



Integration of DNA repair deficiency and immune biomarkers to predict which early stage triple negative breast cancer patients are likely to respond to platinum containing regimens vs. immunotherapy: the neoadjuvant I-SPY 2 TRIAL

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**equal contribution*

Disclosure Information

*AACR, 4/1/2019, mini-symposium #
Denise Wolf*

I have no financial relationships to disclose.

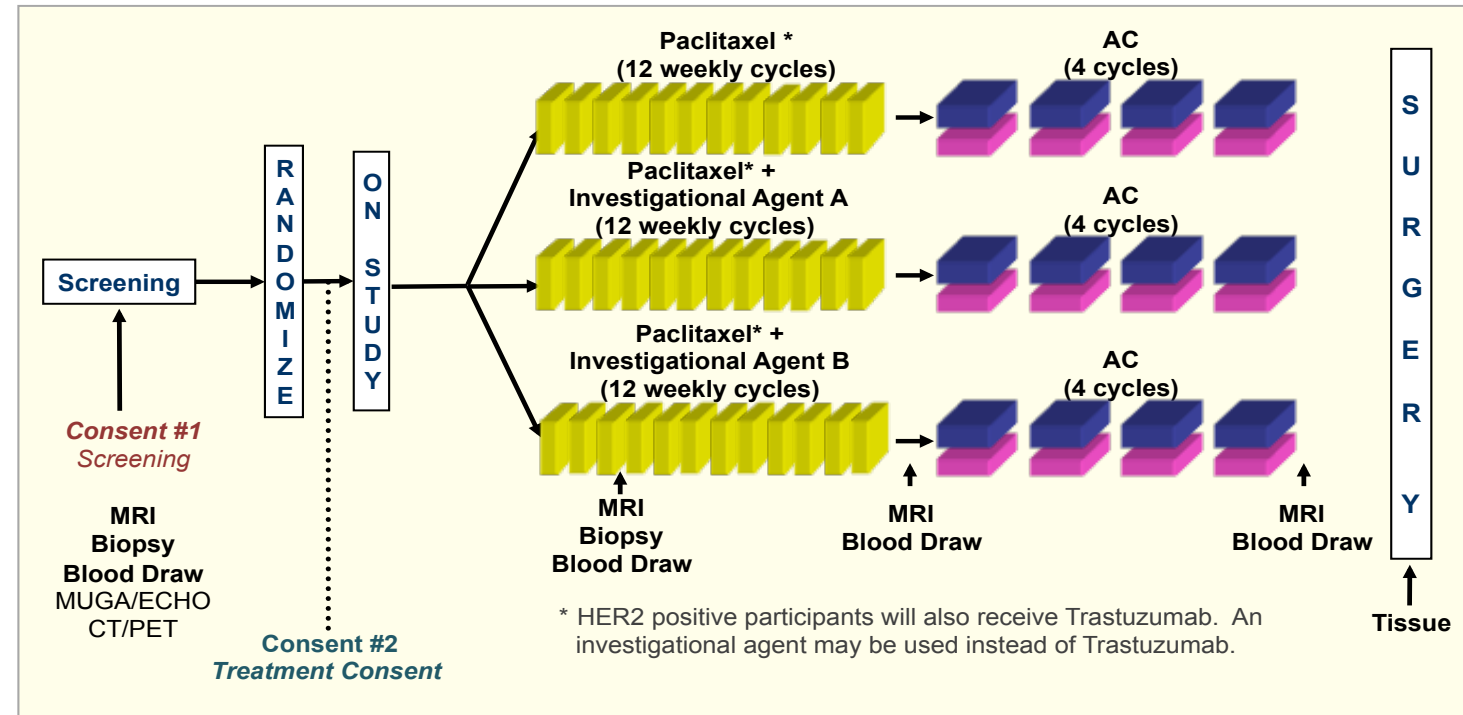
I will not discuss off label use and/or investigational use in my presentation

A changing treatment landscape for triple negative breast cancer

- **HR-HER2- (triple negative TN)**
 - Aggressive breast cancer subtype negative for estrogen receptor and HER2 amplification
- **Historically few treatment options**
 - Standard chemotherapy (anthracycline + taxane)
 - No targeted treatments
- **Multiple recent trials showing increased efficacy!**
 - Platinum-containing regimens (with and without PARP-inhibition)
 - GeparSixto, CALGB 40603, BrighTNess, **I-SPY 2**
 - Immunotherapy-containing regimens
 - **I-SPY 2**; IMpassion130,.. FDA approval - stage IV (atezolizumab); in progress: NeoTRIPaPDL1, KEYNOTE-522

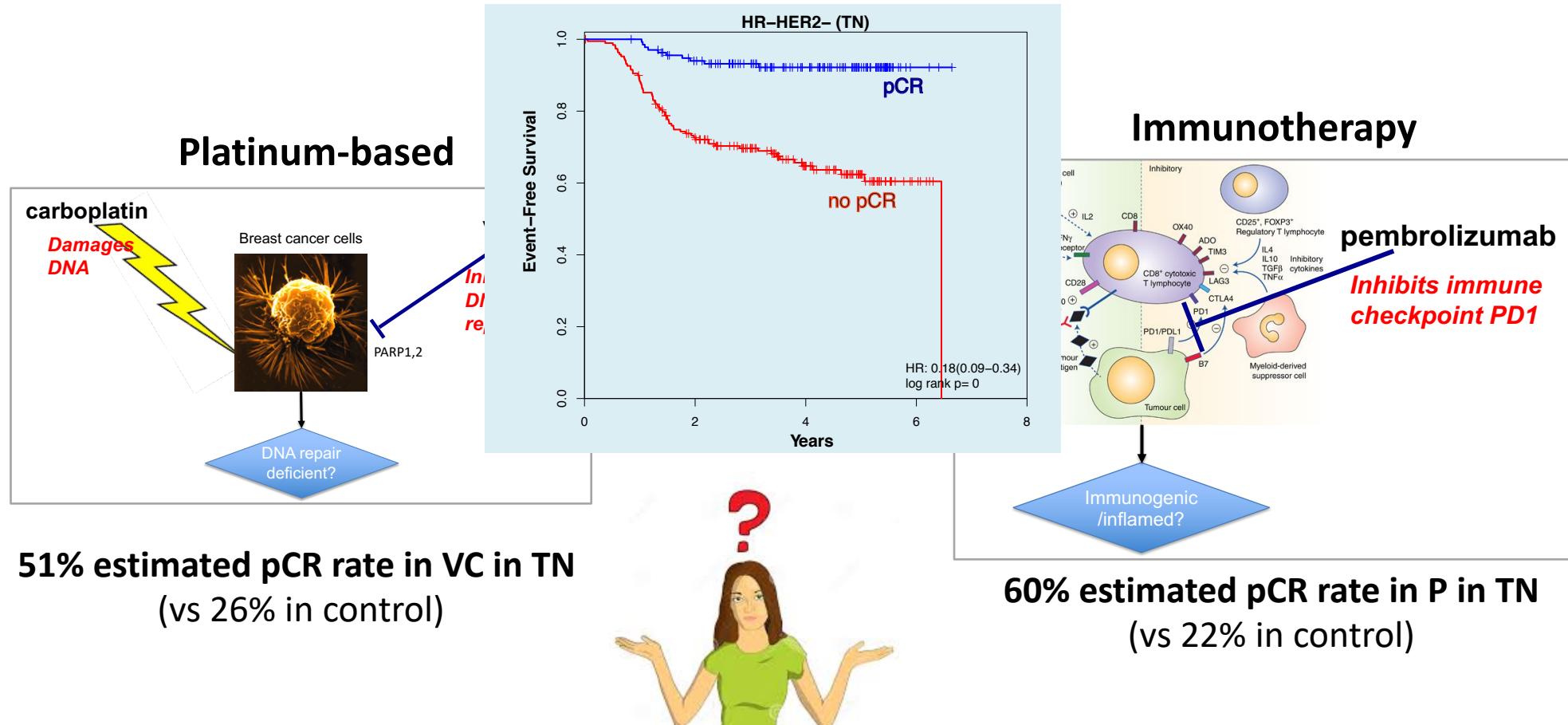
The I-SPY 2 TRIAL Standing Platform for High Risk Early Stage Breast Cancer

- **Phase II, adaptively-randomized neoadjuvant trial**
- **Shared control arm**
 - Standard neoadjuvant chemotherapy
 - HER2+ also gets standard of care for targeted agents
- **Simultaneous experimental arms**
 - Up to four
- **Primary endpoint:** pathologic complete response (pCR)
 - Defined as **no residual invasive cancer in the breast or lymph nodes**



- Agents/combinations **“graduate”** for efficacy = reaching >85% predictive probability of success in a subsequent phase III trial in the most responsive patient subset

BOTH veliparib/carboplatin (VC) combination therapy AND pembrolizumab (P) graduated in the triple negative (TN) subset



- *Who should get what and can we prioritize based on biomarkers to improve outcome?*

I-SPY 2 is a biomarker rich trial

Established

- Level 1 evidence
- FDA cleared or approved or IDE filed
- Used in clinical decision

QUALIFYING

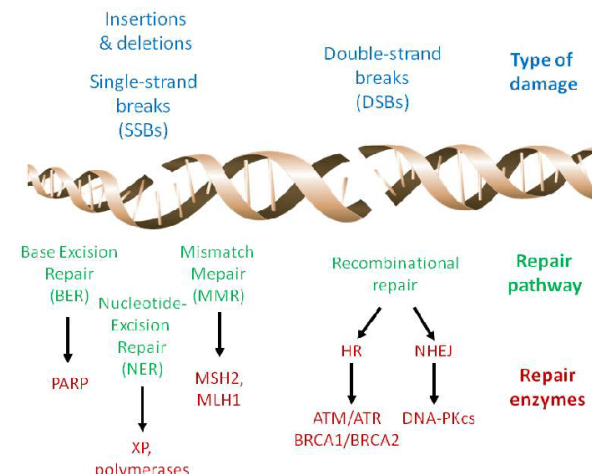
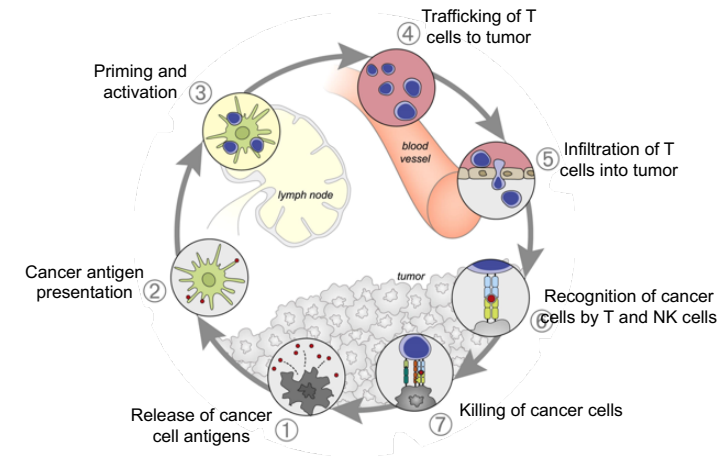
- Level 2 evidence
- Have existing evidence for response prediction
- Evaluated in CLIA setting
- May be based on mechanism of action
- Hypothesis testing
 - Pre-defined biomarkers
 - Pre-specified rigorous statistical framework

EXPLORATORY

- Biomarker discovery
- Hypothesis generation

A growing body of evidence that particular biological tumor classes are more likely to respond to a given class of agent

- For pembrolizumab and other immune checkpoint inhibitors, immune infiltrate/inflamed phenotype is associated with response.
 - Example biomarkers:** TILs, CD8+ T cells, PDL1/PD1 staining, immune expression signatures across cancer types,.. [LOTS of evidence]
- For platinum drugs +/- PARP inhibitors, DNA repair deficiency associated with response.
 - Example biomarkers:** BRCA1/2 germline mutation status, HRD in ovarian/breast cancers,...

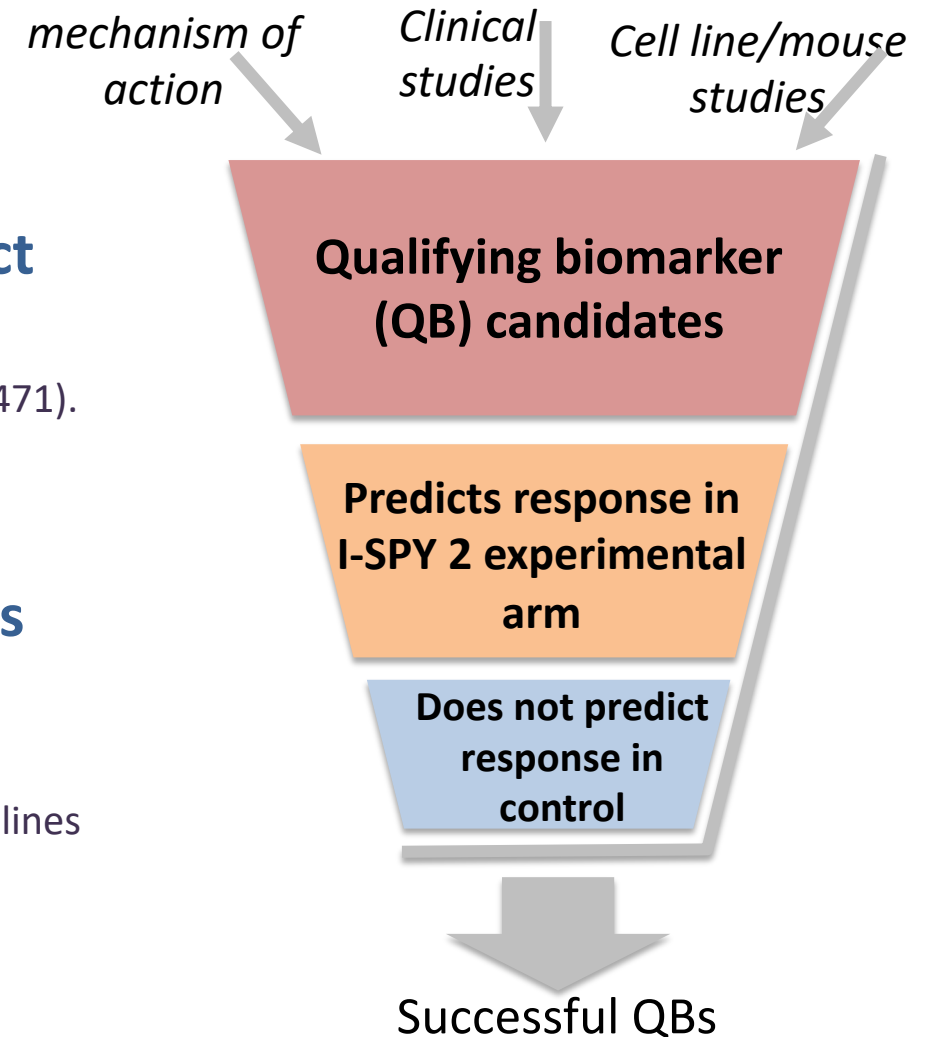


Base Excision Repair (BER) Key Players: APE1, APOBEC1, APTX, MDC1, FEN1, LIG1, LIG3				Key Players: MDC1, MDC1, MDC1, MDC1, MDC1, MDC1, MDC1, MDC1	Key Players: MDC1, MDC1, MDC1, MDC1, MDC1, MDC1, MDC1, MDC1	Key Players: MDC1, MDC1, MDC1, MDC1, MDC1, MDC1, MDC1, MDC1	Key Players: MDC1, MDC1, MDC1, MDC1, MDC1, MDC1, MDC1, MDC1
Mismatch Mediated Repair (MMR) Key Players: EXO1, MSH2, MSH6, LIG1, LIG3, FEN1, LIG1, LIG3				Key Players: MSH2, MSH6, LIG1, LIG3, FEN1, LIG1, LIG3	Key Players: MSH2, MSH6, LIG1, LIG3, FEN1, LIG1, LIG3	Key Players: MSH2, MSH6, LIG1, LIG3, FEN1, LIG1, LIG3	Key Players: MSH2, MSH6, LIG1, LIG3, FEN1, LIG1, LIG3
Nucleotide-Excision Repair (NER) Key Players: CSB, CSA, XPD, XPG, LIG1, LIG3, FEN1, LIG1, LIG3				Key Players: CSB, CSA, XPD, XPG, LIG1, LIG3, FEN1, LIG1, LIG3	Key Players: CSB, CSA, XPD, XPG, LIG1, LIG3, FEN1, LIG1, LIG3	Key Players: CSB, CSA, XPD, XPG, LIG1, LIG3, FEN1, LIG1, LIG3	Key Players: CSB, CSA, XPD, XPG, LIG1, LIG3, FEN1, LIG1, LIG3
Homologous Recombination (HR) Key Players: ATM, ATR, BRCA1, BRCA2, MRE11, EXO1, FANCD1/BRCA1				Key Players: ATM, ATR, BRCA1, BRCA2, MRE11, EXO1, FANCD1/BRCA1	Key Players: ATM, ATR, BRCA1, BRCA2, MRE11, EXO1, FANCD1/BRCA1	Key Players: ATM, ATR, BRCA1, BRCA2, MRE11, EXO1, FANCD1/BRCA1	Key Players: ATM, ATR, BRCA1, BRCA2, MRE11, EXO1, FANCD1/BRCA1
Nonhomologous End-Joining (NHEJ) Key Players: ARTEMIS, ATM, DNA-PKcs, Ku70, Ku80, LIG4				Key Players: ARTEMIS, ATM, DNA-PKcs, Ku70, Ku80, LIG4	Key Players: ARTEMIS, ATM, DNA-PKcs, Ku70, Ku80, LIG4	Key Players: ARTEMIS, ATM, DNA-PKcs, Ku70, Ku80, LIG4	Key Players: ARTEMIS, ATM, DNA-PKcs, Ku70, Ku80, LIG4

Example mechanism-of-action qualifying expression signatures predicting response to pembrolizumab and carboplatin/veliparib

Previously we showed..

- **Immune signatures, including for dendritic cells, predict response to pembrolizumab (P)**
 - 3 gene dendritic cell signature: CCL13, CD209, HSD11B1 (PMID: 28239471). Predicts pCR in I-SPY 2 patients in P arm relative to control (SABCS 2018)
- **DNA repair deficiency (DRD) biomarker PARPi7 predicts response to platinum/PARPi (VC)**
 - 7 gene DNA-repair deficiency signature PARPi-7: BRCA1, CHEK2, MAPKAPK2, MRE11A, NBN, TDG, XPA. Predicts olaparib-sensitivity in cell lines (PMID:22875744) and pCR in I-SPY 2 patients in the VC arm relative to control (PMID: 28948212)



Immune biomarkers: Danaher et. al., J Immunother Cancer. 2017 (PMID: 28239471); Yau et. al., SABCS 2018

DNA repair biomarker: Daemen et.al., Breast Cancer Res Treat. 2012 (PMID:22875744) ; Wolf, Yau, et.al., NPJ Breast Cancer. 2017 (PMID: 28948212)

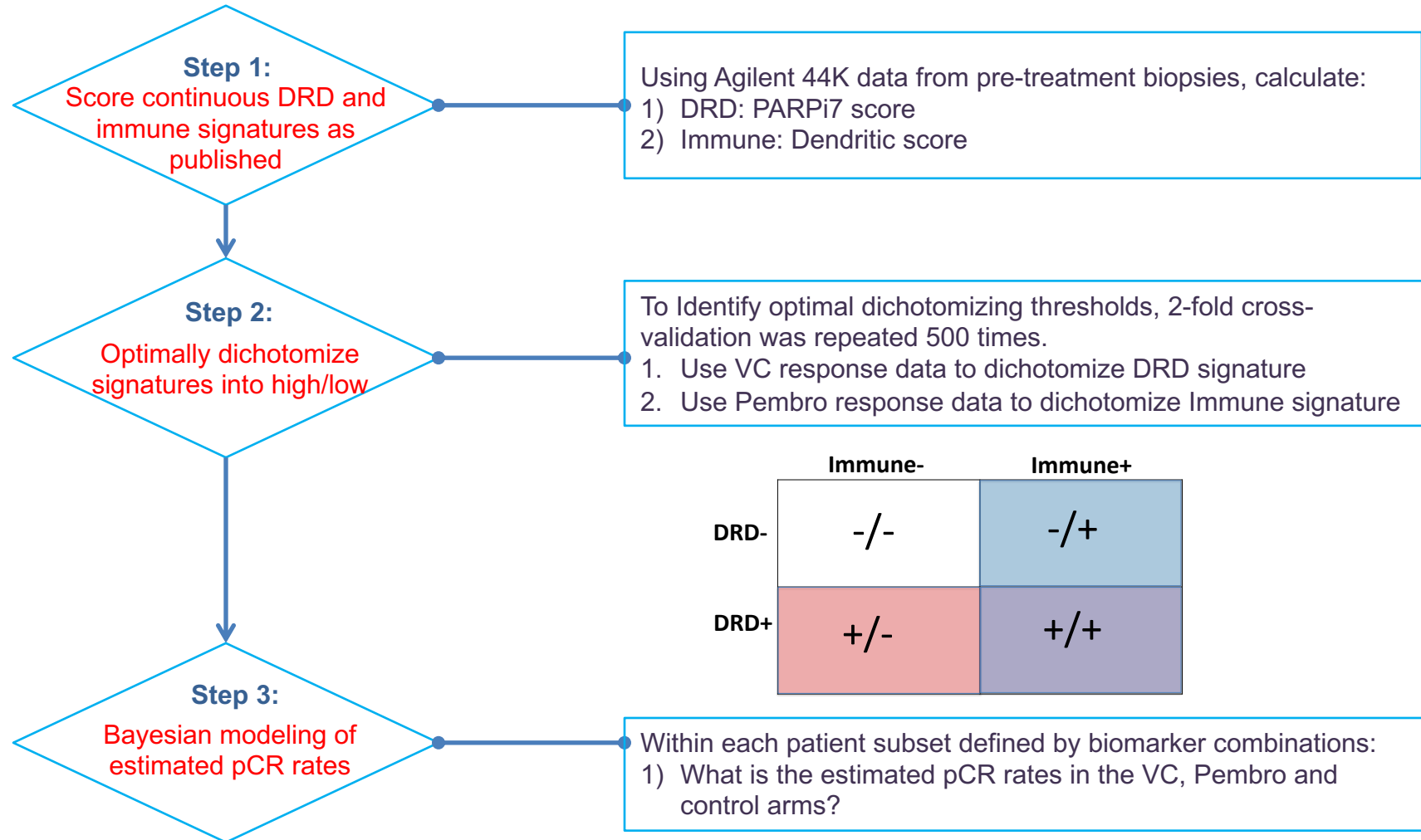
Hypothesis: overlap between Immune and DRD predictive biomarkers
can be used to identify subgroups more likely to respond to
immunotherapy vs. platinum-based therapy

	Immune-	Immune+
DRD-	$-/-$?	$-/+$ <i>anti-PD/L1</i>
DRD+	$+/-$ <i>platinum</i>	$+/+$ <i>anti-PD/L1 <u>or</u> platinum</i>

To test this hypothesis, we used the example qualifying biomarkers: **PARPi7** as our DRD biomarker (**DRD+/-**) and the **dendritic** signature as our Immune biomarker (**Immune+/-**)

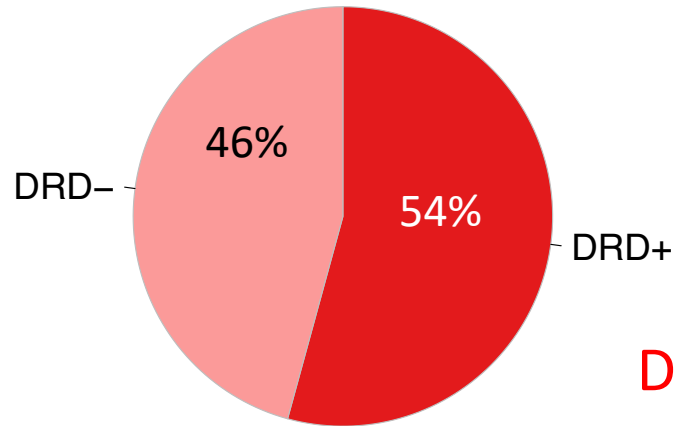
Patients and methods

153 TNBC patients available for analysis in (Control: 85; VC: 39; Pembro:29)

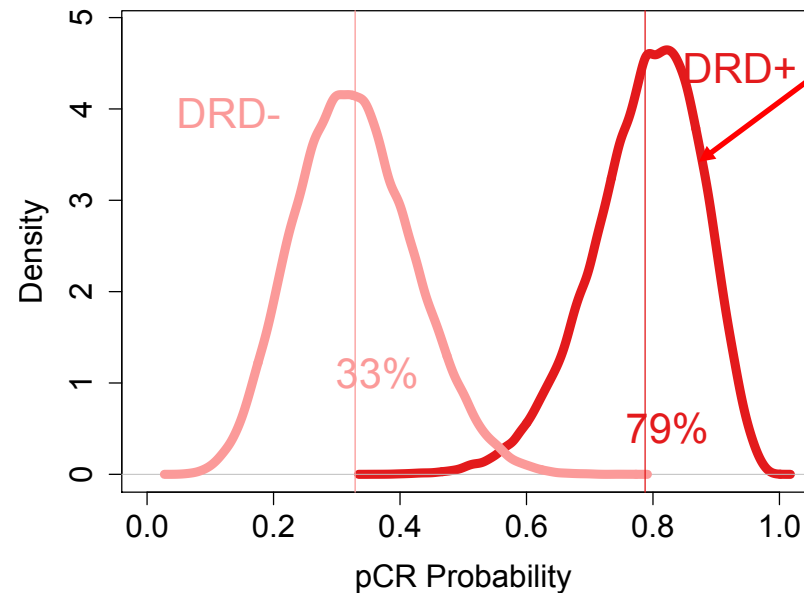


Immune and DRD biomarkers, viewed individually

TN/DRD+



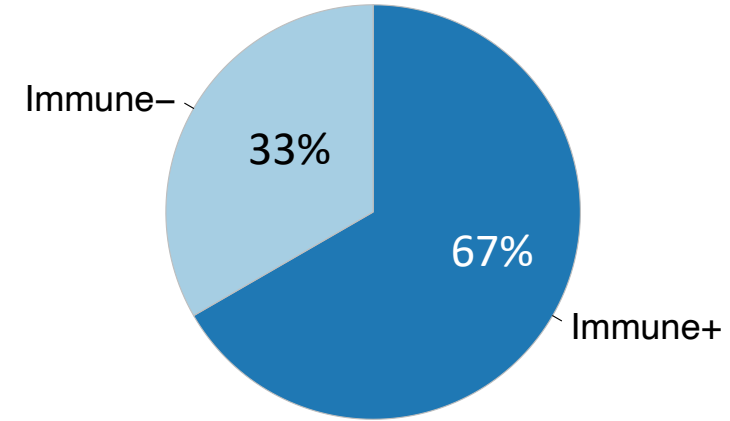
VC response: TN/DRD- vs. TN/DRD+



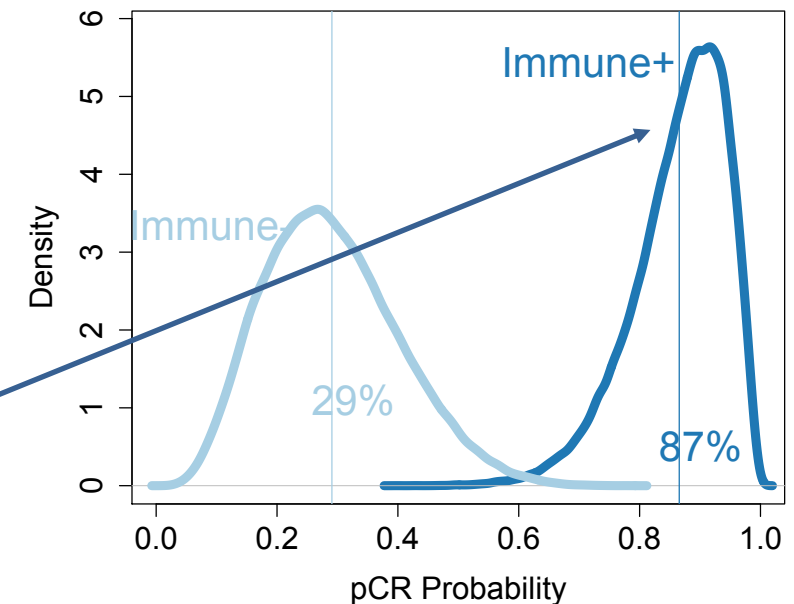
DRD+ patients have a high estimated pCR rate to VC (79%)

Immune+ patients have a high estimated pCR rate to Pembro (87%).

TN/Immune+



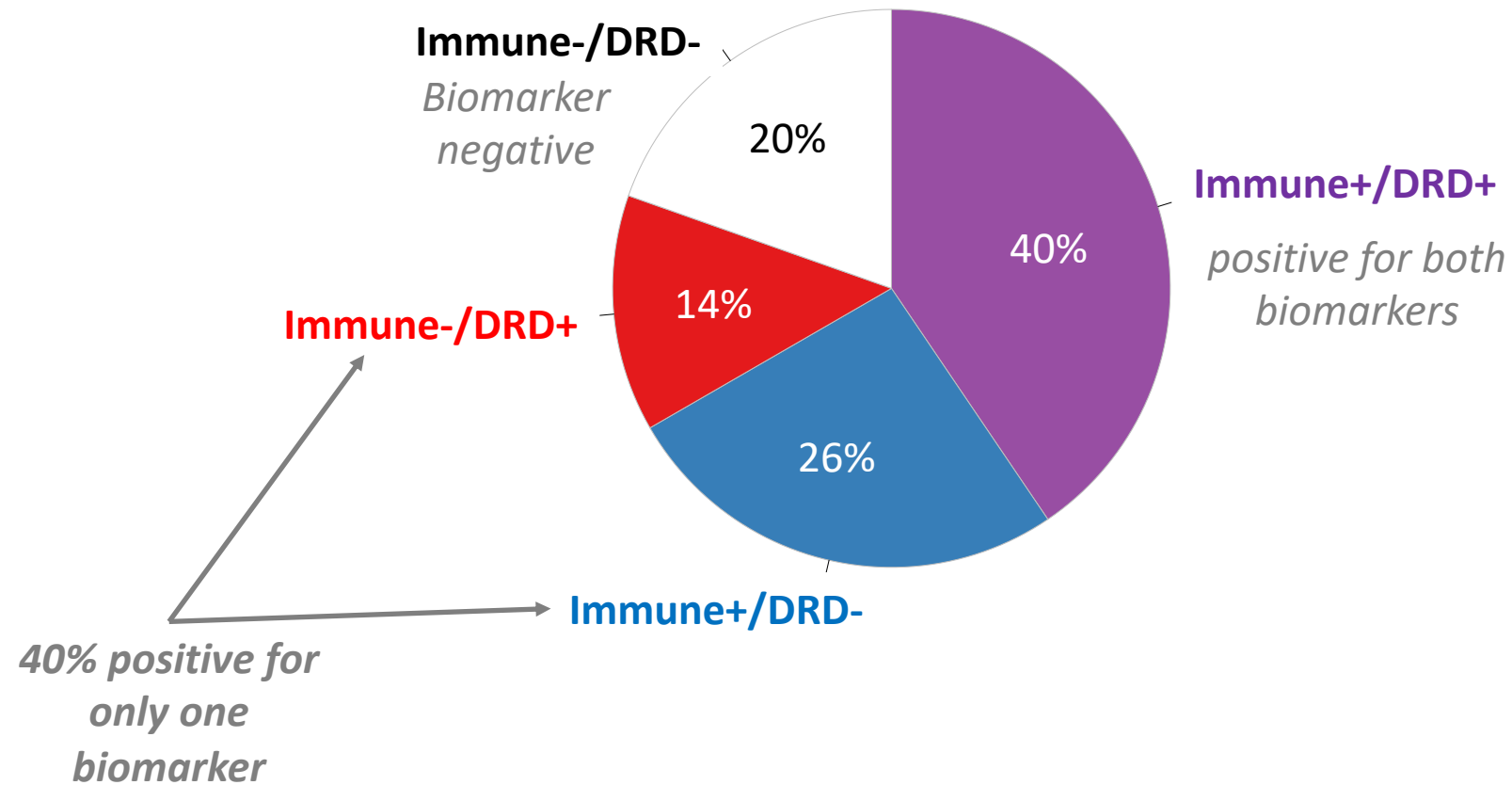
Pembro response: TN/Immune- vs. TN/Immune+



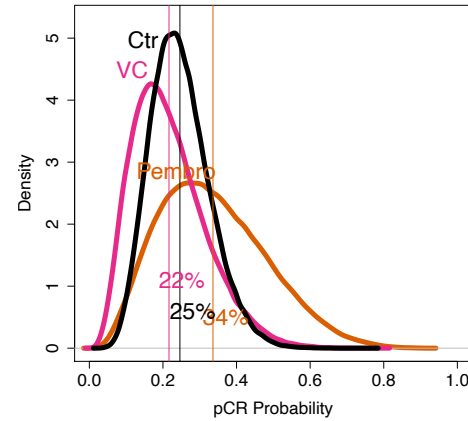
Are these the same patients?

(What is the overlap between Immune+ and DRD+?)

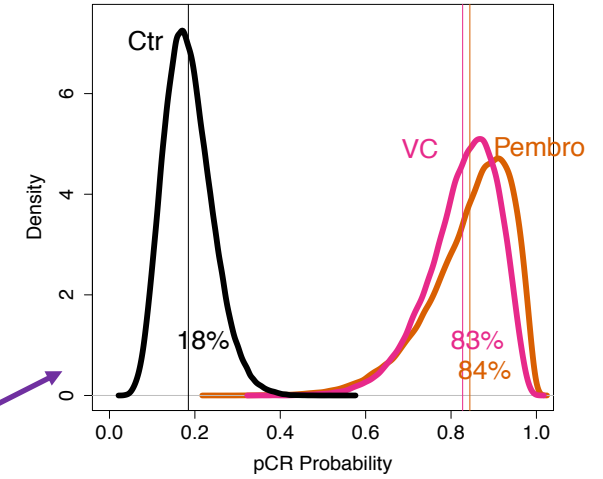
Overlap between immune and DRD predictive biomarkers in TNBC



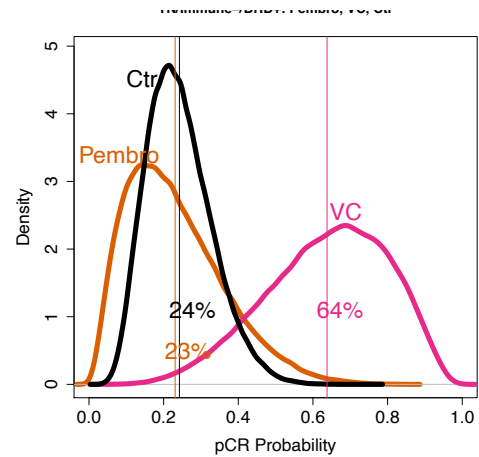
Estimated pCR distributions within biomarker subgroups



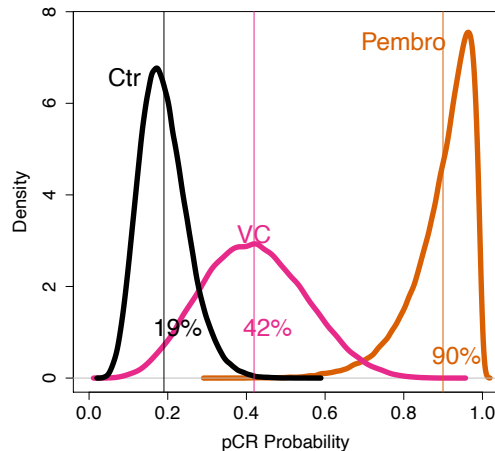
Low pCR in all arms



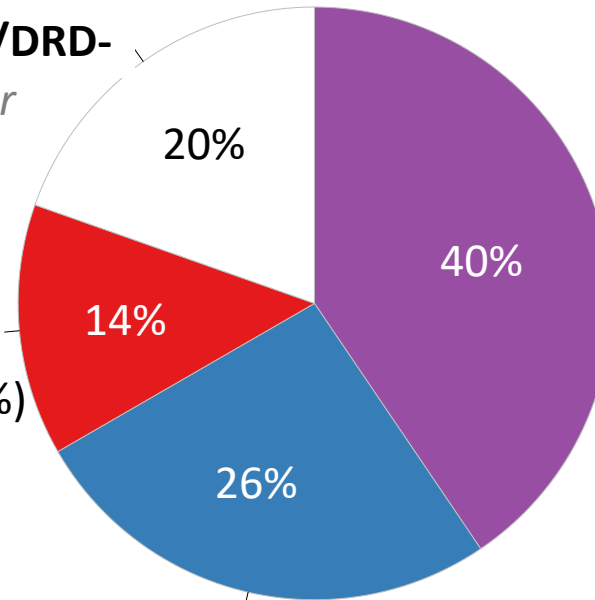
high pCR in Pembro (84%) **and** VC (83%)



← **Immune-/DRD+**
higher pCR in VC (64%)

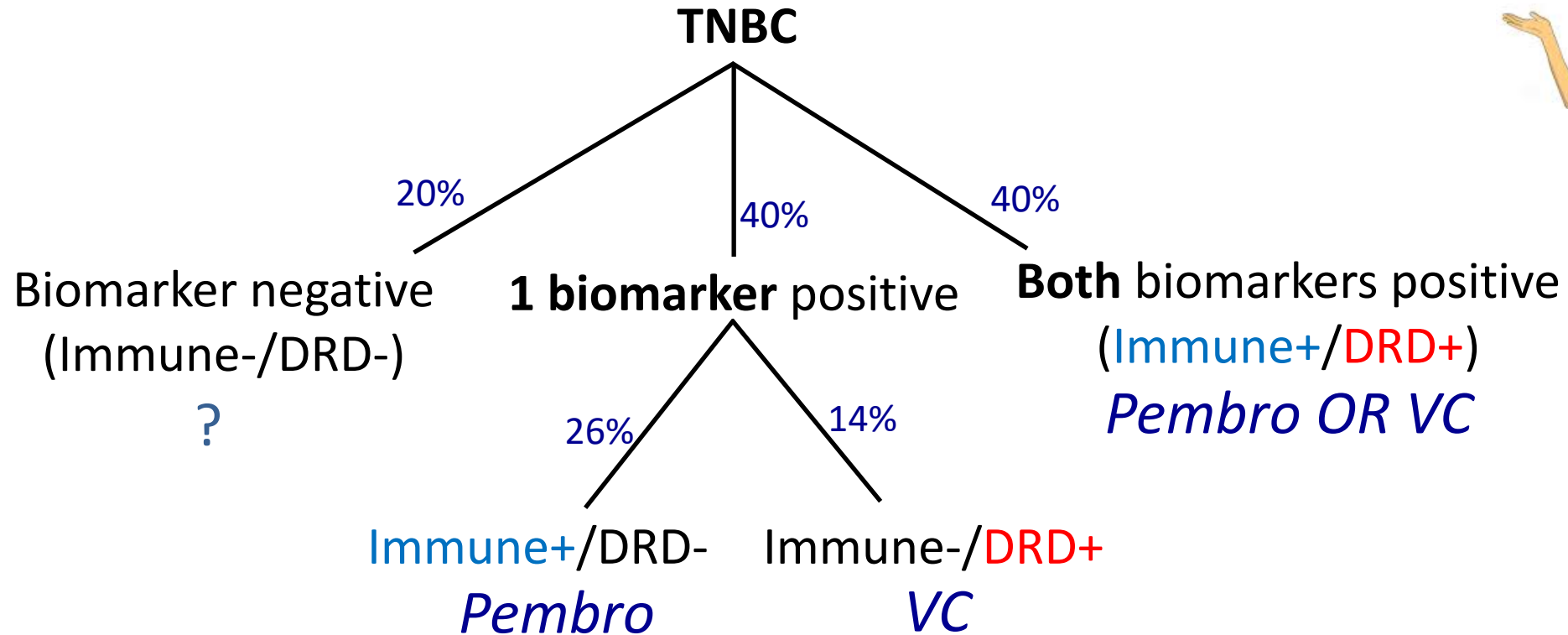


← **Immune+/DRD-**
highest pCR in Pembro (90%)



Immune+/DRD+
positive for both biomarkers

Which drug should be prioritized for whom?



Summary

- TNBC is experiencing a period of optimism, with trials showing increased efficacy for platinum and immunotherapy containing regimens
- **Question:** are patients likely to respond to one treatment also likely to respond to the other, or is there specificity: ***for what percentage does treatment selection matter? How to prioritize?***
- In I-SPY 2, carboplatin/veliparib and pembrolizumab both graduated in the TN subset
- Previously we showed: DRD signatures (e.g. PARPi7) predict response to VC; and immune signatures (e.g., dendritic cell score) predict response to Pembro
- One can use the overlap between Immune and DRD biomarkers to identify patient subgroups more likely to respond to immunotherapy vs. platinum-based therapy
- 40% high in both biomarkers (Immune+/DRD+) => high pCR in both arms (*either treatment good!*)
- 40% high in just one biomarker => highest pCR in Pembro if Immune+/DRD-; highest pCR in platinum if Immune-/DRD+ (*treatment choice matters! Basis for prioritizing?*)
- 20% low in both (Immune-/DRD-). Low pCR rate in both arms. *Alternative approach?*
- Caveat: numbers are small. Validation required.

I-SPY 2 Platform Trial Study Team

Working Group Chairs

PI: Laura Esserman	Operations: Angie DeMichele
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

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Program Management Office

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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

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

Biomarkers: Denise Wolf, Christina Yau, Chip Petricoin, Julia Wulfkuhle, Lamorna Swigert, Gill Hirst, Mark Magbanua & Collaborators

I-SPY 2 Participating Organizations and Funders

