The Changing Landscape in Phase I and Early Phase Clinical Trials

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I have the following financial relationships to disclose:

Consultant for: Boerhinger-Ingelheim (uncompensated), Merck (compensated), Pfizer (compensated), Celgene (compensated)

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Stockholder in: None

Honoraria from: None

Employee of: None

The Traditional Drug Development Paradigm

Phase I Phase II Phase III

- Safety, tolerability
- Pharmacokinetics
- Pharmacodynamics
- Preliminary antitumor activity
- Efficacy
 observed in
 selected tumor
 types, e.g. ORR,
 TTP, PFS
- Meaningful benefit obtained in a randomized setting against existent standard e.g. OS

The Current Drug Development Paradigm

Proof of Mechanism Early Late

- Safety, tolerability on target and off target effects
- Preliminary antitumor activity
- Evidence of target engagement in valid pharmacodynamic biomarkers

- Predictive biomarkers explored
- Antitumor
 activity seen
 using surrogate
 endpoints e.g.
 ORR, TTP or PFS
- Predictive biomarkers confirmed
- Proof of concept using a validated clinical endpoint e.g. OS

Objectives

 Describe features related to the changing nature of phase I clinical trials in the era of novel onco-therapeutics

 Understand the reasons that may have resulted in such changes in phase I trials and their implications in the drug development process

Changing Nature of Phase I Trials

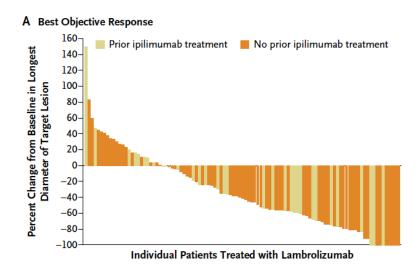
- 1. Trend of increase in the sample size of phase I trials
- Expansion cohorts being conducted for multiple purposes
- Enrichment strategies histology and/or genotype
- 4. Emergence of immuno-oncology era
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- 6. Research biopsies
- 7. Driving go-no-go decisions based on their ability to provide proof-of-concept

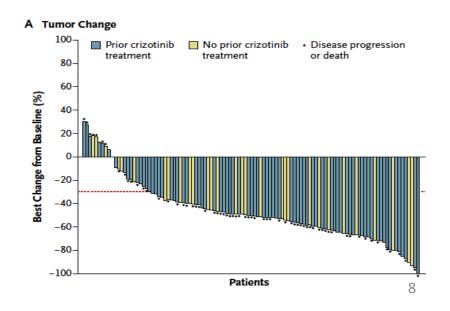
Why Are Phase I Trials Changing?

- 1. Knowledge of molecular biology is accumulating and technology is rapidly advancing
- 2. Molecularly targeted agents and immunooncology agents have become important parts of the oncology therapeutic armamentarium
- 3. Patient and infrastructure resources are limited
- 4. Accelerated regulatory approval is possible for compelling results
- 5. The desire to accelerate the drug development process to bring active compounds to the clinic and improve cancer cures have fueled these changes

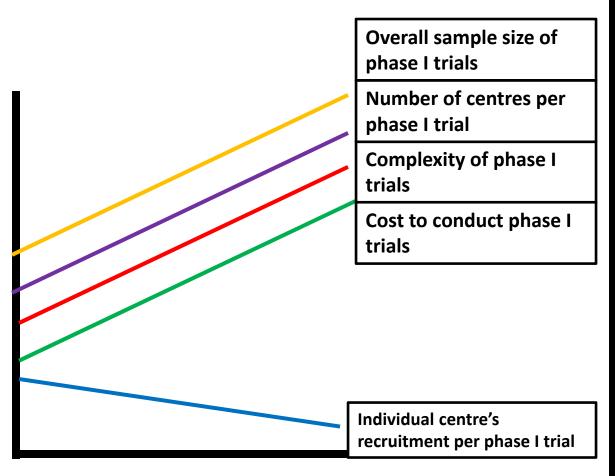
Accelerated Approvals of New Drugs: 2 Examples

Drug	Phase I to Approval by FDA	Time (years)
Pembrolizumab (anti-PD-1 antibody)	February 2011 to September 2014	3.6 years
Ceritinib (ALK inhibitor)	January 2011 to April 2014	3.3 years





Economics and Logistics of Phase I Trials



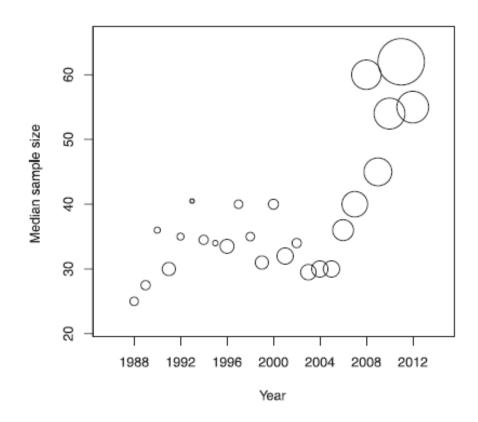
Consequences:

- Each centre needs to open multiple studies to be economically viable
- Greater regulatory burden (protocol amendments, SUSARs, etc)
- Cost per case is increased
- Limited experience being accumulated per centre
- Collection of trial data by sponsor – there must be sharing of toxicity data by grade and frequency on a regular basis throughout protocol conduct

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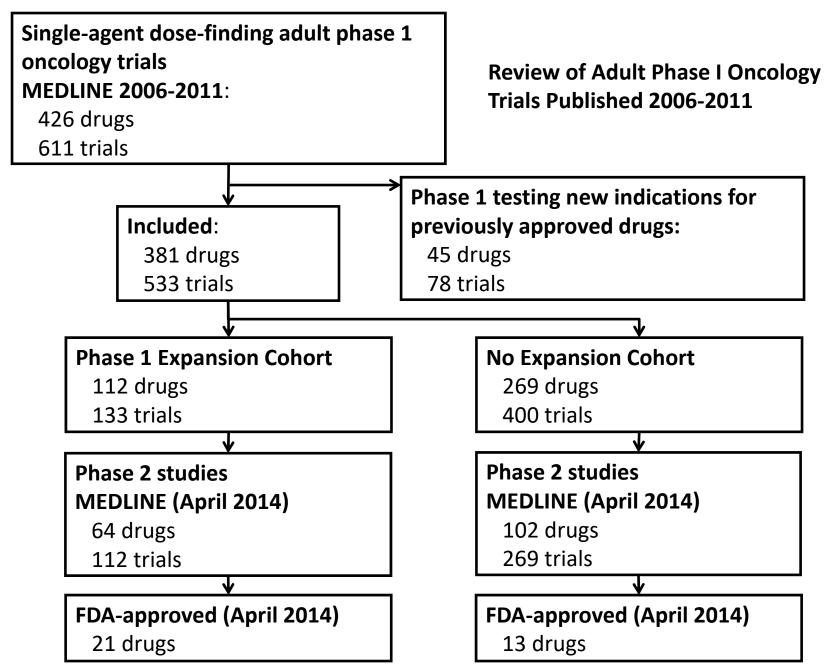
Increase in Overall Sample Size of Phase I Trials



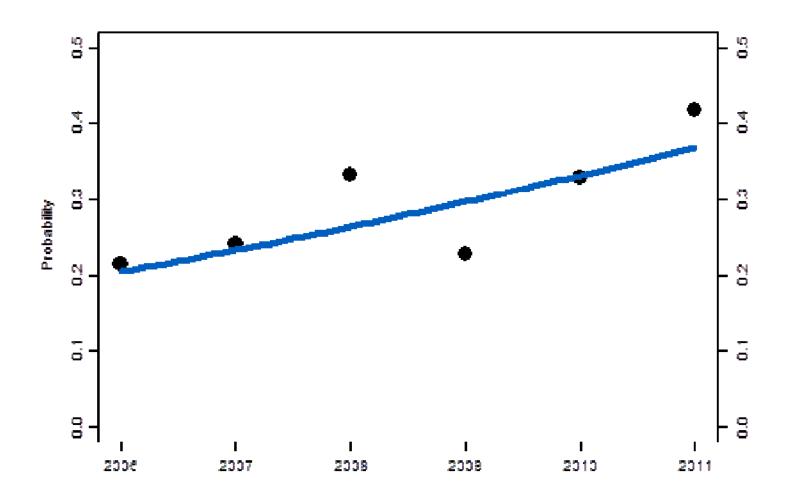
The average sample size of a phase I study has increased from 33.8 patients (1988-1992) to 73.1 patients (2008-2012)

Expansion Cohorts (EC) in Phase I Trials

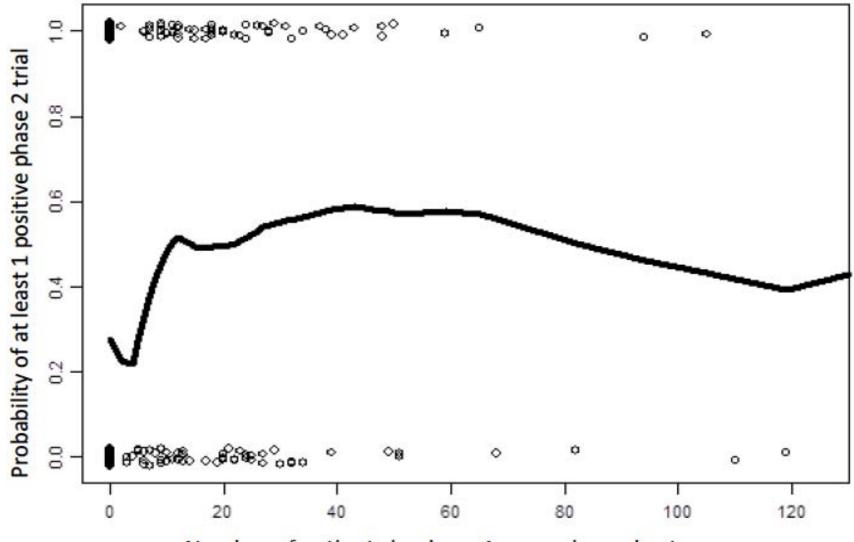
- Systemic review of adult single-agent phase I trials after 2006
- 149 (24%) of 611 trials used ≥ 1 EC, increased from 12% in 2006 to 38% in 2011
- Median number of pts: 22 in dose-escalation cohorts and 17 in EC
- Phase I trials more likely to include EC if multicentre (OR 1.8), non-cytotoxic agents (OR 2.0), industry sponsored (OR 1.6, p = 0.063)
- EC objectives reported in 74% of trials:
 - Safety (80%), efficacy (45%), PK (28%), pharmacodynamics (23%),
 patient enrichment (14%)
 - Among ECs assessing safety, MTD modified in 13% and new toxicities defined in 54%



Probability of Having an Expansion Cohort According to Year of Publication of the Phase I Trial



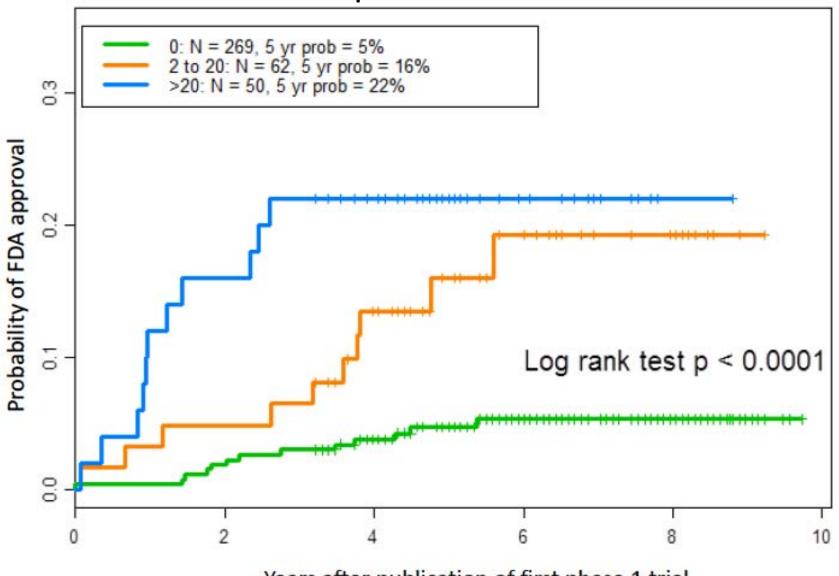
Probability of Success in a Phase 2 Trial Relative to the Size of the Phase 1 Expansion Cohort



Number of patients in phase 1 expansion cohort

Bugano, Hess, Siu, Meric-Bernatam, Razak, Hong, In Press CCR

Probability of FDA Approval and the Number of Patients in the Phase 1 Expansion Cohort



Years after publication of first phase 1 trial

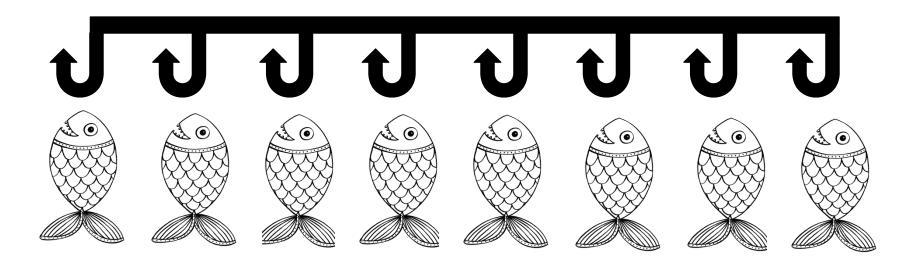
Bugano, Hess, Siu, Meric-Bernatam, Razak, Hong, In Press CCR

Cox Regression Model of Time-to-Drug-Approval

	Comparison	Univariate		Multiva	riate
		HR(95% CI)	р	HR(95%CI)	р
Targeted agent	YvN	0.7(0.3;1.7)	0.42	1.0(0.4;2.4)	0.95
Industry-sponsored	YvN	4.4(1.1;18)	0.4	2.1(0.5;9.5)	0.33
Multicenter	YvN	4.0(1.2;13)	0.02	2.4(0.7;8.5)	0.17
Pub >2008	YvN	1.2(0.6;2.4)	0.57	1.0(0.5;2.2)	0.94
Tumor type	Hematologic v solid	4.0(1.3;12)	0.014	2.4(0.7;8.8)	0.17
	Hem+solid v solid	0.9(0.2;4.0)	0.91	0.6(0.1;4.6)	0.62
	Specific histology v any solid	1.6(0.8;3.4)	0.18	2.1(1.0;4.4)	0.066
Number of patients	21-37 v < 21	0.8(0.3;2.1)	0.67	0.9(0.3;2.6)	0.88
in dose escalation cohort	>37 v < 21	1.7(0.8;3.9)	0.19	1.4(0.6;3.4)	0.46
Number of patients	2-20 vs 0	2.7(1.1;7.0)	0.034	2.1(0.8;5.4)	0.14
in expansion cohort	21-271 vs 0	8.8(4.0;19.0)	<0.0001	6.6(2.9;15)	<0.0001

Single Protocol, Multiple Cohorts Signal-Finding Trials:

Common Design with Immune Checkpoint Inhibitors

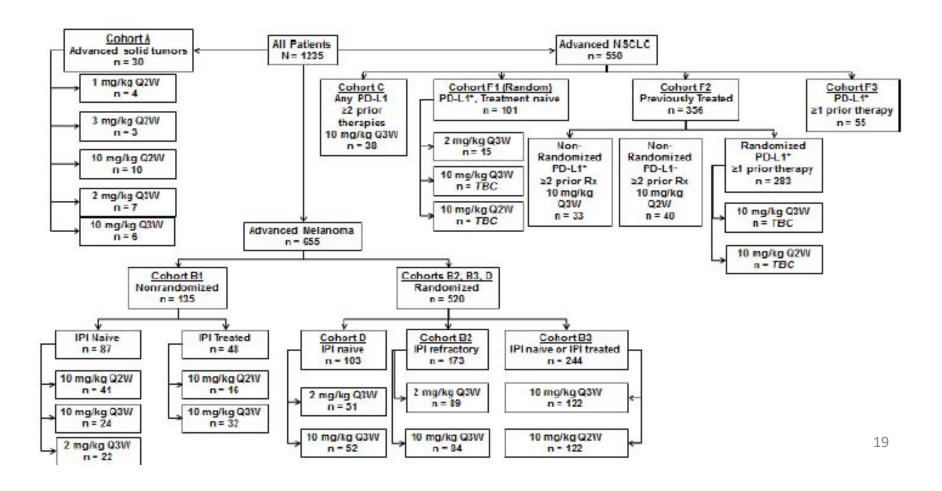


Cancer A Cancer B Cancer C Cancer D Cancer E Cancer F Cancer G Cancer H

Protocol 001 (PN001) First in Human (FIH) to Registration Cohort Expansion

From a small Phase 1-the study expanded to a 655-melanoma patient multi-part study

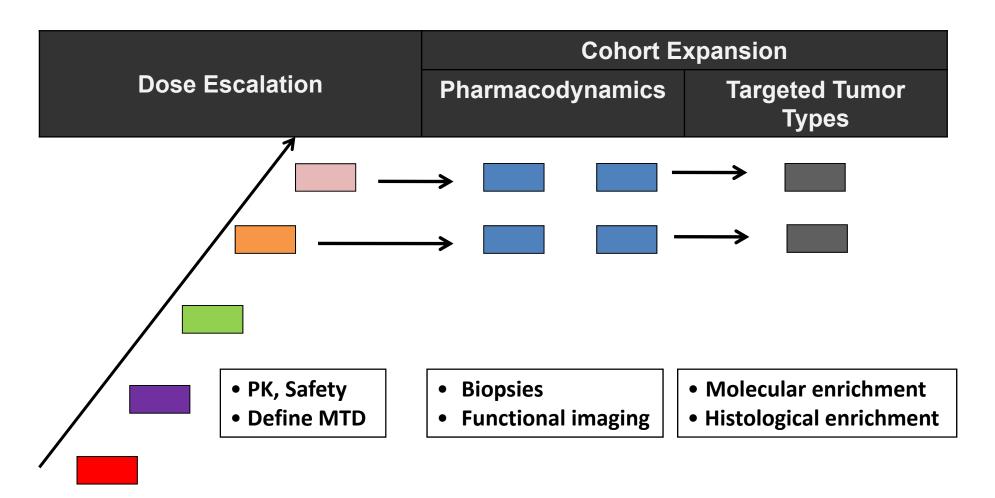
- 5 amendments, between Dec-2011 to Sep-2013, to answer emerging questions
- 4 "phase 2 study-like" parts including 3 randomized dose comparison sub-studies



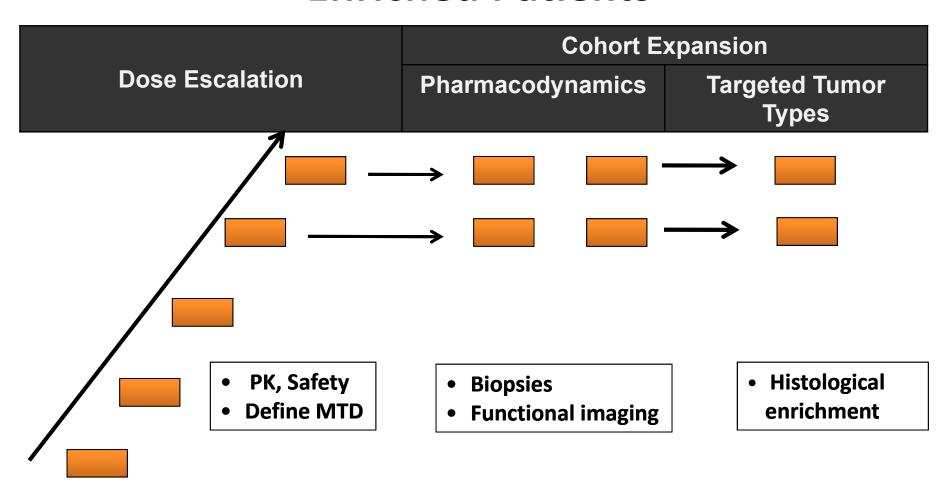
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Phase I Study Design – Unselected Patients in Dose Escalation followed by Specific Expansion Cohorts

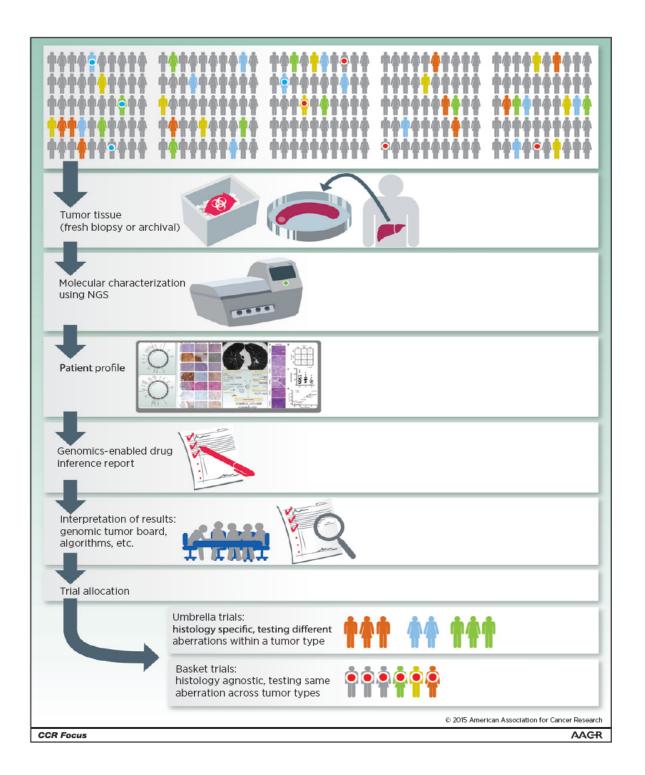


Phase I Study Design – Only Molecularly Enriched Patients



Enrichment and Patient Selection in Phase I Trials

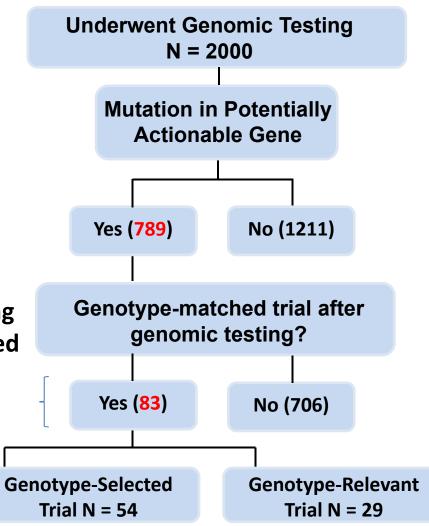
Element	Challenges	Potential Solution
Molecular selection	 Central Screening: Archived tumor tissues requested by multiple sponsors, leading to exhaustion of tissues Turnaround time variable Return of molecular information may lack sufficient annotation Local Screening: Local screening typically not reimbursed Assay may not have been validated in CLIA lab 	 Local laboratory testing using validated multiplexed assay (funding remains an issue)
Identification of rare subsets of patients	↑ screening costs while number of eligible patients ↓, leading to a financial challenge to keep many trials open with few patients recruited per trial	 Support for screening Multiplexed screening Umbrella or Basket protocols



Clinical
Application
of Next
Generation
Sequencing
to Find
Matching
Treatment

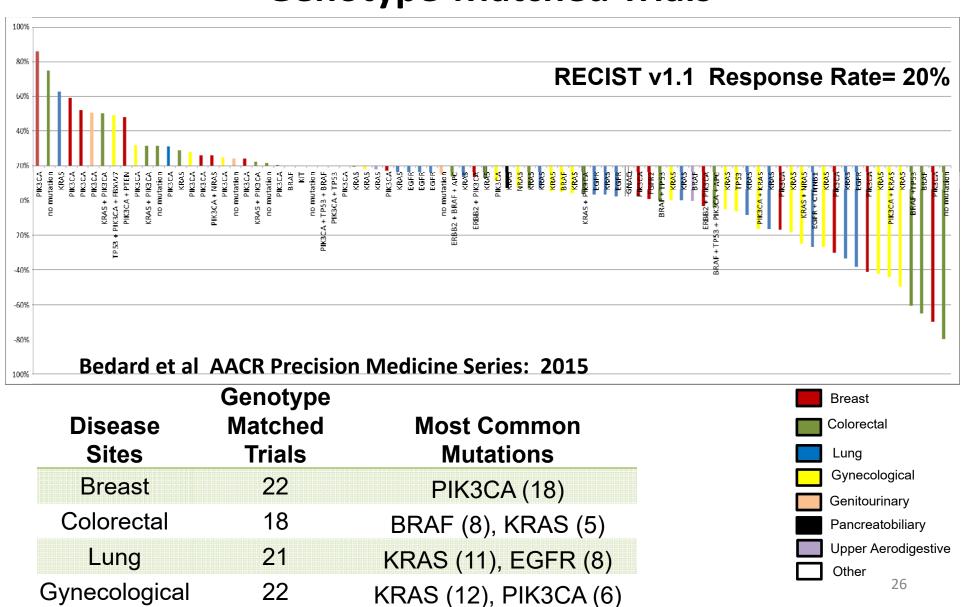
Siu, Conley, Boerner, Lorusso et al. CCR Focus, In Press

MDACC: Enrollment on Genotype-Matched Trials



54/2000 (3%) of pts who underwent genomic testing received genotype-matched treatment

Best Tumor Shrinkage of Patients Enrolled in Genotype-Matched Trials



Characteristics of Therapeutic Trial Patients

		All	Genotype Matched	Genotype Unmatched	p-value
Median Prior	Therapies	4	4	4	p=NS
Range Prior	Therapies	1-18	1-18	1-15	•
Genotyping Platform					
	Sequenom	176	63	113	
Illum	ina TruSeq	101	29	72	p=0.23
lo	n Ampliseq	0	0	0	
≥1 ı	mutation(s)	168	84	84	
no actionab	le mutation	109	8	101	
Trial Phase					
	Phase I	158	74	84	
	Phase II	67	9	58	p<0.001
	Phase III	50	8	42	
Investigational Agent	(s)				
Targeted Mo	onotherapy	112	23	89	
Targeted Drug Combination		86	61	25	- 40 004
Targeted Drug + Chemotherapy		43	7	36	p<0.001
Imm	unotherapy	34	1	33	

Attrition in Molecular Profiling and Genotype-Drug Matching

Enrolled in molecular profiling initiative

- Limitation of resources and personnel to perform profiling
- Screening/eligibility criteria e.g. ECOG, organ functions

Evaluable molecular profiling results

- Insufficient tissue or poor DNA quality/quantity
- Technical, operational or annotation issues

Druggable molecular aberrations

- No druggable/actionable molecular aberrations
- Limited understanding in biological functions of many variants

Genotypematched trial

- Lack of genomic-based trials or approved drugs
- Patients not clinical trial candidates
- Physicians not aware of or not actively seeking out genotype-matching trials

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Selected Molecular Profiling Initiatives and Genotype-Matching to Clinical Trials

Group	Sampl e Size	Platform	Fresh Biopsy vs FFPE	Germ- line Control	Number and % of Patients in Genotype-Matched Clinical Trials
Gustave Roussy	708	30-75 gene panels (Life) + CGH (Agilent)	Fresh biopsy	Yes	140/708 = <mark>19%</mark>
Institut Curie	741	46 gene panel (Life) + CNA (Affymetrix) +IHC	Fresh biopsy	No	195 randomized/741 = 26%
BCCA	100	Whole genome	Fresh biopsy	Yes	1/100 = <mark>1</mark> %
MD Anderson	2,000	11-50 gene panels (Life)	FFPE	No	83/2000 = 4%
Princess Margaret	1,640	23-48 gene panels (Ilumina, Life)	FFPE	Yes	92/1640 = 5.6%

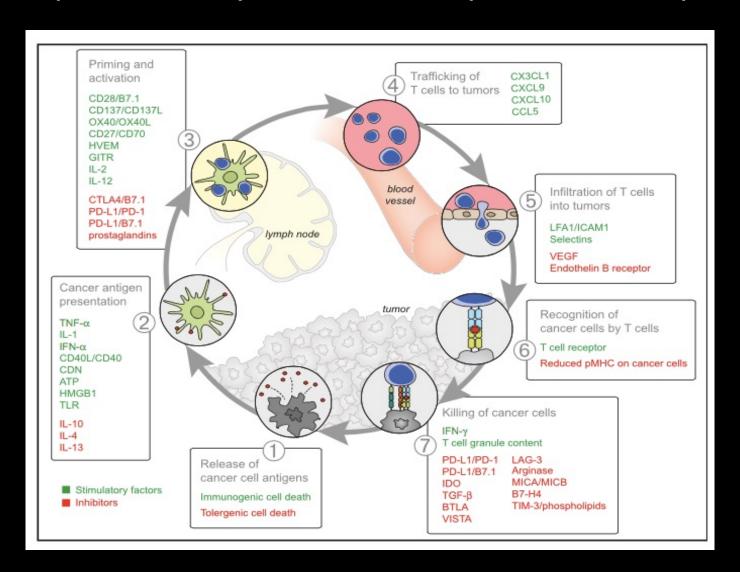
CNA = Copy number alterations; IHC = Immunohistochemistry

Ferte et al. TAT 2015; LeTourneau et al. Lancet Oncol 2015; Laskin J, et al. Cold Spring Harb Mol Stud 2015; Meric-Bernstam et al. J Clin Oncol 2015; Bedard P, et al. AACR Precision Medicine Series 2015.

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Requirements for Spontaneous or Therapeutic Immune Response

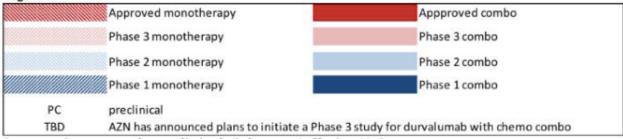


Chen and Mellman, Immunity 2013 39, 1-10DOI: (10.1016/j.immuni.2013.07.012)

Exhibit 25: Heat map of Immuno-oncology development progress by IO class and company (in-house assets)

10 class	Bristol-Myers	Merck & Co.	Roche	AstraZeneca	Pfizer	Novartis	GlaxoSmithKline	Sanofi
PD-1	-							
PD-L1								PC
CTLA-4								
Chemo combo				TBD			1	
IDO	PC	PC						
4-1BB/ CD137								
OX40								
LAG3		"						PC
GITR			.,					PC
CSF-1R								
KIR								

Legend



Source: Company data, clinicaltrials.gov, Jefferies LLC

From @SheffStation

Immunotherapy at Princess Margaret

Approx. 400 patients/ year receive immunotherapy at PM and growing

	Phase I trials: Drug targets Patient No.			Phase I trials: Drug targets	Patient No.
1	PD-1	80	15	TIM3+/-PD-1	5
2	PD-1	34	16	CSF1R+PD-1	5
3	PD-L1	32	17	PD-1	4
4	GITR+/-PD-1	21	18	PD-1+CTLA-4	3
5	PD-1	19	19	PD-1	3
6	PD-L1+OX40	16	20	PD-L1+CD40	2
7	OX40	13	21	PD-1	2
8	LAG+/-PD-1	12	22	CD40+ANG2	2
9	PD-L1+CTLA-4	9	23	4-1BB+PD-L1	2
10	PD-1+VEGF	9	24	CD73+PD-1	2
11	PD-1+CTLA-4 or VEGF	8	25	PD-L1 + MEK	1
12	PD-L1	8	26	TIGIT+PD-L1	1
13	IDO+PD-1	7	27	GITR+PD-L1	1
14	PD-L1	5	28	ICOS+PD-1	1
				Total	307

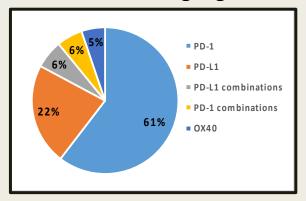
Development of the Princess Margaret Immune Oncology Prognostic Index (PM-IPI): A novel prognostic score for patients treated in immune oncology phase I trials

Results

Baseline patient characteristics

	No. of patients	%
Sex		
Male	107	56%
Female	85	44%
Age	median 57.5 (ra	ange 20.4-84.8)
ECOG PS		
ECOG PS 0	76	40%
ECOG PS 1	116	60%
Primary tumor site		
Melanoma	52	27%
Thoracic	41	21%
Genitourinary	22	11%
Head and neck	20	10%
Sarcoma	14	7%
Gynecologic	13	7%
Gastrointestinal	18	8%
Breast	8	4%
Other	6	3%
No. of prior systemic therapies	median 2 (range 0-8)
No. of metastatic sites	median 3 (range 0-7)
≤2 sites	86	45%
>2 sites	106	55%
Sites of metastasis		
Lung	123	64%
Liver	74	39%
Bone	52	27%
Brain	23	12%

PI IO trials: Drug targets



Patient outcomes (n=192)

· Median PFS: 13.4 weeks

Median OS: 73.6 weeks

90DM: 16%

• ORR: 20% by RECIST 1.1/ irRECIST

Dai et al. ASCO 2016

Results

- Multivariate analysis: Independent prognostic factors
- ECOG PS \geq 1 (HR 3.2, p < 0.001)
- No. of metastatic sites > 2 (HR 2.0, p = 0.003)
- Albumin < lower limit of normal (HR 1.8, p = 0.007)



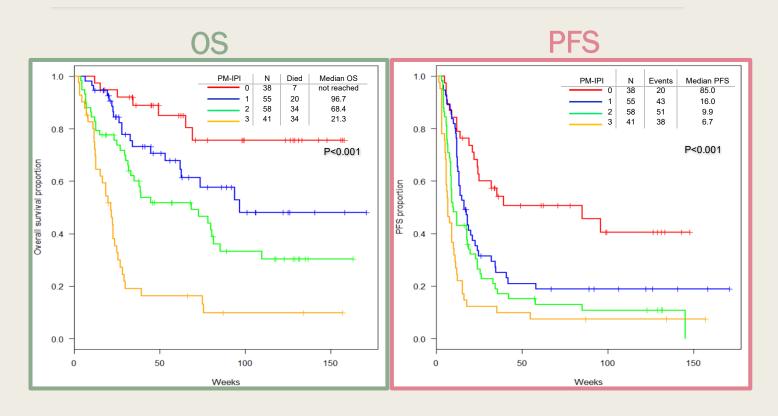
- Patients with a score of 2-3 compared to patients with a score of 0-1:
- Shorter OS (HR 3.4, p < 0.001)
- Shorter PFS (HR 2.3, p < 0.001)
- Higher 90DM (OR 8.1, p < 0.001)
- Lower ORR (OR 0.4, p = 0.019)
- Comparison of PM-IPI with previously published P1 prognostic scores

		PM-IPI	RMI	PMHI	NS	HS
OS	(C-index)	0.71	0.65	0.69	0.59	0.59
PFS	(C-index)	0.66	0.63	0.63	0.58	0.58
90DM	(AUC)	0.75	0.69	0.73	0.68	0.70
ORR	(AUC)	0.64	0.59	0.64	0.58	0.56

RMI: Royal Marsden Index; PMHI: Princess Margaret Hospital Index; NS: Nijmegen Score; HS: Hammersmith Score; AUC: Area under the curve

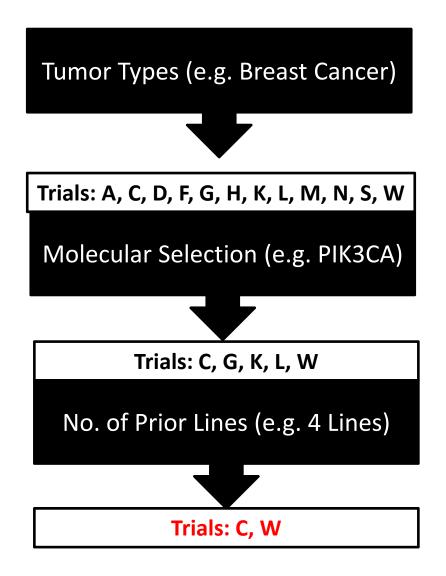
0.5 = no discriminative ability; 1.0 = perfect discriminative ability

Results



Optimization of Phase I Referral Process Developing an App to Assist with Trial Allocation





Mobile Application

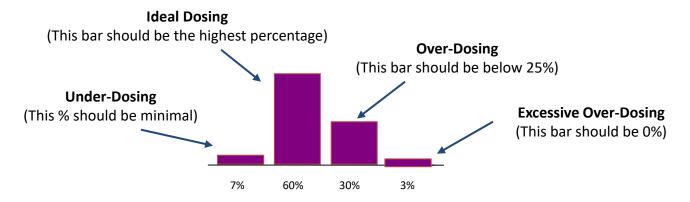


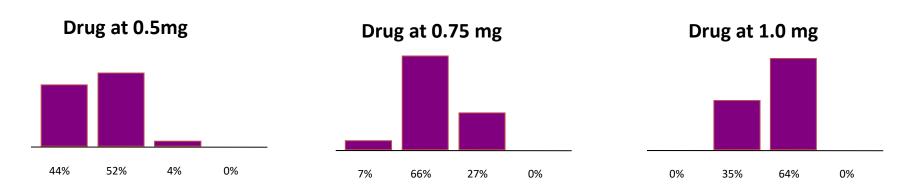
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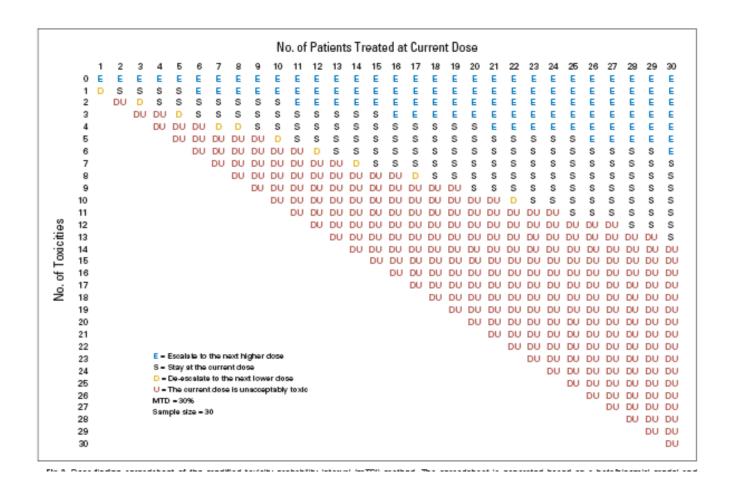
Estimated MTD Based on Bayesian Logistic Method (2-parameter evaluation with over-dose control)

EXAMPLE of Probability of DLTs (Bayesian Design)





Modified Toxicity Probability Interval (mTPI) Design



In Your Opinion, What is the Most Appropriate Dose Escalation Method for these Examples

Example	Most Appropriate Dose Escalation Method
A monoclonal antibody without a valid pharmacodynamic biomarker for optimal biological activity and likely will not have an MTD	
2. Combination of radiation with a new drug with concern for delayed/late toxicity	
3. Combination of radiation with a new drug with minimal concern for interaction or toxicity	
4. A first-in-class new drug with no obvious concerns raised by preclinical data	
5. A first-in-class new drug with likely a narrow therapeutic index	
6. Combination of two drugs each with its own RP2D with unknown risk of interaction	

Purpose of Tumor Biopsies

Diagnostic Tumor Biopsies:

 To establish a clinical diagnosis and to perform validated prognostic or predictive markers for clinical management

Post-Diagnostic Tumor Re-Biopsies:

- To measure a biomarker that can be used to guide clinical management (e.g. integral biomarker – KRAS in CRC):
 - Insufficient tumor from archival sample
 - To obtain current tissue due to concern for clonal evolution
- To perform research (e.g. integrated or exploratory biomarkers – ↓phospho-S6 as a measure of PI3K pathway inhibition)

Patient Attitudes Towards Genomic Testing in Cancer (GTC) (n = 98 patients referred for genomic testing or phase I trials)

Item	Yes	No	Unsure
Would you be interested in leaning more about GTC?	76%	6%	17%
Would you be willing to undergo needle biopsy if required for GTC?	66%	13%	19%
Would you be willing to undergo surgical biopsy if required for GTC?	39% 27%		33%
Do you believe GTC would significantly improve your cancer care?	64%	5%	30%
Would you want disclosure of incidental GTC results			
regarding: a) Inherited familial risk of developing cancer	87%	5%	7%
b) Inherited risk of developing diseases other than cancer	79%	7%	13%
Would you consent to biobank your GTC results and tissue sample for future scientific research?	91%	2%	5%

Patients' Willingness to Undergo Multiple Tests in a Single Trial (n = 61)

Geometric Mean±SD^a

No. of Tests	PET Scan	CT Scan	X-Ray	MRI	Ultrasound	Echocardiogram				
1	9.1±1.4	9.5±1.3	9.7±1.4	7.9±1.9	9.7±1.2	9.4±1.4				
2	8.2±1.6	8.8±1.5	9.0±1.5	6.6 ± 2.2	9.2±1.4	8.4±1.8				
3	7.1±1.9	7.1±2.1	8.1±1.8	5.9 ± 2.3	8.3±1.7	7.8±2.0				
4	$6.4{\pm}2.2$	6.7±2.1	7.3±2.1	5.5 ± 2.4	7.9±1.8	7.7±2.0				
P test for trend	<.001	<.001	<.001	<.001	.003	.010				
	Geometric Mean±SD ^a									
No. of Tests	Skin Biopsy	Tumor Biopsy	Blood Sample	Hair Follicle	Stool Sample	Urine Sample				
1	8.1±1.6	7.6±1.9	9.9±1.2	8.6±1.7	9.1±1.3	9.9±1.2				
2	6.1±2.2	5.8±2.2	9.6±1.3	7.9 ± 1.9	8.0±1.8	9.9±1.2				
3	5.3±2.3	4.6±2.3	9.0±1.5	7.5±2.0	7.4±1.9	9.3±1.6				
4	4.6 ± 2.5	4.0 ± 2.4	8.4±1.7	7.2 ± 2.0	7.3 ± 1.9	9.3±1.6				
P test for trend	.001	.001	.001	.001	.001	.045				

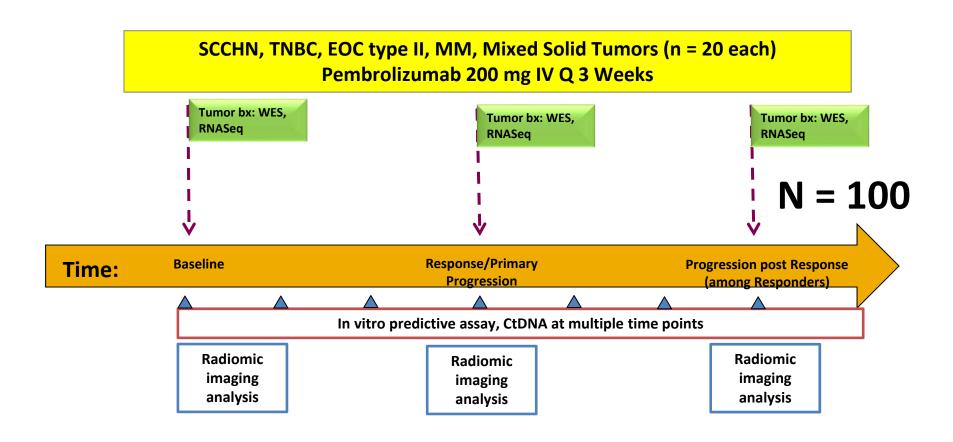
SD indicates standard deviation; PET, positron emission tomography; CT, computed tomography; MRI, magnetic resonance imaging.

^a Answers were scored on a scale from 1 (not willing) to 11 (very willing). The scale range was recoded from a 0-to-10 scale to a 1-to-11 scale by adding 1 to each score to accommodate calculation of the geometric mean and SD.

Research Biopsies

- Identified 22 phase I trials from 2003-2006 which included post-treatment biopsies for PDbiomarkers
- 9/22 studies (41%) tested >4 PD-biomarkers
- Statement on impact on future studies found in 9/22 studies (41%)
- None of the PD-biomarkers impacted phase II/III dose or schedule

INvestigator-initiated Phase II Study of Pembrolizumab Immunological Response Evaluation (INSPIRE)



INSPIRE-A-002

Flow Cytometry Panel #1 (T cell analysis)

CD3 T cells

CD8 CTL

CD4/ CD19 Helper cells / B cells

CD56 NK cells

TcR $\gamma\delta$ $\gamma\delta$ T cells

PD-1 exhaustion

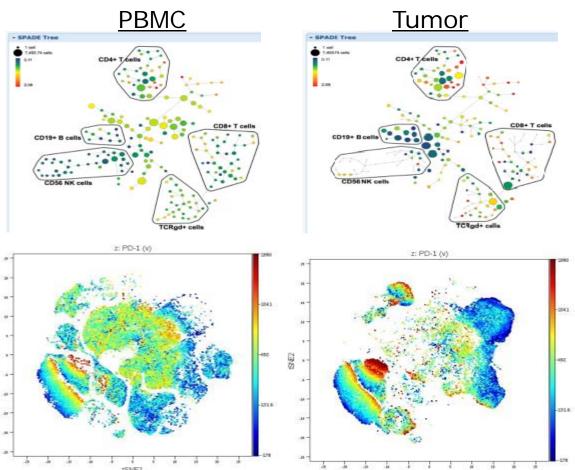
TIGIT exhaustion

PDL1 exhaustion

CTLA4 exhaustion

4-1BB co-stimulation

Unpublished data



The Future Drug Development Paradigm?

Histology + Molecular Selection

Proof of Concept

- Safety, tolerability
- Functional target selection
- Pharmacology
- Antitumor activity

- Substantial efficacy in selected pt populations using innovative trial designs and endpoints
- Trial design accounting for <u>interpatient</u>
 <u>and intratumor heterogeneity</u>

Conclusions

- Phase I trials are playing an increasingly critical role for go-no-go in drug development
- Many emerging features have arisen out of the need to find rare molecular patient subsets, expedite drug development, incorporate promising emerging agents (e.g. IO), while preserving safety in our conduct of phase I trials
- We need to keep key stakeholders (patients, IRB members, referring physicians, study team members) informed and engaged as phase I trials evolve in the drug development paradigm

Phase I Team at the Princess Margaret Cancer Centre

