

Master Protocol in Pediatric Cancer Trials

Jingjing Ye, PhD
BeiGene

ASA Biopharmaceutical Regulatory/Industry Statistical
Workshop 2020

Sep. 24, 2020

The presentation is based on the work while employed at FDA

Disclaimer: The presentation represents the opinion of the presenters, and do not reflect the position of the U.S. Food and Drug Administration, nor BeiGene

Acknowledgement

Gregory Reaman, OCE/FDA

Nita Seibel, NCI

D. Will Parsons, NCI-COG, Baylor College of Medicine

Peter O'Dwyer, ECOG-ACRIN, Abramson Cancer Center, Upenn

Peter Adamson, COG

Slides courtesy from all

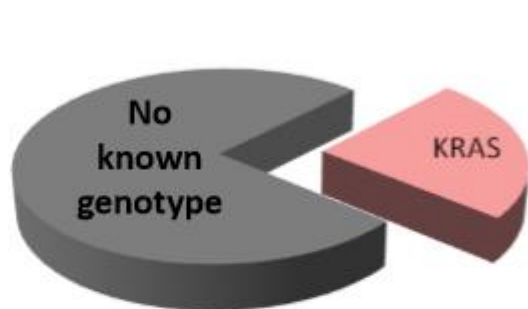
Outline

- Background
 - Precision Medicine and Pediatric Oncology Drug Development
 - Master Protocol Guidance
- NCI-COG Pediatric MATCH Design and Structure
- Challenges and Opportunities
- Summary

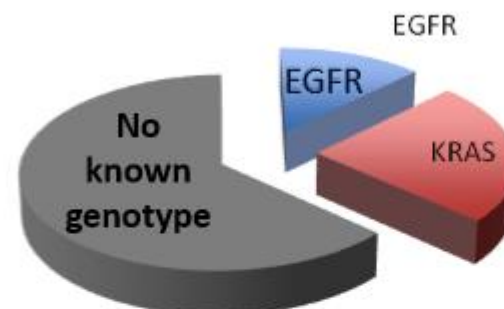
Precision Medicine and Oncology Drug Development

- Precision cancer medicine: **targeted therapy** selection by identifying **key gene variants**
- Evolutionary Paradigm shift: **Human genome** (2003) – wide-spread availability of NGS
- Genomic and proteomic interrogation of individual cancers screened: resulted in creation of **multiple rare subsets** (defined by **molecular phenotype**) of previously common cancers

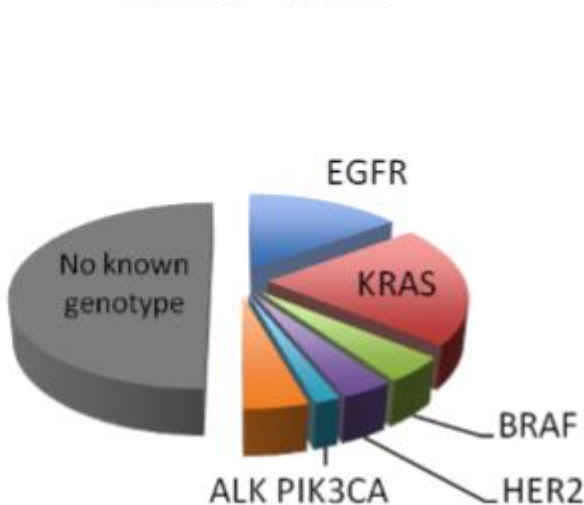
Evolution of Identification of Genomic Alterations in Lung Adenocarcinoma



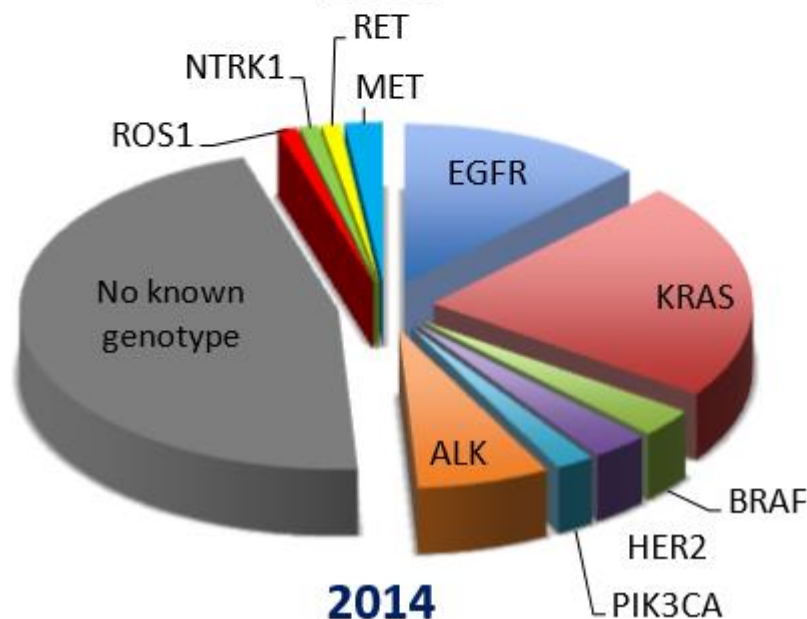
1984 - 2003



2004

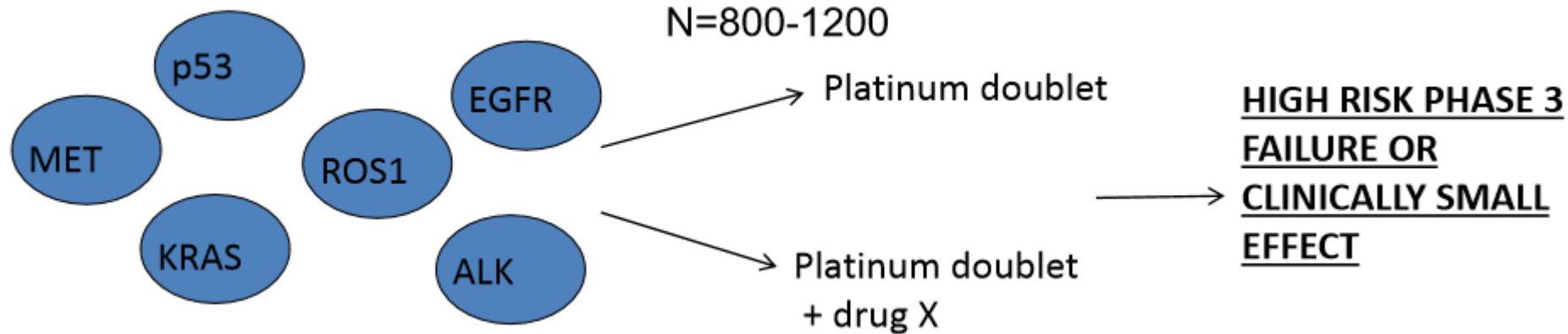


2009

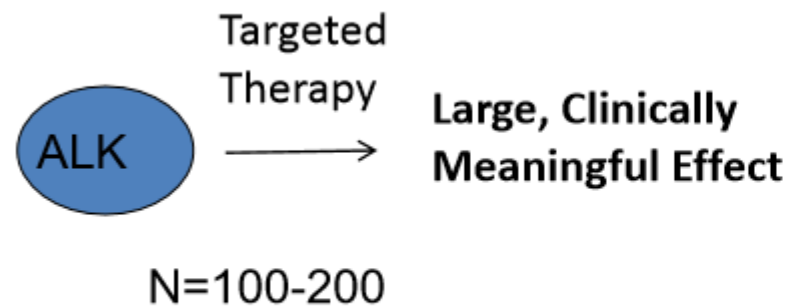


2014

Challenges with “Old Paradigm”



Challenges with “New Paradigm”



- 1% Prevalence of even common tumors: Number needed to screen > 100 patients → need to reduce screen failure rate
- 1 drug/ 1 biomarker per trial unsustainable → Need common multi-analyte platform(s)
- Need Rapid Learning/ Failure/ Confirmation

Precision Medicine in Pediatric Oncology

- Most childhood cancers (embryonal origin) – low mutation frequency
- Initial therapy (H.D. chemo/XRT)
- Some childhood cancers have very few recurrent events
- Post-therapy sequencing of relapse samples accumulate more mutations in targetable oncogenic pathways
- Few opportunities for extrapolation: 5 out of 40 written requests in 2001-2019*

*A review of the experience with pediatric written requests issued for oncology drug products, submitted and under review

Characteristics of an Ideal Master Protocol

- One protocol
- Central governance structure
- Central IRB
- Central DMC
- Central Independent Review Committee
- Central repository of Data and Specimens
- Central screening platform
- Study multiple drugs
 - Targeting more than one marker
 - More than one drug for one marker
- Study multiple markers
 - Overlapping expression of markers
- Leverage common control
- Flexibility to add/remove agents (Adaptive)

Background: NCI-MATCH – Genomically-Driven Trial 2013

A Disease Agnostic Basket Trial: NCI-MATCH

THIS PRECISION MEDICINE TRIAL
EXPLORES TREATING PATIENTS
BASED ON THE MOLECULAR
PROFILES OF THEIR TUMORS

NCI-MATCH¹ IS FOR ADULTS WITH:

- solid tumors (including rare tumors), lymphomas, and myeloma
- tumors that no longer respond to standard treatment



NCI-COG Pediatric MATCH Trial

- Funded by NCI
- Developed jointly by NCI and Children's Oncology Group (COG)
- Conducted by COG
- Refractory and recurrent pediatric Solid tumor, including non-Hodgkins lymphomas and CNS tumors or histiocytosis
- Goal: deliver targeted anticancer therapy that produces a clinically meaningful objective response rate

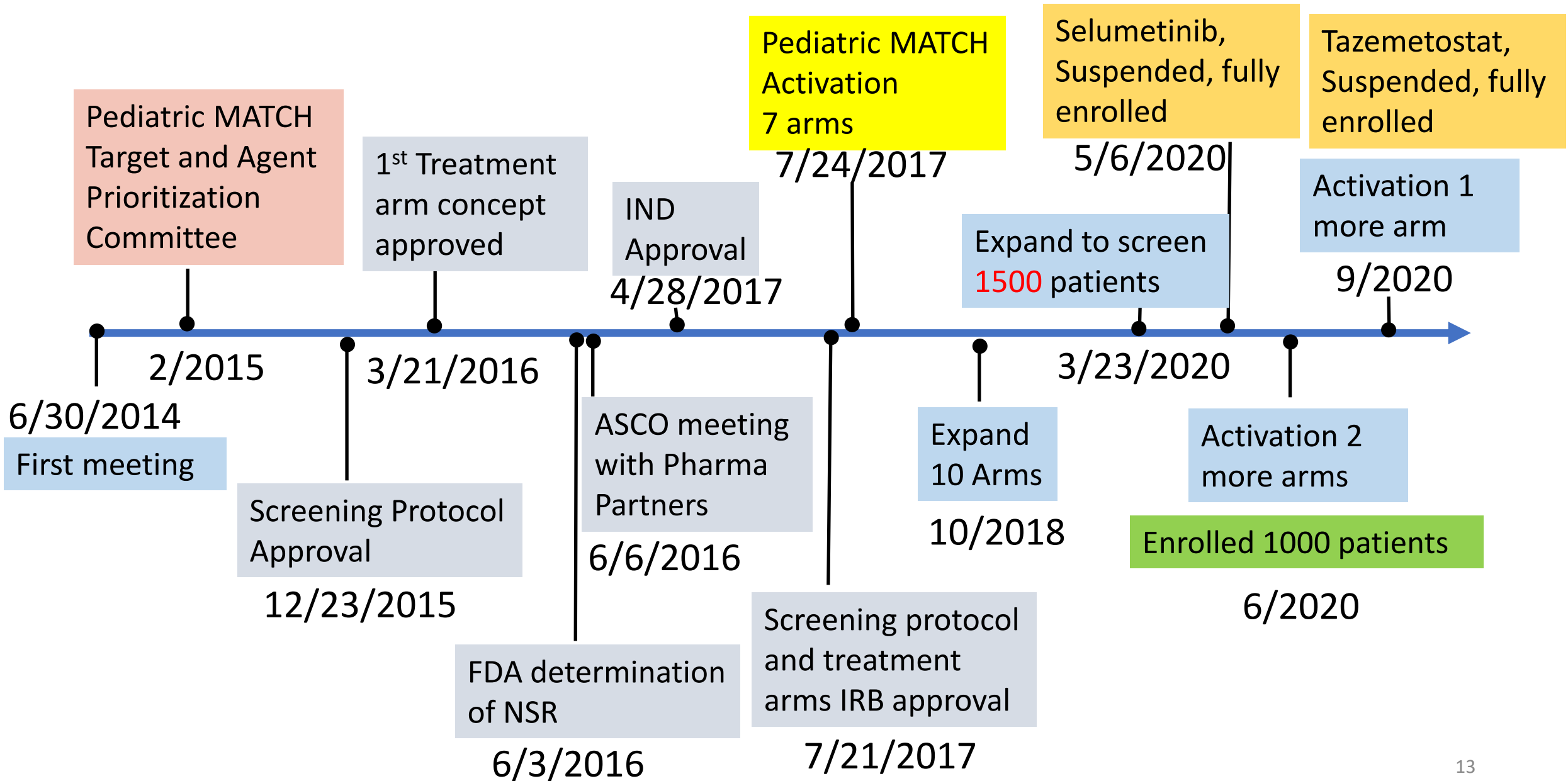
NCI-COG Pediatric MATCH Master Protocol

- Single IND held by CTEP
- Central Governance Structure
- NCI Pediatric Central IRB
- Master Protocol review by CIRB, each marker-drug subprotocol
- Central DMC by COG
- Central screening platform, leverage Adult MATCH trial
- Central Repository of data and specimen

NCI-COG Pediatric MATCH Design Features

- Non-histology driven
- Test many children and adolescents to find widely distributed genetic alterations
- Biopsies from the time of recurrence except for DIPG (from dx)
- Single stage Phase 2 studies
- Inclusion of agents with **adult** RP2D, without formal pediatric phase I testing
- Blood sample acquisition and return of germline sequencing results related to inherited cancer susceptibility

NCI-COG Pediatric MATCH Timeline



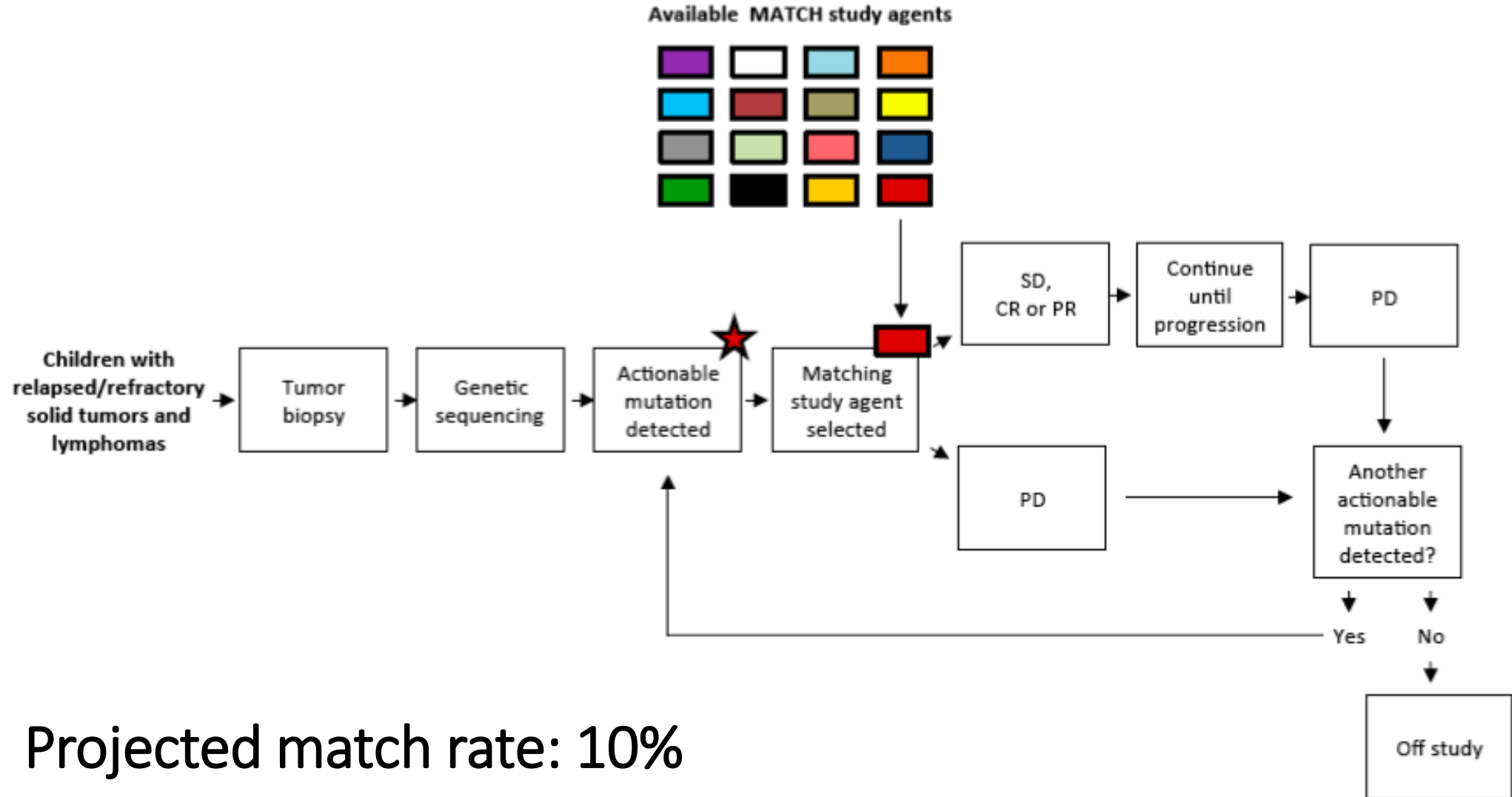
Level of Evidence for **Drugs**

- Level 1: FDA approved for any indication for that target
- Level 2: Agent met a clinical endpoint (objective response, PFS or OS) with evidence of target inhibition
- Level 3: Agent demonstrated evidence of clinical activity with evidence of target inhibition at some level

Level of Evidence for Target Selection

- Level 1: Gene variant credentialed for selection of an approved drug
- Level 2a: Variant is eligibility criteria for an ongoing clinical trial for that drug
- Level 2b: Variant identified in an N of 1 response (s)
- Level 3: Preclinical inferential data
 - Models with variant response; without variant do not
 - Gain of function mutation demonstrated in preclinical model
 - Loss of function (tumor suppressor genes or pathway inhibitor e.g. NF1); stop codon or demonstrated loss of function in preclinical model

NCI-COG Pediatric MATCH Schema



Projected match rate: 10%

Later re-match can occur at physician's request (if slots are available)

Study Designs and Statistical Consideration

- Primary endpoint: Objective Response Rate
- Secondary endpoint:
 - Progression-free survival (PFS)
 - Tolerability
 - PK: erdafitinib, ensartinib, LY3023414, ivosidenib, ulixertinib
- Simon's 2-stage Design, no interim stopping
- Evaluation per arm: ORR and DOR, PFS
- No cross-arm comparison or pooling
- N=20/arm

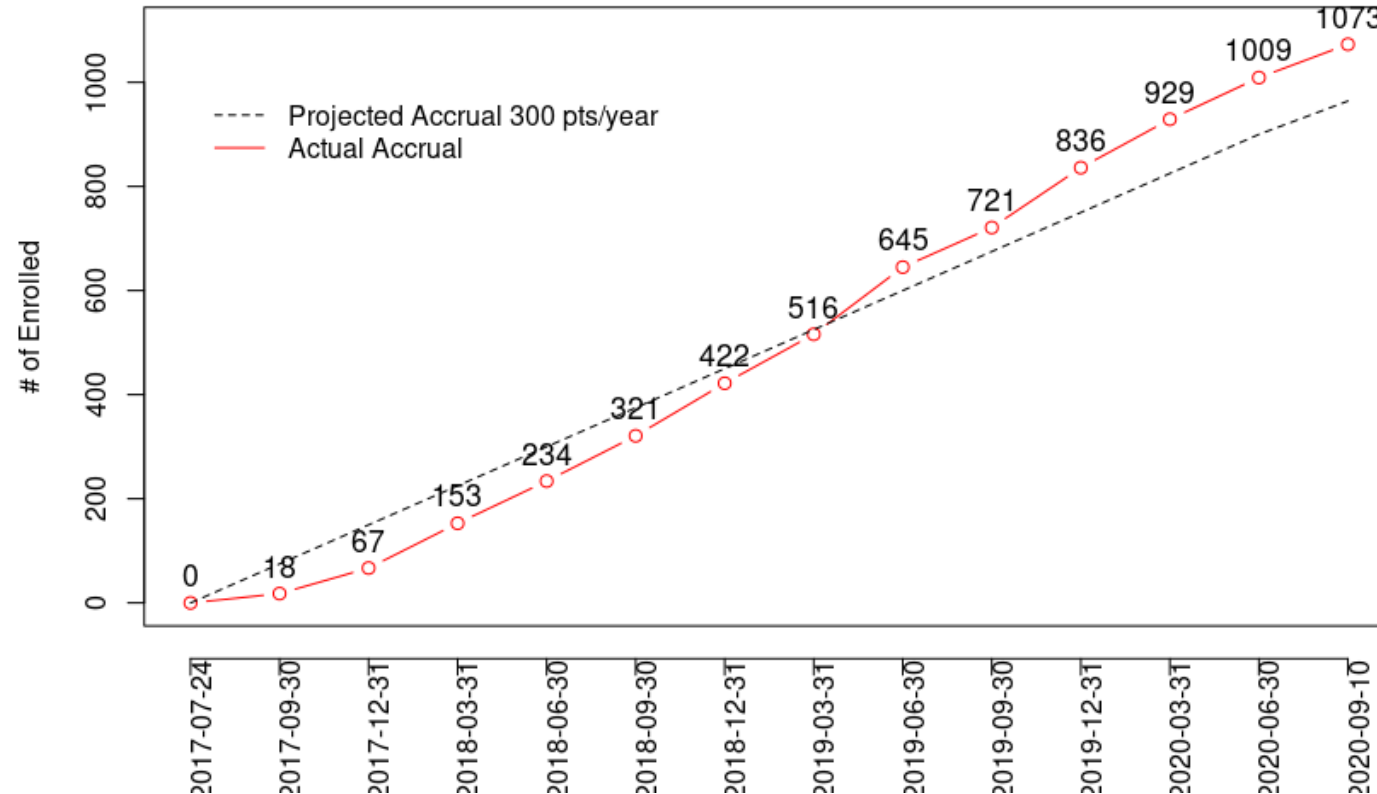
Treatment Subprotocol Status

Protocol ID	Agent	Agent Class	aMOI Frequency	Activation Date	Current Status	Adult Approval
APEC1621-A	Larotrectinib	TRK inhibitor	2-3%	7/24/2017	Recruiting	11/26/2018
APEC 1621-B	Erdafitinib	FGFR inhibitor	2-3%	11/06/2017	Recruiting	4/12/2019
APEC 1621-C	Tazemetostat	EZH2 inhibitor	2-3%	7/24/2017	Suspended, full enrollment – Planned analysis	1/23/2020
APEC 1621-D	LY3023414	PI3K/mTOR inhibitor	5-10%	7/31/2017	Recruiting	
APEC 1621-E	Selumetinib	MEK inhibitor	10-20%	7/24/2017	Suspended, full enrollment – Planned analysis	4/10/2020
APEC 1621-F	Ensartinib	ALK inhibitor	2-3%	7/24/2017	Recruiting	
APEC 1621-G	Vemurafenib	BRAF inhibitor	5%	7/24/2017	Recruiting	8/17/2011
APEC 1621-H	Olaparib	PARP inhibitor	2-3%	7/24/2017	Recruiting	12/19/2014
APEC 1621-I	Palbociclib	CDK4/6 inhibitor	2-3%	6/25/2018	Recruiting	2/3/2015
APEC1621-J	Ulixertinib	ERK1/2 inhibitor	5-10%	10/01/2018	Recruiting	
APEC1621-K	Ivosidenib	IDH1 Inhibitor	1-2%	6/08/2020	Recruiting	7/20/2018
APEC1621-M	Tipifarnib	HRAS Inhibitor	1-3%	7/13/2020	Recruiting	
APEC1621-N	Selpercatinib	RET inhibitor	1-2%	9/14/2020	Recruiting	5/8/2020 ¹⁸

Screening Protocol Enrollment

- 1074 patients from 100 COG sites enrolled between 7/24/2017 and 9/10/2020

Subjects enrolled



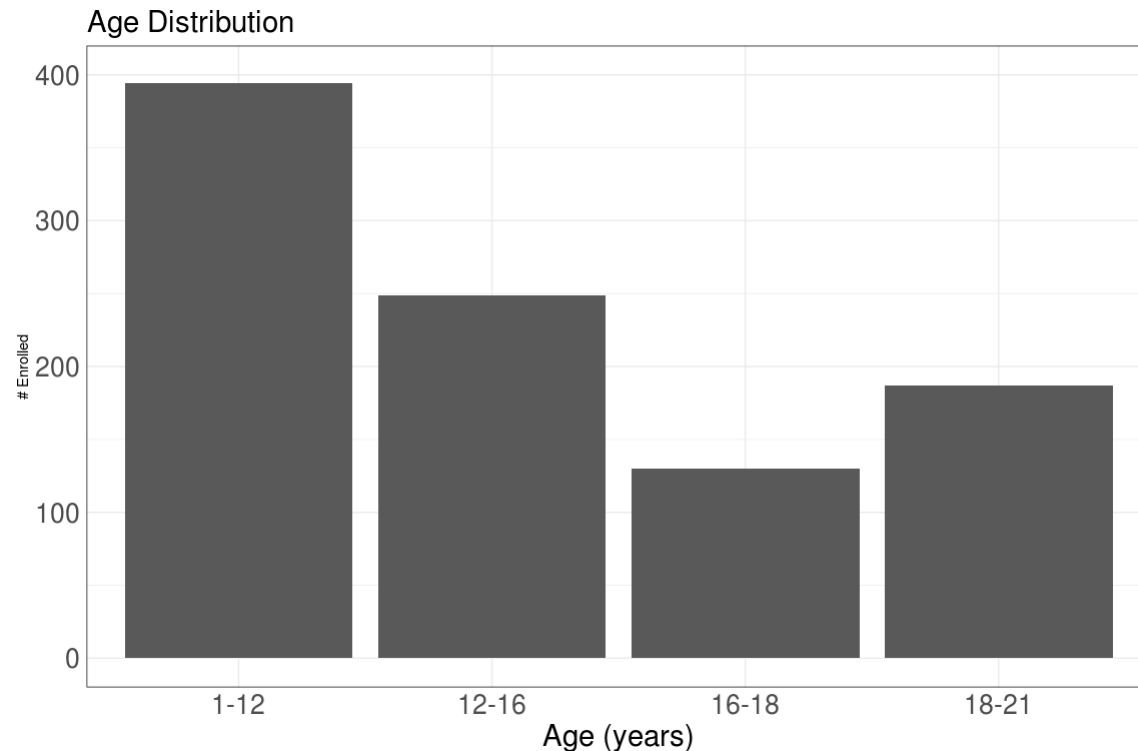
COVID-19

	Dates	N
Q1	7/24/17-9/30/17	18
Q2	10/1/17-12/31/17	49
Q3	1/1/18-3/31/18	86
Q4	4/1/18-6/30/18	81
Q5	7/1/18-9/30/18	87
Q6	10/1/18-12/31/18	101
Q7	1/1/19-3/31/19	94
Q8	4/1/19-6/30/19	129
Q9	7/1/19-9/30/19	76
Q10	10/1/19-12/31/19	115
Q11	1/1/20-3/31/20	93
Q12	4/1/20-6/30/20	80
Q13	7/1/20-9/10/20	64

Screening Protocol Enrollment (as of 4/30/2020)

- Patient age
 - Age range 1 to 21 years (median = 13)
 - 59% of patients from 12-21 years

Patients enrolled



Patient sex, race, ethnicity

Gender	Number (%)
Male	553(58)
Female	407 (42)
Racial	
White	657(68)
Black	134 (14)
Asian	38 (4)
American Indian/Alaskan Native	3 (0.3)
Multiple	15 (2)
Unknown/not reported	105 (11)

Regulatory Agency Discussion in NCI-COG Pediatric MATCH

- Discussion initiated early in the development
- Under one new IND: subprotocol reviewed by different division based on disease
- Alignment on biomarker-driven targeted therapies
- Alignment on testing in pediatric when toxicity acceptable
- Alignment on starting dosing once RP2D in adult determined
- Evaluation by arm
- No IDE required – reviewed as Adult MATCH trial

Challenges and Opportunities

- Existing clinical trial infrastructure
- Abundance of targeted agents
- Biopsy requirement for eligibility
- Evolving standard of care and comparator selection
- Combinations
- Safety oversight and monitoring

Summary

- NCI-COG Pediatric MATCH: collaborative framework for efficient collection, processing and sequencing of refractory pediatric cancers
- ~25% of study patients with tumor submitted assigned to a treatment arm, with 40% enrolled on the trials
- Ability to evaluate a wide spectrum of childhood cancers (from common to ultra-rare)

Summary

- Master protocols expand the promise of Precision Oncology to children
- Efficient mechanism for evaluating novel agents (dose-finding and activity screening)
- Biomarker-driven tissue agnostic cancer drug development strategies must include children
- Early communication with both CDER and CDRH on study design and research use of IVDs and IDE

References

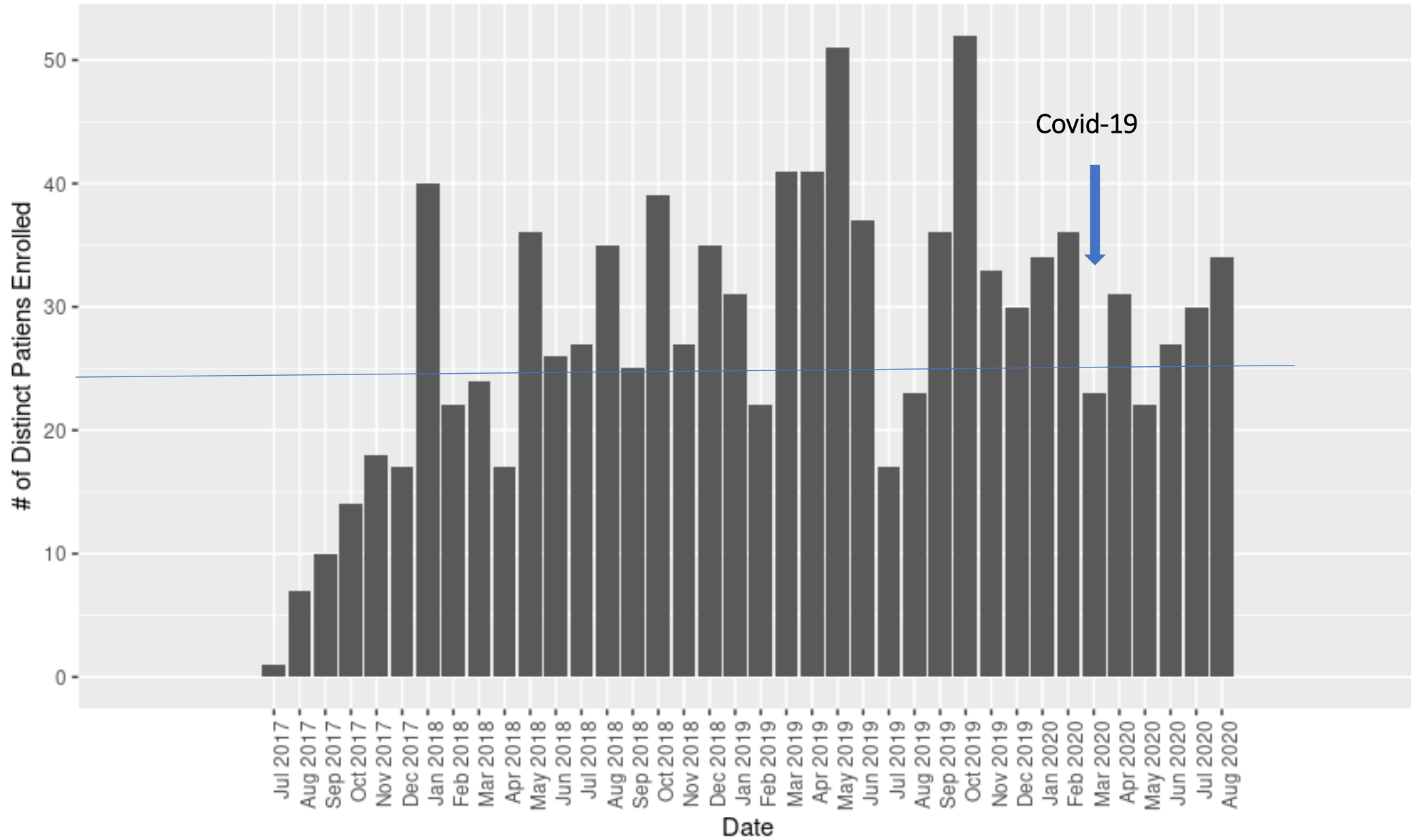
- FDA draft guidance: Master Protocols: Efficient Clinical Trial Design Strategies to Expedite Development of Oncology Drugs and Biologics, 2018
- Vo, K.T., Parsons, W., Seibel, N.L., (2020) Precision Medicine in Pediatric Oncology, *Surg Oncol Clin N Am* 29: 63-72.
- Allen, C.E., et al. (2017) Target and Agent Prioritization for the Children's Oncology Group- National Cancer Institute Pediatric MATCH Trial, *JNCI*, 109 (5): djw274.
- Parsons, et al. (2019) Identification of targetable alterations in the NCI-COG Pediatric MATCH trial, *Journal of Clinical Oncology* 37, no. 15_suppl (May 20, 2019) 10011-10011, ASCO abstract 10011 and presentation
- Adamson, P.C., Parsons, D.W., Seibel, N., (2017), NCI-COG Pediatric MATCH Study, Accelerate Platform 5th Accelerate Paediatric Oncology Conference, <https://www.accelerate-platform.org/wp-content/uploads/sites/4/2017/03/17-Adamson.pdf>
- Mody, R.J., et al. (2015) Integrative clinical sequencing in the management of refractory or relapsed cancer in youth, *JAMA*, 314 (9): 913-925.

References (Cont'd)

- Reaman, G, Master Protocols in Pediatric Oncology: Access to Precision Medicine, Pediatric Master Protocols, FDA-University of Maryland CERSI Co-sponsored Workshop, Sep. 23, 2016
- Harris, M.H., et al. (2016) Multicenter Feasibility Study of Tumor Molecular Profiling to Inform Therapeutic Decisions in Advanced Pediatric Solid Tumors, The individualized cancer therapy (iCat) study, JAMA Oncology, 2(5): 608-615.
- Flaherty, K.T., et al. (2020) The molecular analysis for therapy choice (NCI-MATCH) trial: lessons for genomic trial design, JNCI, doi: 10.1093/jnci/djz245
- Parsons, D.W., et al. (2016) Diagnostic yield of clinical tumor and germline whole-exome sequencing for children with solid tumors, JAMA Oncology, 2(5): 616-624
- O'Dwyer, P.J., (2019) The Evolution of NCI-MATCH: What's next for SWOG and the NCTN, Spring 2019 SWOG Group Meeting, San Francisco, CA, Apr. 24-27, 2019

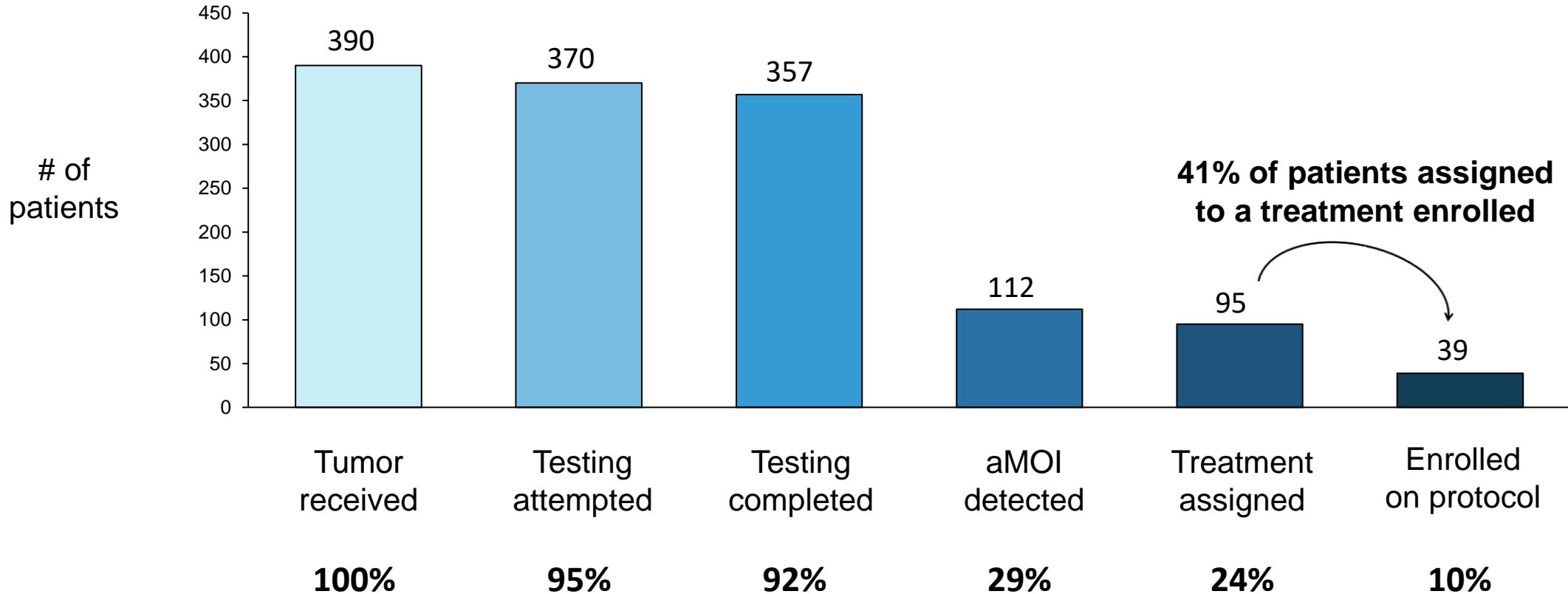
Back-up

Pediatric MATCH Enrollment



Tumor testing and matching (as of ASCO2019)

- Tumor sample was received for 390/422 (92%) enrolled patients, as of 12/31/2018
- Tumor sample was received for 909/960 (95%) enrolled patients, as of 4/30/2020



Median turnaround time (tumor receipt to assignment): 15 days

Subprotocol matching and enrollment

- 95 of 390 (24%) with tumor submitted had at least one match assigned
- 39 of 390 (10%) with tumor submitted enrolled on treatment subprotocol

Protocol ID	Agent	Matched	Enrolled	Enrolled (%)
APEC1621-A	Larotrectinib	3	3	100%
APEC1621-B	Erdafitinib	4	2	50%
APEC1621-C	Tazemetostat	9	4	44%
APEC1621-D	LY3023414	13	4	31%
APEC1621-E	Selumetinib	31	11	35%
APEC1621-F	Ensartinib	8	3	38%
APEC1621-G	Vemurafenib	7	3	43%
APEC1621-H	Olaparib	11	4	36%
APEC1621-I	Palbociclib	8	2	25%
APEC1621-J	Ulixertinib	1	0	0%