

Strategies for Integrating Predictive Biomarkers into **Clinical Trial Design**: A Changing Paradigm

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Disclosures

- **Research Grants: BMS, Genentech, GlaxoSmithKline, Lilly, Merck, Novartis**
- **Consultant: Ariad, AstraZeneca, Boehringer-Ingelheim, BMS, Celgene, Daiichi-Sankyo, GlaxoSmithKline, Genentech, Lilly, Merck, Novartis, Pfizer, Response Genetics, Synta**

Prognostic versus Predictive Biomarkers

Prognostic Marker

**Information about
disease outcome
independent of treatment**



Predictive Marker

**Information on disease
outcome related to a
specific treatment**

Example : EGFR Mutation in NSCLC

Mutation +: better prognosis

Mutation - : worse prognosis

Example : EGFR Mutation in NSCLC

*Mutation + : ~70% probability of response
to EGFR TKI therapy*

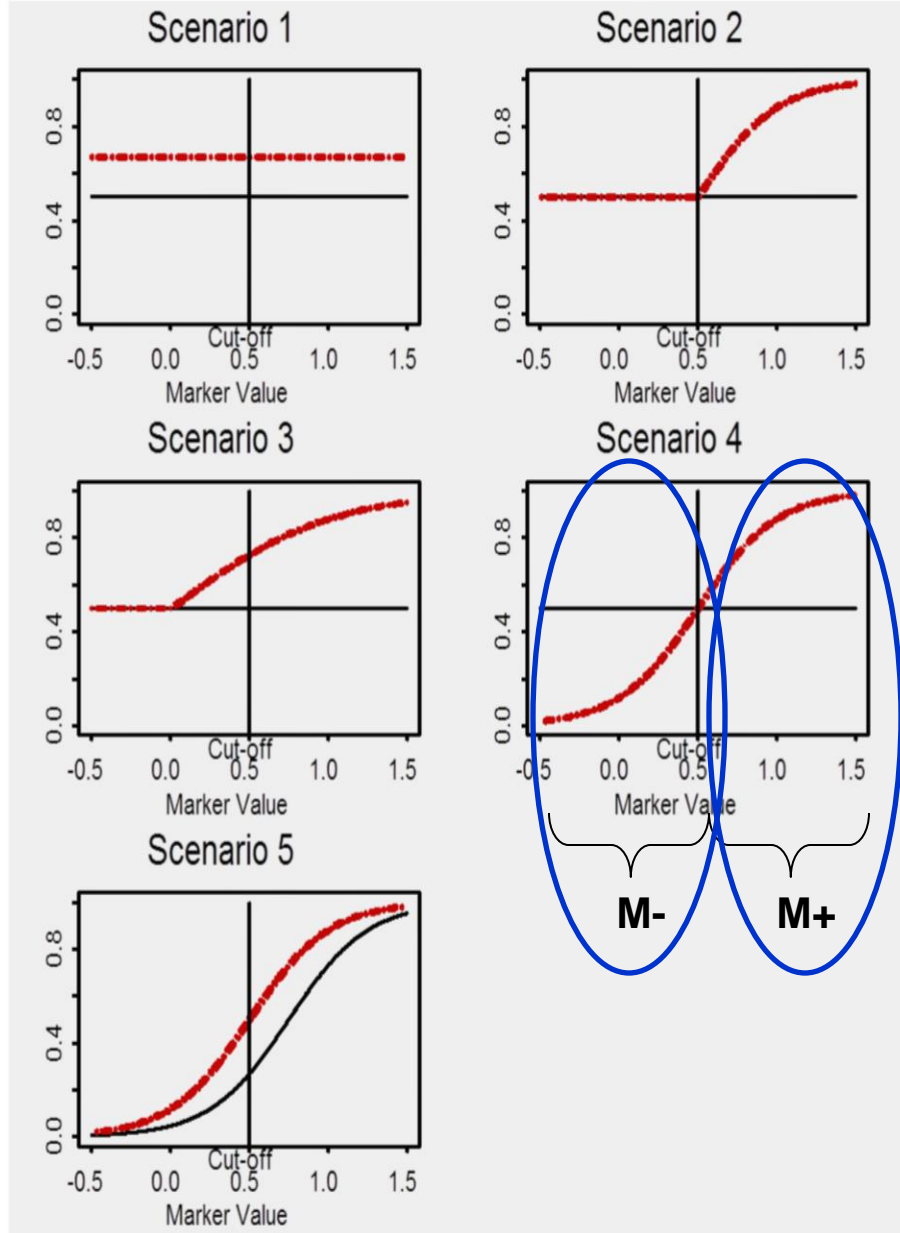
*Mutation - : <5% probability of response
to EGFR TKI therapy*

Some biomarkers are both prognostic & predictive

**Only predictive biomarkers can be used to indicate
“which patients should be treated with which drug”
(a Targeted Therapy)**

**Predictive biomarkers can also identify patients who may be harmed
by “targeted therapy”**

Possible Outcome Scenarios: Marker+ versus Marker-



M+ : Marker positive, Marker value > cut-point

M- : Marker negative, Marker value < cut-point

— T1: Standard Therapy

- - T2: New (Experimental) Therapy

Scenario 1: biomarker is neither prognostic nor predictive

Predictive Markers:

Scenario 2: T2 benefits M+ pts, but not M- pts

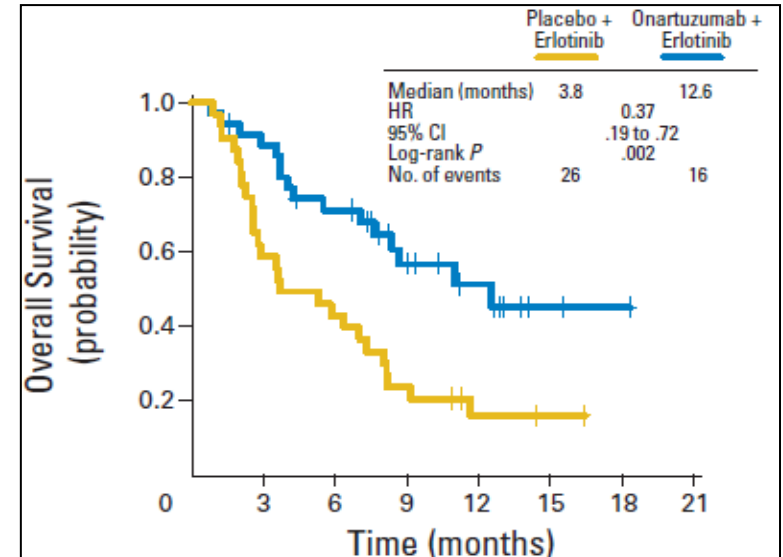
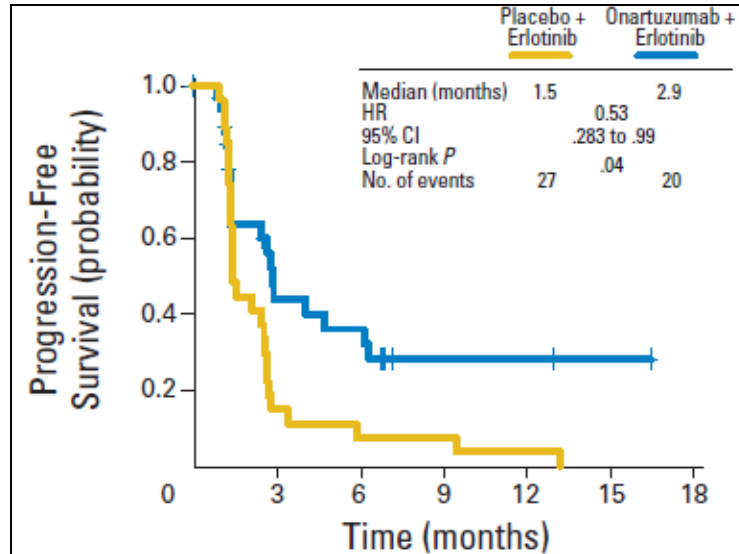
Scenario 3: T2 benefits M+ & M- pts, but effect on M+ pts is more

Scenario 4: T2 benefits M+ pts, but is harmful to M- pts (total interaction)

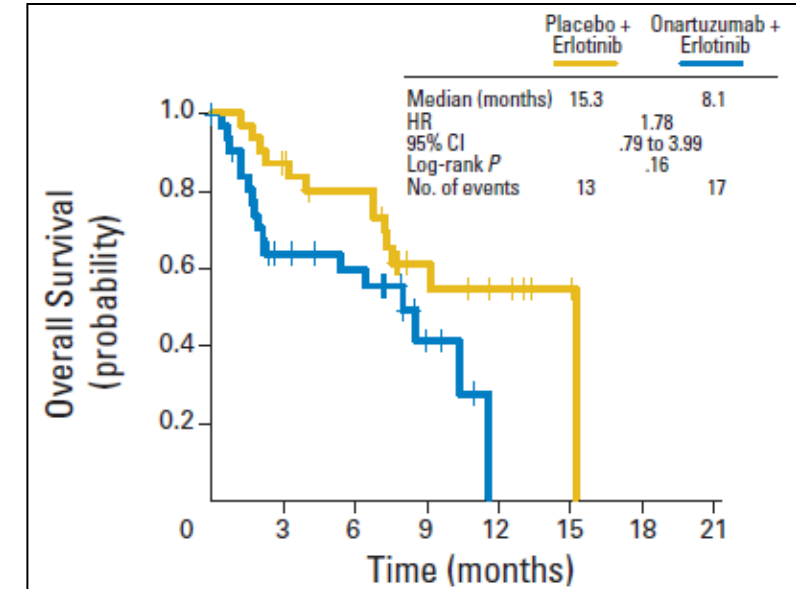
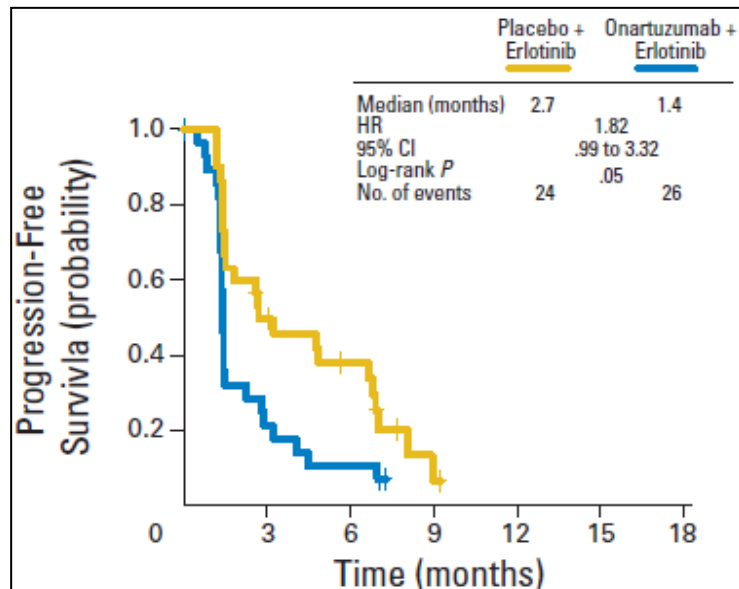
Scenario 5: Prognostic Marker (no predictive value)

Phase II Trial of Erlotinib +/- MetMab: PFS & OS

Met
High

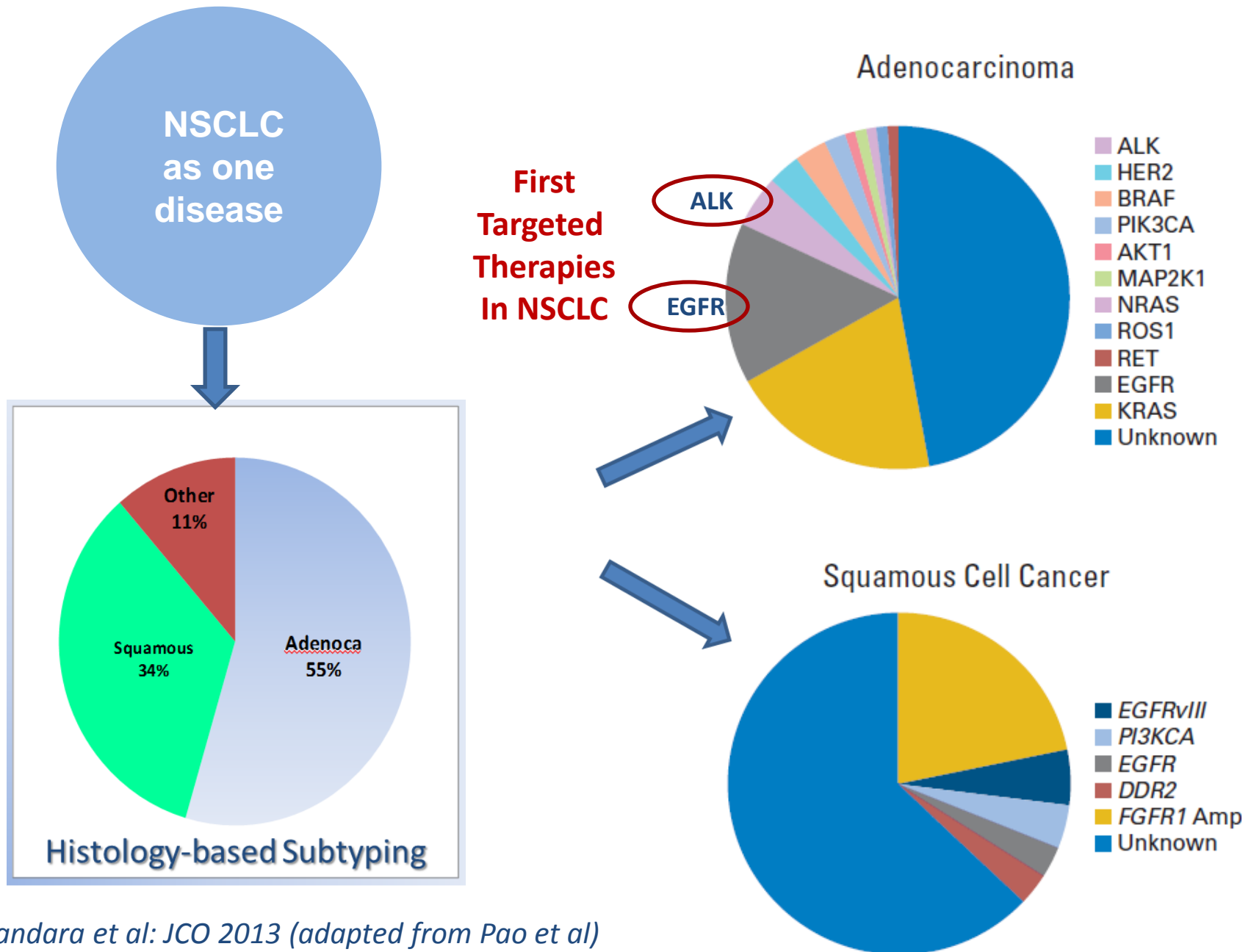


Met
Low

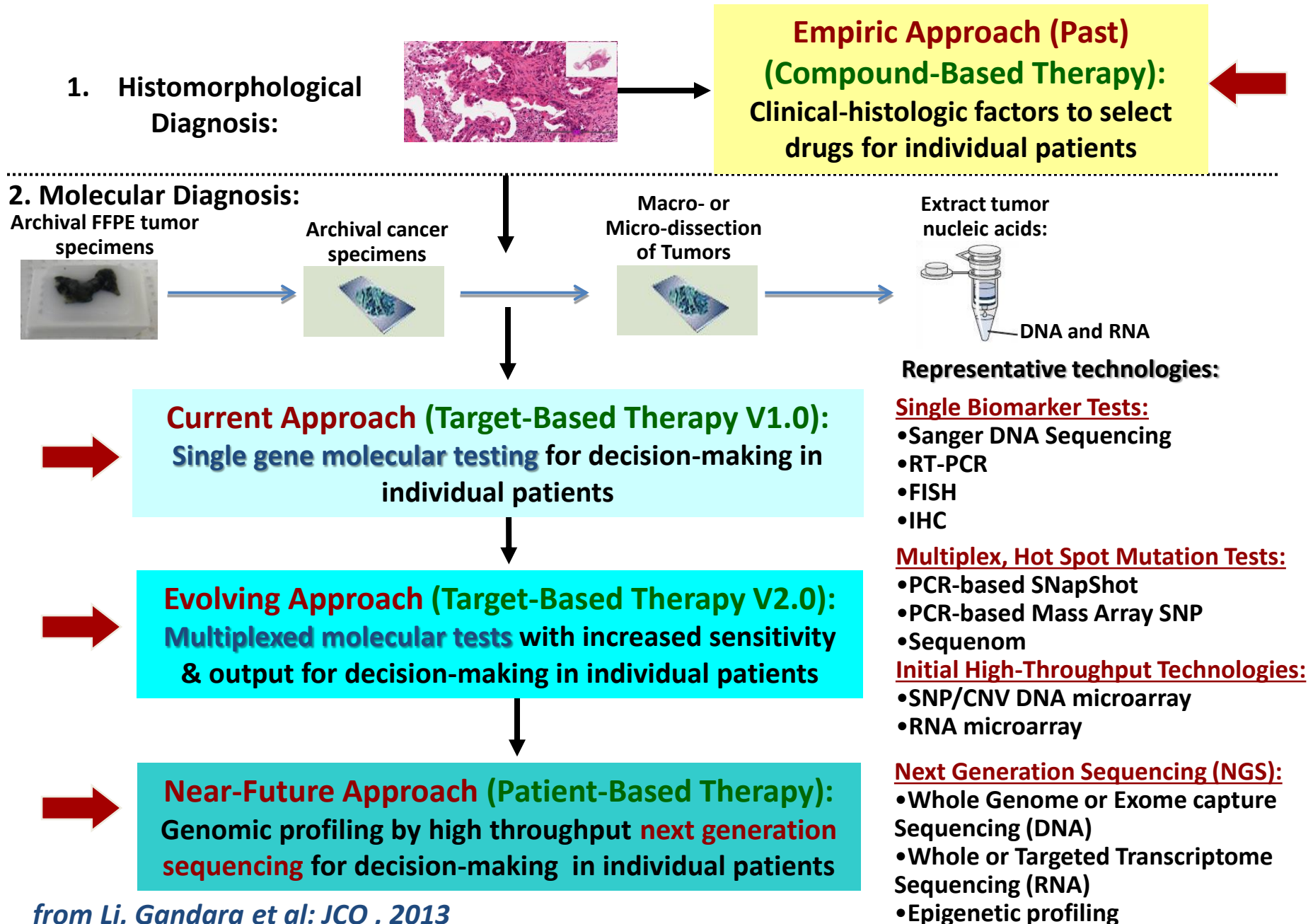


Targeted Agents can do harm in the wrong patient population

Evolution of NSCLC Subtyping from Histologic to Molecular-Based



Integration of Biomarkers into **Clinical Practice**: Past, Current & Future



Need for Paradigm Shift in Targeted Therapy Clinical Trial Design (Presumes Biomarker Potential)

“All Comer” Phase III Design adding Targeted Therapy to Chemotherapy



- When Marker not known or not validated (analytical)
- Marker (if known) can be retrospectively assessed
- Cautionary Tale: Most Phase III “All Comer” trials in NSCLC targeted therapy fail
- May be random differences in Marker+ and Marker- proportions per arm

Classic RCT Design (Unselected): Phase III Trials of Chemotherapy +/- Targeted Agent* in 1st-line Therapy of Advanced Stage NSCLC

Target	Agent	Survival Benefit
MMPs	Prinomastat, Others	No
EGFR TKI	Gefitinib or Erlotinib	No
Farnesyl Transferase (RAS)	Lonafarnib	No
PKC α	ISIS 3521	No
RXR	Bexarotene	No
VEGFR (TKI)	Sorafenib	No
VEGF (Mab)	Bevacizumab	Yes
EGFR (Mab)	Panitumumab	No
TLR9 Agonist	PF-351	No
EGFR (Mab)	Cetuximab	Yes**
IGR1-R	Figitumumab	No
VDA	ASA-404	No

***In combination with platinum-based chemotherapy versus chemotherapy**

****EGFR IHC positive**

from Gandara et al: Clin Lung Cancer, 2012

Biomarker-driven Clinical Trial Designs (selected)

Examples:

Trastuzumab in Breast CA

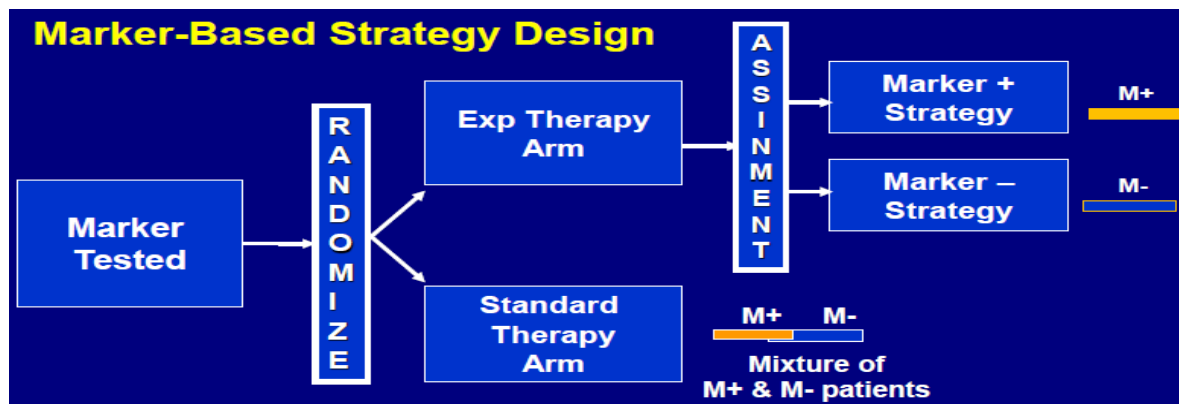
EGFR TKIs in EGFR MT+ NSCLC



Example:

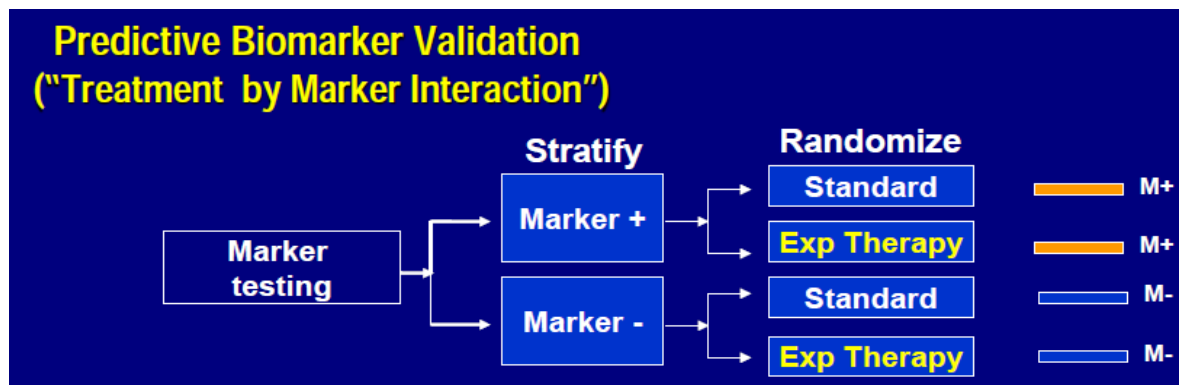
SLCG trial of

ERCC1-driven therapy

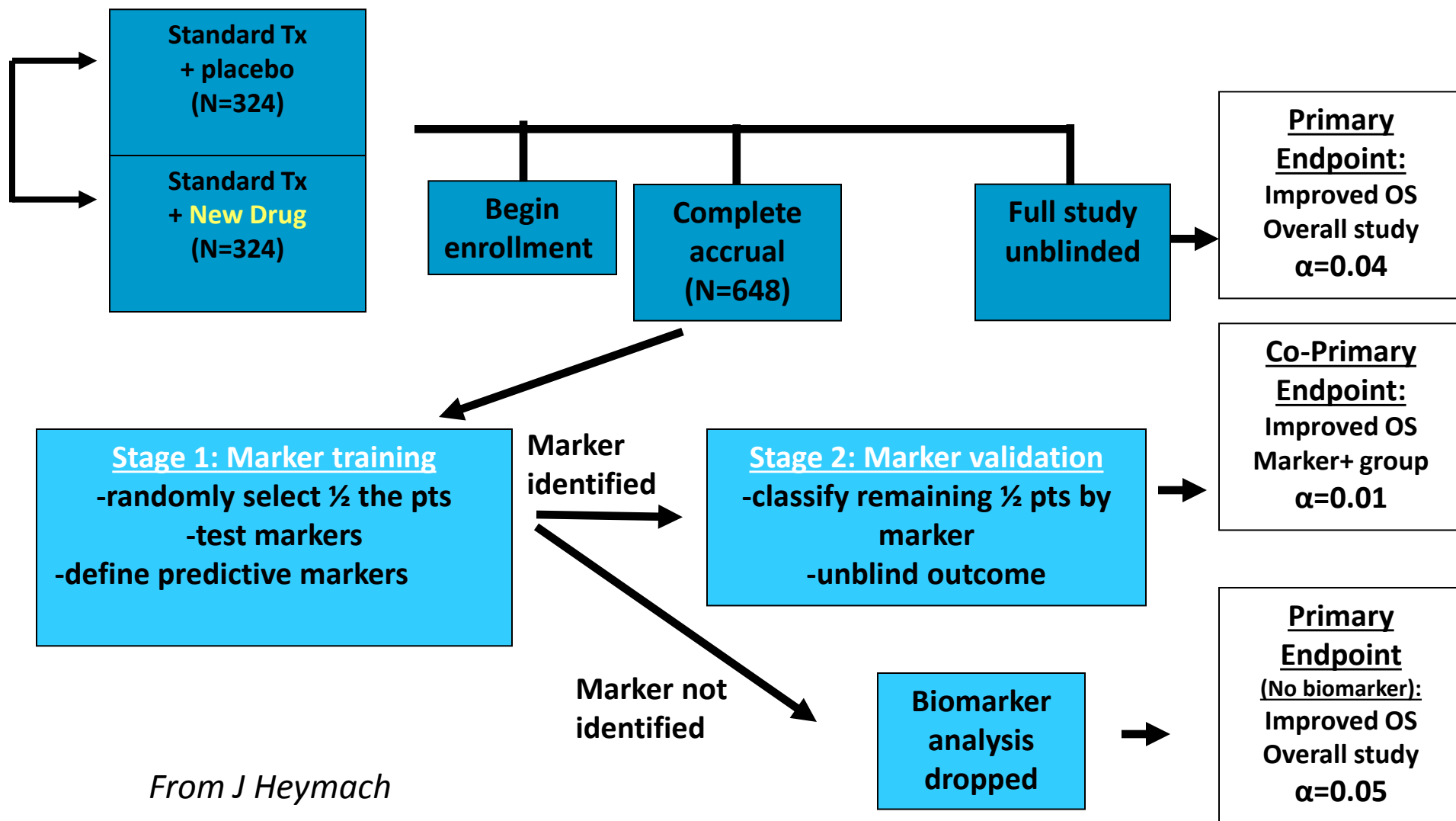


Example:

NCI-MARVEL trial



Phase III Embedded Biomarker Testing & Validation



Integrated **New Drug-New Biomarker** Development Paradigm:

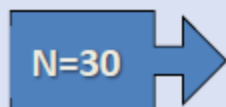
Phases of Development of a New Drug

Pre-clinical



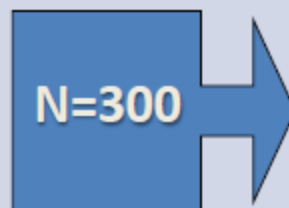
~18 mo.

Phase I



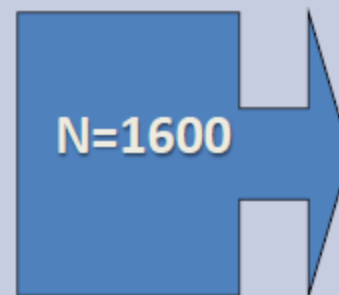
~18 mo.

Phase II



~18 mo.

Phase III



~36 mos

**Drug
Approval**

Total Time
~90 mos
(7.5 years)

Phases of Development of **New Biomarker** linked to New Drug

**Confirm
Target**

Assay
Development

**Integrate
Biomarker**

Assay
Performance

**Biomarker
Informative?**

Assay
Validation

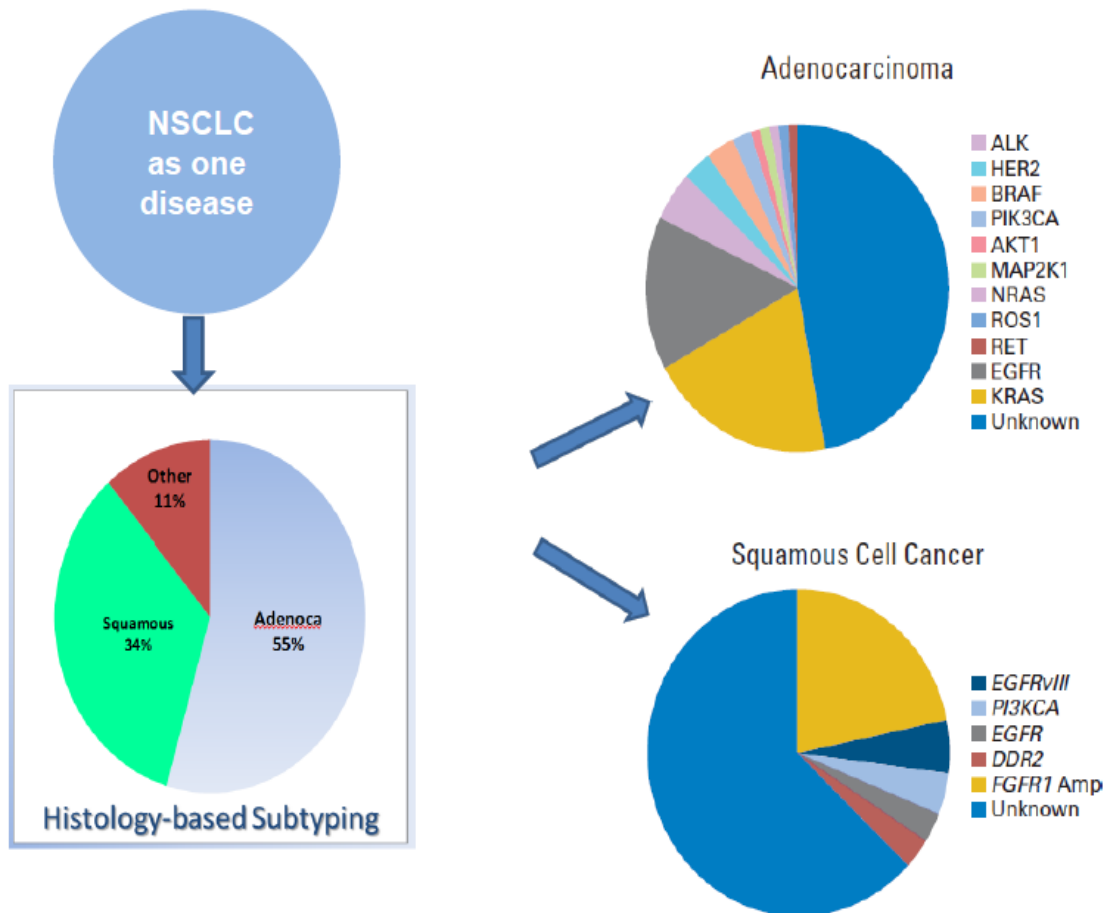
**Clinical
Validation**

Co-Primary
Endpoint

**Clinical
Application
of
Biomarker**

Unmet Needs in Future NSCLC Clinical Trials when viewed as a **Multitude of Genomic Subsets**

Evolution of NSCLC → Histologic Subsets → Genomic Subsets



Unmet Needs in Clinical Trials:

- How to speed drug development for uncommon-rare genotypes?
- How to apply broad-based genomic screening (NGS)?
- How to account for both inter- and intra-tumor heterogeneity in clinical trials?
- How to design trials to circumvent or prevent acquired resistance in oncogene-driven subsets?

“Strategies for Integrating Biomarkers into Clinical Development of New Therapies for Lung Cancer”

**A Joint NCI Thoracic Malignancies Steering Committee-FDA Workshop
Bethesda MD – February 2-3, 2012**

- **Trial Design Challenges in the Era of Biomarker-driven Trials**
 - Innovative Statistical Designs
 - Challenges for Community Oncology Practice participation
 - The Patient Perspective
- **Drug & Biomarker Co-Development in Lung Cancer**
 - Need for Early Co-Development
 - Need for Improved Pre-Clinical Models with clinical relevance
- **Development of Future Lung Cancer Trials**
 - **TMSC Master Protocol Task Force in NSCLC**
 - Biomarker-driven trial designs in both early stage adjuvant therapy & advanced stage NSCLC
 - Account for inter-patient tumor heterogeneity & genomic complexity of NSCLC

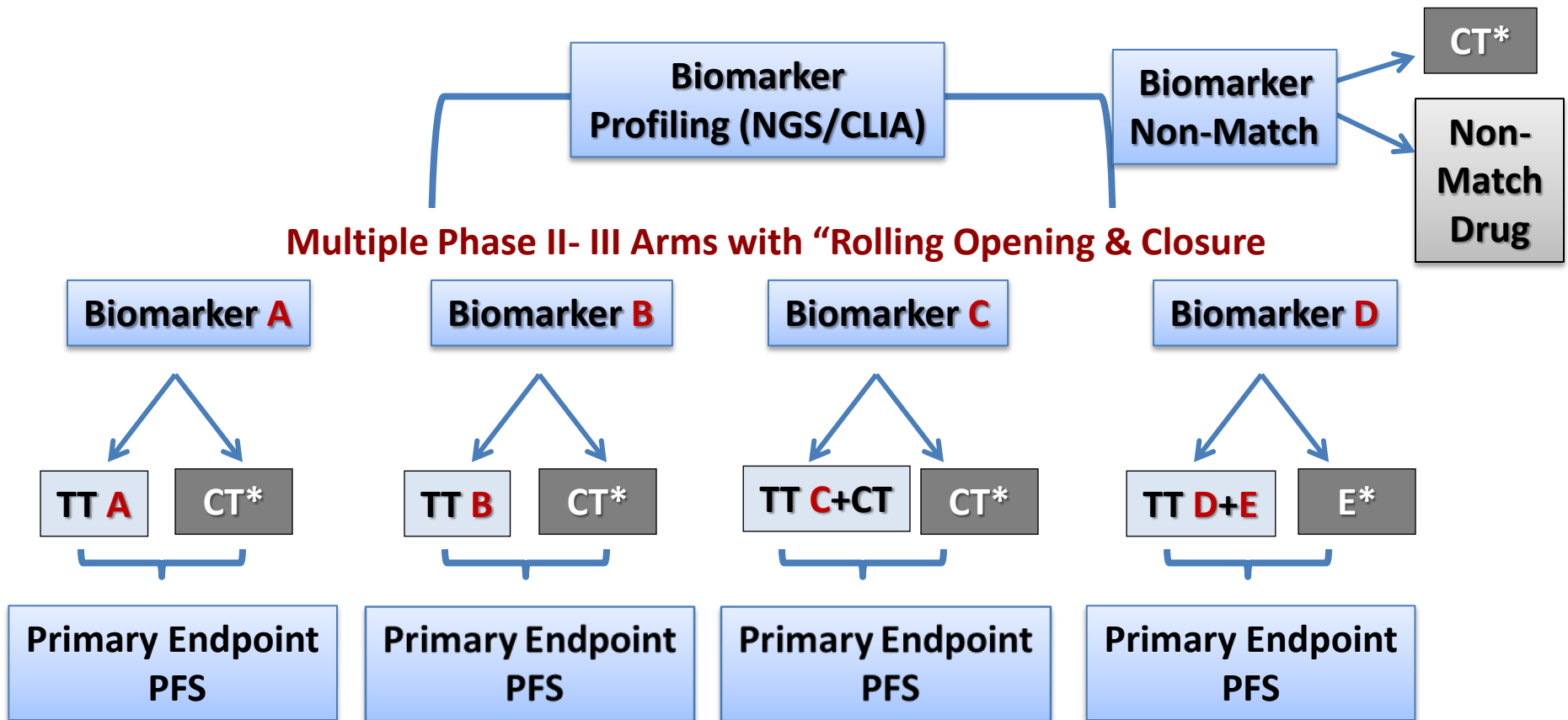
Selection of Therapeutic Targets for SCCA: Rationale for S1400 Master Lung Protocol

- **SCCA** represents an unmet need
- **Candidate targets** are available from results of TCGA project & other studies, identified by a **biomarker**
- **Drugs** (investigational) are now available for many of these targets
- Trials can be designed to **allow testing of multiple new drug-biomarker combinations at the same time** (“MASTER PROTOCOL” concept)

Therapeutic targets
SCCA-TCGA 2012

Gene	Event Type	Frequency
<i>CDKN2A</i>	Deletion/Mutation/ Methylation	72%
<i>PI3KCA</i>	Mutation	16%
<i>PTEN</i>	Mutation/Deletion	15%
<i>FGFR1</i>	Amplification	15%
<i>EGFR</i>	Amplification	9%
<i>PDGFRA</i>	Amplification/Mutati on	9%
<i>CCND1</i>	Amplification	8%
<i>DDR2</i>	Mutation	4%
<i>BRAF</i>	Mutation	4%
<i>ERBB2</i>	Amplification	4%
<i>FGFR2</i>	Mutation	3%

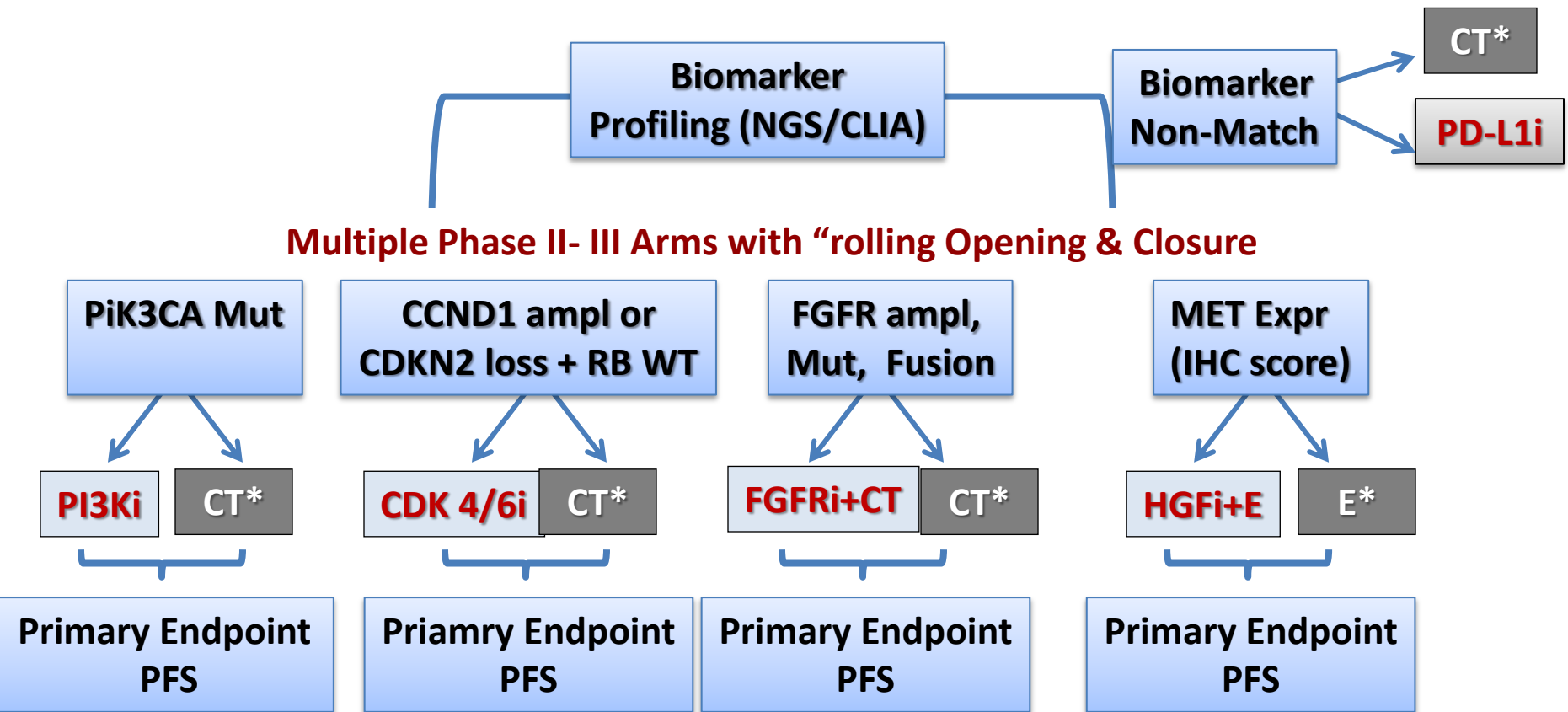
S1400: MASTER LUNG-1: Squamous Lung Cancer- 2nd Line Therapy



TT=Targeted therapy, CT=chemotherapy (docetaxel or gemcitabine), E=erlotinib

*Project Chair: V. Papadimitrakopoulou
Steering Committee Chair: R. Herbst
SWOG Lung Chair: D. Gandara*

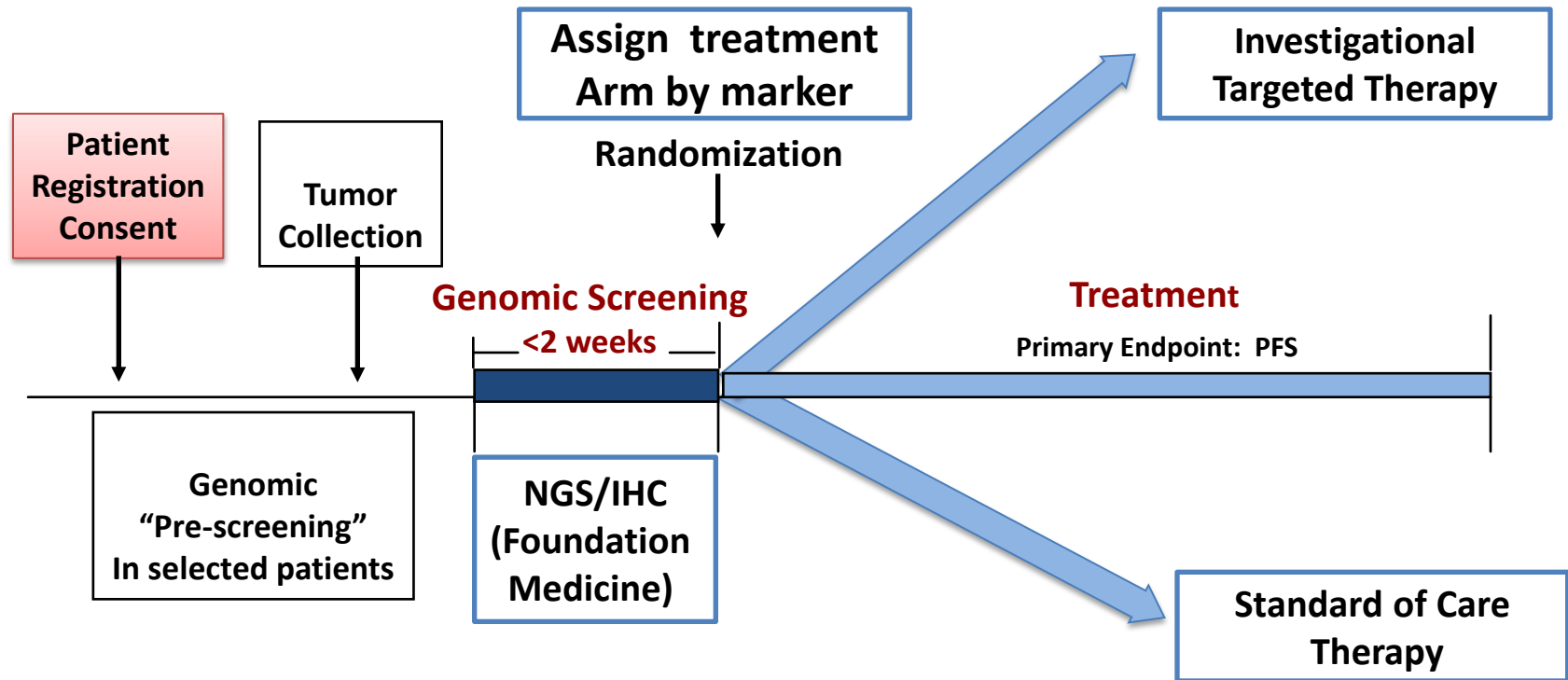
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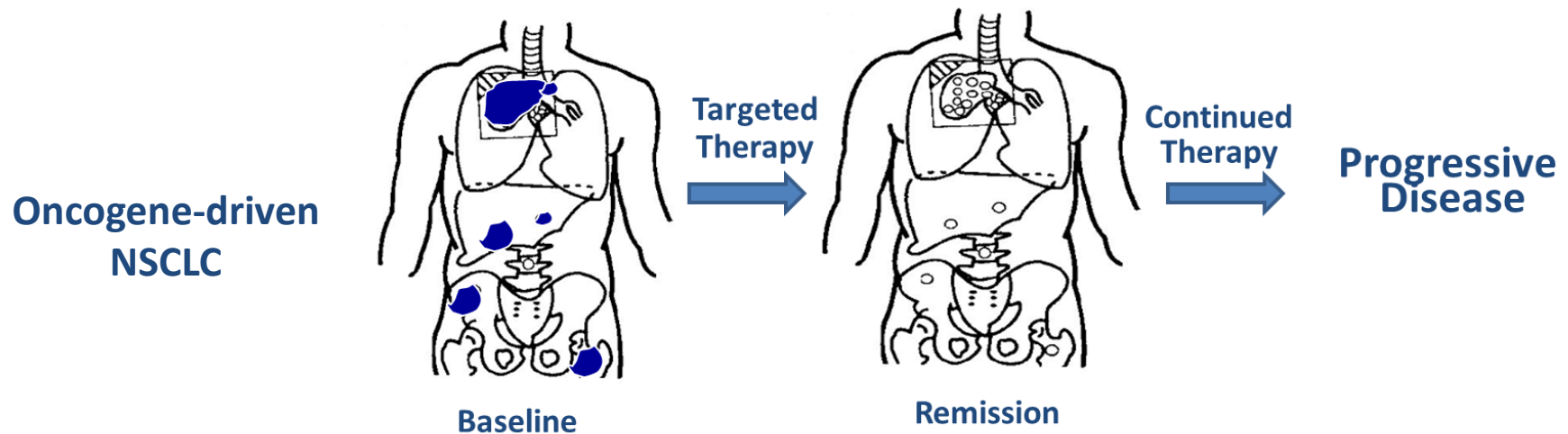
S1400 (MASTER LUNG-1) Squamous Lung Cancer- 2nd Line Therapy



- Organizers: FOCR, NCI-TMSC, FDA, FNHI
- Participants: Entire North American Lung Intergroup (SWOG, Alliance, ECOG-Acrin, NRG, NCI-Canada)
- Screening: **~1,000 patients/year**
- With 6 arms open simultaneously, anticipate a “hit rate >60% in matching a patient with a drug/biomarker arm

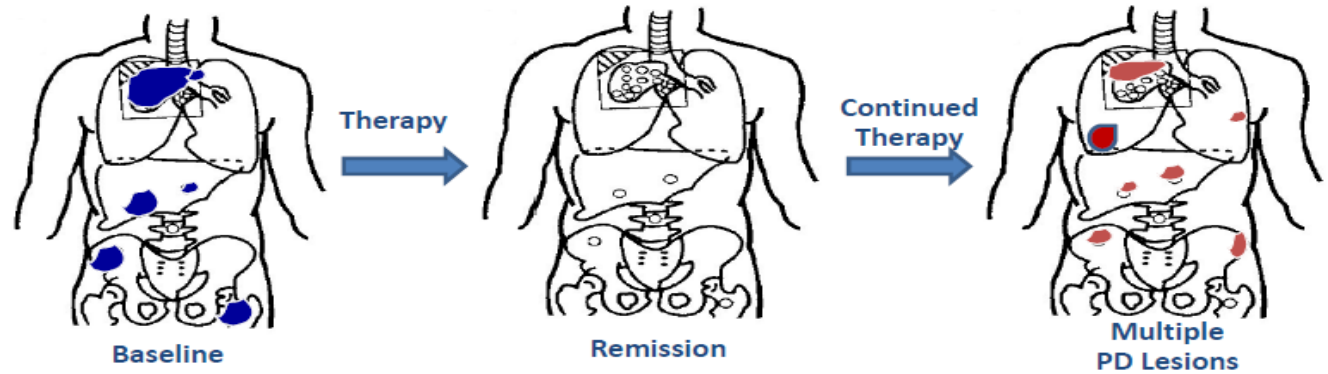
Acquired Resistance to Targeted Therapies in Oncogene-Driven NSCLC: Clinical Practice & Clinical Trials

- **Targeted Therapies** against Oncogene-Driven Cancers [EGFR mutation+ (**Erlotinib**) or ALK fusion+ (**Crizotinib**)] improve response and PFS when compared with chemotherapy
- Even in these most sensitive cancers, **acquired resistance is ~universal**, with PFS averaging ~10-14 months
- The “**subtype**” of **progressive disease (PD)** in individual patients varies greatly (**Systemic-PD, Oligo-PD and CNS-PD**)
- **Hypothesis:** “Best” management options at the time of PD varies greatly dependent on the PD subtype (also true for clinical trial designs)

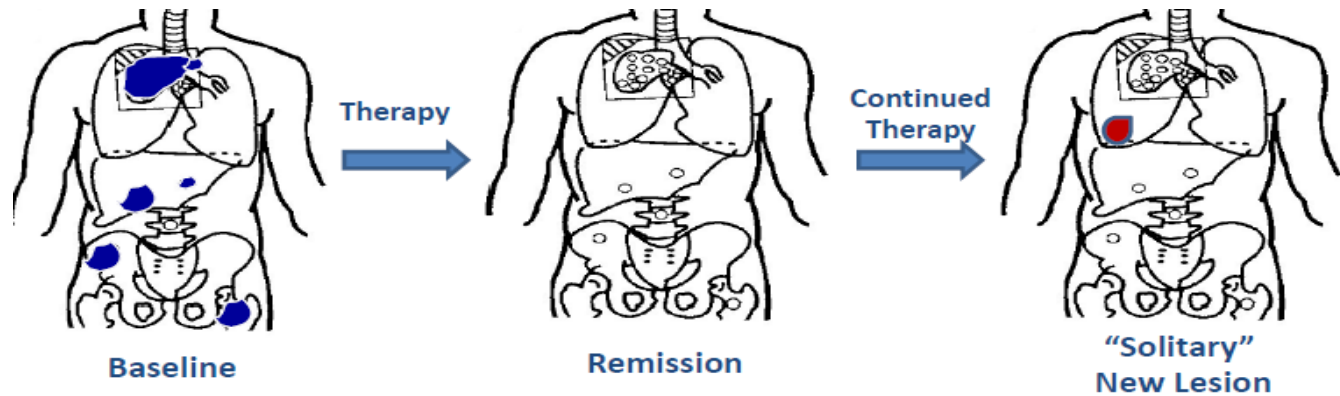


Acquired Resistance to Targeted TKIs: PD Subtype influences Clinical Practice & Clinical Trial Design

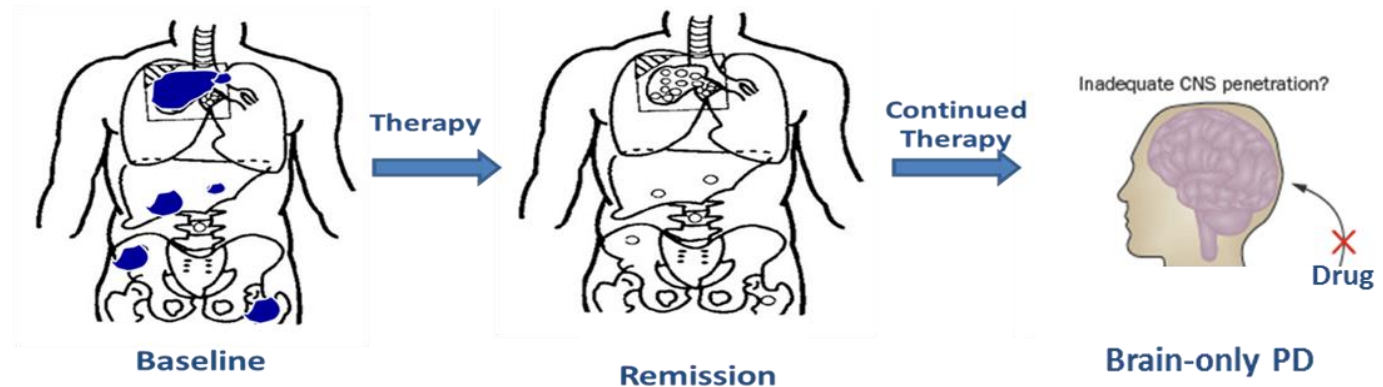
Systemic-PD



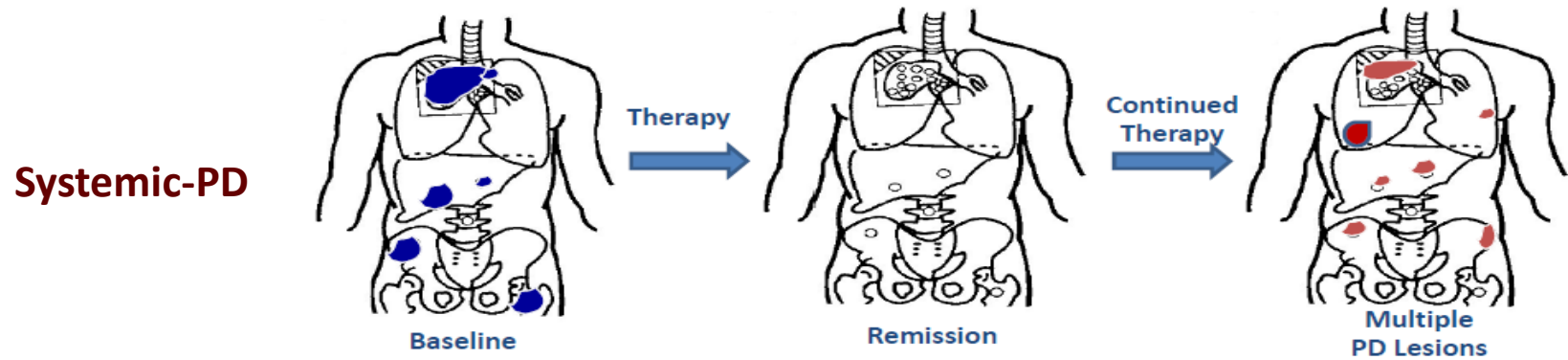
Oligo-PD



CNS-PD (Sanctuary)



Clinical Trial Designs addressing **Acquired Resistance** in Oncogene-Driven NSCLC with **Systemic PD**



Advanced
NSCLC with
Oncogene-driven
Cancer

-EGFR Mutation
-ALK Fusion

Targeted
TKI

RECIST
Response

Subsequent
Systemic PD

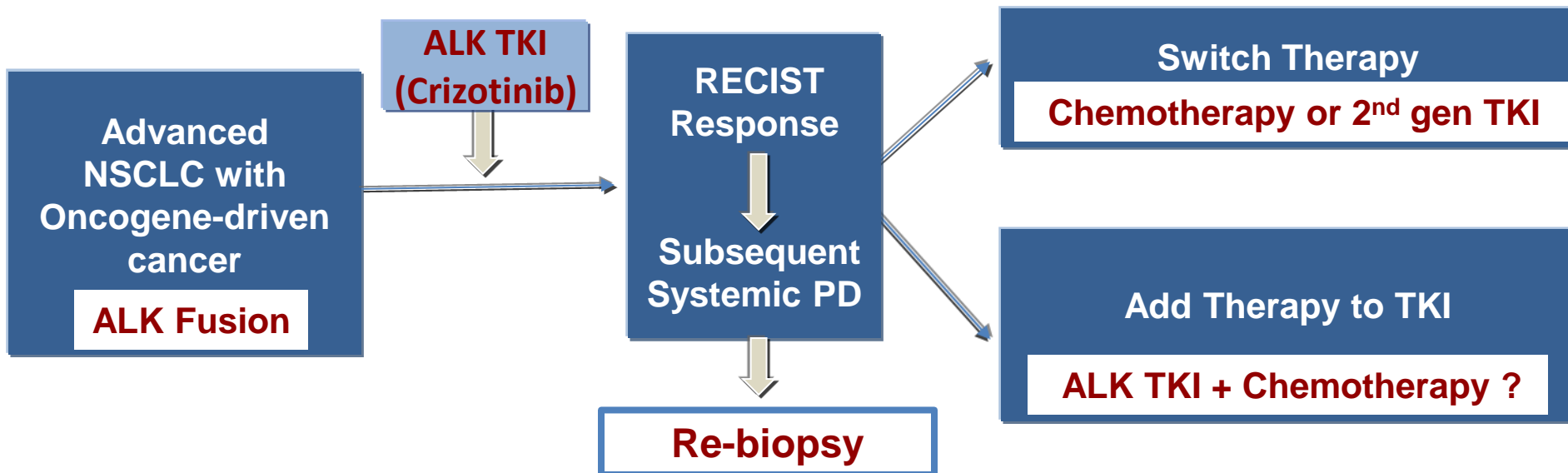
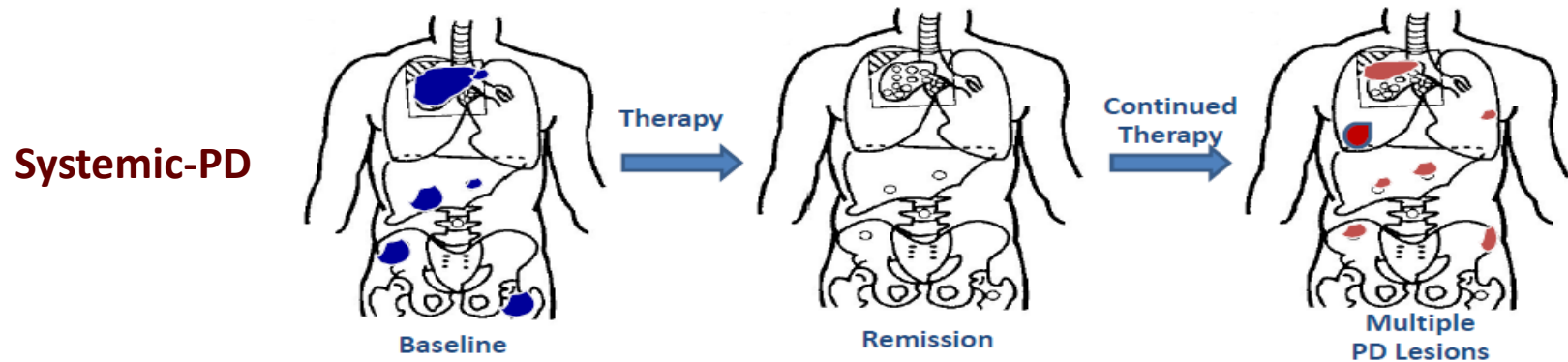
Re-biopsy

Switch Therapy
(Chemotherapy or 2nd gen TKI)

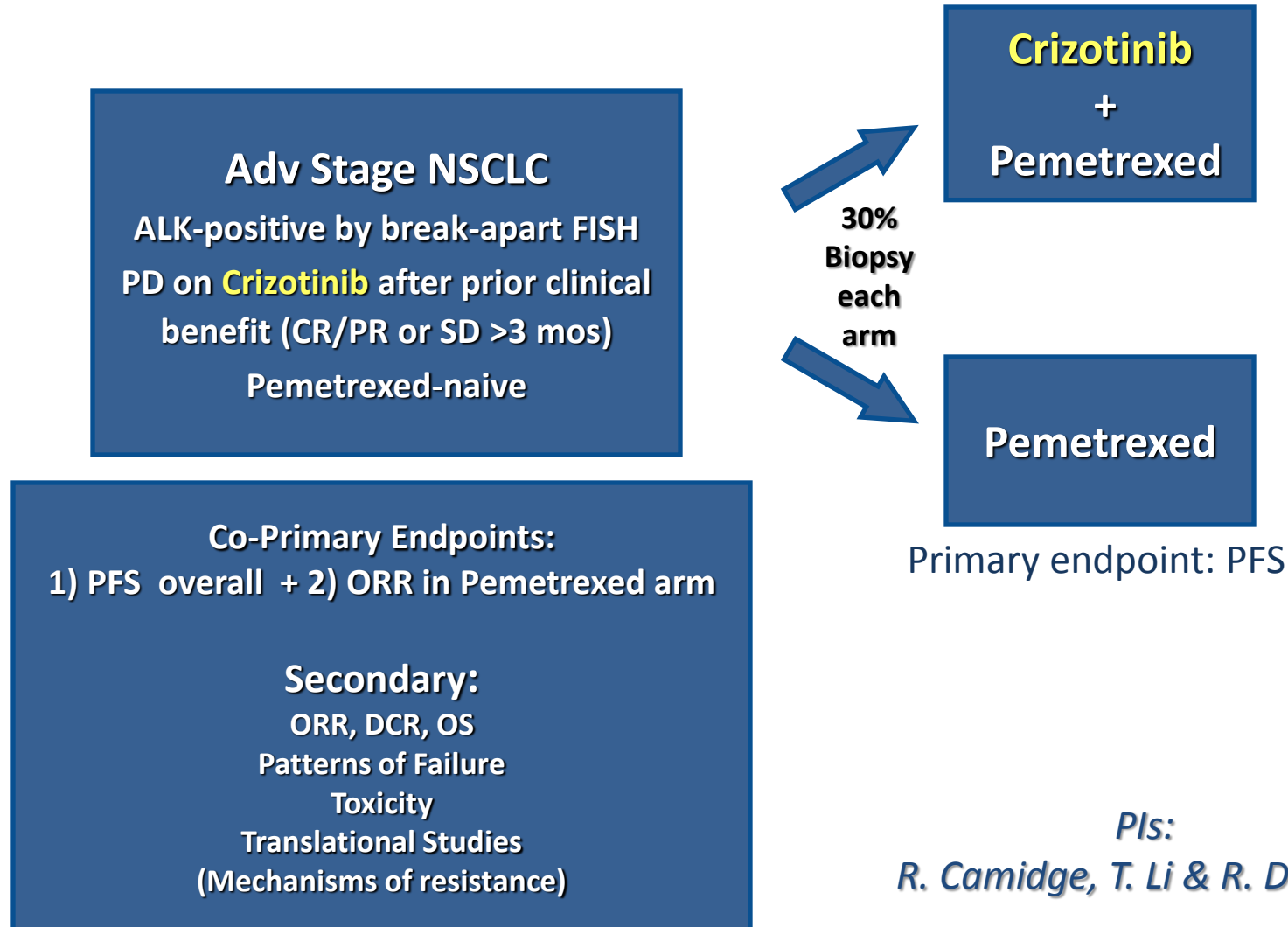
Continue same 1st gen TKI alone
(to “slow progression”)

Add Therapy to 1st gen TKI
-Chemotherapy ?
-Another Targeted Agent?

Clinical Trial Designs addressing **Acquired Resistance** in Oncogene-Driven NSCLC with Systemic PD



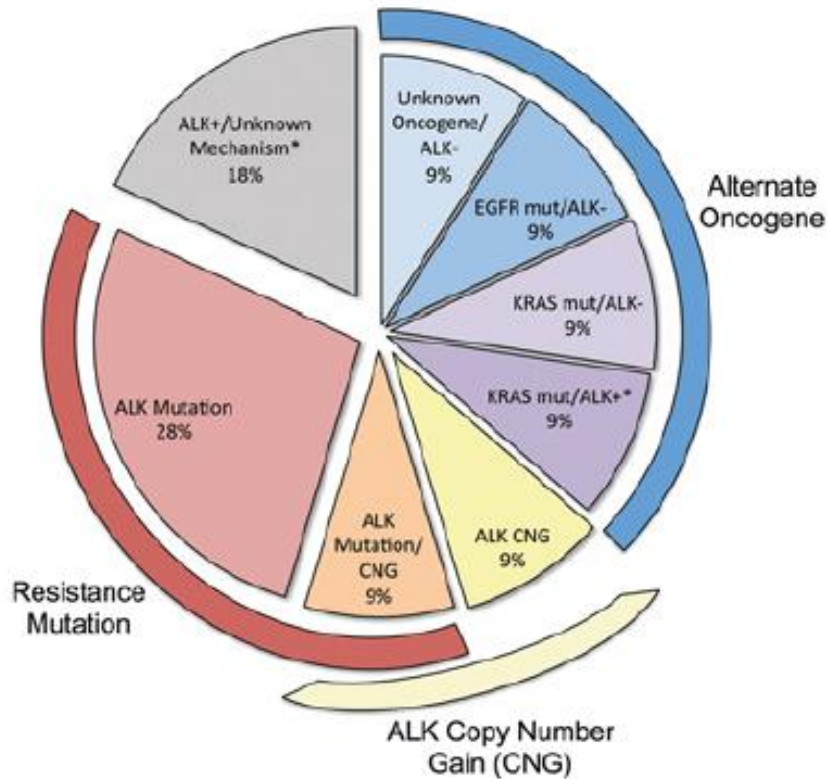
S1300: SWOG/Intergroup Phase II Trial in ALK-positive NSCLC progressive after Crizotinib



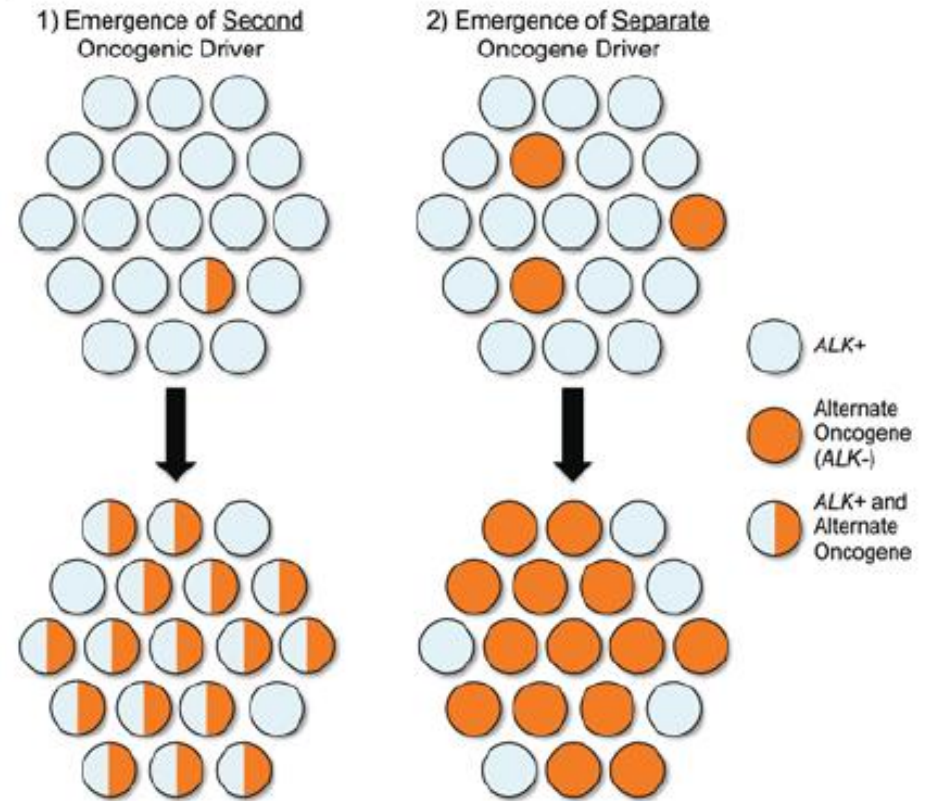
PIs:
R. Camidge, T. Li & R. Doebele

Emergence of ALK Resistance Mechanisms after Crizotinib

A



B



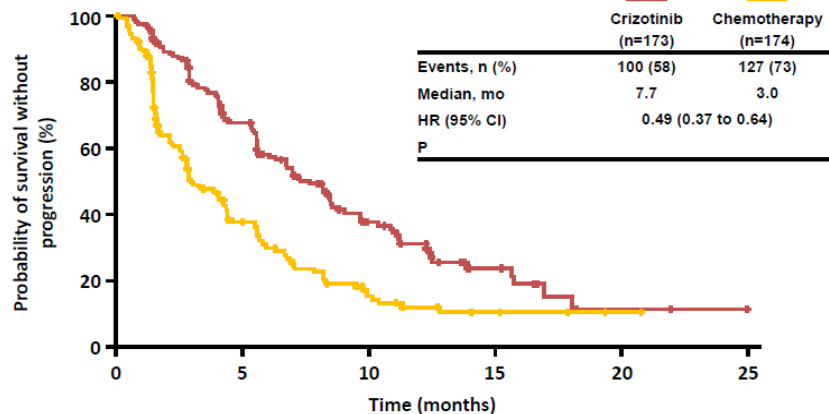
- Secondary resistance ALK mutations
- ALK Gene copy number increase
- Transition to EGFR mutation
- Transition to KRAS mutation

Consistent with mathematical models
of Evolutionary Biology

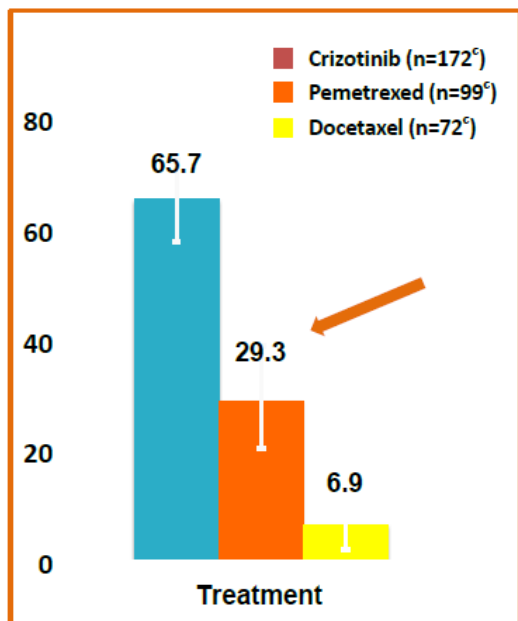
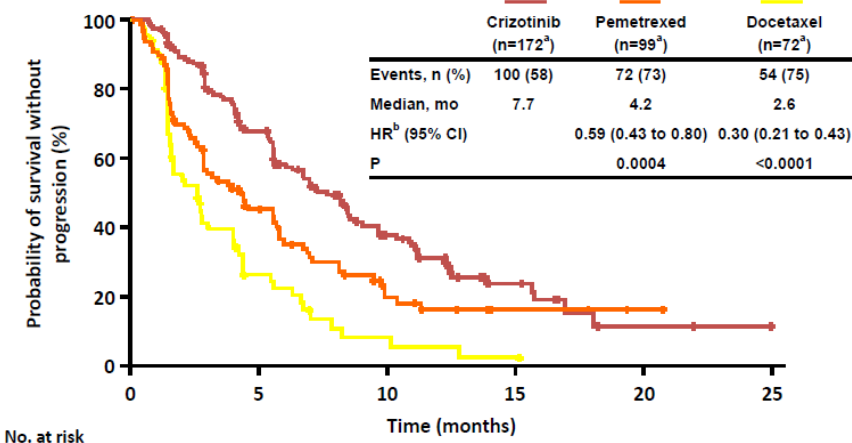
Doeble, Camidge et al: CCR 2012

Trial 1007: Crizotinib vs Chemotherapy in ALK+ NSCLC

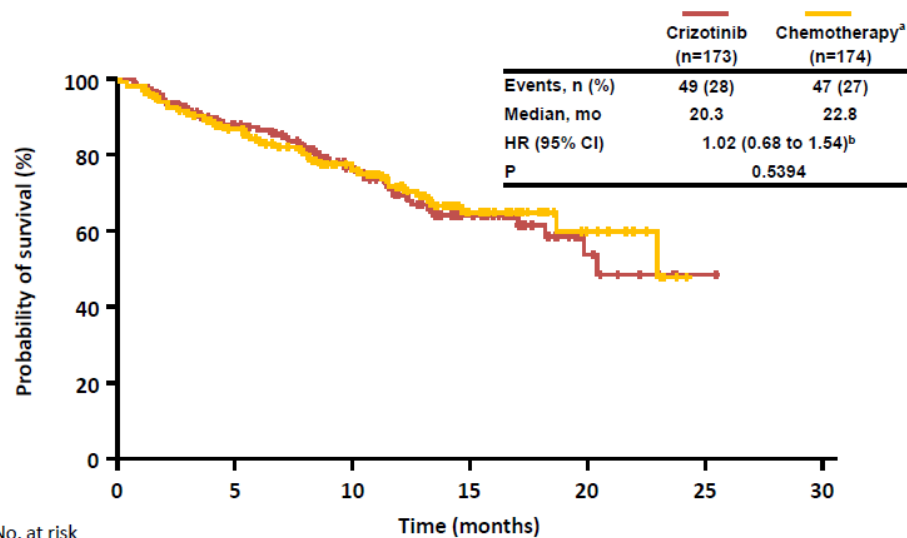
Primary Endpoint: PFS



PFS of Crizotinib vs Pemetrexed or Docetaxel



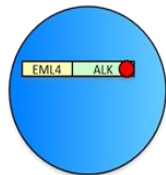
Interim Analysis of OS



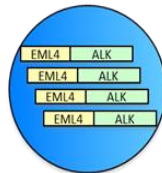
S1300: SWOG/Intergroup Phase II Trial in ALK-positive NSCLC progressive after Crizotinib

ALK dominant mechanisms of resistance

Resistance Mutations

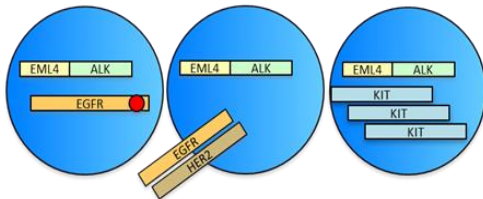


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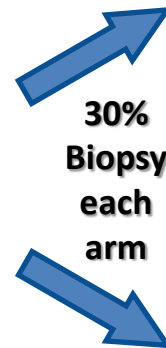
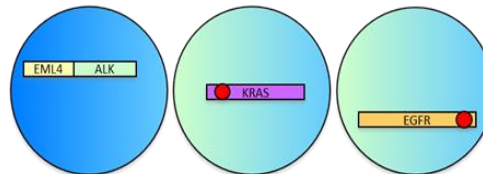


ALK non-dominant mechanisms of resistance

Second Oncogene (partially ALK dependent)



Separate Oncogene (ALK-independent)



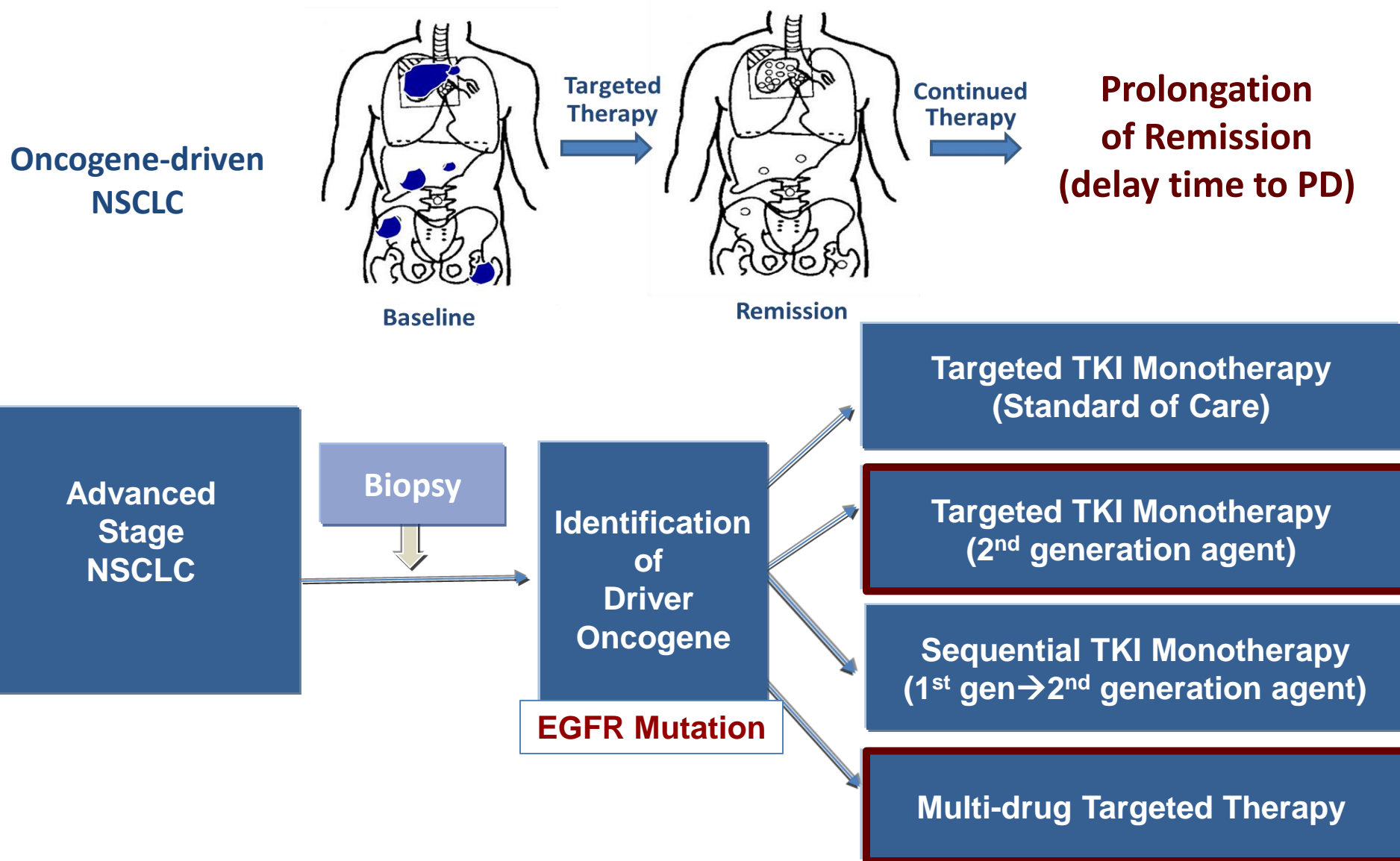
Crizotinib
+
Pemetrexed

Pemetrexed

Secondary:
ORR, DCR, OS
Patterns of Failure
Toxicity
Translational Studies
(Mechanisms of resistance)

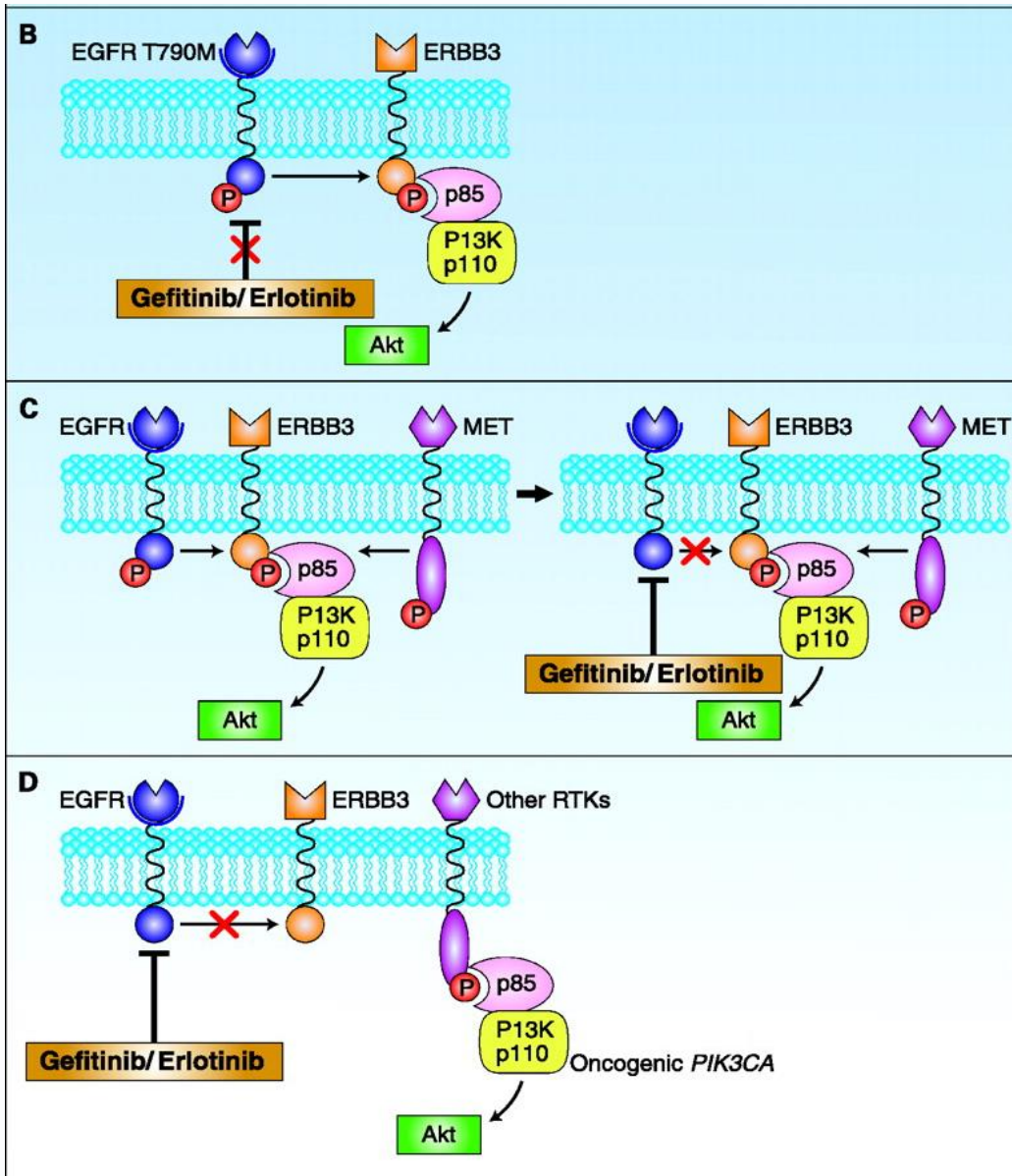
PIs:
R. Camidge, T. Li & R. Doebele

Clinical Trial Designs for Circumvention (Prevention or Delay) of Acquired Resistance in Oncogene-Driven NSCLC



from Gandara, Redman et al: Clin Lung Cancer 2013 (in Press)

Mechanisms of EGFR TKI Resistance (Selected)



•Secondary EGFR mutation (i.e. T790m)

↳ 2nd Gen EGFR TKIs
i.e. Afatinib

3rd Gen- **Afatinib/Cetuximab**
AZ9291, CO1686

•Bypass signaling via ERBB3

↳ Anti-ERBB3 drugs
i.e. MM151 MoAB

•MET over-expression

↳ MET Inhibitors
i.e. MET-Mab (MoAB)
ARQ197 (TKI)

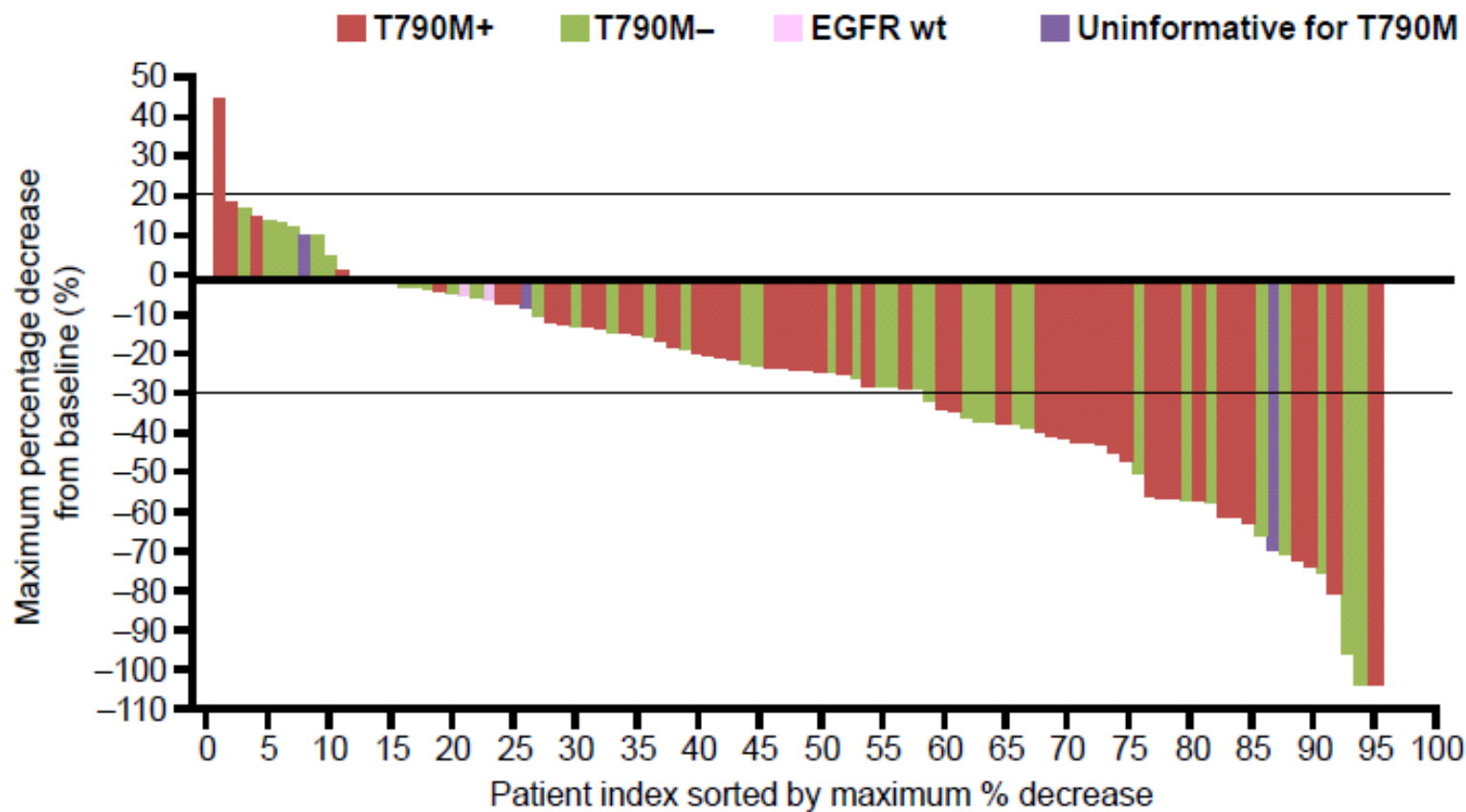
•PIK3CA Mutation/AKT

↳ i.e. BKM120 (PIK3CA)
i.e. MK2206 (AKT)

& Others → HSP inhibitors
i.e. Ganetespib
AUY922

adapted from Engelman et al

Afatinib + Cetuximab in EGFR-mutated NSCLC refractory to EGFR TKI



Response rate: 30%
Clinical benefit (DCR): 75%

Janjigian, et al. ESMO 2012

Developing Randomized trials: Afatinib +/- Cetuximab in EGFR mutation+ NSCLC (North American Intergroup)

PI:
Goldberg
(SWOG-
S1403)

Circumvention
of Resistance

Stage IIIB-IV
Adenocarcinoma with
EGFR mutation+
1st Line
EGFR TKI naïve

R
A
N
D
O
M
I
Z
A
T
I
O
N

Afatinib*

Afatinib + Cetuximab*

*at PD: Biopsy for genomic study
& PDX development (selected)

PI:
Pao
(ECOG-
coordinated)

Reversal
of Resistance

Stage IIIB-IV
Adenocarcinoma with
EGFR mutation+
**EGFR TKI pre-treated
& resistant**

R
A
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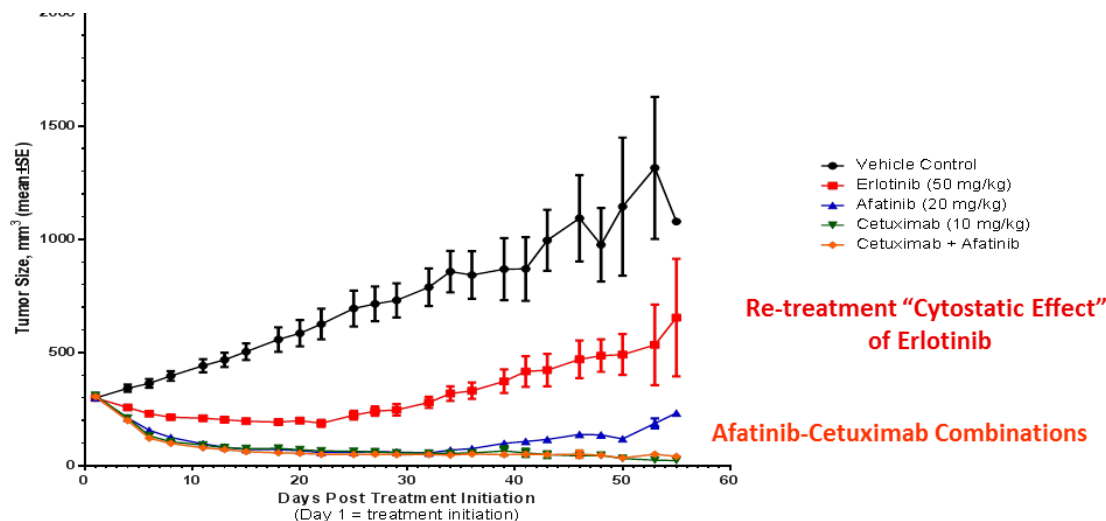
Afatinib

Afatinib + Cetuximab

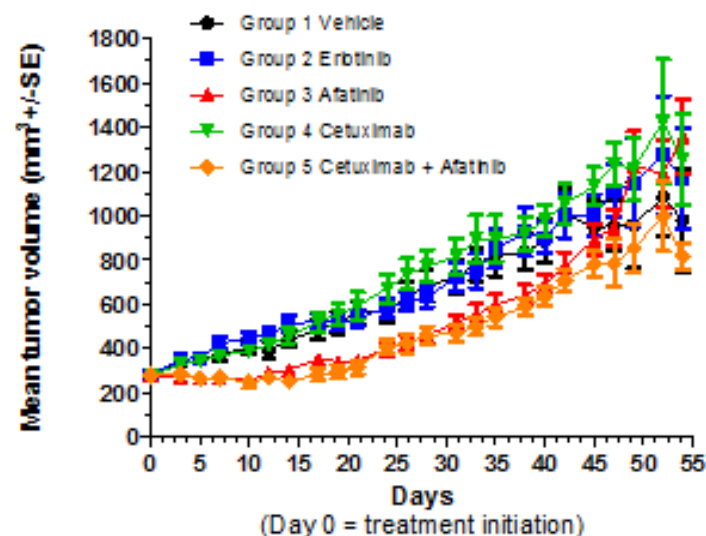
Afatinib-Cetuximab in EGFR mutant & Erlotinib Acquired Resistance PDX Models:

Results in PDX models mimic the clinical response to Afatinib-Cetuximab

LG0703: EGFR TKI resistant (L858R + High MET. T790M negative)



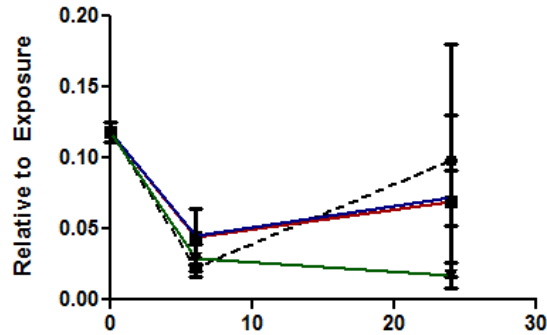
LG1049: EGFR TKI resistant (E19del + T790M positive)



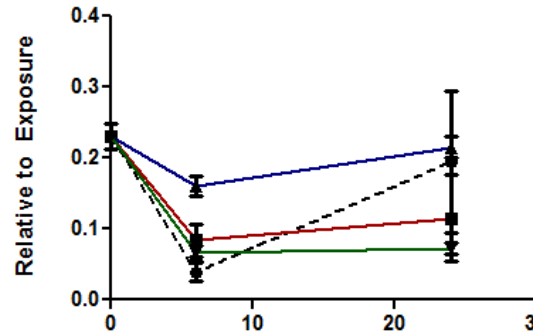
**Next Generation Sequencing & Signaling Pharmacodynamics
Pre- & Post-Therapy
to determine & characterize Mechanisms of Sensitivity/Resistance**

Time-dependent treatment effects on signaling pharmacodynamics (LG703)

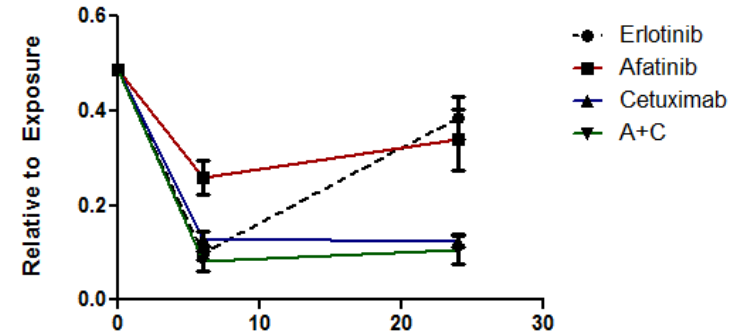
AKT1 expression



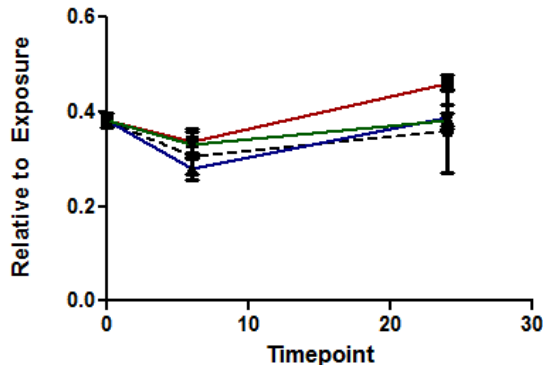
AKT2 expression



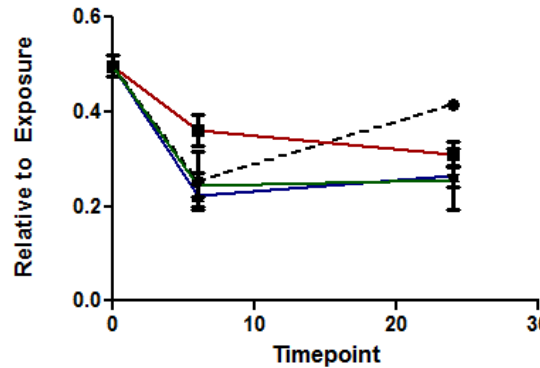
ERK1



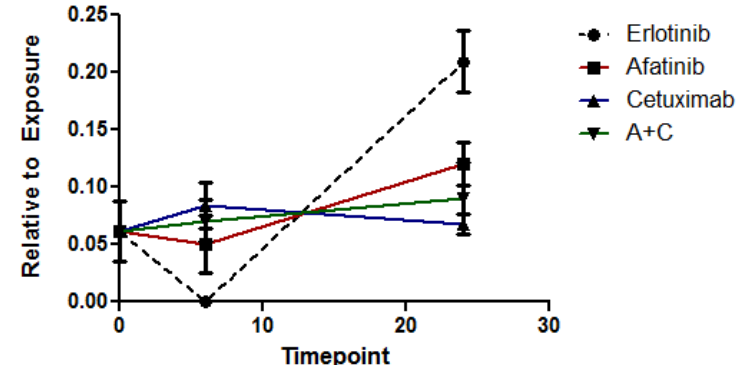
ERK2



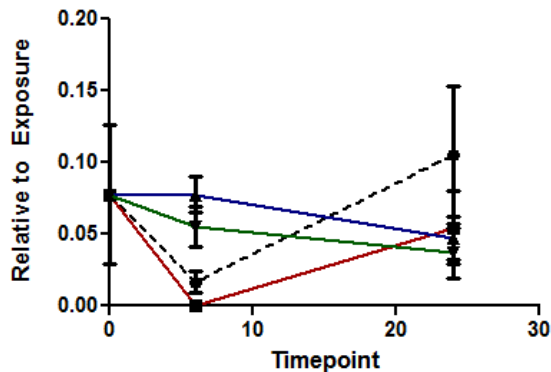
p38a



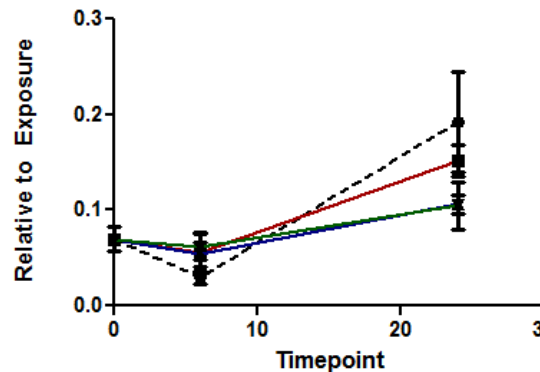
MSK2



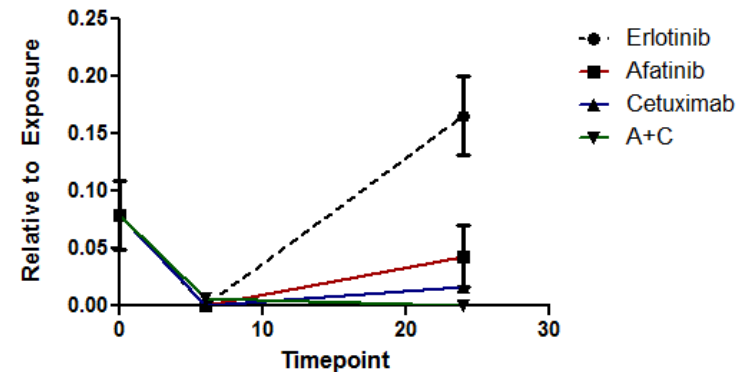
RSK1



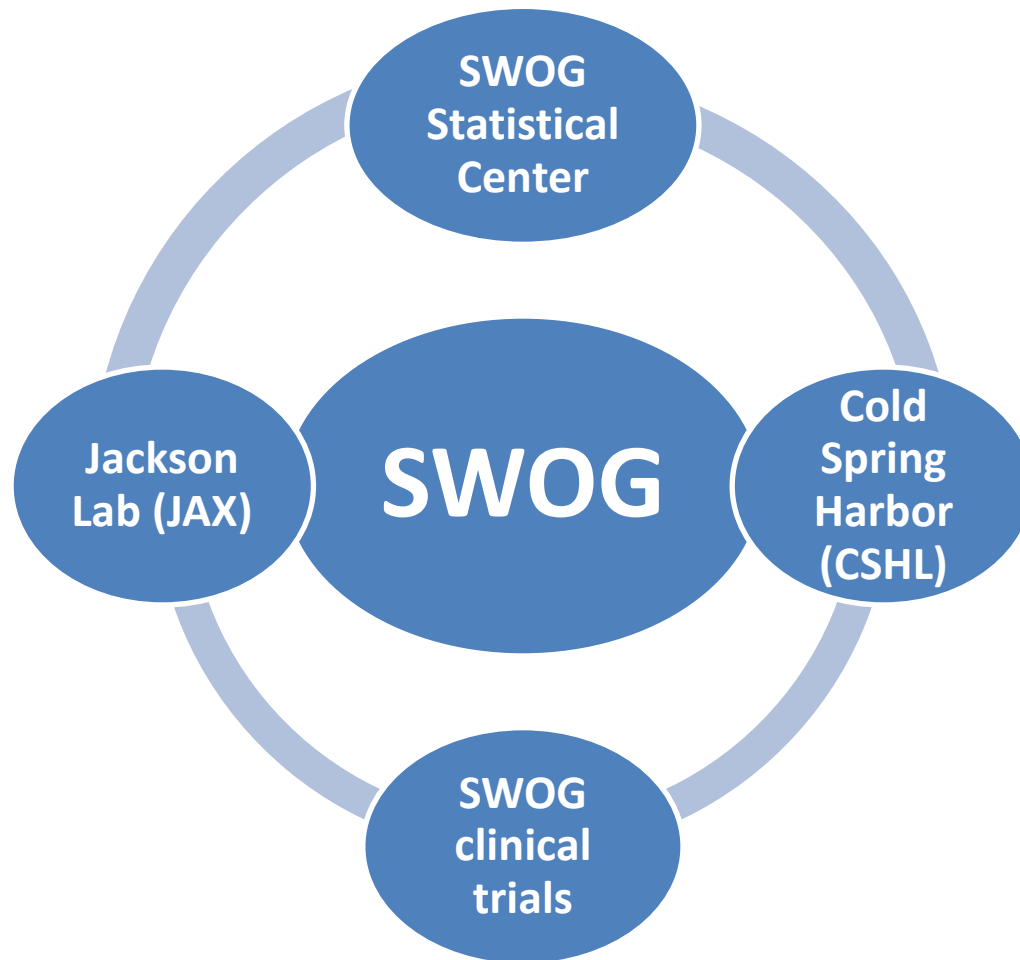
JNK pan



p70S6K



SWOG Translational Science Center: Pilot PDX Project in S1403



Summary: Integrating Predictive Biomarkers in Clinical Trial Design

- Master Protocol Designs may provide **operational efficiencies to speed up drug-biomarker development & approval**, including for drugs directed against uncommon genotypes
- **Despite advances with targeted TKIs in Oncogene-driven NSCLC no patients are cured and acquired/adaptive resistance is ~universal**
- **Subtyping PD into clinically relevant categories** should assist in both clinical trial design & day-to-day patient management
- **Methods to identify mechanisms of acquired resistance & how to overcome them (or circumvent them)** are needed
- Clinical Trial designs will need to account for **inter- and intra-patient tumor heterogeneity & the most likely mechanisms of resistance**
- **Clinically & genomically annotated PDX resources** may assist in achieving this latter goal