



# I SPY 2: The right drug, the right patient, the right time

*Using Biology to Adaptively Guide Treatment for Early Breast Cancer and Predict Response*

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I-SPY 2 co-Principal Investigator and Biomarker Chair

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I-SPY 2 TRIAL PI's:

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## Disclosure

- Co-founder & stockholder Agendia BV
- No other disclosures

## Basic Principles of I-SPY

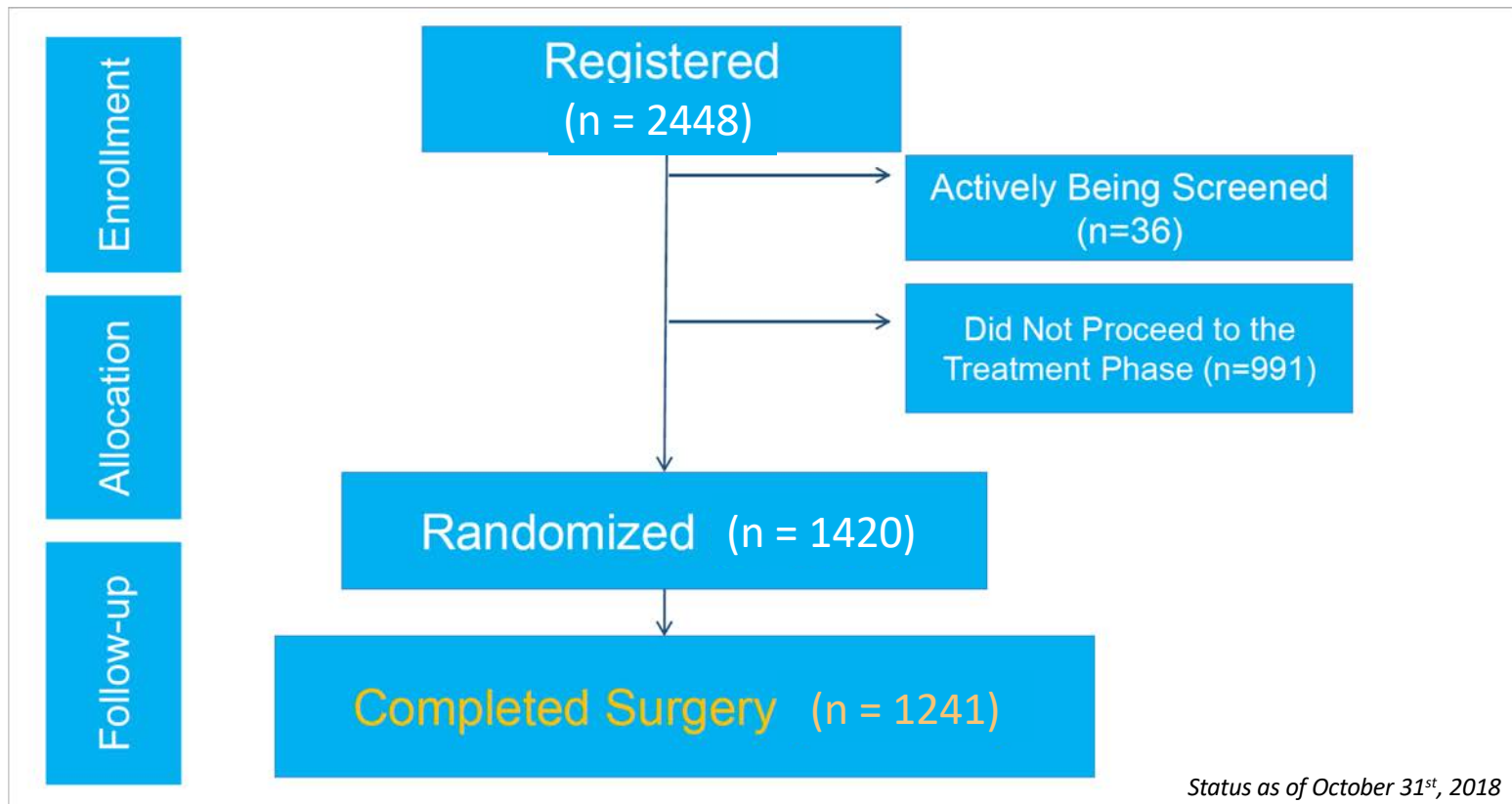
- Test new drugs where they matter most
  - **Early stage** rather than metastatic disease
- Change the order of therapy: learn about response early in the course of care
  - Neoadjuvant setting (**systemic therapy before surgery**)
  - Primary Endpoint is complete response to therapy (**pCR**)
- Build an efficient engine to evaluate drugs, accelerate knowledge turns
  - Master Protocol, Adaptive Design
- Use imaging and biomarker guidance
  - Focus on the population of patients who are at high risk for EARLY recurrence
  - Insights about who responds to what agents
  - “Graduation” for efficacy = threshold predictive probability of success in next phase III trial
- Collaborative by Design:
  - FDA, IRBs, Pharma, Biotech, Academics, Community Cancer Ctrs, Advocates

# I-SPY 2 Participating Sites

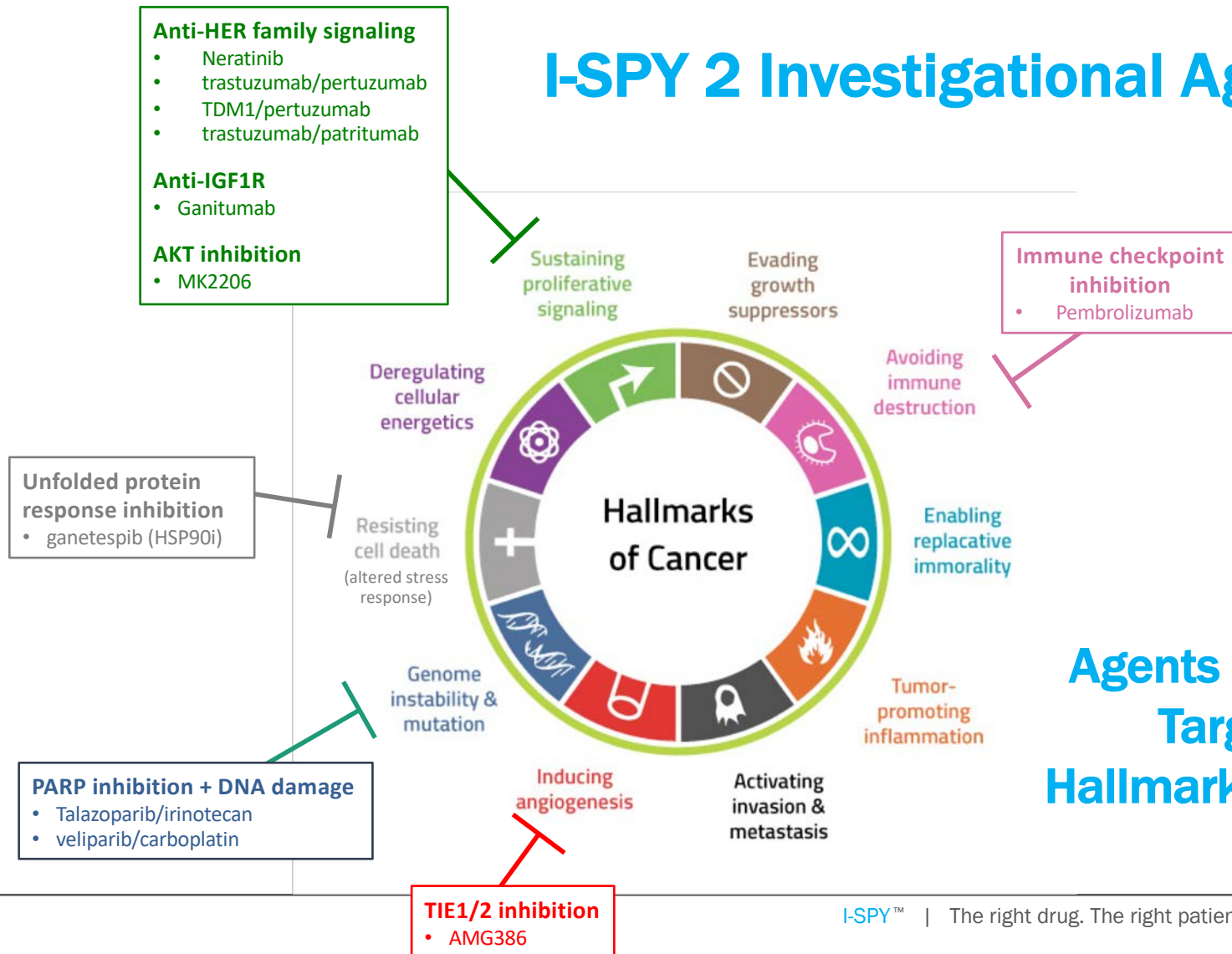
16 Sites Open and Enrolling  
+ 3 Opening in Q3/Q4 2018



# Trial Patient Enrollment Overview

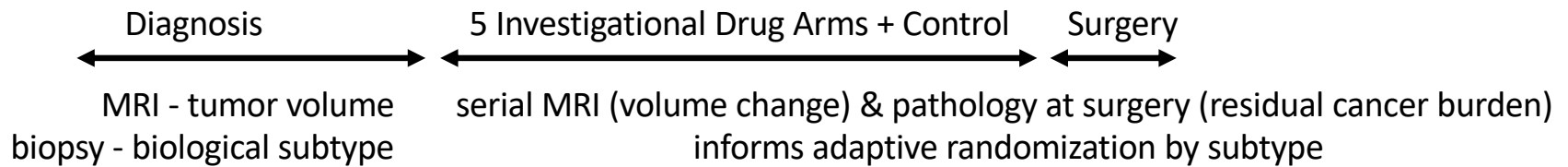


# I-SPY 2 Investigational Agents



**Agents Developed Targeting Hallmarks of Cancer**

# I-SPY 2 Framework for Early High Risk Breast Cancer: Biomarkers Guide Enrichment of Neoadjuvant Drug Arm with Responding Subtype

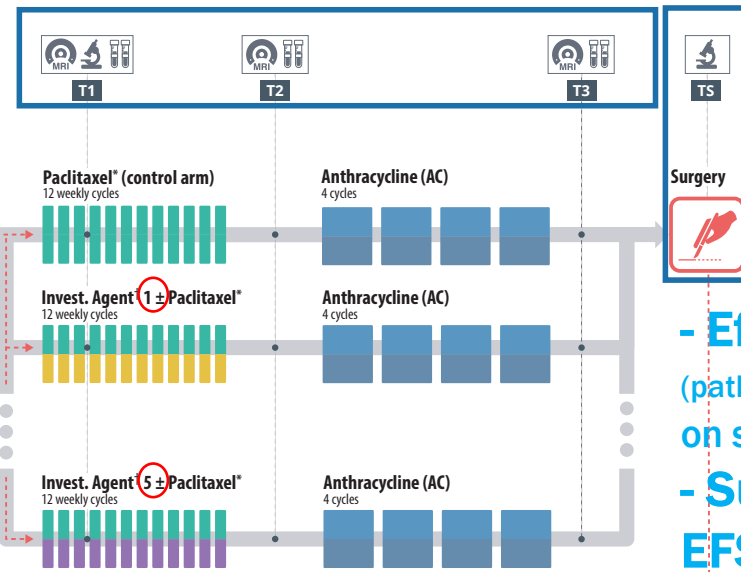
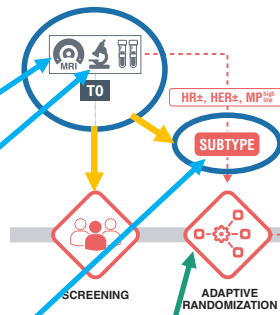
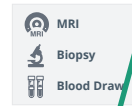


## Biomarkers:

- Imaging
- Pathology
- Molecular Biology

### 8 subtypes by:

- Hormone Receptor +/-
- HER2 +/-
- MammaPrint High1/High2 (HR+/MammaPrint low excluded)



- Efficacy endpoint: pCR (pathological Complete Response) on surgery specimen
- Survival endpoint: EFS/DRFS at 3y & 5y (Event/Distant Recurrence Free survival)

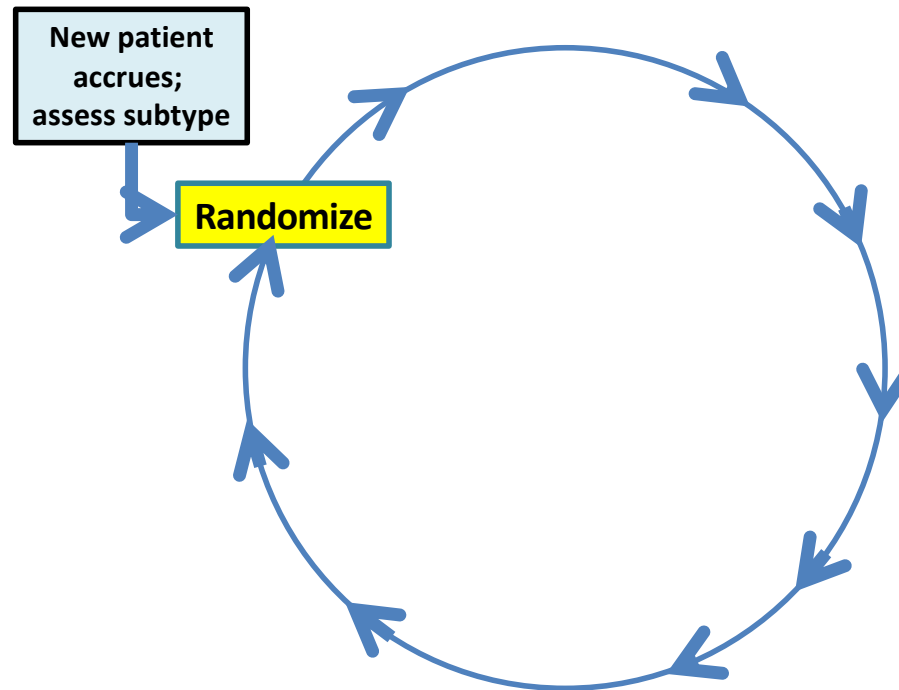
\* Patients who are HER2+ may also receive trastuzumab (Herceptin)

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

Adaptive Randomization

I-SPY™ | The right drug. The right patient. The right time.™

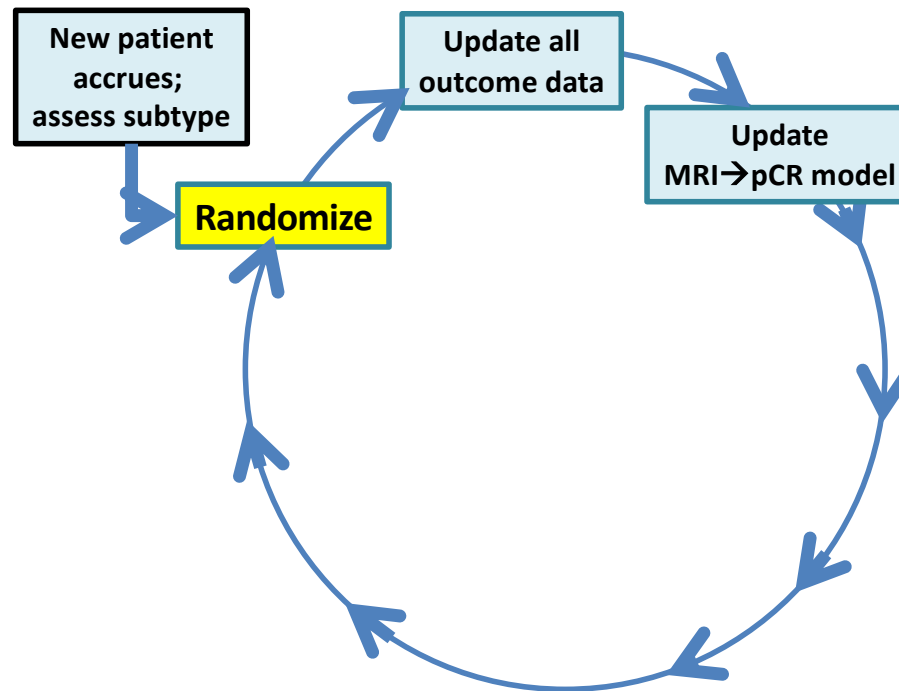
# I-SPY 2 Adaptive Randomization



Adaptive randomization based on 8 subtypes  
(hormone receptor (HR) +/-, HER2 +/-, MammaPrint-High 1 or High 2;  $2^3=8$ )



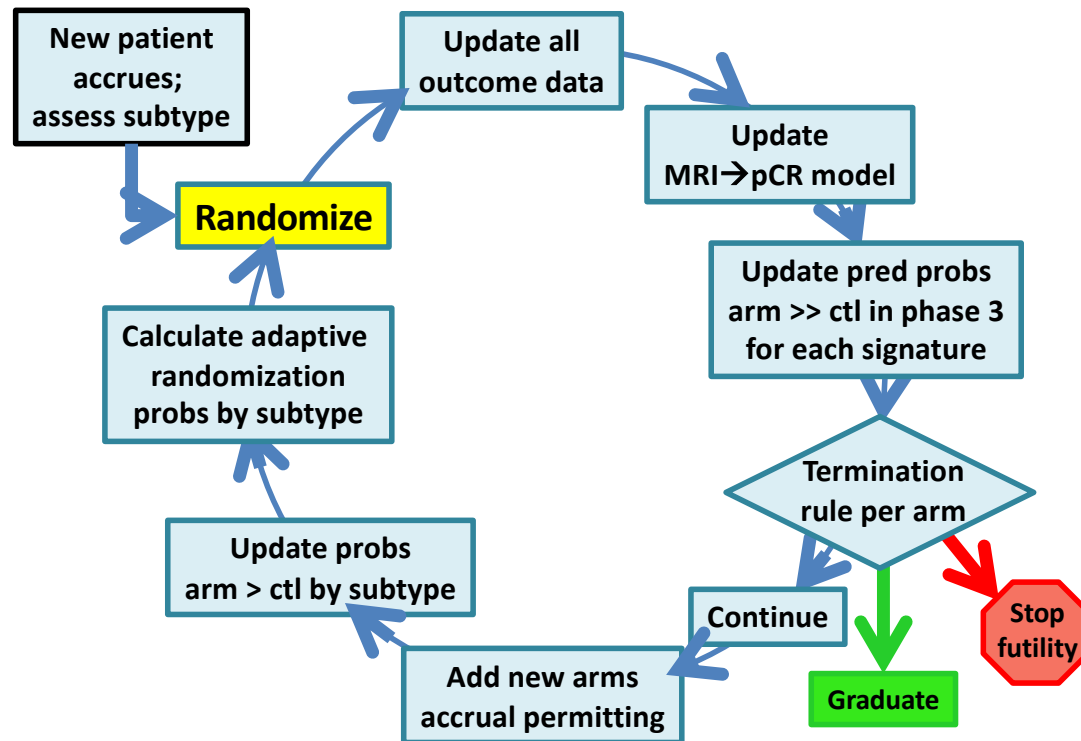
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# I-SPY 2 Adaptive Randomization

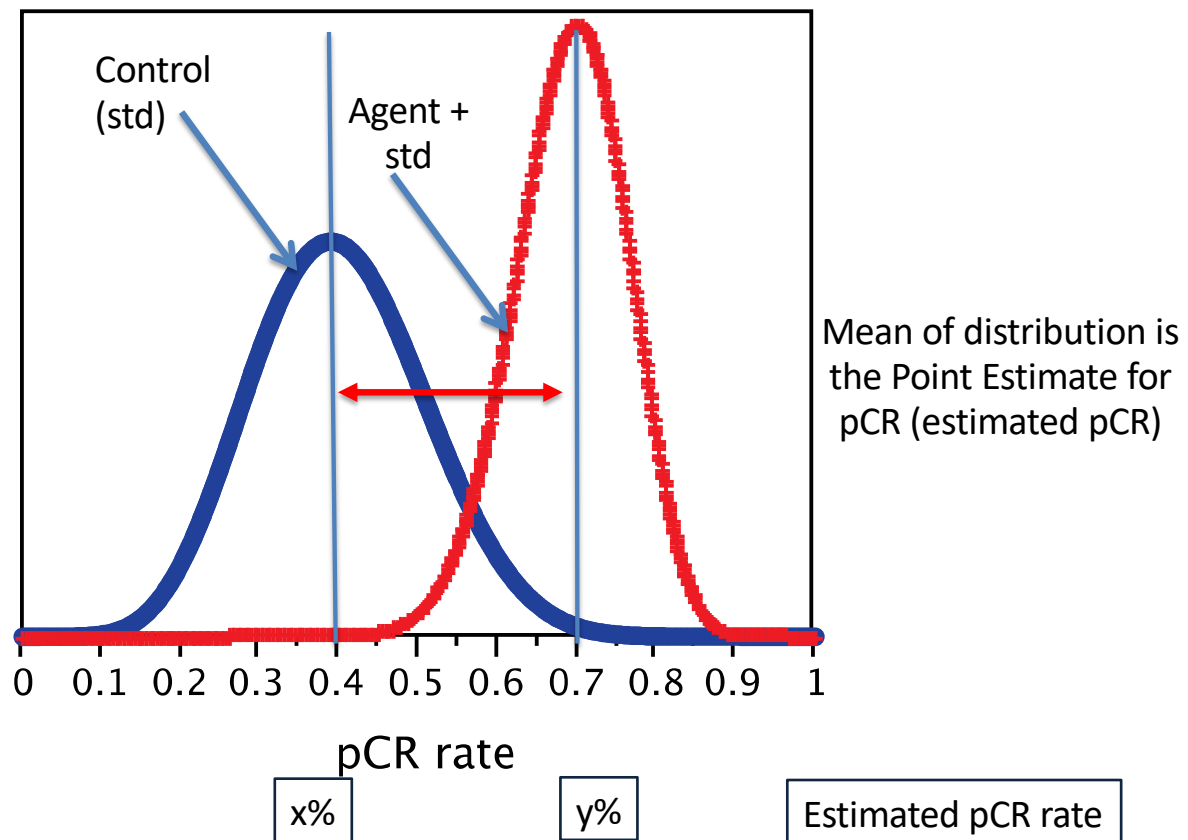
Adaptive randomization based on 8 subtypes



Graduation based on 10 signatures (combinations of subtypes)

# Methods for Estimating Response Probability (pathological Complete Response = pCR)

Distribution of pCR rates



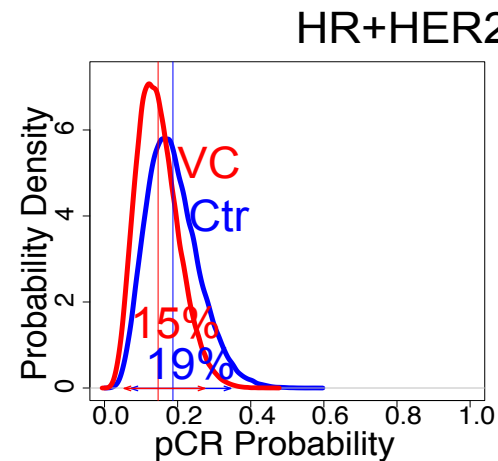
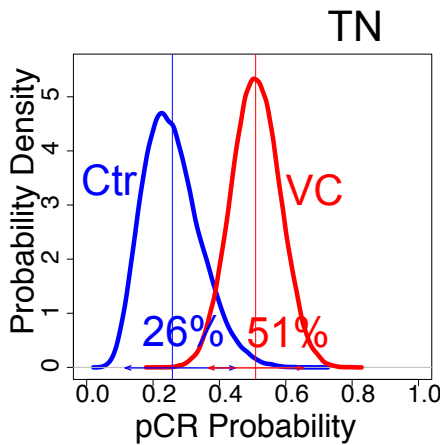
# I-SPY 2 Framework:

## Biomarkers Guide Enrichment of Drug Arm with Responding Subtype

Example: Veliparib (PARP-inhibitor)/Carboplatin

Biomarkers indicated while arm was ongoing:

- response in Triple-Negative (TN) Breast Cancer > 'graduation'
  - no response in Hormone receptor positive Breast cancer (HR+/HER2-)
- and the adaptive randomization enriched the VC arm with TN Breast Cancer



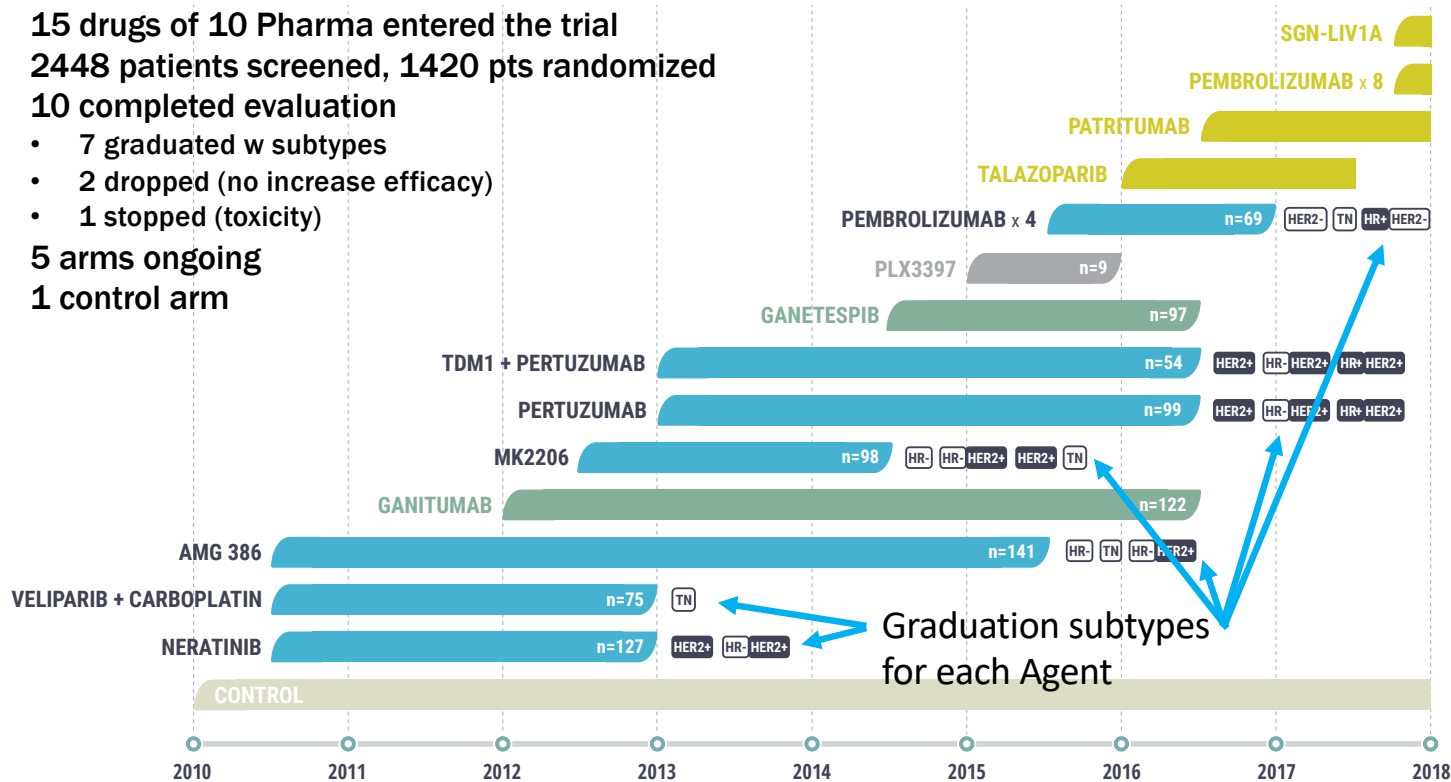
# Timeline of Investigational Drugs and Graduating Subtypes

## Biomarkers Guide Enrichment of Drug Arm with Responding Subtype

15 drugs of 10 Pharma entered the trial  
 2448 patients screened, 1420 pts randomized  
 10 completed evaluation

- 7 graduated w subtypes
- 2 dropped (no increase efficacy)
- 1 stopped (toxicity)

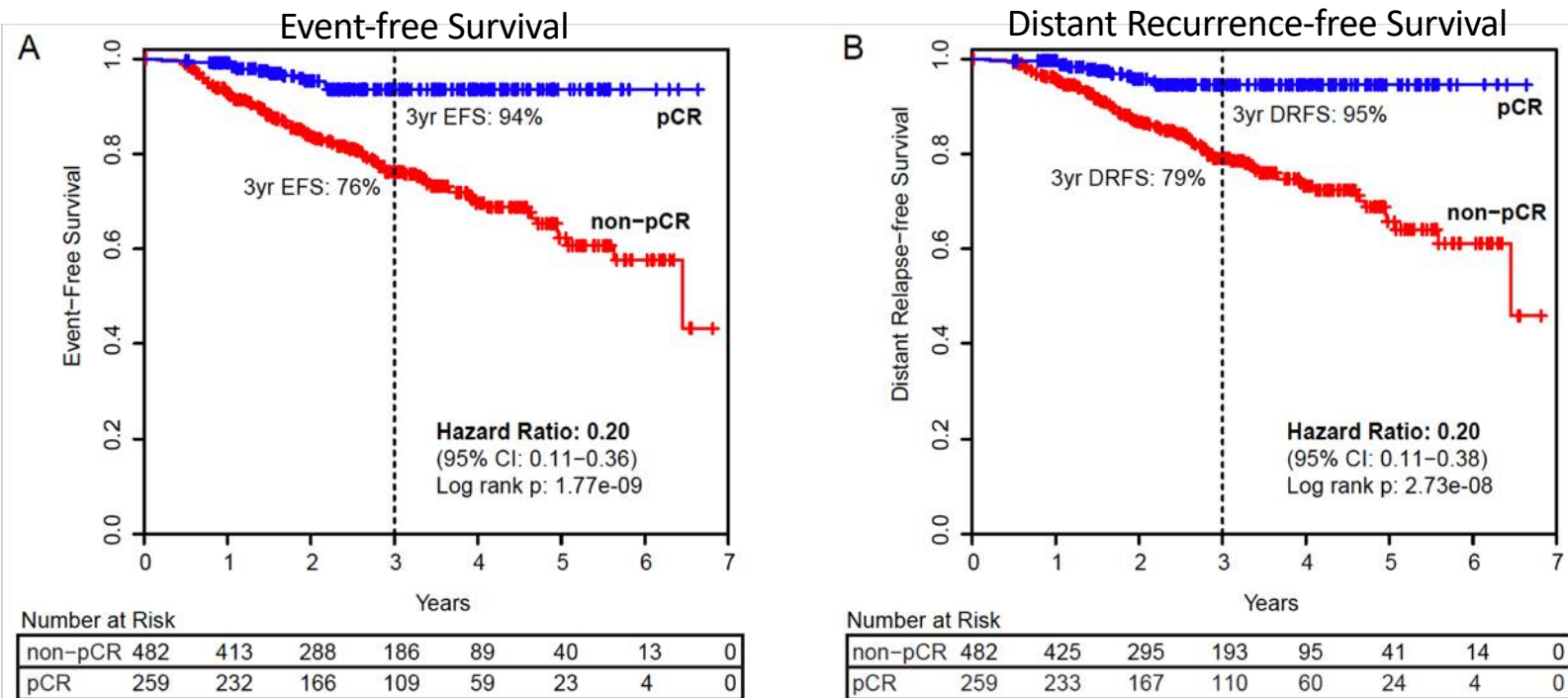
5 arms ongoing  
 1 control arm



- Efficacy endpoint:  
 Seven graduating drugs  
 Response rate at least  
 doubled vs. control treatment  
 (20% > 40% pCR), some  
 drug/subtype 65% pCR

# pCR relates to survival regardless of treatment

10 treatment arms, 741 patients, minimal 2 yr and median 2.7 yr follow-up



pCR gives 94-95% 3 yr survival, regardless of drug vs. no-pCR 76-79% 3 yr survival

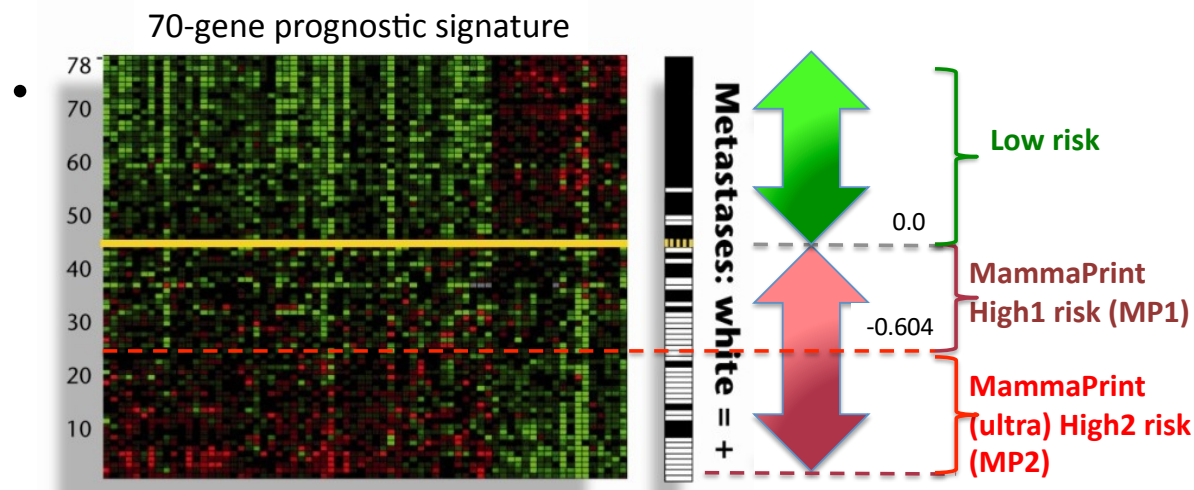
**Figure 3:** Association between pCR and Survival Outcomes (A) Kaplan Meier curves of EFS by pCR; (B) Kaplan Meier curves of DRFS by pCR.

(poster PB-11 DeMichele et al)

## Qualifying Biomarkers to improve response prediction

- Important to get every patient to pCR (increased probability of survival)
- I-SPY 2 randomizes by 8 subtypes (HR +/-, HER2+/-, MammaPrint High1/High 2)
- How can biology further identify responders?
  
- I-SPY 2 tests ‘Qualifying Biomarkers’, which have existing evidence for response prediction
  - Biology of Targeted agent, eg DNA repair deficiency, HER2 signaling, immune signatures, biology subtyping
- Presented here: 70-gene signature (MammaPrint) High1 versus High2 (high risk and very high risk for recurrence, and 80-gene molecular subtyping signature (Blueprint) which identifies luminal-, basal- and HER2-type

## 70-gene High1 and High2 risk as biomarker of response prediction



MammaPrint 70-gene expression signature identifies patients at low risk and high risk for recurrence.

Here we use a High-risk1 and High-risk2 (ultra-high) sub-classification

986 patients I-SPY 2 patients with MPHigh1/High2 class assessments (49% MP1, 51% MP2)

**Control arm:** paclitaxel (with trastuzumab (H) in HER2+), followed by doxorubicin/cytoxan (AC) (ctr treatment)

**9 Experimental arms:** veliparib/carboplatin (VC); neratinib (N); MK2206; Ganitumab; Ganetespib; AMG386;

TDM1/pertuzumab(P); H/P; and Pembrolizumab; + ctr treatment

Assessment of association of MP1/2 class and pCR:

**Univariate:** Logistic model and **Multivariate:** Logistic model adjusting for HR and HER2 status, and treatment arm as covariates. Significance threshold: p value < 0.05

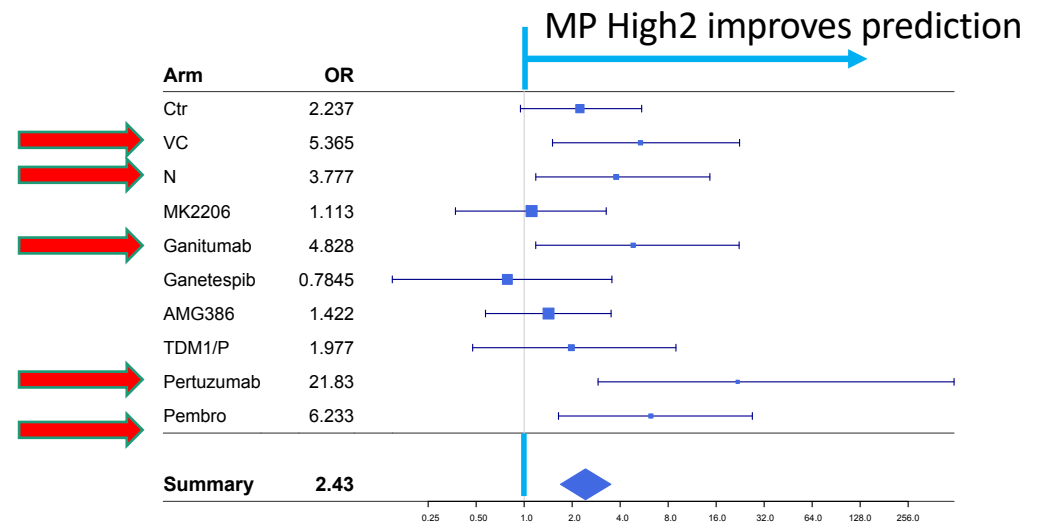
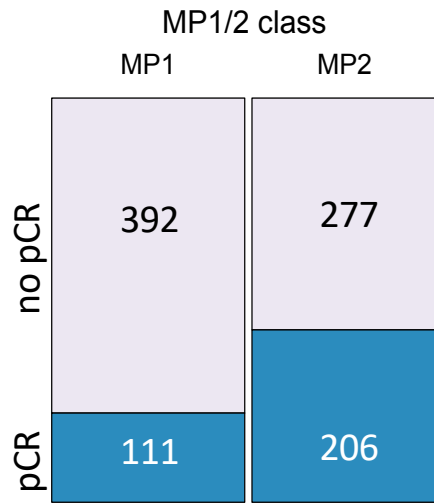


Denise Wolf, PhD  
Computational Scientist



# MPHigh1/High2 predicts ‘chemo-sensitivity’

- 986 I-SPY 2 patients across and within 10 treatment arms
- Association of MP High1/High2 with pCR across all, and within 5 arms



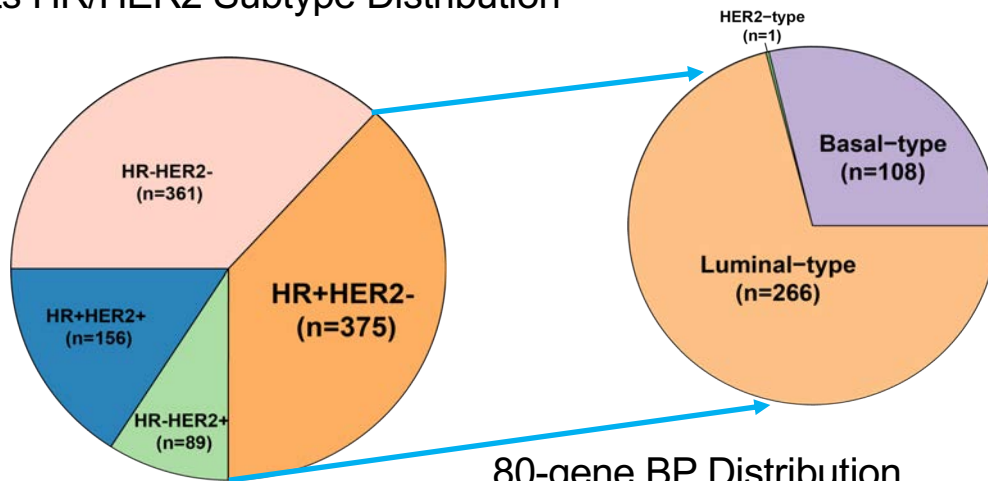
Across all arms combined, **MP High2 associates with pCR** (OR=2.43; p=1.31E-06) in a model adjusting for treatment arm, HR, and HER2 status

**MP High2 associated with pCR in half the arms** (*Veliparib-carbo, Neratinib, Ganitumab, Trastuzumab/Pertuzumab and Pembrolizumab*) in a model adjusting for HR and HER2 status (OR 2.43)  
 - most strongly in HR+/HER2- (OR 3.62; p=1.18E-0.5) (data not displayed)

## 80-gene Molecular subtype 'basal' as biomarker of response prediction

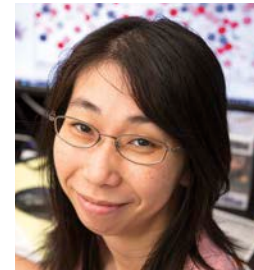
- BluePrint molecular subtype identifies functional luminal-, basal- and HER2-type

986 patients HR/HER2 Subtype Distribution



80-gene BP Distribution within HR+HER2- patients

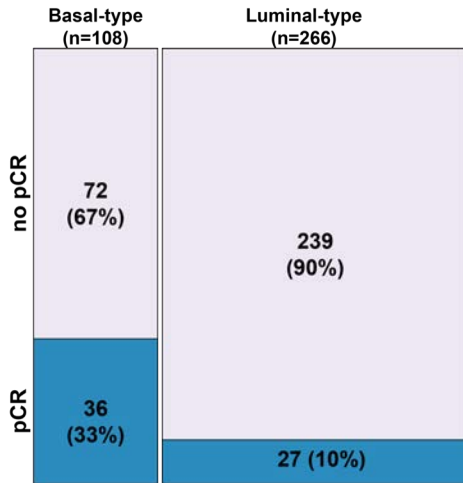
While the majority of **HR+HER2- patients** are Luminal (71%), **29% are Basal-type**



Christina Yau, PhD  
Computational Scientist

## HR+/HER2- with Basal subtype predicts ‘chemo-sensitivity’

- 375 I-SPY 2 HR+/HER2- patients across 8 treatment arms
- Association of molecular BluePrint basal subtype with pCR



Across all arms combined, **BP basal** associates with pCR (OR= 4.98, p<0.0001) in a model adjusting for treatment arm, HR, and HER2 status

Estimated pCR Rate (95% CI) by Subtype By Arm

Arm	BP-Luminal	BP-Basal
A	10% (4%-16%)	32% (20%-44%)
B	7% (0%-15%)	34% (18%- 50%)
C	9% (2%-15%)	35% (21%-50%)
D	10% (2%-17%)	29% (15%-43%)
E	10% (3%-17%)	32% (9%- 46%)
F	15% (1%- 29%)	31% (16%-46%)
G	9% (0%-19%)	32% (17%- 48%)
H	17% (5%- 29%)	41% (21%-62%)

Within treatment arms, the estimated **pCR** rates among **HR+HER2- Basal** patients ranged from **29%-41%**, compared to **7%-17%** in **HR+HER2- Luminal** patients

## I SPY 2: Learning, Innovating, and Evolving

- **Patient Centered**
  - Adaptive randomization, they get the best agent for their subtype
- **Maximizes chance of pCR and cure for each patient**
  - pCR results in 95% 3 yr disease-free survival (no-pCR 76-79%)
- **Qualifies predictive biomarkers to identify responders (ENA 2018)**
  - MammaPrint High1/High2, Blueprint molecular subtypes
  - Can prioritize treatment in subsequent trials (I-SPY 2.2 trial design)
- **Increases chance of pCR and cure for the high risk population**
  - Learn, approve drugs and combinations that are effective and less toxic
- **A design that patients like, that investigators like, where industry will participate- speeds the chance that patients will survive**
- **Advances regulatory science**

# I-SPY 2 TRIAL Study Team

## Working Group Chairs

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

## 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

<b>Executive Director:</b> Smita Asare	<b>I-SPY 2 Biomarkers/Specimens:</b>
<b>Program Administration:</b> Kat Steeg, Lorena Kanu, Julie LeDuc, Jill Parker, Melanie Hanson	Lamorna Brown-Swigart, Gillian Hirst, Denise Wolf, Chip Petricoin, Julie Wulfkuhle
<b>Safety:</b> Sausan Abouharb, Linda Doody, Monina Angeles, CCSA	<b>I-SPY Imaging Lab:</b> Jessica Gibbs, Melanie Regan
<b>Data Analysis &amp; IT:</b> Christina Yau, Adam Asare, Garry Peterson, Amy Wilson, Tim Fu	<b>Business Development:</b> Julie Sudduth-Klinger, Dan Dornbusch
<b>Operations Manager:</b> Ruby Singhrao	<b>Grants:</b> Jeff Matthews

## Site PIs

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

## Sponsor:


**Quantum Leap Healthcare Collaborative**  
Dave Mandelkern, Nancy Lisser, Mike Bankert, Adam Asare, Smita Asare

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and all of the investigators, staff, our DSMB and  
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**Biomarkers:** Denise Wolf, Christina Yau, Chip Petricoin, Julia Wulfkuhle, Lamorna Swigert, Gill Hirst & Collaborators  
**NKI:** Sabine Linn, Tessa Severson, Daniel Vis, Lodewyk Wessels, Rene Bernards, Emile Voest

**Qualifying Biomarker Process:** Denise Wolf, Christina Yau et al, Nature Partner Journals Breast Cancer, 2017

# I-SPY 2 Participating Organizations

<p><b>Sponsor</b></p>  <p>A Healthcare Collaborative</p> <p><b>Funders, Operations</b></p> 	<p><b>Investigational Agent Providers</b></p> 
<p><b>Biomarker Platforms</b></p> 	

NIH I-SPY 2.2 Program Project 2017-2010