

Precision Oncology:
The Australian Genomic Cancer Medicine Centre



AGCMC's platform enables Precision Oncology

- AGCMC offers a molecular signal-seeking "live" drug development platform
- Master protocol of novel "basket" clinical trial design
- Identification of new targets and populations via high-throughput live matching of patients to cognate drugs
- Accelerated protocol development and trial completion
- World-class biospecimen-rich correlative science

Drug development Trials

- Molecular Screening
- Basket Trial Protocol Design
- Pre Phase 2 Trials
- Registrational Trials

Data Collection
Germline Trial and Data
Real world evidence
Correlative science

Government Policy Innovation
Pan Cancer Trial Design
Trial completion
Patient Advocacy

Medical education Immersion in Precision Oncology Program Fellowships



AGCMC is a risk sharing collaboration between industry, cancer research and government

Australian Genomic Cancer Medicine



- Patient centred national eco-system supported by state and federal governments
- 8 Academic Centres of Excellence & 3 medical research centres across Australia
- Focus is rare, less common, early-onset and poor outcomes cancers

Academia















Royal Adelaide Hospital Royal Darwin Hospital Royal Hobart Hospital

Princess Alexandra Hospital

Government and Societies











AGCMC is a "one stop shop" platform that collaborates with all stakeholders in precision oncology from research to patients

Value proposition for each Cancer Stakeholder

Government

- 1. Screen and treat Australians with rare cancer
- 2. Support policy in screening & treatment of cancer

Industry

- 1. Rapid molecular signal seeking drug dev platform
- 2. Rich correlative science to inform pipeline planning

Precision
Oncology Cos

- 1. Consolidated access to experts, capabilities, data & patients
- 2. End to end beta platform in precision oncology

Patient Advocacy

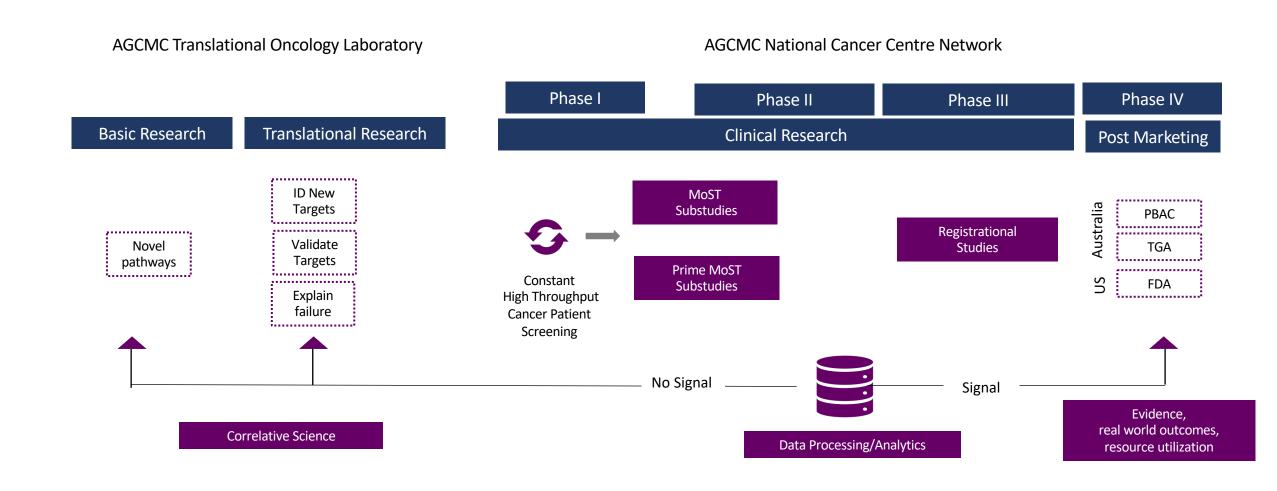
- 1. Access to screening & treatment for Australians
- 2. Informing policy to improve cancer care

Corporate Sponsors

National program that addresses a significant unmet need for Australians with rare cancer



Platform capability extends beyond research to inform discovery, expedite drug development & support approvals





MoST study is a "live" Precision Medicine Platform for biomarker driven signal seeking clinical studies

Study Rationale

Expedite testing of novel therapeutics with platform that is constantly screening patients

Progress to date in POC

- N= 1200+ Screened
- 509/806 recommended treatment
- 8 Studies
- 9 Compounds studies

Details

- 80% Rare Cancers
- Includes common cancers
- Leverages Master Protocol
- Multiple studies running on platform
- Combination studies
- Objective: Identify potential signals of activity for novel therapeutics
- Population is patients with confirmed advanced or metastatic solid cancer, failed other therapies
- Screening for molecular features
- Various panels tested
- Tests broad range of therapeutics to increase number of patients treated
- Target N= 3000 + screened



AGCMC Data enables basic research and translational research

- Biospecimen-rich correlative science driving drug development
- National capacity for 'real world' registries to support regulatory approval

Data

WGS germline data



Clinical cohort data



Tumour genomics



Correlative science



Clinical trials data



Data Processing/analytics



Final Report

Patient Profile Overview

Patients Qualify for MoST

Alternative clinical trial

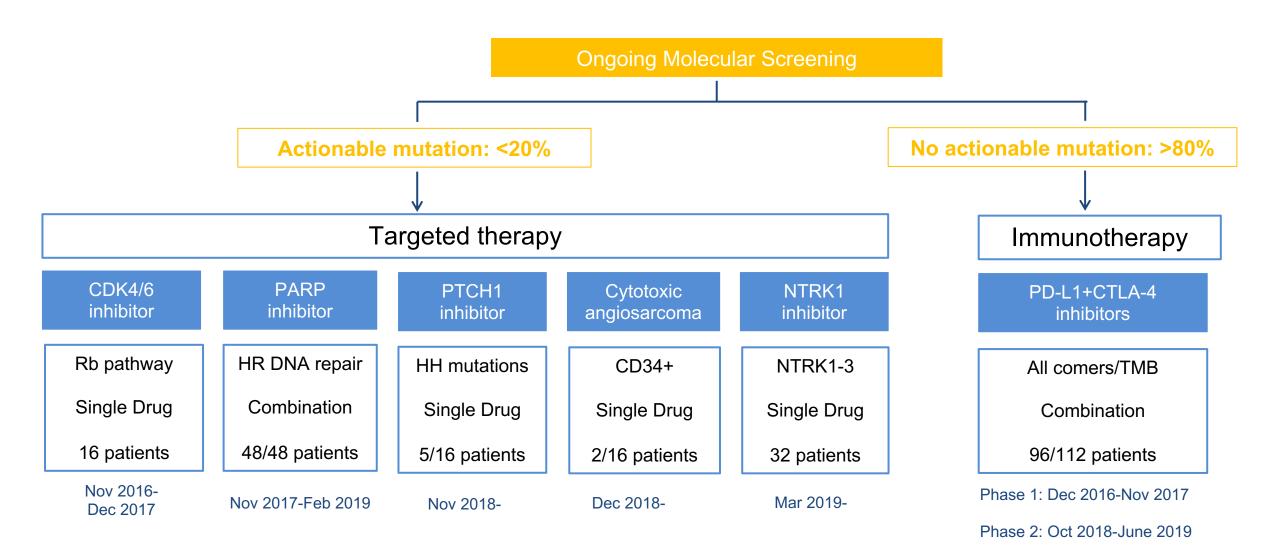
Analytics report

Genetic risk score



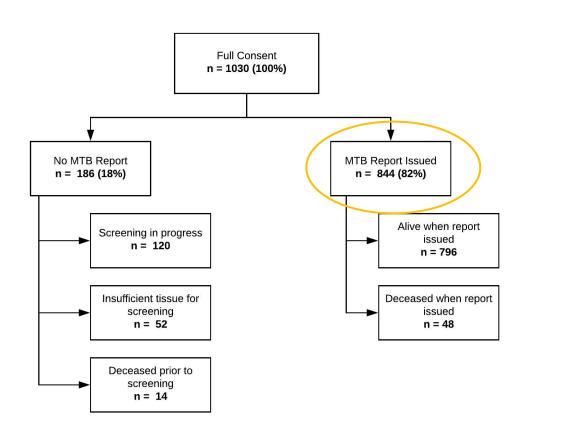


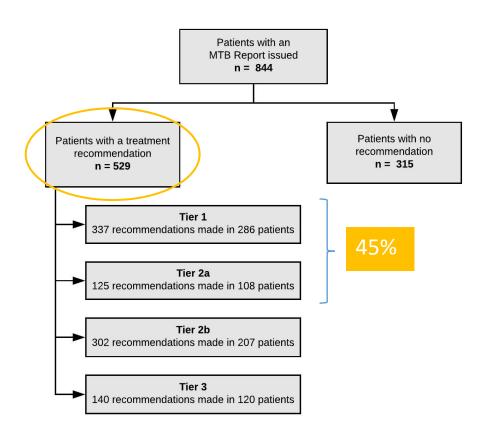
MoST launched in 2016 as a POC at one academic centre then scaled to a national Australian Precision Oncology Platform





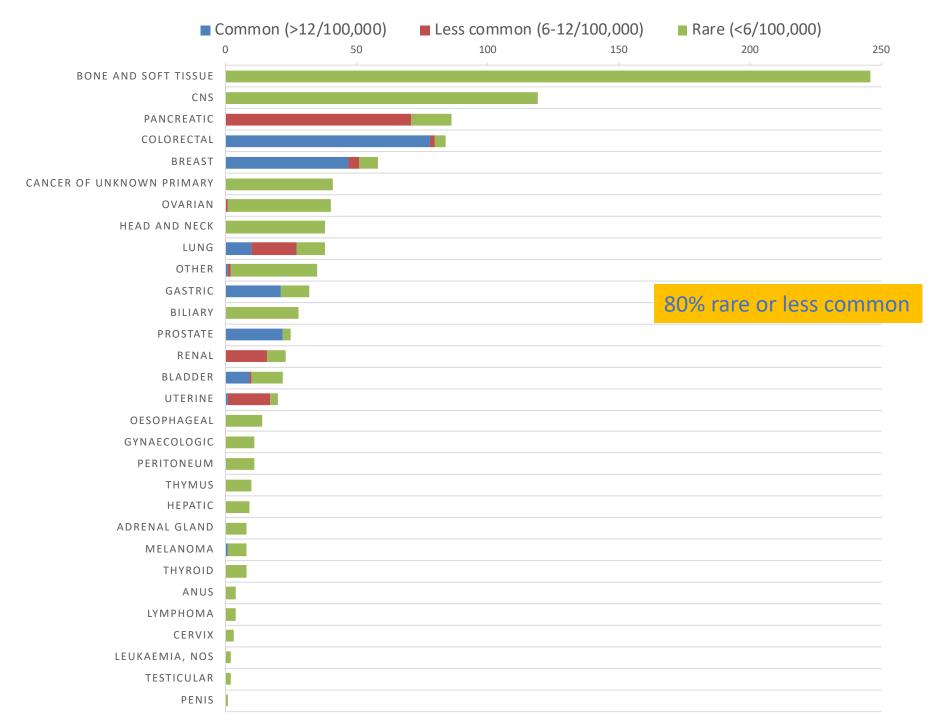
MoST progress in first 2 years





FFPE-based panel systems under evaluation:

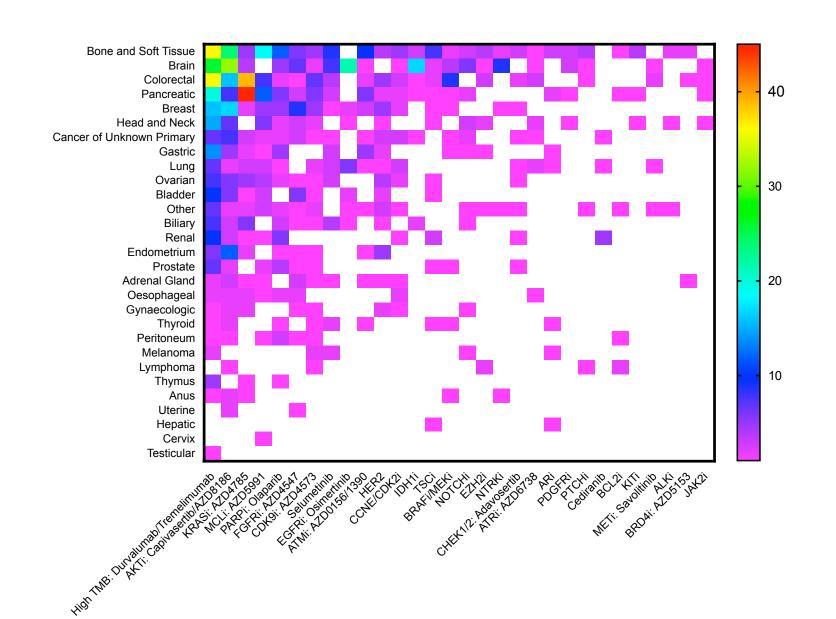
- In-house
- TST170
- Oncomine





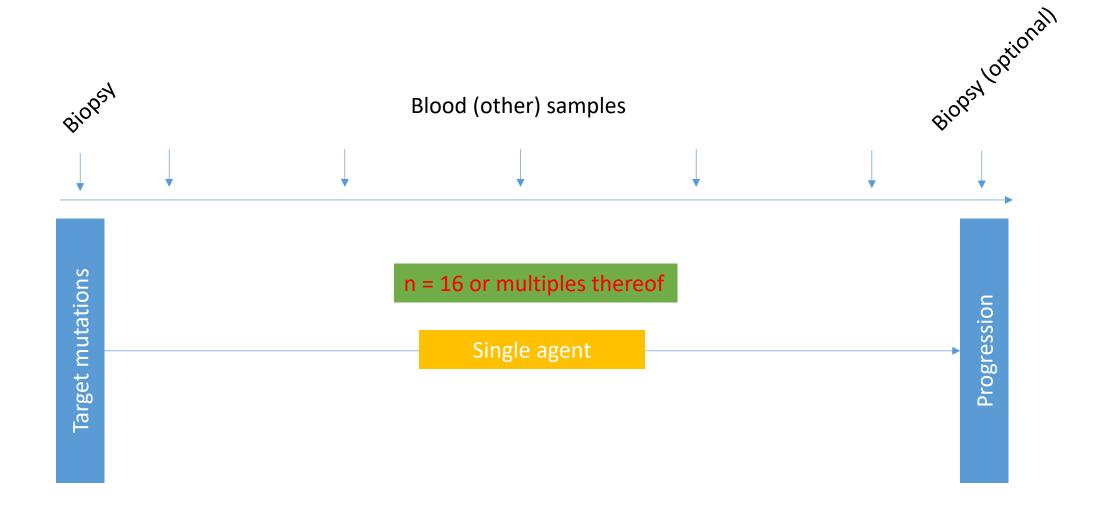
DRUG:TARGET MATRIX IN MOST PHASE 1 (n=900)





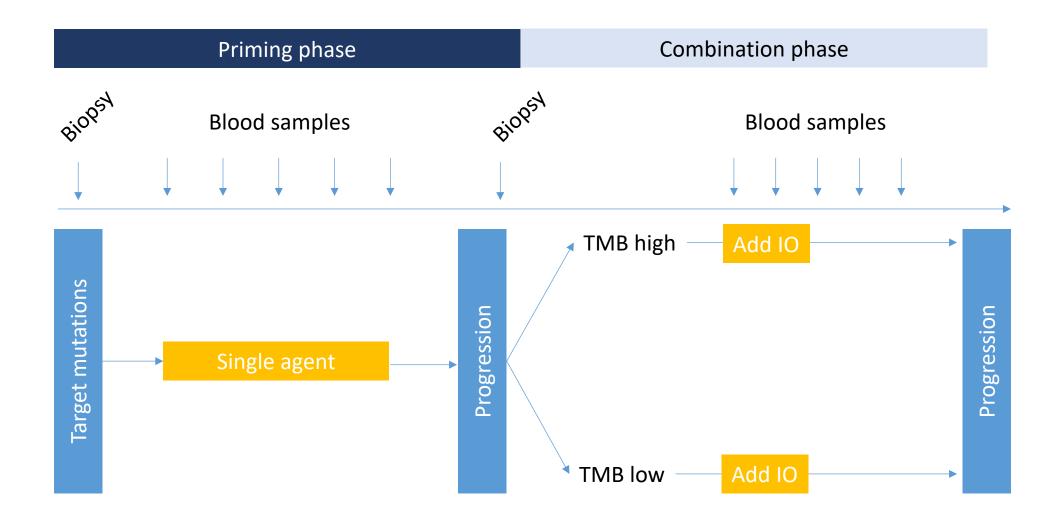


MoST design





Prime-MoST is a novel trial design for combination drug development and will launch via the platform in 2020







- Biospecimen processing and handling
- Genomics
 - DNA/RNAseq
 - TCR/BCR clonotyping
 - Long-read capacity
 - Bioinformatics
 - Microbiomics
- Proteomics
 - PCT-SWATH mass spectroscopy
- Cell-based assays



CELLULAR GENOMICS





Summary of services provided leveraging national ecosystem of cancer research centres, clinics and labs

Services	End Users	
Clinical Trials Ph 1 & 2 (Protocol Design & Dev, Monitoring, Safety)	 Industry Government Medical Community (Referrals) Patient Organizations (Referrals) 	
Registrational Trials	IndustryMedical Community (Referrals)Patient Organizations (Referrals)	
Translational Research	• Industry	
Basic Research	• Industry	
Real World Data	IndustryGovernment	
Data Processing/Analytics	 3rd Party Precision Onc Co Industry 	
Policy Support	IndustryGovernmentPatient Organization	
Education/Thought Leadership	IndustryMedical Community	



AGCMC expansion includes adding more molecules, registrational trials, and policy support

Objective	Signal Seeking	Registration Trials	PBAC/Managed Access
Key Activity	Repurposed drugsNew compoundsNew combinations	Pan Cancer Registrational Trials	Build use caseSupport with data/analyticsKOL support/Pt Advocacy
Target Outcomes	New TargetsNovel trial designsCorrelative Science	Novel Therapeutic Indications	 Expedited approval process Contextual Data Package Confirmed for Rapid Access
3-5 Year Horizon	Scale Platform to Asia:	FDA Registration	Upgrade data package:

Data Collection/Correlative Science _____

Publications

CapTCR seq for hybrid capture TCR repertoire profiling

Australian
Genomic
Cancer
Medicine

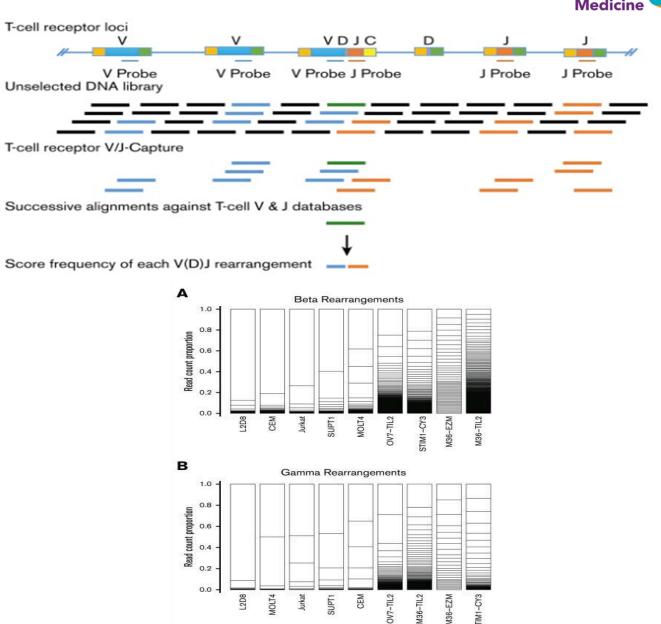
Identification of clonal population of T-cells that possess common rearrangements T-cell receptor

Method developed by Prof Trevor Pugh, Blood Advances 2018, Dec 11: 2(23) 3506

Investigate baseline PBMCs, baseline tumor and 4 week post Tx PBMCs

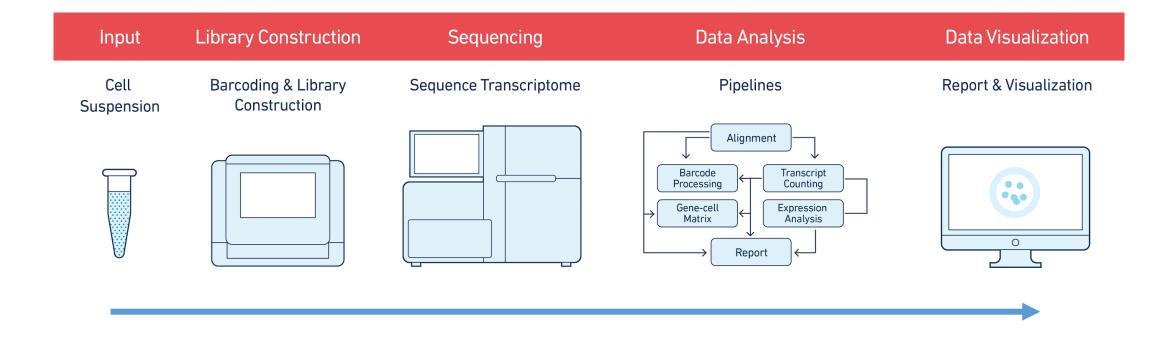
A recent study shows that peripheral blood TCR repertoire may facilitate patient stratification for ICB in melanoma, Hogan, SA et al Cancer Immunology Res, 2018.

Sequencing has been done, analysis underway





Established processed for generating single cell sequence data



We have currently sequence on average 150,000 cells per week

We have capacity to generate data for 50 patient samples per week, although this can be easily increased with additional staffing