

Outsmarting cancer, together

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From genomics to clinical impact,  
and a sustainable future for precision oncology



## Molecular screening & therapeutics

Tumour profiling to evaluate biomarker-driven treatments for patients.

## Health system reform

Leading health system reform through evidence.



Omico

A national precision oncology network.  
Connecting leading specialists, researchers,  
government, industry partners and community.  
An innovative model for research-lead  
cancer care. Improving cancer  
outcomes for all Australians.

## Personalised risk management

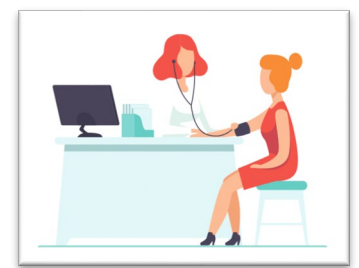
Using heritable genetic information to assess cancer predisposition and investigate clinical risk management.

## Patient support & advocacy

Supporting patients and families today and planning the health system for tomorrow.



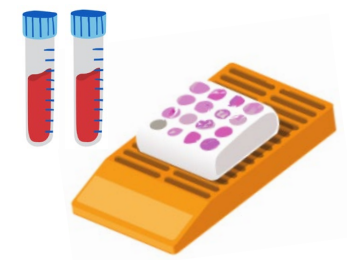
# Accelerating access to Comprehensive Genomic Profiling



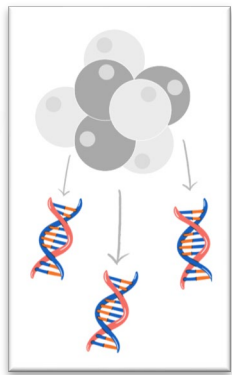
Referral from oncologist



Patient consent



Biospecimen retrieval

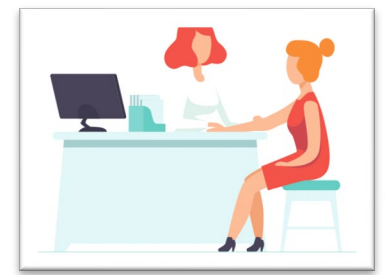


## Clinician

## Omico Central

## NATA Laboratory

Comprehensive genomic profiling  
TSO500  
Foundation CDx  
IHC-based screening



Discuss results with oncologist



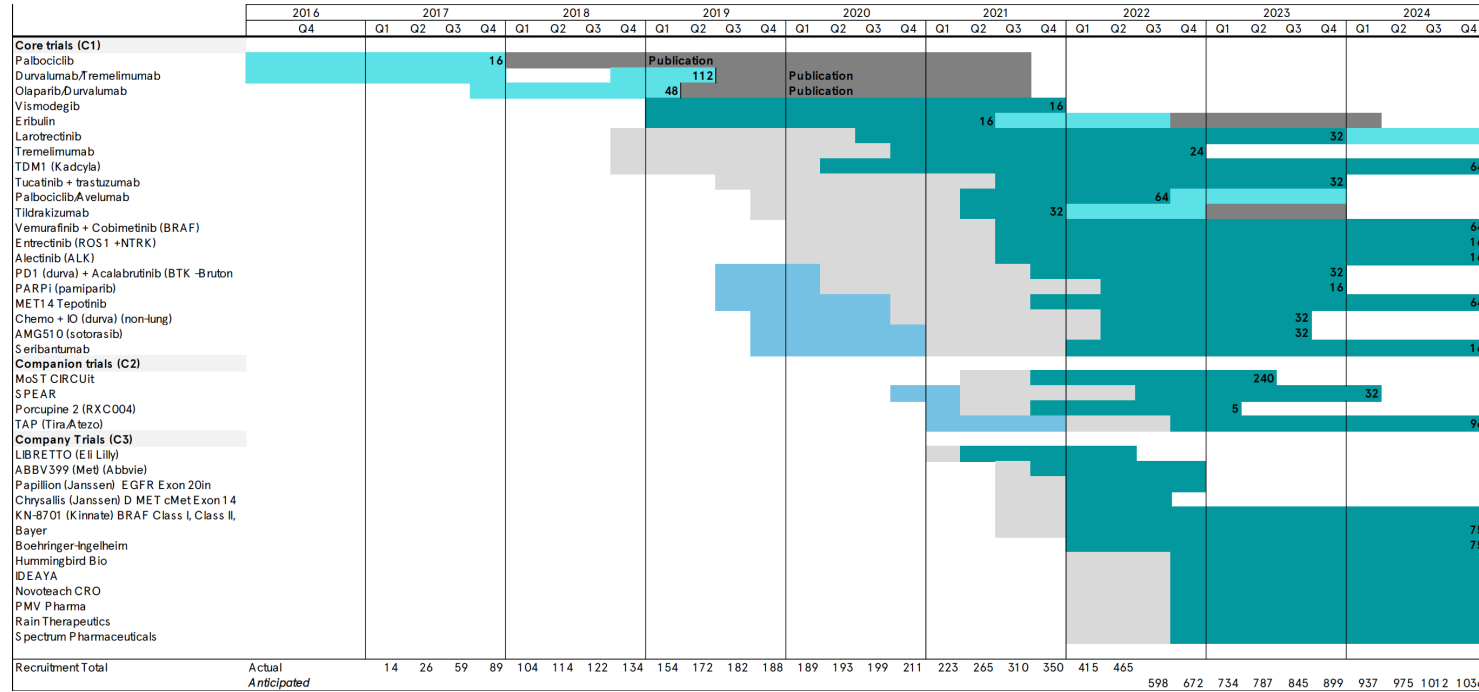
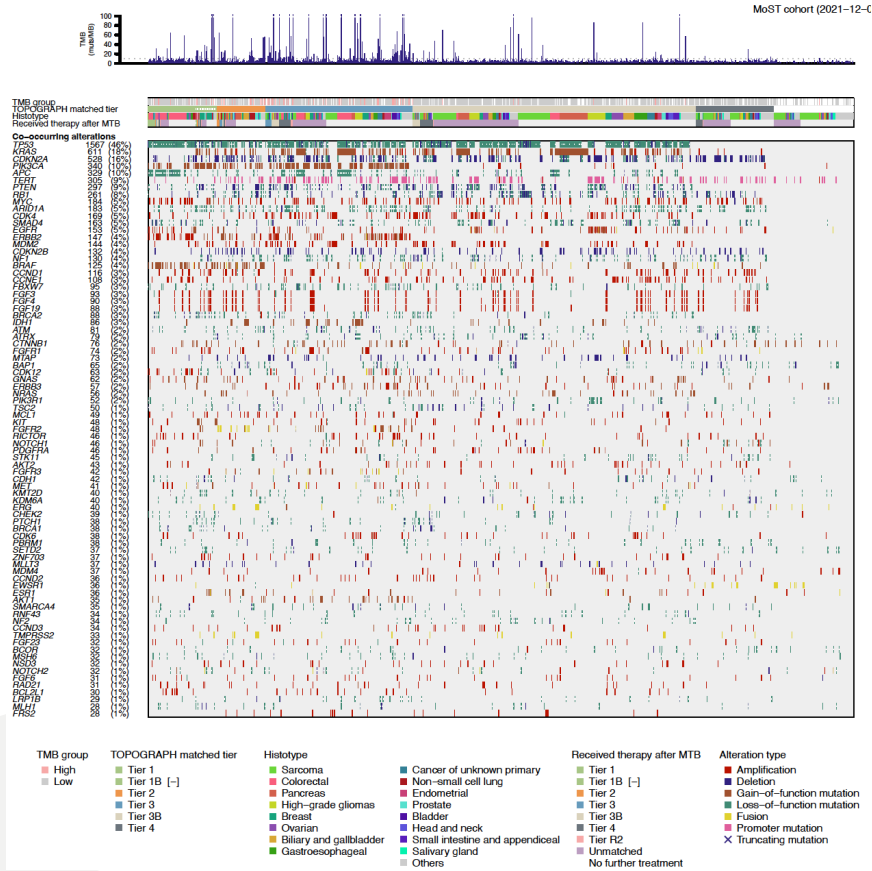
Molecular Tumour Board and report issued



Clinical trials matching



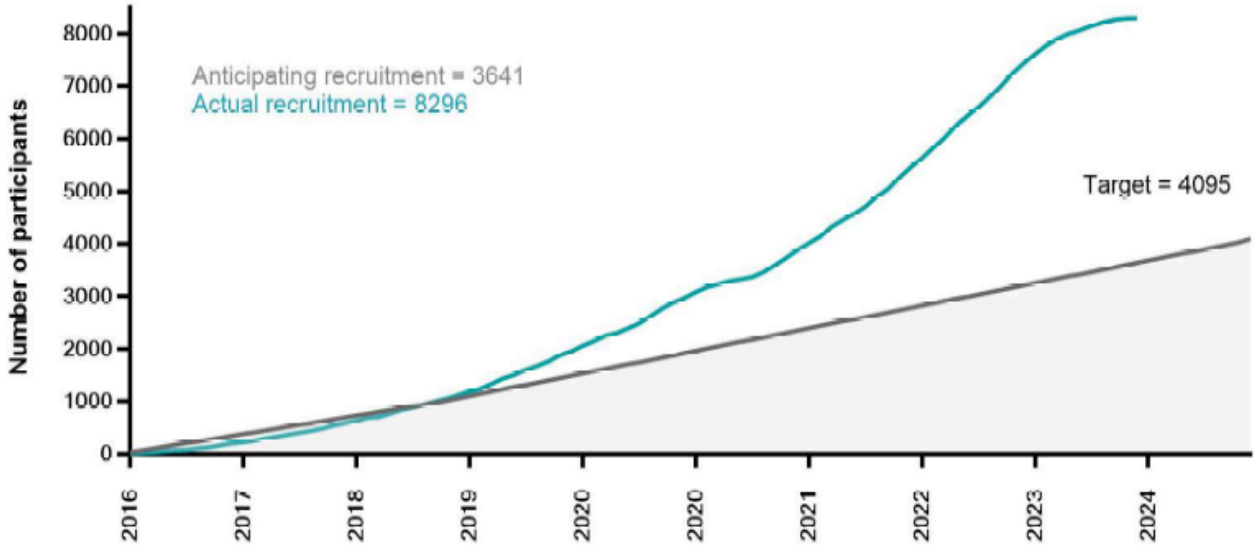
# Trial portfolio: >900 patients enrolled onto biomarker-matched trials



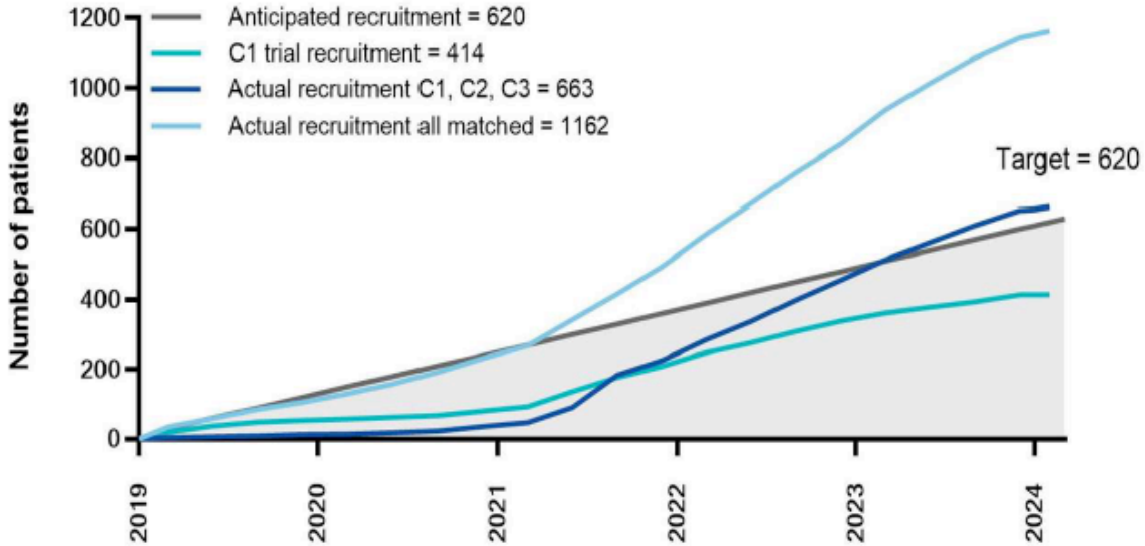
# Demand for precision medicine

>800 clinicians in Australia have referred onto MoST

70% rare



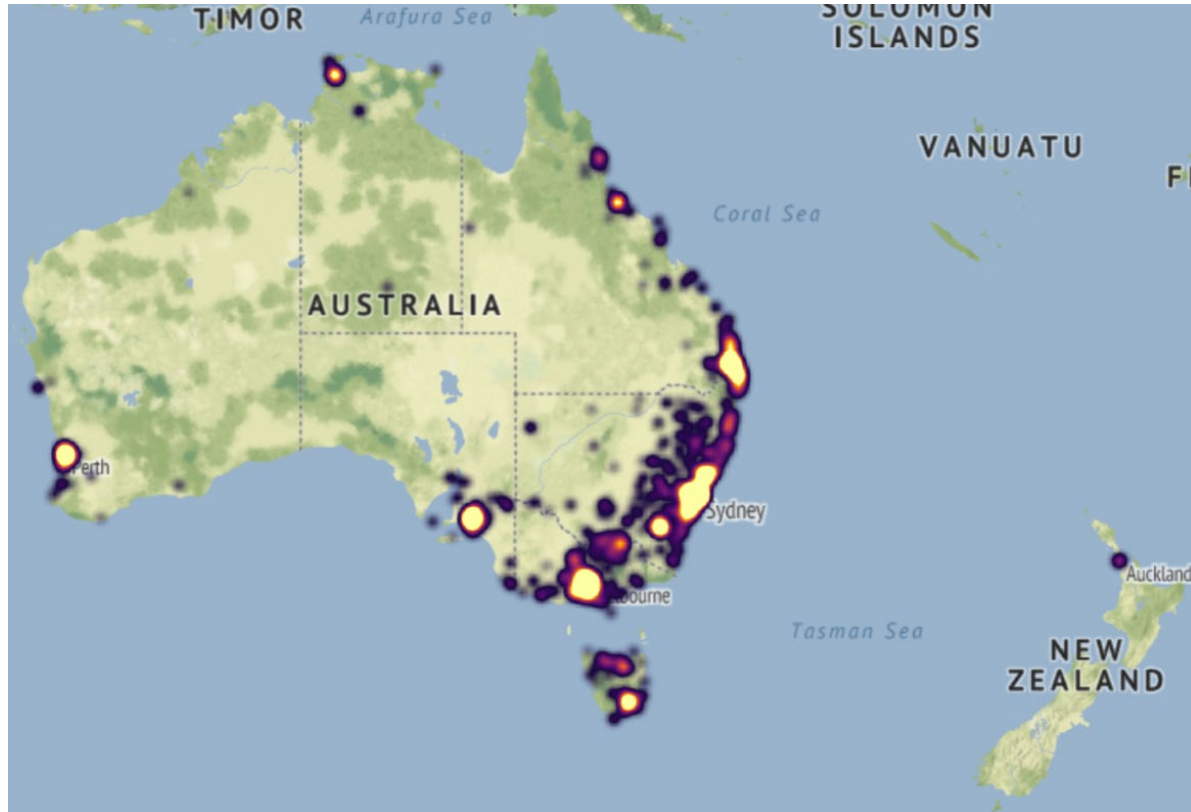
Patients with tumor profiling



Patients receiving matched therapy

<sup>1</sup> <https://hwd.health.gov.au/resources/publications/factsheet-mdcl-medical-oncology-2016.pdf>

# Omico's national population screening and clinical trial network



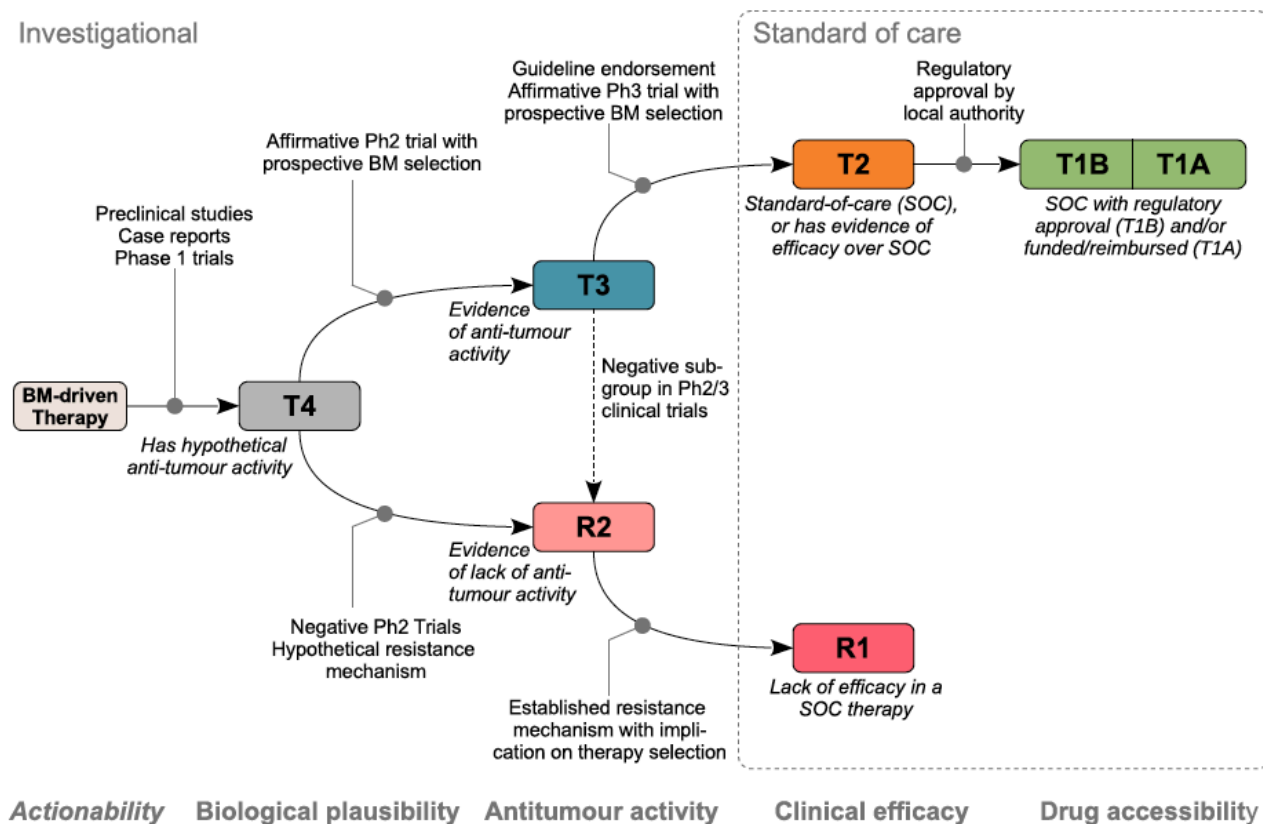
Omico Clinical Screening Recruitment  
**13,000 and increasing**



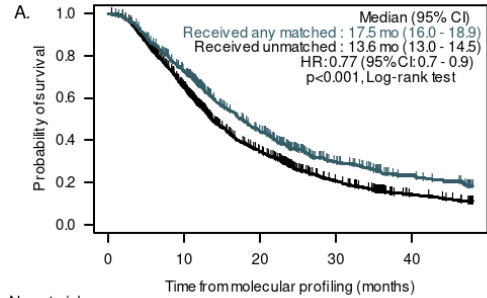
Omico National Clinical Trial Network  
**49 sites to date, and growing**

# Criteria-based curation of a therapy-focused compendium to support treatment recommendations in precision oncology

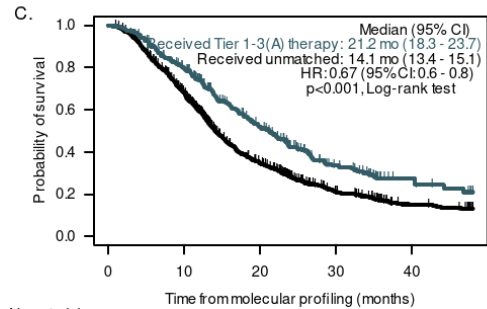
Frank P. Lin<sup>1,2,3</sup>, Subotheni Thavaneswaran<sup>2,3,4</sup>, John P. Grady<sup>3,4</sup>, Mandy Ballinger<sup>3,4</sup>, Maya Kansara<sup>3,4</sup>, Samantha R. Oakes<sup>3,4</sup>, Jayesh Desai<sup>5</sup>, Chee Khoon Lee<sup>2,6</sup>, John Simes<sup>2</sup> and David M. Thomas<sup>3,4</sup>



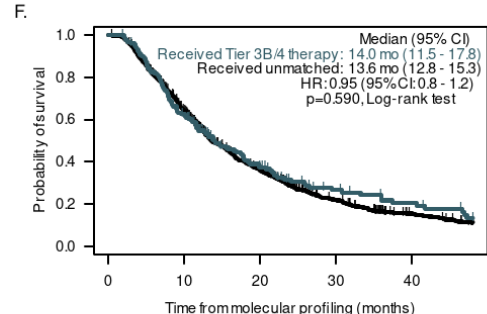
# Outcomes: improved survival (update to K7700)



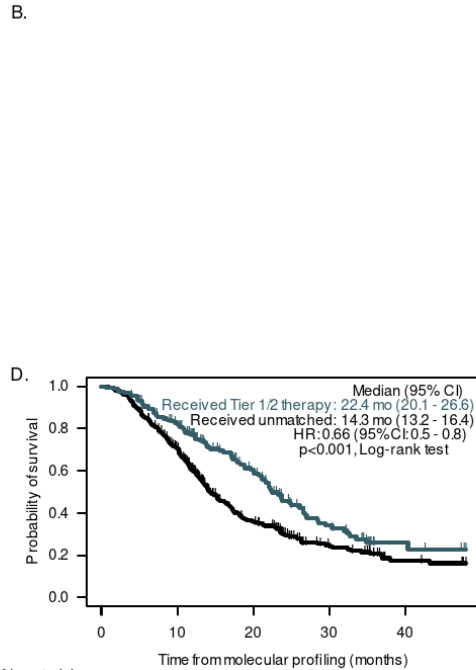
No. at risk	Time from molecular profiling (months)				
	0	10	20	30	40
Matched	722	498	240	108	53
Unmatched	1588	982	411	177	78



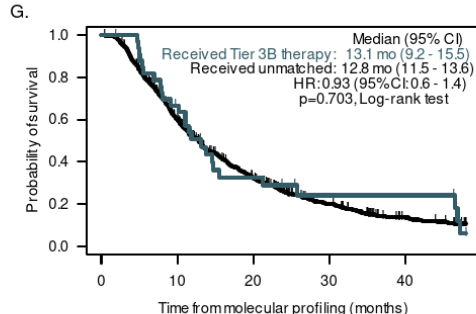
No. at risk	Time from molecular profiling (months)				
	0	10	20	30	40
Matched	283	217	115	53	32
Unmatched	946	607	233	94	40



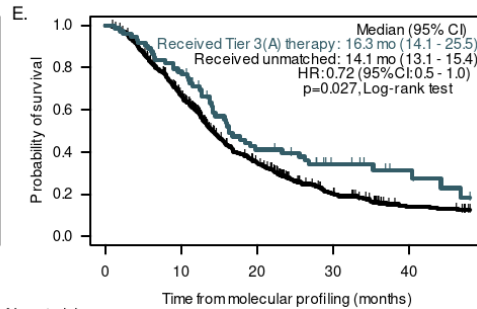
No. at risk	Time from molecular profiling (months)				
	0	10	20	30	40
Matched	163	101	50	25	17
Unmatched	918	560	256	115	57



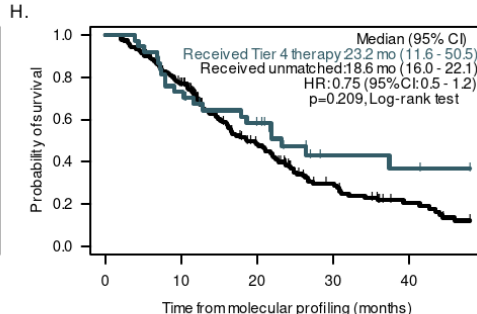
No. at risk	Time from molecular profiling (months)				
	0	10	20	30	40
Matched	160	124	71	29	18
Unmatched	377	252	93	39	16



No. at risk	Time from molecular profiling (months)				
	0	10	20	30	40
Matched	34	23	10	6	6
Unmatched	815	466	208	96	49



No. at risk	Time from molecular profiling (months)				
	0	10	20	30	40
Matched	73	58	27	21	12
Unmatched	619	392	159	64	27



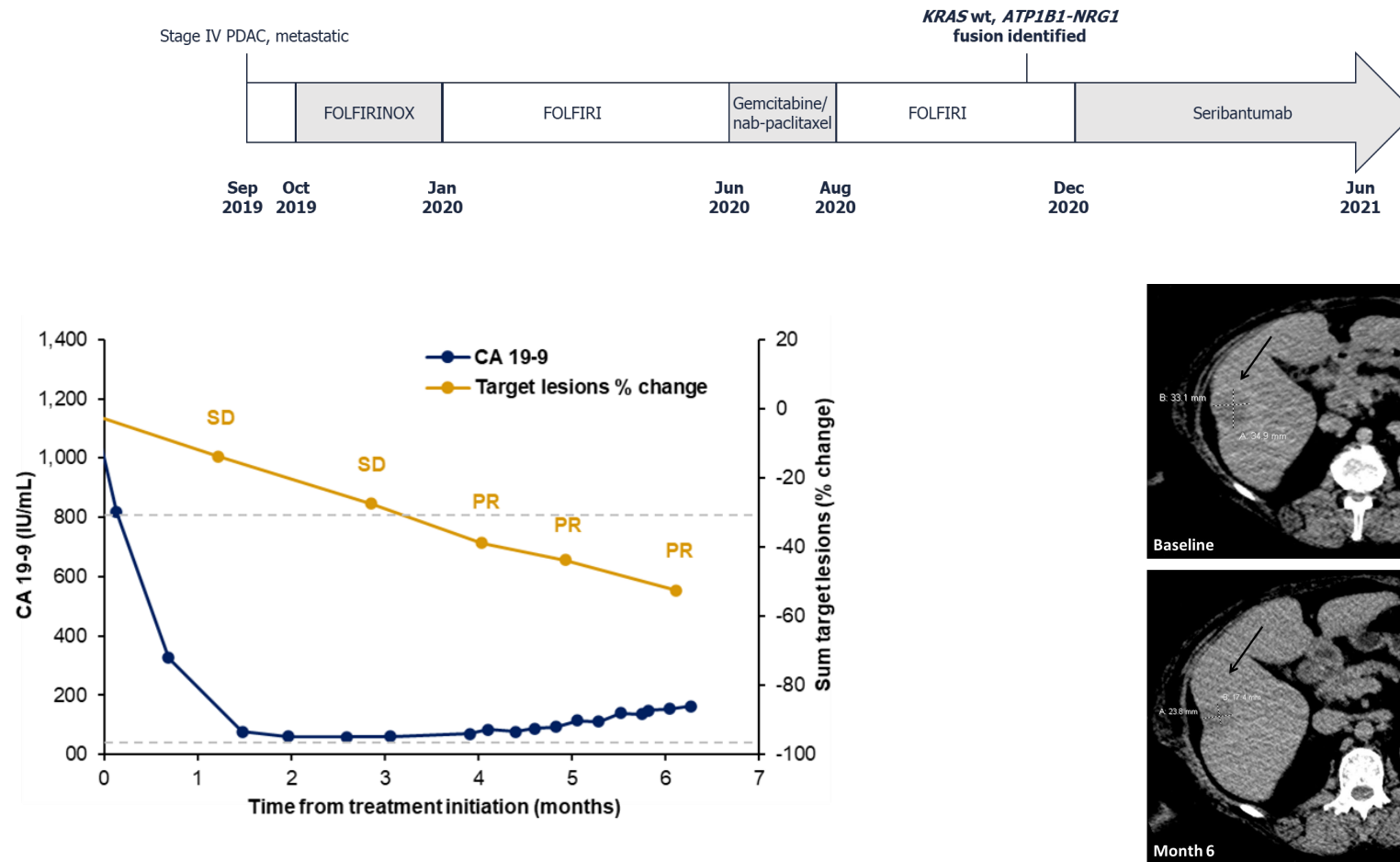
No. at risk	Time from molecular profiling (months)				
	0	10	20	30	40
Matched	38	28	20	11	7
Unmatched	194	146	70	34	16

Data from MoST\*

- **> 60%** of patients have had a matched treatment identified
- Of these, **37.5% have a drug target (T1-3)** which appears to result in a **doubling of survival**, provided the drug is accessible.



# Case studies: NRG1 fusion in pancreas cancer



# PrOSPeCT is a \$190m project that will establish Australia as a regional hub for cancer drug development and build the digital economy in health analytics

PrOSPeCT (Precision Oncology Screening Platform enabling Clinical Trials) is an Omico project that will...

1



Provide broad-based access to comprehensive genomic profiling for non-curative solid or haematologic cancer patient populations – **conducting large-scale genomic screening of more than 23,000 patients**

2



Maximise opportunities in clinical trials by identifying patients for specific biomarker-dependent clinical trials – **driving the efficiency of patient recruitment and decreasing trial costs**

3



Enhance biomarker-dependent drug development by enabling advanced molecular analyses – **creating a real-world dataset and analytics platform with national scale comprising more than 23,000 patients**



PrOSPeCT will be delivered alongside a public-private consortium of partners from across the Australian genomics and clinical trials ecosystem



**Major Sponsor\***



**Commercial Collaboration Partners**



**Delivery partners**



Note: \*Funding through Modern Manufacturing Initiative

# CaSP Monthly Dashboard (to end August 2024)

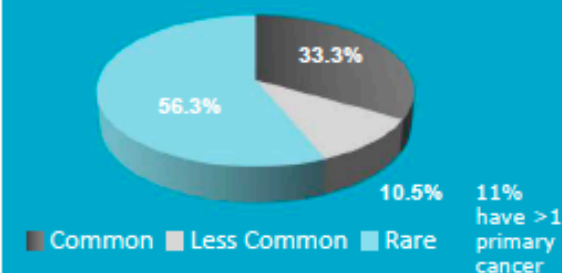
## Patient Recruitment & Screening

8,399	Patients referred (CaSP and Zero)
7,059	Patients consented for screening (CaSP and Zero)
4,352	MOB (Molecular Oncology Board) reports issued
3,217 (73.9%)	Patients have had a matched therapy recommendation (Based on those with MOB report issued to date)
849	Clinicians have referred patients to CaSP

## Patient Demographics & Cancer Classification

### Consented pts / state (CaSP only)

State	#	Regional & rural pts
ACT	99	36%
NSW	2,337	
NT	65	
QLD	1,023	
SA	414	
TAS	331	
VIC	1,773	
WA	62	



Gender	Female	52%
Age at Diagnosis	Average	59y
	Range	9-94y

## Time from patient consent to MOB report issue

	Planned	Actual (median)
Standard cases:	8-10 weeks	8.1 weeks
Urgent cases expedited:	5-6 weeks	5.9 weeks



## Omico National Clinical Trial Network (ONCTN)

Total sites

61



State	Total	Regional
ACT	2	
NSW	26	11
NT	1	1
QLD	7	2
SA	6	1
TAS	2	
VIC	14	5
WA	3	

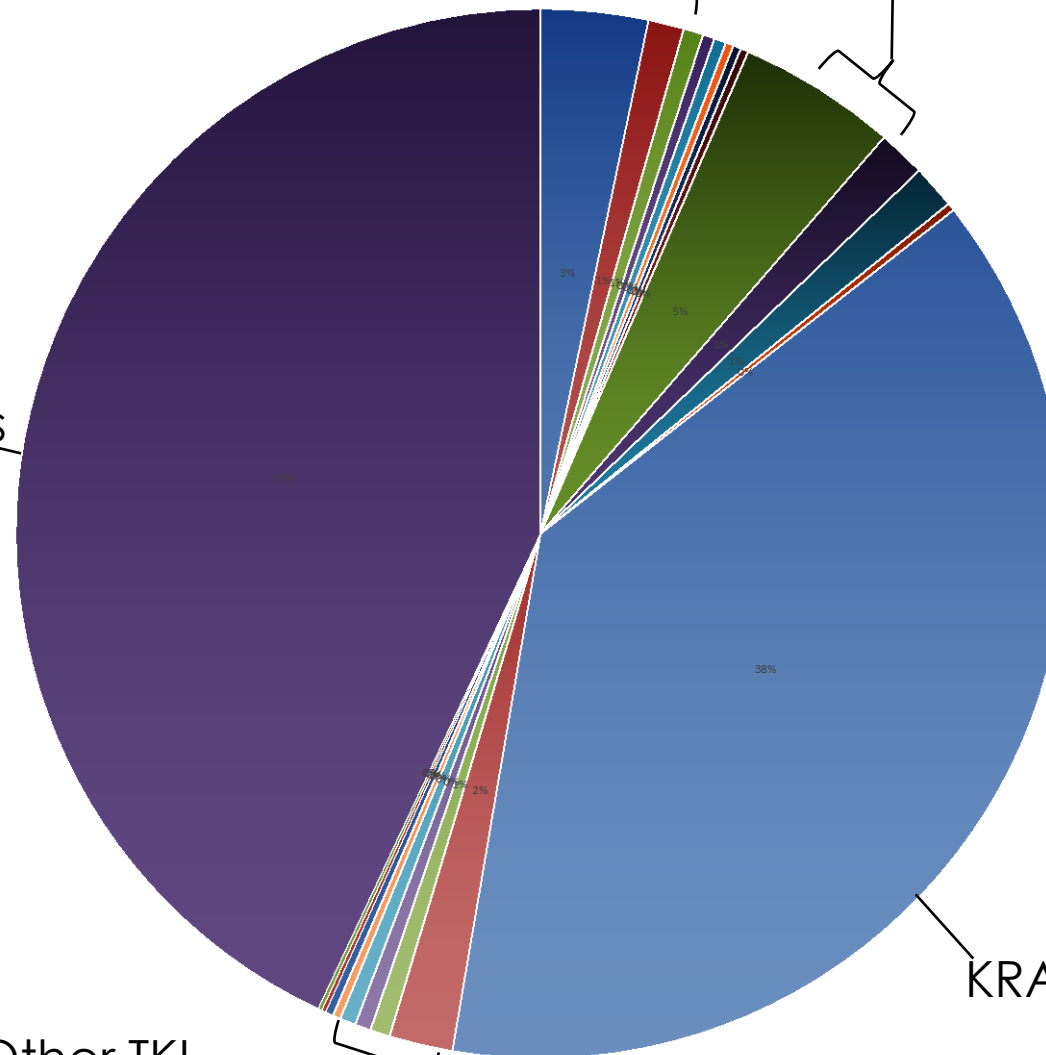
# Pancreatic cancer

Patients = 466/855 (57%)  
Drug targets = 818

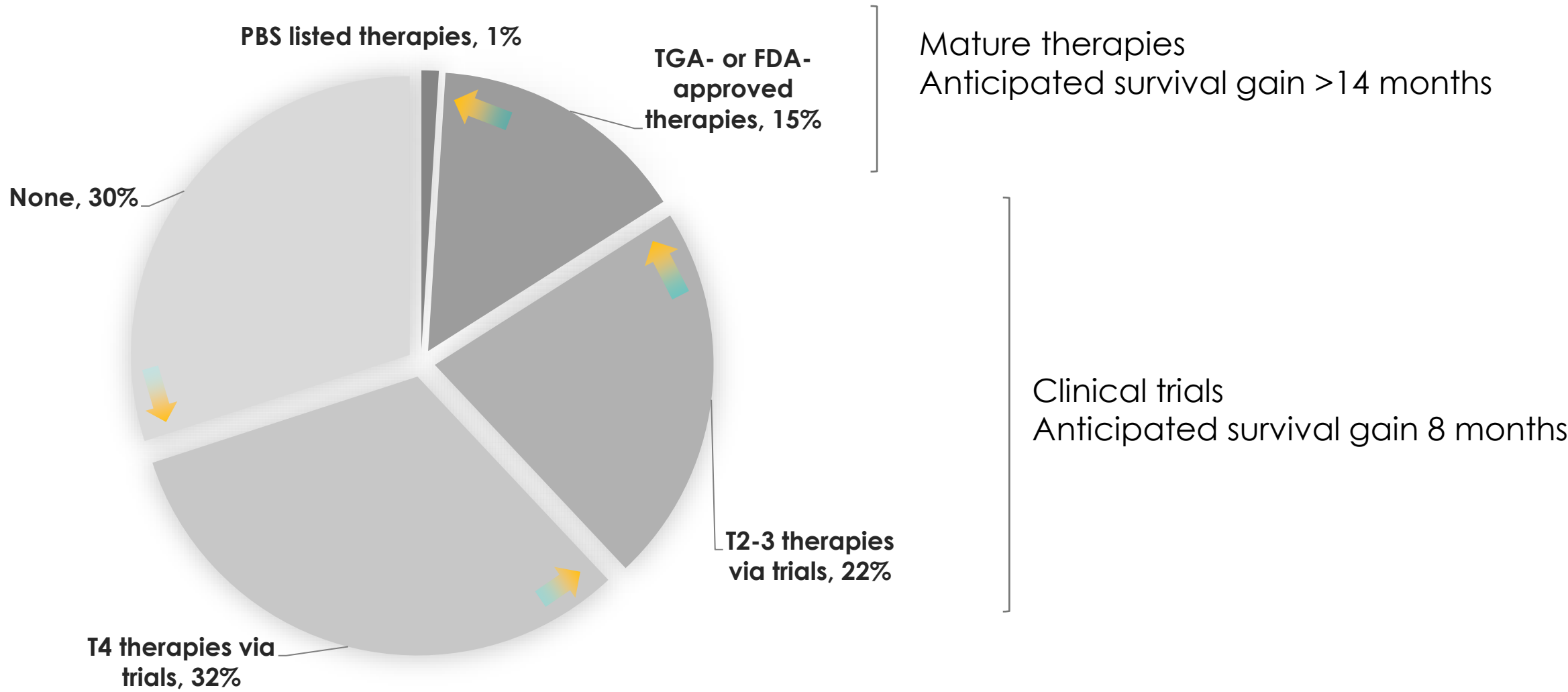
MTAP-directed therapies

Immunotherapy

PARPi



KRAS G12C/D



# Health systems are challenged to deliver equity, certainty and sustainability in the face of rapid scientific advances

## Public policy papers



## Timely, equitable and affordable access

- Long timelines between TGA and PBS approvals exacerbated by accelerated drug development
- High unmet needs with growing and united community calls for social justice and structural equity

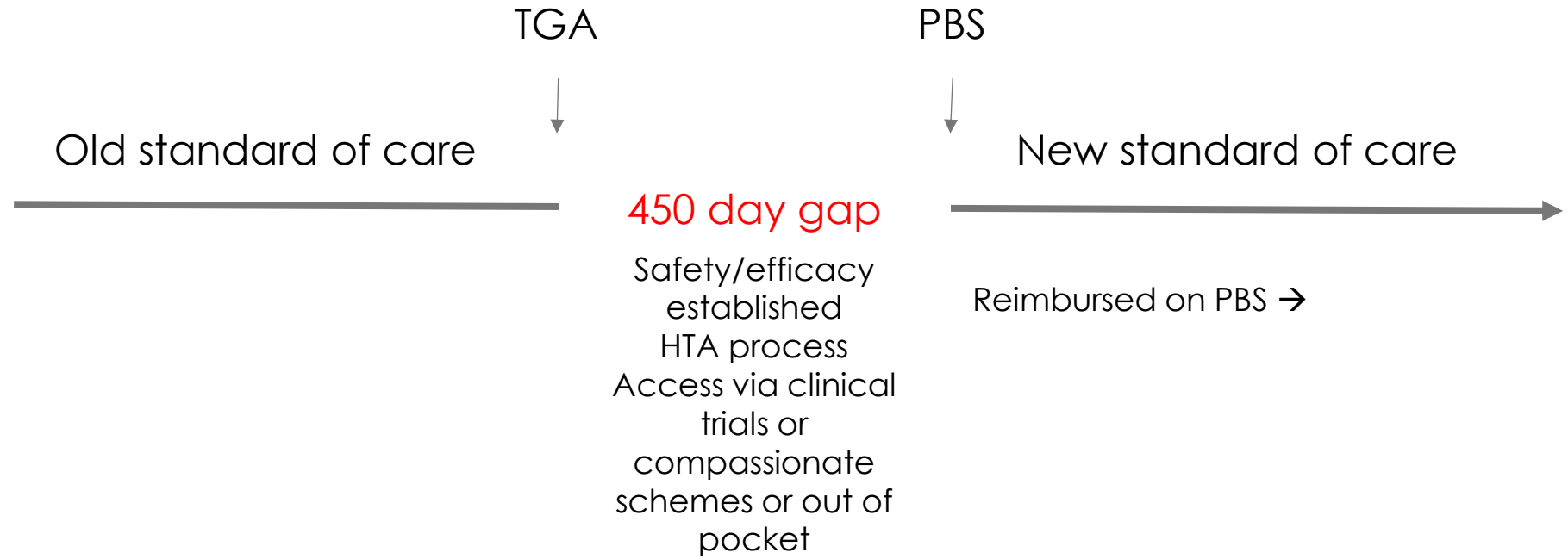
## Strengthening HTA by reducing uncertainty

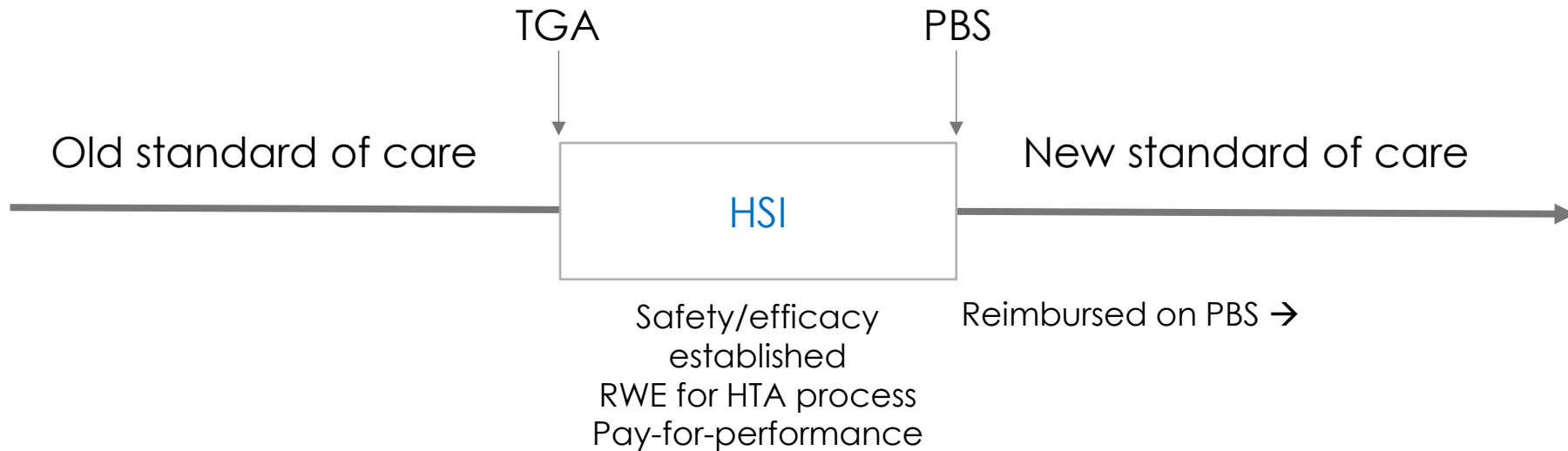
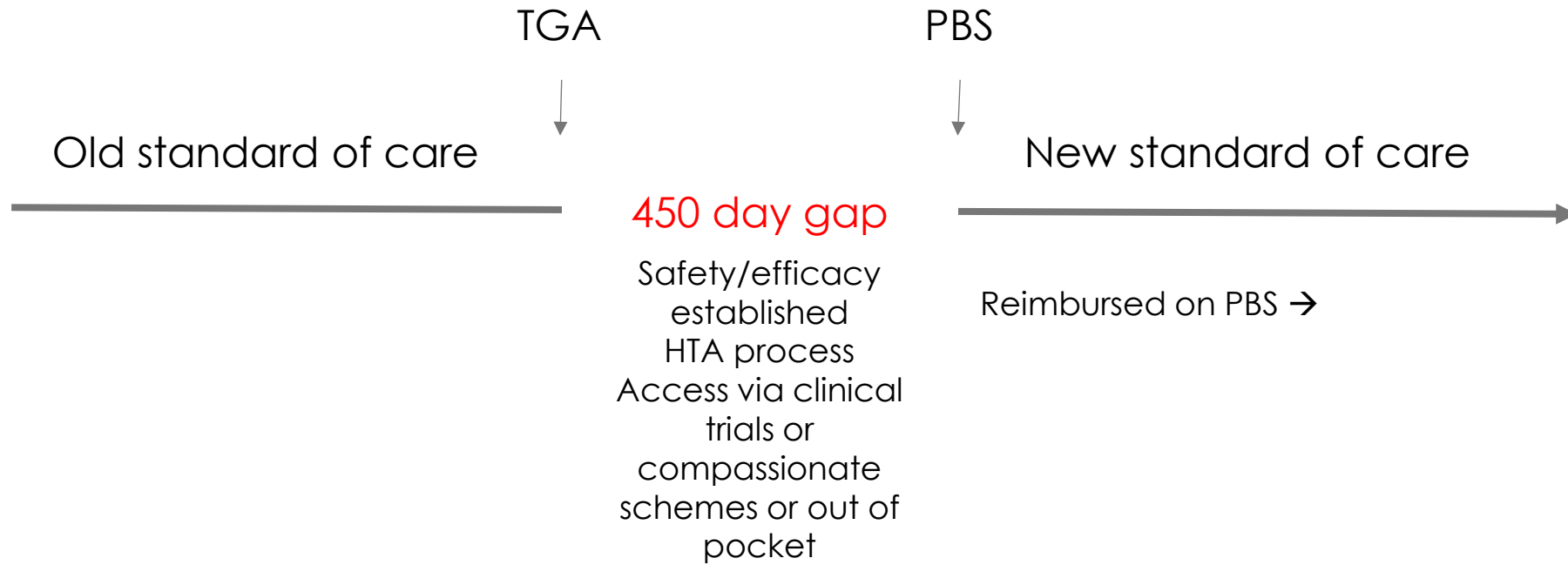
- Establish robust clinical safety, efficacy and cost-effectiveness data beyond traditional RCT paradigm
- Manage administrative and cost burden associated with increasing complexity of health innovation

## Economic growth from R&D investment

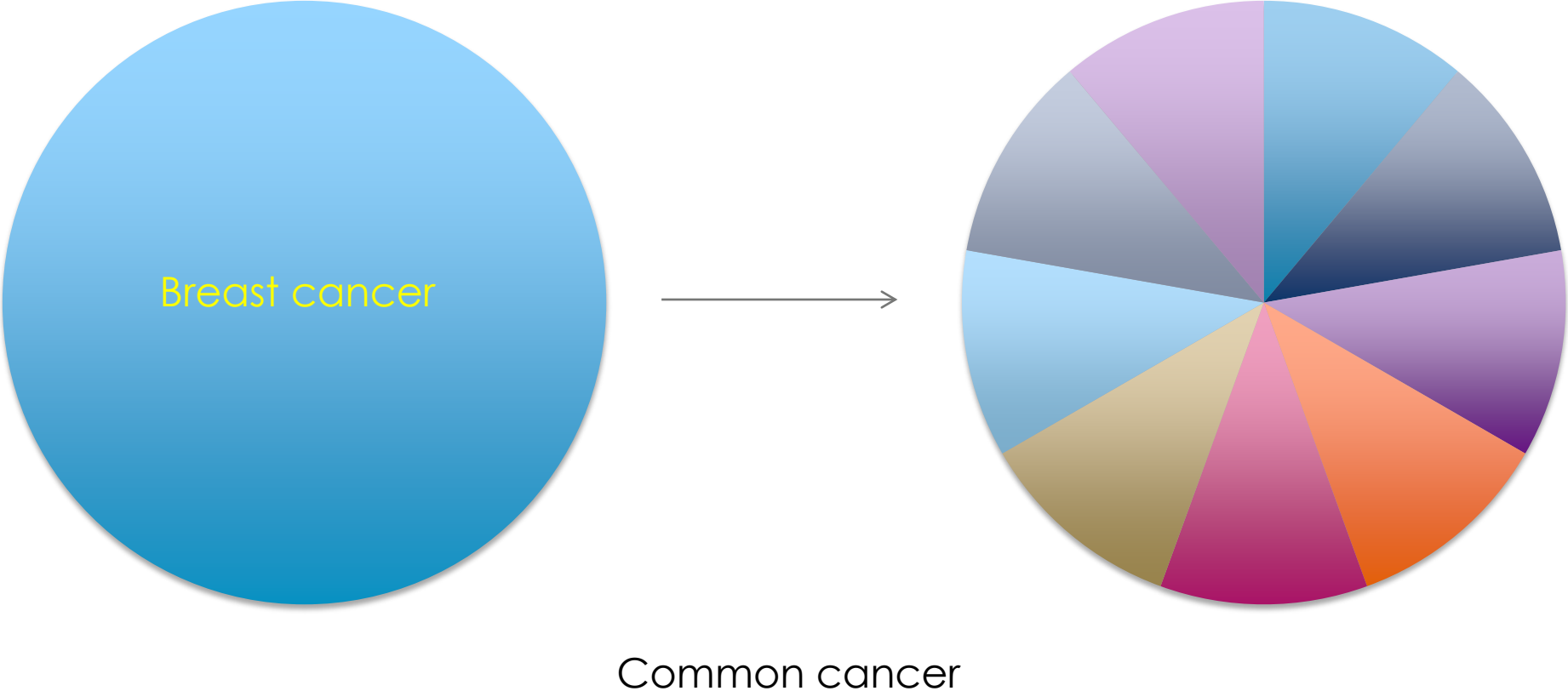
- Drive economy through investment in manufacturing, clinical trials and commercialization of research
- Maintain Australia as a first-launch country for health innovation

Sources: Senate Inquiry into equitable access to diagnosis and treatment for individuals with rare and less common cancers. (2023); Australian Cancer Plan, Cancer Australia (2023); Rare Cancers, Rare Diseases and Unmet need Clinical Trials initiative (2024), HTA Policy and Methods Review (2023), Intergenerational Report (2023); Medical Science Co-investment Plan (2024); MTP Connect Australia's Clinical Trial Sector (2024).

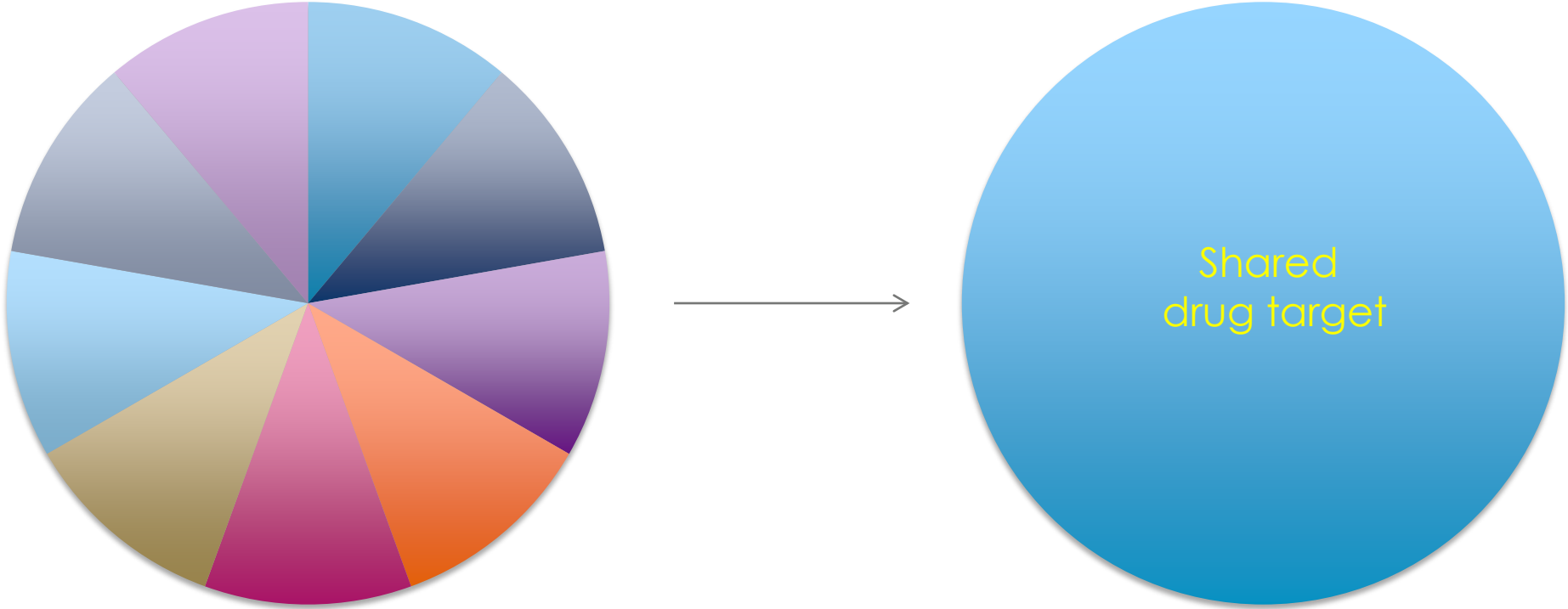




# Genomics is reshaping the cancer landscape

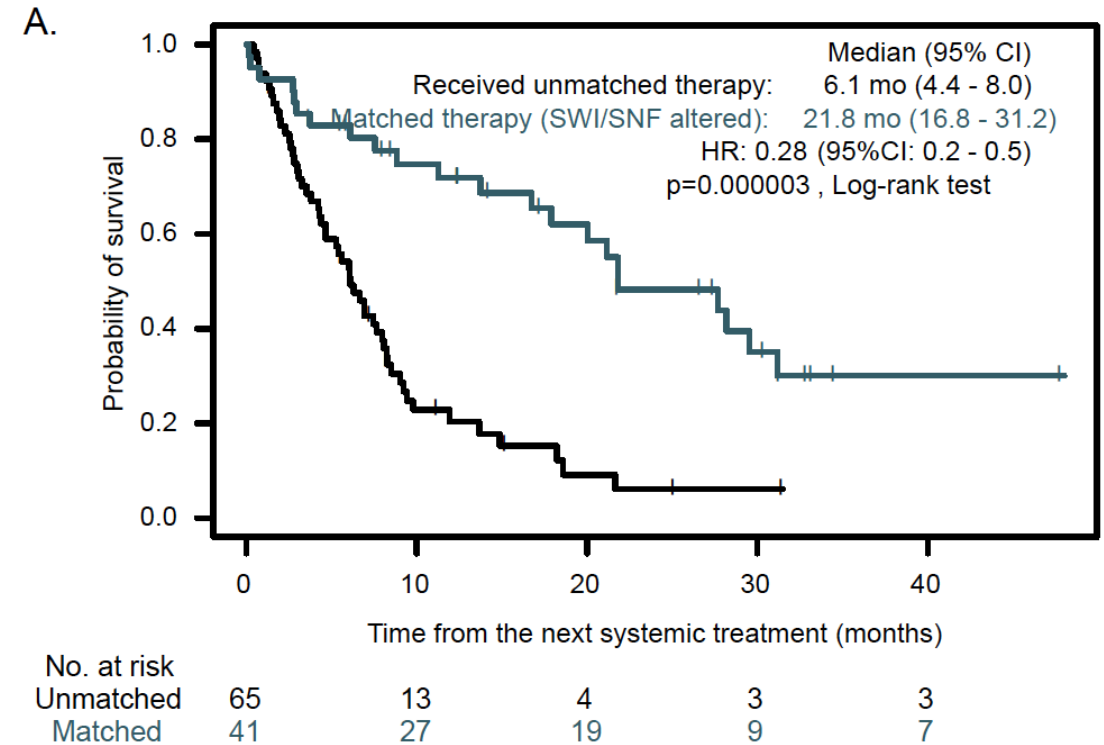
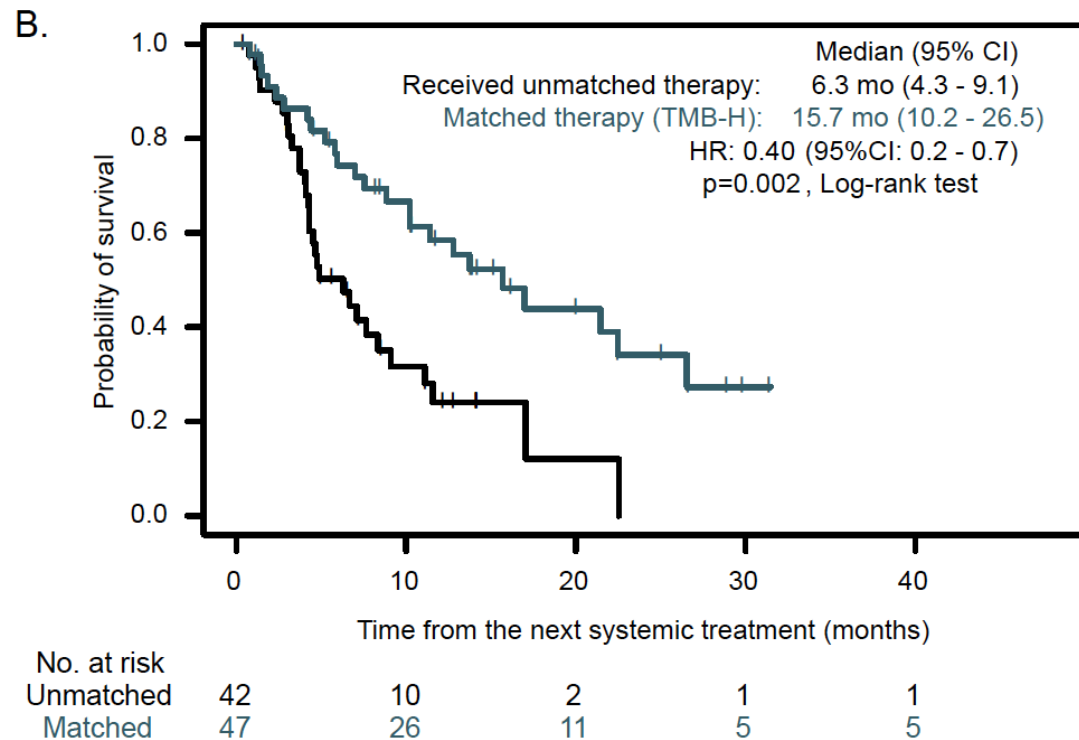


# Genomics is reshaping the cancer landscape

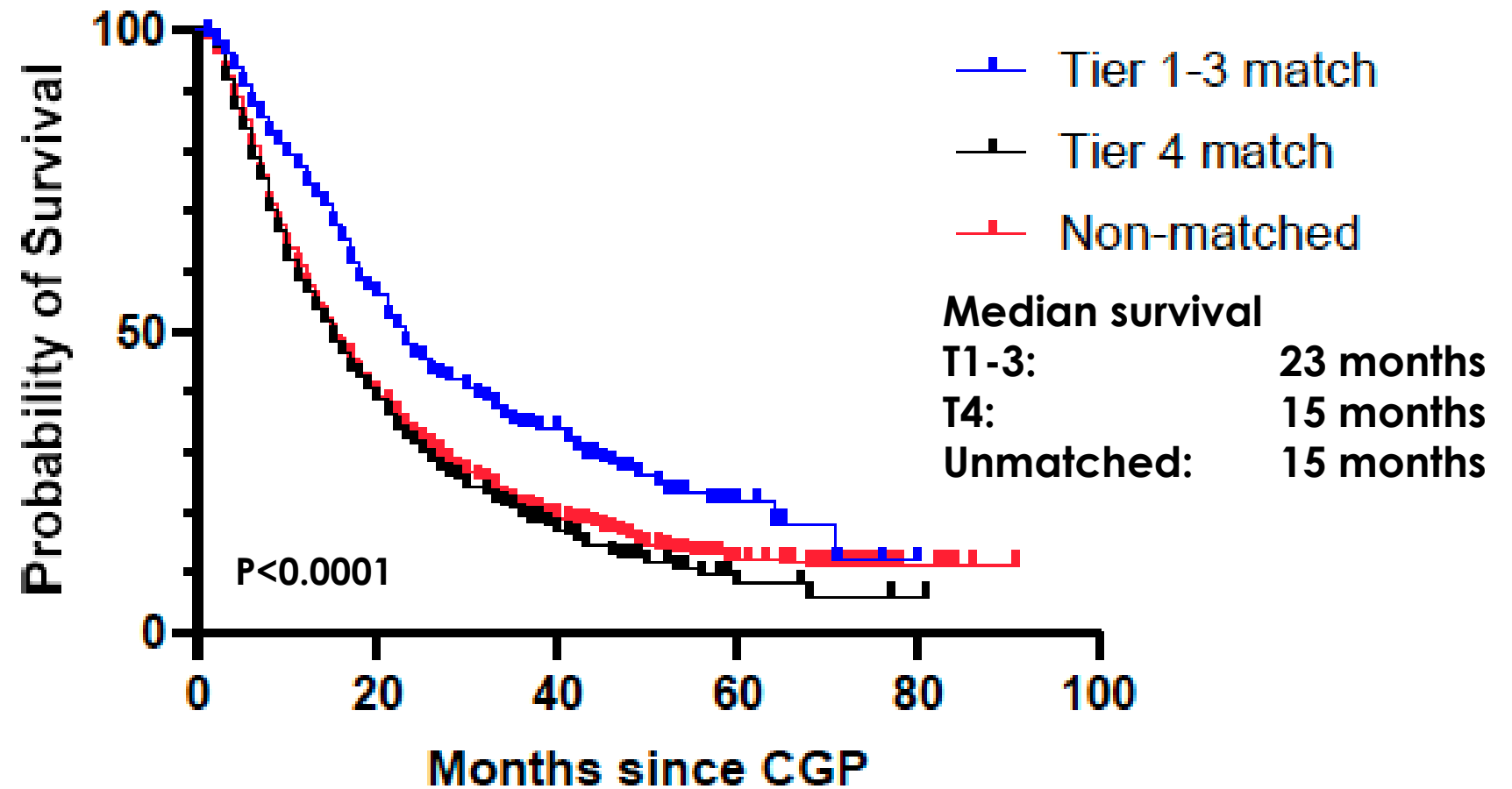


Rare cancers

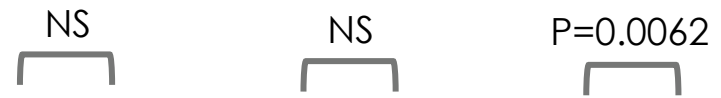
# Pan-cancer biomarkers: a strategy for rare cancers



# Outcomes for rare cancers in MoST

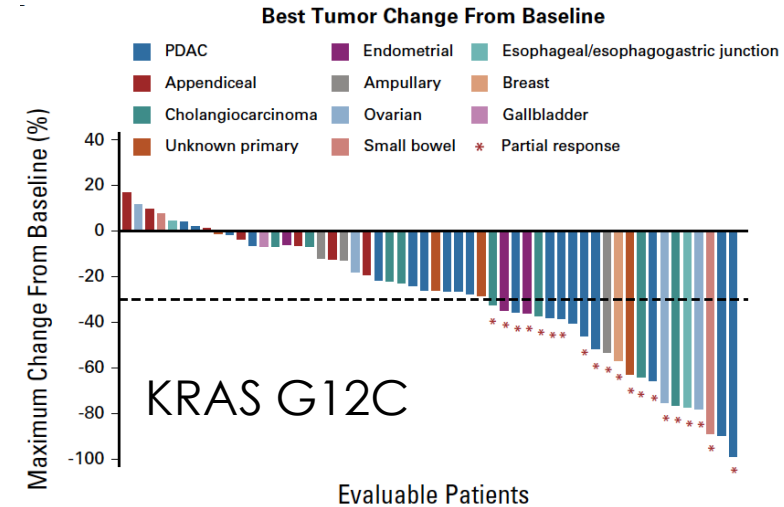
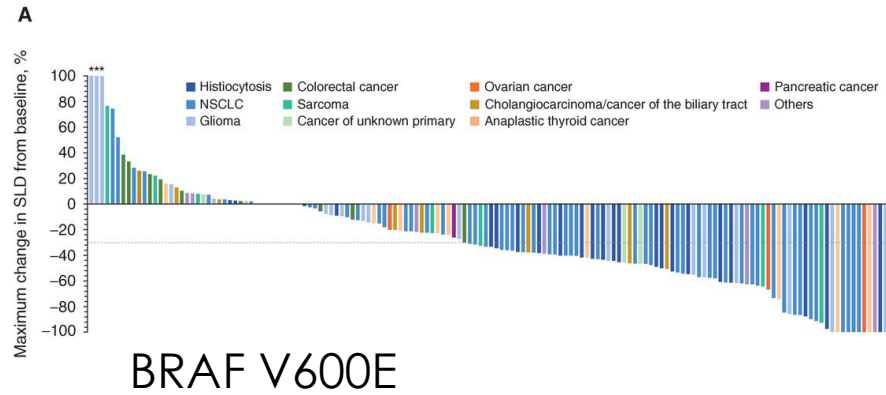
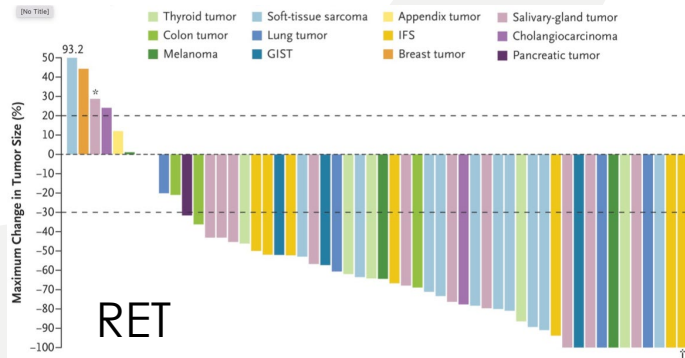
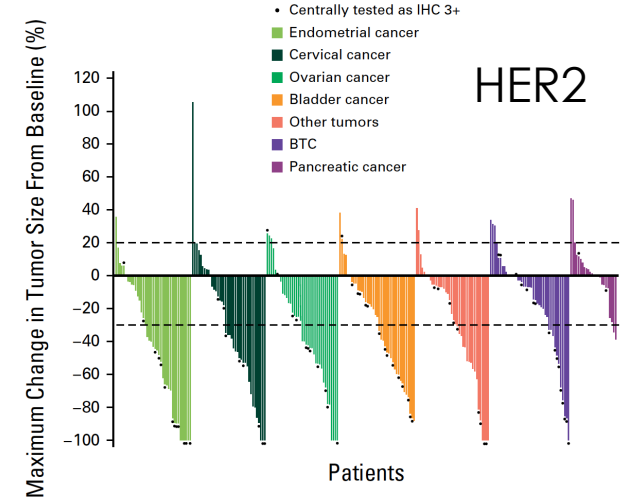
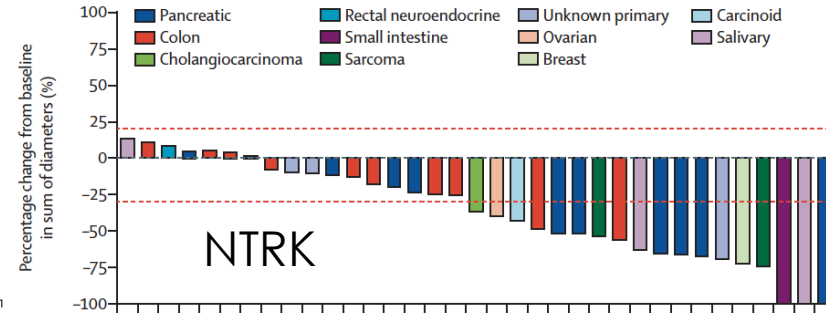
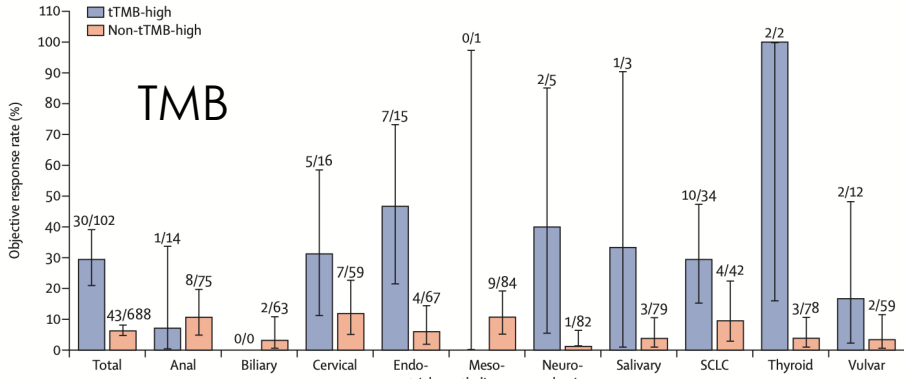


# Health-related Quality of Life



# Health resource utilisation

# Cancer agnostic therapies

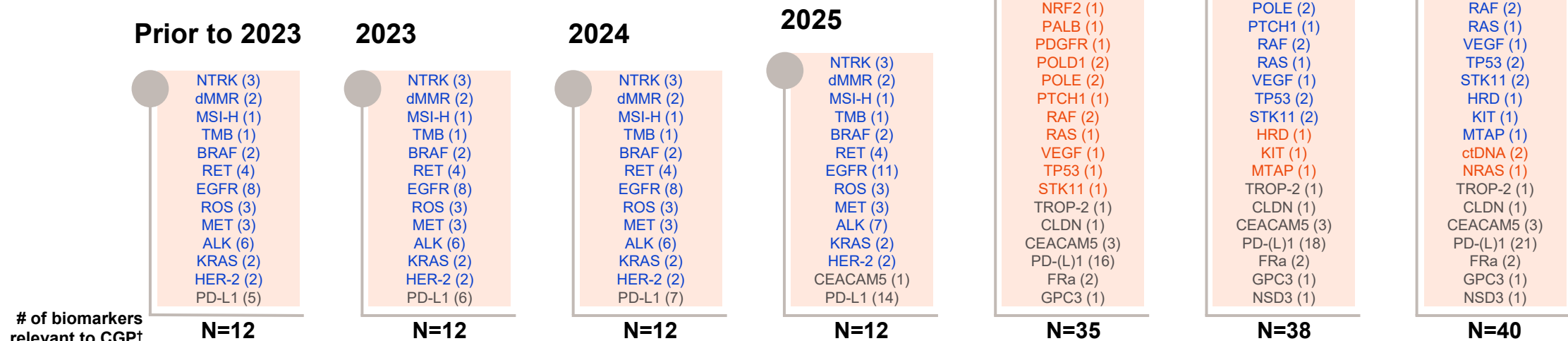


# Biomarker-dependent drugs are becoming increasingly important in oncology

Biomarkers likely to have approved targeted therapies with tumour agnostic indications\*

Analysis includes Phase 1/2, Phase 2 & Phase 3 trials initiated before April 1, 2023

- = Established biomarker suitable for CGP
- = New biomarker suitable for CGP
- = Biomarker without immediate relevance for CGP (other detection methods more appropriate†)



\*Multiple secondary sources used to cross validate information, including Trialstrove, clinicaltrials.gov, EudraCT, ChiCTR; FDA approval timeline estimation based on Phase 3 primary completion date + 8 months review; analysis based on current Phase 1/2, Phase 2 and Phase 3 trials with inclusion criteria requiring patient selection based on alterations to specific biomarkers; assumption made that all ongoing trials will lead to approval; †Biomarkers without immediate relevance for CGP are primarily those for which detection of protein expression via IHC or other immunology approaches is more appropriate (e.g., PD-L1). CGP: comprehensive genomic profiling; ctDNA: circulating tumour DNA; dMMR: DNA mismatch repair deficiency; HRD: homologous recombination deficiency; IHC: immunohistochemistry; MSI-H: microsatellite instability-high; NSCLC: non-small cell lung cancer; TMB: tumour mutational burden.

Currently, 7 drug targets which are FDA and TGA-approved for tumour-agnostic indications

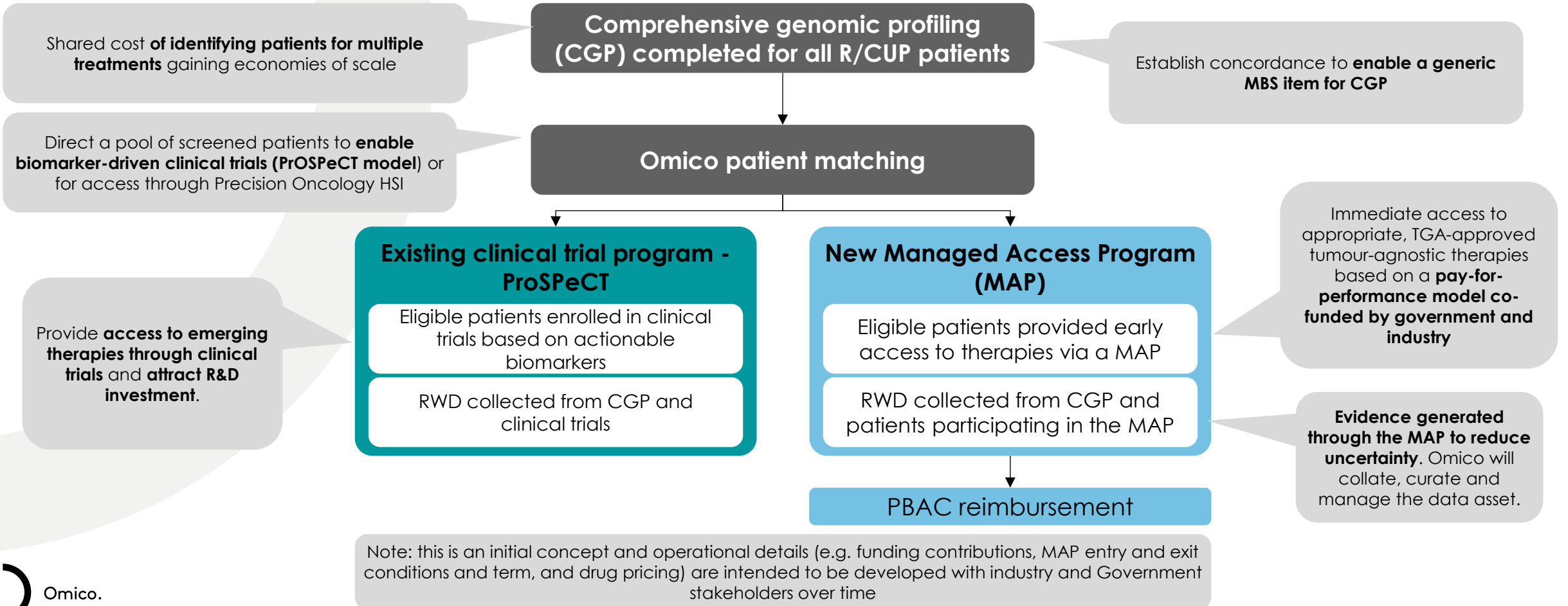
### TGA-approved pan-cancer therapies and their targets

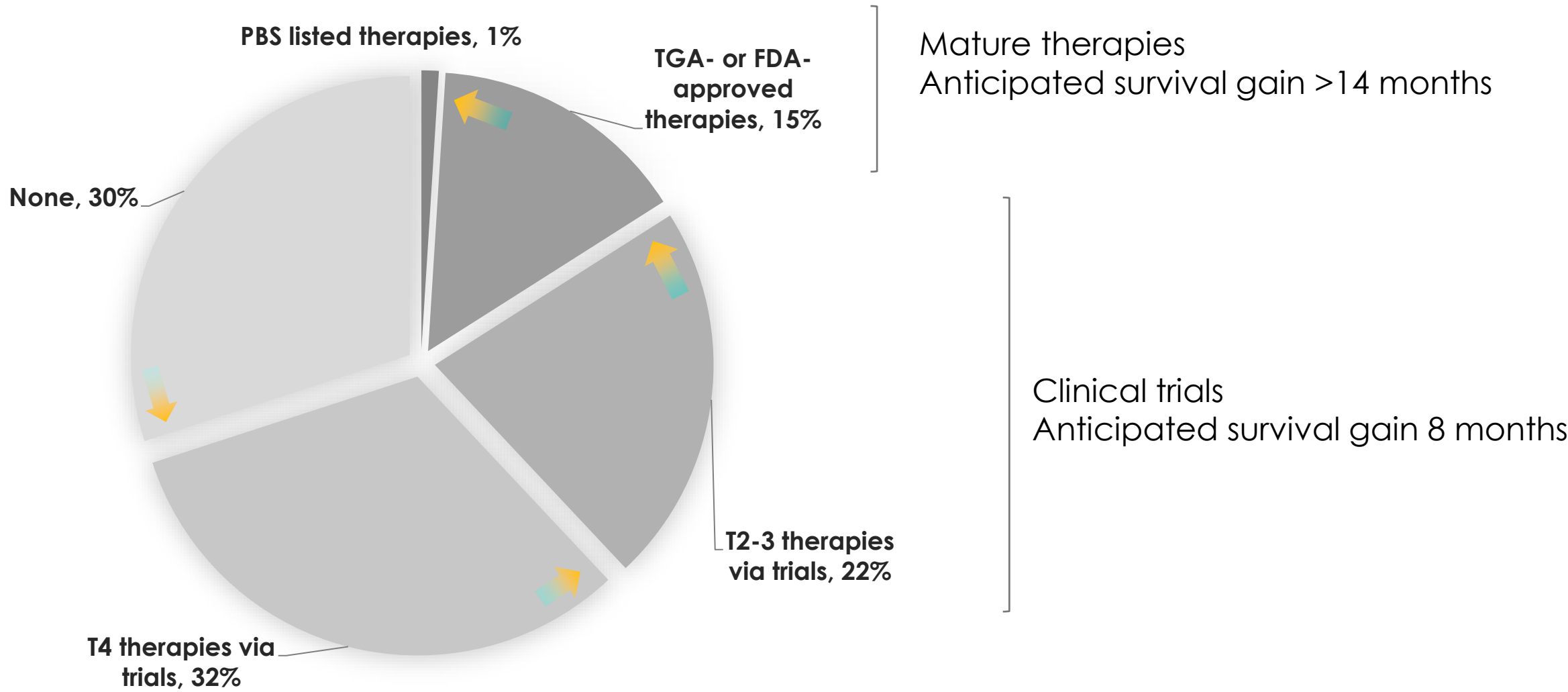
Biomarker / target	Frequency of target*	ORR (%)	Approved drug(s)	Year of FDA approval	Current Australian subsidy
dMMR	1.03%	40%	Pembrolizumab	2017	No submission to PBAC for tumour agnostic indication
			Dostarlimab	2022	As above
MSI-High	1.12%	40%	Pembrolizumab	2017	As above
TMB-high	7.91%	33%	Pembrolizumab	2020	As above
BRAF <sup>V600E</sup>	1.62%	41%	Dabrafenib + Trametinib	2022	2019 (PBS) No submission to PBAC for tumour agnostic indication
RET fusions	0.08%	44%	Selpercatinib	2022	No
NTRK fusions	0.23%	75%	Larotrectinib	2018	2022 (PBS) for paediatric NTRK tumours
			Entrectinib	2019	No submission to PBAC for tumour agnostic indication
HER2 amp.	2.71%	61%	Deruxtecan	2024	Being submitted to TGA

# The Precision Oncology – HSI: a unified, contained ecosystem for evaluating complex health innovations as routine care

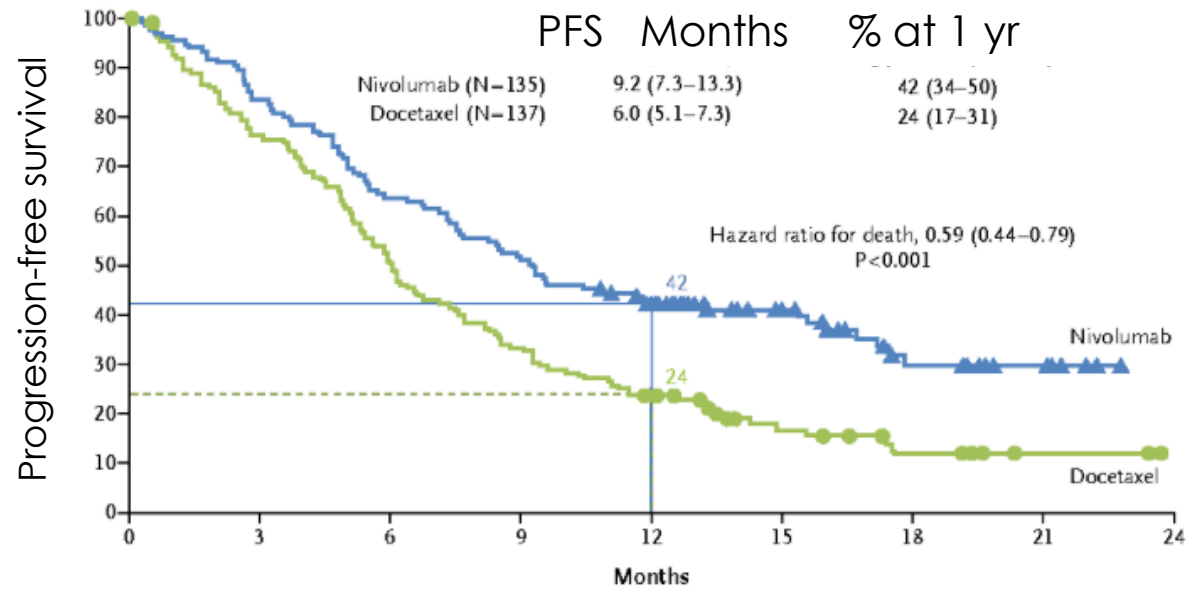
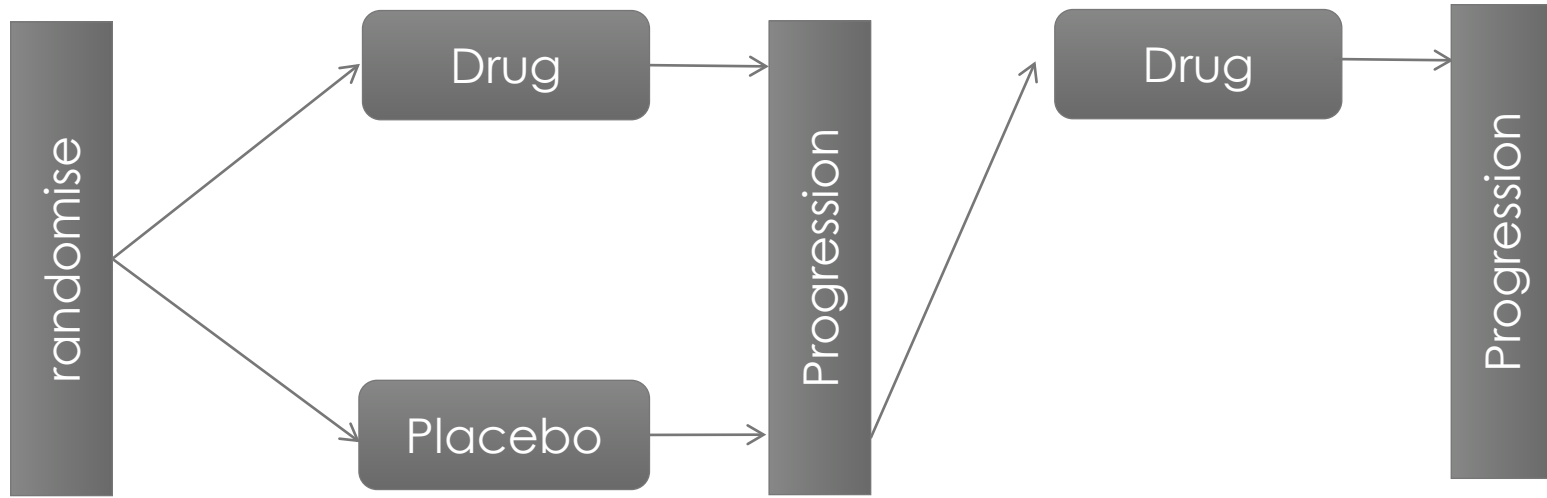
## Precision Oncology Health System Incubator ecosystem

**Target patient population: rare cancer / cancer of unknown primary (R/CUP) patients (c.5,000 new patients screened / year)**



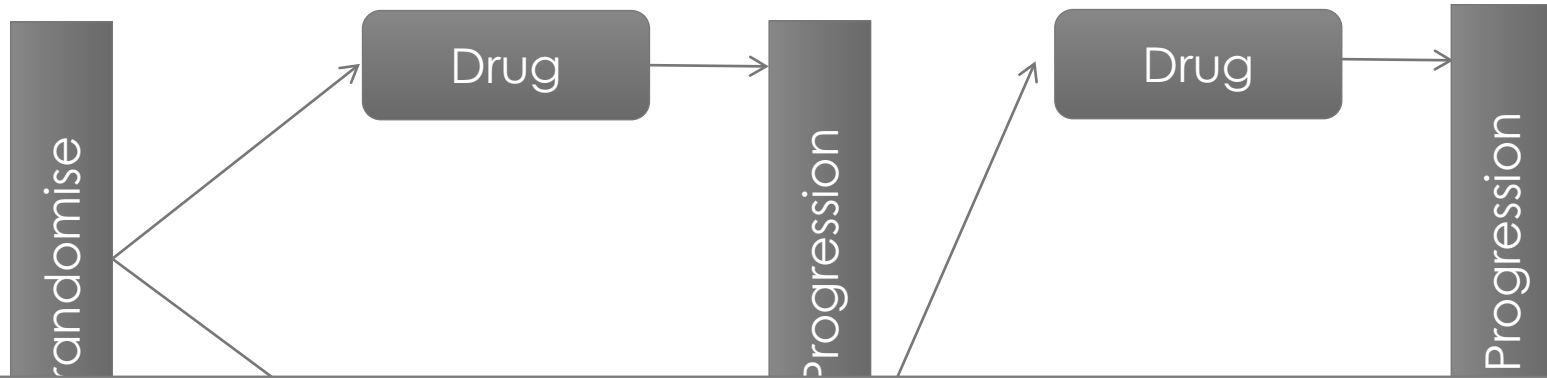


NSCLC

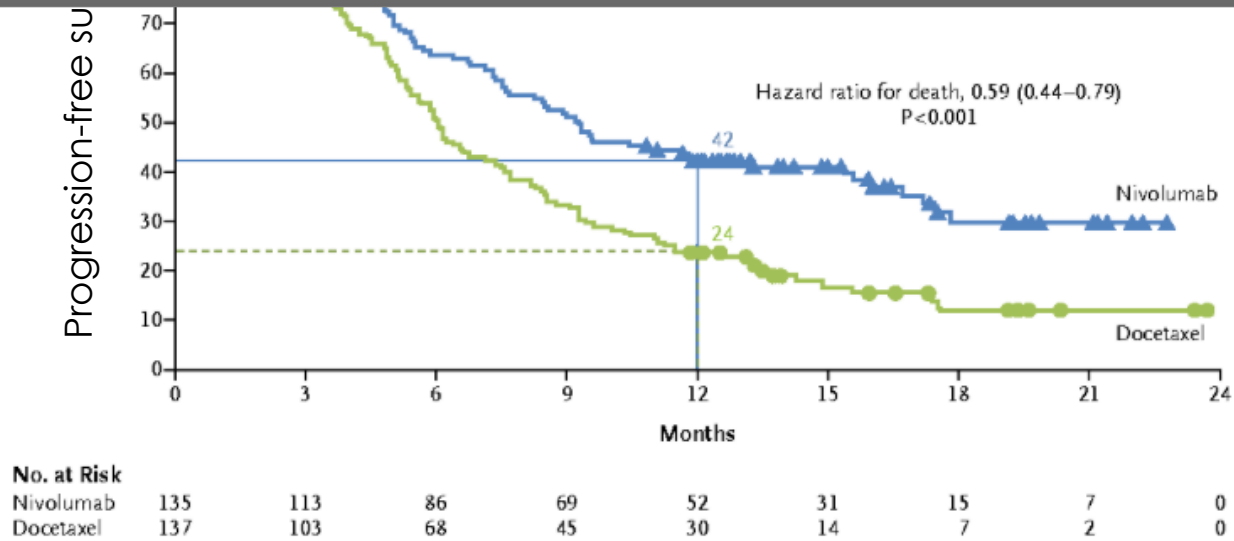


No. at Risk		0	3	6	9	12	15	18	21	24
Nivolumab	135	113	86	69	52	31	15	7	0	0
Docetaxel	137	103	68	45	30	14	7	2	0	0

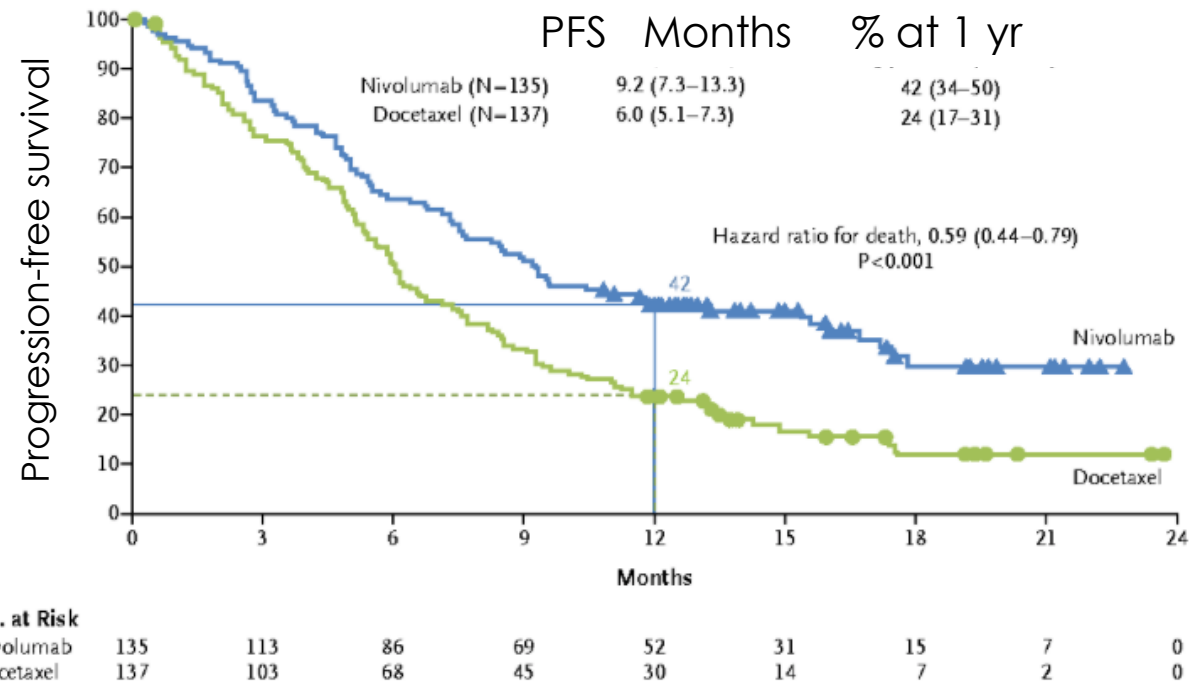
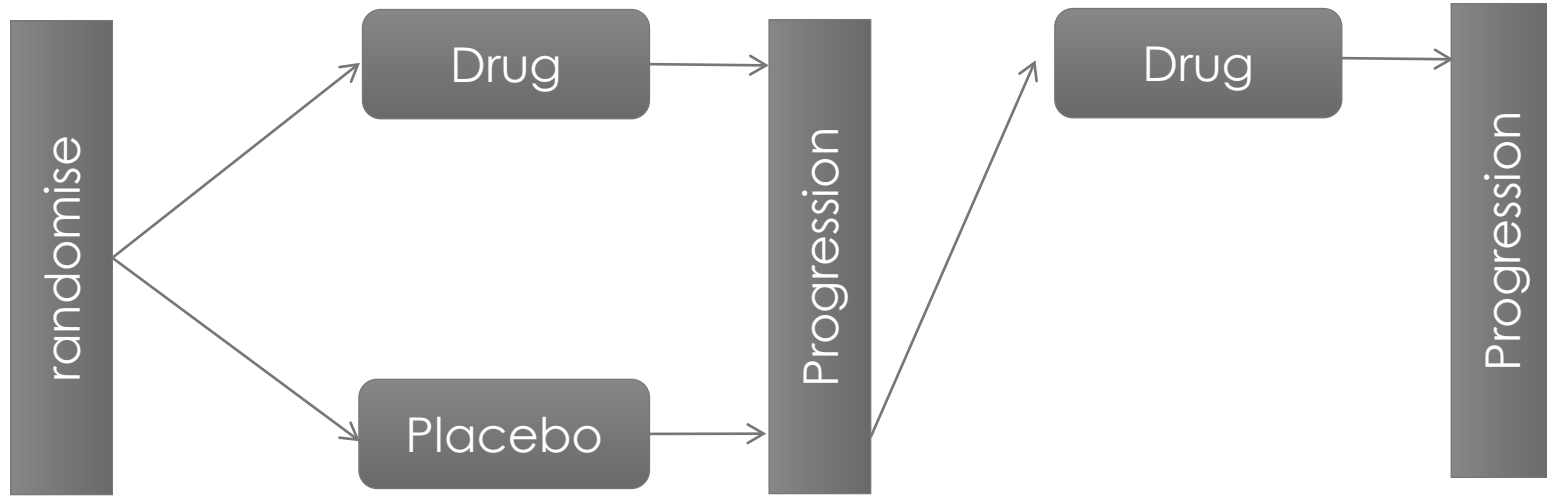
NSCLC



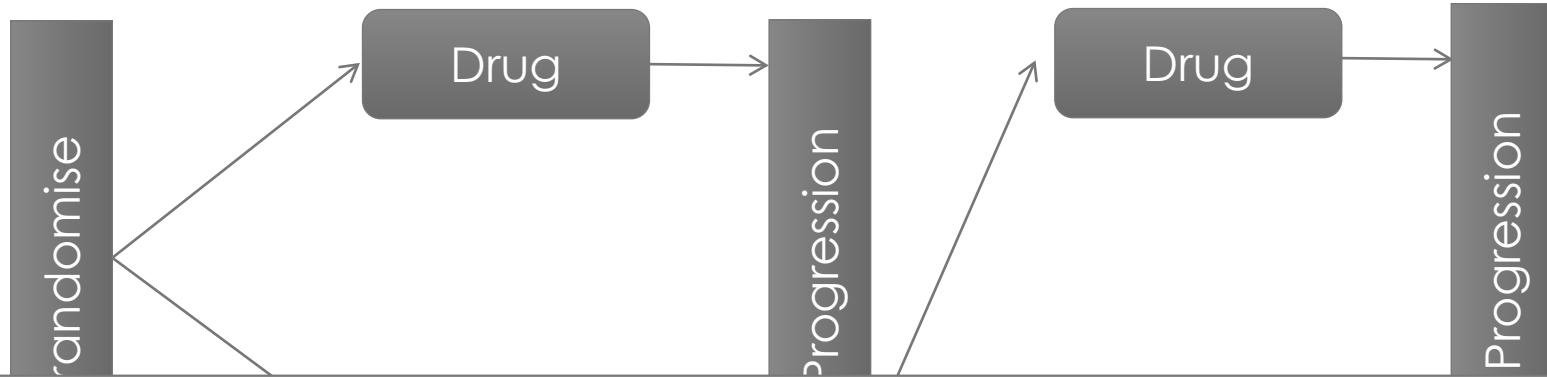
\$1B PBS reimbursement and drug access for 1,000 Australians with NSCLC



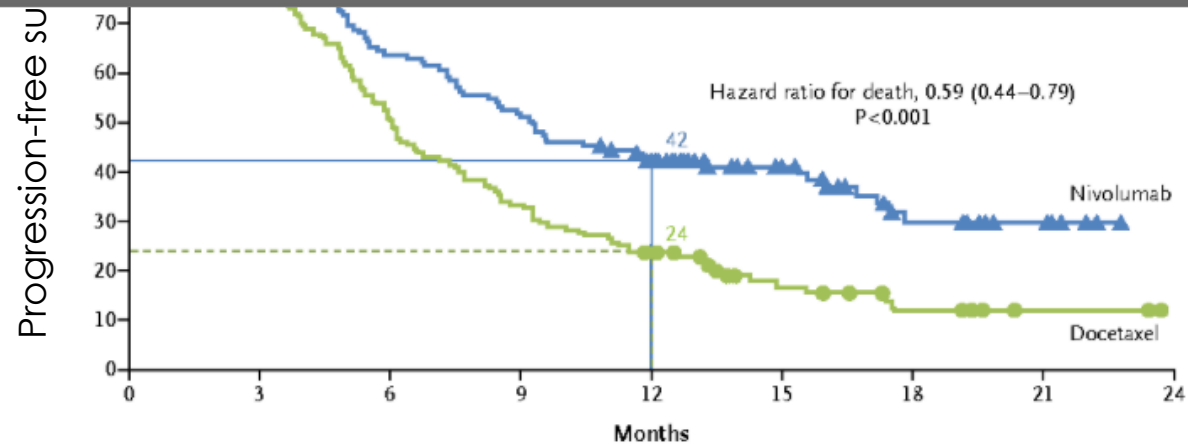
Rare pan-cancer  
TMB high



Rare pan-cancer  
TMB high



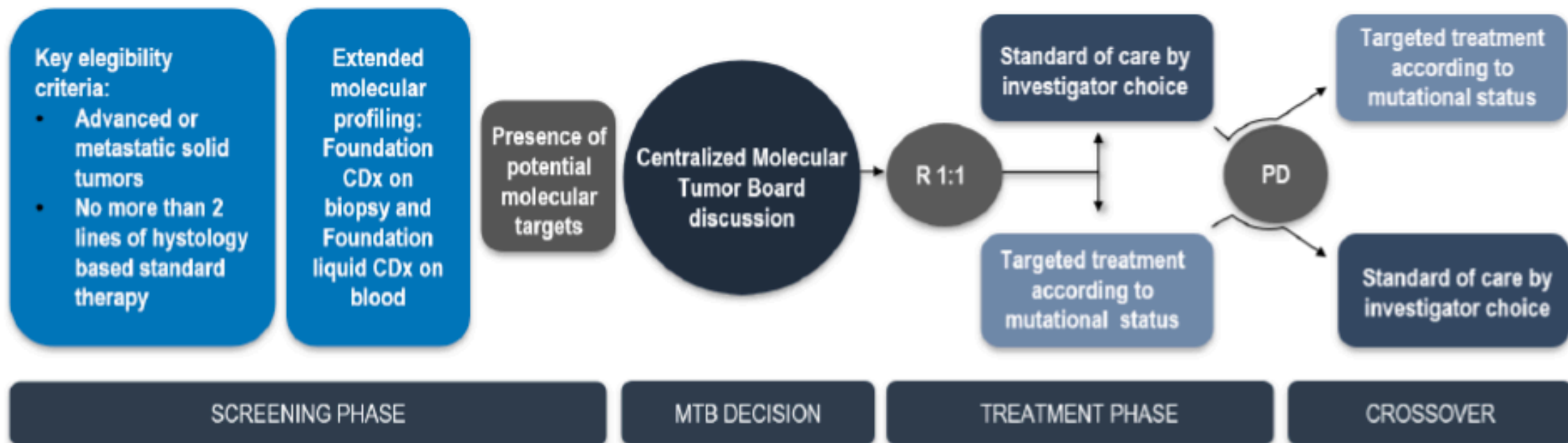
1,200 Australians with rare TMB-high cancers: we don't randomise pan-cancer



No. at Risk	0	3	6	9	12	15	18	21	24
Nivolumab	135	113	86	69	52	31	15	7	0
Docetaxel	137	103	68	45	30	14	7	2	0

# The ROME Trial:

A Randomized, Multi-basket, Phase 2, Multicentric Trial (NCT04591431)



- ✓ For each patient, we performed CGP using NGS with Foundation One, both on tissue samples and liquid biopsies. Historical samples were allowed but had to be collected within three months of patient screening
- ✓ Once molecular alterations were identified in tumor tissue or liquid biopsy, that could potentially be targeted by drugs available within the study the clinical case was reviewed by the Molecular Tumor Board (MTB)
- ✓ The MTB then decided whether to enroll the patient and proceed with randomization or to classify the case as a screening failure
- ✓ At PD crossover was planned

**PRIMARY ENDPOINT:** ORR  
**SECONDARY ENDPOINT:** PFS, TTF, TTNT, safety profile, OS.  
**EXPLORATORY ENDPOINTS:** PFS in homogenous subgroups of pts

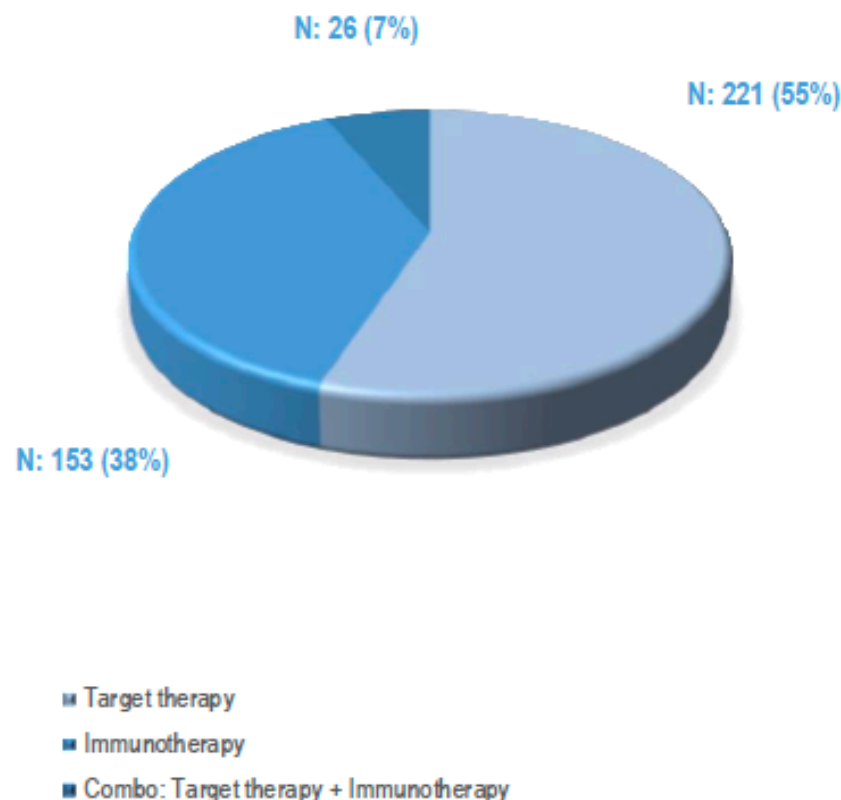
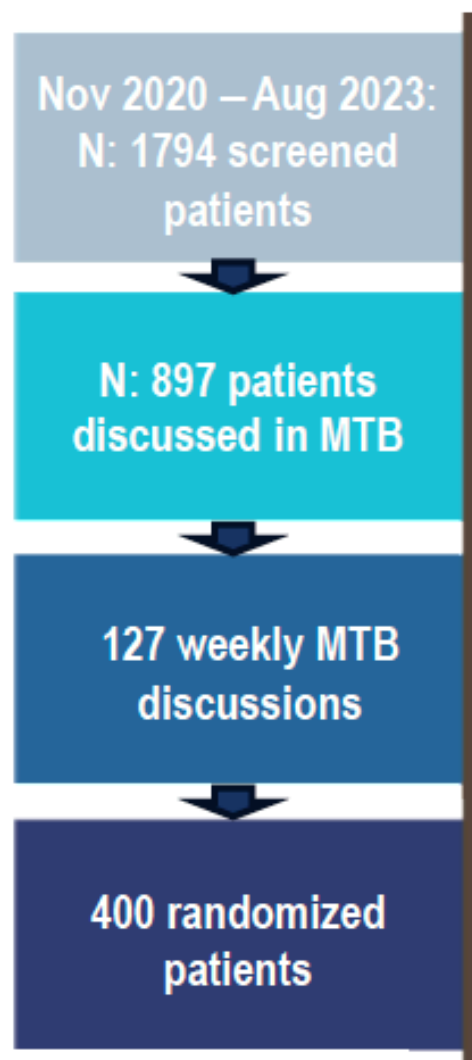
# The ROME trial: Available targeted treatments and pathways

Targeted Therapy	Target pathways
Erlotinib	EGFR mutations
Trastuzumab, pertuzumab, TDM1, lapatinib	ERBB2 amplification/mutations
Everolimus	mTOR mutations, AKT mutations
Vemurafenib, cobimetinib	BRAF mutations
Alectinib, brigatinib	ALK traslocations, RET alterations
Pralsetinib*, selpercatinib*	RET alterations
Palbociclib	CDK4/6, CDKN2A/B alterations
Ponatinib	Bcr-abl traslocations
Vismodegib	SMO/PTCH1 mutations
Itacitinib	JAK mutations
Pemigatinib	FGFR1/2/3 alterations
Ipatasertib, alpelisib*	Pi3k, AKT, PTEN alterations
Entrectinib*	NTRK1/2/3 fusions, ROS1
Tepotinib	METex14, MET alterations
Talazoparib	BRCA 1/2 mutations, HRD
Immunotherapy	Biomarkers
Nivolumab, ipilimumab, atezolizumab	MSI, high TMB**

\* Drugs included in the V4 protocol amendment in 10/2022

\*\* high TMB defined as >10 mut/mb in solid and/or liquid NGS

# The ROME trial: treatments assigned



Alteration group	Therapy assigned	n	
ALK/ROS1	Alectinib	4	
	Entrectinib	2	
Cell cycle	Palbociclib	5	
Erb	TDM1	31	
	Pertuzumab + Trastuzumab	12	
	Trastuzumab + Lapatinib	11	
	Pertuzumab + TDM1	2	
	TDM1 + Atezolizumab	1	
	Trastuzumab + Everolimus	1	
FGFR	Pemigatinib	33	
HRD	Talazoparib	7	
JAK/STAT	Itacitinib	1	
MET	Tepotinib	1	
Other	Everolimus	2	
	Ipatasertib	66	
	Atezolizumab + Ipatasertib	24	
	Alpelisib	6	
PIK3CA/AKT/PTEN	Everolimus	3	
	PTCH1	Vismodegib	2
	RAS/RAF/MAPK	Vemurafenib + Cobimetinib	16
Cobimetinib		14	
Cobimetinib + Atezolizumab		1	
RET	Pralsetinib	1	
	Selpercatinib	1	
Signature	Ipilimumab + Nivolumab	148	
	Nivolumab	5	

# The ROME trial: characteristics of ITT population

Characteristic	Overall N: 400 (%)	Arm A: Standard of Care N: 200 (%)	Arm B: Targeted Therapy N: 200 (%)
<b>Age (years)</b>			
median (range)	61 (22-85)	60 (34-84)	62 (22-85)
<b>Gender</b>			
Male	192 (48)	100 (50)	92 (46)
Female	208 (52)	100 (50)	108 (54)
<b>Ethnicity</b>			
Caucasian	393 (98)	197 (98)	196 (98)
Oriental	2 (0.5)	0	2 (1)
Hispanic	3 (0.8)	2 (1)	1 (0.5)
Afro-american	1 (0.1)	1 (1)	0
Other	1 (0.1)	0	1 (0.5)
<b>Primary tumor</b>			
Colorectal	65 (16)	31 (16)	34 (17)
Breast	39 (10)	19 (9)	20 (10)
Gastric	36 (9)	19 (9)	17 (9)
Glioblastoma	36 (9)	21 (10)	15 (7)
Biliary tract/gallbladder	36 (9)	10 (5)	26 (12)
NSCLC	35 (9)	18 (9)	17 (9)
Ovarian	18 (5)	8 (4)	10 (5)
Pancreatic	17 (4)	10 (5)	7 (4)
melanoma	13 (3)	8 (4)	5 (2)
anus	11 (2)	5 (3)	6 (3)
Other	94 (24)	51 (26)	43 (22)
<b>Enrollment line</b>			
Second	204 (51)	99 (50)	105 (52)
Third	196 (49)	101 (50)	95 (48)

# Primary endpoint: ORR in ITT population

Best response	Arm A Standard of Care N: 200 (%)	Arm B Targeted Therapy N: 200 (%)	p value
CR	0	5 (2.5%)	
PR	19 (9.5%)	29 (14.5%)	
SD	35 (17.5%)	36 (18.0%)	
PD	147 (73.0%)	130 (65.0%)	
ORR	9.5% [5.8-14.4]	17.0% [12.1-22.9]	0,027

Global ORR  
13.3%

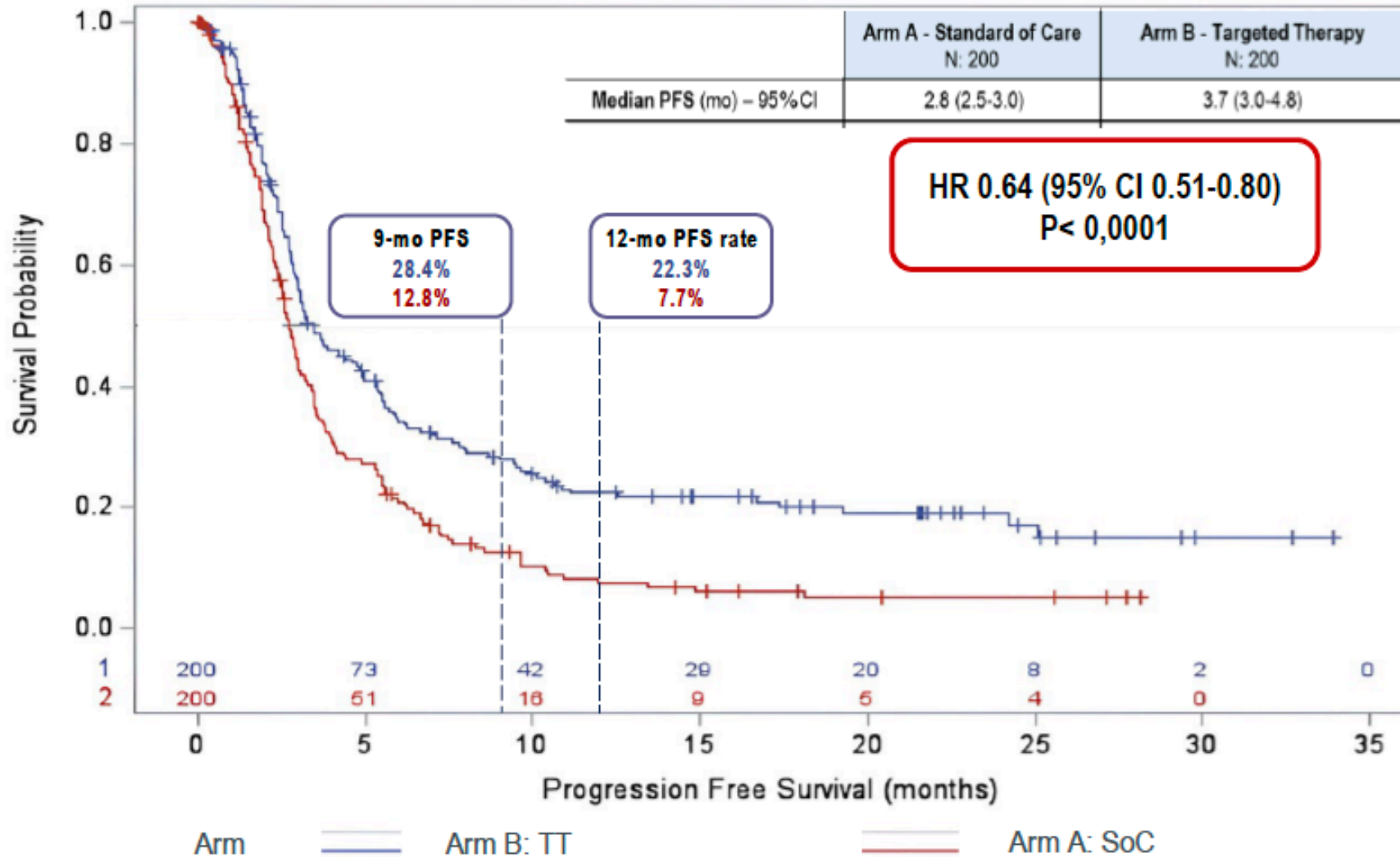
9.5%  
19/200

SOC

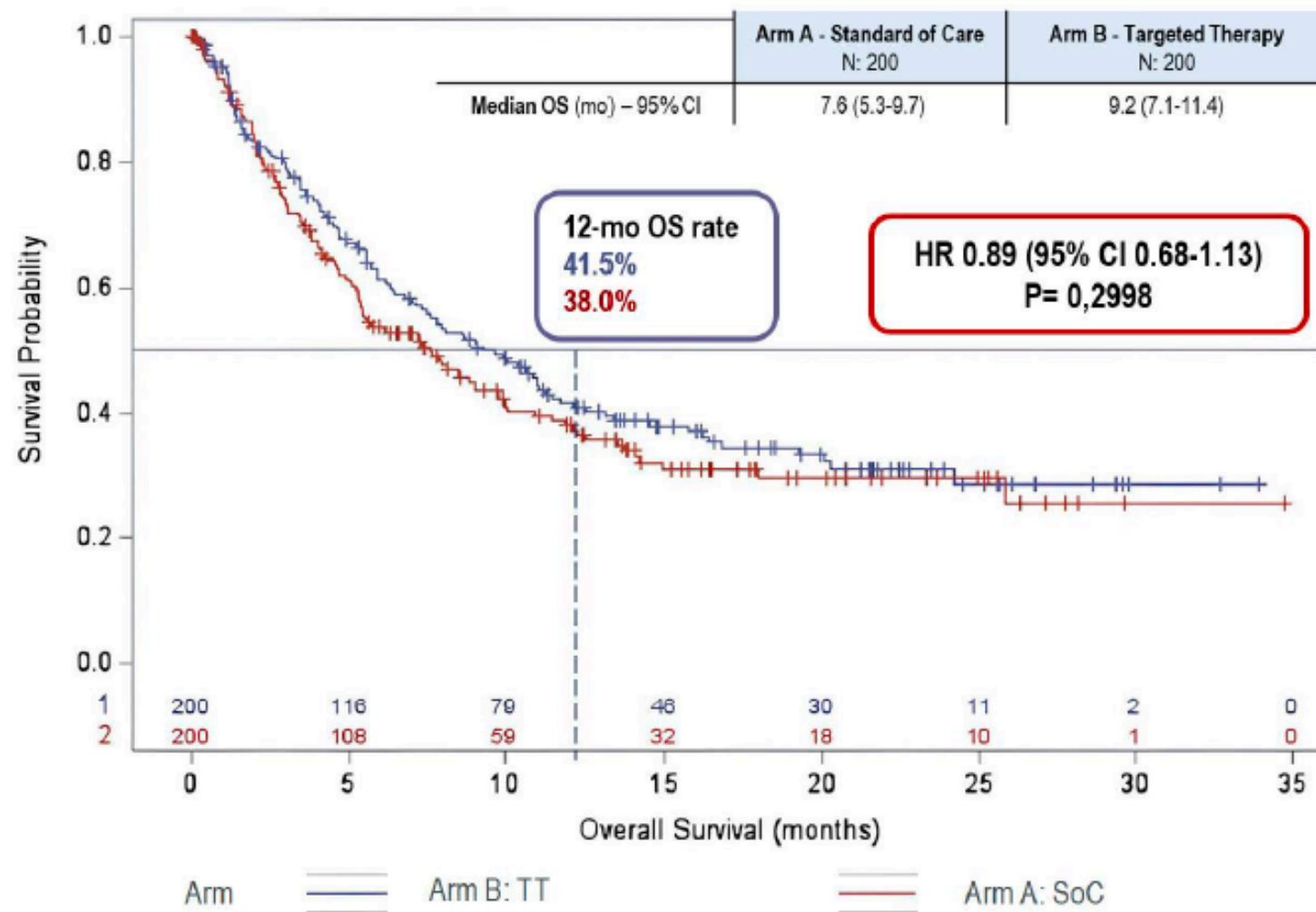
17.0%  
34/200

TT

# Secondary endpoint: PFS in ITT population



# Secondary endpoint: OS in ITT population





Thank you!

