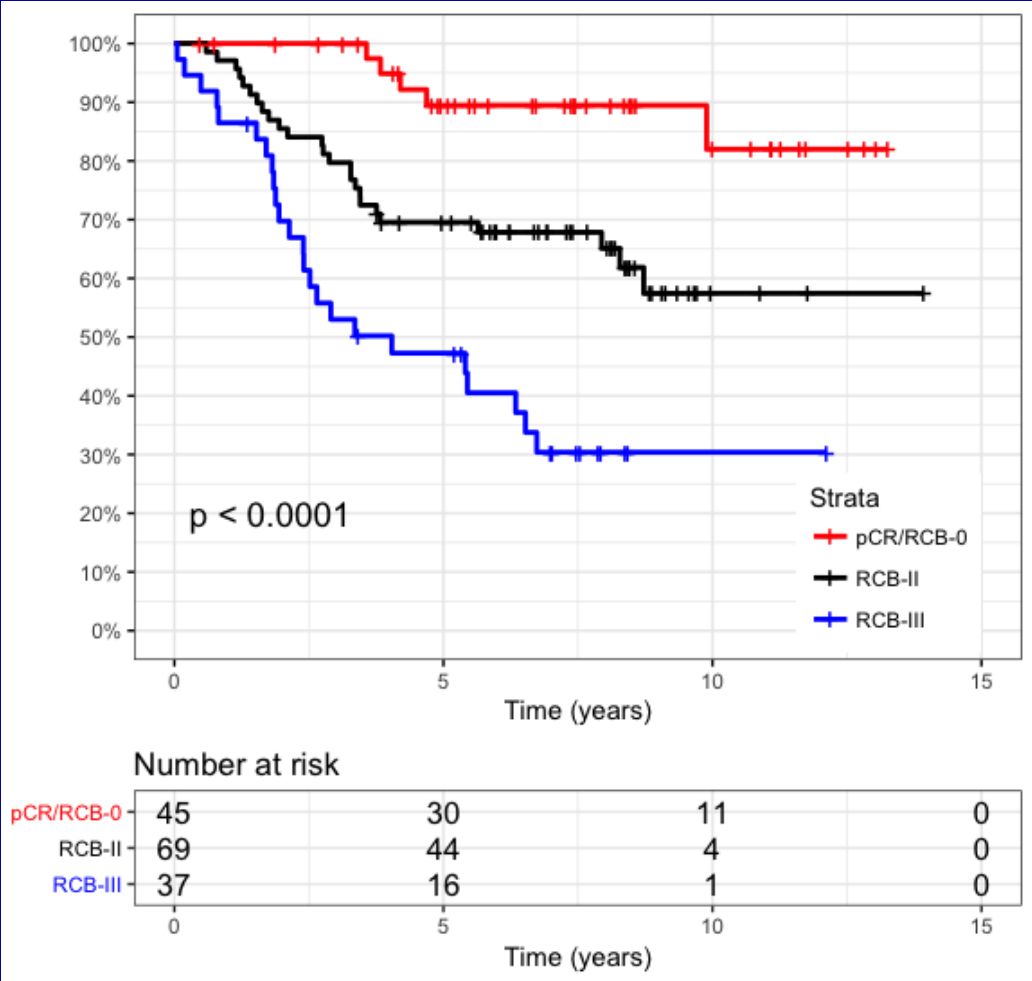
43%

High SET2,3 (low risk, n= 114)



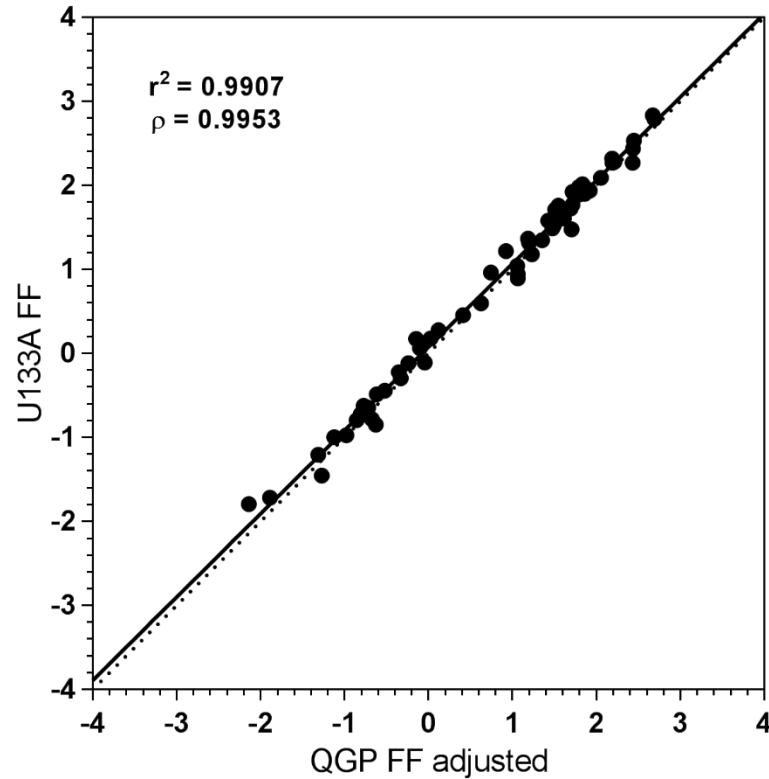
57%

Low SET2,3 (high risk, n= 151)

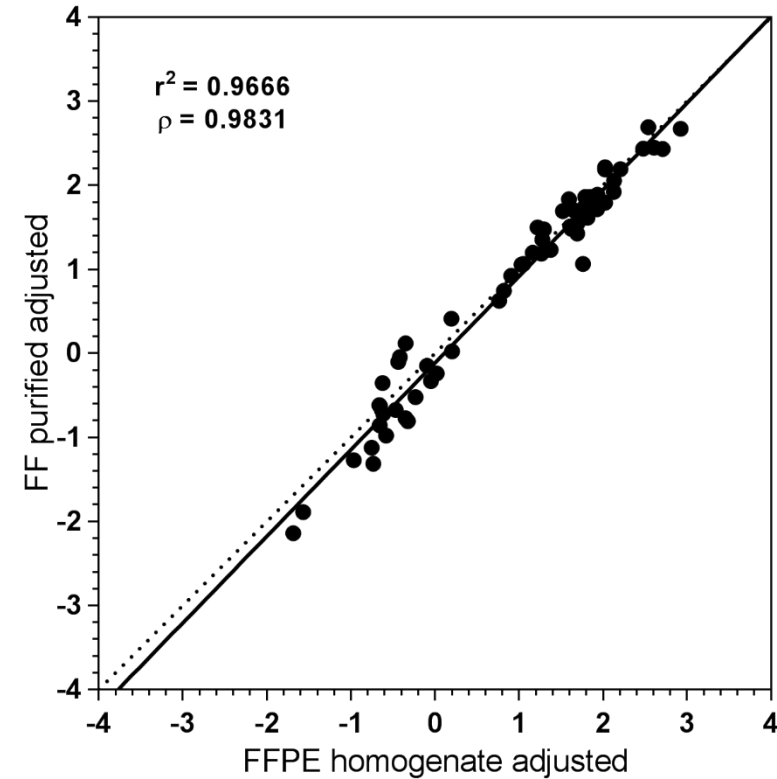


# Translation of SET2,3 to a Customized Test Format QGP

Inter-Platform  
Using Fresh Frozen Samples



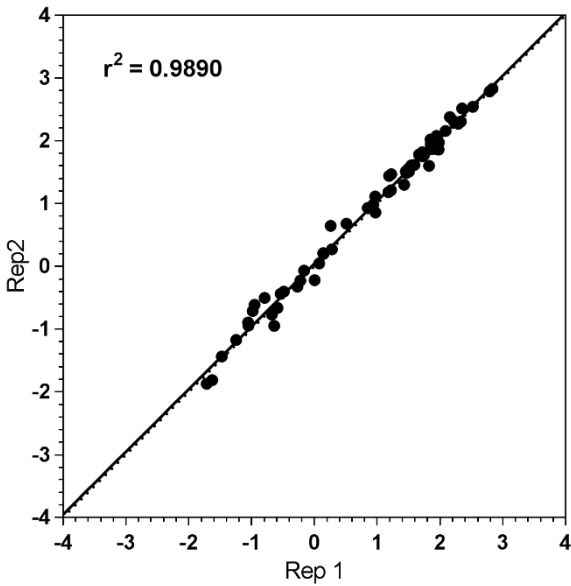
Sample Fixation  
Fresh Frozen vs FFPE



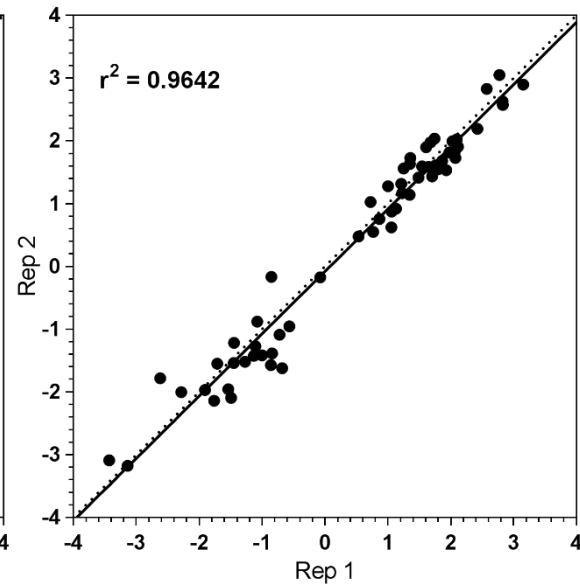
QuantiGene Plex (QGP) assay using Luminex MagPix device, lysis homogenate from unstained tissue section

# Technical Reproducibility of SET2,3 Assay

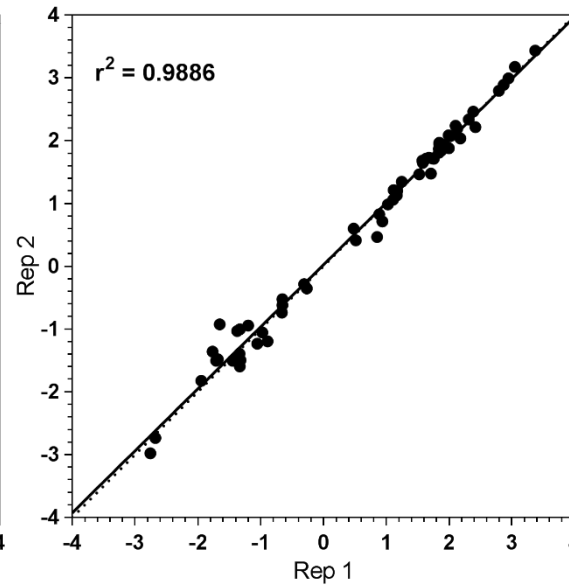
U133A Microarrays  
RNA from fresh frozen tissue



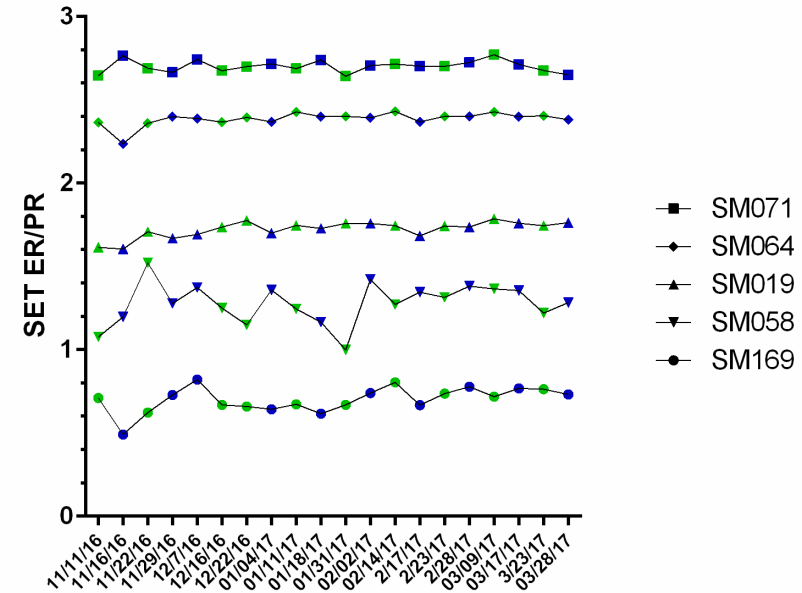
NANOSTRING  
RNA from FFPE tissue



QuantiGene Plex  
FFPE tissue slides

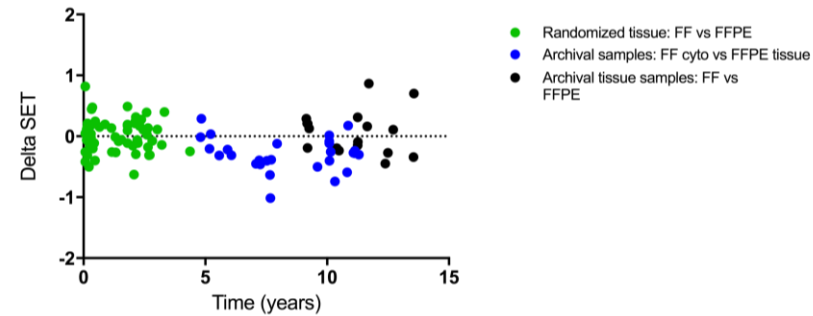


Timeline Reproducibility of QGP

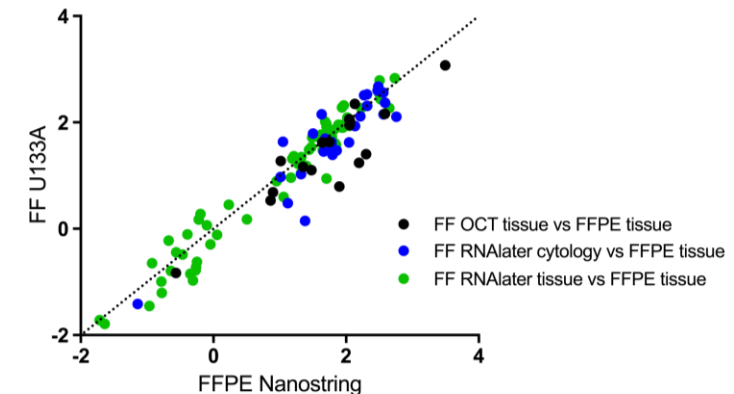
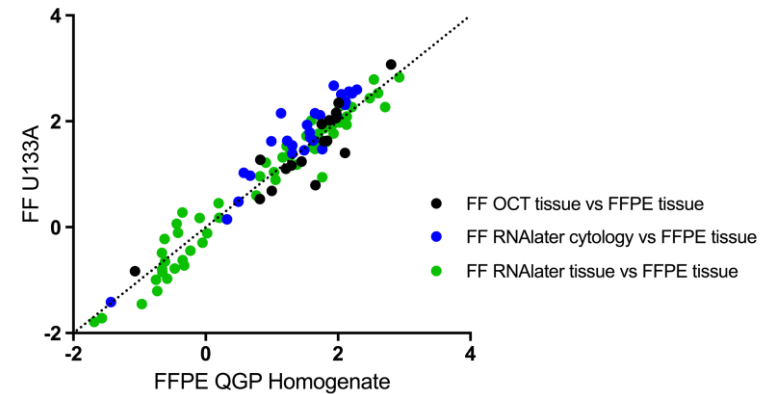
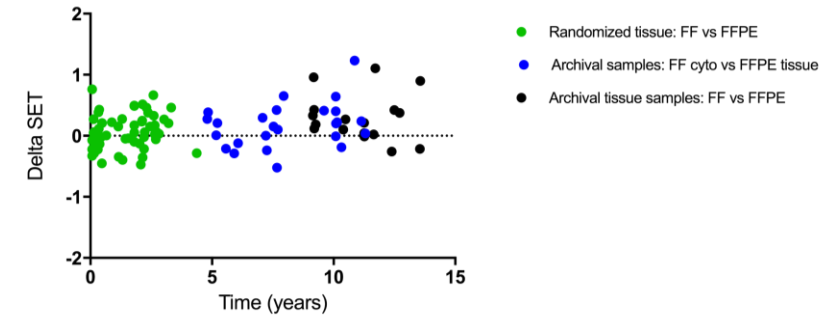


# SET2,3 Assay Performance: Older FFPE & Frozen Samples

QuantiGene Plex using FFPE  
Duration of tissue storage  
Difference (delta) from frozen tissue U133A

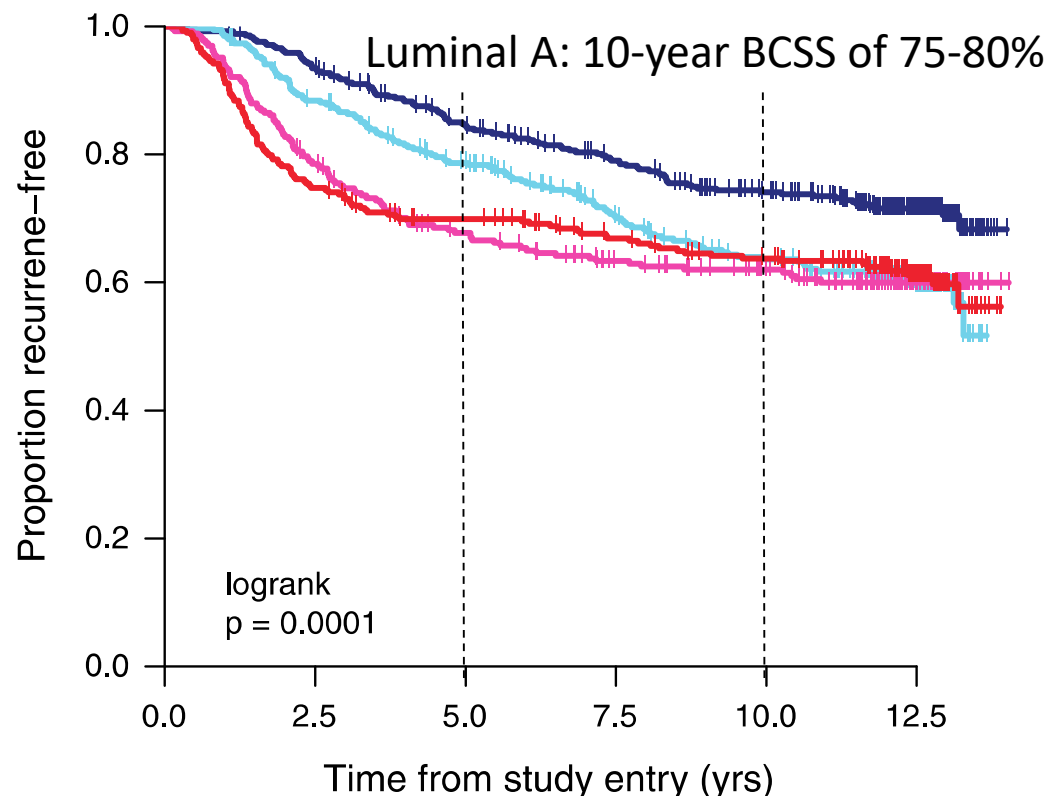


NANOSTRING using FFPE  
Duration of tissue storage  
Difference (delta) from frozen tissue U133A

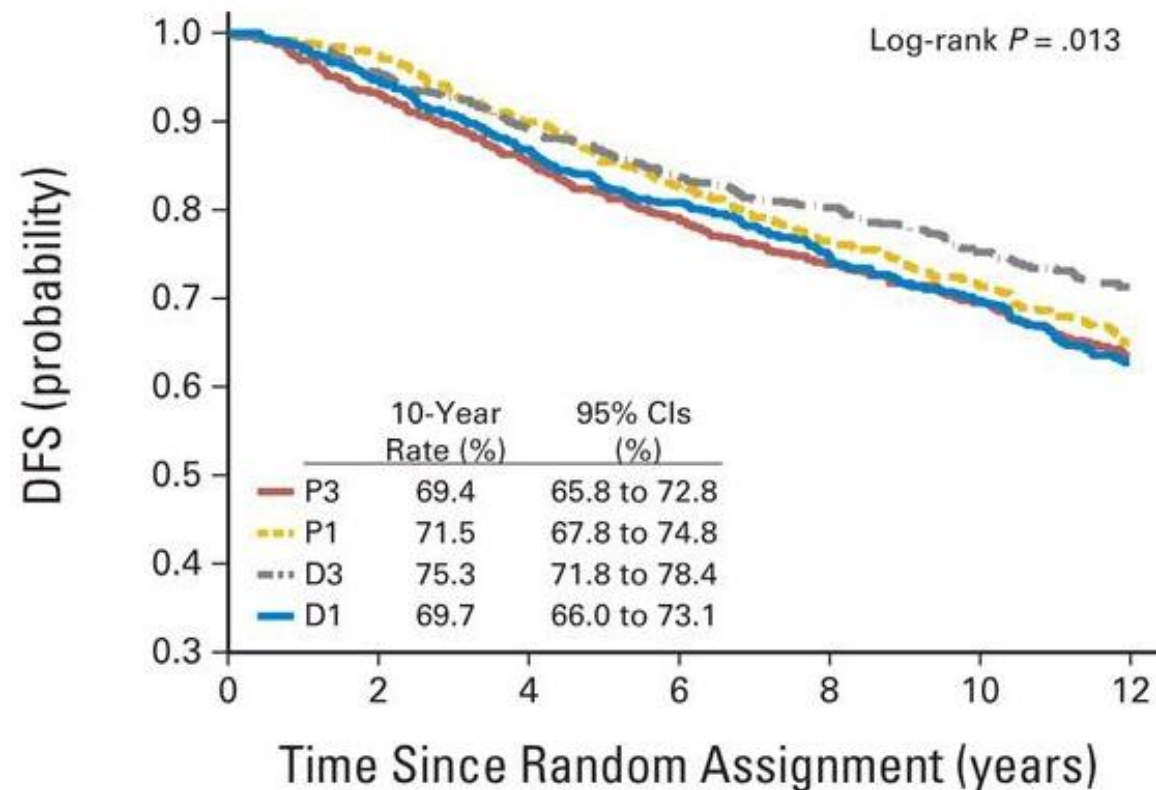


# Validation Strategy: Stratify Long-Term Risk

**CALGB-9741, AC/T chemo, all subtypes**



**ECOG E-1199, AC/T chemo, HR+/HER2-**



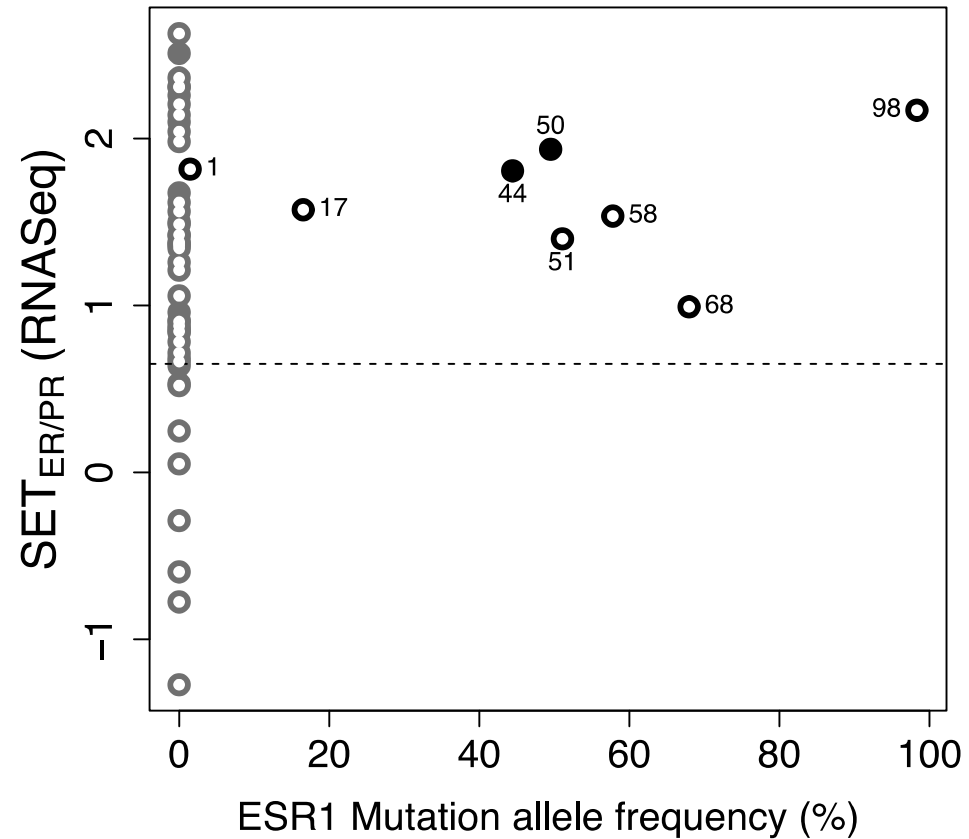
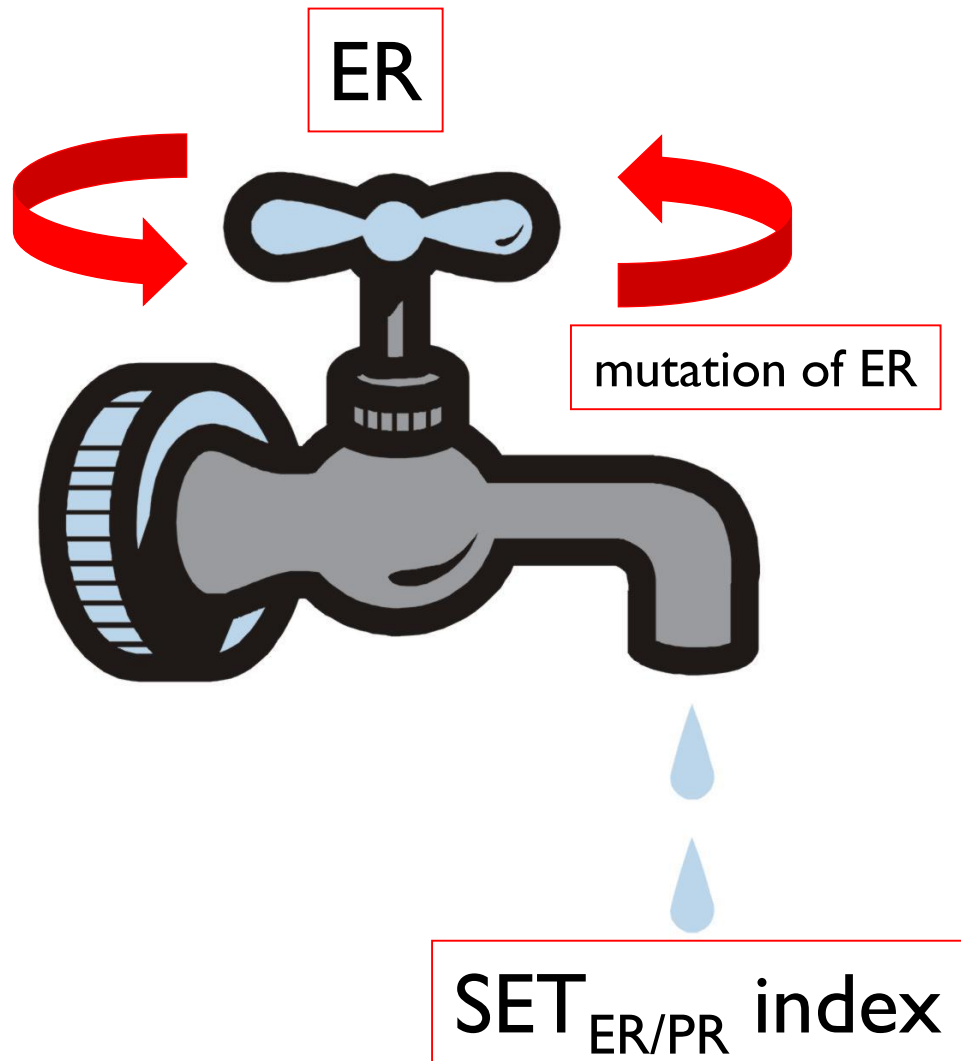
	Number at risk					
	0	2.5	5.0	7.5	10.0	12.5
LuminalA	414	385	335	293	256	125
LuminalB	338	295	253	203	172	81
Her2Enriched	266	207	171	146	126	57
BasalLike	293	219	195	173	159	69

	No. at risk						
	0	2	4	6	8	10	12
P3	715	653	572	505	452	377	197
P1	706	673	600	525	458	385	189
D3	731	683	621	562	513	436	223
D1	735	683	599	523	458	381	187

## The next wave of treatments for Stage II-III HR+ breast cancer:

- Concurrent cdk4/6 inhibition with endocrine therapy
- Concurrent inhibition of PI3-kinase or pathway with endocrine therapy
- De-escalation to hormonal therapy alone
- Less intense / less toxic chemotherapy regimens (calibrated to lower risk)

# Customized Transcriptomic Assay: Quantify Mutant Transcript Load + Gene Expression Activity



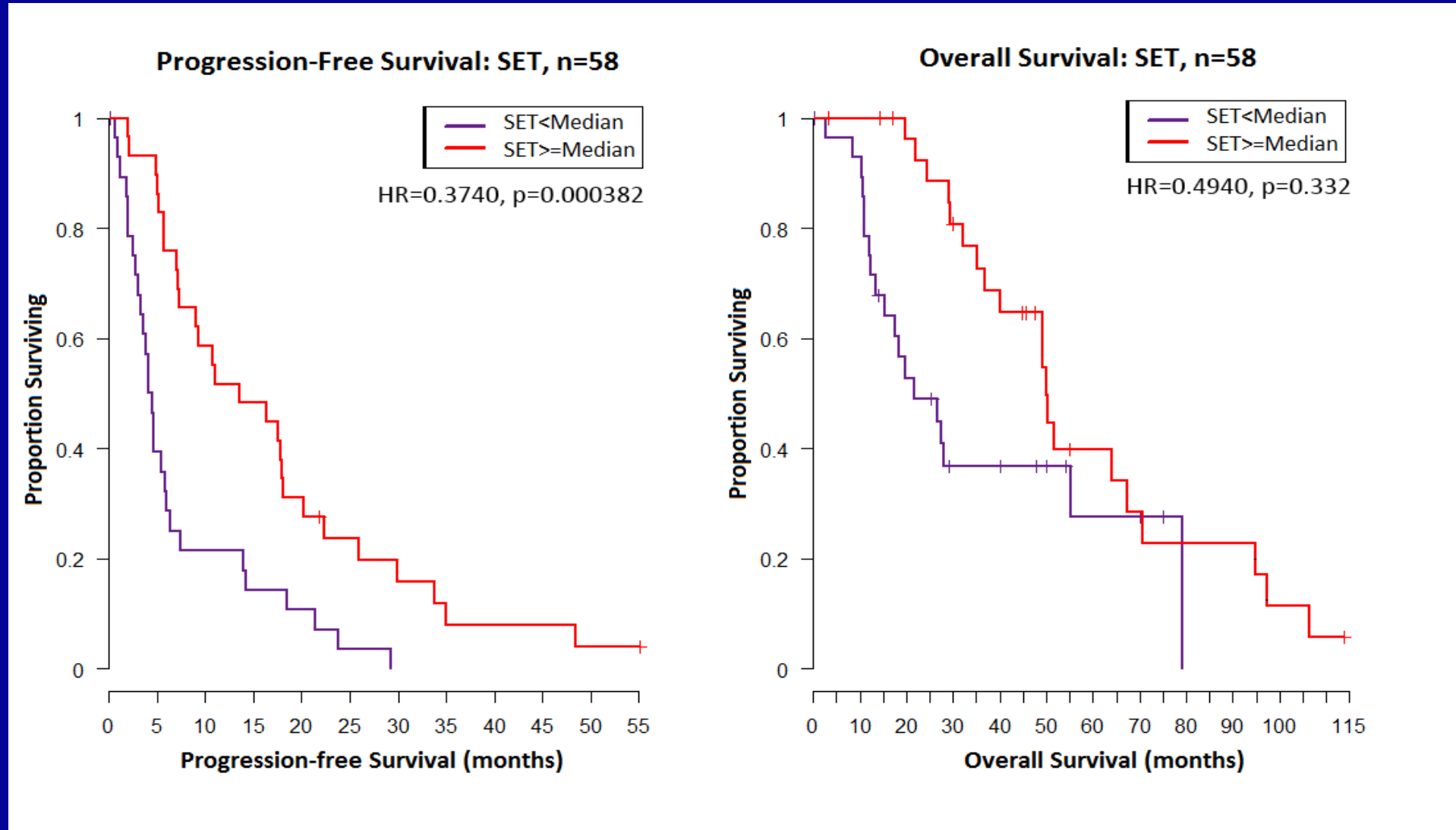
*ESR1* LBD  
SET index GES (n=18)

*PIK3CA* hotspots  
PI3K index GES (n=10)

*AKT1* hotspot  
*ERBB2* hotspot  
*FGFR1* expression

Reference genes (n=10)

# Prognostic: Endocrine Treatment of Metastatic Breast Cancer



Independent of standard clinical and pathologic variables

*Lau et al. Am Assoc Cancer Res 2016*



# Summary

**Pathologic response to chemotherapy is meaningful in all subtypes**

- **Standardized measurement of RCB is prognostic and generalizable**

## **HR+/HER2-**

- **Residual risk is defined by c-Stage, prognostic biology, sensitivity to chemotherapy, endocrine therapy, other treatments**
  - **persists over the long-term**
- **Sequential synergy derived from chemotherapy followed by endocrine therapy**
- **Endocrine-related transcription is clinically relevant in all stages of disease**
  - **Biospecimen research helped us to define a robust biomarker for clinical studies**

## **TNBC**

- **Extent of residual disease after chemotherapy is the most important prognostic information**
- **Predicting response/resistance to chemotherapy-based treatments is challenging**

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