



How to design trials to meet the perspectives of personalized cancer medicine

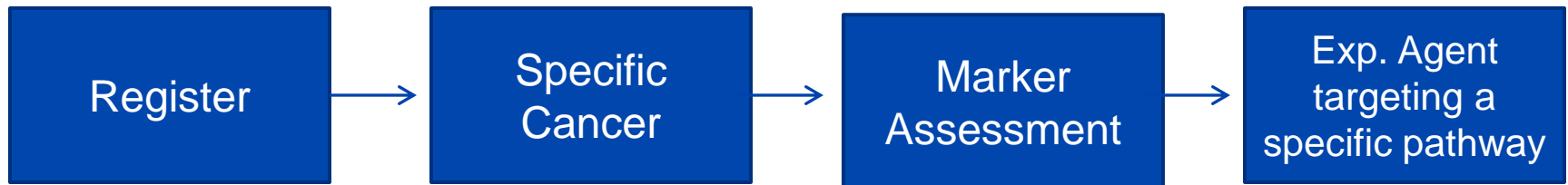
Sumithra J. Mandrekar
Professor of Biostatistics, Mayo Clinic

Molecular Profiling: Challenges and Perspectives
Special Symposium
ESMO Congress, September 28, 2014

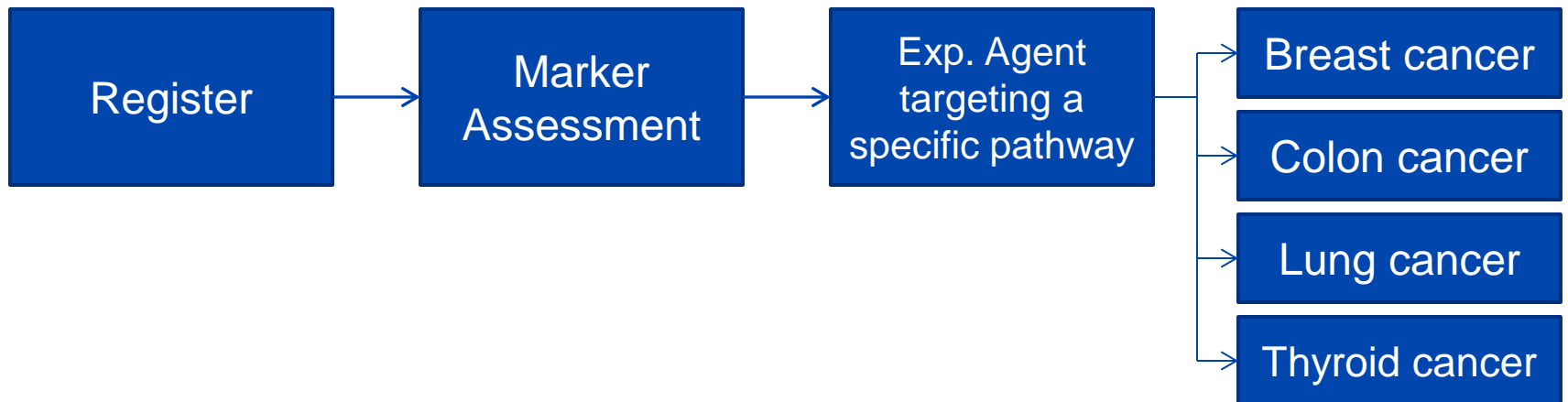
Disease-based Trials



Disease-based and Marker-based Trials



Marker-based Trials



Early Phase Design Considerations

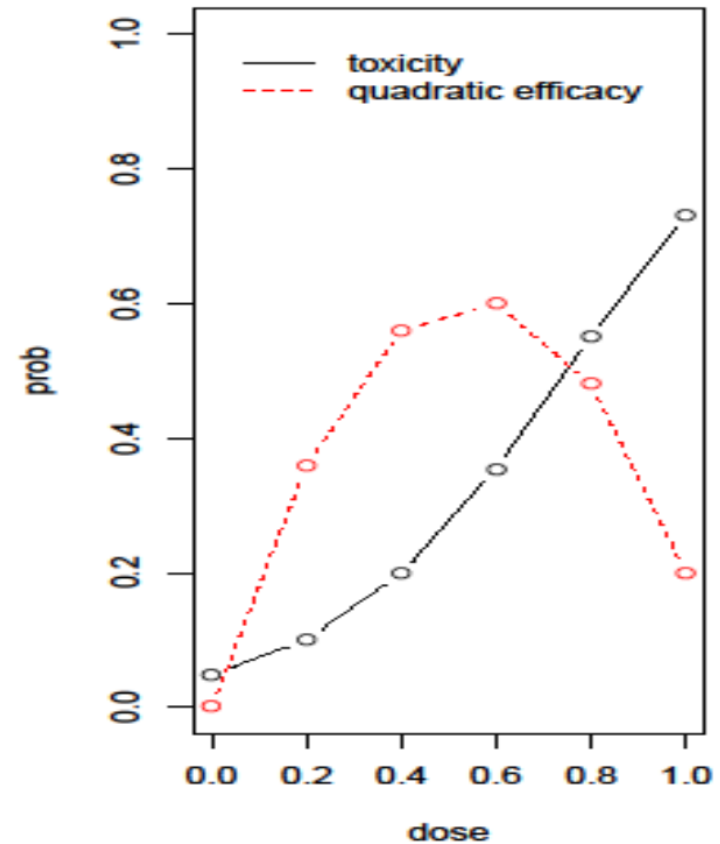
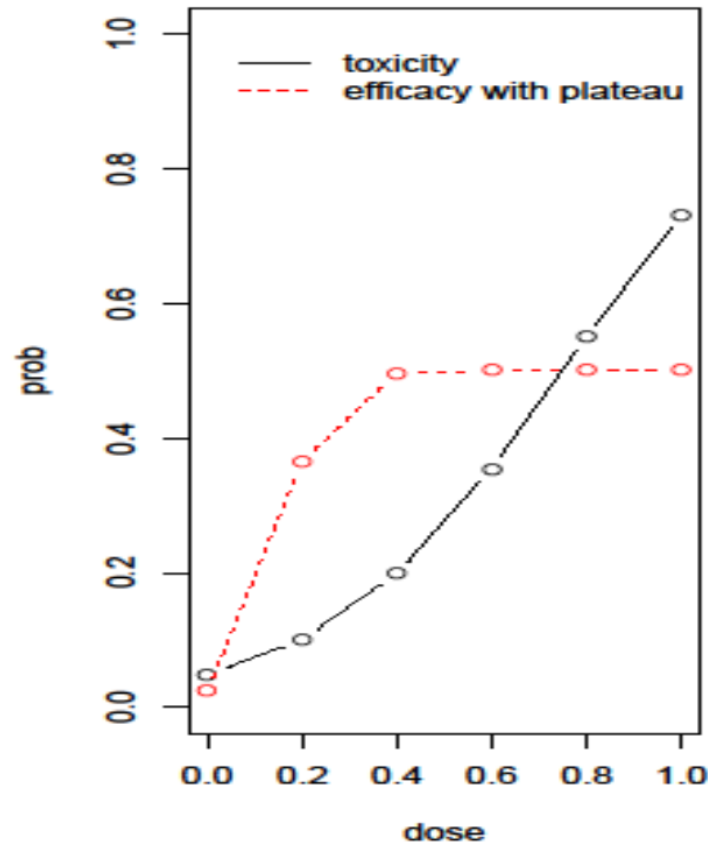
- Establish preliminary efficacy signal, in addition to understanding safety
 - Choice of endpoints
 - Model-based design algorithms
- Identify subsets of patients most likely to benefit from the new treatment
 - Enrichment strategies
 - Expansion cohorts

Choice of Endpoints

- **M**aximum **T**olerated **D**ose
 - Highest safe dose is the most efficacious dose?
- **M**aximum **E**ffective **D**ose
 - Incorporate a measure of efficacy in addition to safety assessment – Dual Endpoints.
 - Toxicity; and Biomarker or clinical response
- Challenges with biomarker assessments:
 - Correlation between biomarker and clinical outcome established?
 - Assay characteristics and performance?
 - Assessment time points?

Model Based Designs

Dynamic estimation of the dose-toxicity and dose-efficacy curves



Patient Selection

- Enrichment strategies: pros
 - Identify subsets who benefit most from treatment
 - Increase feasibility of trials in rare genotypes
 - Examples:
 - Crizotinib for ALK positive NSCLC
 - Vemurafenib for BRAF mutation melanoma
- Enrichment strategies: caution!
 - Valid assays?
 - Real time assessment?
 - Complete understanding of tumor biology?
 - Complete understanding of drug metabolism pathway?

Expansion Cohorts

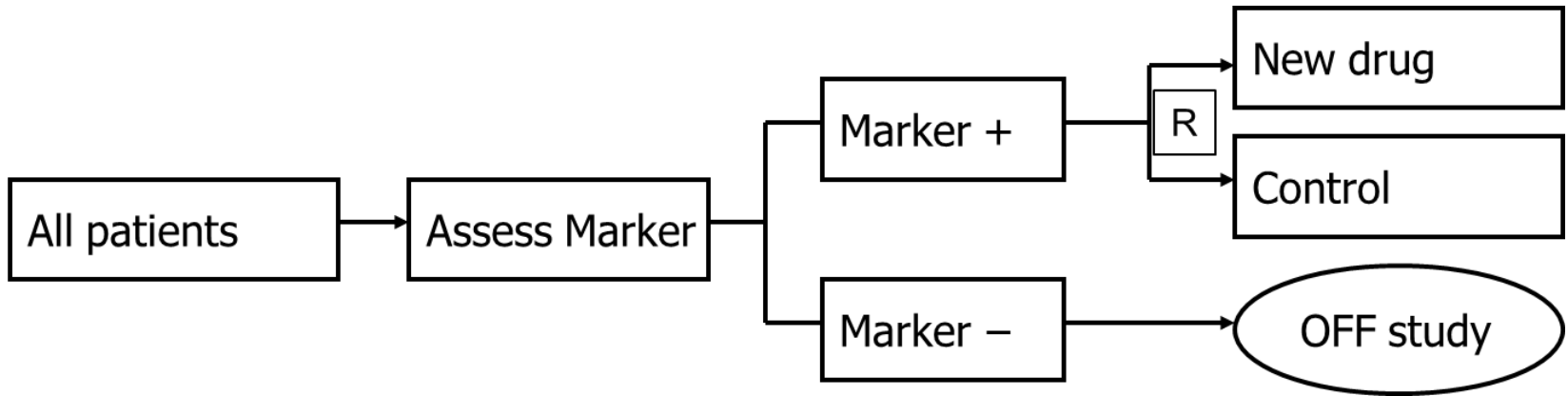
- Studying safety profile
 - Exploring neighboring dose levels for BOD/MED
 - Performing PK/PD studies
 - Assessing efficacy in enriched subgroups

 - Design of expansion cohorts:
 - Rigorous: pre-defined hypotheses etc.
- OR
- Exploratory – refine assay, cut points, patient subset identification

Phase II and III Design Considerations

Single marker case

Enrichment or targeted trial design



- Randomize marker positive patients only

Enrichment Design example: Vemurafenib in Melanoma with BRAF V600E Mutation

- Compelling evidence: Prior phase I and phase II trials demonstrated response rates of more than 50% in patients with metastatic melanoma with the BRAF V600E mutation.
 - 5 patients with WT did not respond in Phase I to therapeutic doses of Vemurafenib
- Phase III trial: Patients with BRAF V600E mutation were randomized 1:1 to vemurafenib with dacarbazine
- Central testing: At one of five central laboratories in the United States, Germany, and Australia.
- Vemurafenib was associated with a relative reduction of 63% in the risk of death and of 74% in the risk of either death or disease progression, as compared with dacarbazine ($P < 0.001$ for both comparisons).

Using markers to restrict trial eligibility: beware

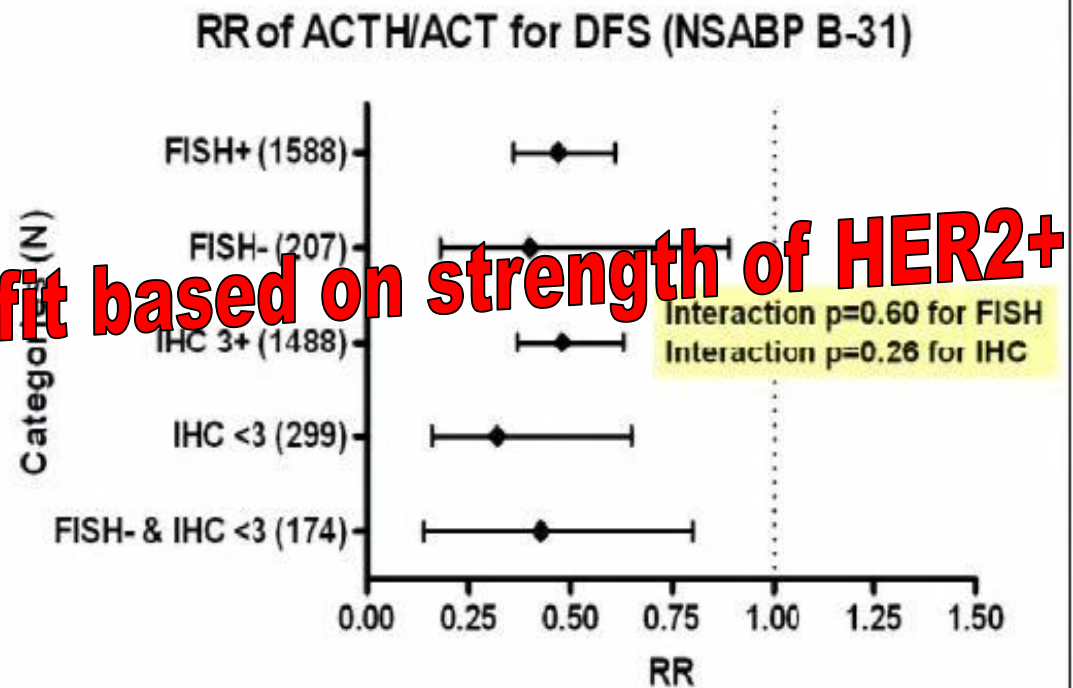
Table 1. Relative Risks of Disease Progression and Death among Patients in the ACT with the ACT Group.*

End Point and Central HER2 Assay†	ACT no. of events/total no. of events	ACTH	Relative Risk (95% CI)
Disease progression			
HER2-positive	163/875	85/804	0.47 (0.37–0.62)
HER2-negative	20/92	7/82	0.34 (0.14–0.80)
Death			
HER2-positive	55/875	38/804	0.66 (0.43–0.99)
HER2-negative	10/92	1/82	0.08 (0.01–0.64)

* The 95% confidence intervals (CI) and P values were adjusted according to the number of events from the univariate Cox proportional-hazards model for each subgroup in the Adjuvant Breast and Bowel Project B-31 trial. ACT denotes doxorubicin, cyclophosphamide, and epirubicin; ACTH denotes doxorubicin, cyclophosphamide, and epirubicin plus trastuzumab.

† Central HER2 assay results were defined as negative if they were negative by both fluorescence in situ hybridization (FISH) and immunohistochemical analysis (IHC) and were positive if they were positive by either assay.

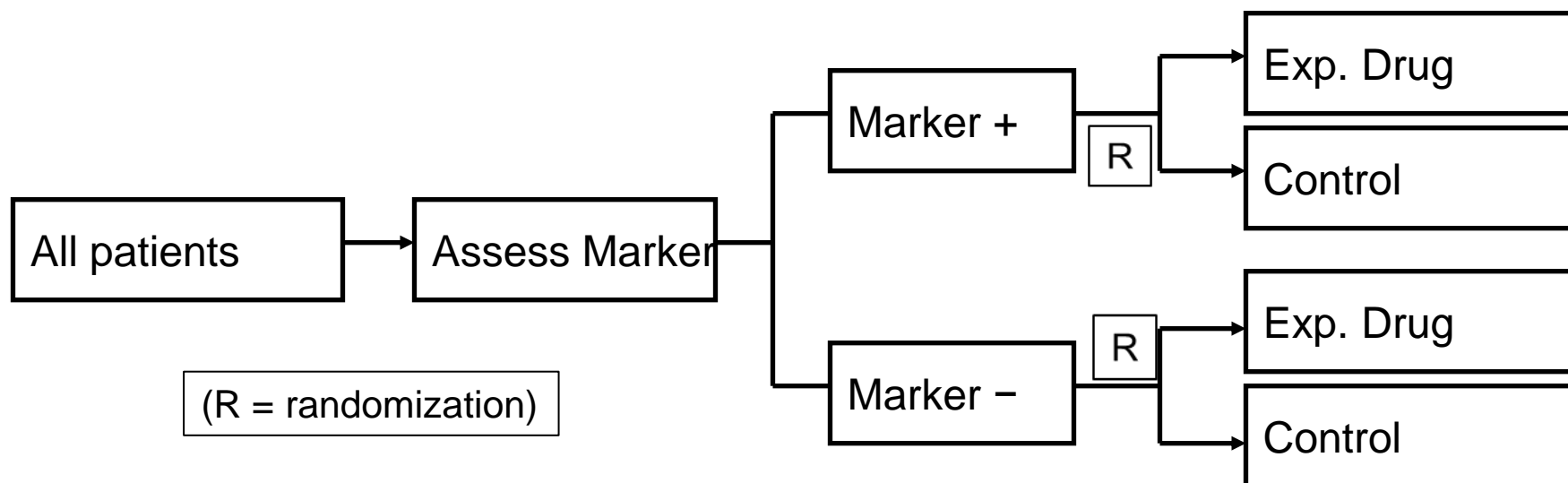
No difference in benefit based on strength of HER2+



Ongoing study of Herceptin in patients with low (1+ or 2+) HER2-positive BC.

Paik et al, NEJM 2008
Hayes et al., NEJM 2011

Marker by treatment interaction Design, AKA Biomarker Stratified Design



Randomize all patients, stratified by marker status.

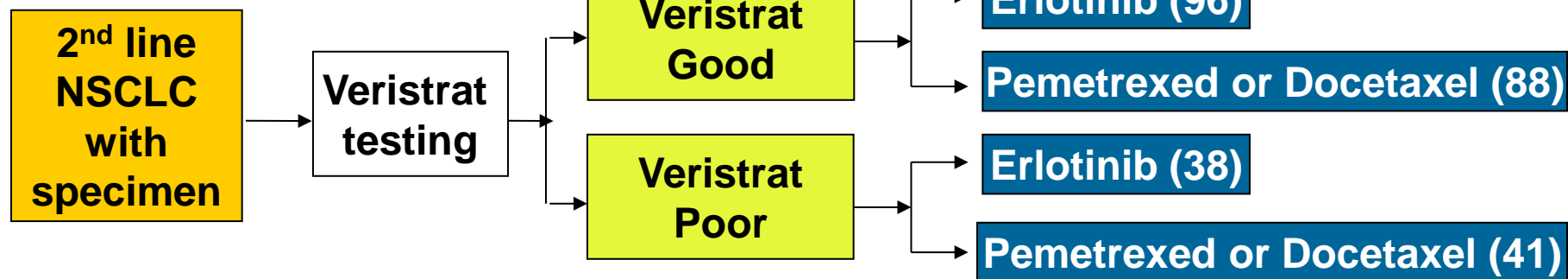
Mostly used in settings with two approved regimens.

Randomized Proteomic Stratified Study of Second-Line Erlotinib versus Chemotherapy in Patients with Inoperable Non-Small Cell Lung Cancer

PROSE Trial

VeriStrat is a serum based protein assay.

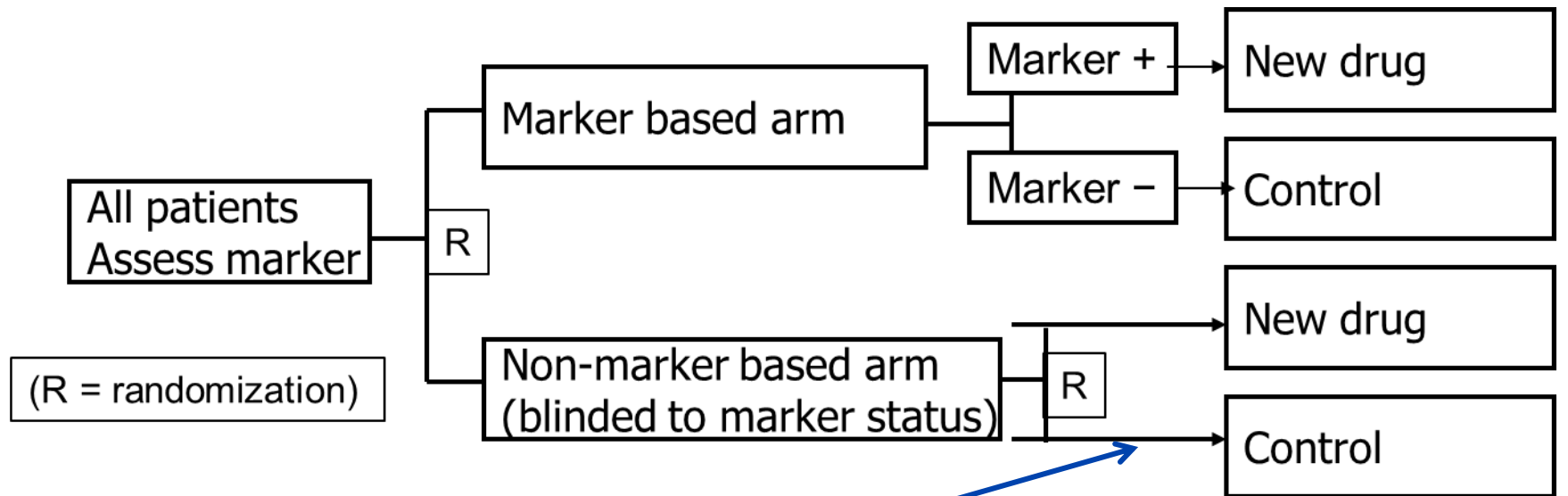
Initial Registration



Primary Endpoint: Overall Survival; Secondary: PFS, RR

Good group: No difference in OS; Poor Group: Chemo better than Erlotinib
Significant interaction between treatment and veristrat classification (p=0.037)

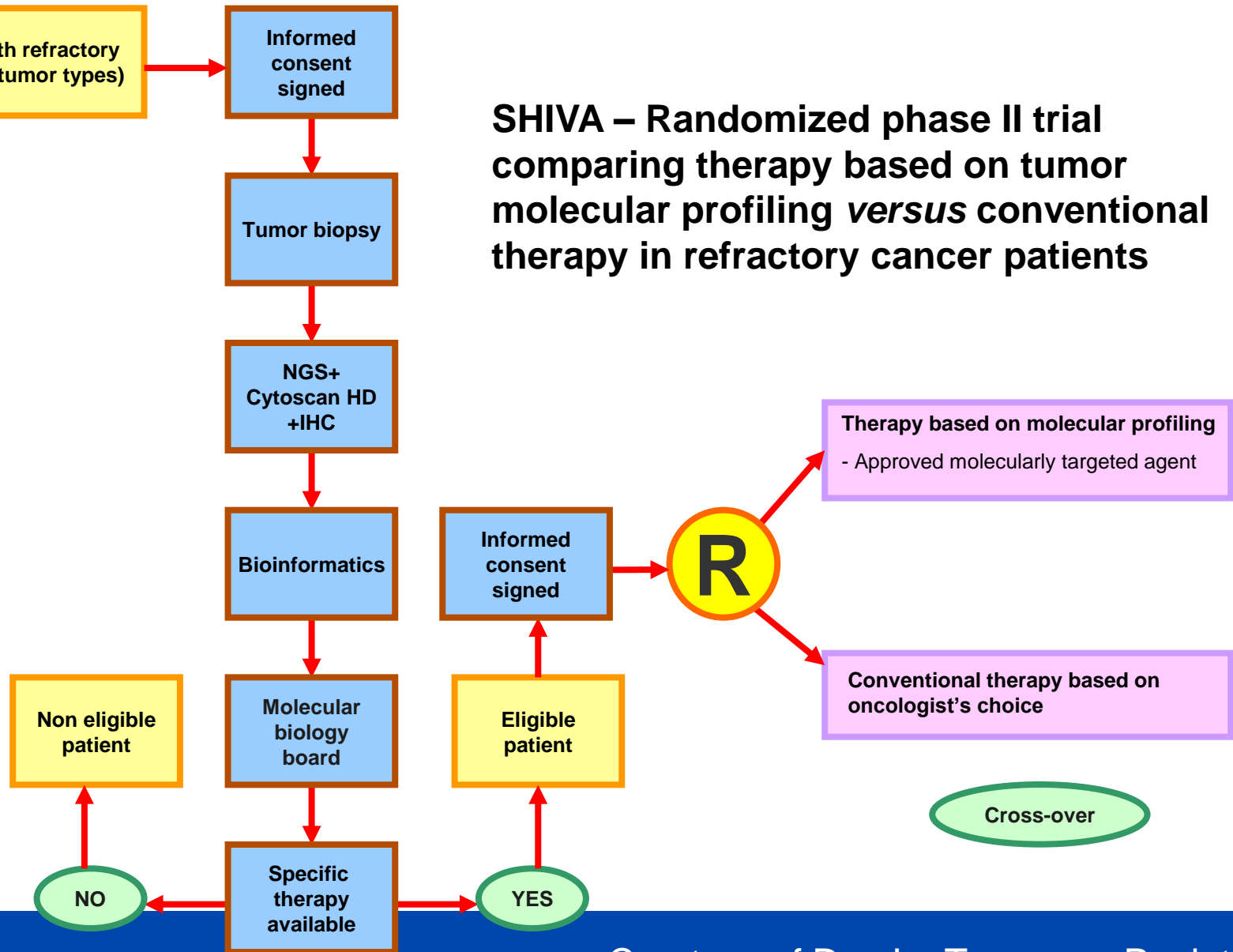
Marker Strategy design



Or assign all to control, or use physician's choice

- %Overlap in treatments on both arms – dilutes the ability to distinguish treatment from marker effect!
- Special considerations needed for the randomization ratio to marker prevalence in the non-marker based arm

Enrichment followed by “modified” marker strategy design



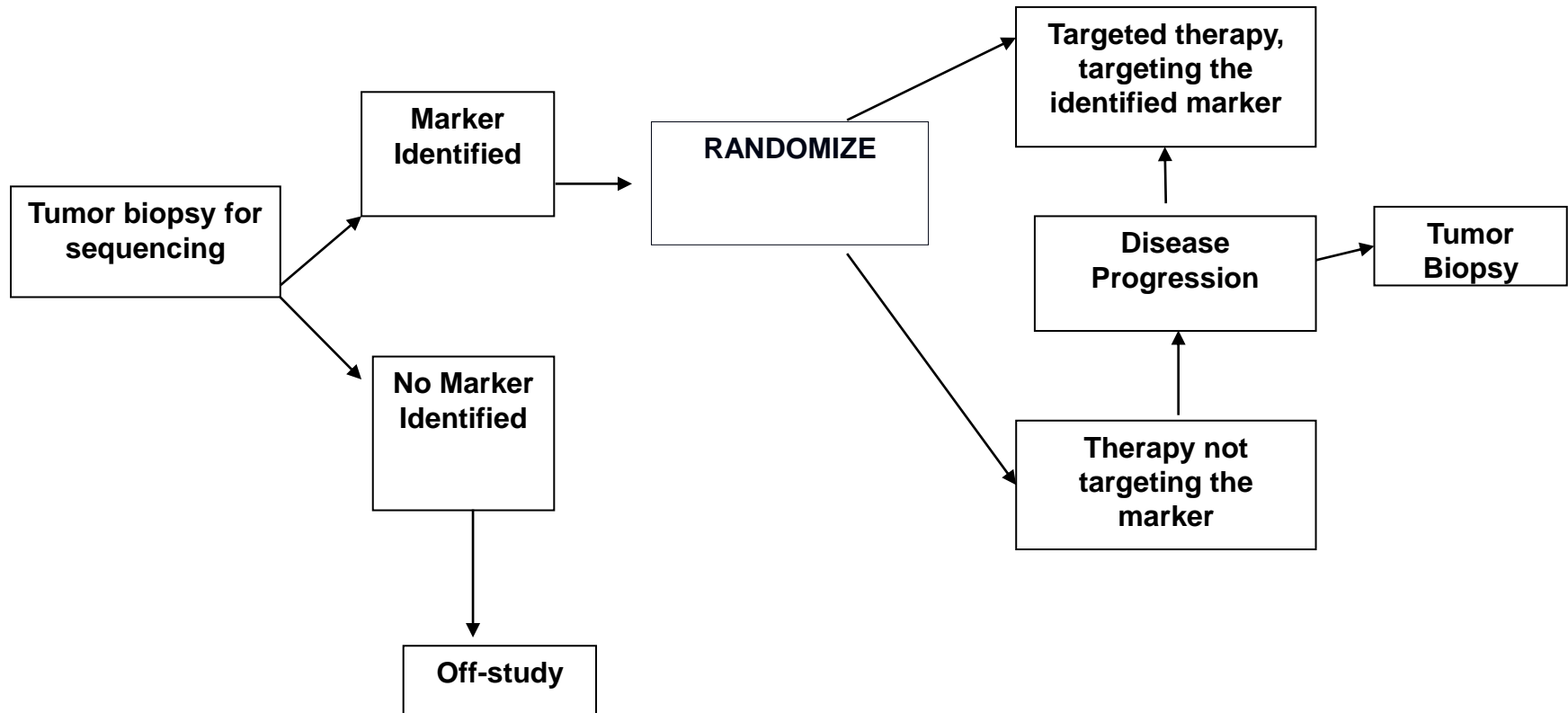
SHIVA Design Details

- Endpoint: 6 months PFS rate
 - Hypothesis: 15% for cytotoxic agents versus 30% in the experimental arm ($HR = 1.6$)
- → 142 events; 2-sided type 1 error of 5%, power of 80%
- → 200 patients to be randomized (~1000 screened)

NCI Precision Medicine Initiatives

Enrichment followed by modified marker strategy design

M-PACT Trial



Endpoints: response rate and 4-month progression-free survival

Single marker case: Prevalence Considerations

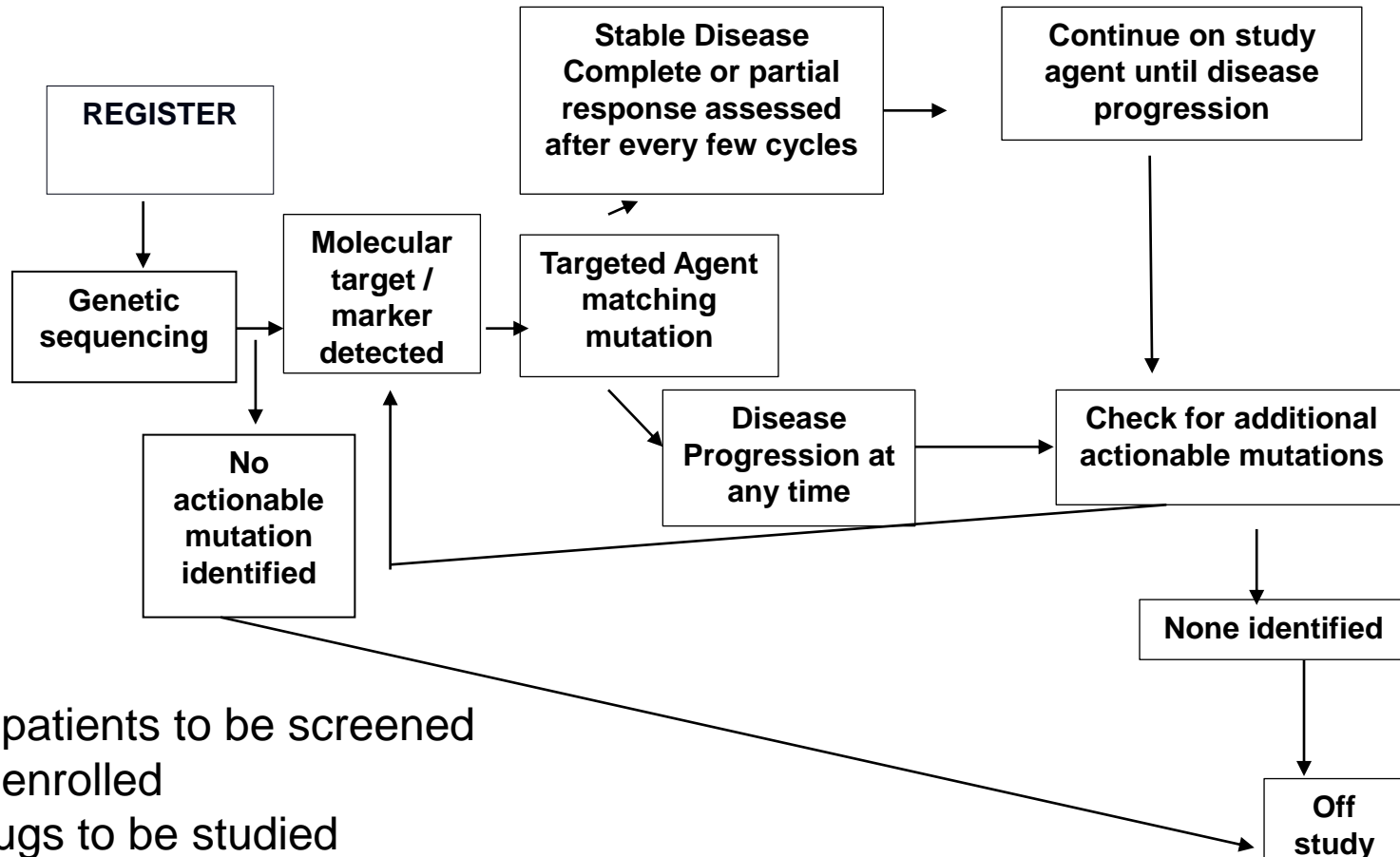
- Low ($< 20\%$): Consider enrichment designs
- High ($> 50\%$): All-comers with retrospective marker subgroup assessments or adaptive designs
- Moderate ($20\%-50\%$):
 - Stratified by marker, primary hypothesis in one marker subset; but enroll all to confirm no benefit in the other subgroup

Rare populations: N of 1?

- Enroll and treat few subjects
- Examine genomic profiles
 - Match treatment to genomic profile?
- Assess safety and efficacy within:
 - Certain tumor types, and/or
 - Certain genomic profiles
- Basket Trial Designs
 - 1st phase: unselected population -- Identify patients who benefit using genomic profiling, NGS etc.
 - 2nd phase: prospectively screen patients with that profile, parallel phase 2 studies

NCI Precision Medicine Initiatives

MATCH Trial (ECOG-ACRIN)



~3000 patients to be screened

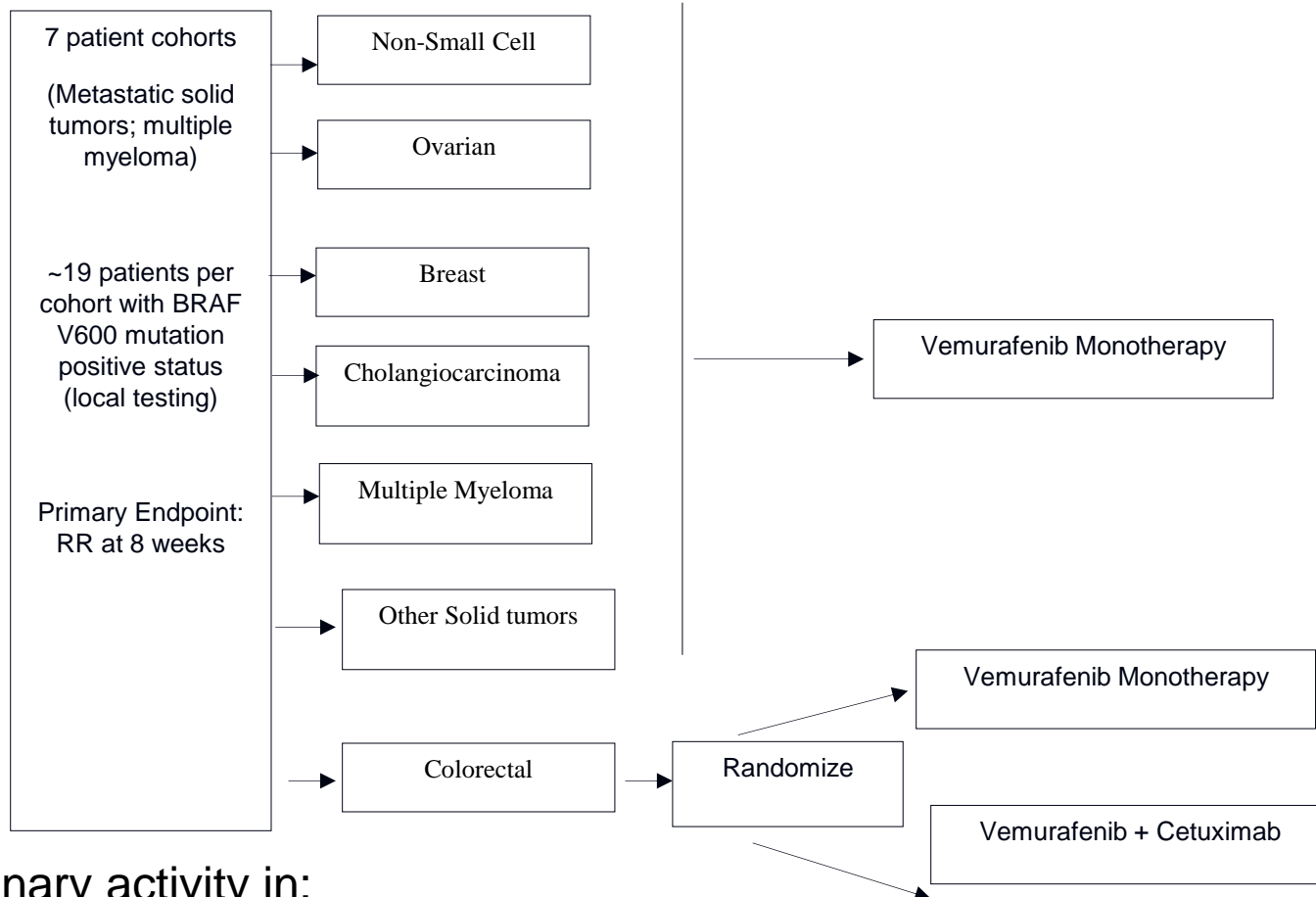
~1000 enrolled

~15 drugs to be studied

Endpoints: Response Rate and 6-month Progression-free Survival Rate

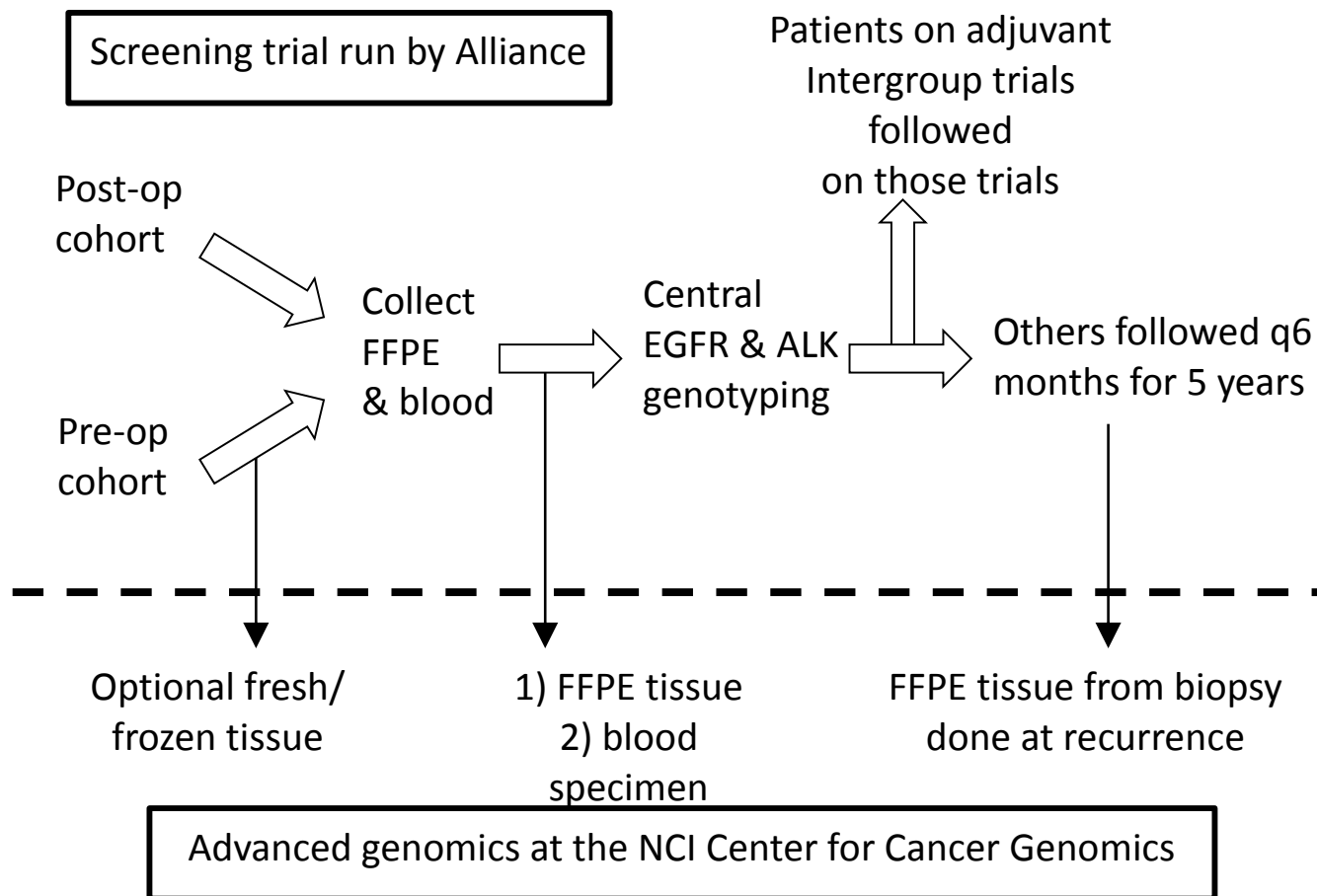
Success: > 25% RR, and/or > 35% 6-month PFS rate

Vemurafenib Basket trial (VE-BASKET): Non-Melanoma BRAF V600-mutation positive tumors



Preliminary activity in:
NSCLC, Orphan tumors, Cholangiocarcinoma

Adjuvant Lung Cancer Enrichment Marker Identification and Sequencing Trial (ALCHEMIST)

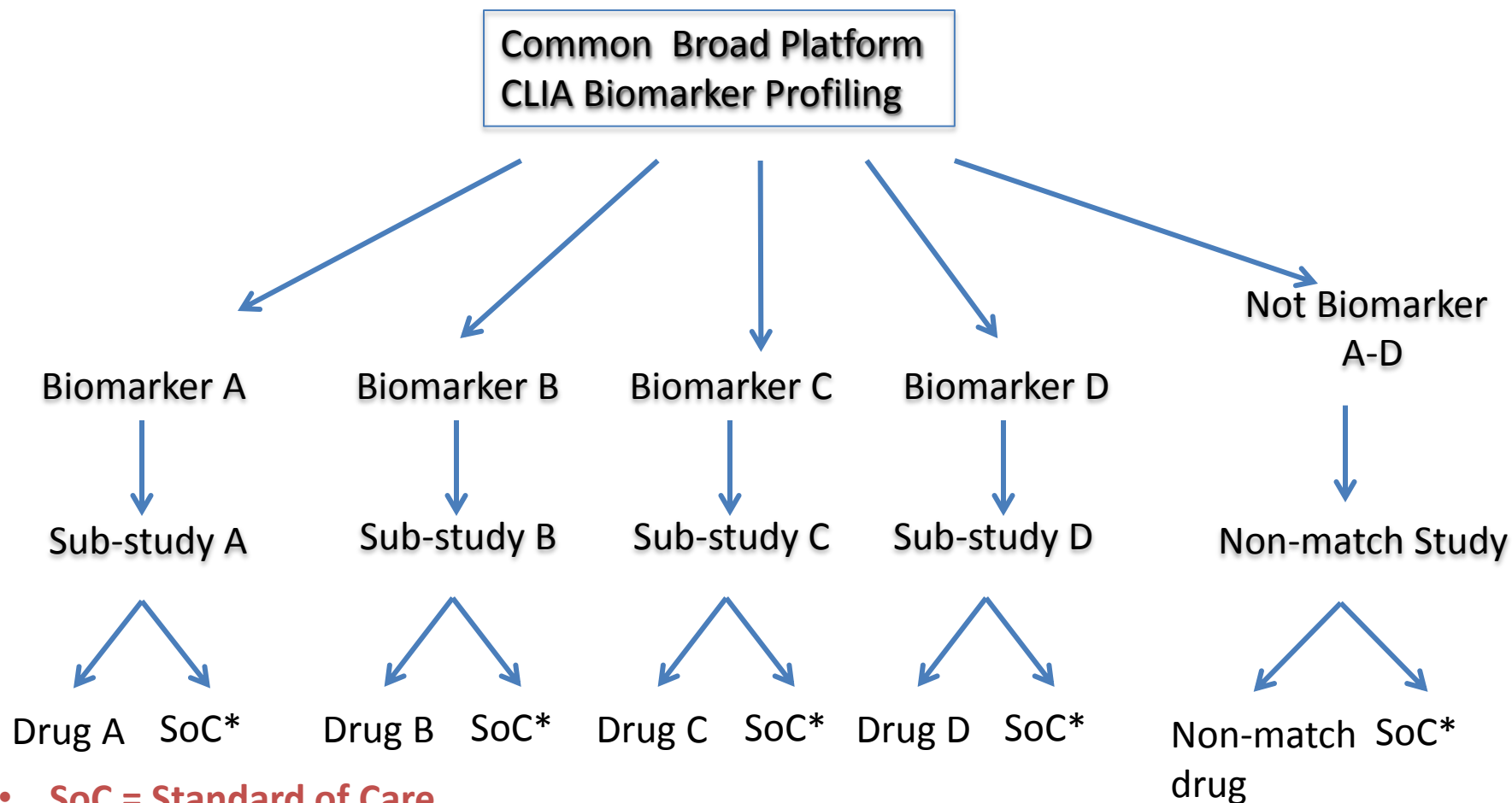


ALK-prevalence ~ 5%; EGFR mutation prevalence ~10-15%

Phase II and III Design Considerations

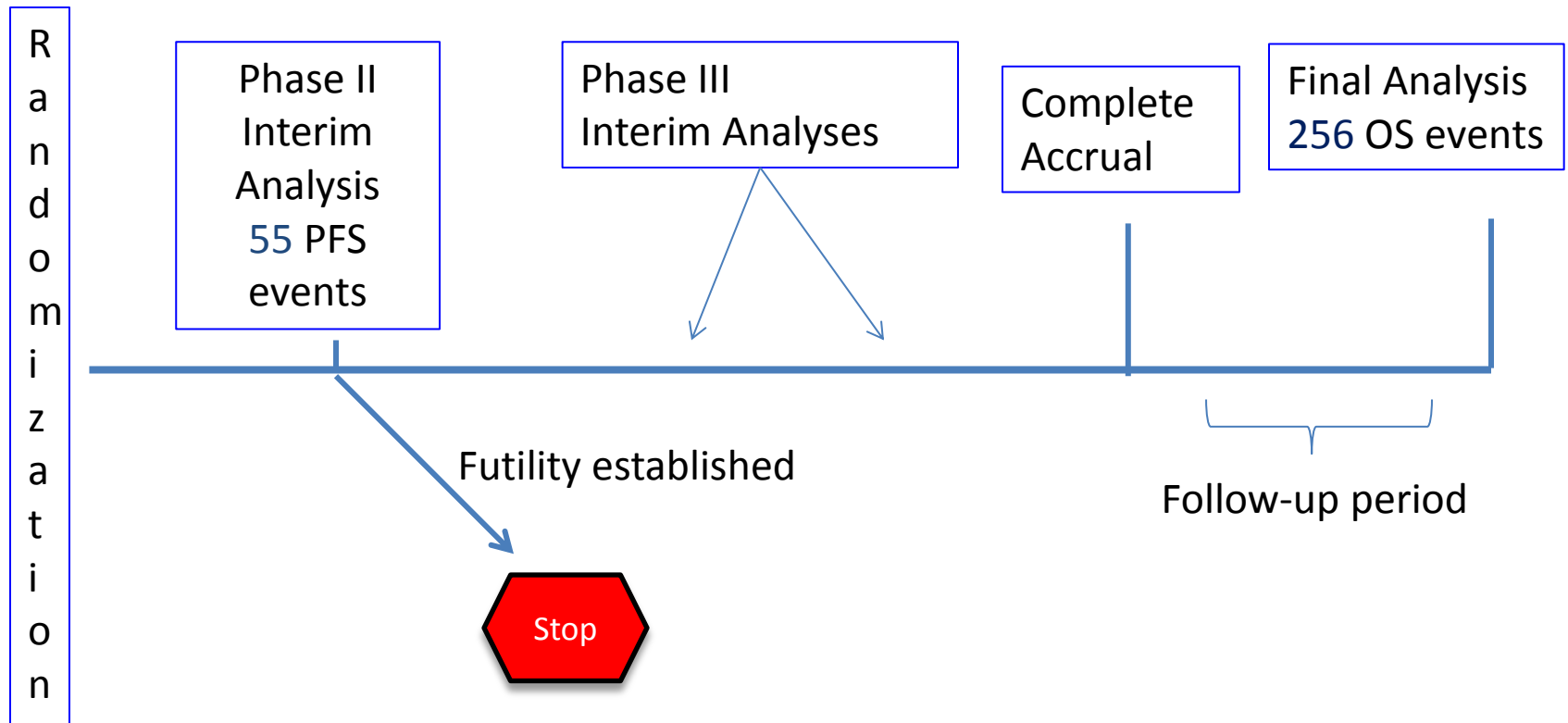
Multi marker scenario

SWOG S1400 Master Lung Protocol Design: Lung-MAP



- **SoC = Standard of Care**
- **Experimental drug could be single agent or a combination; SoC can vary by biomarker.**
- **Patients with multiple markers assigned randomly to a sub study: randomization ratio matching marker prevalence**

Phase II/III Design: SWOG S1400



PFS: Primary endpoint for Phase II
OS: Primary endpoint for Phase III

The NEW ENGLAND JOURNAL *of* MEDICINE

ESTABLISHED IN 1812

MARCH 8, 2012

VOL. 366 NO. 10

Intratumor Heterogeneity and Branched Evolution Revealed by Multiregion Sequencing

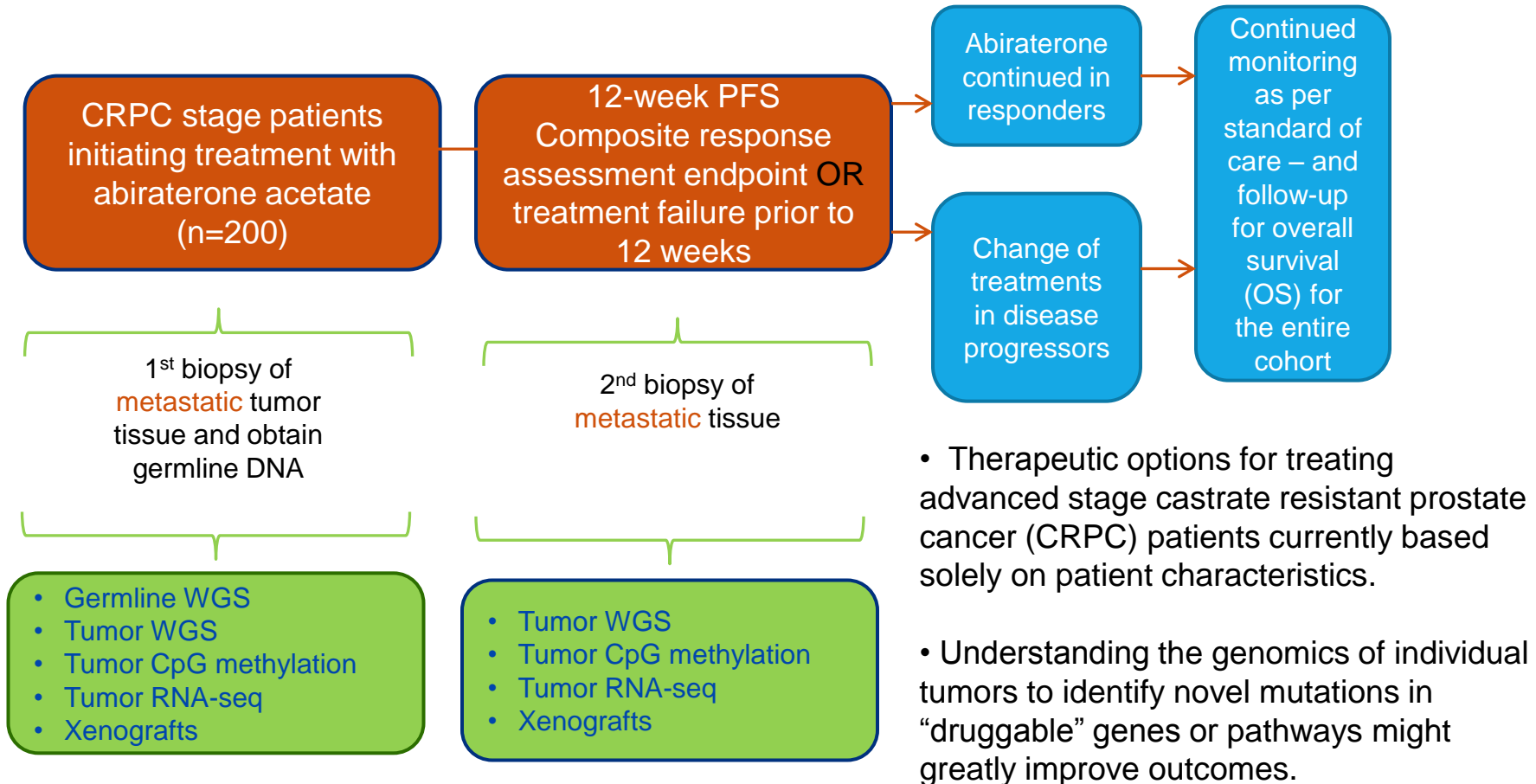
Marco Gerlinger, M.D., Andrew J. Rowan, B.Sc., Stuart Horswell, M.Math., James Larkin, M.D., Ph.D., David Endesfelder, Dip.Math., Eva Gronroos, Ph.D., Pierre Martinez, Ph.D., Nicholas Matthews, B.Sc., Aengus Stewart, M.Sc., Patrick Tarpey, Ph.D., Ignacio Varela, Ph.D., Benjamin Phillimore, B.Sc., Sharmin Begum, M.Sc., Neil Q. McDonald, Ph.D., Adam Butler, B.Sc., David Jones, M.Sc., Keiran Raine, M.Sc., Calli Latimer, B.Sc., Claudio R. Santos, Ph.D., Mahrokh Nohadani, H.N.C., Aron C. Eklund, Ph.D., Bradley Spencer-Dene, Ph.D., Graham Clark, B.Sc., Lisa Pickering, M.D., Ph.D., Gordon Stamp, M.D., Martin Gore, M.D., Ph.D., Zoltan Szallasi, M.D., Julian Downward, Ph.D., P. Andrew Futreal, Ph.D., and Charles Swanton, M.D., Ph.D.

Intratumor heterogeneity may foster tumor evolution and adaptation and hinder personalized-medicine strategies that depend on results from single tumor-biopsy samples.

Once you start studying medicine you never get through with it
--Dr. Charles H. Mayo

PROstate cancer Medically Optimized genome enhanced Therapy – I (PROMOTE)

PI: Dr. Kohli



PROMOTE-II Design based on PROMOTE-I

- Compelling evidence: Enrichment design
- Fairly strong, but not compelling evidence: Biomarker Stratified design
- Evidence preliminary and exploratory: adaptive design

Treatment of Platinum resistant Ovarian Cancer (PI: Dr. Haluska)

- Avatars generated at the time of surgery.
- Upon engraftment, the avatars would be expanded in platinum-chemotherapy to develop platinum resistant disease.
- Upon regrowth (typically 4-8 weeks), randomized to one of 4 salvage regimens.

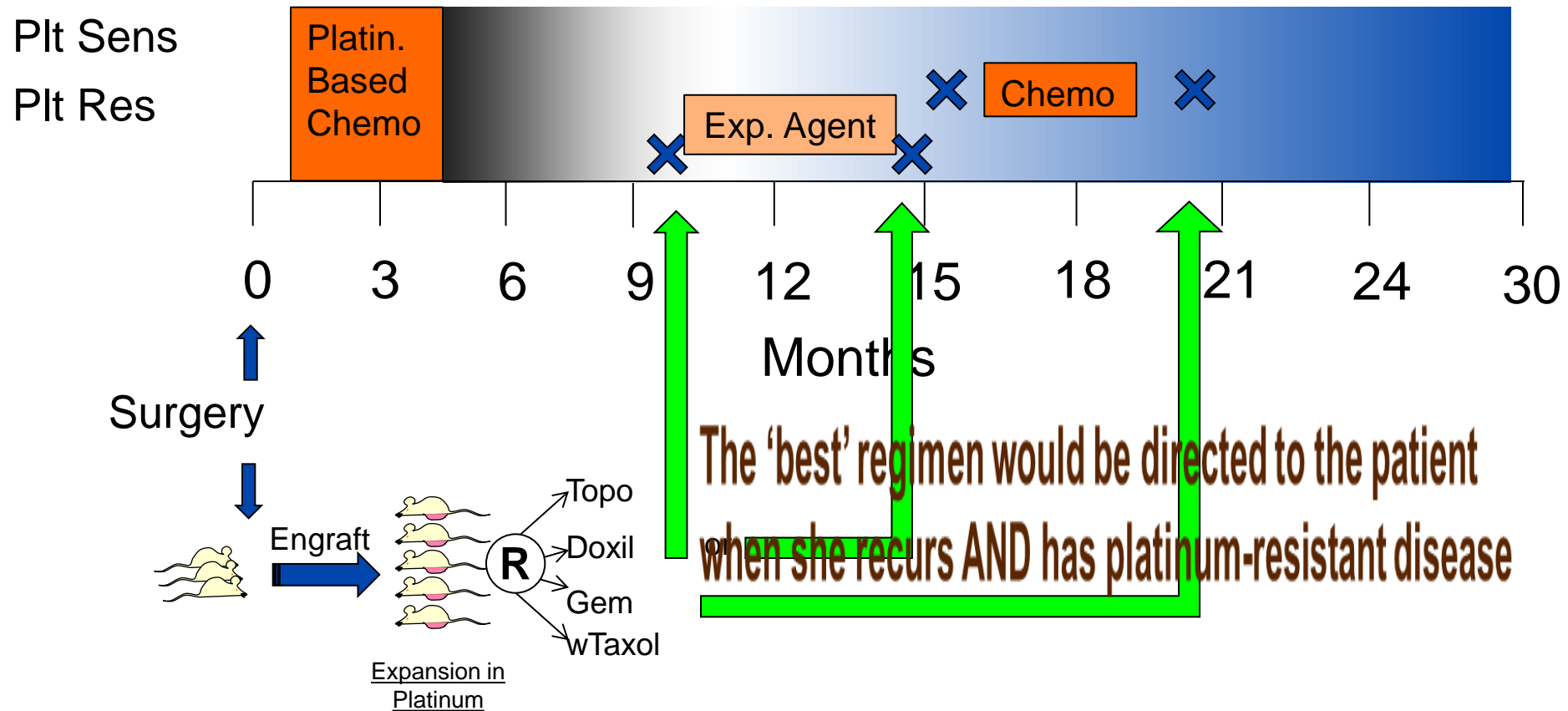


Figure 1. Marker Strategy Design
 Figure 1. Schematic of hypotheses for prognostic and predictive behavior of Avatar

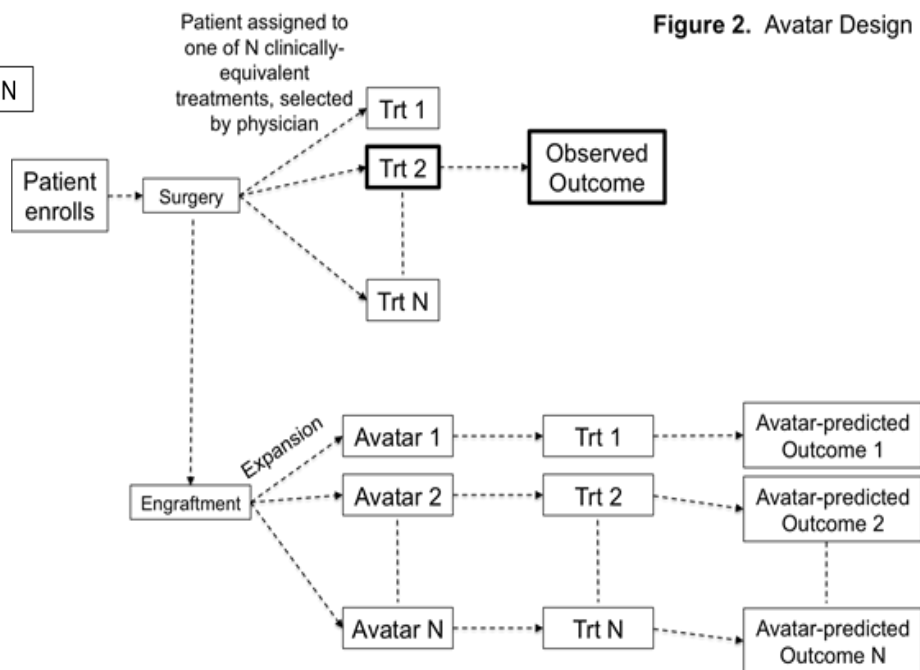
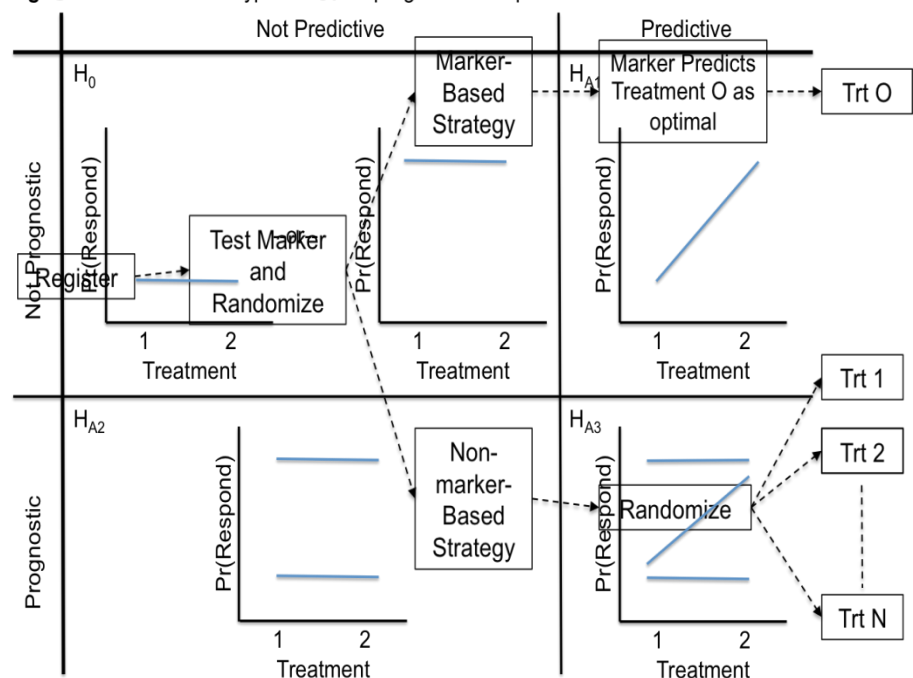


Figure 2. Avatar Design

Overall Design Strategy Recommendations

- Phase I: No restrictions
 - Use expansion cohorts to further understand marker-subgroup effects, endpoints etc.
- Phase IIa (optional): Single arm, enriched
 - Proof of concept
- Phase IIb: Randomized phase II unselected
 - Primary comparison: Marker (+)
 - Randomize enough Marker (-) to demonstrate lack of benefit
 - Consider adaptive designs
- Phase III: Based on randomized phase II
 - Enrichment, all-comers, marker-stratified, marker-strategy, adaptive

Important Considerations

Integral Biomarker Studies

- Strength of pre-clinical evidence of the marker
 - Restrict patients based on marker status or enroll all patients regardless of the marker status?
- Reproducibility and validity of assays
 - Local versus Central Testing
- Prevalence of the marker
 - Low versus moderate
 - Threshold for cut offs; detection limits?
- Feasibility and timing of biomarker assessments
 - Multiple biopsies: pre and post treatment
- **Key Message: You cannot have many moving parts or unknowns in the design of a trial**



THANK YOU FOR YOUR ATTENTION

Sumithra J. Mandrekar
mandrekar.sumithra@mayo.edu