

Design Concept for a Confirmatory Basket Trial

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 - **Cong Chen—led group; co-led concept development; led all statistical and simulation work for initial design**
 - Zoran Antonijevic, Amgen
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- Pathway design subgroup, additional members:
 - Christine Gause, Merck
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 - Sammy Yuan, Merck
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 - Advisor: Sue-Jane Wang
- Pathway design subgroup is one of 5 working subgroups of the **DIA Small Populations Workstream**, a group of 50 statisticians and clinicians from industry, academia, and national health authorities (FDA and EMEA)
- Small populations workstream is part of **DIA Adaptive Design Scientific Working Group (ADSWG)**, a group of > 200 statisticians and clinicians from industry, academia, and national health authorities (FDA and EMEA)

Joining DIA working groups

- Adaptive design scientific working group:
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- Bayesian scientific working group: email Fanni Natanegara: natanegara_fanni@lilly.com

Non-Oncology Rare Diseases

- Estimates of rare diseases
 - Up to 7000 different rare diseases
 - Estimated 30 million sufferers in US alone
 - About half are children
 - Many of these diseases are progressive, debilitating, and lethal

Small Populations Within A Common Disease, Cancer

- ▶ The increasing discovery of molecular subtypes of cancer leads to small subgroups that actually correspond to orphan or “niche” indications, even within larger tumor types
- ▶ Enrolling enough patients for confirmatory trials in these indications may be challenging.
- ▶ The shift to a molecular view of cancer requires a corresponding paradigm shift in drug development approaches
- ▶ Exclusive use of “one indication at a time” approaches will not be sustainable

Approaches to development based on predictive biomarkers

- ▶ Optimized co-development of a single drug and its companion diagnostic
 - Gives a clear hypothesis and answer and still has a role in selected instances
 - Will be challenging to do in niche indications
- ▶ “Umbrella” trials
 - One tumor type with multiple drugs and predictive biomarkers
 - Patients are matched to drugs based on predictive biomarkers
 - Cooperation among multiple sponsors
 - Examples: BATTLE, I-SPY, Lung-MAP

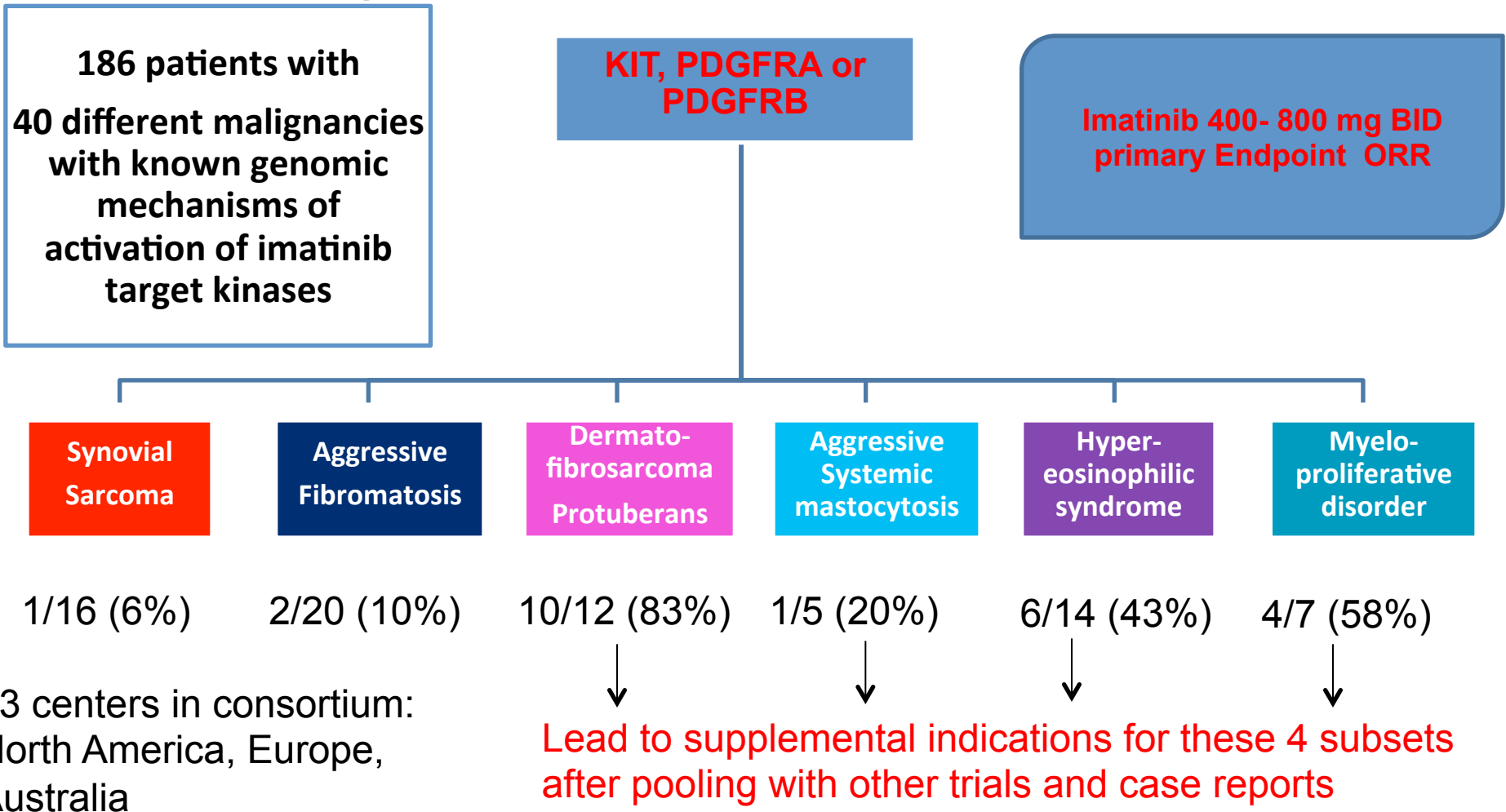
Basket Trials

- ▶ Multiple tumor types with one drug and predictive biomarker
- ▶ Evaluation often based on pooled analysis
 - In some designs, pooling can be partial, based on Bayesian hierarchical model. Degree of pooling can be adjusted based on data
 - In some designs, indications are considered individually. Basket is then more of an operational tactic
- ▶ Premise is that molecular subtype is more fundamental than histology
- ▶ Can be single sponsor or consortium
- ▶ Opportunity for multiple indications for the sample size of one: dramatic potential patient and resource savings

Agenda

- ▶ Introduction
- ▶ General Design Concept for a Confirmatory Basket Trial
- ▶ Challenges and Recommendations for Overcoming Them
- ▶ Performance Simulations and Design Considerations
- ▶ Conclusions

The Original Basket: Imatinib B2225



13 centers in consortium:
North America, Europe,
Australia

Pembrolizumab Approval in MSI-High in Solid Tumors

- MSI High is a DNA repair defect that leads to an increased mutational burden within affected tumor cells
- Enhanced mutational burden is believed to create neoantigens, leading to enhanced immunogenicity and greater chance of benefit from a PD-1/PD-L1 antagonist
- 5 single arm studies were pooled, 149 total patients
- Primary endpoint was ORR: 39.6%, 78.8% duration > 6 months
- Most patients had colorectal cancer
- 10 other solid tumor types had between 1 and 5 responses
- Approval was for any solid tumor that was MSI-High

Features of These Designs

- A similar design to Imatinib B2225 was endorsed at a Brookings/Friends Conference in 2011
- Common features:
 - Exploratory and opportunistic in nature
 - Single-arm trials with ORR as primary endpoint
 - Often intend to use pooled population for primary analysis to gain broader indication across tumor types (individual tumor type is not adequately powered)
 - Involve possibly transformative medicines in patients with great unmet need and seemingly exceptionally strong scientific rationale

Issues

- Clinical data to support pooling may be limited, and treatment effect may differ between tumor types
 - Vemurafenib works in melanoma with BRAF V600E mutation but not colorectal cancer with same mutation
- Not all drugs hoped to be transformational live up to this promise
- Response rate may not predict overall survival
- Single arm trials are subject to patient selection bias
- Predictive effect of a biomarker is confounded with the prognostic value which is often unknown
- Health authorities can be non-committal upfront

DIA Small Population Pathway Subteam

- Can we develop a **generalizable confirmatory basket design concept with statistical rigor?**
 - Applicable not only to exceptional cases, but to all effective medicines in any line of therapy
 - Follow existing accelerated and standard approval pathways to increase **potential approvability**
- This would have multiple benefits
 - Increase and accelerate access to effective medicines for patients in niche indications
 - Provide sponsors with cost-effective options for development in niche indications
 - Provide health authorities with more robust packages for evaluation of benefit and risk
 - **MOST OF DRUG DEVELOPMENT RESOURCES ARE SPENT IN THE CONFIRMATORY PHASE**

Most of Drug Development Resources Are Spent in the Confirmatory Phase



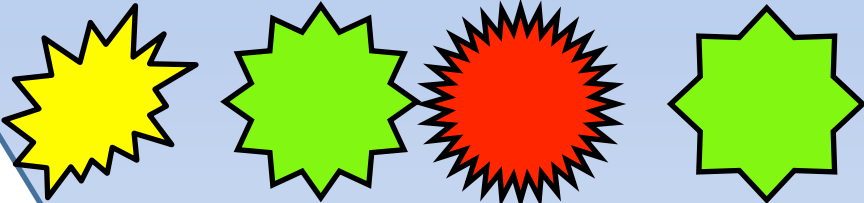
GENERAL DESIGN CONCEPT

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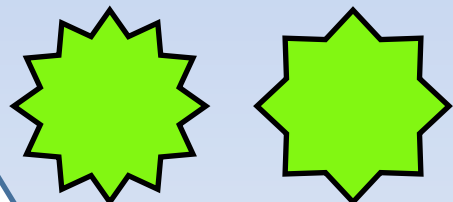
SELECTION



PRUNING
(External Data)



PRUNING
(Interim endpoints)



Accelerated
Approval
Option

Consistent trend in
definitive endpoint

FULL APPROVAL
(Pooled analysis of
definitive endpoint)

Features of the Design (I)

- ▶ Tumor histologies are grouped together, each with their own control group (shared control group if common SOC)
- ▶ Randomized control is preferred
 - Single arm cohorts with registry controls may be permitted in exceptional circumstances as illustrated by imatinib B225 and others
- ▶ In an example of particular interest, each indication cohort is sized for accelerated approval based on a surrogate endpoint such as progression free survival (PFS)
 - This may typically be 25-30% of the size of a Phase 3 study
- ▶ In another approach, an interim evaluation of partial information on the definitive endpoint may be used
- ▶ Initial indications are carefully selected as one bad indication can spoil the entire pooled result

Features of the Design (II)

- ▶ Indications are further “pruned” if unlikely to succeed, based on:
 - External data (maturing definitive endpoint from Phase 2; other data from class)
 - Internal data on surrogate endpoint OR partial information on definitive endpoint
- ▶ Sample size of remaining indications may be adjusted based on pruning
- ▶ Type I error threshold will be adjusted to control type I error (false positive rate) in the face of pruning
 - Pruning based on **external** data does not incur a statistical penalty
 - Discussed in more detail later in talk
- ▶ Study is positive if pooled analysis of remaining indications is positive for the primary definitive endpoint
 - Remaining indications are eligible for full approval in the event of a positive study
 - Full pooling chosen for simplicity
 - Some of the remaining indications may not be approved if they do not show a trend for positive risk benefit as judged by definitive endpoint

CHALLENGES OF BASKET DESIGNS AND RECOMMENDATIONS FOR OVERCOMING THEM

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Challenge #1: Having a Control Group

- In some settings, a control group is not ethical
- Resolution: randomized trial should be applied, if and only if:
 - There is clinical equipoise between the two randomized arms
 - Experimental agent is not expected to be transformational, only beneficial
 - There is an SOC for control
 - Example: steroids +/- rituximab for refractory autoimmune diseases
- Current generation of non-randomized basket studies for transformational agents provides SOC baseline for future randomized trials

Challenge 2: Risks of Pooling

- ▶ One of more bad indications can lead to a failed study for all indications in a basket
- ▶ Histology can affect the validity of a molecular predictive hypothesis, in ways which cannot always be predicted in advance
 - Vemurafenib is effective for BRAF 600E mutant melanoma, but not for analogous colorectal cancer (CRC) tumors
 - This was not predicted in advance but subsequently feedback loops leading to resistance were characterized

Addressing challenge 2

- ▶ Basket trials are recommended primarily after there has been a lead indication approved (by optimized conventional methods) which has validated the drug, the predictive biomarker hypothesis, and the companion diagnostic
 - Example, melanoma was lead indication preceding Brookings trial proposal in V600E mutant tumors
- ▶ Indications should be carefully selected
- ▶ Indications should be pruned in several steps before pooling

Challenge 3: Different Indications May Have Different Endpoints

- Less of an issue for oncology
- Even in auto-immune diseases, generalized interim endpoints can be created across diverse diseases:
 - Interim: improvement (response)
 - Final: time to worsening

Challenge 4: Timescales of endpoint development may differ

- Resolution:
 - what matters is ***relative*** improvement
 - If necessary, TTE data may be normalized to the medians on control arms of the different indications
 - study completes when data is mature on all arms

Challenge 5: SOC may differ between arms

- Resolution:
 - what matters is ***relative*** improvement ***in a redefined disease entity based on a molecular biomarker***
 - *safety must be assessed both as an individual analysis relative to individual control **and** as a pooled analysis relative to pooled control*
 - *safety data should be available from reference indications and from phase 2 studies*

Challenge 6: Threshold for Approval May Differ Between Arms

- Resolution: study is judged by pooled result of *relative* improvement with statistical and clinical significance
 - Thresholds for such criteria are well known

Challenge 7: Clinical validity of the predictive biomarker hypothesis

- ▶ The clinical validity of the predictive biomarker can only be verified by inclusion of “biomarker negative” patients in the confirmatory study
- ▶ Addressing the challenge
 - Recommend a smaller pooled, stratified cohort for biomarker negative patients, powered on surrogate endpoint
 - Would need to expand the biomarker negative cohort (to evaluate definitive endpoint) if surrogate endpoint shows possible benefit
 - Prior evidence should permit this if:
 - An approved lead indication has already provided clinical evidence for the predictive biomarker hypothesis
 - Prior phase 2 studies support the predictive biomarker hypothesis in other indications

Challenge 8: Adjusting for Pruning

- ▶ Pruning indications that are doing poorly on surrogate endpoints may be seen as cherry picking
 - This can inflate the false positive rate, an effect termed “random high bias”
- ▶ Addressing the challenge:
 - Emphasize use of **external data**, especially maturing Phase 2 studies, for pruning
 - Pruning with external data does not incur a penalty for random high bias
 - Apply statistical penalty for control of type I error when applying pruning using **internal data**
 - Methods for calculating the penalty are described in stat methods papers (see key references)
 - Rules for applying penalty must be prospective
 - Penalty is not large enough to offset advantages of design

Type I error control under global null hypothesis

- k tumor indications each with sample size of N and all with 1:1 randomization
- An interim analysis is conducted at information fraction t for each tumor indication and a tumor will not be included in the pooled analysis if $p\text{-value} > \alpha_t$
- The pooled analysis will be conducted at α^* so that the overall Type I error is controlled at α when there is no treatment effect for any tumor (H_0)
- What is α^* ?

Solving for adjusted alpha (α^*)

- Let Y_{i1} be the test statistics based on information fraction t , and Y_{i2} be the test statistics based on the final analysis of data in the i -th cohort ($i=1, 2, \dots, k$)
- Suppose that m cohorts are included in the final analysis ($m \geq 1$), and let V_m be the corresponding test statistics. The probability of a positive outcome in pooled analysis is

$$Q_0(\alpha^* | \alpha_t, m) = \Pr_{H_0} (\cap \{Y_{i1} > Z_{1-\alpha_t} \text{ for } i=1, \dots, m\}, \cap \{Y_{j1} < Z_{1-\alpha_t} \text{ for } j=m+1, \dots, k\}, V_m > Z_{1-\alpha^*})$$

or
$$Q_0(\alpha^* | \alpha_t, m) = \Pr_{H_0} (\cap \{Y_{i1} > Z_{1-\alpha_t} \text{ for } i=1, \dots, m\}, V_m > Z_{1-\alpha^*}) (1 - \alpha_t)^{(k-m)}$$

- α^* is solved from below where $c(k, m) = k! / ((k-m)! m!)$

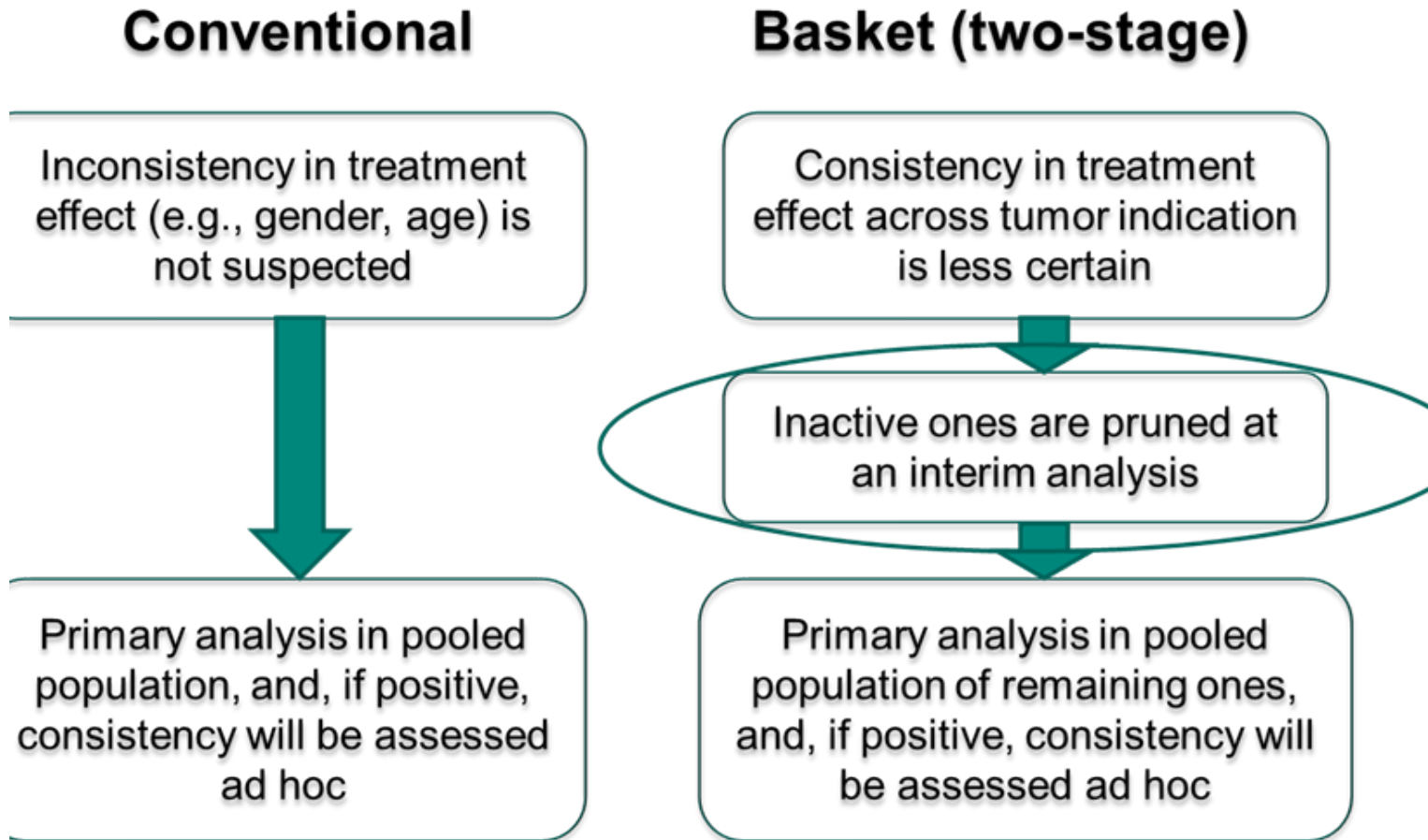
$$\sum_{m=1}^k c(k, m) Q_0(\alpha^* | \alpha_t, m) = \alpha$$

Challenge #9: Strong Control of FWER

- This problem is still open
- Challenge:
 - One or more strongly positive indications can drive an overall pooled positive result and negative indications are carried along
 - Simulation involves a large number of cases and the degree to which active indications are active affects the results
- A recent MSKCC study* simulated a popular Bayesian basket trial design (using a Bayesian hierarchical model) and found FWER of up to 57%.
 - Authors advocate characterization of FWER by simulation

*Cunanan K et al. Specifying the True- and False-Positive Rates in Basket Trials, *J Clin. Onc. Precision Onc.* , published online November 3, 2017

Should Basket Trials Control FWER by Indication?



Other FWER Considerations

- A basket trial with k indications replaces k independent trials that collectively would have a family-wise error rate of approximately $k * 0.025$
- Should we therefore allow approximately $k*0.025$ for FWER of a basket trial?
- Under would conditions would FDR be a better measure than FWER?

Challenge #10: Availability of tissue

- ▶ Tissue sampling and processing are variables that can greatly affect the outcome of a study based on a predictive biomarker
- ▶ Basket studies will require cooperation and uniformity across departments organized by histology
- ▶ Addressing the challenge:
 - The sponsor must have extensive contact with the pathology department and relevant clinical departments at all investigative sites and provide standard methods for tissue sampling, handling, and processing
 - The sponsor should engage an expert pathologist who is dedicated to training prior to trial start, and troubleshooting during the trial

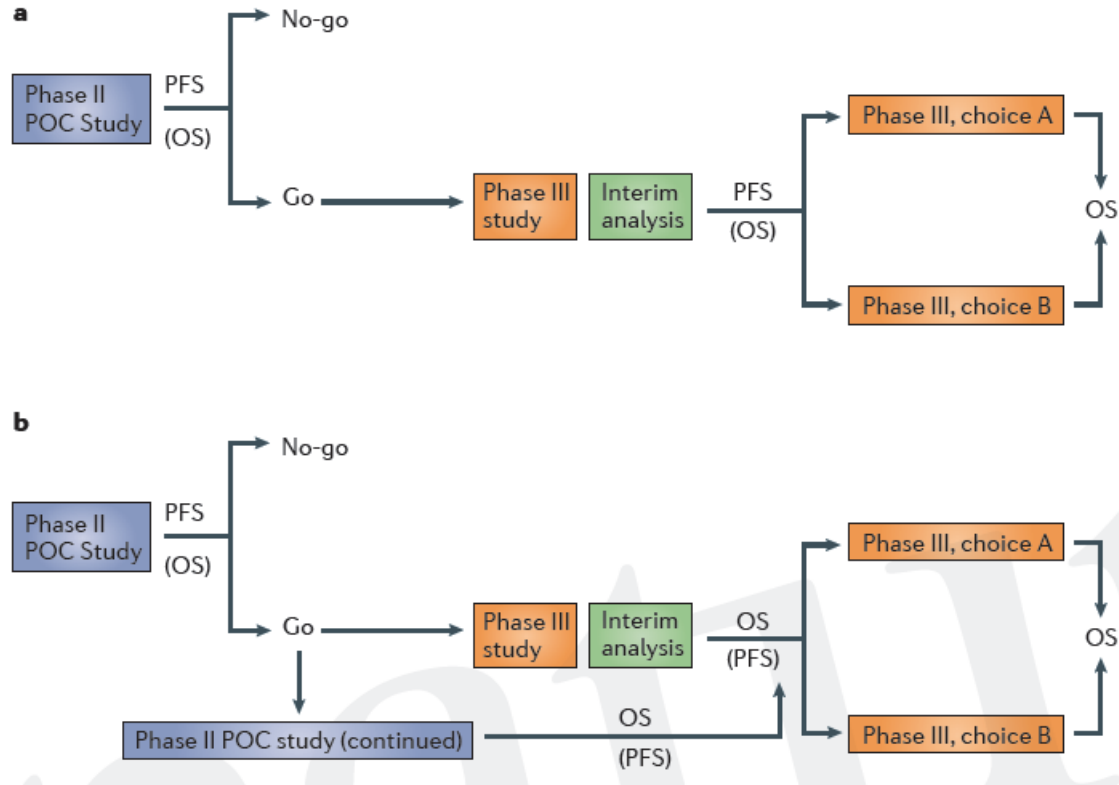
Challenge #11: High Screen Failure Rate

- ▶ Pro: patients will have access to tailored therapy
- ▶ Con: patient has a high risk of being a screen failure if biomarker positive subgroup is low prevalence
- ▶ Addressing the challenge:
 - Study should provide a broad-based test like NGS which will give the patient some guidance on alternative therapies if they are screen failures for basket study

Challenge #12: Interim endpoints may not predict definitive endpoints

- ▶ Addressing the challenge:
 - Prefilter indications based on maturing definitive endpoint data from phase 2 or other external data
 - See Figure 2
 - Require consistent trend in definitive endpoint for final full approval

Phase 2 Influencing Phase 3 Adaptation: The Phase 2+ Method



Beckman, R.A., Clark, J. & Chen, C. Integrating predictive biomarkers and classifiers into oncology clinical development programmes. *Nature Reviews Drug Discovery* **10**, 735-748 (2011)

Another Possible Source of External Data

- Real World Data (RWD) from Off-Label Use
- Impact of RWD on basket trial performance is currently under study in a project led by postdoctoral fellow Daphne Guinn



PERFORMANCE SIMULATIONS AND DESIGN CONSIDERATIONS

December 4, 2018

Comparison of operating characteristics

- $k=6$ tumor indications with total planned event size (kN) ranging from 150-350
 - The true treatment effect is $-\log(0.6)$, or hazard ratio of 0.6 in a time-to-event trial
- Pruning occurs at when half of the events have occurred
- Number of active indications (g) with target effect size ranges from 3 to 6, with remaining ones inactive

Study power and sample sizes under different pruning and pooling strategies

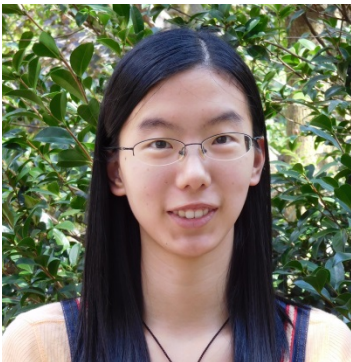
Planned events	Number of active tumors	Power (%) for a positive study				Exp. number of events for pooled population			Exp. number of events for overall study		
		D0	D1	D2	D3	D0/D2	D1	D3	D0/D3	D1	D2
200	6	95	85	95	93	200	157	179	200	179	221
200	5	85	75	91	86	200	144	172	200	172	228
200	4	67	62	82	76	200	131	166	200	166	234
200	3	44	45	68	61	200	119	159	200	159	240
300	6	99	96	99	99	300	254	277	300	277	323
300	5	96	81	98	96	300	232	266	300	266	334
300	4	84	81	94	91	300	209	255	300	255	345
300	3	60	64	84	79	300	187	244	300	244	356

An Application of Special Interest

- A randomized controlled basket trial with 1:1 randomization in 6 tumor indications, each targeting a hazard ratio of 0.5 in PFS with 90% power at 2.5% alpha for global null hypothesis
 - 88 PFS events and 110 patients planned for each indication
 - PFS analysis is conducted when all are enrolled
- D2 is applied to keep total sample size at 660 in pooled population targeting 430 death events
 - The study has ~90% power to detect a hazard ratio of 0.7 in OS at 0.8% alpha (after taking the penalty) assuming $\rho=0.5$
 - Observed hazard ratio ~0.79 or lower for a positive trial in pooled population (vs ~0.84 under D0) for alpha control **under global null**
- Potential to gain approvals in 6 indications based on comparable sample size to a conventional Phase 3 trial

Characterization of Performance Constrained by FWER (ongoing)

- Team includes Yuru Ren, Valeriy Korostyshevskiy, and Sammy Yuan
- Currently studying single TTE endpoint with normally distributed hazard ratios, mean of 1.0 for inactive, 0.7 for active
- Simulate different scenarios of how many indications in basket are inactive. Maximum Type I error (worst case scenario) is FWER
- What power is achievable when FWER must be $\leq k * 0.025$?



Current Approaches

- In order to control FWER, we must add an additional post-correction step
- Each indication is tested up to twice individually*
 - at interim information time t $[0,1]$ at significance level $\alpha-t$ to govern pruning, AND
 - if part of a successful pool, in a post check at significance level α -post

*Beckman R and Loeb LA. Multistage Proofreading in DNA Replication. Quarterly Reviews of Biophysics 26: 225-331 (1993)

Preliminary Results

- $k = 6$; HR = 0.7, nominal power of pool = 95%; $t = 0.5$, **alpha t = 0.4, alpha post = 0.1:**

6	0.5276	0.9467	0.0000
5	0.5262	0.8954	0.0631
4	0.5079	0.8032	0.1111
3	0.4654	0.6675	0.1432
2	0.3780	0.4739	0.1451
1	0.2316	0.2402	0.0986

- $k = 3$; HR = 0.7, nominal power of pool = 95%; $t = 0.5$, **alpha t = 0.3, alpha post = 0.1:**

3	0.6940	0.9534	0
2	0.6815	0.8435	0.0493
1	0.5813	0.5871	0.0731

Future Plans

- Further parameter optimization
- Application of heterogeneity detection methods (Simon)
- Study of application with using surrogate interim endpoint
- Application of RWD to study design

Challenge #14: The Standoff

- Health authorities “understandably” won’t commit until given a real example to consider
- Sponsors “understandably” cautious about being first to innovate in confirmatory space
- Resolution:
 - FDA, under **PDUFA VI pilot program**, will be engaging with selected sponsors to bring forward complex innovative designs
 - We must take this risk for our patients



Conclusions

- ▶ It is feasible to create a general design concept for a basket study that is suitable for many agents
- ▶ Multiple challenges can be addressed with careful planning
- ▶ Benefits include:
 - Increased and earlier patient access to targeted therapies for small subgroups
 - Cost-effective methods for sponsors to develop targeted agents in small subgroups
 - More robust datasets for health authorities to assess benefit-risk in these small patient groups

Key References

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