



ESMO Preceptorship Programme

Breast Cancer

Multidisciplinary management, standards of care, therapeutic targets and future perspectives

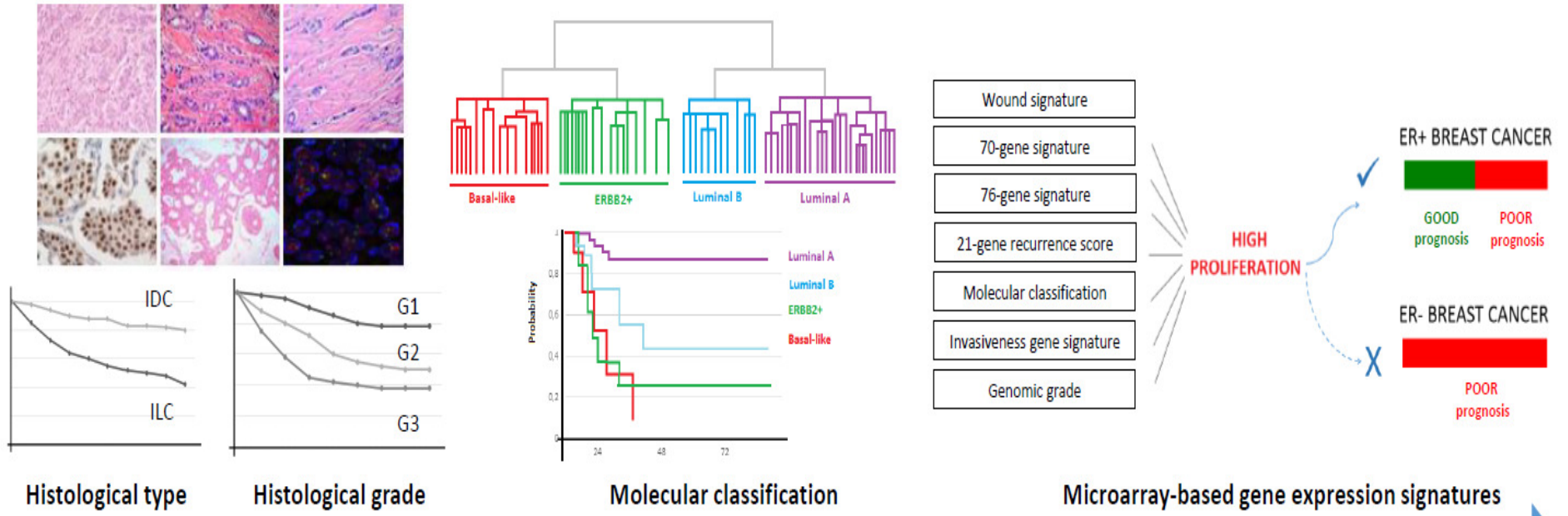
Lisbon, Portugal

16-17 September 2016



BREAST CANCER CLASSIFICATION: TRADITIONAL PATHOLOGY AND MOLECULAR SUBTYPES

Prof. Fernando Schmitt
Director of Department of Pathology and Medicine
Laboratoire National de Santé, Luxembourg
General-Secretary of the International Academy of Cytology



12th St. Gallen International Breast Cancer Conference
 Primary Therapy of Early Breast Cancer with Treatment Consensus Update
 16-19 March 2011
 St. Gallen, Switzerland

Approximately 500 GENES Intrinsic gene list

4 IHC MARKERS

- ER
- PR
- HER2
- Ki67

Surrogate for molecular classification

13th St. Gallen International Breast Cancer Conference
 Primary Therapy of Early Breast Cancer Evidence, Controversies, Consensus
 13-16 March 2013
 St. Gallen, Switzerland

Intrinsic subtype	Pathologic definition
Luminal A	ER >1% PR >1% Ki-67 <14%? <20%? HER2 negative
Luminal B	ER >1% PR >1% Ki-67 >20%? HER2 negative

MOLECULAR TESTS

Recurrence score(s)

Molecular tests for risk prediction

St Gallen 2015 Viena

Gene expression data

Copy number data

HER2 mutations

IntClusts

NGS ...

TCGA results: IntClust, mutation profile *et al.*



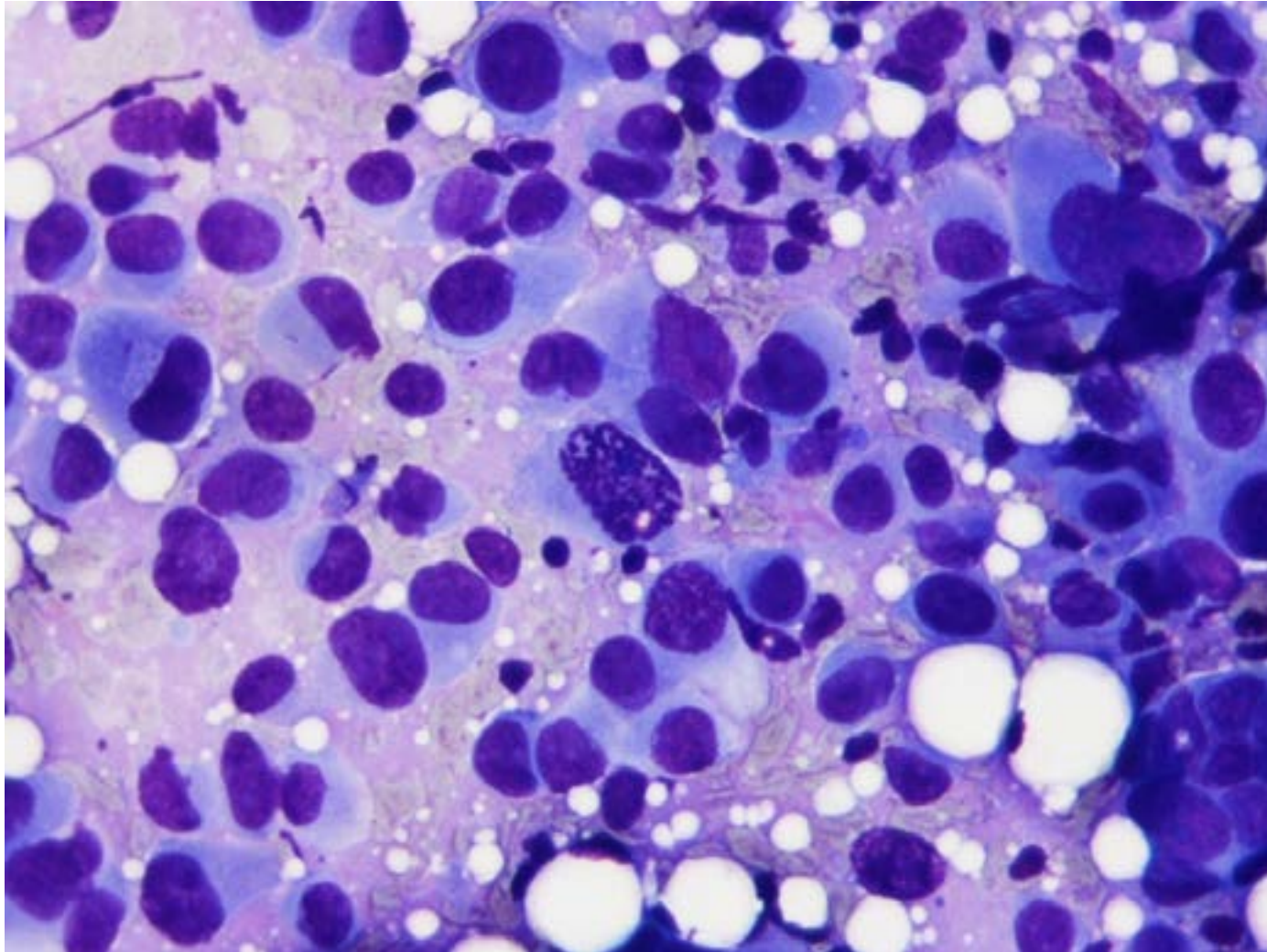
Why do we need a classification?

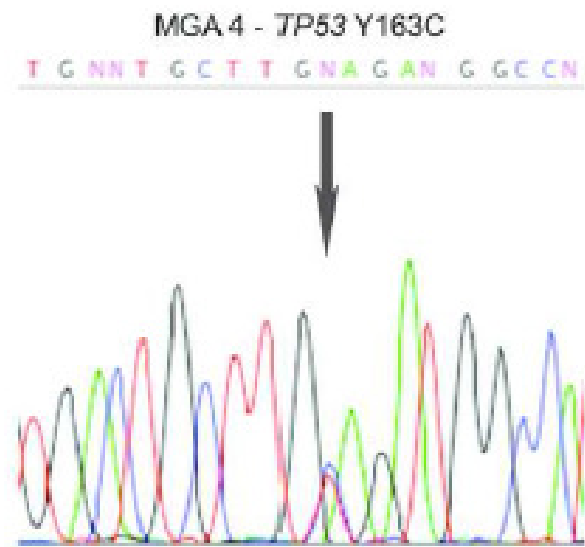
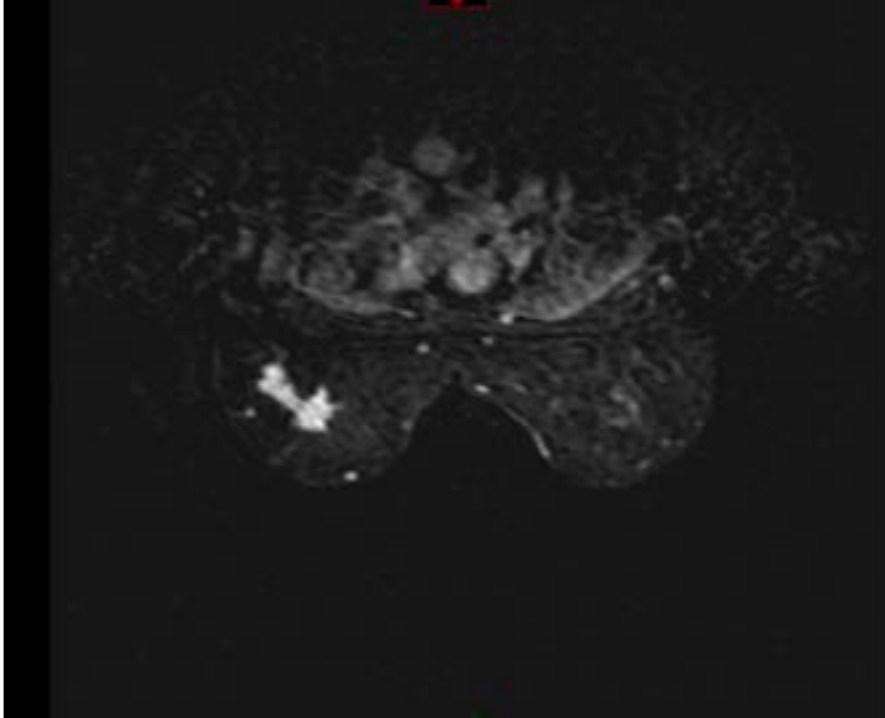
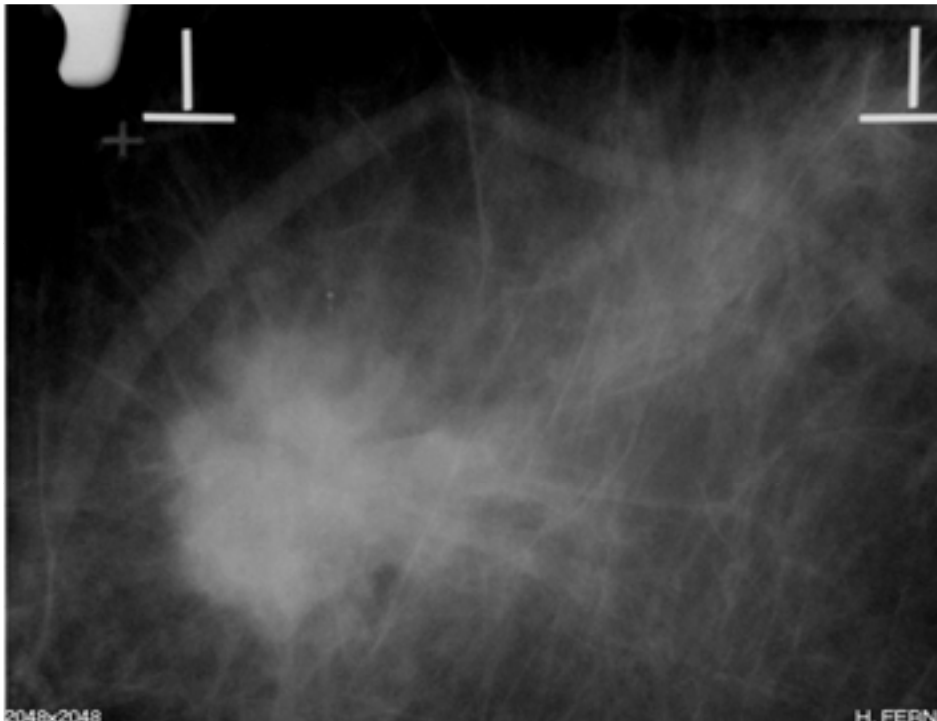
Aim 1: **Diagnosis**

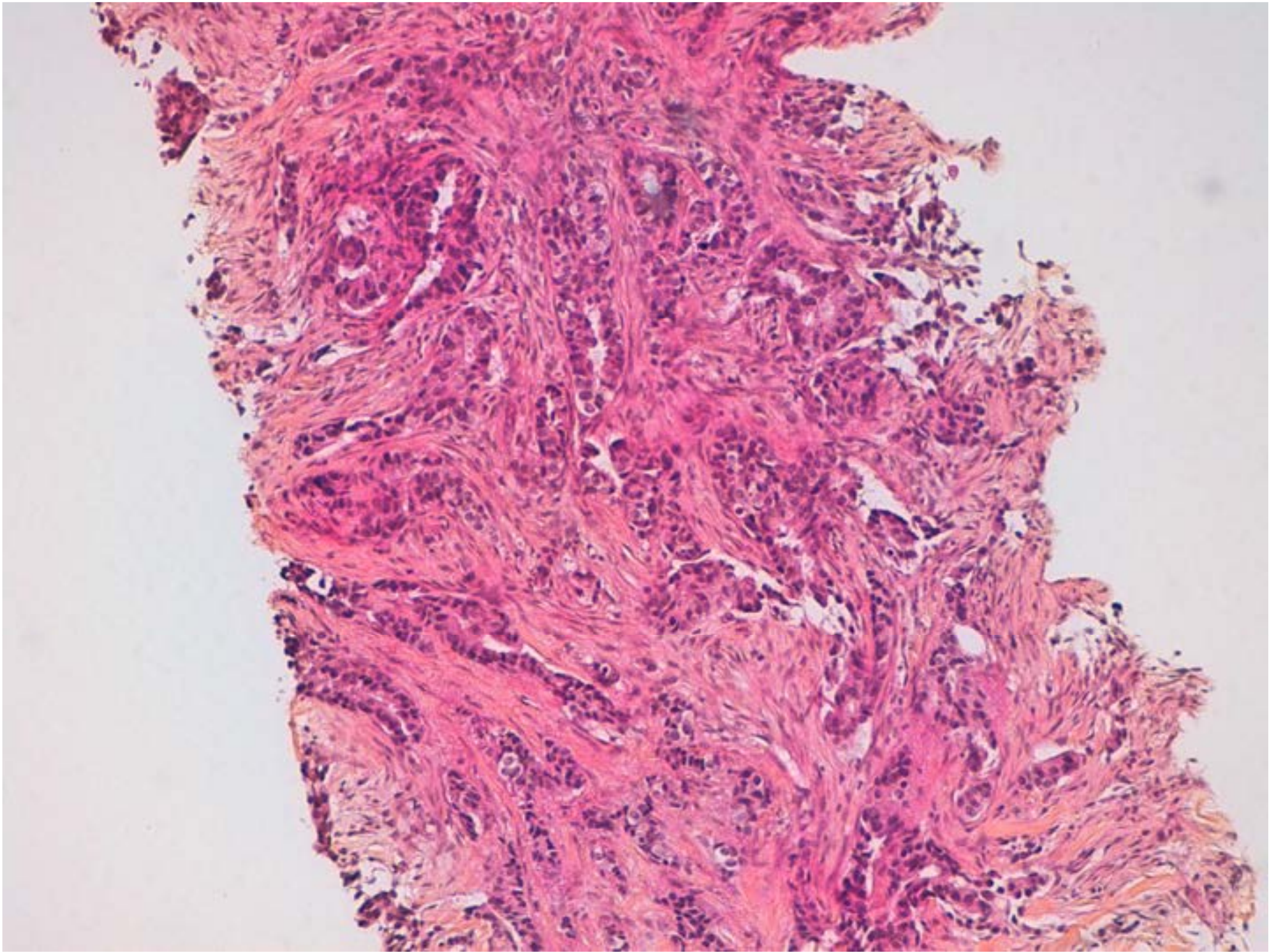
Aim 2: **Prognosis**

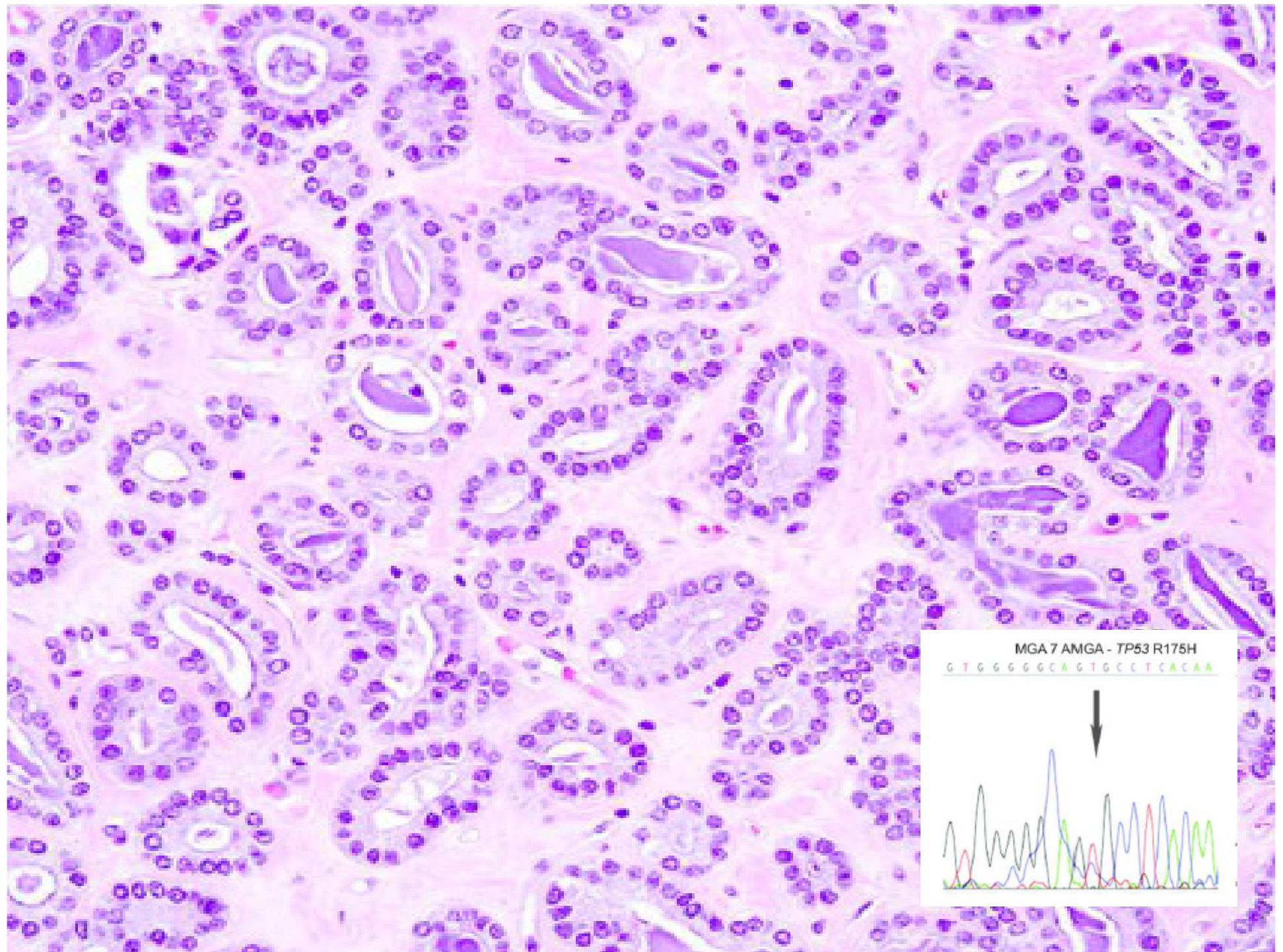
Aim 3: **Prediction**

Breast cancer diagnosis is morphological



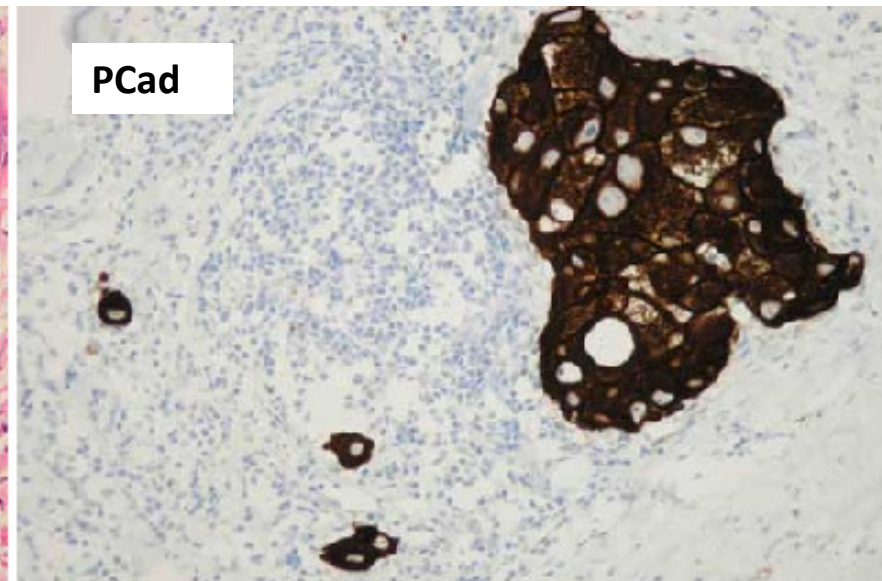
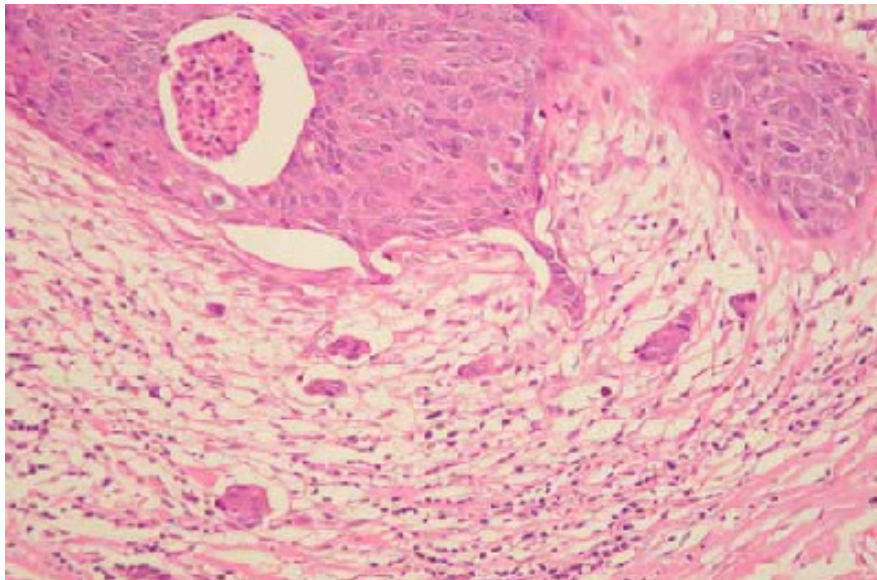






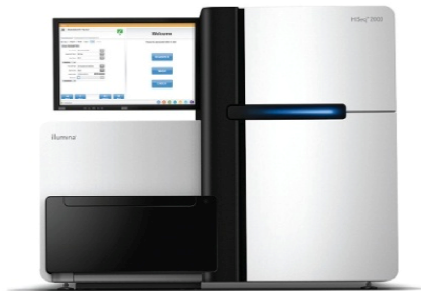
Breast cancer diagnosis is morphological

Microinvasion

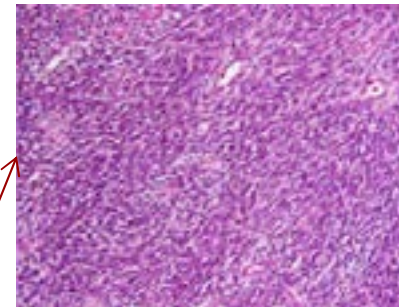
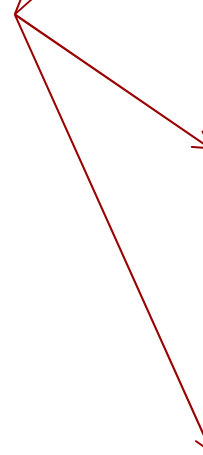


Molecular results without pathology can be messy...

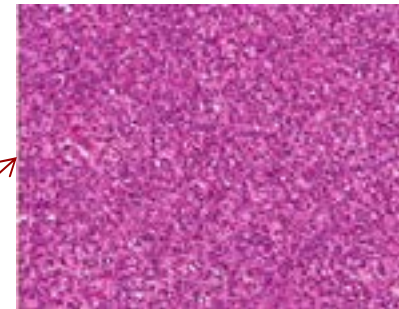
Biopsy



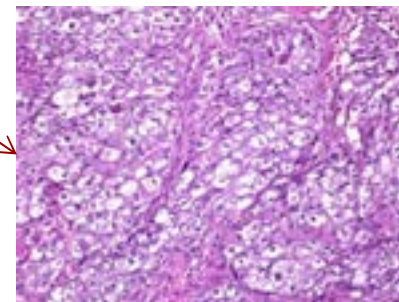
ETV6-NTRK3



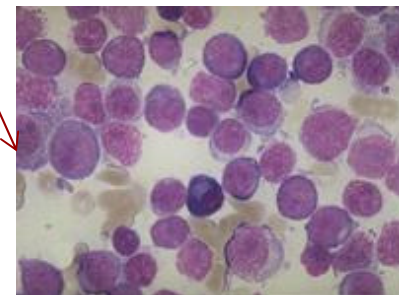
Congenital fibrosarcoma



Cellular mesoblastic nephroma

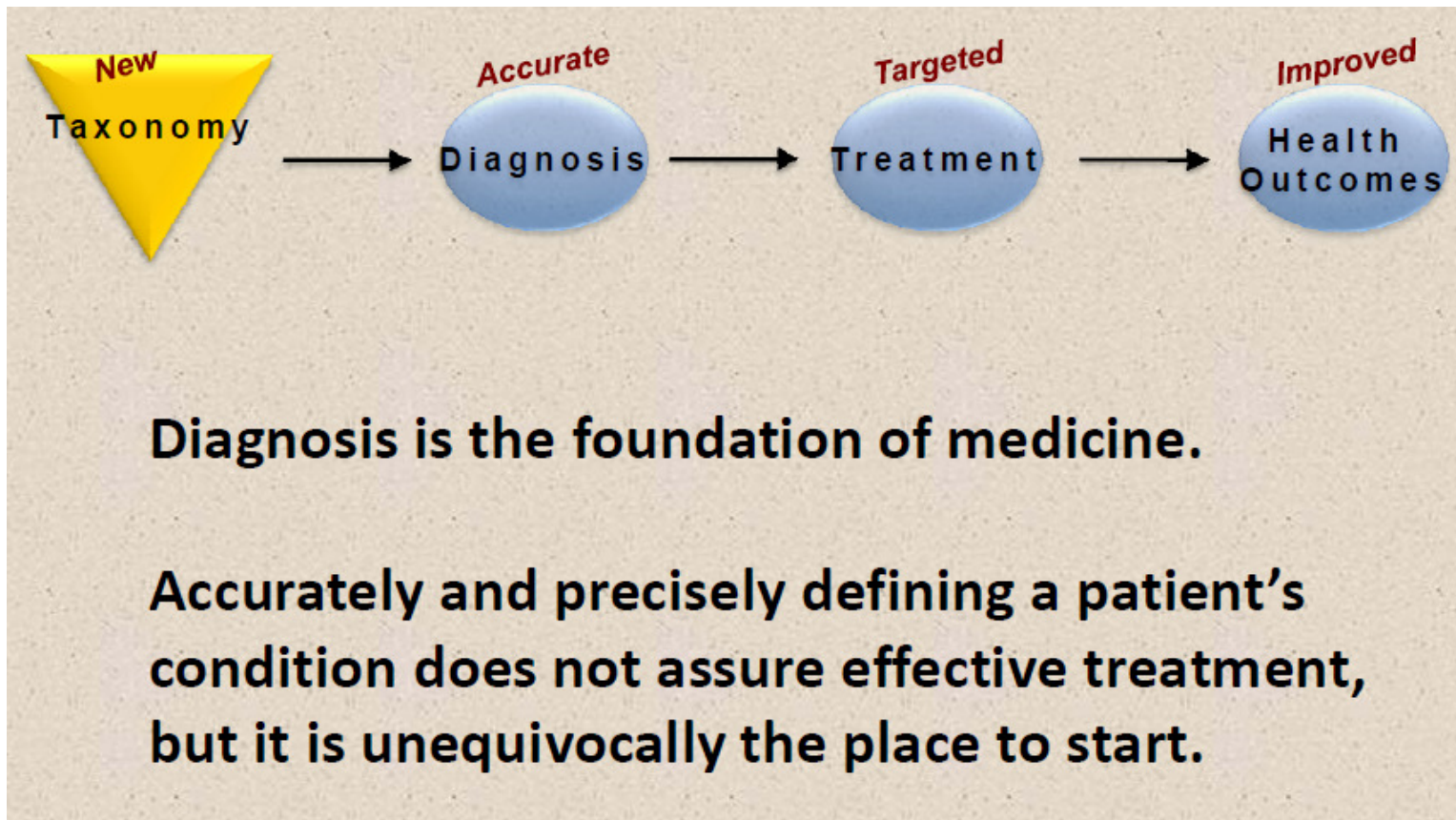


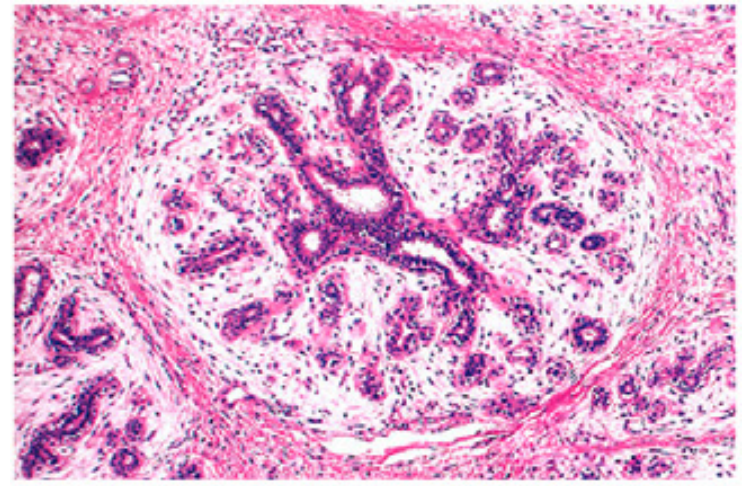
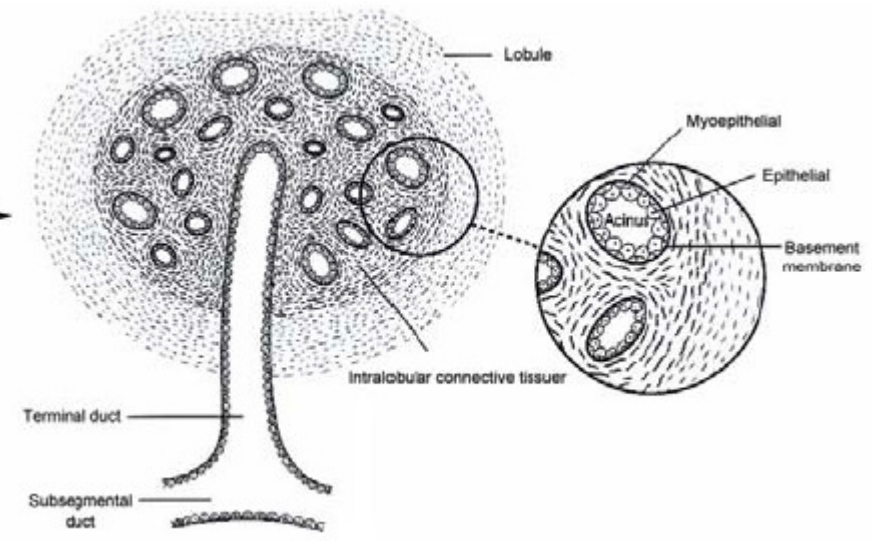
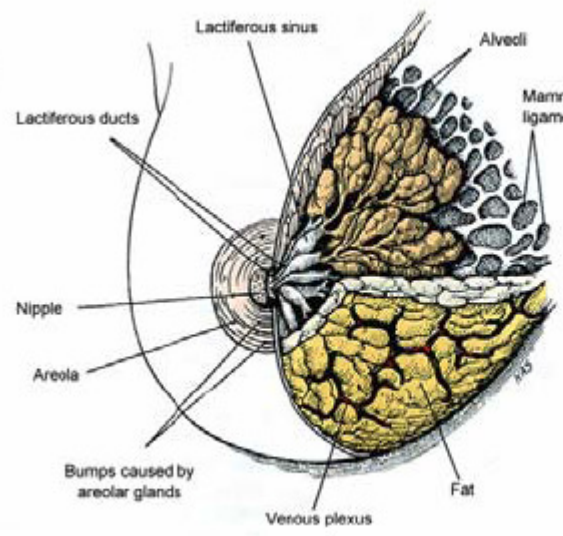
Secretory carcinoma



Acute myeloid leukaemia

Precision Medicine





Putative Model to explain Breast Cancer Molecular Signatures

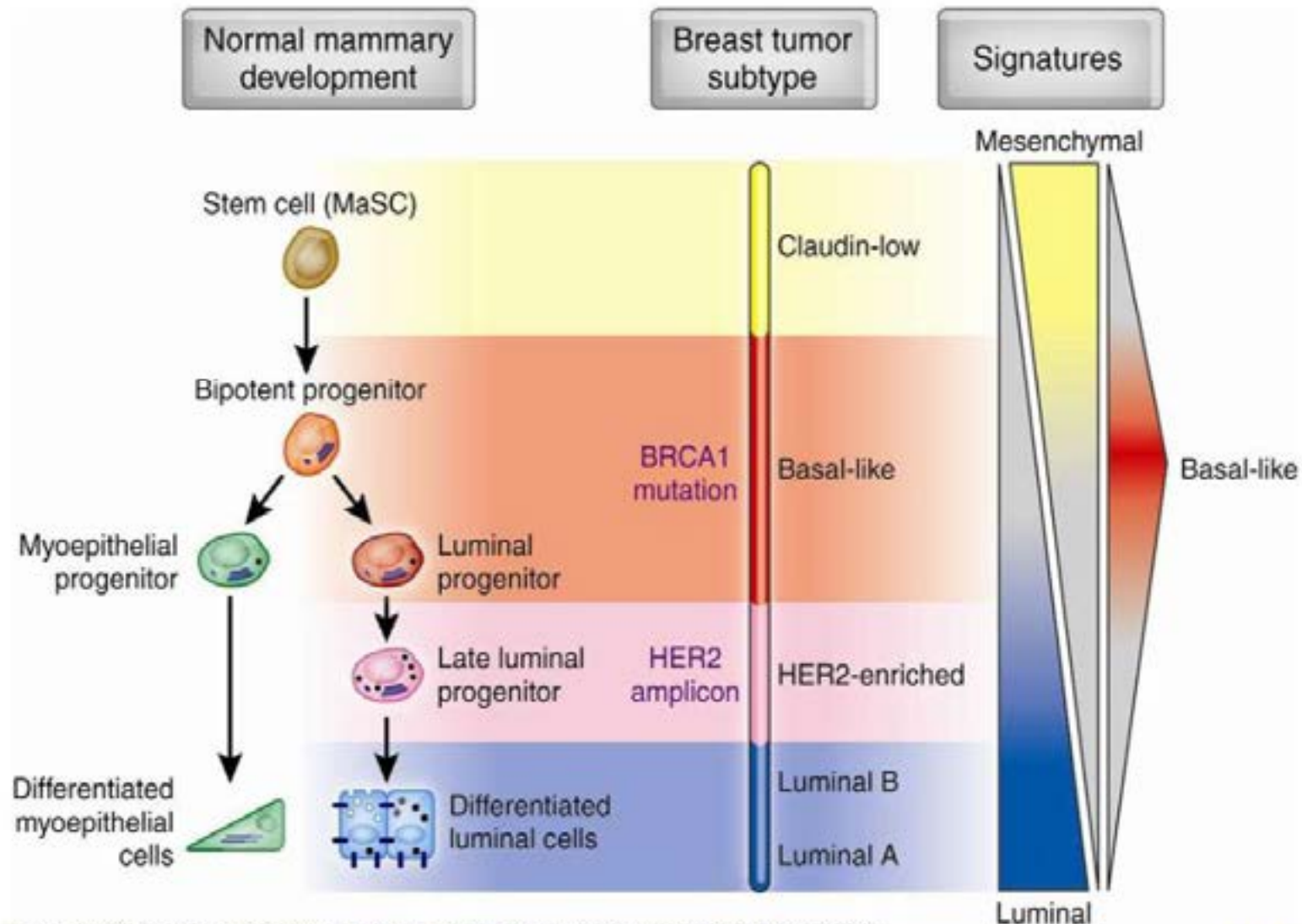
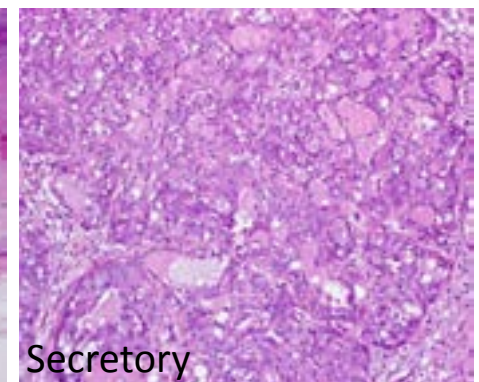
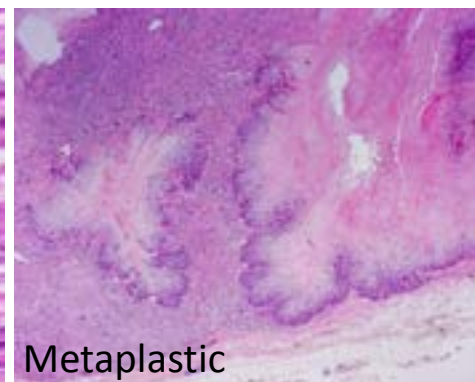
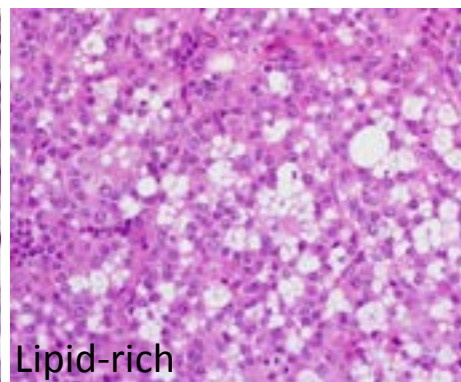
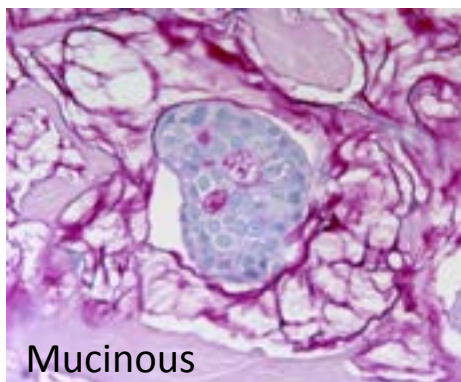
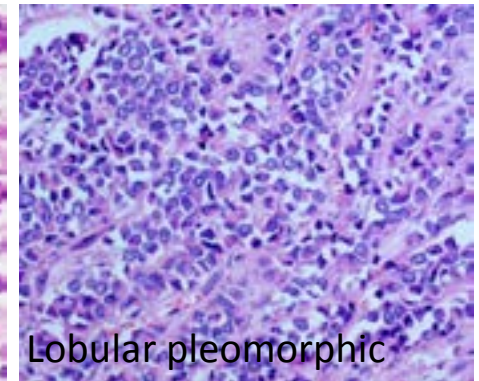
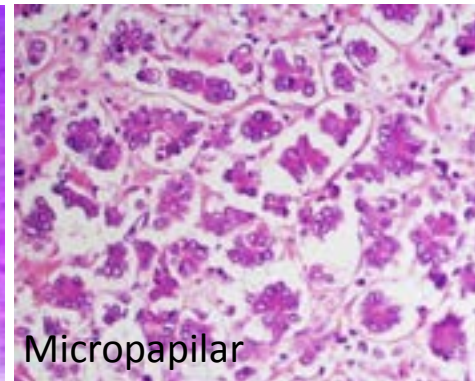
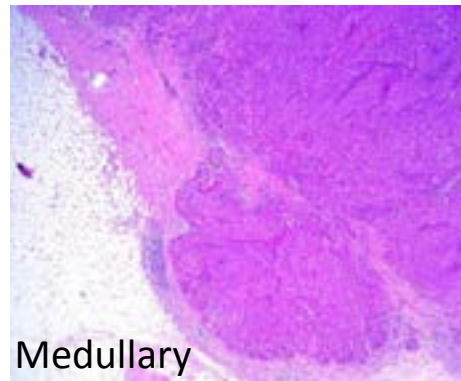
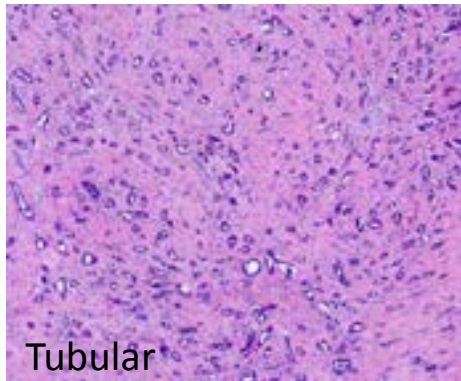
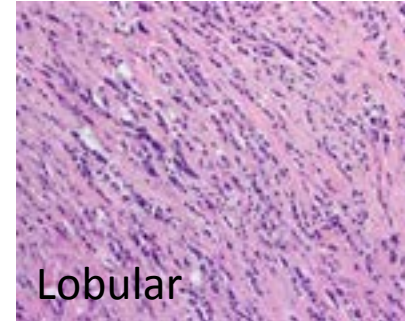
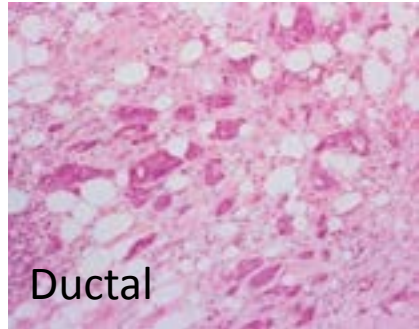


Image by Katie Vicari, from Prat and Perou, Nature Medicine, Aug;15(8):842-4 (2009)

Histological types of breast carcinoma



WHO Classification of Tumours of the Breast

(Edited by David R. Lakhani, Lee D. Ellis, Stewart J. Schnitt, Pradyumn Kumar, Marc J. van de Vijver)

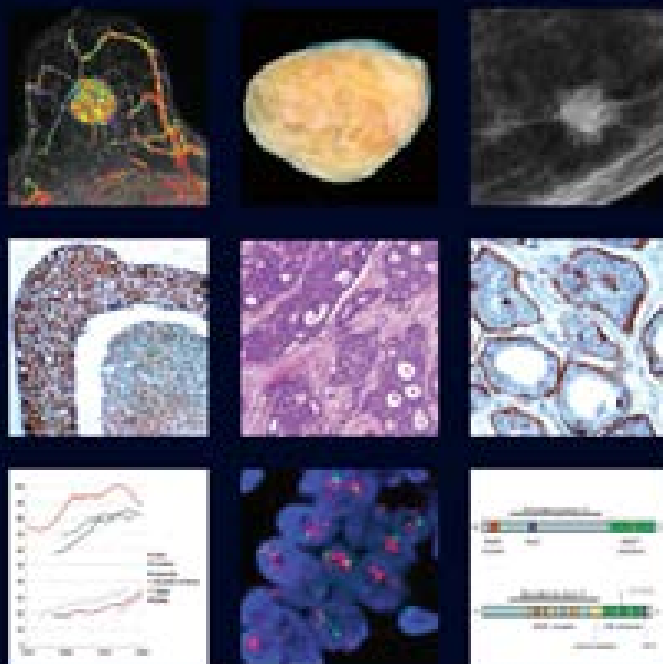
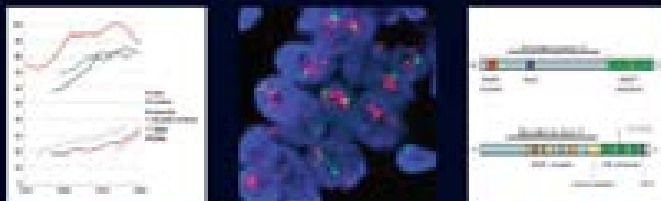
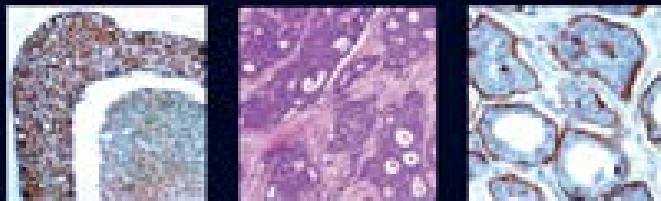


Table 1. Invasive breast carcinomas (without microinvasive carcinoma and invasive papillary lesions)

Type	Classification
Invasive carcinoma of no special type (NST)	8500/3
Pleomorphic carcinoma	8522/3
Carcinoma with osteoclast-like stromal giant cells	8035/3
Carcinoma with choriocarcinomatous features	
Carcinoma with melanotic features	
Invasive lobular carcinoma	8520/3
Classic lobular carcinoma	
Solid lobular carcinoma	
Alveolar lobular carcinoma	
Pleomorphic lobular carcinoma	
Tubulolobular carcinoma	
Mixed lobular carcinoma	
Tubular carcinoma	8211/3
Cribriform carcinoma	8201/3
Mucinous carcinoma	8480/3
Carcinoma with medullary features	
Medullary carcinoma	8510/3
Atypical medullary carcinoma	8513/3
Invasive carcinoma NST with medullary features	8500/3
Carcinoma with apocrine differentiation	
Carcinoma with signet-ring-cell differentiation	
Invasive micropapillary carcinoma	8507/3
Metaplastic carcinoma of no special type	8575/3
Low-grade adenosquamous carcinoma	8570/3
Fibromatosis-like metaplastic carcinoma	8572/3
Squamous cell carcinoma	8070/3
Spindle cell carcinoma	8032/3
Metaplastic carcinoma with mesenchymal differentiation	
Chondroid differentiation	8571/3
Osseous differentiation	8571/3
Other types of mesenchymal differentiation	8575/3
Mixed metaplastic carcinoma	8575/3
Myoepithelial carcinoma	8982/3
<i>Epithelial-myoepithelial tumors</i>	
Adenomyoepithelioma with carcinoma	8983/3
Adenoid cystic carcinoma	8200/3
<i>Rare types</i>	
Carcinoma with neuroendocrine features	
Neuroendocrine tumor, well-differentiated	8246/3
Neuroendocrine carcinoma poorly differentiated (small cell carcinoma)	8041/3
Carcinoma with neuroendocrine differentiation	8574/3
Secretory carcinoma	8502/3
Invasive papillary carcinoma	8503/3
Acinic cell carcinoma	8550/3
Mucoepidermoid carcinoma	8430/3
Polymorphous carcinoma	8525/3
Oncocytic carcinoma	8290/3
Lipid-rich carcinoma	8314/3
Glycogen-rich clear cell carcinoma	8315/3
Sebaceous carcinoma	8410/3

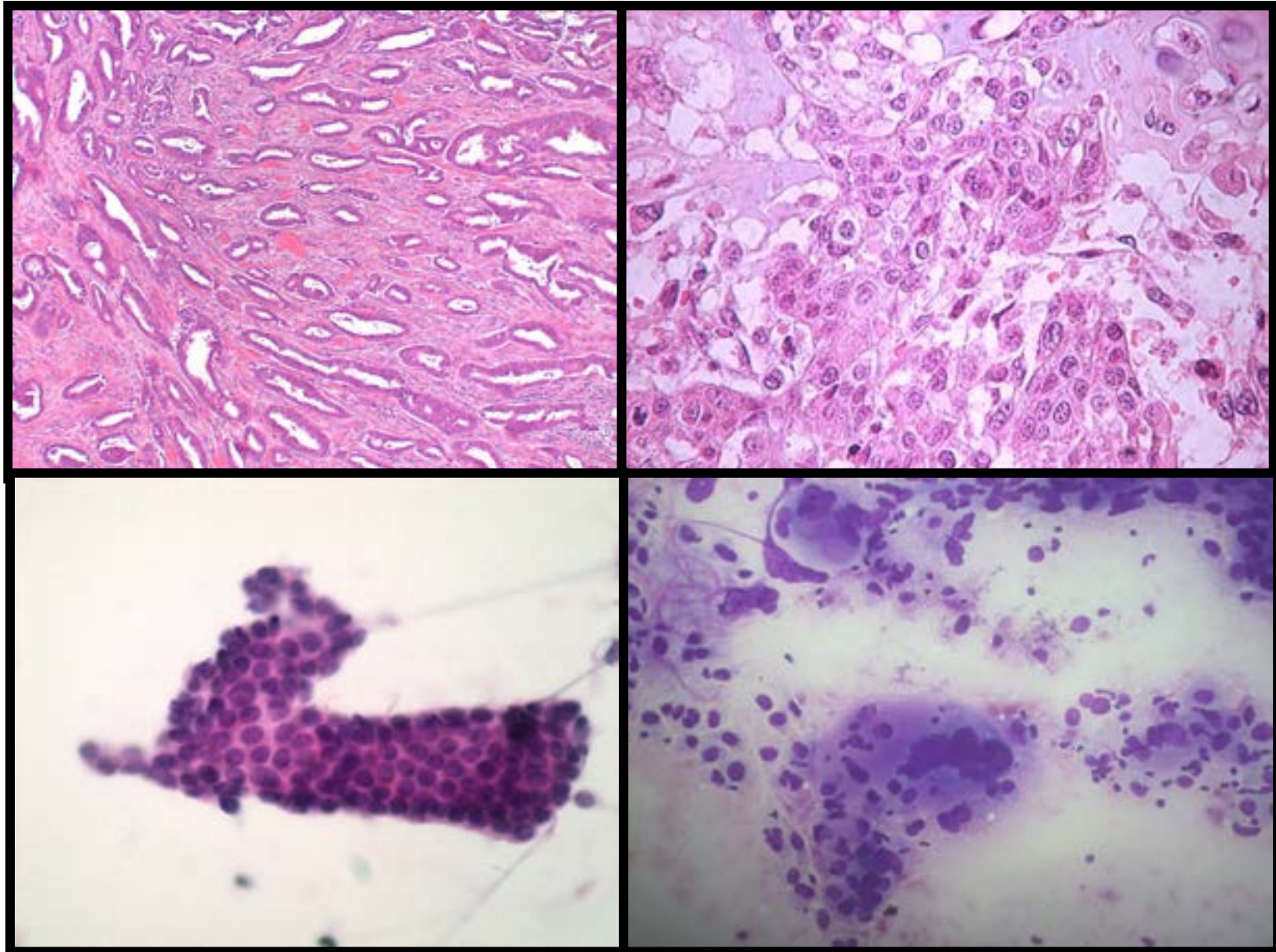
WHO Classification of Tumours of the Breast

Edited by David N. Lakhani, Ian O. Ellis, Stewart J. Schnitt, Puay Hoon Tan, Marc J. van de Vijver

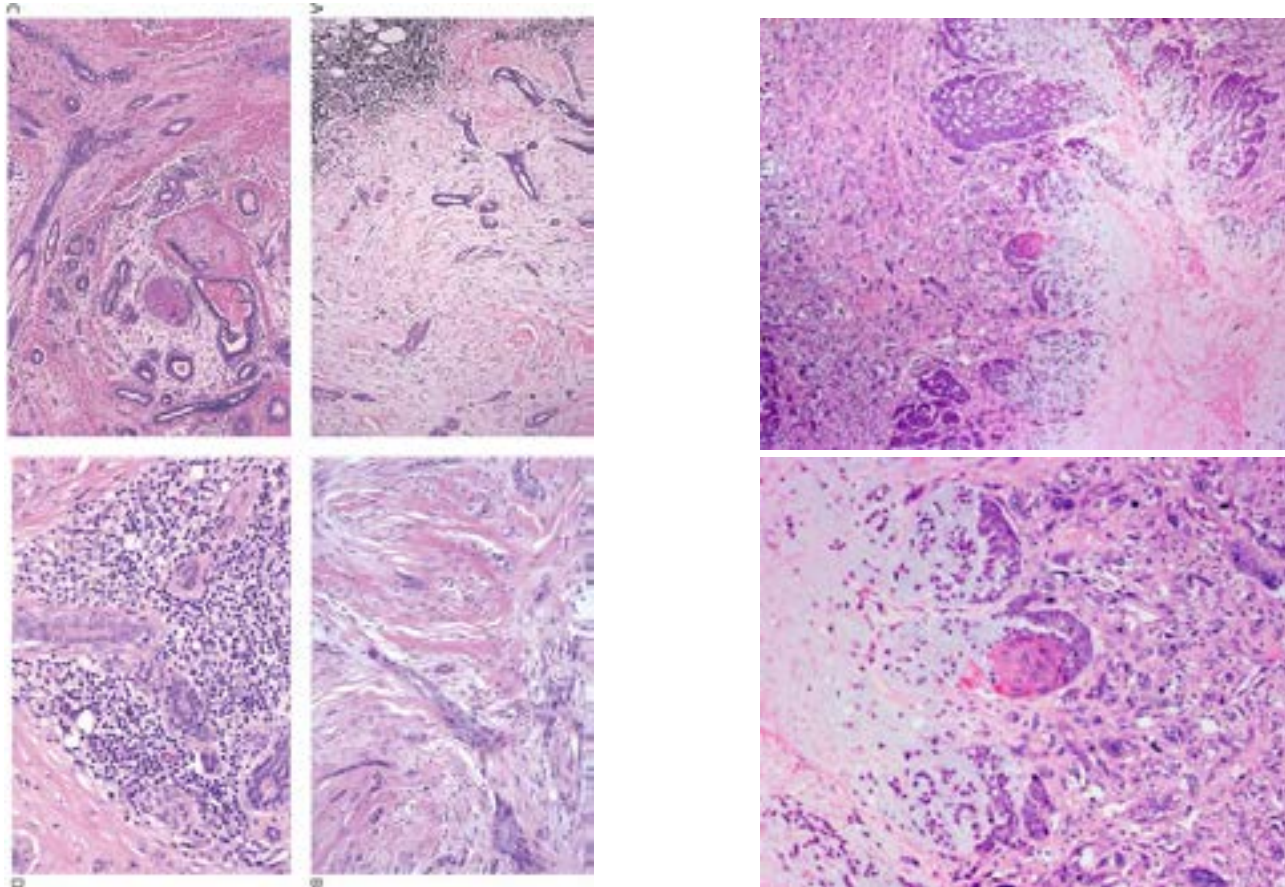


Type	Classification
Precursor lesions	
Ductal carcinoma in situ	8500/2
Lobular neoplasia	
Lobular carcinoma in situ	
Classic lobular carcinoma in situ	8520/2
Pleomorphic lobular carcinoma in situ	8519/2*
Atypical lobular hyperplasia	
Intraductal proliferative lesions	
Usual ductal hyperplasia	
Columnar cell lesions including flat epithelial atypia	
Atypical ductal hyperplasia	
Papillary lesions	
Intraductal papilloma	8503/0
Intraductal papilloma with atypical hyperplasia	8503/0
Intraductal papilloma with ductal carcinoma in situ	8503/2*
Intraductal papilloma with lobular carcinoma in situ	8520/2
Intraductal papillary carcinoma	8503/2
Encapsulated papillary carcinoma	8504/2
Encapsulated papillary carcinoma with invasion	8504/3
Solid papillary carcinoma	
In situ	8509/2
Invasive	8509/3

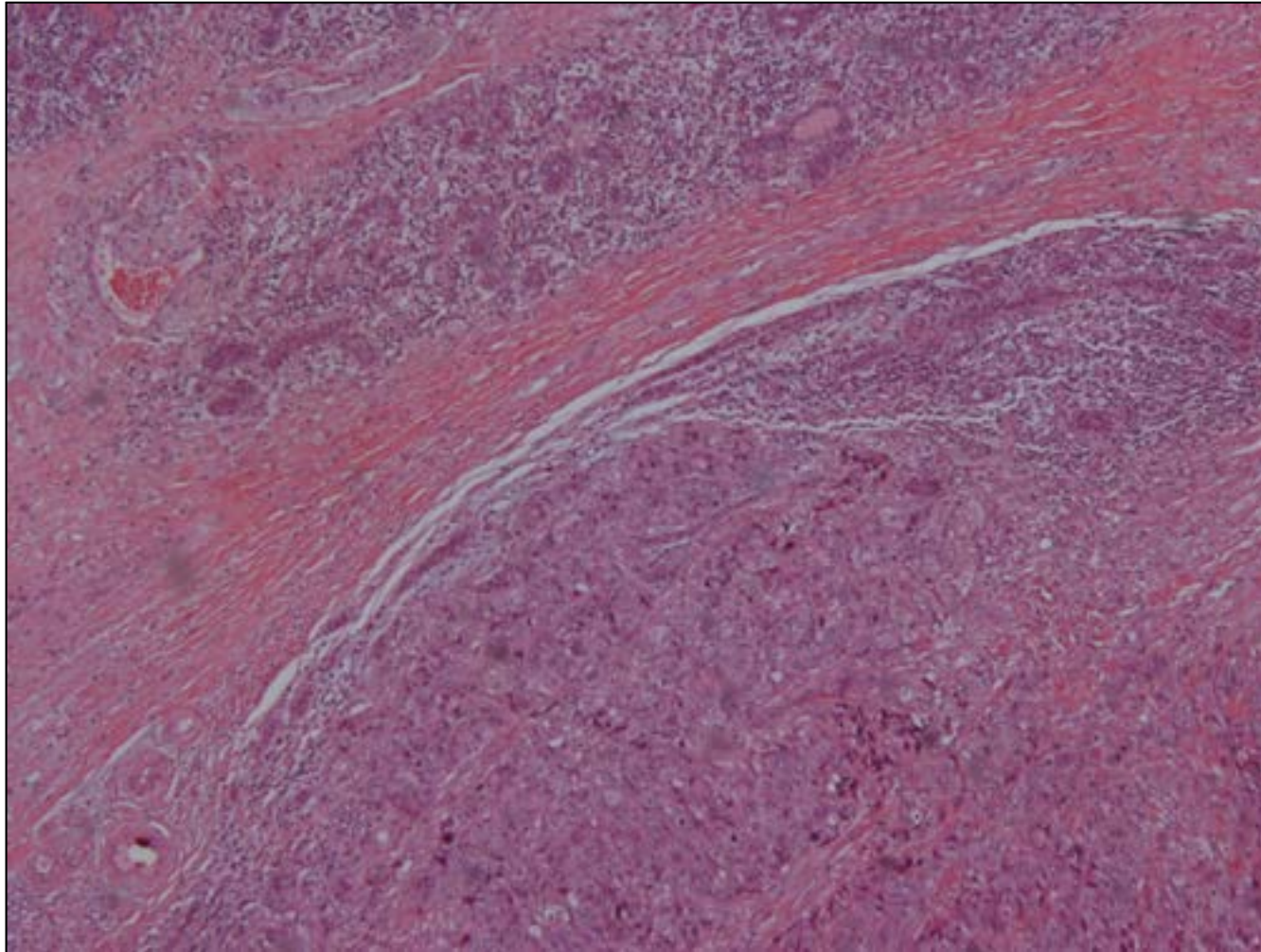
Breast cancer classification and prognosis

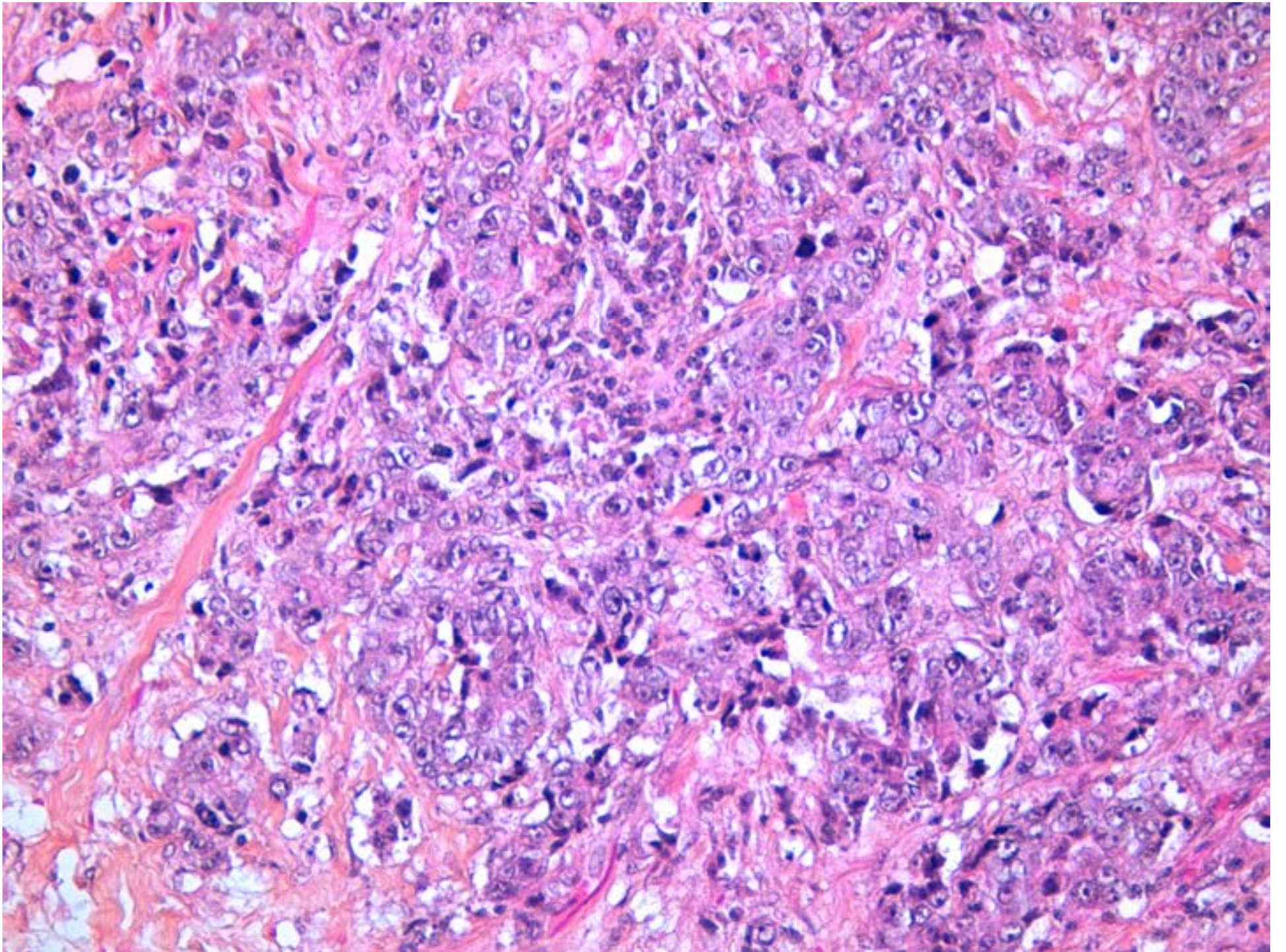


Breast cancer classification and prognosis

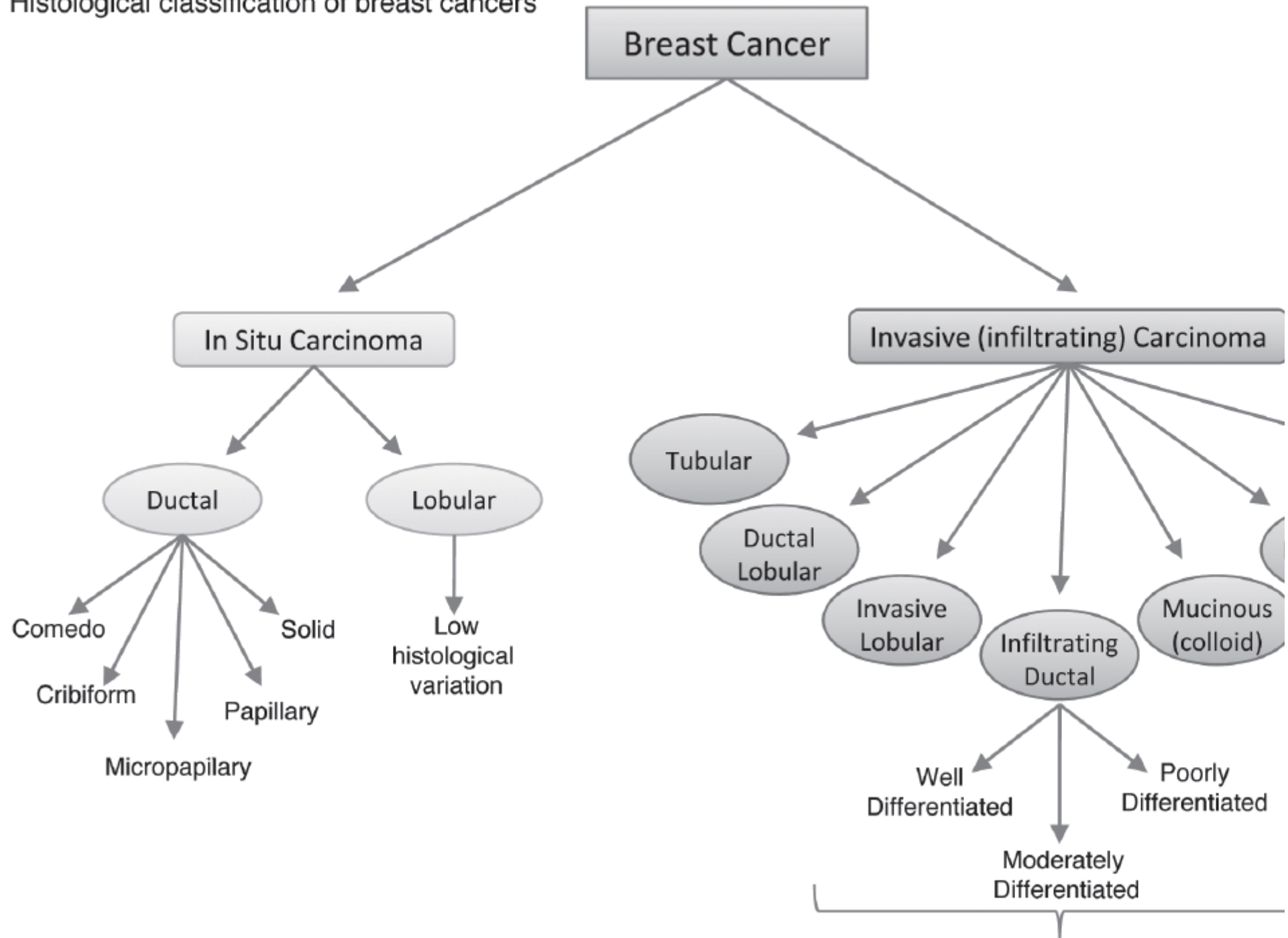


Breast cancer classification and prognosis

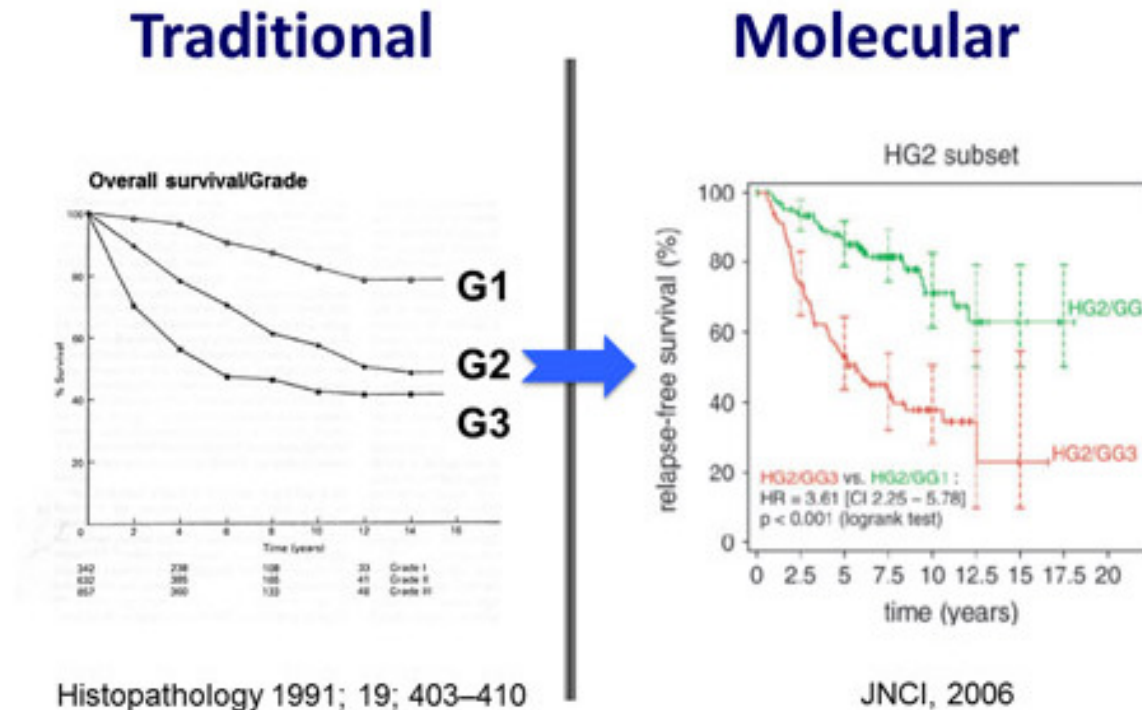




Histological classification of breast cancers

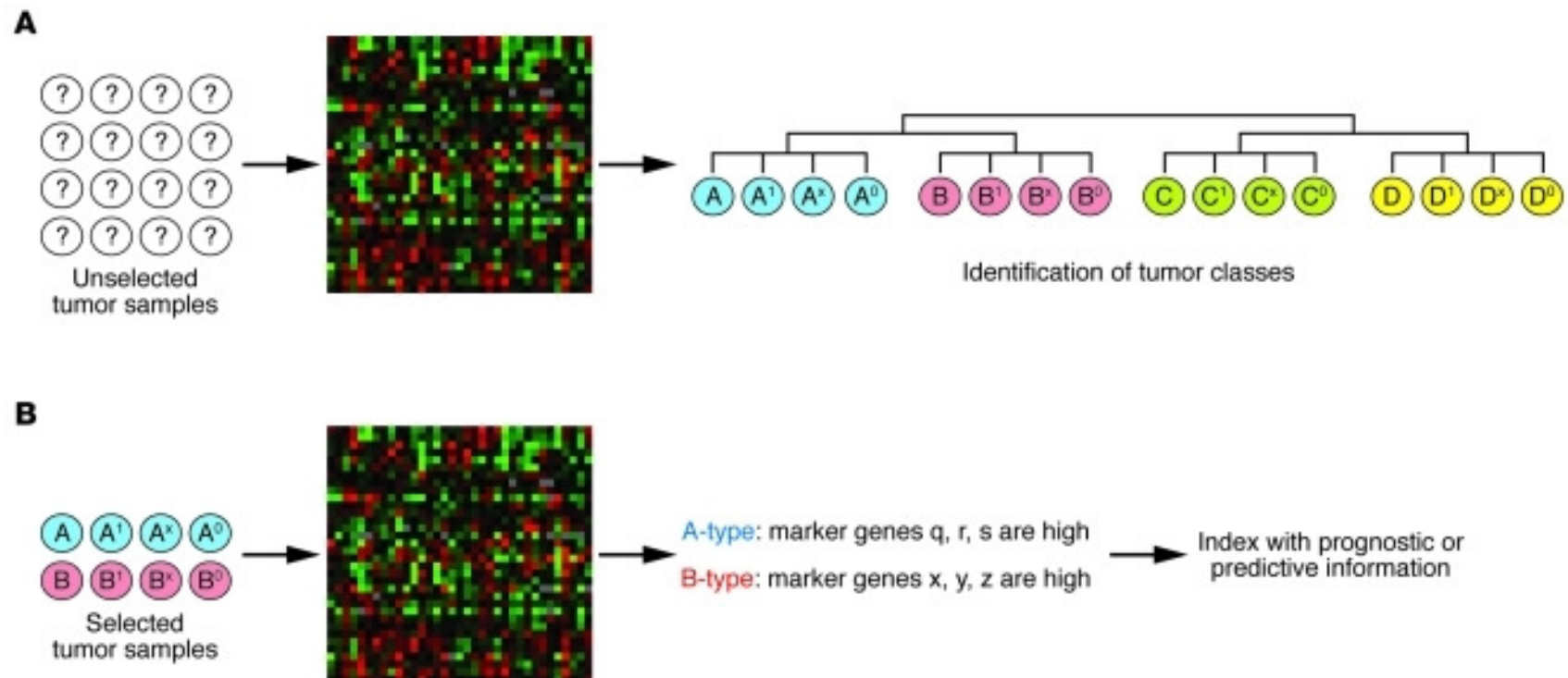


Breast cancer classification and prognosis



Oncotype DX and PAM 50 approximately split this group in half when classified as low risk RS (56%) and Luminal A (63%) approximately.

Gene-expression profiling (microarray-based)



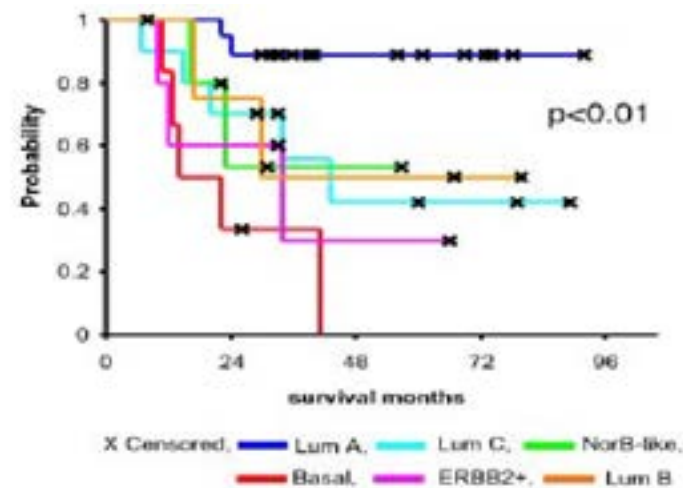
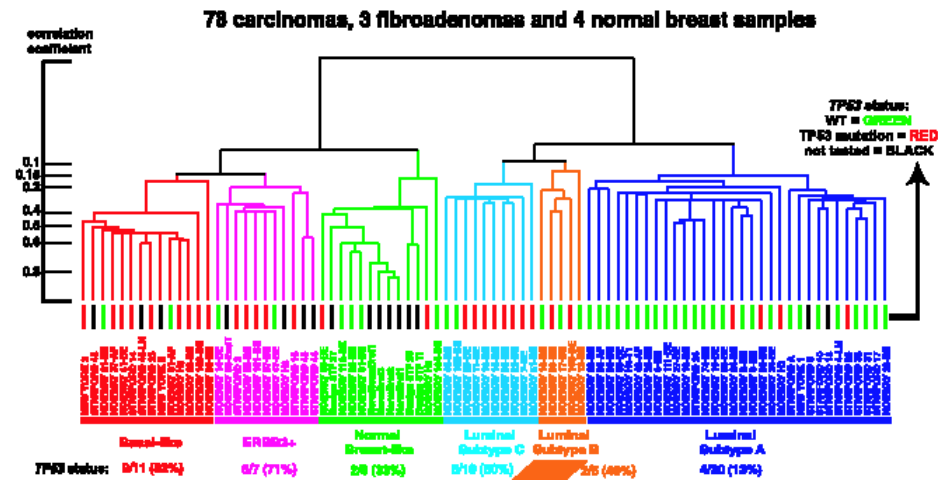
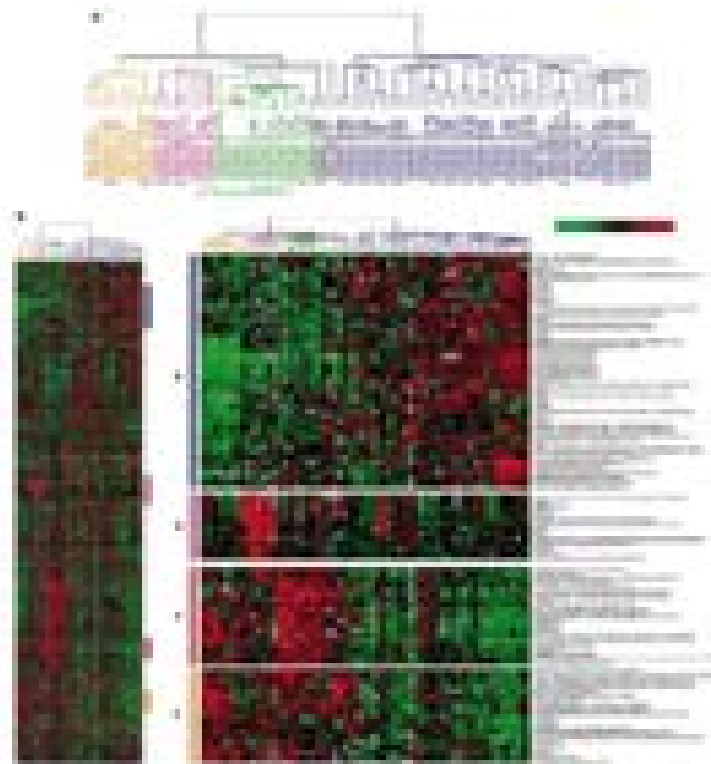
letters to nature

Gene expression patterns of breast carcinomas distinguish tumor subclasses with clinical implications

Therese Sorlie^{1,2,3}, Charles M. Perou^{1,2}, Robert Tibshirani⁴, Turid Aas¹, Stephanie Geisler⁵, Hilde Johnsen⁶, Trevor Hastie⁴, Michael B. Eisen⁷, Matt van de Rijn⁸, Stefanie S. Jeffrey⁹, Thor Thorsen⁶, Hanne Quirf⁶, John C. Matese¹, Patrick O. Brown¹⁰, David Botstein¹, Per Eystein Lønning¹, and Anne-Lise Borresen Dale^{1,2}

Molecular portraits of human breast tumours

Charles M. Perou¹, Therese Sorlie^{1,2}, Michael B. Eisen⁷, Matt van de Rijn⁸, Stefanie S. Jeffrey⁹, Thor Thorsen⁶, Hanne Quirf⁶, John C. Matese¹, Patrick O. Brown¹⁰, David Botstein¹, Per Eystein Lønning¹, Anne-Lise Borresen Dale^{1,2}, Robert Tibshirani⁴, Turid Aas¹, Stephanie Geisler⁵, Hilde Johnsen⁶, Trevor Hastie⁴, Michael B. Eisen⁷, Matt van de Rijn⁸, Stefanie S. Jeffrey⁹, Thor Thorsen⁶, Hanne Quirf⁶, John C. Matese¹, Patrick O. Brown¹⁰, David Botstein¹, Per Eystein Lønning¹, and Anne-Lise Borresen Dale^{1,2}



Molecular Classification of Breast Cancer

Pu et al. *Breast Cancer Research* 2016, 18:168
<http://breast-cancer-research.com/content/18/1/168>

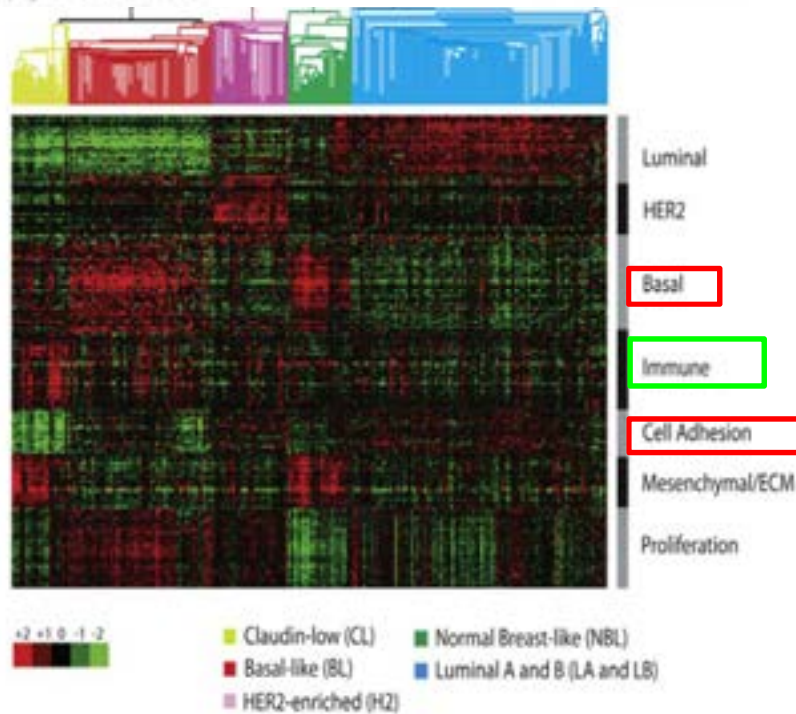


RESEARCH ARTICLE

Open Access

Phenotypic and molecular characterization of the claudin-low intrinsic subtype of breast cancer

Alex Prat^{1,2,3}, Joel S Parker^{1,2}, Olga Karginova^{1,2,3*}, Cheng Fan¹, Chad Livasy^{1,3}, Jason I Herschkowitz⁴, Xiaoping He^{1,2,3}, Charles M Perou^{1,2,3*}



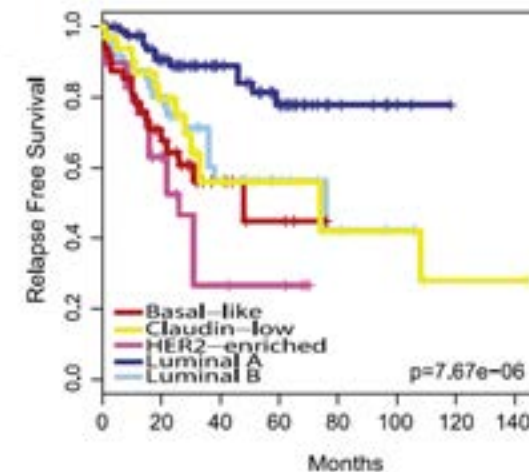
LUMINAL A: ER+/PgR+/HER2-

LUMINAL B: ER+/PgR+/HER2+ and or Ki67+

HER-OE: ER-/PgR-/HER2+

BASAL-LIKE: ER-/PgR-/HER2-/Basal Markers

CLAUDIN-LOW: ER-/Pg-/HER2-/Claudin^{low}



Molecular Classification of Breast Cancer

ER +

80%

Luminal A
Luminal B

ER -

20%

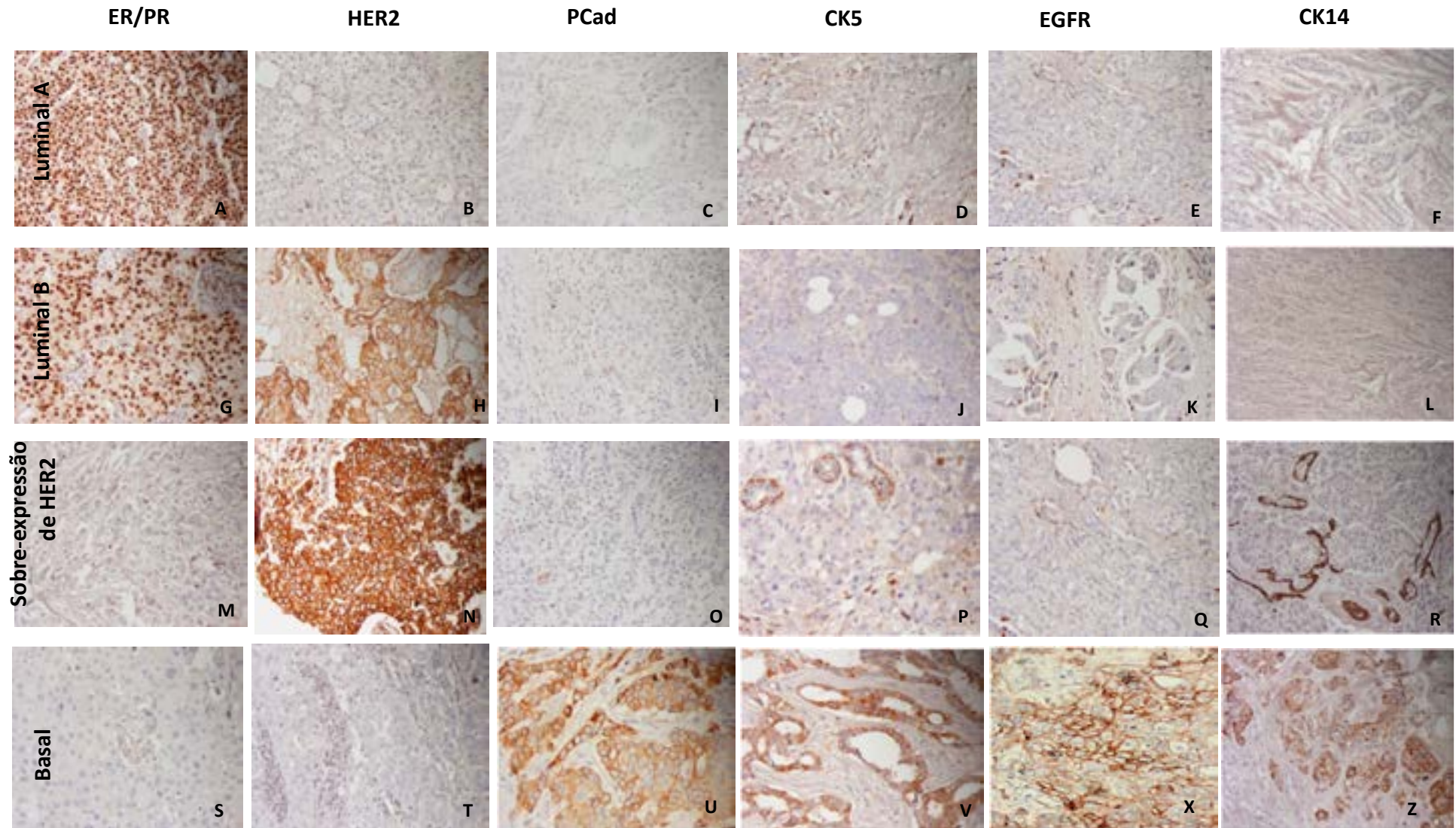
HER2
Basal
Claudin-low

.....

Irina Mates · Rozany Daltro · Marcelo Alvarenga ·
Luiz Carlos Zeferino · Fernando Schmitt

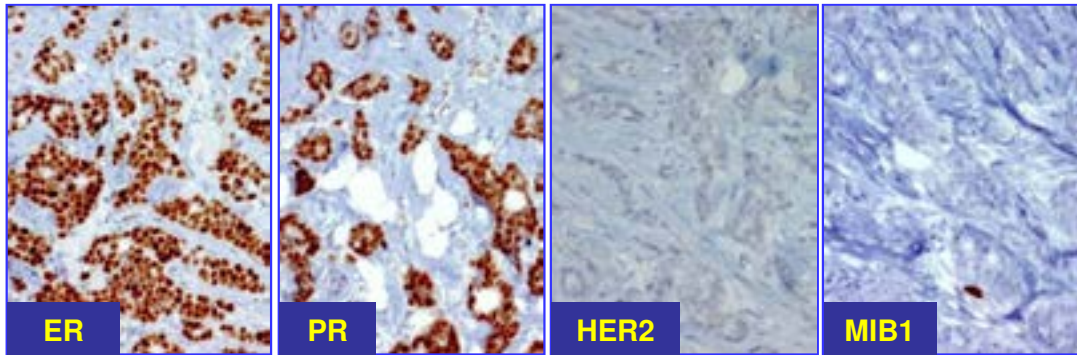
**p63, cytokeratin 5, and P-cadherin: three molecular markers
to distinguish basal phenotype in breast carcinomas**

IHC TRANSLATION OF MOLECULAR CLASSIFICATION

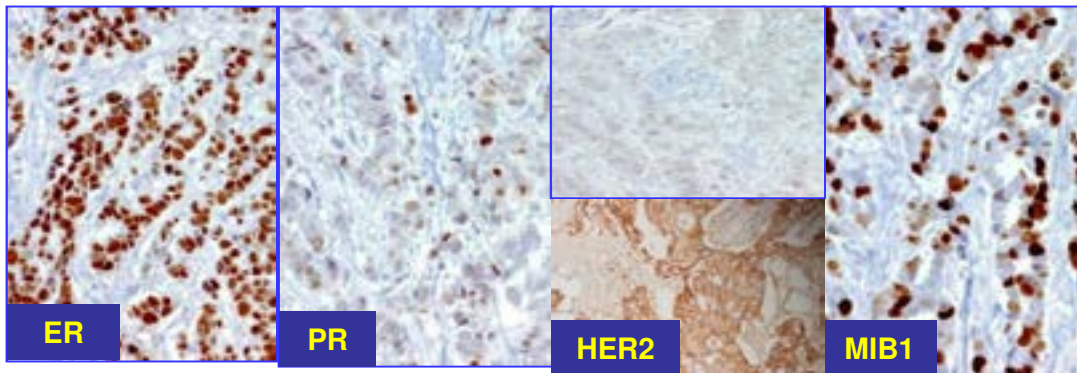


ER Positive Breast Cancer

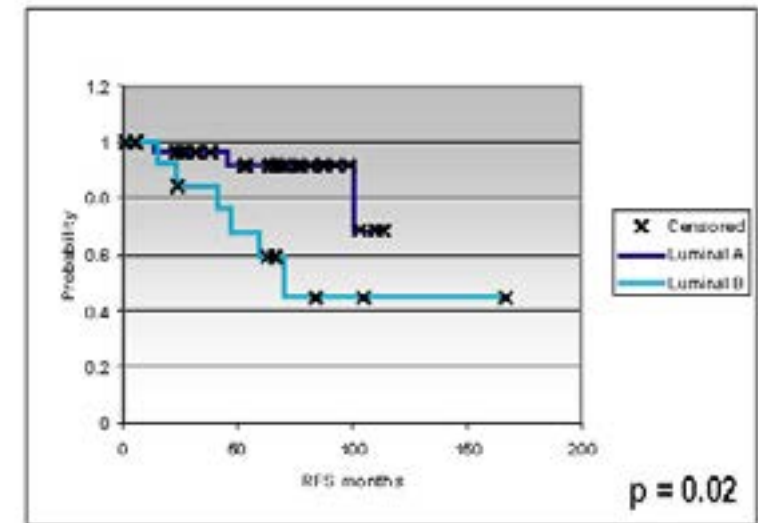
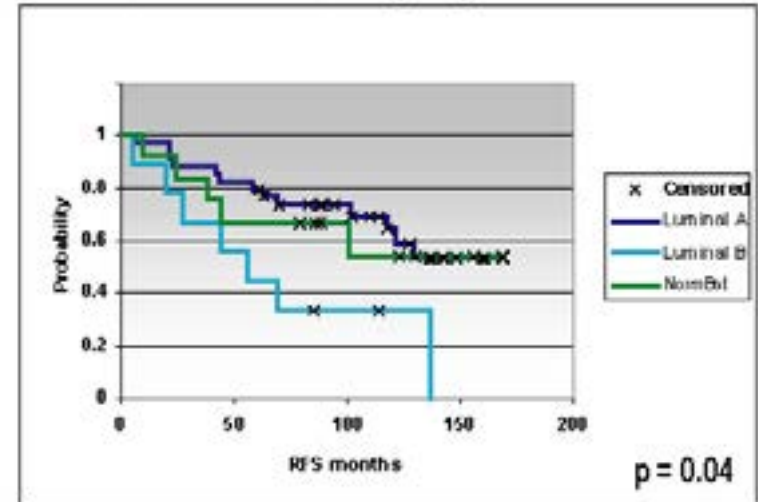
Luminal A



Luminal B

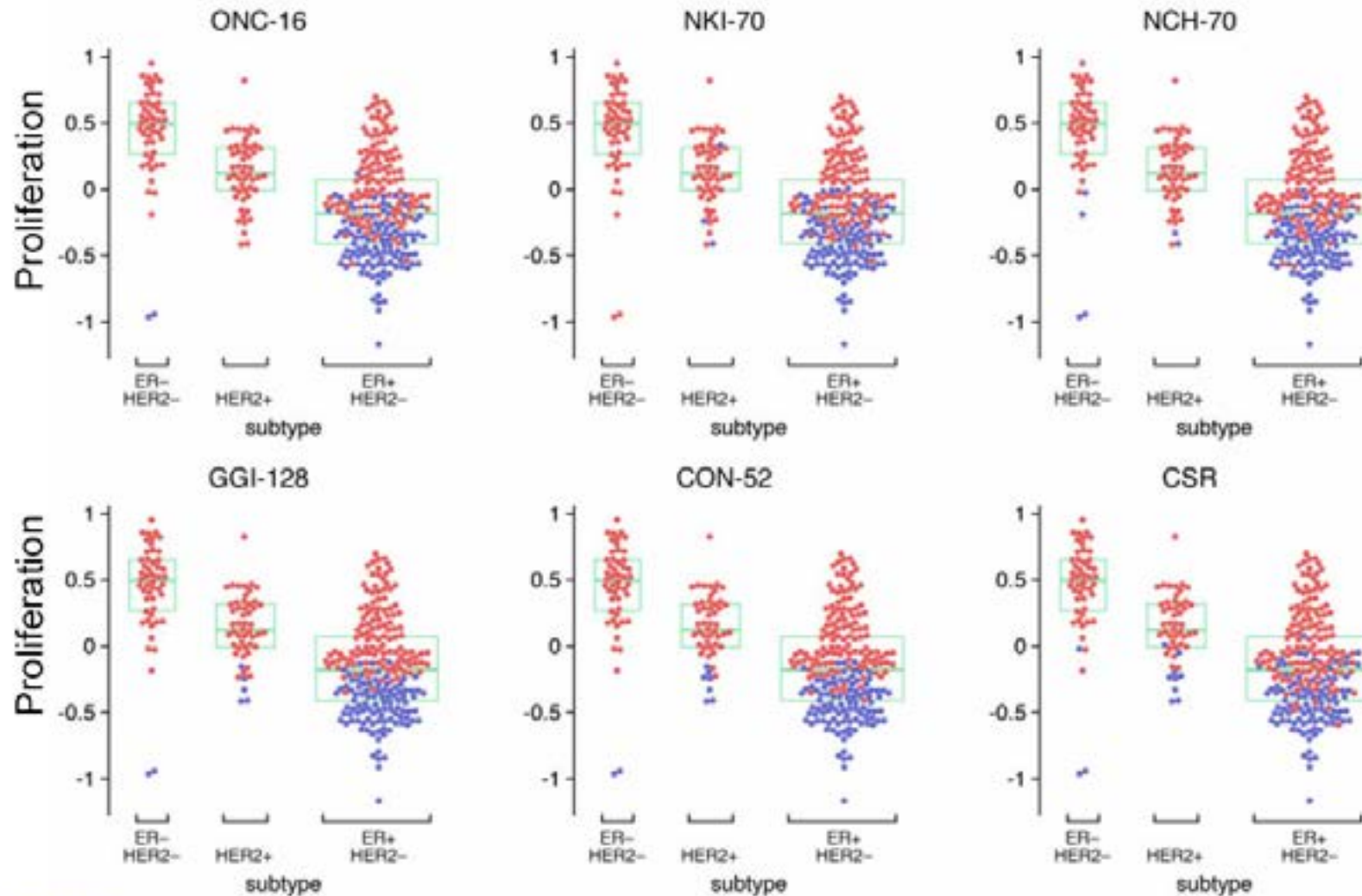


60 Sample ER+ Tamoxifen-Treated Test Set
Ma et al., Cancer Cell 5, 1-10 (2004).



45 Tamoxifen Treated Test Set #2
Chang et al., PNAS 102, 3738-43 (2005) + UNC

Meta-Analysis – Gene signatures



Blue dots: good prognosis

Red dots: poor prognosis

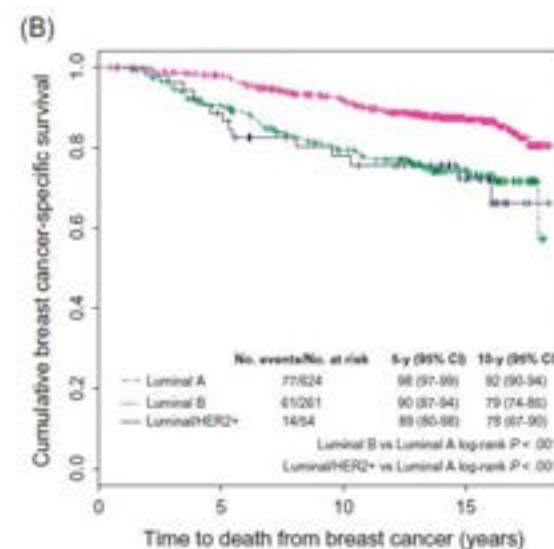
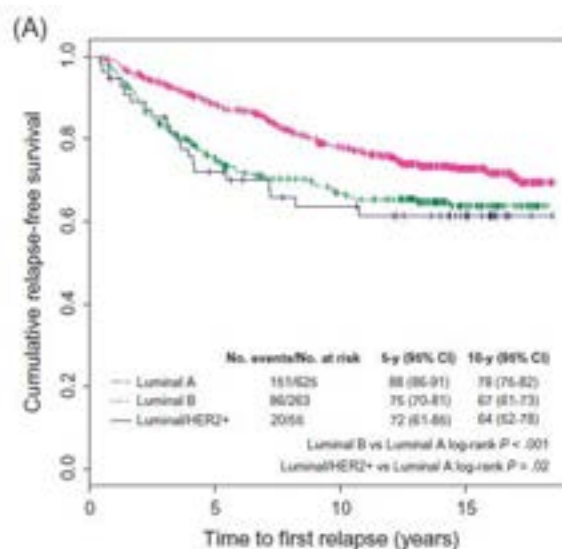
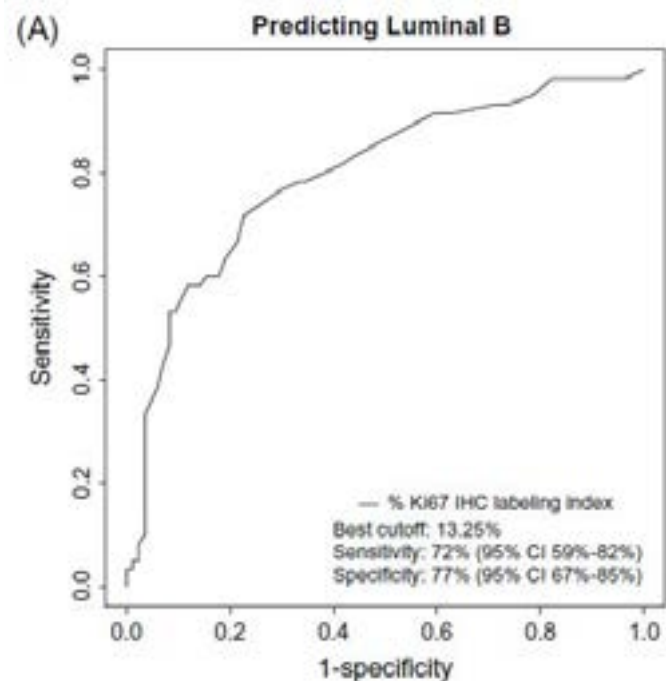
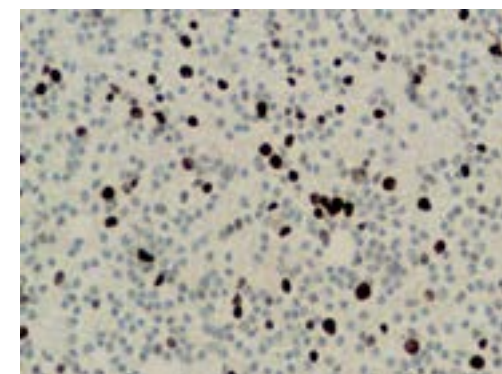
Wirapati et al. Breast Cancer Res 2008;10:R65

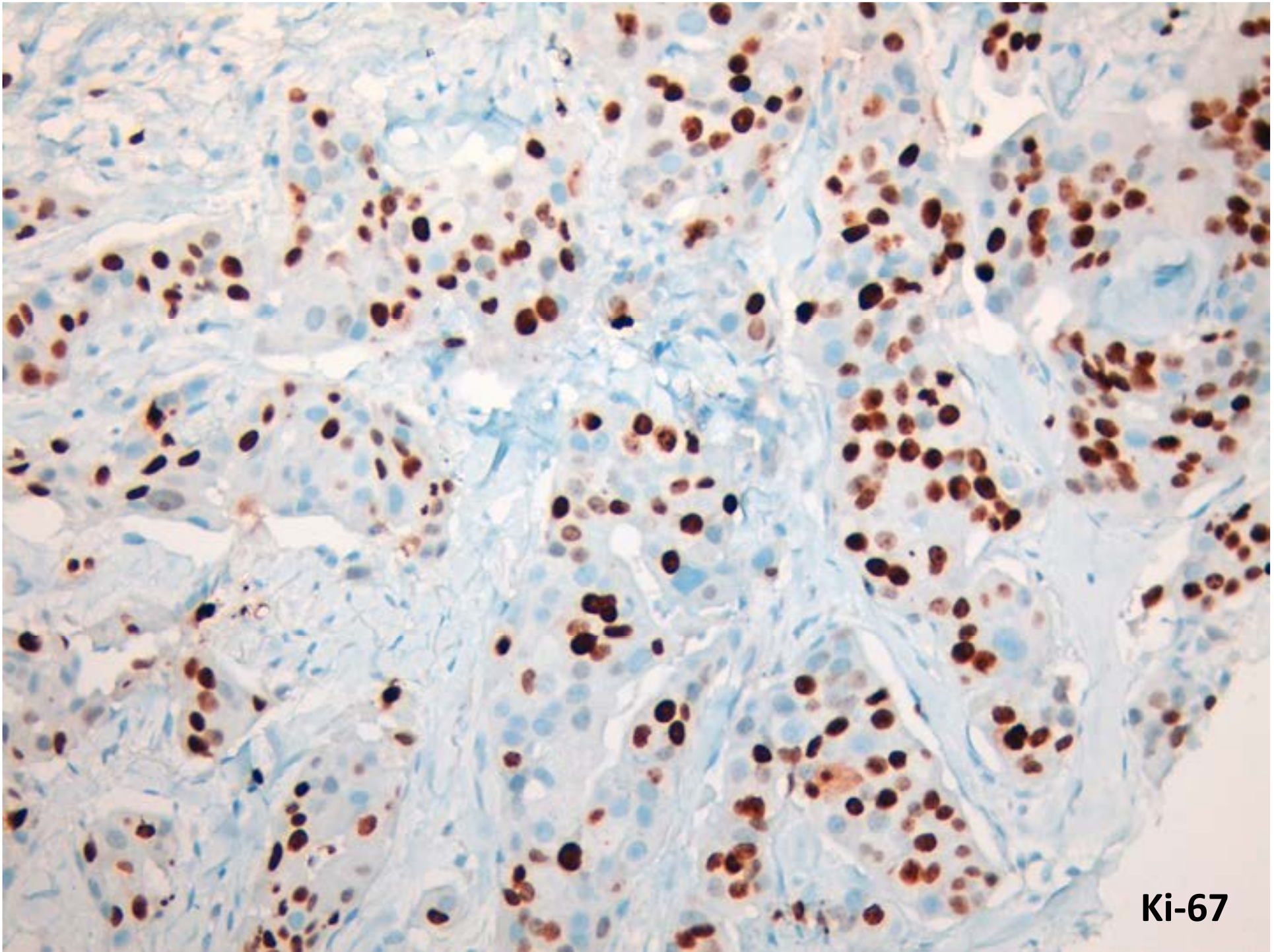
Intrinsic subtype	Clinico-pathologic surrogate definition	Notes
Luminal A	<p>'Luminal A-like'</p> <p><i>all of:</i></p> <ul style="list-style-type: none"> ER and PgR positive HER2 negative Ki-67 'low' Recurrence risk 'low' based on multi-gene-expression assay (if available)^b 	<p>The cut-point between 'high' and 'low' values for Ki-67 varies between laboratories.^a A level of <14% best correlated with the gene-expression definition of Luminal A based on the results in a single reference laboratory [23]. Similarly, the added value of PgR in distinguishing between 'Luminal A-like' and 'Luminal B-like' subtypes derives from the work of Prat et al. which used a PgR cut-point of $\geq 20\%$ to best correspond to Luminal A subtype [24]. Quality assurance programmes are essential for laboratories reporting these results.</p>
Luminal B	<p>'Luminal B-like (HER2 negative)'</p> <ul style="list-style-type: none"> ER positive HER2 negative and <i>at least one of:</i> Ki-67 'high' PgR 'negative or low' Recurrence risk 'high' based on multi-gene-expression assay (if available)^b <p>'Luminal B-like (HER2 positive)'</p> <ul style="list-style-type: none"> ER positive HER2 over-expressed or amplified Any Ki-67 Any PgR 	<p>'Luminal B-like' disease comprises those luminal cases which lack the characteristics noted above for 'Luminal A-like' disease. Thus, either a high Ki-67^a value or a low PgR value (see above) may be used to distinguish between 'Luminal A-like' and 'Luminal B-like (HER2 negative)'.</p>
Erb- B2 overexpression	<p>'HER2 positive (non-luminal)'</p> <ul style="list-style-type: none"> HER2 over-expressed or amplified ER and PgR absent 	
'Basal-like'	<p>'Triple negative (ductal)'</p> <ul style="list-style-type: none"> ER and PgR absent HER2 negative 	<p>There is an 80% overlap between 'triple-negative' and intrinsic 'basal-like' subtype. Some cases with low-positive ER staining may cluster with non-luminal subtypes on gene-expression analysis. 'Triple negative' also includes some special histological types such as adenoid cystic carcinoma.</p>

Ki67 Index, HER2 Status, and Prognosis of Patients With Luminal B Breast Cancer

Maggie C. U. Cheang, Stephen K. Chia, David Voduc, Dongxia Gao, Samuel Leung, Jacqueline Snider, Mark Watson, Sherri Davies, Philip S. Bernard, Joel S. Parker, Charles M. Perou, Matthew J. Ellis, Torsten O. Nielsen

J Natl Cancer Inst 2009;101:736-750



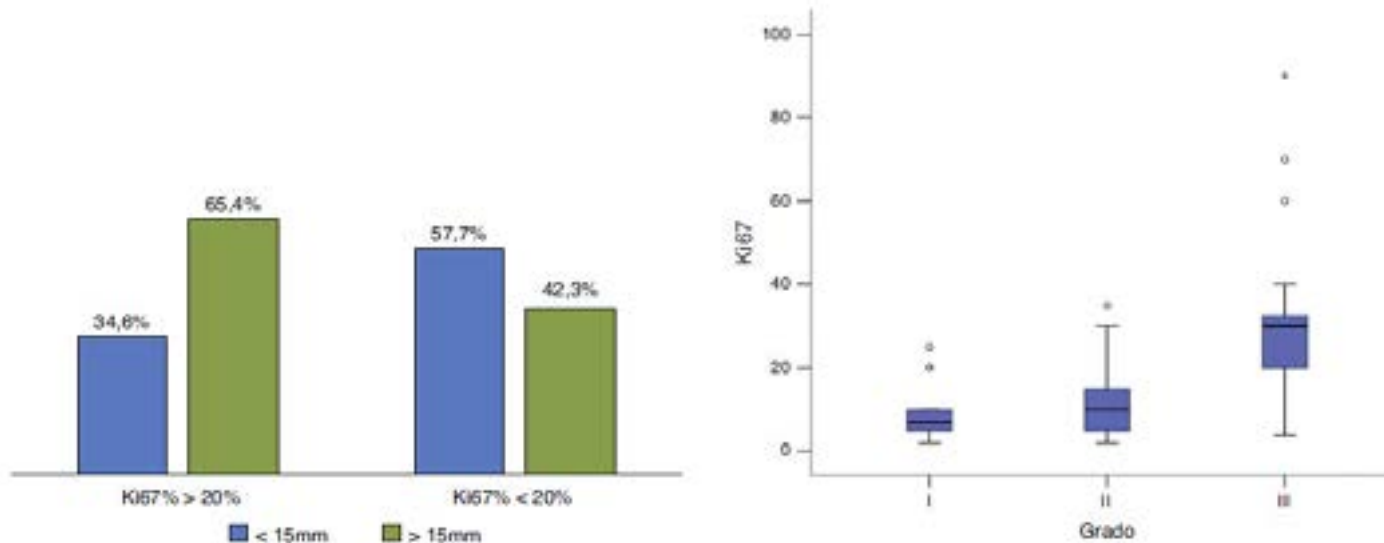


Ki-67

St Gallen Conference 2015

Distinction between Luminal A-like and Luminal B-like (HER2 neg) can be:

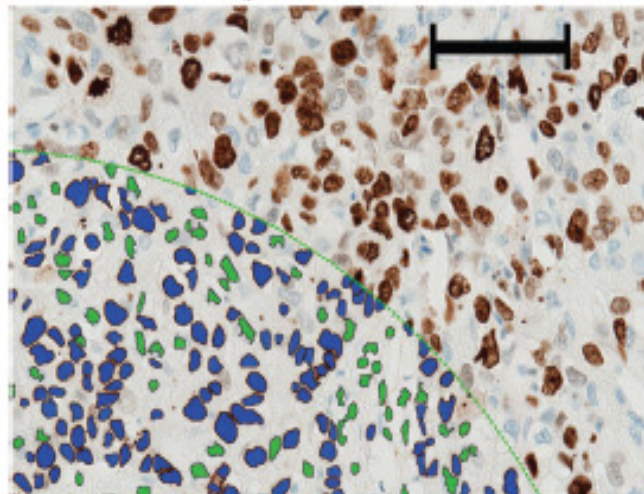
- Der
- If us



- Only appropriately determined by multi-gene classifiers: No 66.7%
- Subtype need not be determined since it can be replaced by risk scores derived from multi-gene tests: No 59.5%

Digital image analysis outperforms manual biomarker assessment in breast cancer

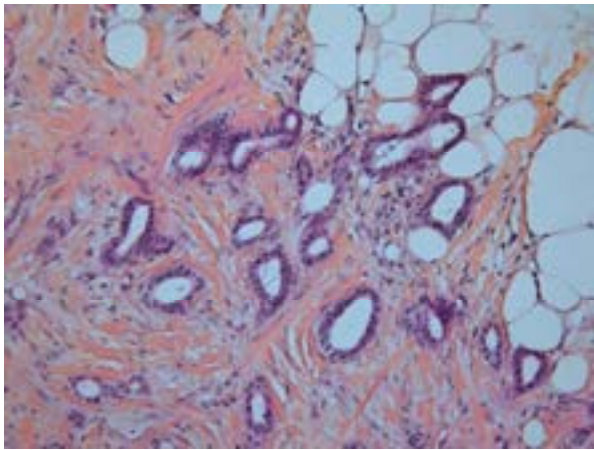
Gustav Stålhammar^{1,2}, Nelson Fuentes Martinez^{1,3}, Michael Lippert⁴, Nicholas P Tobin⁵,
 Ida Mølholm^{4,6}, Lorand Kis⁷, Gustaf Rosin¹, Mattias Rantalainen⁸, Lars Pedersen⁴,
 Jonas Bergh^{1,5,9}, Michael Grunkin⁴ and Johan Hartman^{1,5,7}



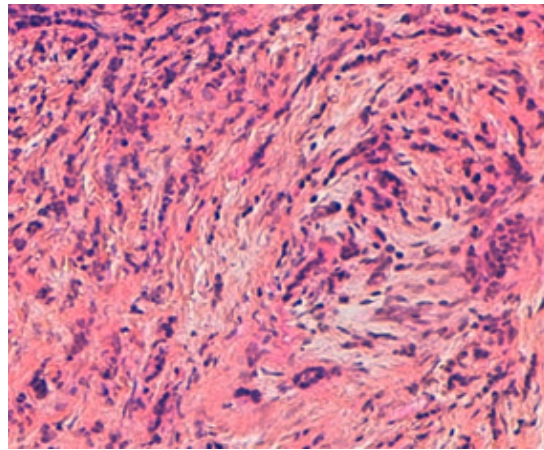
<i>Ki67 scoring method</i>	<i>Sensitivity for PAM50 Luminal B vs A</i>	<i>Specificity for PAM50 Luminal B vs A</i>
<i>DIA invasive margin</i>		
Cutoff $\geq 20\%$	84%	78%
Cutoff $\geq 20.2\%^*$	82%	79%
<i>DIA hot spot</i>		
Cutoff $\geq 20\%$	90%	65%
Cutoff $\geq 25.2\%^*$	86%	77%
<i>DIA average</i>		
Cutoff $\geq 20\%$	60%	90%
Cutoff $\geq 15.5\%^*$	80%	83%
<i>Manual</i>		
Cutoff $\geq 20\%$	75%	70%
Cutoff $\geq 22.5\%^*$	74%	75%

Do we still need a morphological classification?

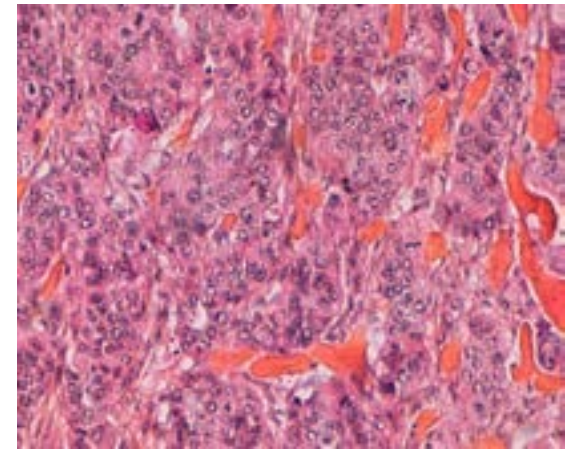
“ER-positive” breast carcinomas



Tubular carcinoma

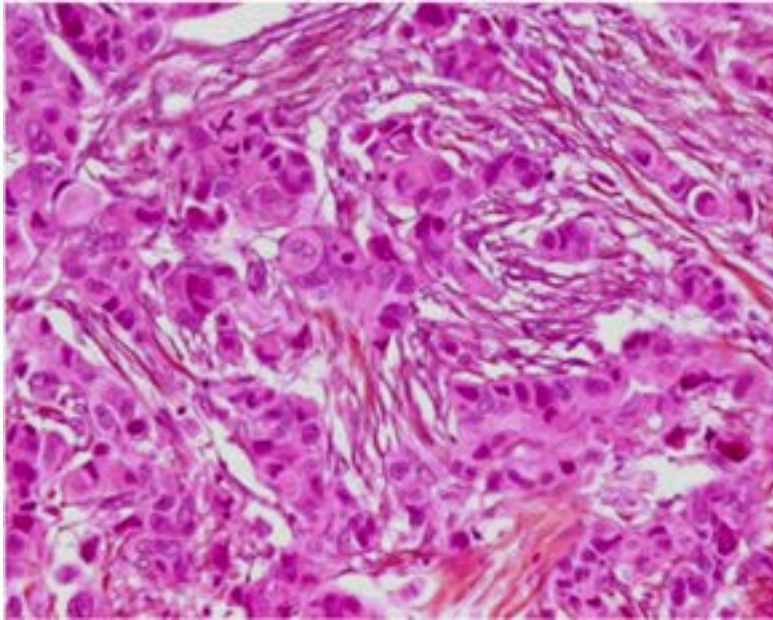


Lobular carcinoma

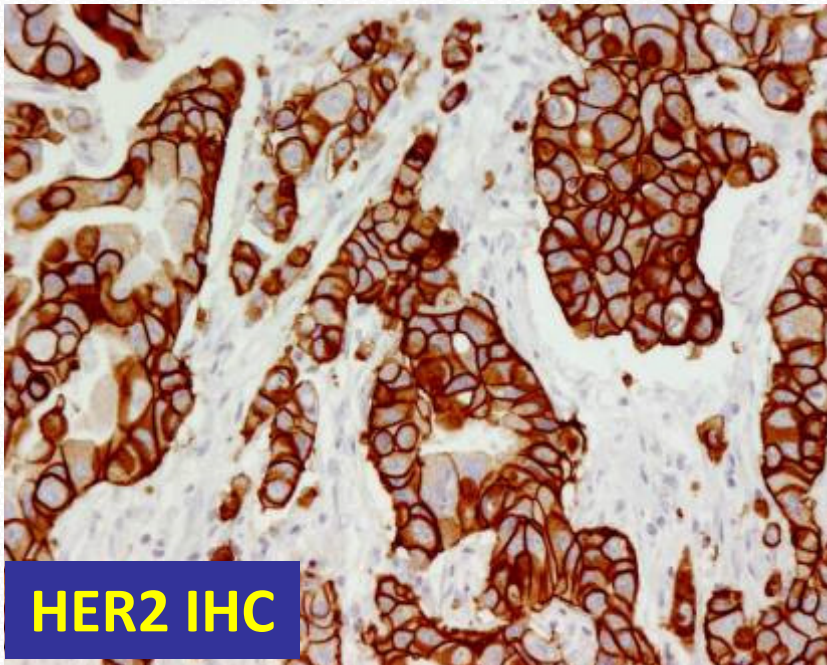
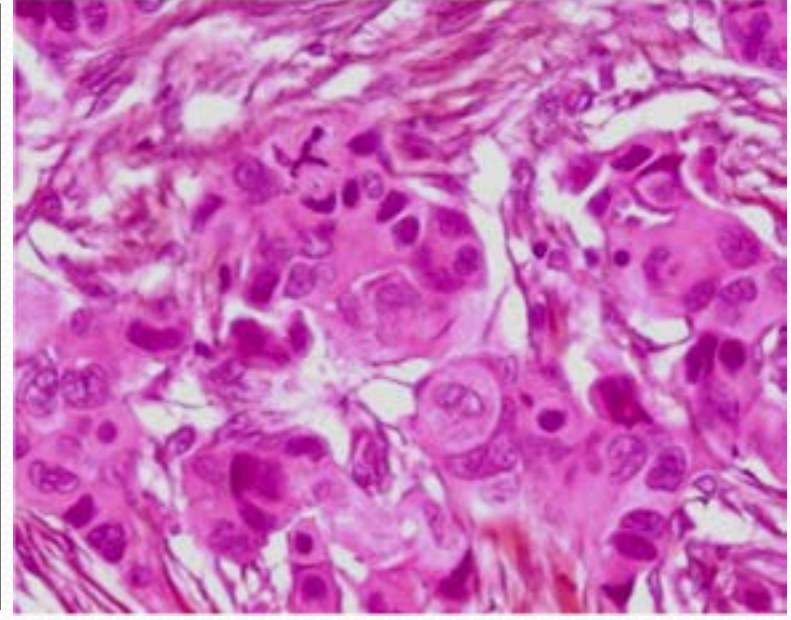


IDC Grade III

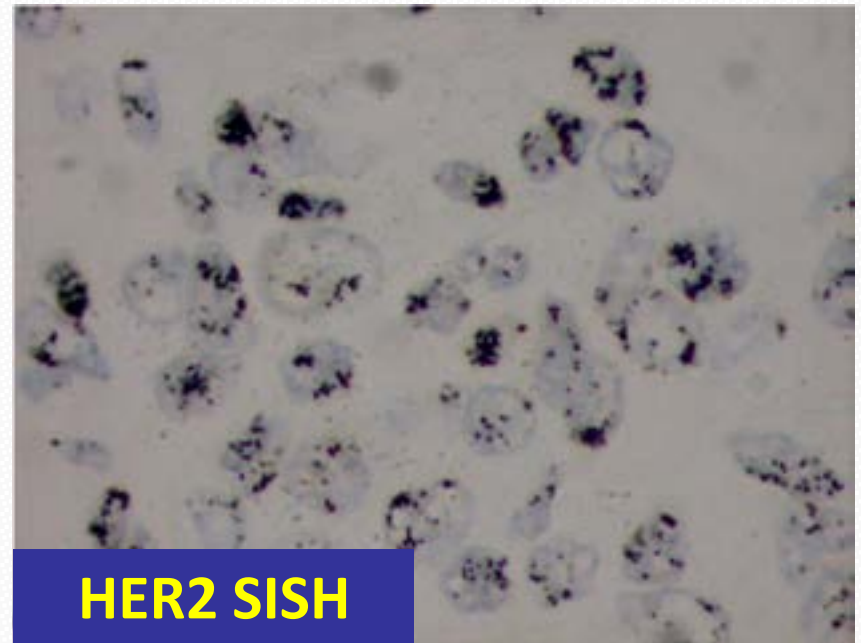
HER 2- OE BREAST CANCER



HER 2 +



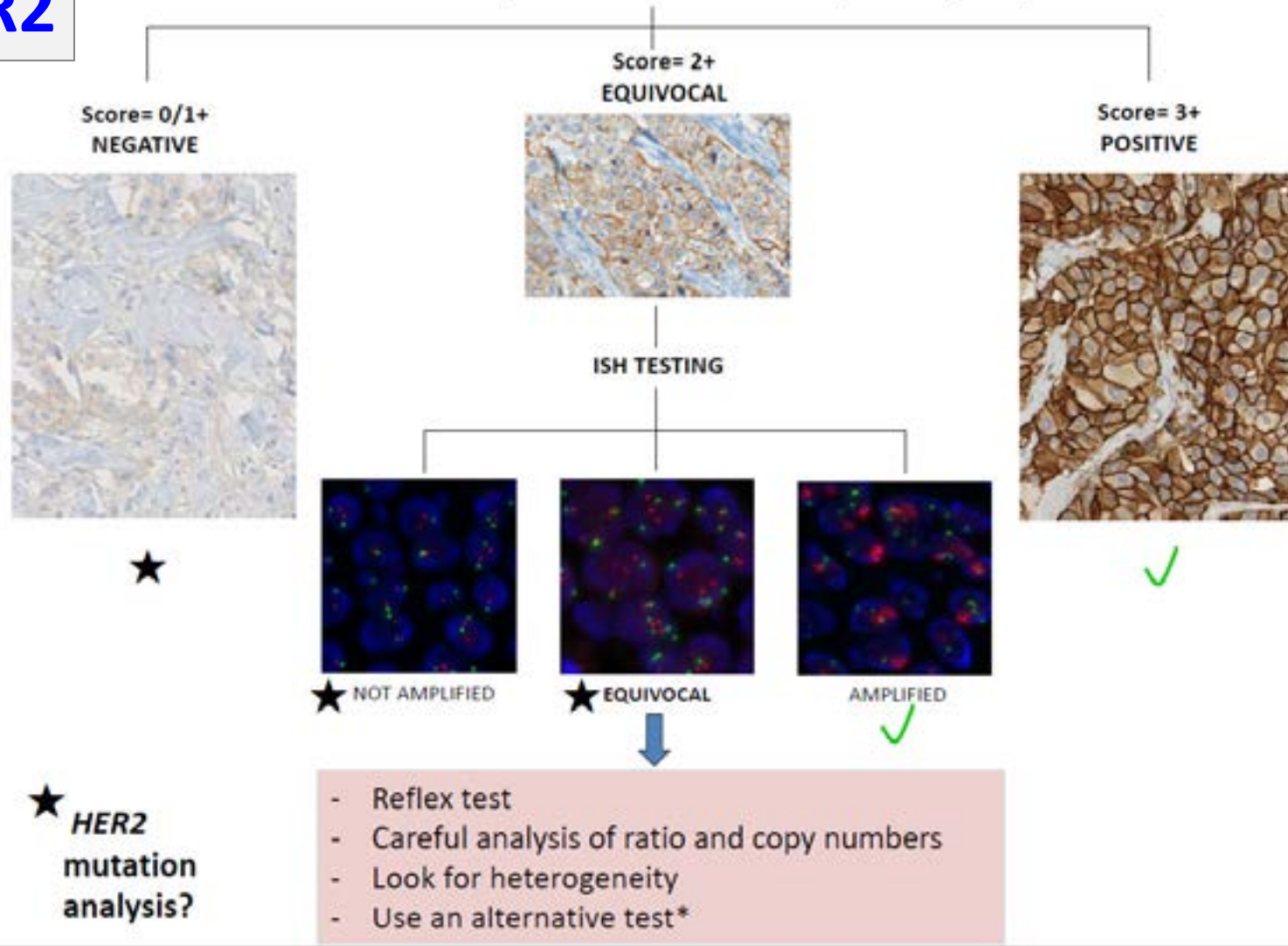
HER2 IHC



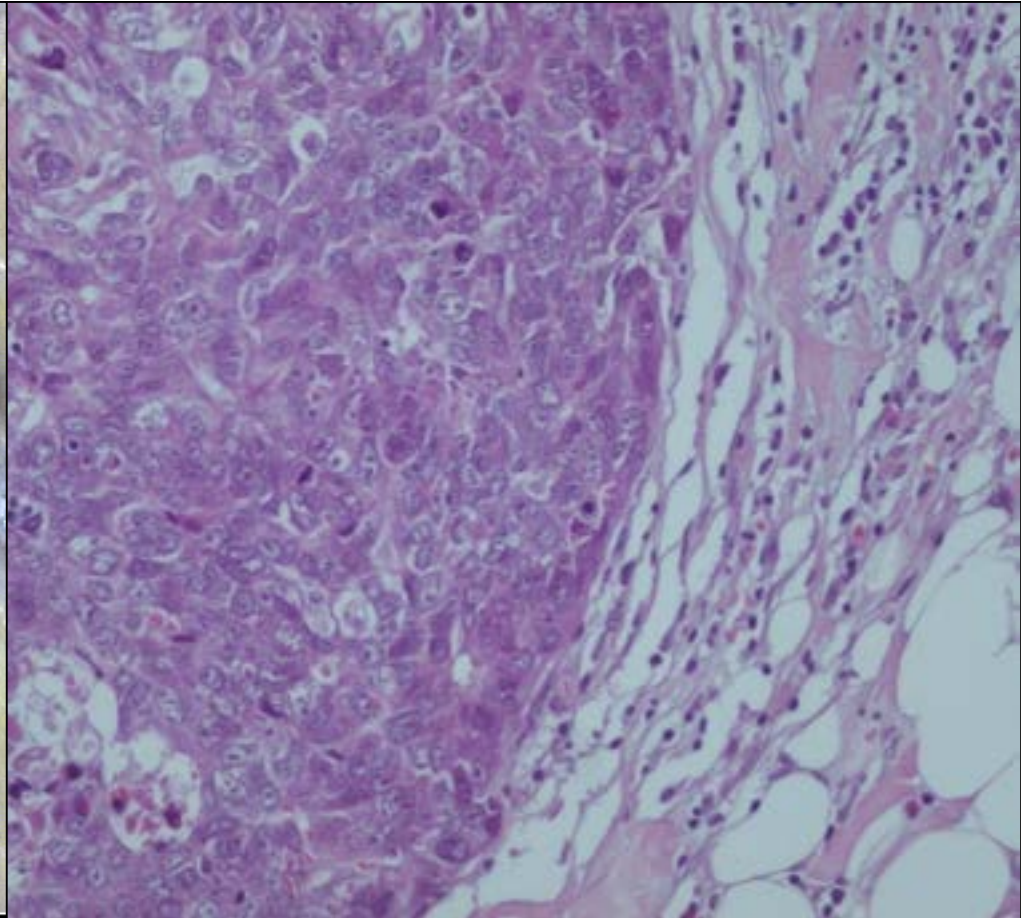
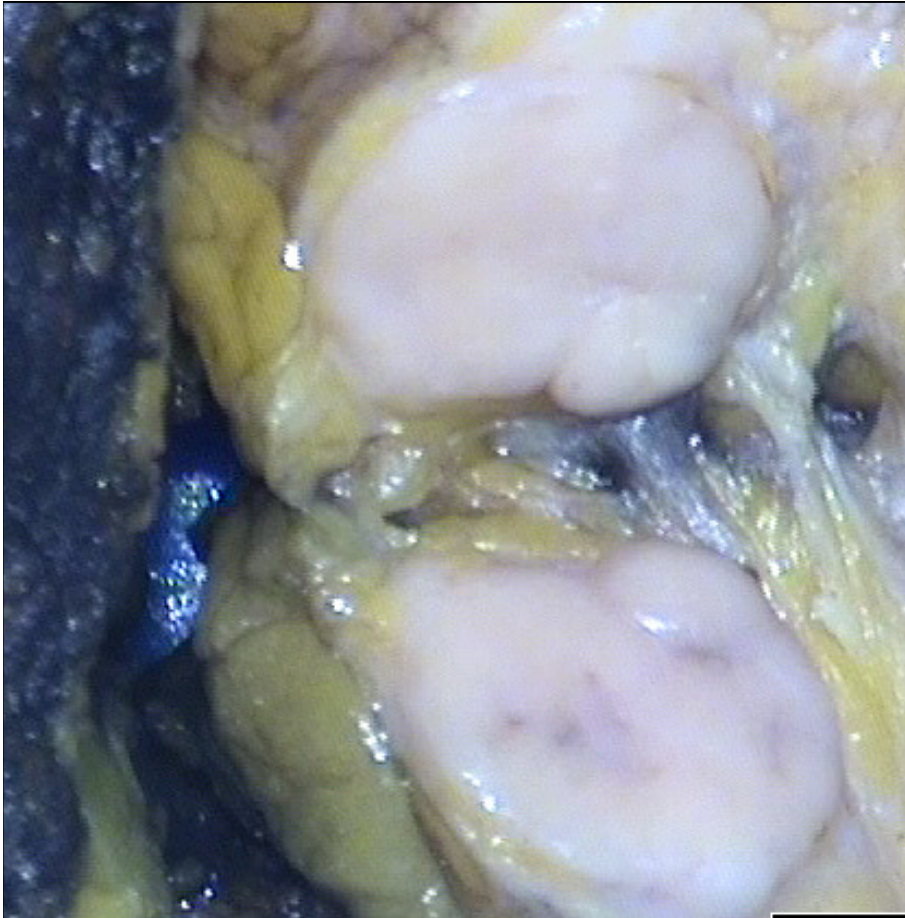
HER2 SISH

HER2

Invasive breast carcinoma processed with reliable pre-analytical procedures



1994



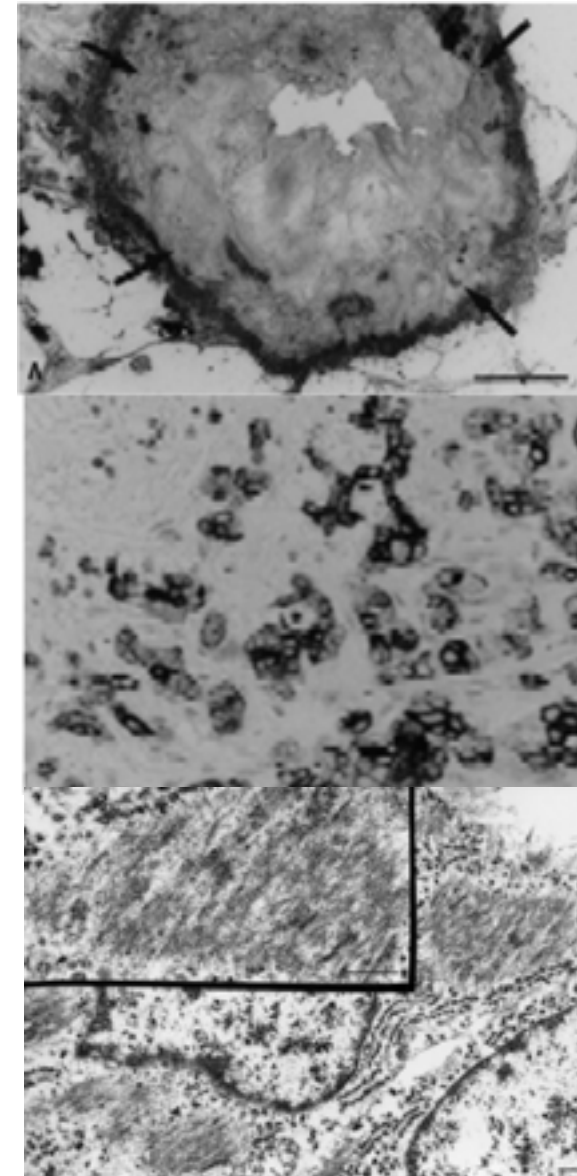
Large, Central Acellular Zones Indicating Myoepithelial Tumor Differentiation in High-Grade Invasive Ductal Carcinomas as Markers of Predisposition to Lung and Brain Metastases

Hitoshi Tsuda, M.D., Teruko Takarabe, C.T., Fumio Hasegawa, M.T., Takashi Fukutomi, M.D., and Setsuo Hirohashi, M.D.

TABLE 3. Effect on patient prognosis and preferential metastasis sites of IDCs with large central acellular zones by Cox's univariate analysis model

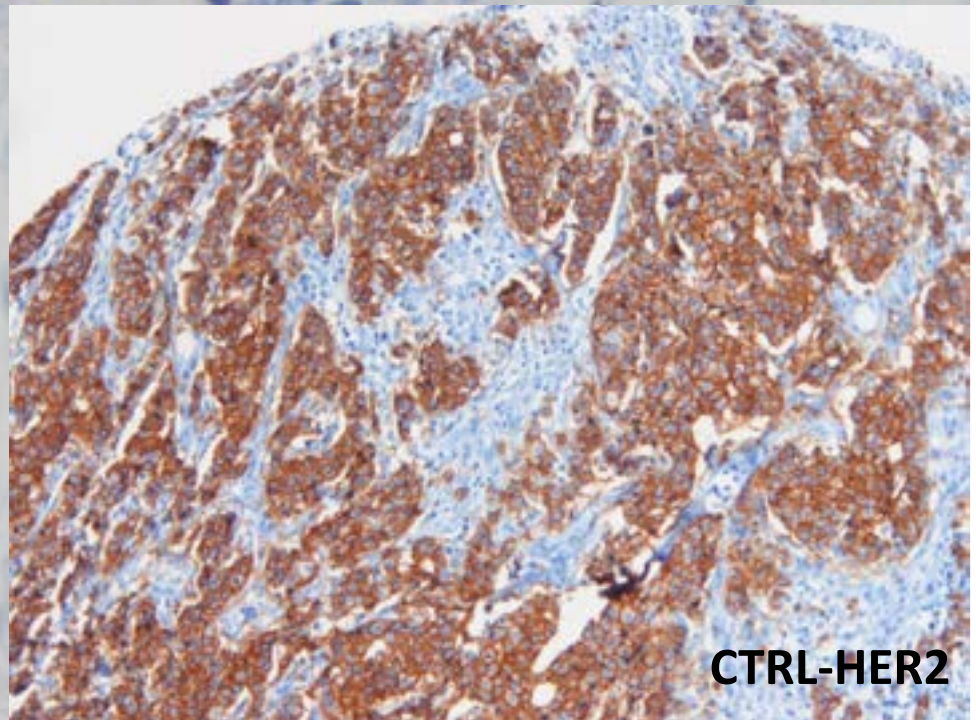
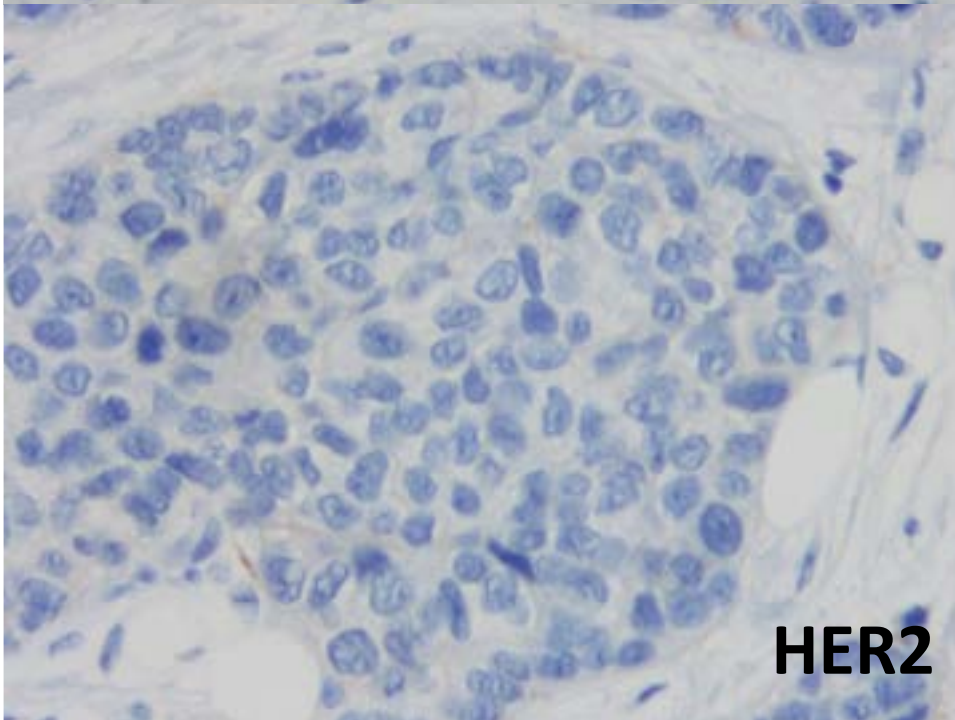
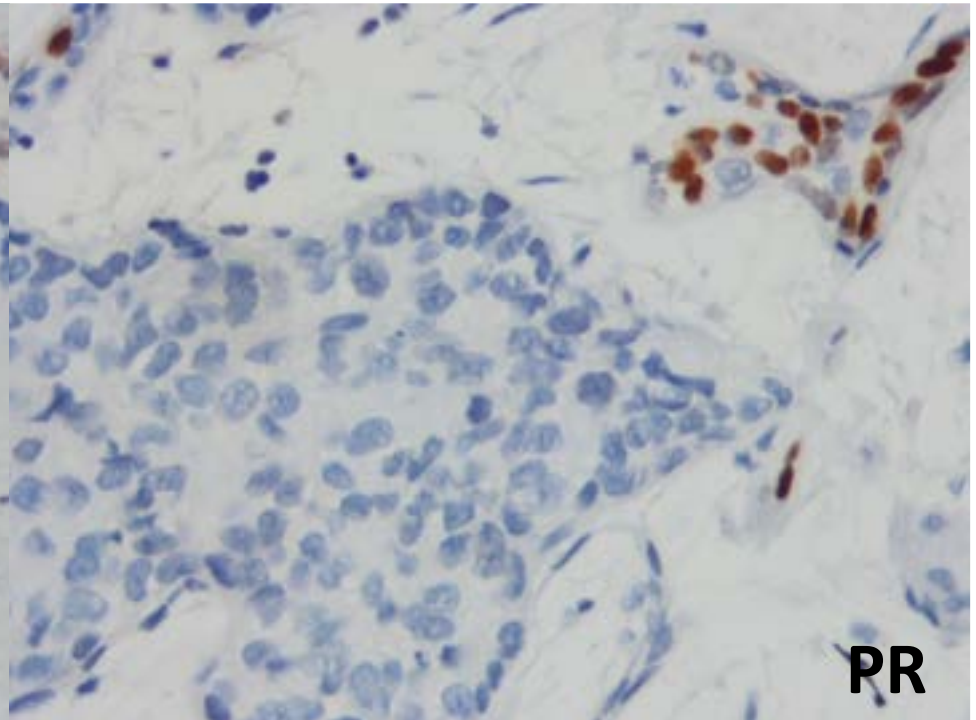
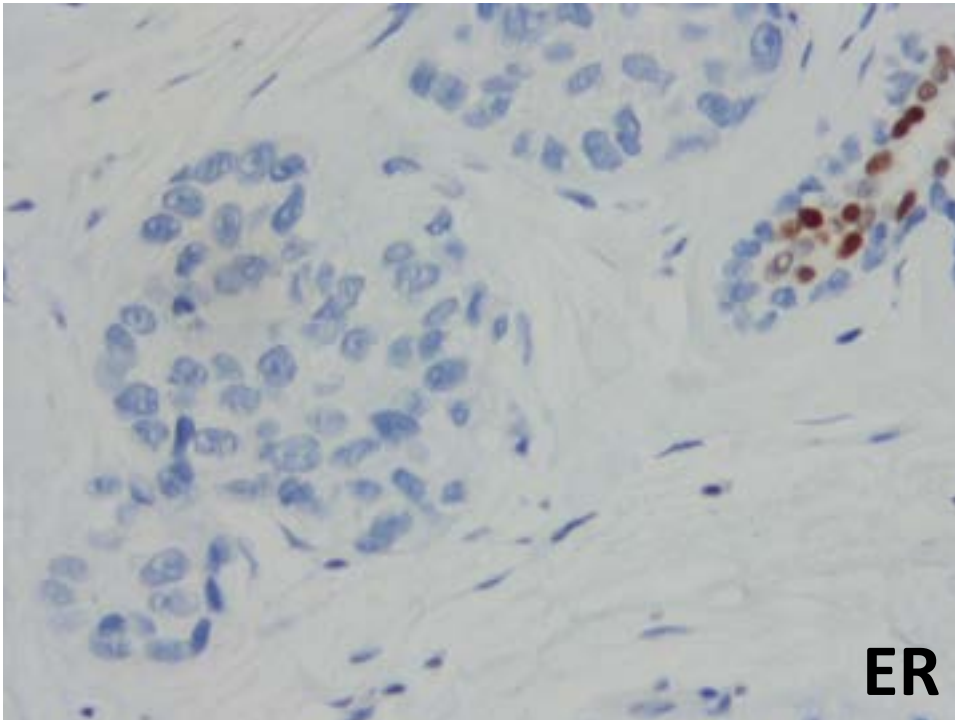
	No. of tumors with metastasis (%)	Risk ratio	95% confidence interval	p value*
A. Metastasis				
1. Metastasis to any organ				
Cases (n = 20)	13 (65)	2.74	1.28-5.86	0.0096
Control subjects (n = 40)	14 (35)			
2. Brain metastasis				
Cases (n = 20)	6 (30)	3.77	1.14-12.45	0.030
Control subjects (n = 40)	5 (13)			
3. Lung metastasis				
Cases (n = 20)	9 (45)	3.67	1.40-9.61	0.008
Control subjects (n = 40)	8 (20)			
4. Bone metastasis				
Cases (n = 20)	4 (20)	1.18	0.36-3.86	NS
Control subjects (n = 40)	5 (13)			
5. Locoregional recurrence				
Cases (n = 20)	4 (20)	1.41	0.42-4.74	NS
Control subjects (n = 40)	8 (20)			
6. Liver metastasis				
Cases (n = 20)	1 (5)	0.78	0.087-7.08	NS
Control subjects (n = 40)	4 (10)			
B. Death by cancer				
Cases (n = 20)	10 (50)	3.78	1.48-9.63	0.0054
Control subjects (n = 40)	8 (20)			

IDC: Invasive ductal carcinoma



Triple-negative breast cancer

- Tumour cells negative for ER,PR and HER2
- 10 to 15% of sporadic breast cancer cases
- Characteristics include:
 - higher prevalence among premenopausal African-American patients
 - high nuclear grade and proliferative indices
 - frequently abnormalities on p53 and BRCA 1 genes
 - chemosensitive but poor prognosis
 - peak risk of recurrence is between first and third years and the majority of deaths occur in the first 5 years following therapy.



REVIEW ARTICLE

CURRENT CONCEPTS

Triple-Negative Breast Cancer

William D. Foulkes, M.B., B.S., Ph.D., Ian E. Smith, M.D.,
and Jorge S. Reis-Filho, M.D., Ph.D.

N ENGL J MED 363:20 NEJM.ORG NOVEMBER 11, 2010

Review

Basal-like and triple-negative breast cancers:
a critical review with an emphasis on the
implications for pathologists and oncologists

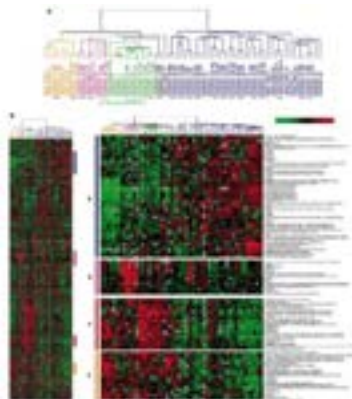
Sunil Badve¹, David J Dabbs², Stuart J Schnitt³, Frederick L Baehner⁴, Thomas Decker⁵,
Vincenzo Eusebi⁶, Stephen B Fox⁷, Shu Ichihara⁸, Jocelyne Jacquemier⁹, Sunil R Lakhani¹⁰,
José Palacios¹¹, Emad A Rakha¹², Andrea L Richardson¹³, Fernando C Schmitt¹⁴,
Puay-Hoon Tan¹⁵, Gary M Tee¹⁶, Britta Weigelt¹⁷, Ian O Ellis¹² and Jorge S Reis-Filho¹⁸

- There is still no internationally accepted definition for basal-like breast cancers and how best to define these tumours is a matter of controversy and ongoing debate.

letters to nature

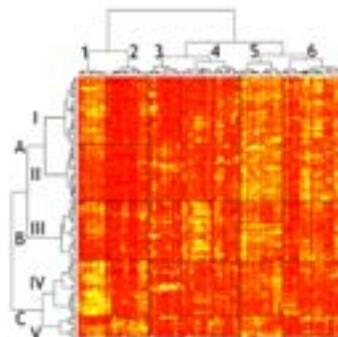
Molecular portraits of
human breast tumours

Charles M. Perou¹, Thorsteinn Sørlie¹, Michael B. Eisen²,
Mark van de Vijver³, Stefano G. Jeffrey⁴, Christian A. Lee⁵,
Jeffrey S. Perou⁶, Douglas T. Ross⁷, Mike Johnson⁸,
Lars H. Ståle⁹, Sigrun Haegler¹⁰, Alexander Pangloss-Medendorp¹¹,
Christel Wilshire¹², Gertine S. Zlob¹³, Per S. Lønning¹⁴,
Anne-Lise Børresen-Rasmussen¹⁵, Patrick S. Broome¹⁶ & Svend Støerli¹⁷



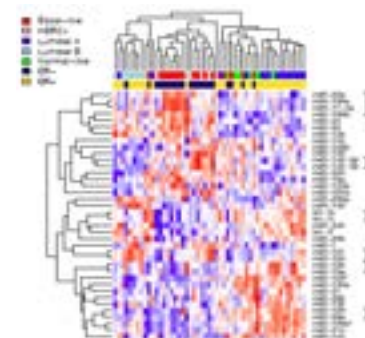
Surface-enhanced laser desorption/ionization time-of-flight
proteomic profiling of breast carcinomas identifies
clinicopathologically relevant groups of patients similar to
previously defined clusters from cDNA expression

Kristyna Brodskova¹, Eva Budinská¹, Pavel Bouchal^{1,2}, Lenka Heryšková¹, Dana Křoflíková¹,
Dalibor Vaňk¹, Rostislav Vyzula¹, Borivoj Vojtesek¹ and Rudolf Nemeš¹



MicroRNA expression profiling of human breast
cancer identifies new markers of tumour subtype

Cherie Blencowe^{1,2,3,4*}, Leonard D Goldbaum^{1,2,3*}, Natalie P Thomas^{1,2,3}, Immaculada
Spicer^{1,2}, Sun-Feng Chan^{1,2}, Mark J Dunning^{1,2}, Nuno L Barbosa-Morais^{1,2}, Andrew
E Tenchovskoff^{1,2}, Andrew R Green⁵, Ian O Ellis⁶, Suresh Tavakoli^{1,2,3}, Carlos
Caldas^{1,2,3}, Eric A Miska^{1,4*}



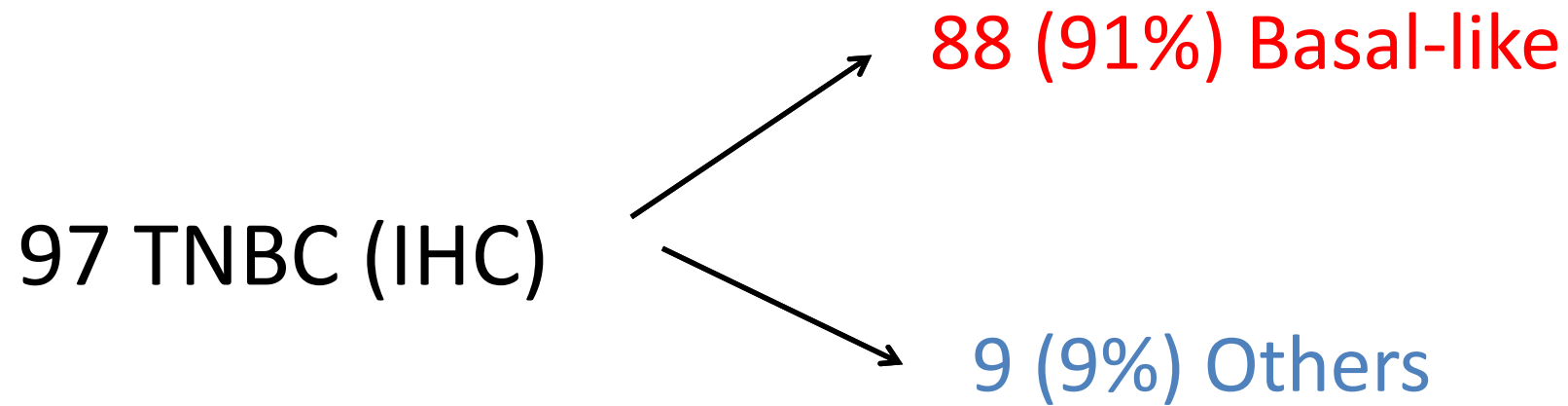
Basal-like breast carcinomas

Table 1 | Highlights of genomic, clinical and proteomic features of subtypes

Subtype	Luminal A	Luminal B	Basal-like	HER2E
ER ⁺ /HER2 ⁻ (%)	87	82	10	20
HER2 ⁺ (%)	7	15	2	68
TNBCs (%)	2	1	80	9
TP53 pathway	<i>TP53</i> mut (12%); gain of <i>MDM2</i> (14%)	<i>TP53</i> mut (32%); gain of <i>MDM2</i> (31%)	<i>TP53</i> mut (84%); gain of <i>MDM2</i> (14%)	<i>TP53</i> mut (75%); gain of <i>MDM2</i> (30%)
PIK3CA/PTEN pathway	<i>PIK3CA</i> mut (49%); <i>PTEN</i> mut/loss (13%); <i>INPP4B</i> loss (9%)	<i>PIK3CA</i> mut (32%) <i>PTEN</i> mut/loss (24%) <i>INPP4B</i> loss (16%)	<i>PIK3CA</i> mut (7%); <i>PTEN</i> mut/loss (35%); <i>INPP4B</i> loss (30%)	<i>PIK3CA</i> mut (42%); <i>PTEN</i> mut/loss (19%); <i>INPP4B</i> loss (30%)
RB1 pathway	Cyclin D1 amp (29%); <i>CDK4</i> gain (14%); low expression of <i>CDKN2C</i> ; high expression of <i>RB1</i>	Cyclin D1 amp (58%); <i>CDK4</i> gain (25%)	<i>RB1</i> mut/loss (20%); cyclin E1 amp (9%); high expression of <i>CDKN2A</i> ; low expression of <i>RB1</i>	Cyclin D1 amp (38%); <i>CDK4</i> gain (24%)
mRNA expression	High ER cluster; low proliferation	Lower ER cluster; high proliferation	Basal signature; high proliferation	HER2 amplicon signature; high proliferation
Copy number	Most diploid; many with quiet genomes; 1q, 8q, 8p11 gain; 8p, 16q loss; 11q13.3 amp (24%)	Most aneuploid; many with focal amp; 1q, 8q, 8p11 gain; 8p, 16q loss; 11q13.3 amp (51%); 8p11.23 amp (28%)	Most aneuploid; high genomic instability; 1q, 10p gain; 8p, 5q loss; <i>MYC</i> focal gain (40%)	Most aneuploid; high genomic instability; 1q, 8q gain; 8p loss; 17q12 focal <i>ERRB2</i> amp (71%)
DNA mutations	<i>PIK3CA</i> (49%); <i>TP53</i> (12%); <i>GATA3</i> (14%); <i>MAP3K1</i> (14%)	<i>TP53</i> (32%); <i>PIK3CA</i> (32%); <i>MAP3K1</i> (5%)	<i>TP53</i> (84%); <i>PIK3CA</i> (7%)	<i>TP53</i> (75%); <i>PIK3CA</i> (42%); <i>PIK3R1</i> (8%)
DNA methylation	–	Hypermethylated phenotype for subset	Hypomethylated	–
Protein expression	High oestrogen signalling; high MYB; RPPA reactive subtypes	Less oestrogen signalling; high FOXM1 and MYC; RPPA reactive subtypes	High expression of DNA repair proteins, PTEN and <i>INPP4B</i> loss signature (pAKT)	High protein and phospho-protein expression of EGFR and HER2

Percentages are based on 466 tumour overlap list. Amp, amplification; mut, mutation.

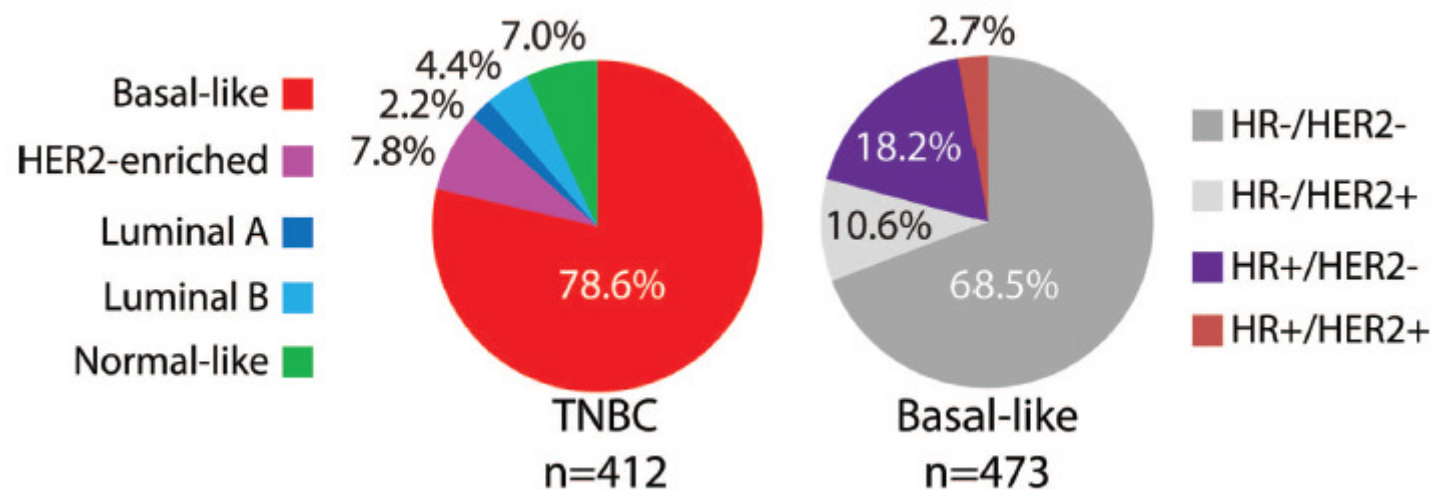
Not all TN are basal-like!



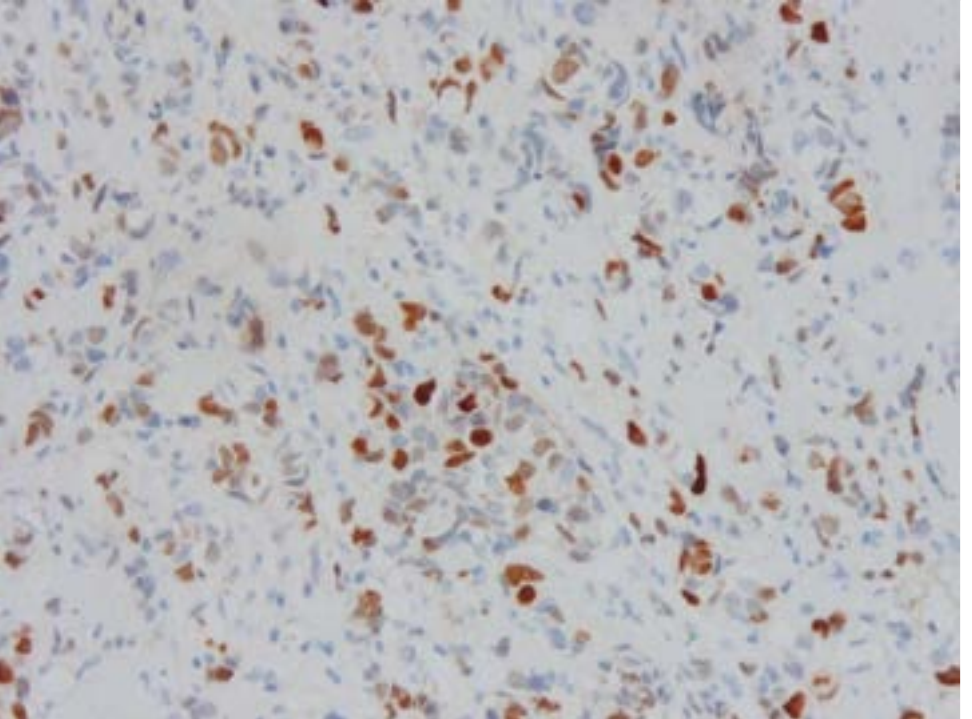
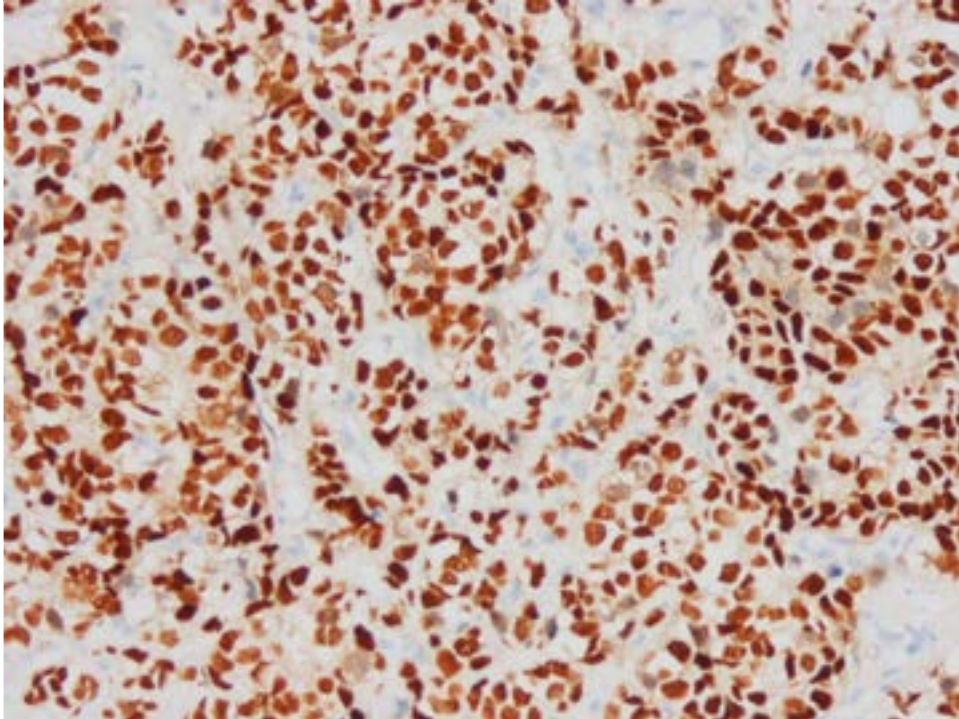
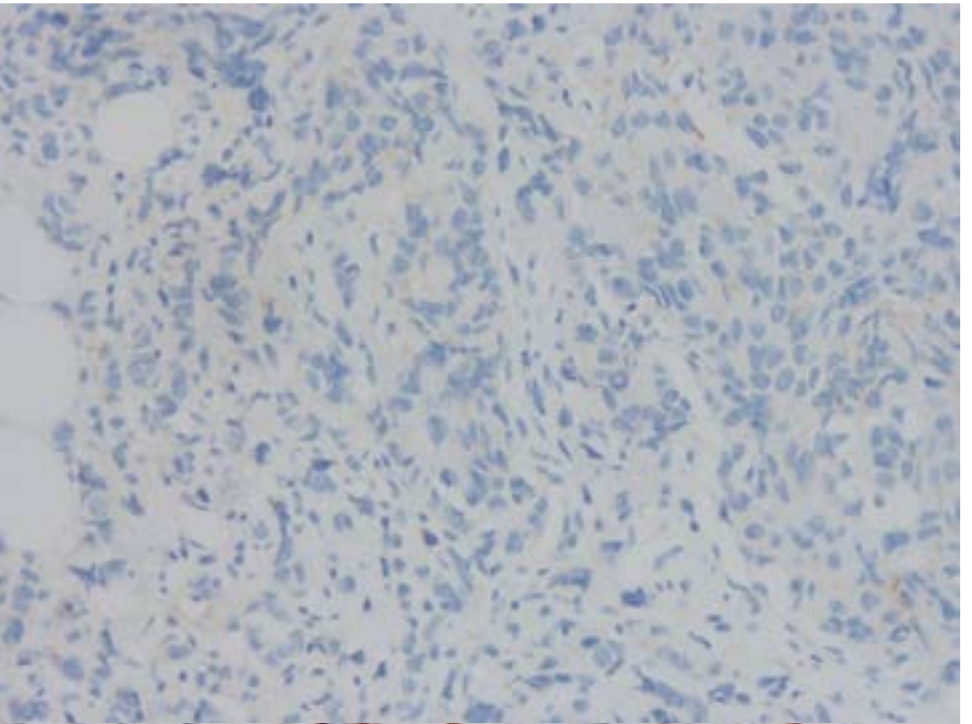
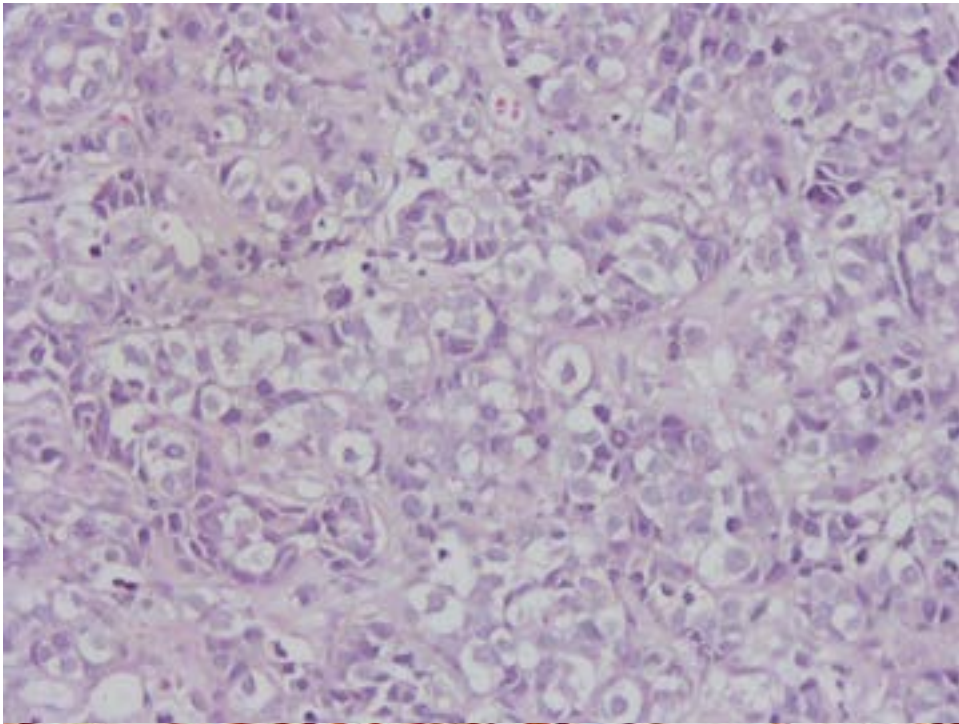
Molecular Characterization of Basal-Like and Non-Basal-Like Triple-Negative Breast Cancer

ALEIX PRAT,^{a,b,c} BARBARA ADAMO,^{b,c} MAGGIE C.U. CHEANG,^d CAREY K. ANDERS,^d LISA A. CAREY,^d CHARLES M. PEROU^{d,e,f}

The Oncologist 2013;18:123–133

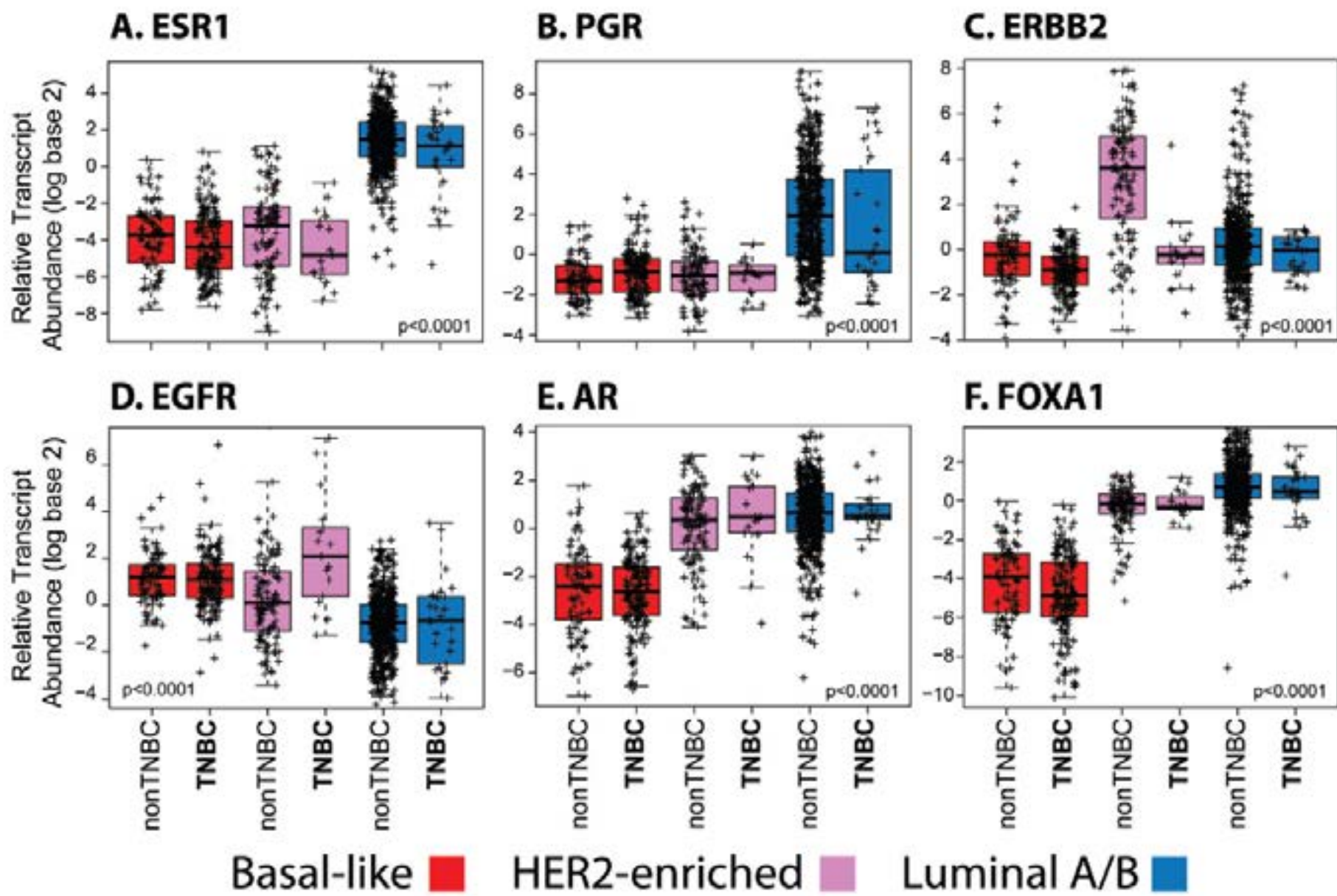


There are limitations to use IHC for Receptors as Surrogates for Molecular Subtype



TN and basal-like definitions should not be considering synonymous because considerable discordance exists (~25%)

- False-positivity or false-negativity of the IHC-based assays for determining HR and HER2 status, because these tests are challenged by interlaboratory and intermethod discordance rates.
- Assessment in different areas of the tumour ? Unlikely that two different subtypes coexist in the same tumour enough to explain the discordance rate.
- Gene expression measures a large number of related genes, compared with the 3 individual biomarkers used to define TN disease. For example, a TN tumour that has low levels of ESR1 and PGR might be luminal due to the expression of other luminal-related genes (GATA3 and/or FOX1A).

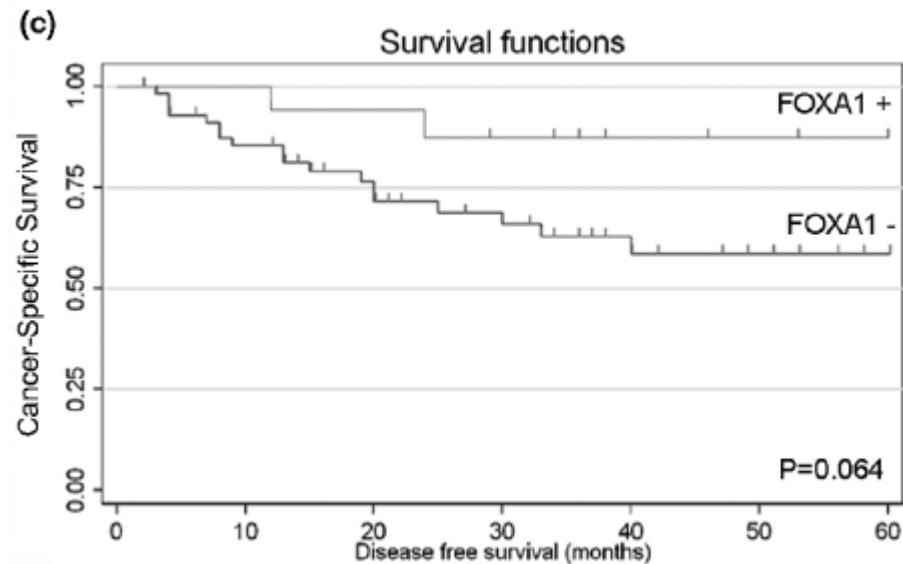


Expression of FOXA1 and GATA-3 in breast cancer: the prognostic significance in hormone receptor-negative tumours

André Albergaria^{1,2}, Joana Paredes², Bárbara Sousa², Fernanda Milanezi², Vitor Carneiro³, Joana Bastos^{4,5}, Sandra Costa¹, Daniella Vieira⁶, Nair Lopes², Eric W Lam⁷, Nuno Lunet^{4,5} and Fernando Schmitt^{2,8}

Breast Cancer Research 2009, **11**:R40 (doi:10.1186/bcr2327)

ER NEGATIVE TUMOURS

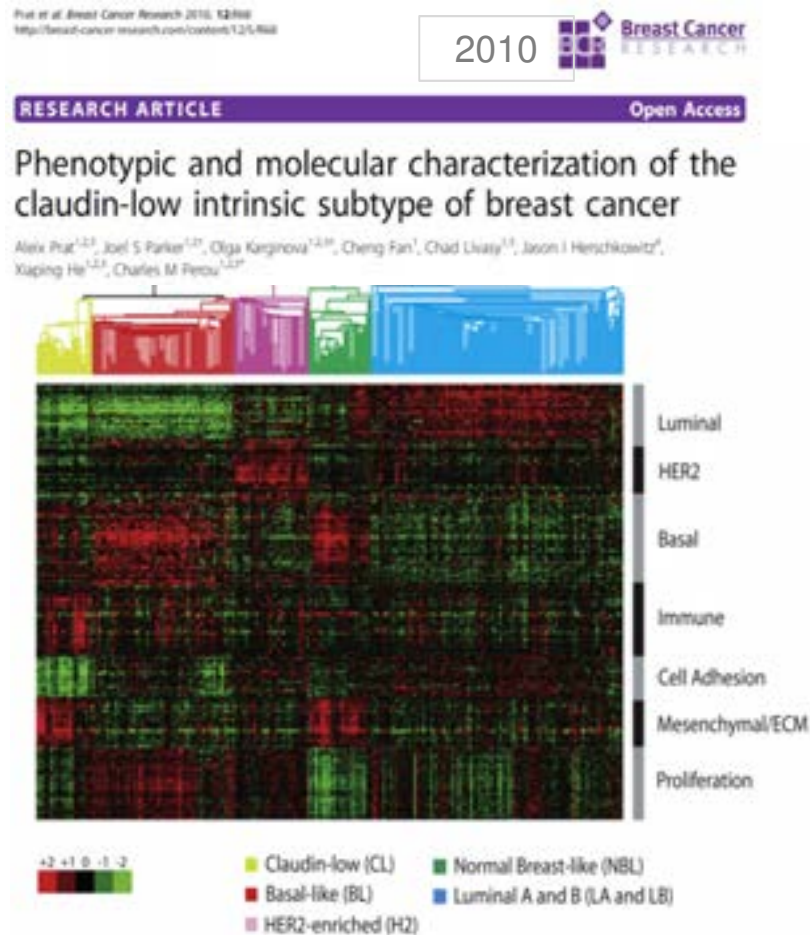


Triple-negative breast cancer is a heterogeneous clinical entity

- Gene expression profile classification revealed an heterogeneous group of breast malignancies:
 - Basal-like (EGFR and/or CK5/6 and /or CK14 and/or PCad)
 - Claudin-low (low/absent expression of adhesion molecules)
 - Molecular apocrine
 - Other intrinsic molecular subtypes
 - Normal-breast like (normal adipose tissue and other non epithelial and basal epithelial) ???

Claudin-low carcinomas

New molecular subgroup, sorted from the triple negative breast cancer group



- Low expression of genes involved in tight junctions and cell-cell adhesion:

- *Claudins 3, 4, 7,*
- *Occludin*
- *Ecadherin*

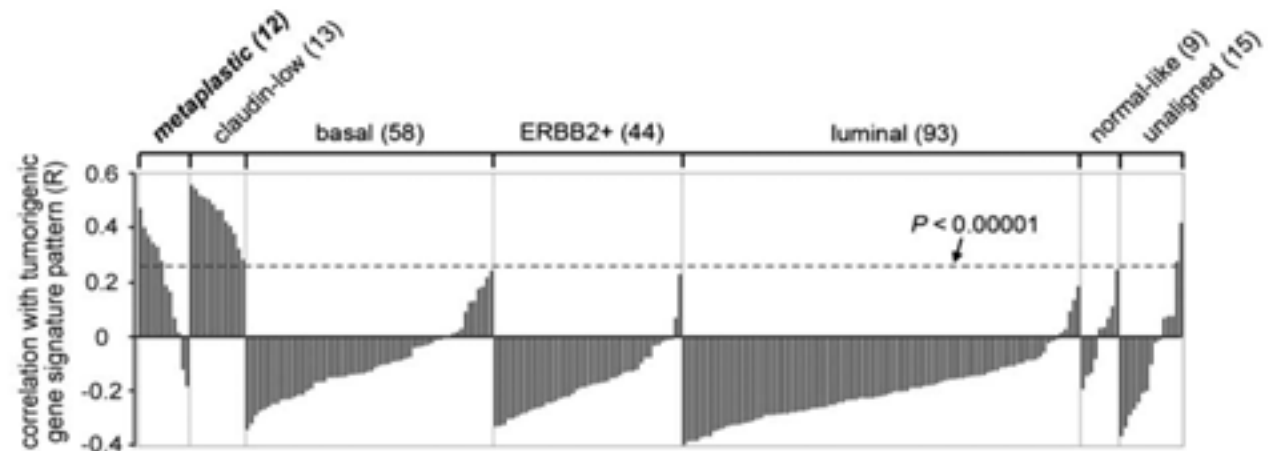
- Low expression of luminal genes,
- Inconsistent basal gene expression
- High expression of lymphocyte and endothelial cell markers

Characterization of a Naturally Occurring Breast Cancer Subset Enriched in Epithelial-to-Mesenchymal Transition and Stem Cell Characteristics

Bryan T. Hennessy,^{1,2*} Ana-Maria Gonzalez-Angulo,^{1,2*} Katherine Stenik-Hale,^{2*} Michael Z. Ciletti,¹ Savitri Krishnamurthy,¹ Ju Seog Lee,¹ Jane Fridlyand,¹ Aysegül Sahin,¹ Roshan Agarwal,¹ Corwin Jos,¹ Hsinbin Liu,¹ David Stivers,¹ Keith Baggerly,¹ Mark Carey,¹ Ana Iltis,¹ Carlos Montezano,¹ Xuping He,¹ Victor Weigman,¹ Cheng Fan,¹ Juan Palacios,¹ Gabriel N. Hortobagyi,¹ Laura K. Nolden,¹ Nicholas J. Wang,¹ Vicente Valero,¹ Joe W. Gray,¹ Charles M. Perou,¹ and Gordon B. Mills^{1,2}

Departments of ¹Genetics Medical Oncology, Systems Biology, Breast Medical Oncology, Pathology and ²Stem Cell Biology and ³Comparative Biology and ⁴Walter Center for Molecular Medicine, The University of Texas M. D. Anderson Cancer Center, Houston, Texas; ⁵Lawrence Berkeley National Laboratory, Berkeley, California; ⁶Ohio State Hospital and ⁷University of Illinois, Urbana, Illinois; ⁸Kingdon Comprehensive Cancer Center, Chapel Hill, North Carolina; and ⁹Department of Pathology, Thomas Jefferson University, Philadelphia, Pennsylvania

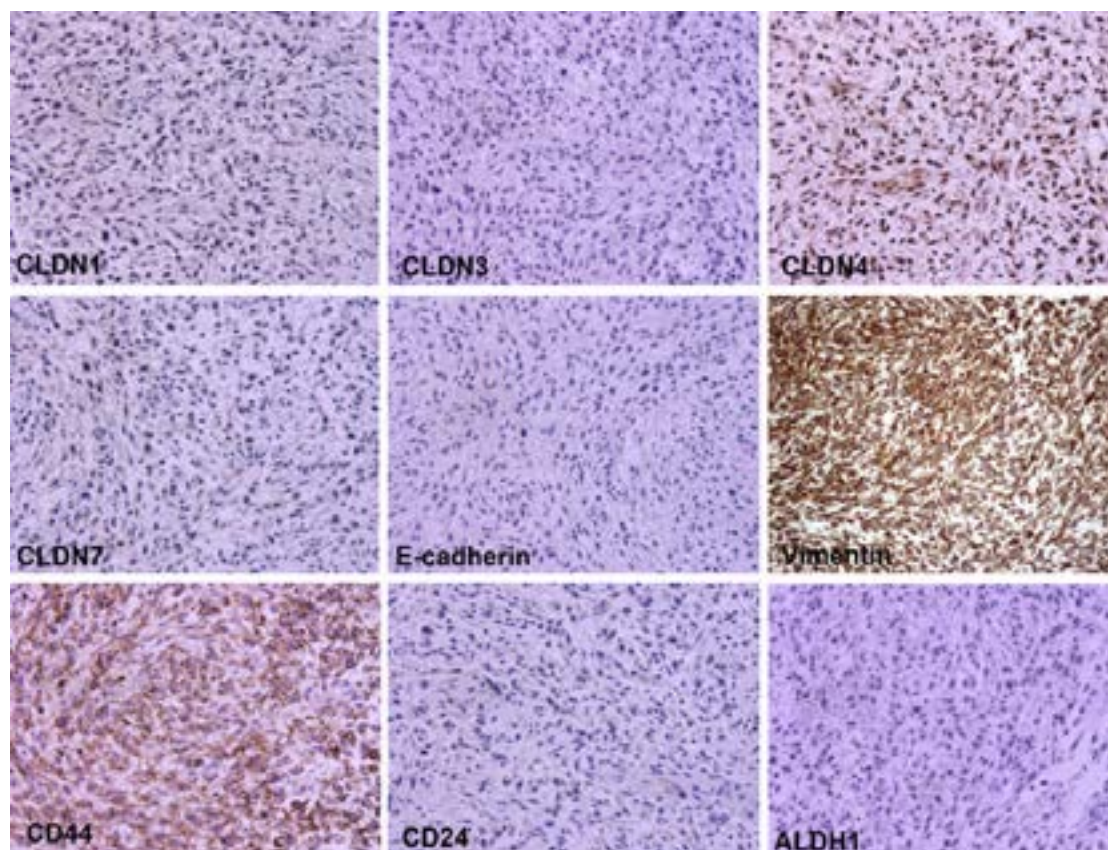
CD44+/CD24-/low phenotype



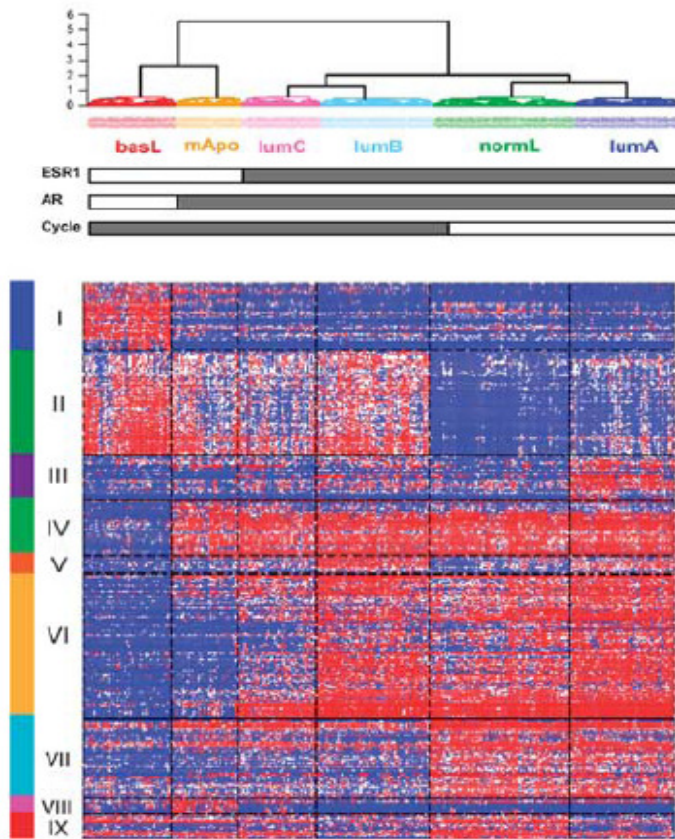
MBCs and Claudin-low tumors present similar transcriptional profiles and are enriched in stem cell characteristics

Original article

Immunohistochemical features of claudin-low intrinsic subtype in metaplastic breast carcinomas

Renê Gerhard^{a,g}, Sara Ricardo^{a,b,g}, André Albergaria^a, Madalena Gomes^a, Alfredo Ribeiro Silva^c,
Ângela Flavia Logullo^d, Jorge F. Cameselle-Teijeiro^e, Joana Paredes^{a,f}, Fernando Schmitt^{a,f,*}^aIPATBUP – Institute of Molecular Pathology and Immunology of Porto University, Porto, Portugal^bICBAS – Abel Salazar Biomedical Science Institute, Porto, Portugal^cDepartment of Pathology, Medical Faculty, University of São Paulo, Ribeirão Preto, Brazil^dDepartment of Pathology, School of Medicine, Federal University of São Paulo, São Paulo, Brazil^eComplejo Hospitalar Universitario de Vigo (CHUVI), Vigo, Spain^fMedical Faculty of Porto University, Porto, Portugal

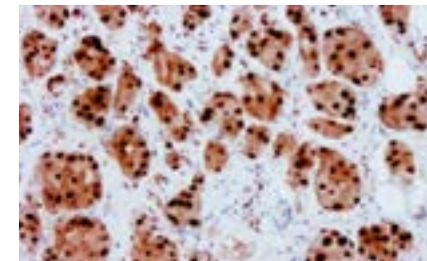
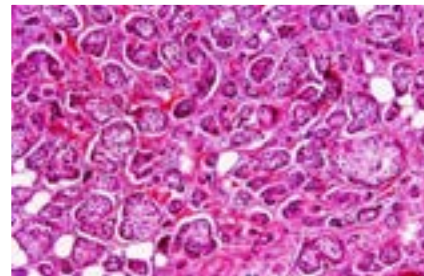
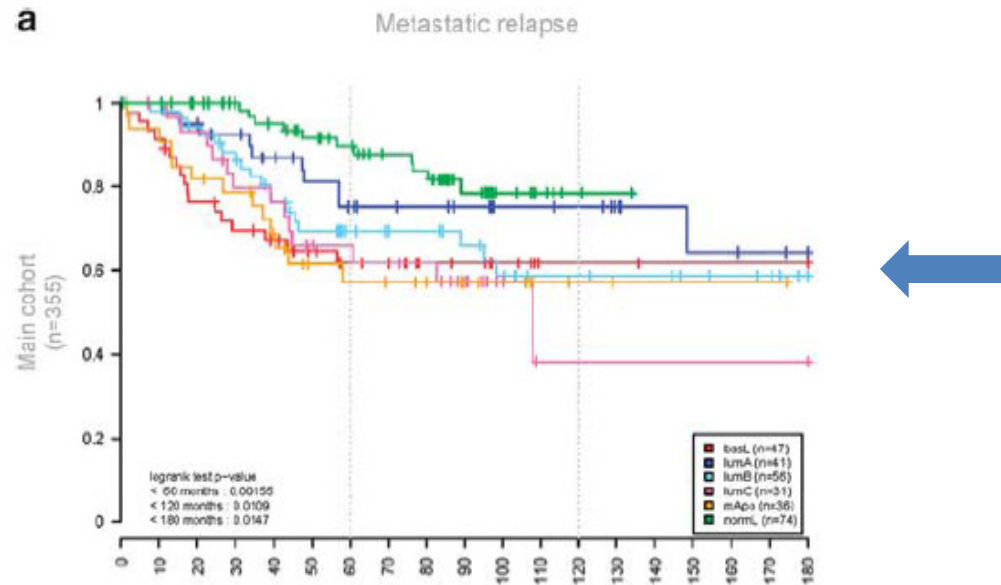
Molecular Apocrine



Benign and malignant apocrine lesions of the breast

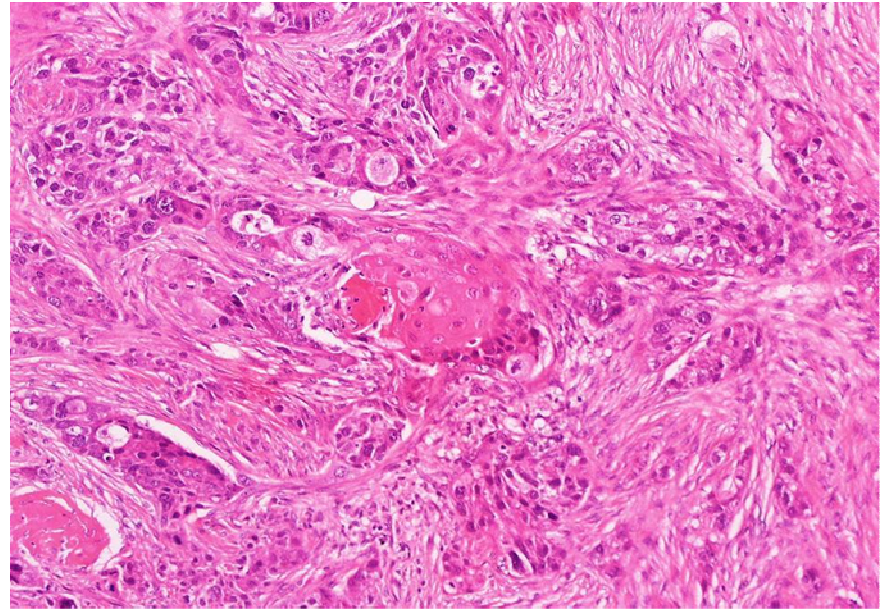
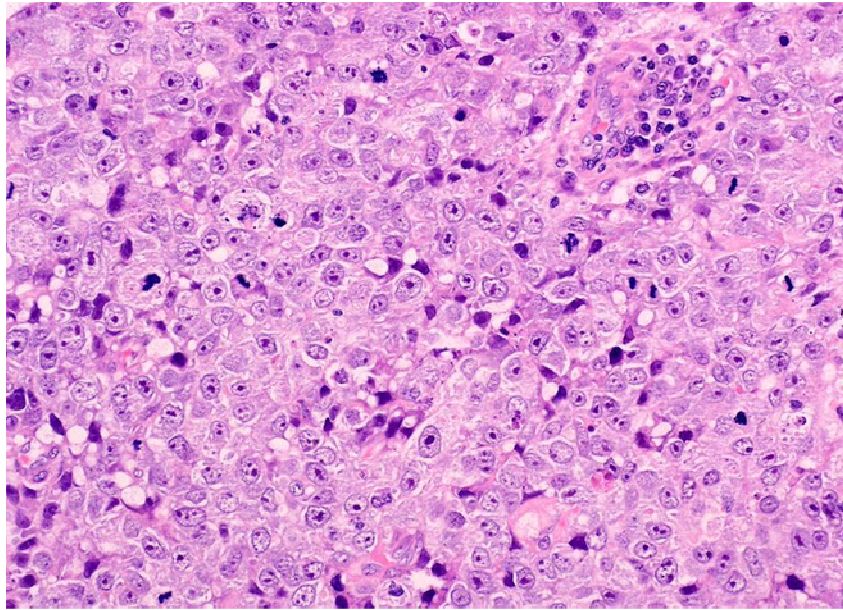
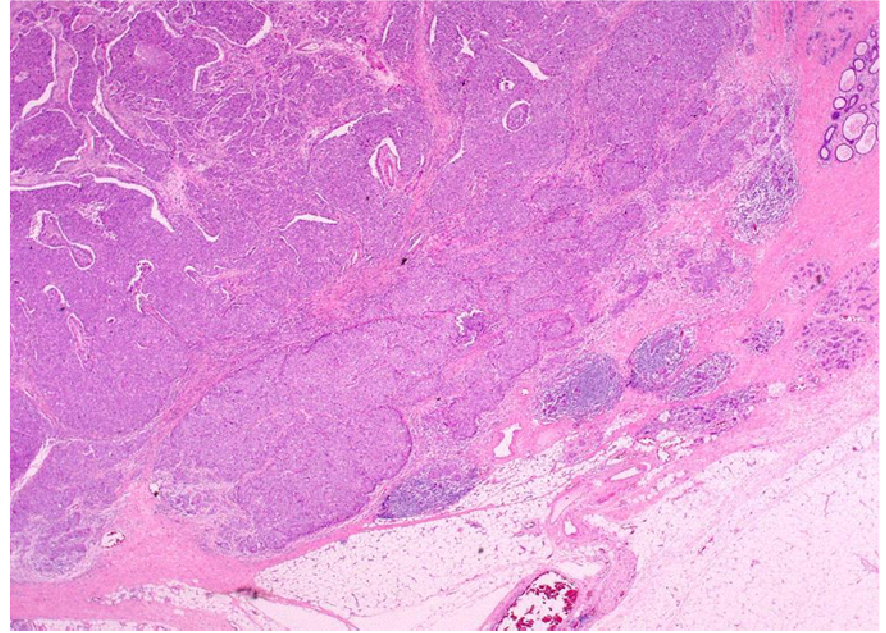
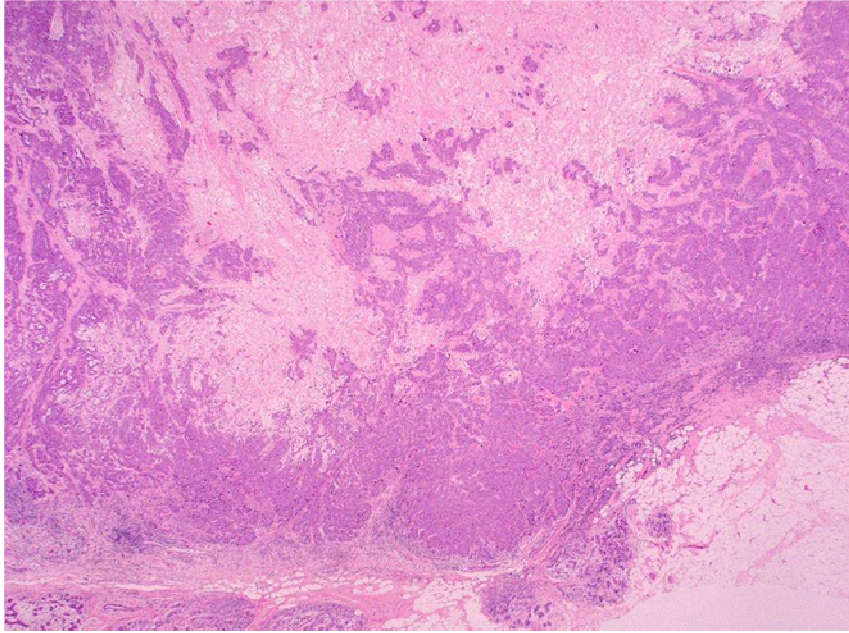
Expert Rev. Anticancer Ther. 12(2), 215–221 (2012)

Renê Gerhard¹,
 José Luis Costa¹ and
 Fernando Schmitt^{*1,2}



Histology of Basal-Like Cancers Identified By Expression Profiling

- Histologic grade 3 (100%)
- Solid architecture
- No tubule formation, high density of cells with no intervening stroma
- Pushing border (61%)
- Stromal lymphocytic infiltrate (56%)
- High mitotic rate (100%)
- Geographic zones of necrosis (74%)
- Medullary-like features
- *(Central fibrotic/acellular zone)*
- *(Little or no associated DCIS)*



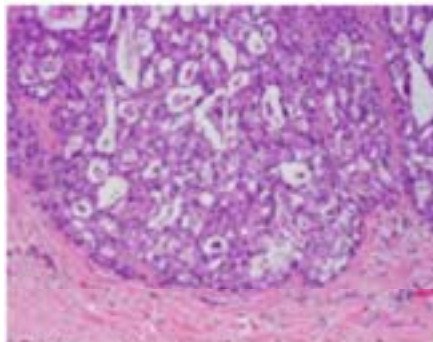
Do we still need a morphological classification?

“Triple-Negative” breast carcinomas

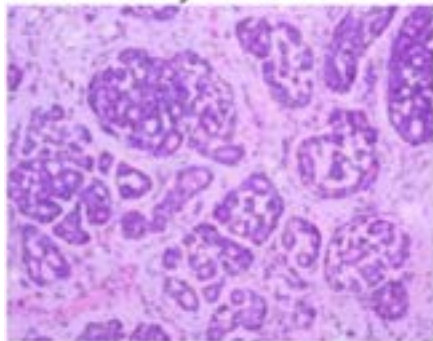
Low grade tumours

High grade tumours

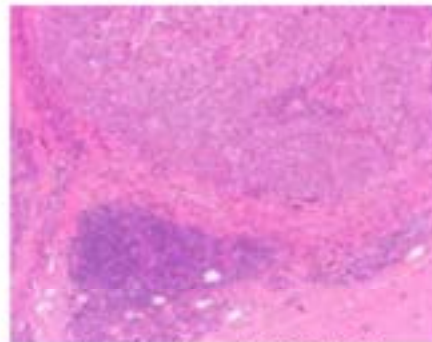
Secretory carcinoma



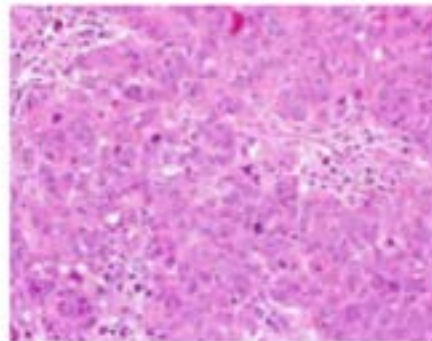
Adenoid cystic carcinoma



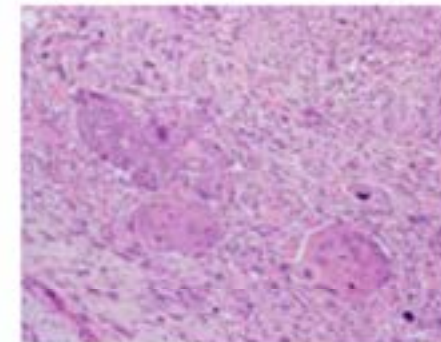
Medullary breast cancer



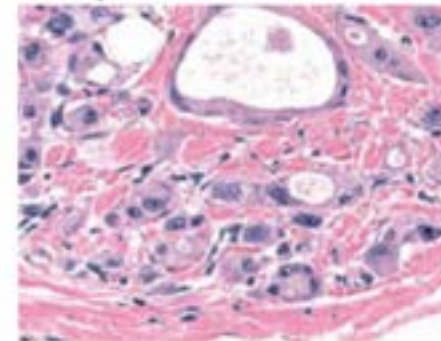
Grade 3 – IDC-NST



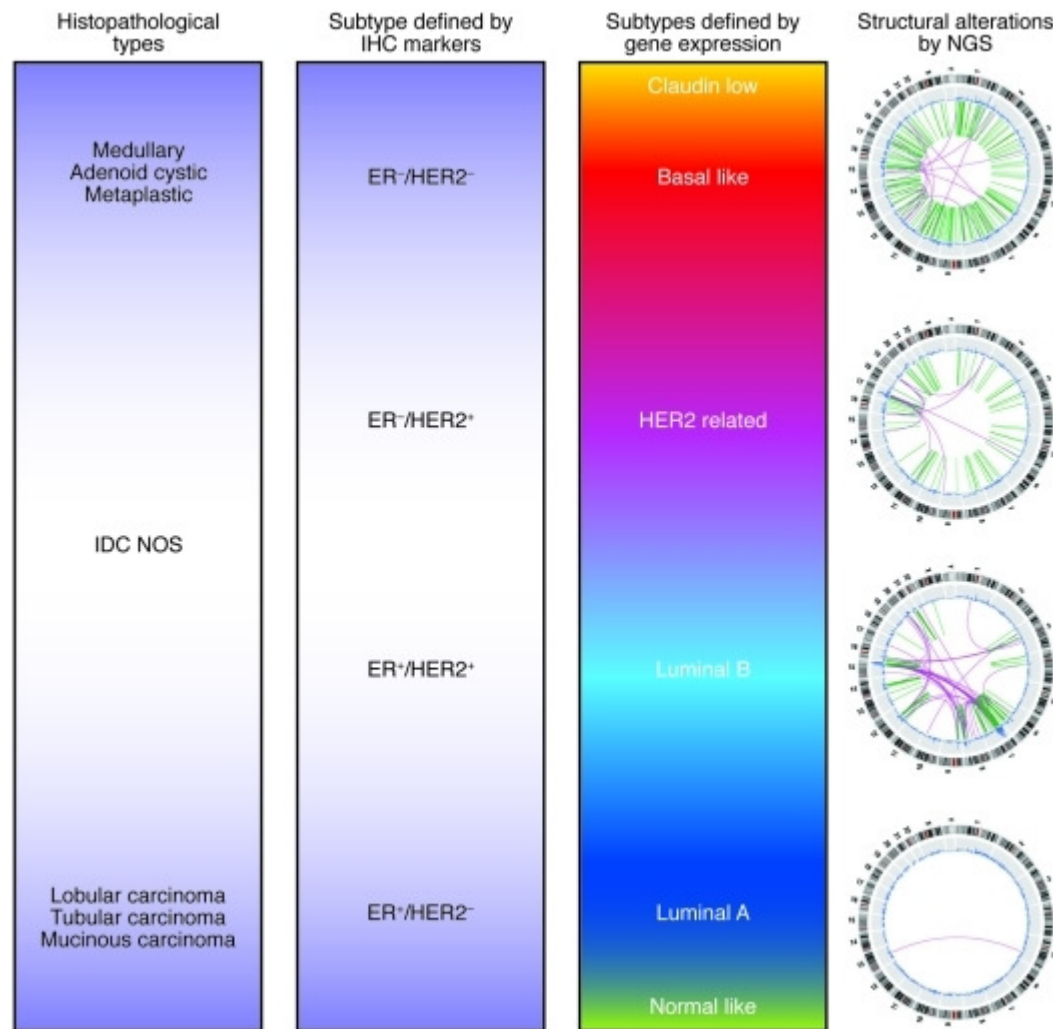
Metaplastic breast cancer



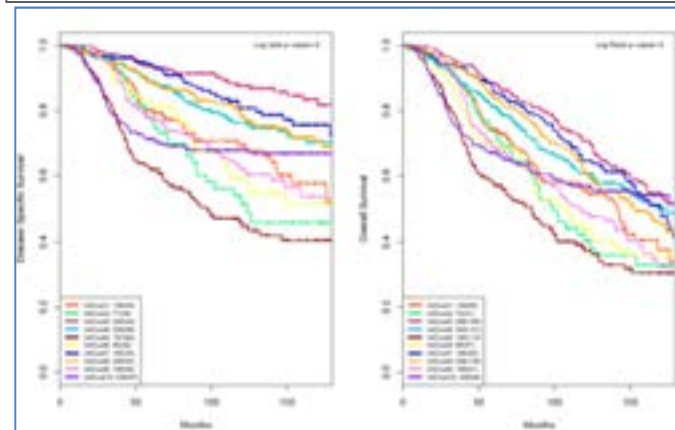
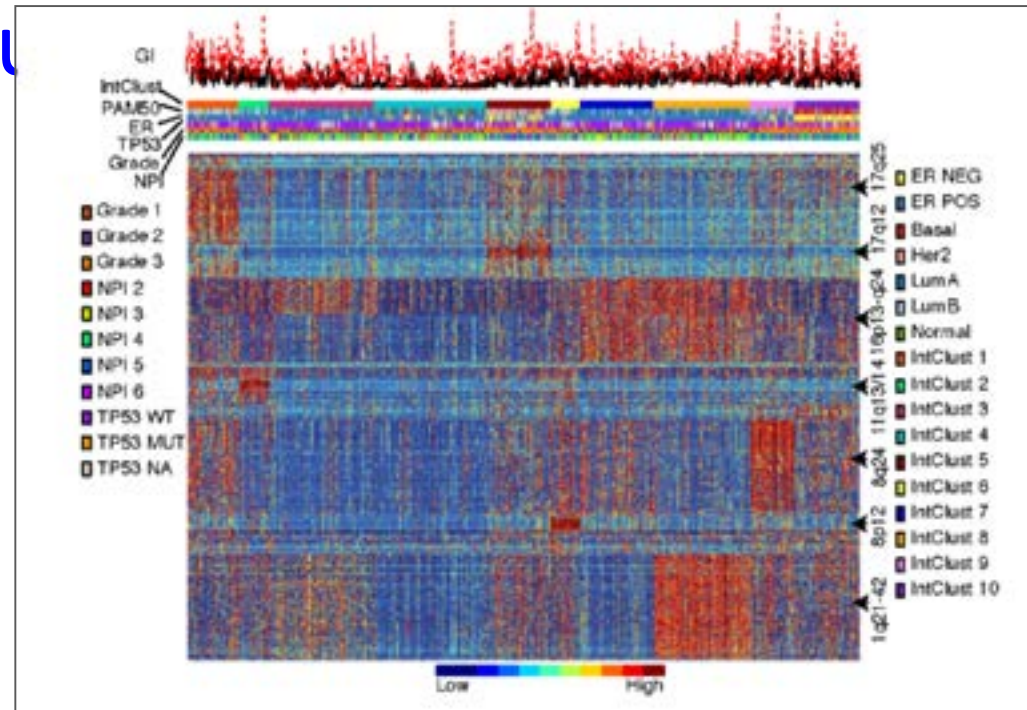
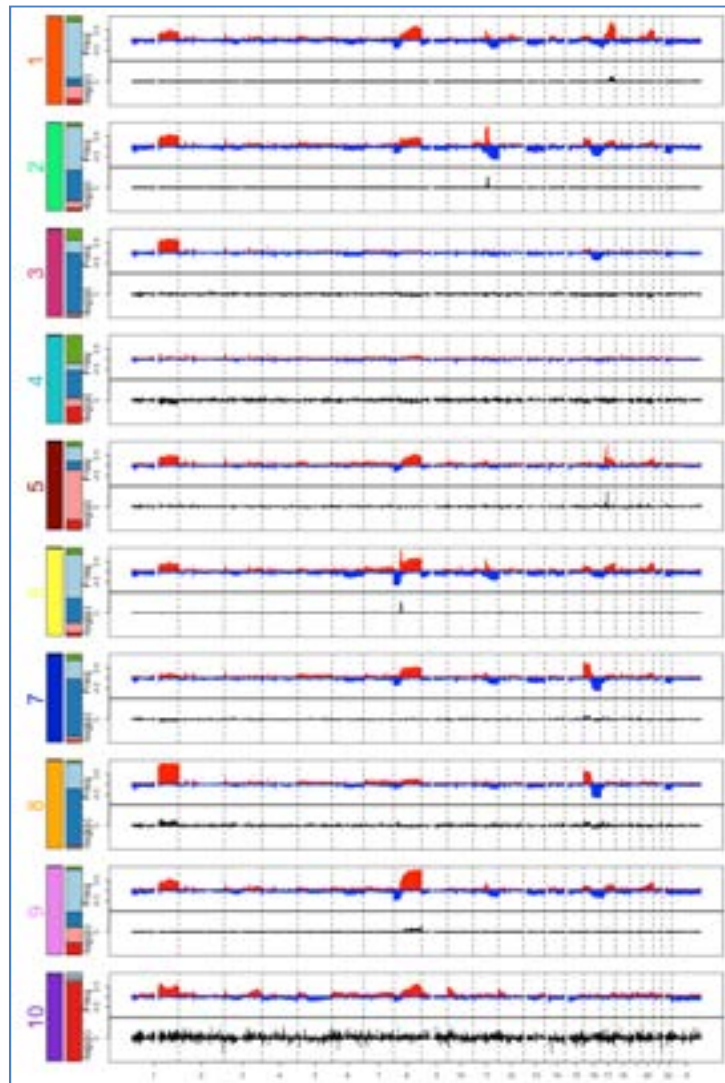
Apocrine Carcinoma



Breast cancer classification

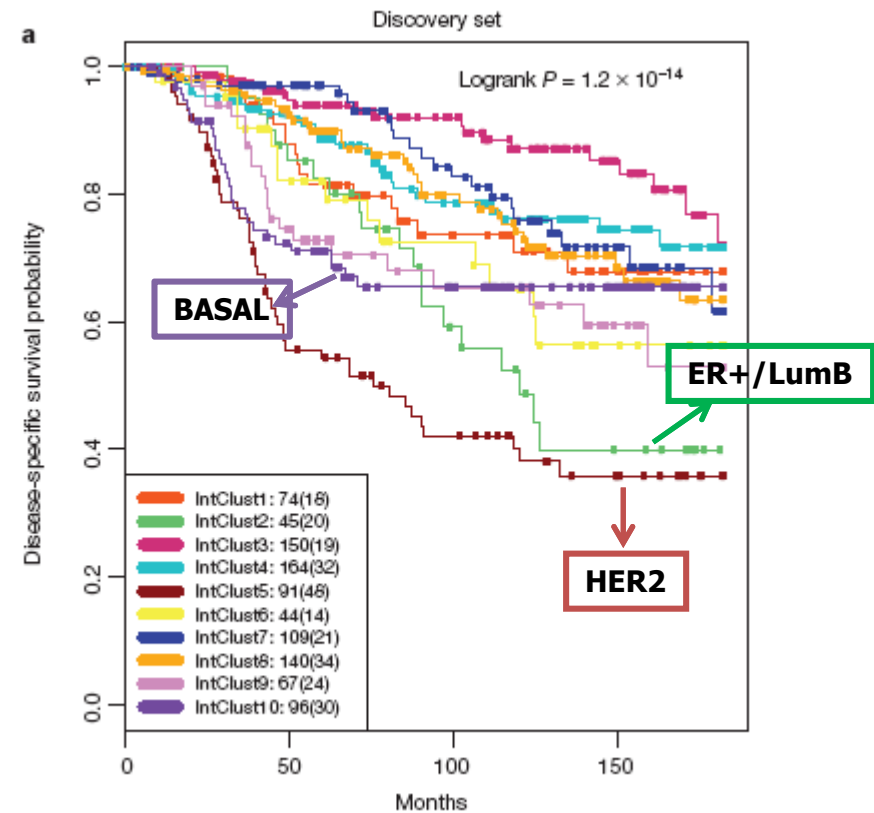
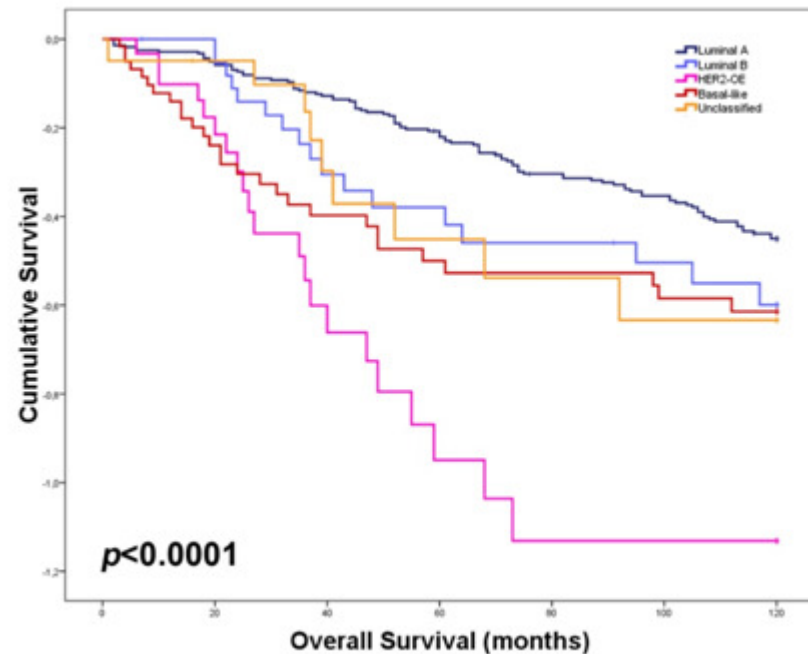


The genomic and transcriptomic architecture of 2,000 breast tumours reveals novel



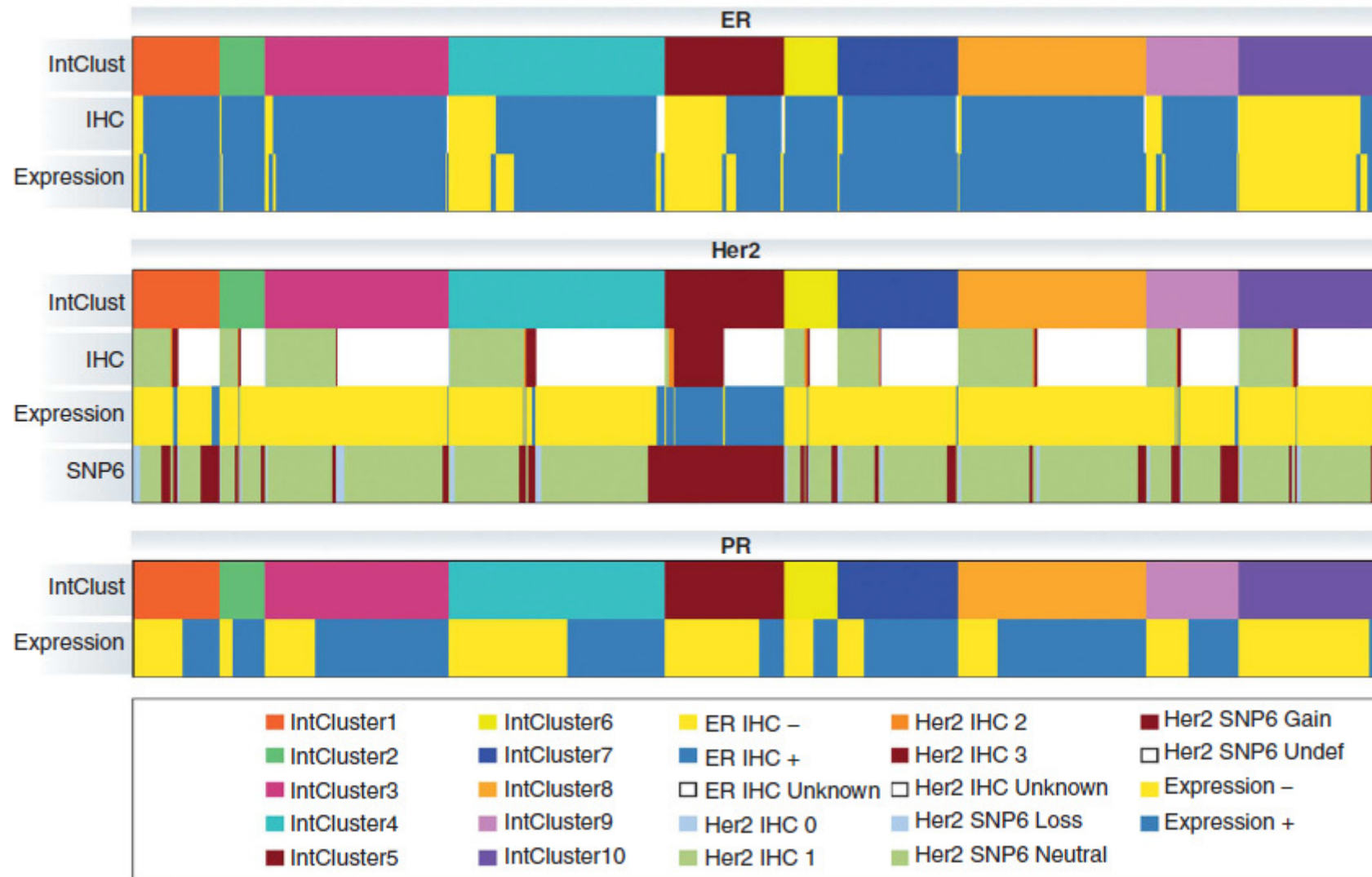
Curtis C et al. Nature

Integrative clusters and survival



The open question:
How can we integrate these subtypes into
daily clinical work?

A new genome-driven integrated classification of breast cancer and its implications



Landscape of somatic mutations in 560 breast cancer whole-genome sequences

Serena Nik-Zainal^{1,2}, Helen Davies¹, Johan Staaf³, Manasa Ramakrishna¹, Dominik Glodzik¹, Xueqing Zou¹, Inigo Martincorena¹, Ludmil B. Alexandrov^{1,4,5}, Sancha Martin¹, David C. Wedge¹, Peter Van Loo^{1,6}, Young Seok Ju¹, Marcel Smid⁷, Arie B. Brinkman⁸, Sandro Morganella⁹, Miriam R. Aure^{10,11}, Ole Christian Lingjærde^{11,12}, Anita Langerød^{10,11}, Markus Ringnér³, Sung-Min Ahn¹³, Sandrine Boyault¹⁴, Jane E. Brock¹⁵, Annegien Broeks¹⁶, Adam Butler¹, Christine Desmedt¹⁷, Luc Dirix¹⁸, Serge Dronov¹, Aquila Fatima¹⁹, John A. Foekens⁷, Moritz Gerstung¹, Gerrit K. J. Hooijer²⁰, Se Jin Jang²¹, David R. Jones¹, Hyung-Yong Kim²², Tari A. King²³, Savitri Krishnamurthy²⁴, Hee Jin Lee²¹, Jeong-Yeon Lee²⁵, Yilong Li¹, Stuart McLaren¹, Andrew Menzies¹, Ville Mustonen¹, Sarah O'Meara¹, Iris Pauporté²⁶, Xavier Pivot²⁷, Colin A. Purdie²⁸, Keiran Raine¹, Kamna Ramakrishnan¹, F. Germán Rodríguez-González⁷, Gilles Romieu²⁹, Anieta M. Sieuwerts⁷, Peter T. Simpson³⁰, Rebecca Shepherd¹, Lucy Stebbings¹, Olafur A. Stefansson³¹, Jon Teague¹, Stefania Tommasi³², Isabelle Treilleux³³, Gert G. Van den Eynden^{18,34}, Peter Vermeulen^{18,34}, Anne Vincent-Salomon³⁵, Lucy Yates¹, Carlos Caldas³⁶, Laura van't Veer¹⁶, Andrew Tutt^{37,38}, Stian Knappskog^{39,40}, Benita Kiat Tee Tan^{41,42}, Jos Jonkers¹⁶, Åke Borg³, Naoto T. Ueno²⁴, Christos Sotiriou¹⁷, Alain Viari^{43,44}, P. Andrew Futreal^{1,45}, Peter J. Campbell¹, Paul N. Span⁴⁶, Steven Van Laere¹⁸, Sunil R. Lakhani^{30,47}, Jorunn E. Eyfjord³¹, Alastair M. Thompson^{28,48}, Ewan Birney⁹, Hendrik G. Stunnenberg⁸, Marc J. van de Vijver²⁰, John W. M. Martens⁷, Anne-Lise Børresen-Dale^{10,11}, Andrea L. Richardson^{15,19}, Gu Kong²², Gilles Thomas⁴⁴ & Michael R. Stratton¹



Minions Language



English:
Hello!
Goodbye!
Thank you!
For you
Marriage
Apples
Ice-cream
I'm sorry
I'm hungry
Ugly!
I swear...
Fire
We love you
I hate you!
What
Cheers
Kiss kiss

Minions:
Bellol
Poopayel
Tank yul
Para tu
La boda
Papples
Gelato
Bi-do
We want bananal
Bananoninal
Underwear...
Bee-do-bee-do-bee-do
Tulaliloo ti amo
Tatata-bala-tu
Po-ka
Kampai
Muak muak muak



Massively Parallel Sequencing-based studies of Breast Cancer

- The collection of genetic aberrations found in breast cancer is complex with a limited number of genes that are frequently mutated in unselected cases.
- The number of genes mutated in small minorities of breast cancer is vast.
- The repertoire of mutations in luminal and basal-like breast cancer is rather different.
- There is no gene or mutation that defines a subtype of breast cancer.
- These studies led to the identification of novel driver genes and that genes that encodes ER alpha (ESR1) and HER2 can be targeted by activating mutations.

Molecular Classification

Conclusions

- GEP studies have provided significant advances in the molecular classification and prognostication of breast cancer, and has given new insights regarding therapeutic prediction.
- The clinical management of patients is still based on the assessment of morphology, ER,PR, HER2 and Ki67.
- New avenues for discovering and validating prognostic and predictive biomarkers are being developed through NGS approaches.

Breast Cancer: prognostication and therapy prediction

First Generation Gene Signatures

Systemic Therapy

Second Generation Gene Signatures

Predictive gene signatures

Novel avenues for prognostication and therapy prediction

ER POSITIVE

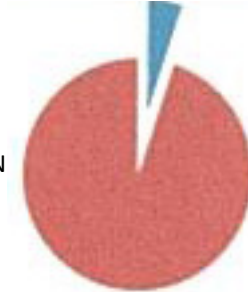
ER NEGATIVE

PROGNOSTIC SIGNATURES



LOW PROLIFERATION

HIGH PROLIFERATION



LOW PROLIFERATION	HIGH PROLIFERATION
HIGHER ENDOCRINE THERAPY BENEFIT	HIGHER CHEMOTHERAPY BENEFIT

HIGH PROLIFERATION
HIGHER CHEMOTHERAPY BENEFIT

Stromal signatures

Immune-response signatures

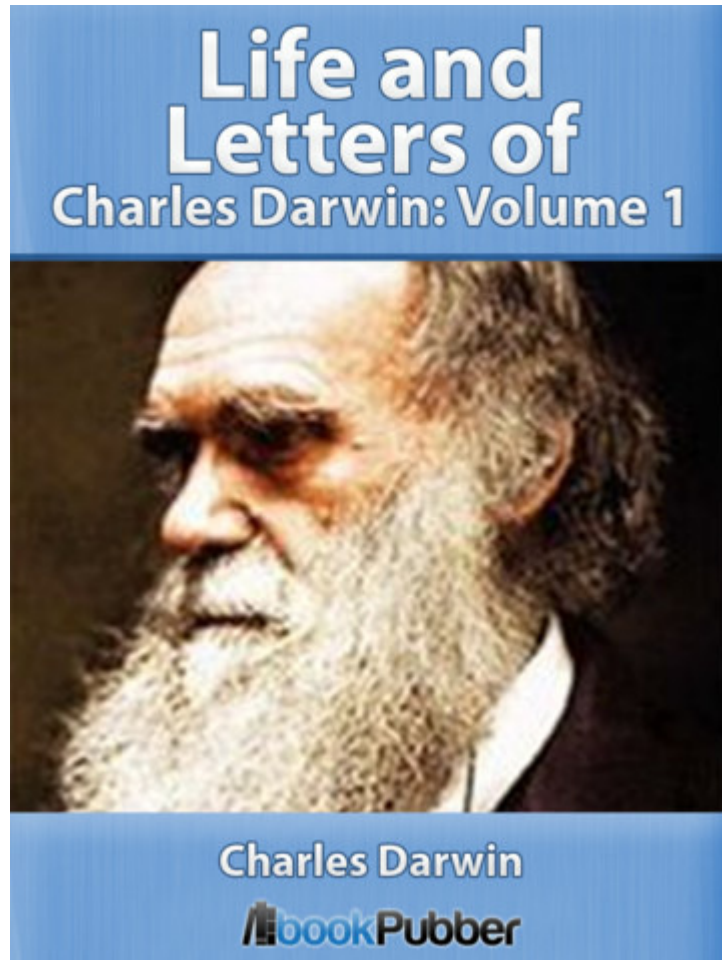
Predict signature (SET index)

LOW IMMUNE-RESPONSE	HIGH IMMUNE-RESPONSE
LOW BENEFIT OF CHEMOTHERAPY	HIGH BENEFIT OF CHEMOTHERAPY

Predict signatures

MASSIVE PARALLEL SEQUENCING

Taxonomy dilemma: *lumpers vs splitters*



- “Those who make many species are splitters, and those who make few are the lumpers”.
- In Medicine, this divide is exacerbated when a clear mechanistic understanding of a disease entity is incomplete.

Taxonomy of breast cancer based on normal cell phenotype predicts outcome

Sandro Santagata,¹ Ankita Thakkar,² Ayse Ergonul,² Bin Wang,² Terri Woo,¹ Rong Hu,^{3,4} J. Chuck Harrell,⁵ George McNamara,² Matthew Schwede,⁶ Aedin C. Culhane,⁶ David Kindelberger,¹ Scott Rodig,¹ Andrea Richardson,¹ Stuart J. Schnitt,⁷ Rulla M. Tamimi,^{3,4} and Tan A. Ince²

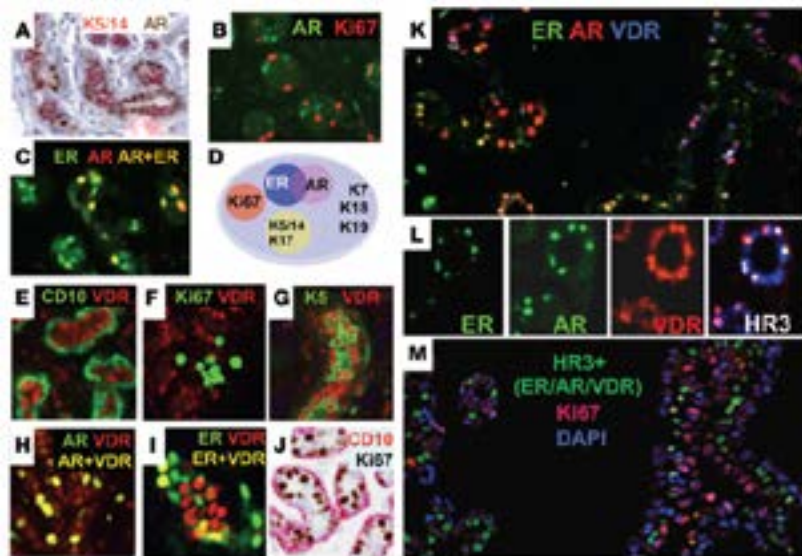
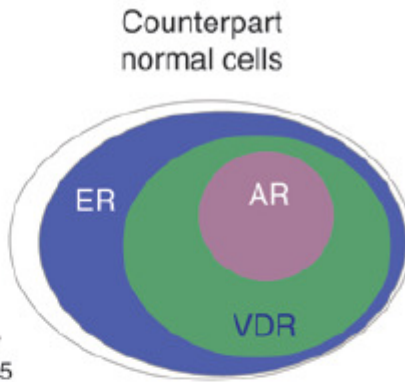
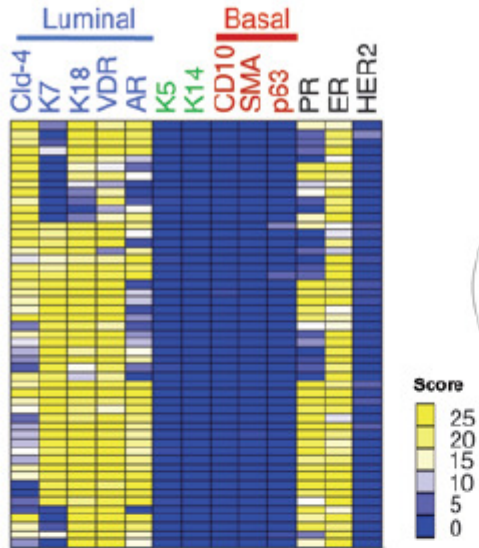


Table 1
Cellular differentiation states in normal human breast lobules

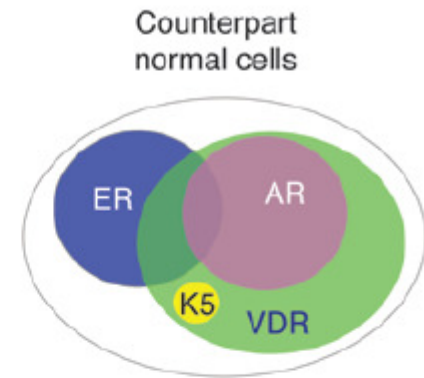
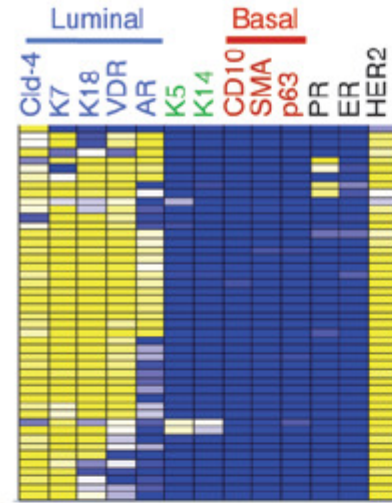
Cell type	ER	AR	VDR	KS/14/17	Ki67	Clc-4	K7/8/18	CD10/SMA/p63
Luminal								
L1 (HR0)	Ki67+	-	-	-	+	+	+	-
L2 (HR0)	Ki67+	-	-	-	-	+	+	-
L3 (HR0)	KS-	-	-	+	-	+	+	-
L4 (HR1)	ER+	+	-	-	-	+	+	-
L5 (HR1)	AR+	+	-	-	-	+	+	-
L6 (HR1)	VDR+	-	+	-	-	+	+	-
L7 (HR1)	KS-VDR+	-	+	+	-	+	+	-
L8 (HR2)	ER-AR-	+	-	-	-	+	+	-
L9 (HR2)	ER-VDR+	+	+	-	-	+	+	-
L10 (HR2)	AR-VDR+	+	+	-	-	+	+	-
L11 (HR3)	ER-AR-VDR+	+	+	-	-	+	+	-
Myoepithelial								
My1	CD10+	-	-	-	-	-	-	+
My2	KS+	-	-	+	-	-	-	+



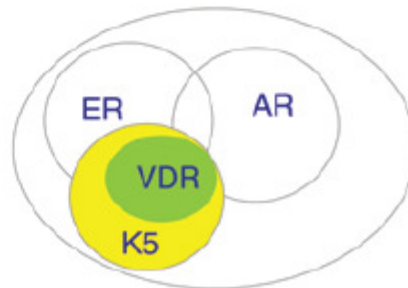
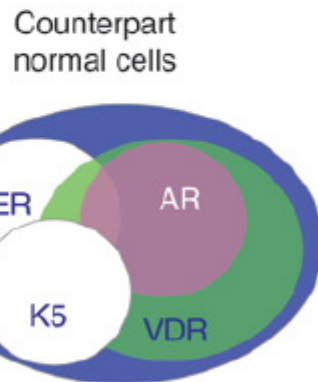
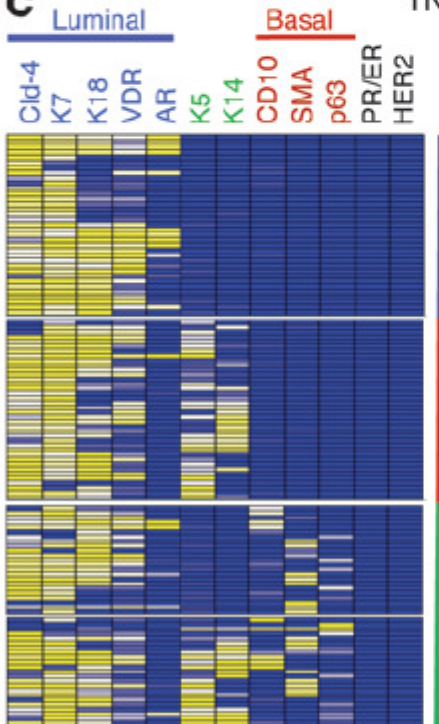
A ER⁺ tumors



B HER⁺ tumors



C TNBC tumors



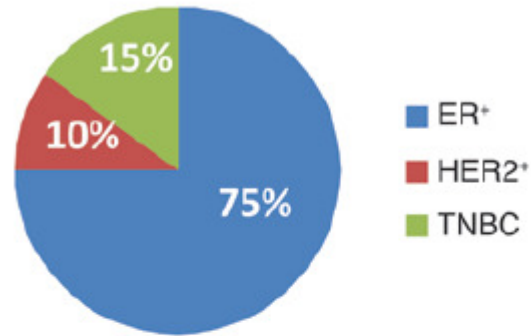
HR0 (ER-AR-VDR-)

HR1 (ER+,VDR+, or AR+)

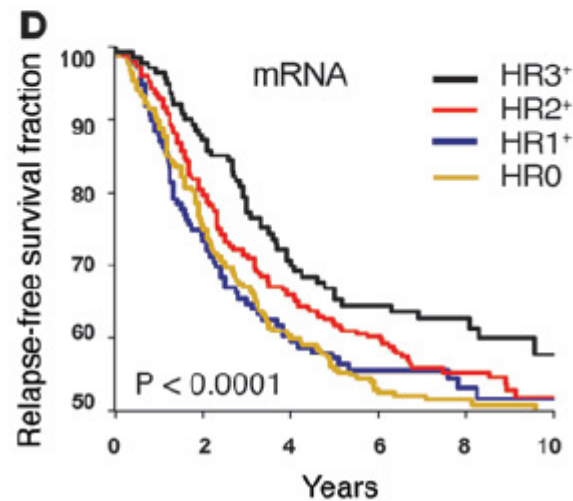
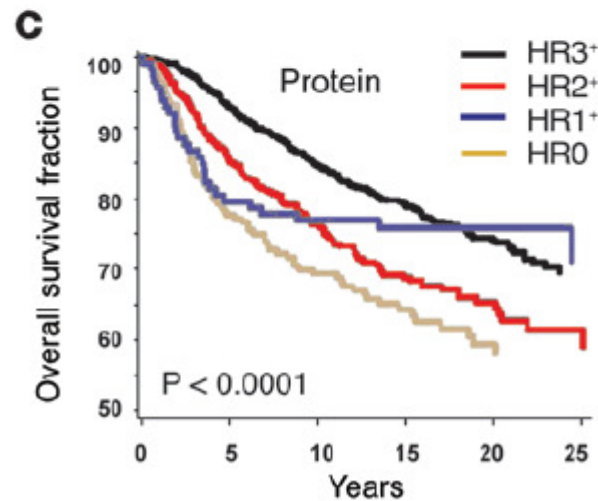
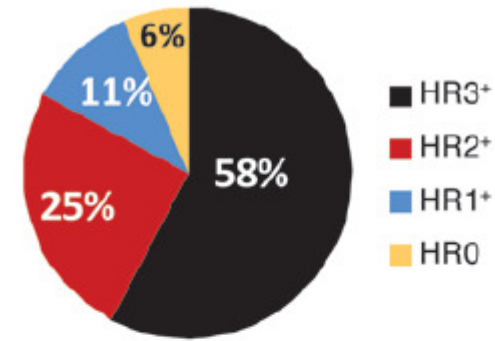
HR2 (ER+AR+,AR+VDR+ or ER+VDR+)

HR3 (ER+VDR+AR+)

A ER/PR/HER2 classification



B HRO–HR3 classification



Examination of 3,157 human breast tumors revealed that these HR subtypes were distinct from the current classification scheme, which is based on ER, PR and HER2. Patient outcomes were best when tumors expressed all three hormone receptors (HR3) and worst when they expressed none (HR0)

Balancing between classic morphology and molecular classification



- There will be no morphology versus molecular but personalized medicine is based on a combined morphological-molecular pathology report including classical morphology (HE/IHC/ISH) and diverse molecular analyses.

Where are we today (at least at our Institution)?

- ER, PR and HER2 status are the major drivers of clinical decision making regarding the type of systemic therapy.
- These 3 biomarkers in conjunction with histologic grade/mitotic count could be used to infer luminal, HER2 and TN subtypes .
- But given current options for systemic therapy, need to subclassify beyond ER,PR and HER2 in clinical practice is debatable.
- Clinicians are increasingly thinking about breast cancers by their molecular subtype.

