



SAN ANTONIO
BREAST CANCER
SYMPOSIUM®

Henry B. Gonzalez Convention Center,
San Antonio, Texas, USA

2017 DEC. 5-9

I have no financial relationship(s) with commercial interests to disclose.

Pathological Complete Response Predicts Event-Free and Distant Disease Free Survival in the I-SPY 2 TRIAL

Douglas Yee, MD

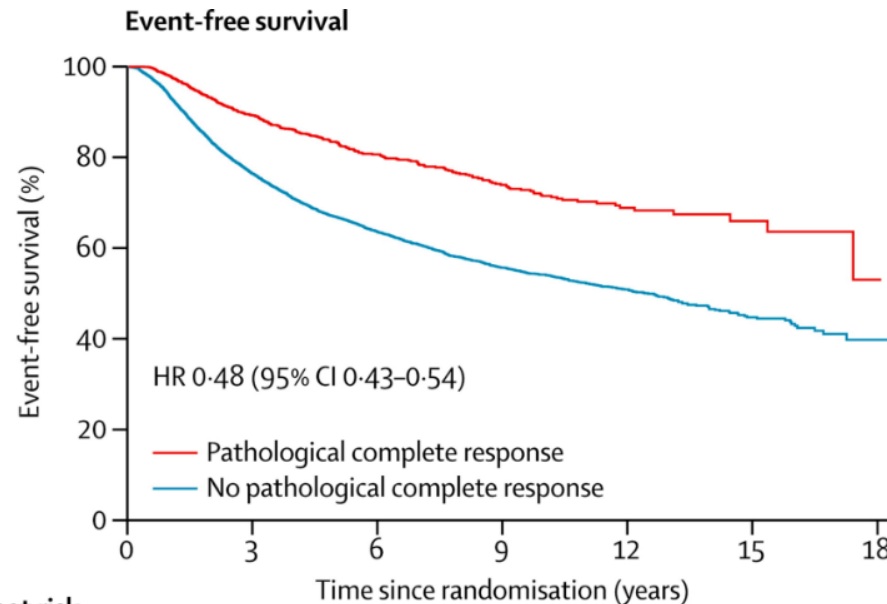
Masonic Cancer Center, University of Minnesota

On behalf of I-SPY2 Investigators and authors:

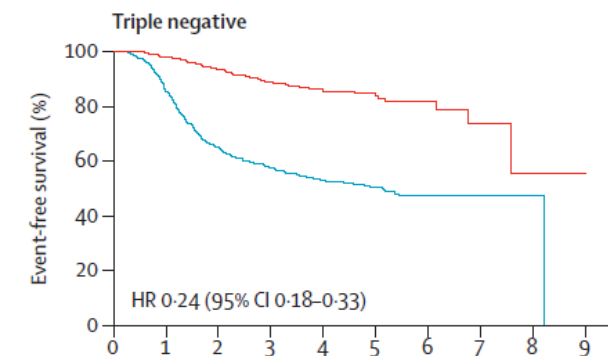
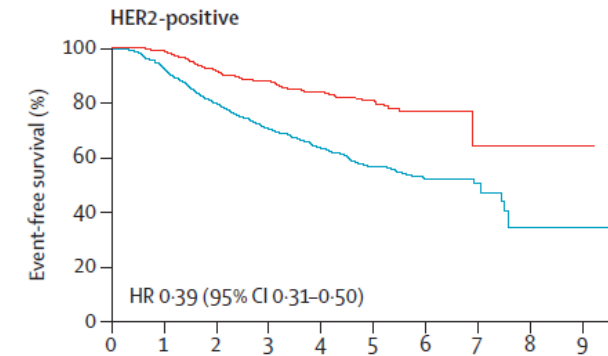
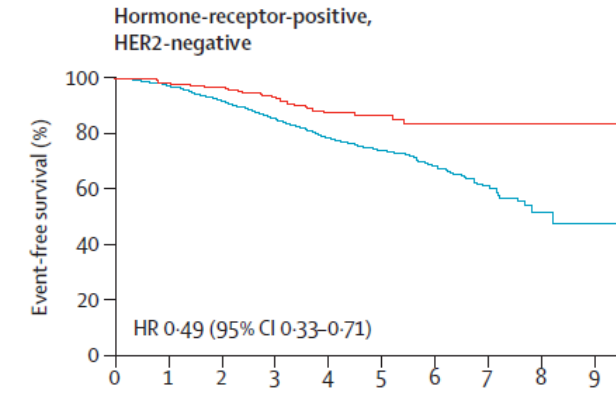
Yee D, DeMichele A, Isaacs C, Symmans F, Yau C, Albain KS, Hylton NM, Forero-Torres A, van't Veer LJ, Perlmutter J, Rugo HS, Melisko M, Chen Y-Y, Balassanian R, Krings G, Datnow B, Hasteh F, Tipps A, Weidner N, Zhang H, Tickman R, Thornton S, Ritter J, Amin K, Klein M, Chen B, Keeney G, Ocal T, Feldman M, Klipfel N, Sattar H, Mueller J, Gwin K, Baker G, Kallakury B, Zeck J, Duan X, Ersahin C, Gamez R, Troxell M, Mansoor A, Grasso LeBeau L, Sams S, Wisell J, Wei S, Harada S, Vinh T, Stamatakis MD, Tawfik O, Fan F, Adams A, Rendi M, Minton S, Magliocco A, Sahoo S, Fang Y, Hirst G, Singhrao R, Asare SM, Wallace AM, Chien AJ, Ellis ED, Han HS, Clark AS, Boughey JC, Elias AD, Nanda R, Korde L, Murthy R, Lang J, Northfelt D, Khan Q, Edmiston KK, Viscusi R, Haley B, Kemmer K, Zelnak A, Berry DA, Esserman LJ.

pCR and EFS

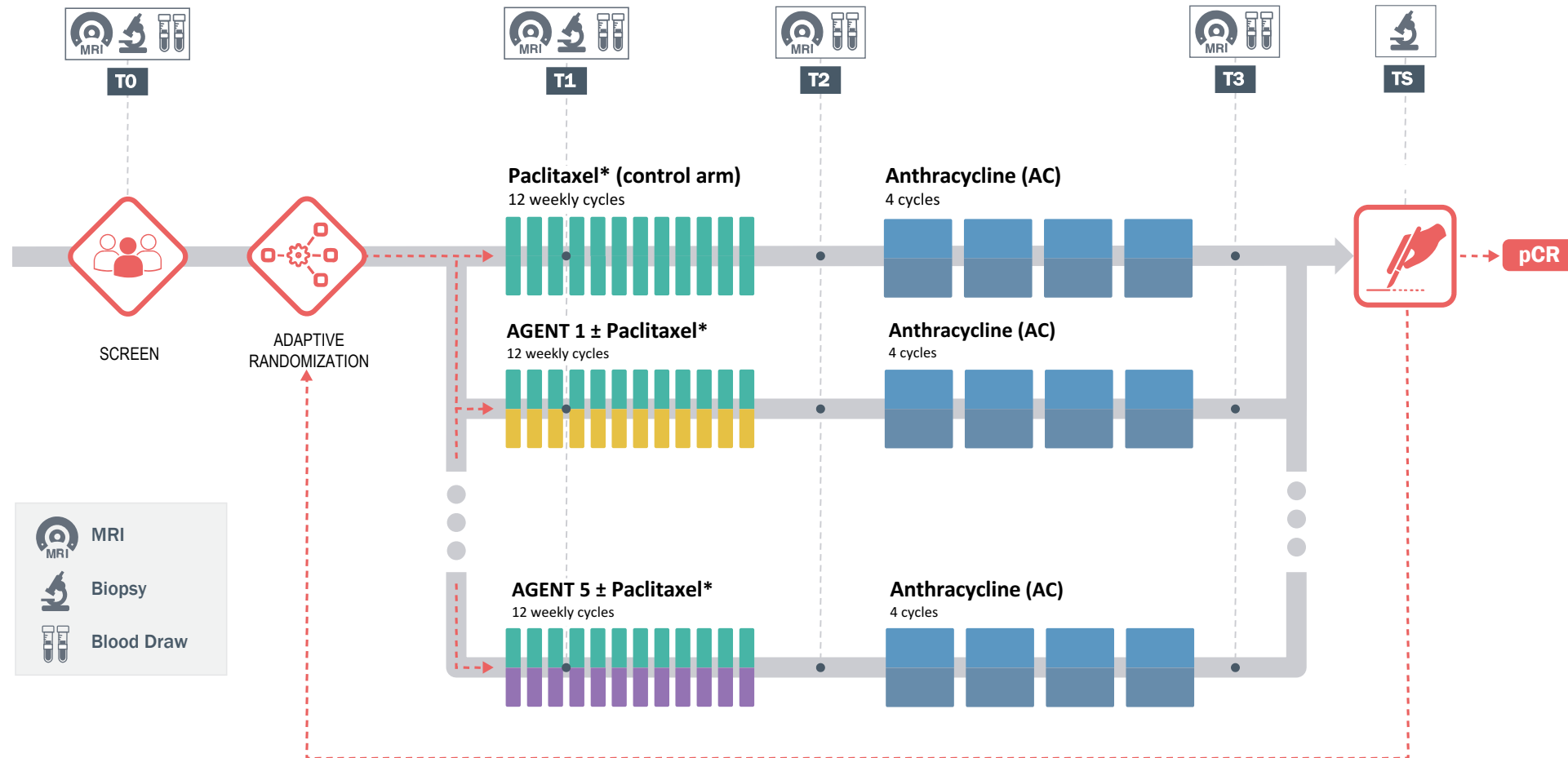
- FDA Meta Analysis (Cortazar et al, Lancet 2014)
 - >11K patients from 12 neoadjuvant trials
 - Median follow-up for EFS: 5.4 years



Number at risk							
Pathological complete response	2131	1513	583	337	124	35	2
No pathological complete response	9824	6169	2674	1523	525	165	1



Study Design

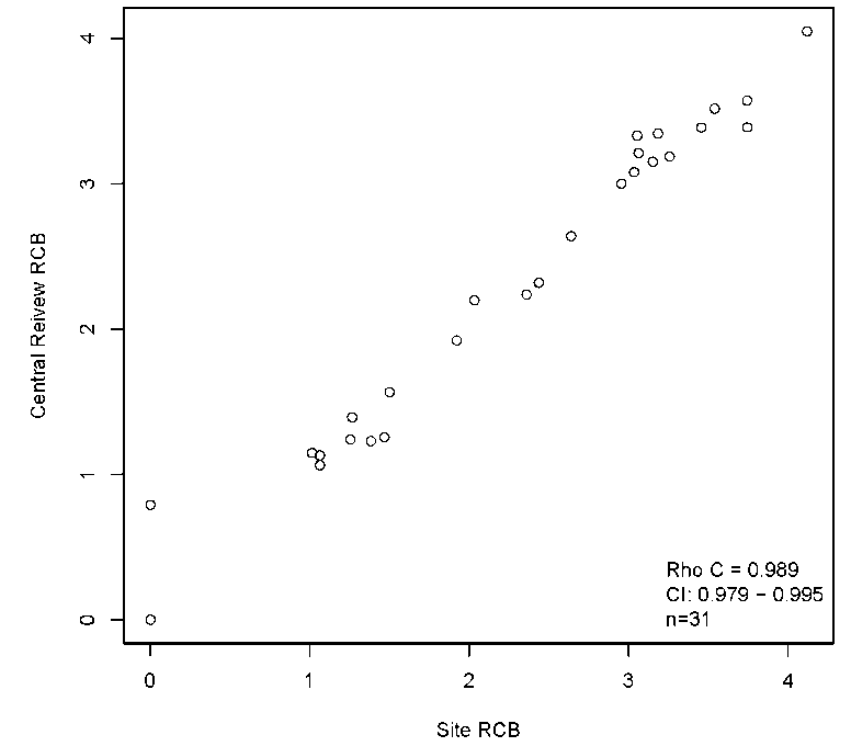


HR+/HER2- patients with low-risk MammaPrint Scores are not enrolled in I-SPY2

Analysis

- **Primary Endpoint:**
 - Pathological complete response (pCR)
 - Defined as no residual invasive cancer in breast or lymph nodes
 - Assessed using the Residual Cancer Burden (RCB) method*
 - Highly reproducible between local and central pathologist review
- **Intent-to-treat:**
 - Patients who did not complete assigned therapy are considered non-pCR (withdrew, left the institution, received non-protocol therapy, or progressed).
- **Secondary endpoints:**
 - RCB
 - EFS
- **I-SPY 2 To Date**
 - >1000 patients completed surgery
 - 11 investigational agents/combinations

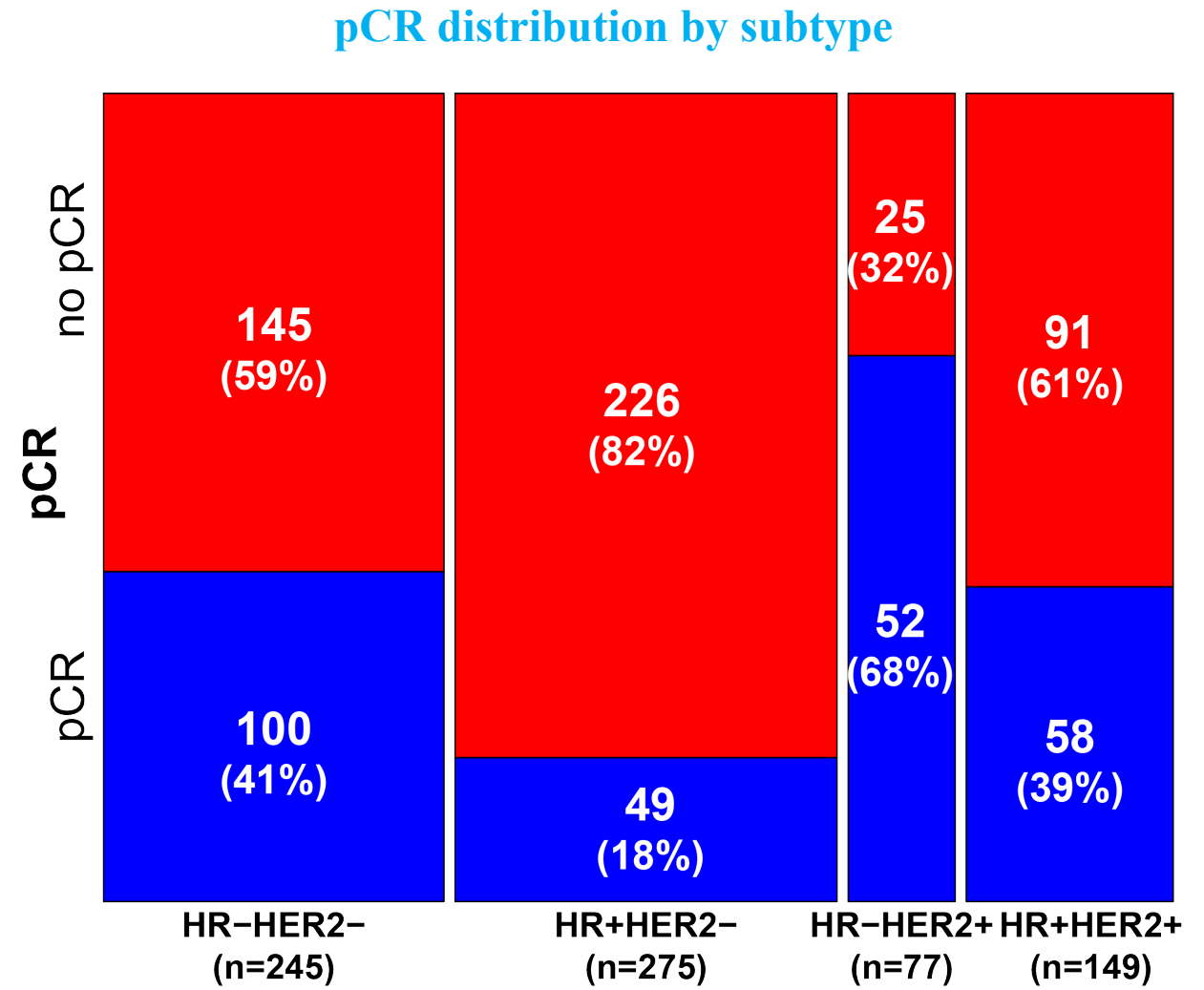
Scatterplot of RCB index entered by Site vs. Central Review



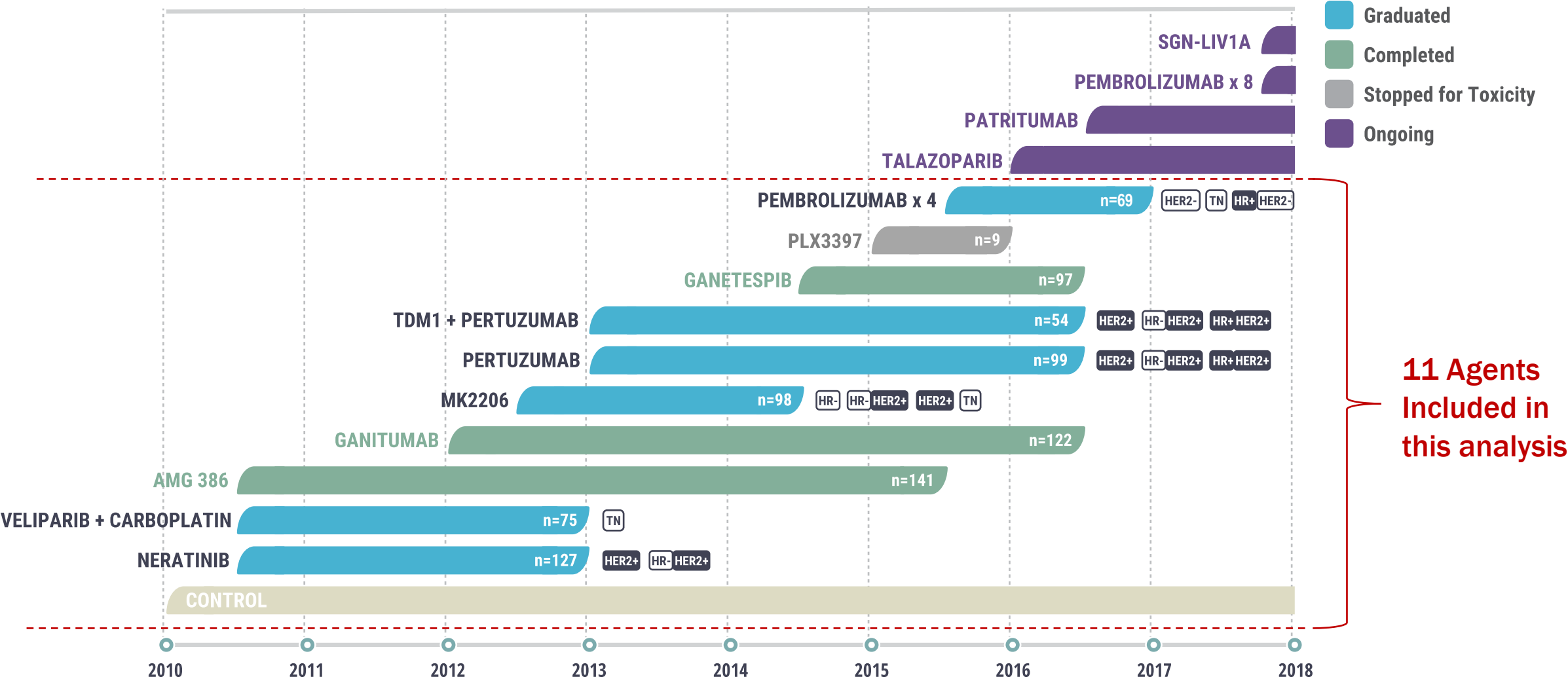
*Symmans, et al. J Clin Oncol 25:4414 2007 PMID: 17785706

EFS Dataset

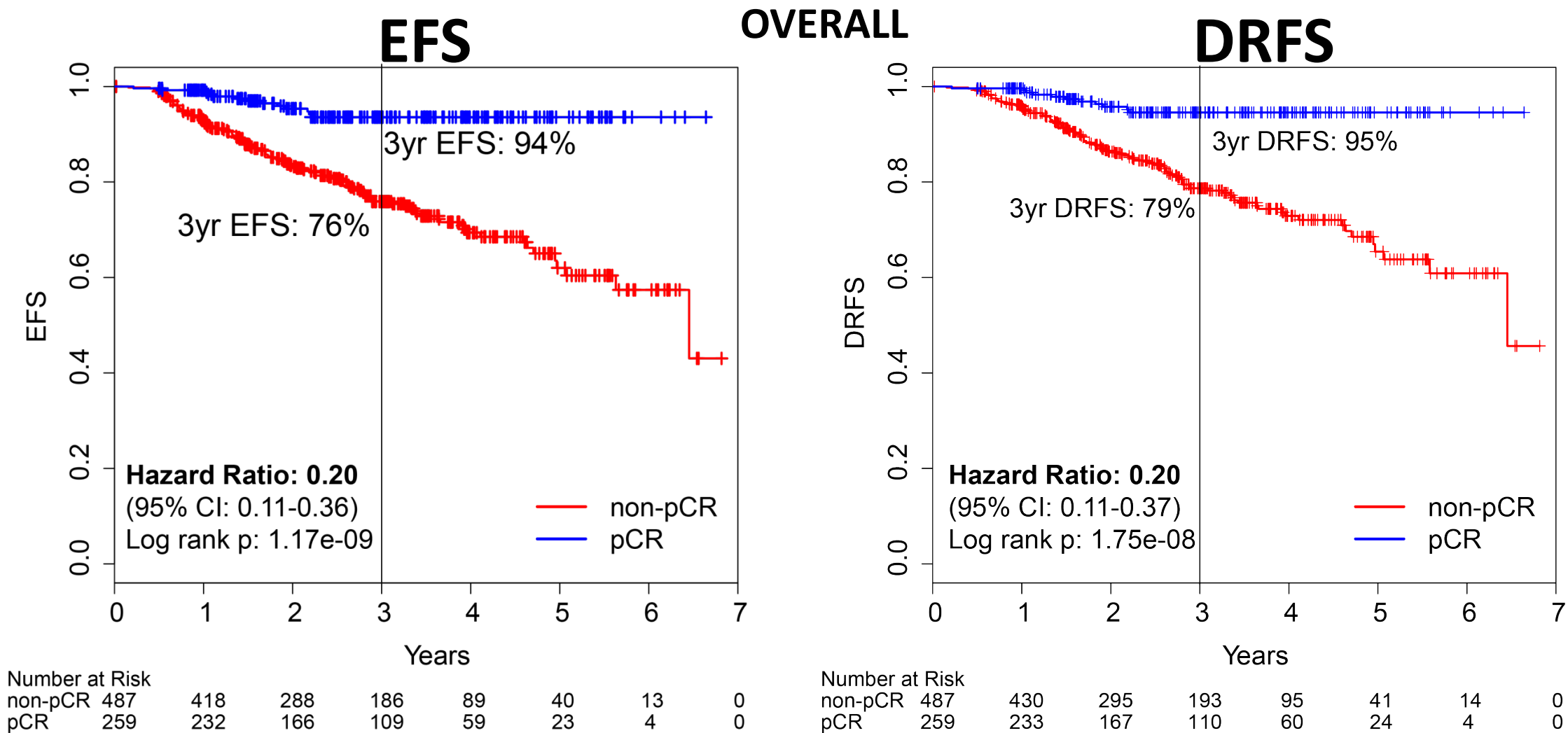
- Evaluable population: 746
 - 259 (35%) pCR, 487 (65%) non-pCR
 - Median follow-up: 2.7 yrs (0.02-7.2)
 - 126 EFS events, 109 DRFS events
- 12 patients did not go to surgery
 - considered non-pCR per protocol



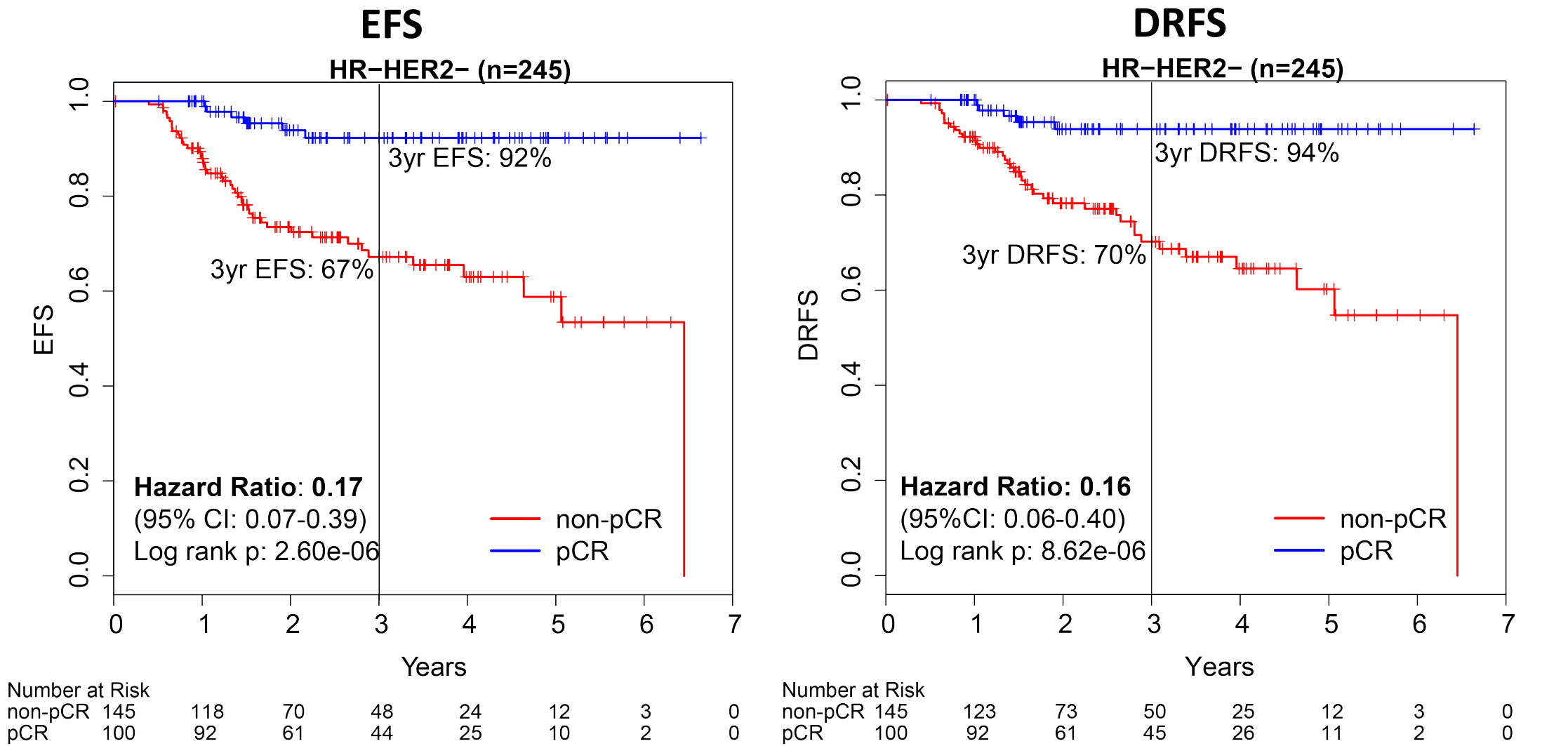
Agent Timeline



pCR is a highly significant predictor of EFS and DRFS

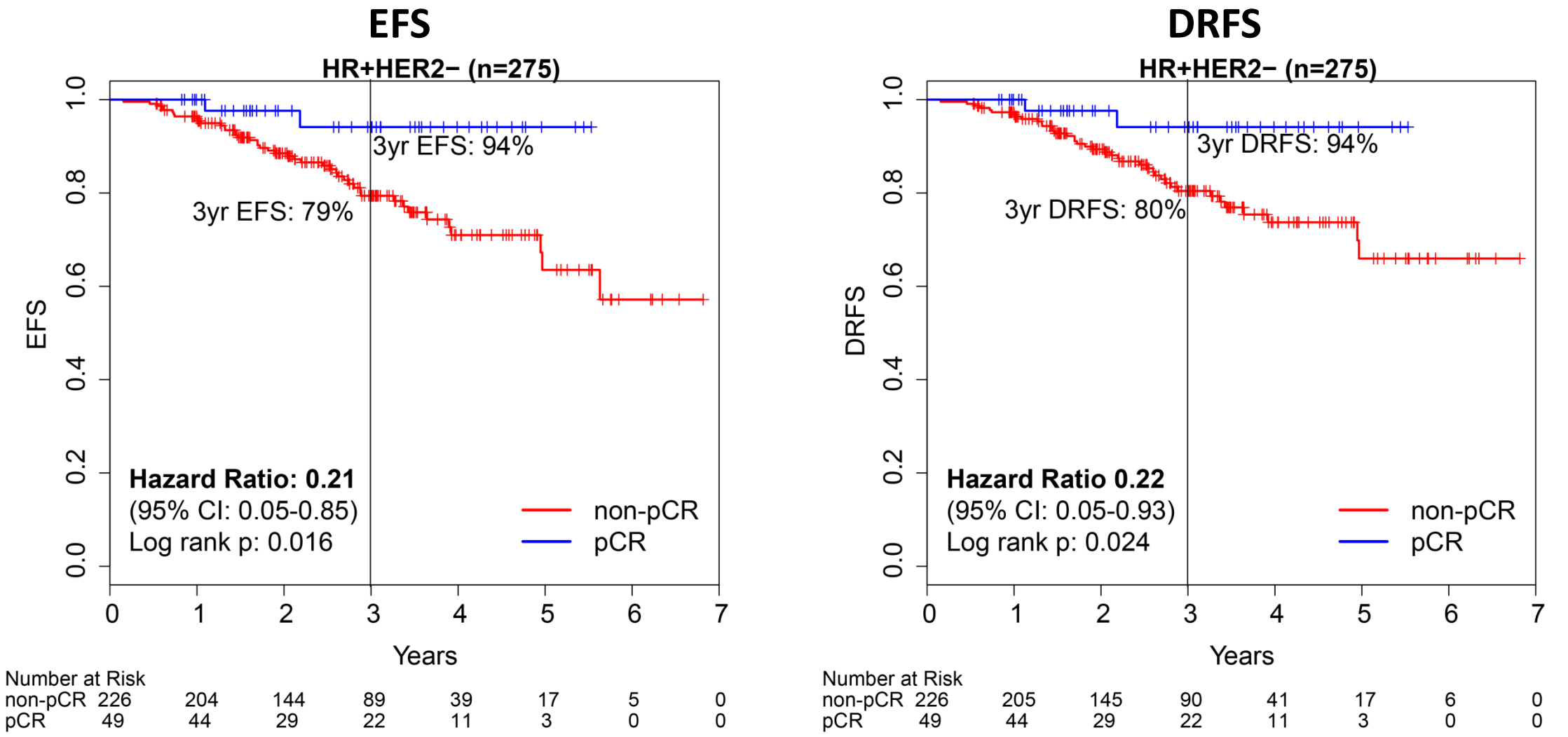


pCR is predictive of EFS and DRFS in TNBC

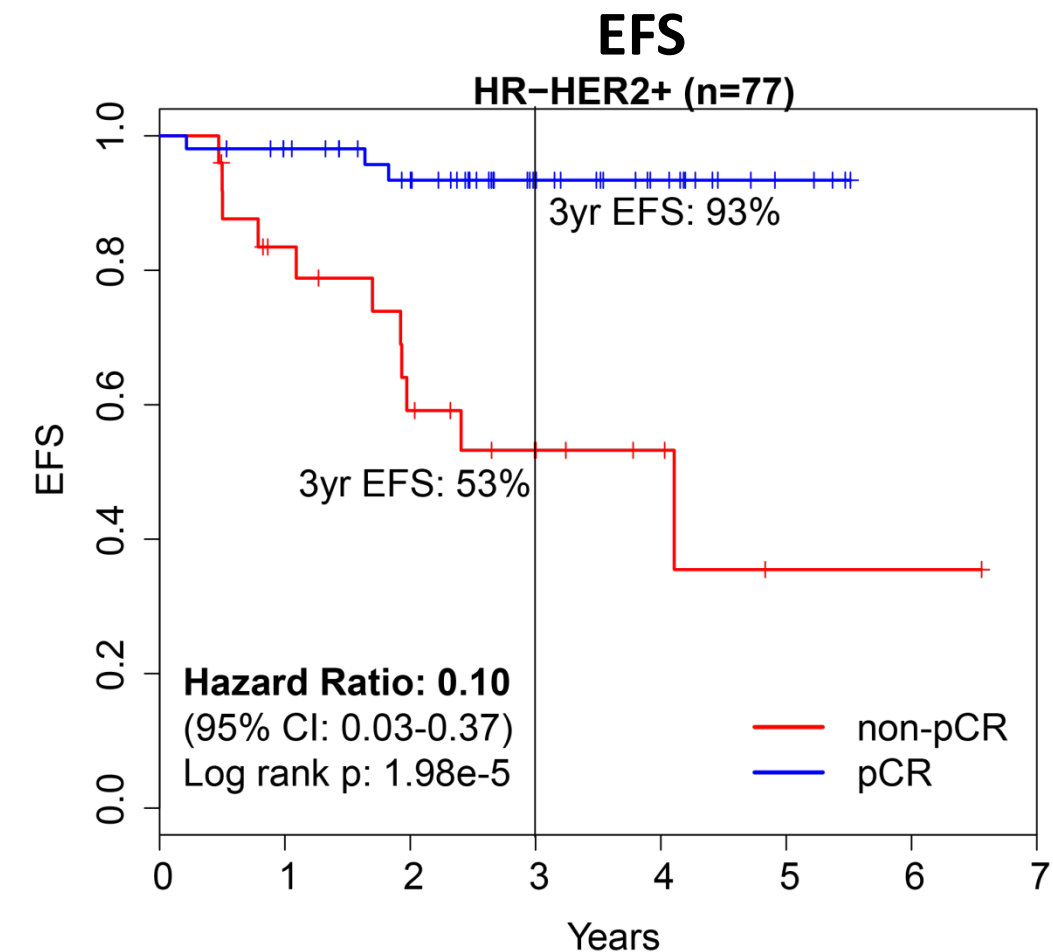


This presentation is the intellectual property of Douglas Yee.
Contact yeex006@umn.edu for permission to reprint and/or distribute.

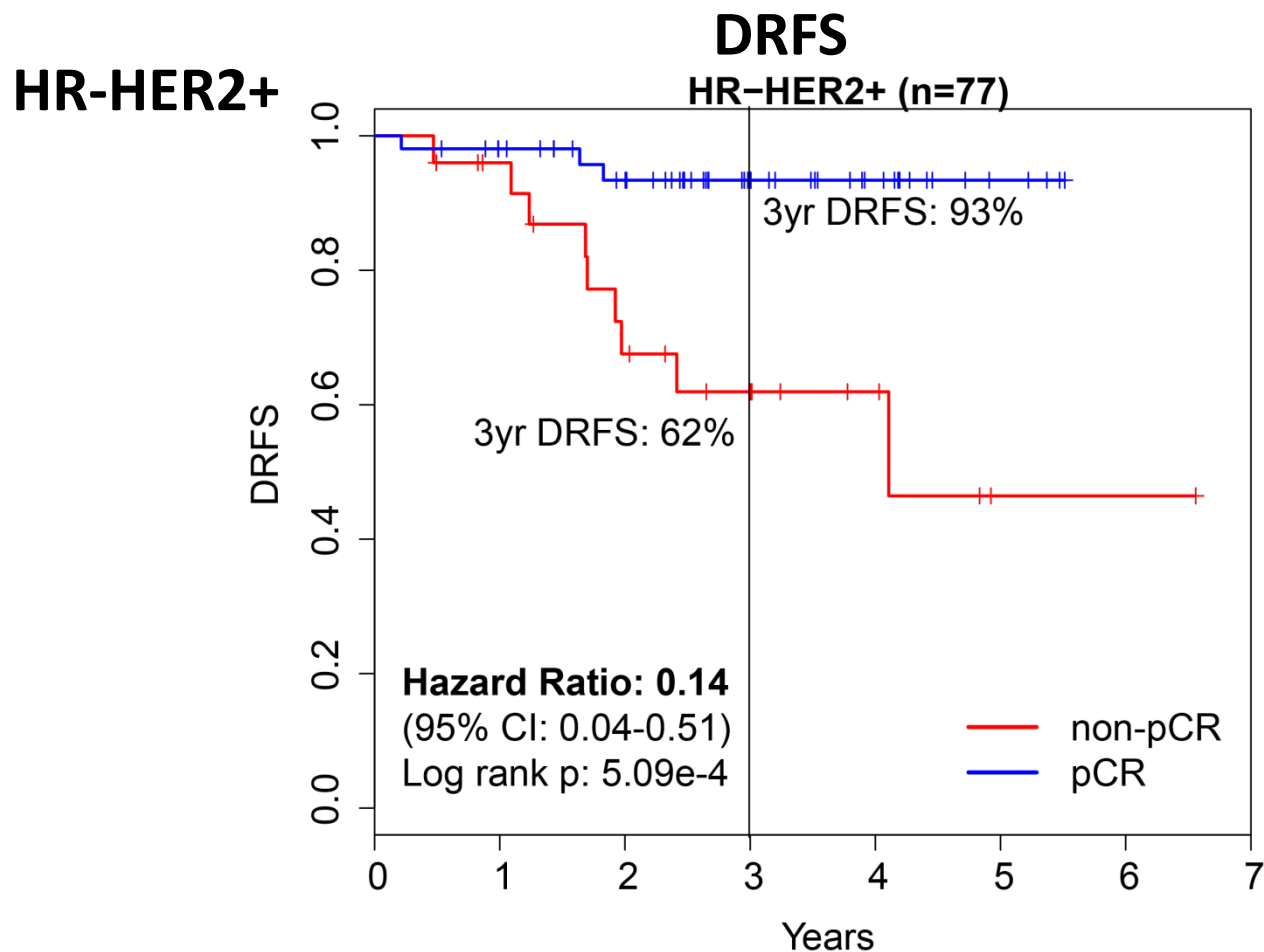
pCR is predictive of EFS and DRFS in HR+/HER2-



pCR is predictive of EFS and DRFS in HR-/HER2+

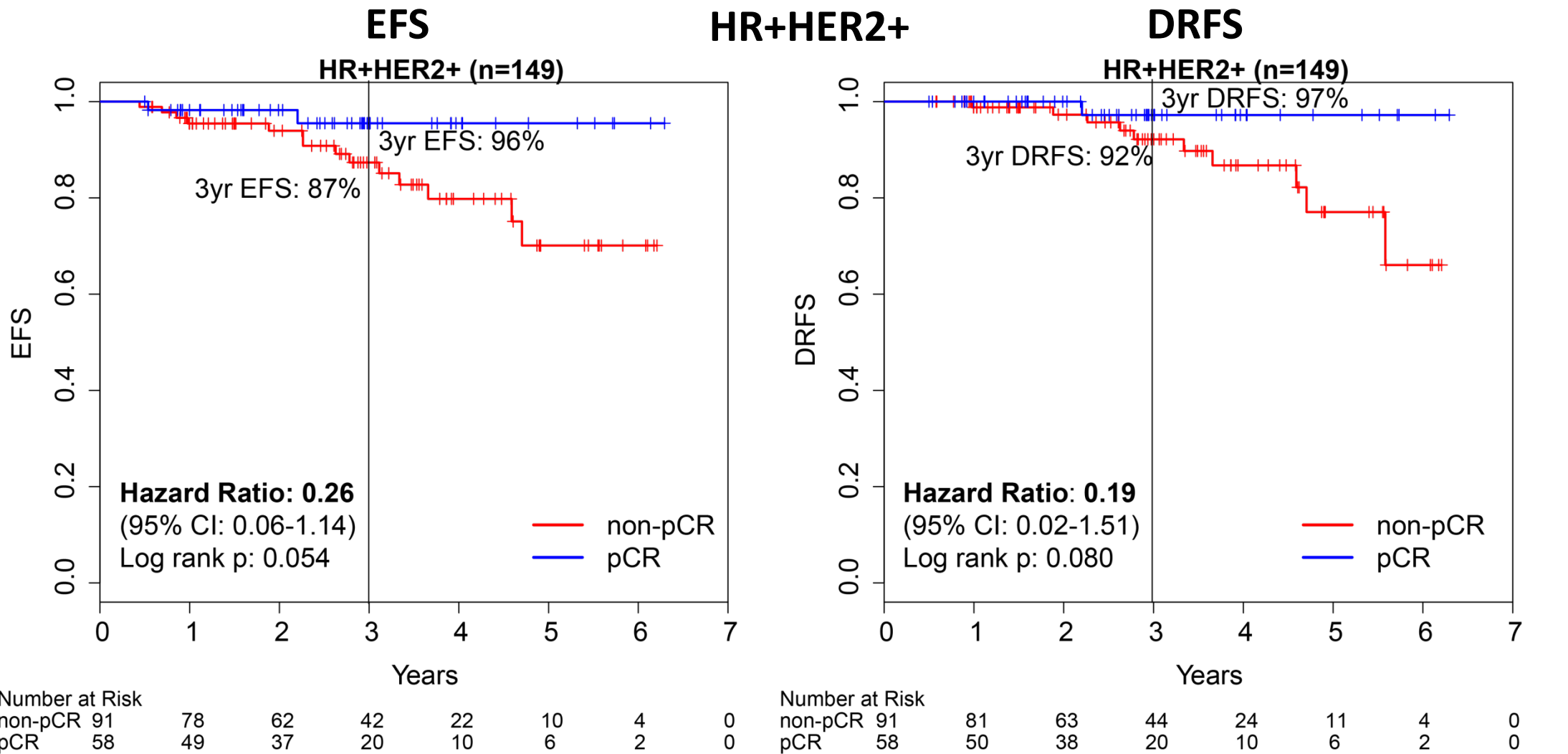


Number at Risk								
non-pCR	25	18	12	7	4	1	1	0
pCR	52	47	39	23	13	4	0	0

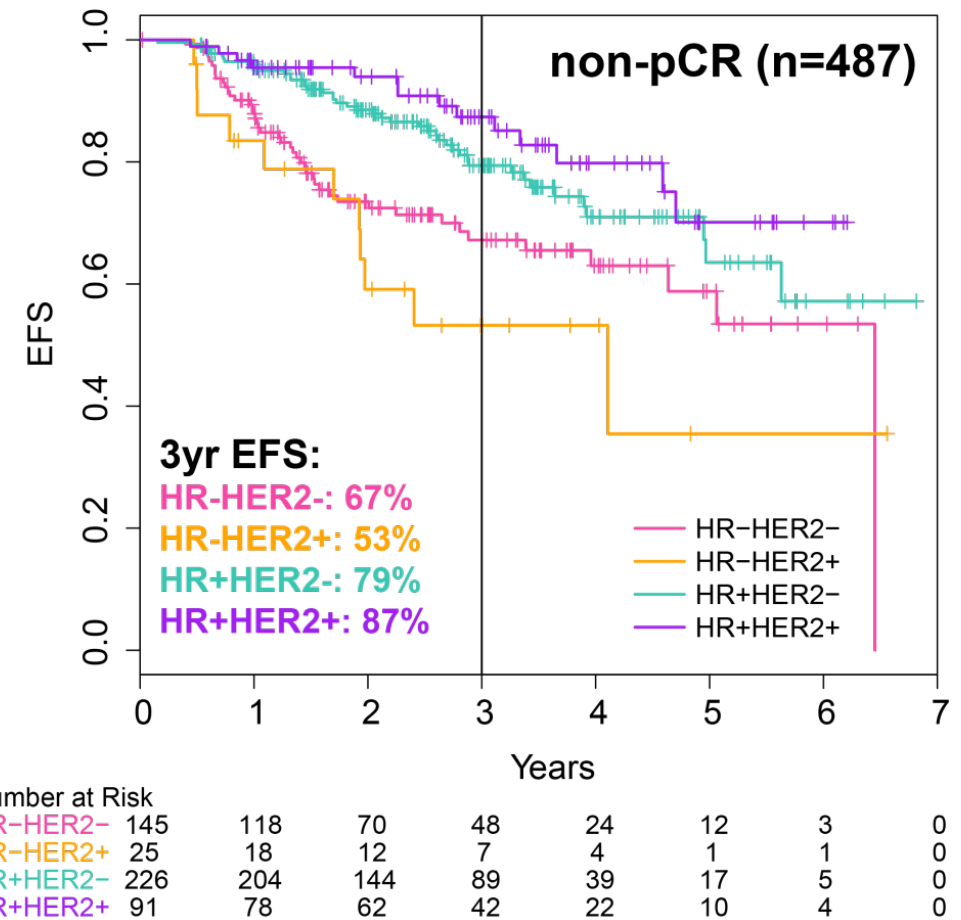
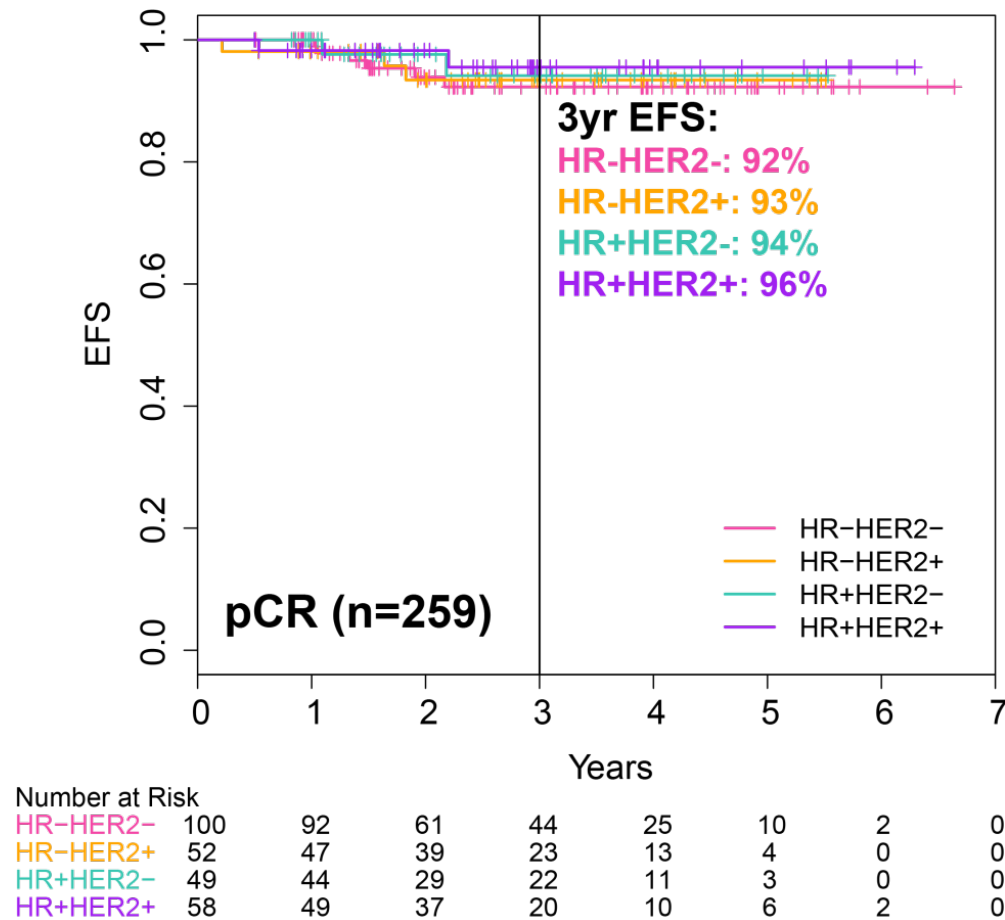


Number at Risk								
non-pCR	25	21	14	9	5	1	1	0
pCR	52	47	39	23	13	4	0	0

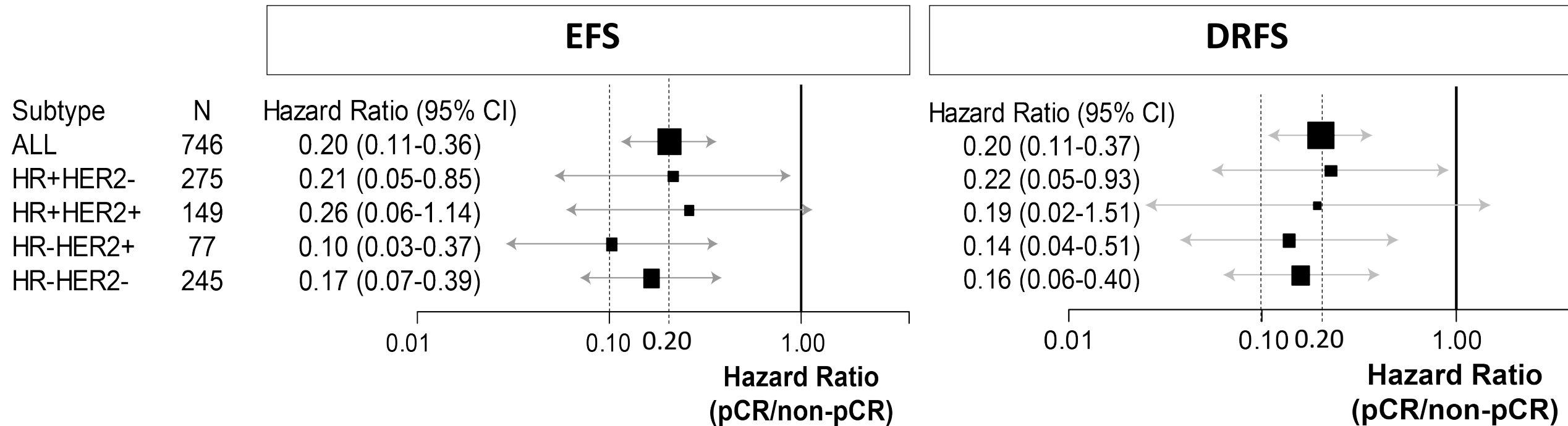
pCR is predictive of EFS and DRFS in HR+/HER2+



EFS by pCR & non-pCR by Subtype



EFS and DRFS Hazard Ratio for pCR vs non-pCR



I-SPY2 EFS Hazard Ratio for pCR/non-pCR compared to FDA meta-analysis and cooperative group results

	I-SPY 2	Cortazar Meta-analysis	Cooperative Group CALGB 40603
Overall	0.20 (0.11-0.36)	0.48 (0.43-0.54)	
*HR+HER2-	0.21 (0.05-0.85)	0.49 (0.33-0.71)	
HER2+	0.21 (0.08-0.55)	0.39 (0.31-0.50)	
HR-HER2-	0.17 (0.07-0.39)	0.24 (0.18-0.33)	0.30 (0.19-0.45)

*Mammaprint low patients excluded

Summary

- pCR is a strong predictor of EFS and DRFS in the setting of a multiple agent platform trial that includes:
 - Standards for eligibility
 - *high risk for early recurrence (MP low risk, HR+Her2- excluded)*
 - *exclusion of metastatic disease*
 - All chemotherapy given before pCR determination
 - Standards for pathology assessment and multidisciplinary identification (surgeons, radiologists, pathologists)
 - Long term follow-up of patients over time (correlation of early, intermediate, and late endpoints)
- pCR is equally predictive across all tumor subsets
- pCR as an endpoint enables rapid evaluation of novel therapy combinations and can accelerate the identification of effective and potentially less toxic regimens

The Future of I-SPY 2

- Achieving pCR through any therapy for any subtype is a sufficient endpoint
- Develop minimally invasive techniques (MRI and biopsy) to identify pCR prior to definitive surgery
 - Validate robust MRI and tissue predictors of pCR
 - Deescalate toxic therapy (AC) if pCR obtained early
- Re-assign patients to new therapies if pCR is not predicted
 - Validate robust MRI and tissue predictors of non-PCR
 - Assign new therapies based on molecular profiling of tumor and link to investigational agents

Acknowledgements

WORKING GROUP CHAIRS

PI:	Laura Esserman	Operations:	Angie DeMichele
Co-PI:	Don Berry	Biomarkers:	Laura van 't Veer
Imaging:	Nola Hylton	Pathology:	Fraser Symmans
Agents:	Doug Yee	Advocates:	Jane Perlmutter
Safety:	Hope Rugo	PRO/QOL:	Michelle Melisko

SITE PRINCIPAL INVESTIGATORS

Columbia:	Kevin Kalinsky	UAB:	Andres Forero-Torres
Denver:	Anthony Elias	UChi:	Rita Nanda
Gtown:	Claudine Isaacs	UCSD:	Anne Wallace
Loyola:	Kathy Albain	UCSF:	Jo Chien
Mayo:	Judy Boughey	UMinn:	Doug Yee
Moffitt:	Heather Han	UPenn:	Amy Clark
OHSU:	Kathleen Kemmer	USC:	Julie Lang
Swedish:	Erin Ellis	Yale:	Tara Sanft

SPONSOR

Quantum Leap Healthcare Collaborative

Dave Mandelkern, Nancy Lisser, Mike Bankert, Adam Asare, Smita Asare, Kristen Zeitzer

PROJECT OVERSIGHT

Anna Barker/ASU, Gary Kelloff/NCI, Janet Woodcock/FDA, Richard Pazdur/FDA, Robert Becker/FDA, ShaAvhree Buckman/FDA,CDER, Steve Gutman, David Wholley/FNIH

PROGRAM MANAGEMENT OFFICE

Executive Director:

Smita Asare

Program Administration:

Kat Steeg, Lorena Kanu, Julie LeDuc, Jill Parker, Melanie Hanson

Safety:

Sausan Abouharb, Linda Doody, Monina Angeles, CCSA

Data Analysis & IT:

Christina Yau, Adam Asare, Garry Peterson, Amy Wilson, Tim Fu

Operations Manager:

Ruby Singhrao

Biomarkers/Specimens:

Lamorna Brown-Swigart, Gillian Hirst, Denise Wolf, Chip Petricoin, Julie Wulfkuhle

Imaging Lab:

Jessica Gibbs, Melanie Regan

Business Development:

Julie Sudduth-Klinger, Dan Dornbusch

Manuscripts/Strategy:

Jeff Matthews

PRIOR COLLABORATORS and STAFF

Larissa Korde, Rashmi Murthy, Donald Northfelt, Qamar Khan, Kirsten Edmiston, Rebecca Viscusi, Barbara Haley, Amelia Zelnak, Meredith Buxton, Melissa Paolini, Julia Lyanderes,

Thank you to the remarkable patients and families, our amazing advocates,
all of the investigators, staff, and our DSMB for supporting the trial

Participating Organizations

FUNDING PARTNERS

William K Bowes, Jr. Foundation
Give Breast Cancer the Boot
University of California San Francisco (UCSF)
The Biomarkers Consortium

Quintiles
The Breast Cancer Research Foundation
Safeway, an Albertsons Company

INVESTIGATIONAL AGENT PROVIDERS

Seattle Genetics
AstraZeneca
Daiichi-Sankyo
Merck
Pfizer
Puma Biotechnology
AbbVie

Synta Pharmaceuticals
Genentech
Amgen
Plexxikon

STUDY SPONSOR

Quantum Leap Health Care Collaborative

BIOMARKER PLATFORMS & DATA SUPPORT

Berry and Associates
CCS Associates
salesforce
Agendia
Natera

Hologic
Novella Clinical
Oregon Health & Science University (OHSU)
UCSF
The Translational Genomics Research Institute (TGen)