

OPTIMISE: Optimal Precision Therapies to Customise Care in Childhood and Adolescent Cancer

Daniel A Morgenstern¹, Rebecca Deyell², Sarah Cohen-Gogo¹, Monia Marzouki³, Denise Connolly¹, Jordan Staunton¹, Marion Mateos⁴, David Ziegler⁴
 1) Hospital for Sick Children, Toronto; 2) BC Children's Hospital; 3) CHU Ste Justine; 4) CCI, University New South Wales

Goal

- OPTIMISE is a precision oncology platform study designed to leverage findings from ZERO and PROFYLE national sequencing initiatives to provide treatment options for patients with relapsed/refractory solid tumours and brain tumours

Objectives

Primary aims

- To use molecular sequencing to allocate children, adolescent and young adult (CAYA) patients with advanced solid tumours to specific treatment arms of molecularly-targeted agents;
- To determine the recommended phase II dose of a novel single agent or combination in the CAYA patient population;
- To determine the objective response rates (ORR; complete and partial response) in CAYA patients with advanced solid tumours with defined specific genomic alterations treated with molecularly-targeted agents.

Secondary aims

- To estimate the overall clinical benefit rate (CBR; complete and partial response and stable disease) in CAYA patients with advanced solid tumours with defined specific genomic alterations treated with molecularly-targeted agents;
- To estimate the progression free survival (PFS) in this CAYA patient population receiving molecularly-targeted agents;
- To obtain information about the tolerability and safety of molecularly-targeted agents and/or novel combinations;
- To obtain estimates of pharmacokinetics of molecularly-targeted agents and/or novel combinations in this CAYA patient population.

Exploratory aims

- To explore correlation between symptoms reported by patients and/or caregivers and AEs reported by clinicians.
- To assess the impact of study participation on symptoms reported by patients and/or their caregivers.
- To explore engagement in self-reporting of symptoms by patients and/or caregivers

Study method

- Multi-arm platform study open across Australia and Canada
- ANZCHOG is international sponsor with C17 Council regulatory coordinator for Canada
- Patient's must be pre-enrolled on ZERO, PROFYLE or equivalent comprehensive sequencing program
- Study arm-specific eligibility criteria apply

Results

- Study is ongoing, no results currently publicly available

Impact/outcomes

- Study activated in both Australia and Canada
- Hospital for Sick Children, CHU Ste Justine and BC Children's Hospital open for Arm A
- CHEO and Stollery in process of opening
- Planning to activate IWK for Arm A via CRAFT decentralised model
- Arm C activated in Australia, awaiting activation at SickKids and then other sites

Project timelines

- Initial Australian Medical Research Future Fund Grant 2021
- Grants from Cancer Research Society and CIHR for Canada in 2022
- Initial regulatory approval from Health Canada September 2024
- Study is ongoing

The challenge



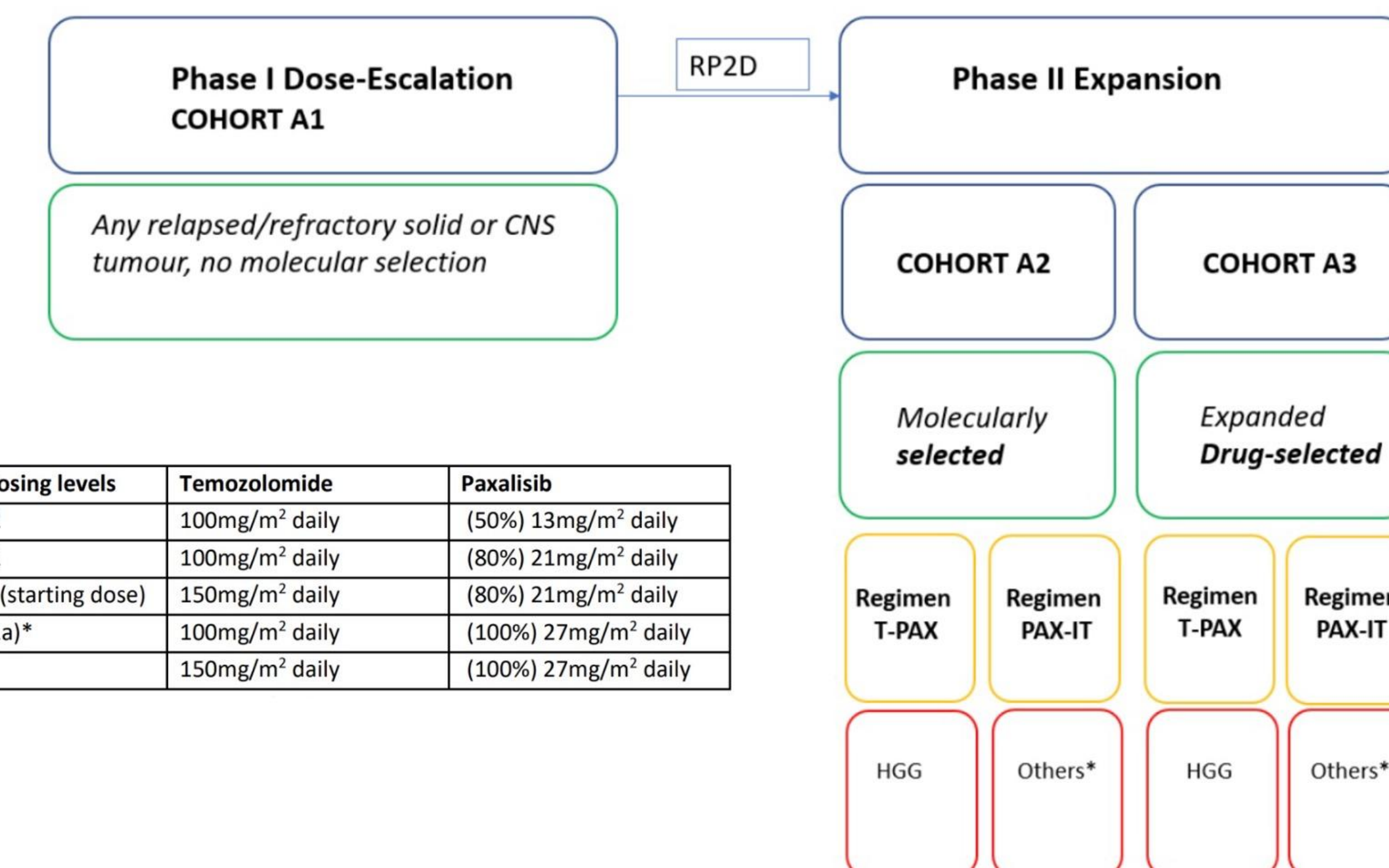
PROFYLE

Comprehensive access to cancer sequencing across Canada for CAYA patients

- No 'built in' access to targeted therapeutic interventions
- Lack of available clinical trials
- Lack of access to clinical trials/targeted agents

Arm A: paxalisib plus chemotherapy

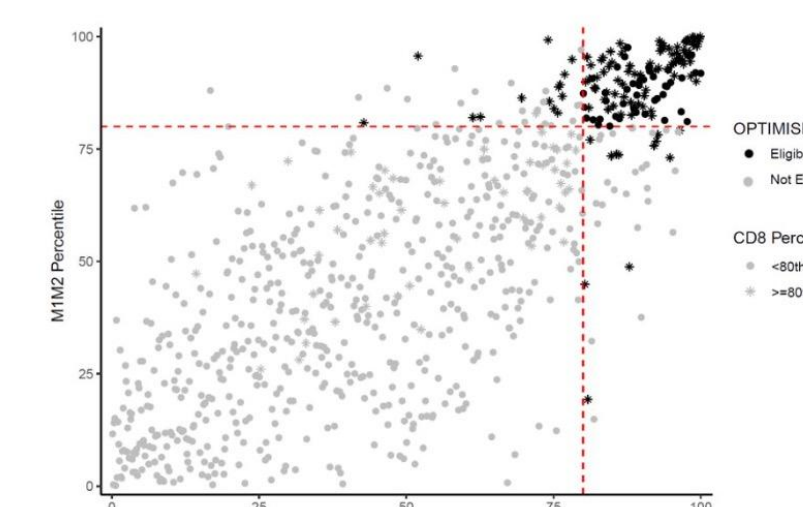
Paxalisib is a CNS-penetrant PI3K/mTOR inhibitor
 Combination with temozolomide ± irinotecan



Arm C: nivolumab + relatlimab (Opduolag)

Combined PD1 and LAG3 inhibition

- Cohort C1 Immune infiltrated
 ≥80th centile for 2 of 3 scores:
- CD8 T-cell
 - M1/M2 macrophage
 - iPASS



- Cohort C2 RRD deficient
 failed to respond or relapsed after anti-PD(L)1

- Combined PD1/LAG3 inhibition (Opduolag)
- Enrolment limited to patients able to receive fixed adult dose
- Patient selection based on immune infiltration determined by RNAseq
- Pre-screening through existing national sequencing programs in Canada and Australia (PROFYLE/ZERO)
- Pilot study for initial signal of efficacy (12 patients)
- Additionally allow enrolment of patients with RRD tumours previously treated with anti-PD1 monotherapy (≤12 patients)

Overall study schema

