Clinical and molecular characterization of IDH1/2 mutant cholangiocarcinoma

Florian Castet^{1,2}, Queralt Serra-Camprubí², Carles Fabregat-Franco^{1,2}, Helena Verdaguer^{1,2}, Gloria Castillo¹, Tian V.Tian², Teresa Macarulla^{1,2}*

¹ Gastrointestinal Cancer Unit, Oncology Department, Vall d'Hebron University Hospital, 08035, Barcelona, Spain ² Preclinical and Translational Research Program, Vall d´Hebron Institute of Oncology (VHIO), 08035 Barcelona, Spain. [†] Joint first authors *Corresponding author; tmacarulla@vhio.net

• Cholangiocarcinoma (CCA) is the second most common liver cancer and a leading cause of cancer-related death worldwide.

• 15-20% of CCA tumours harbour **IDH1/2 mutations (IDH**^{mt}) that can be targeted with IDH inhibitors, such as ivosidenib, which improve progression-free survival in second-line treatment.

GOAL: Provide a **clinical characterization of IDH**^{mt} **CCA** in routine clinical practice and **generate a collection of patient-derived xenografts (PDXs)** to perform drug efficacy evaluation.

PART 1: Clinical characterization of IDHmt CCA

 Key eligibility criteria Histologically confirmed advanced CCA Genomic profiling of primary or metastatic tumour. 	IDH ^{mt} CCA N=77
	IDH ^{wt} CCA N=254

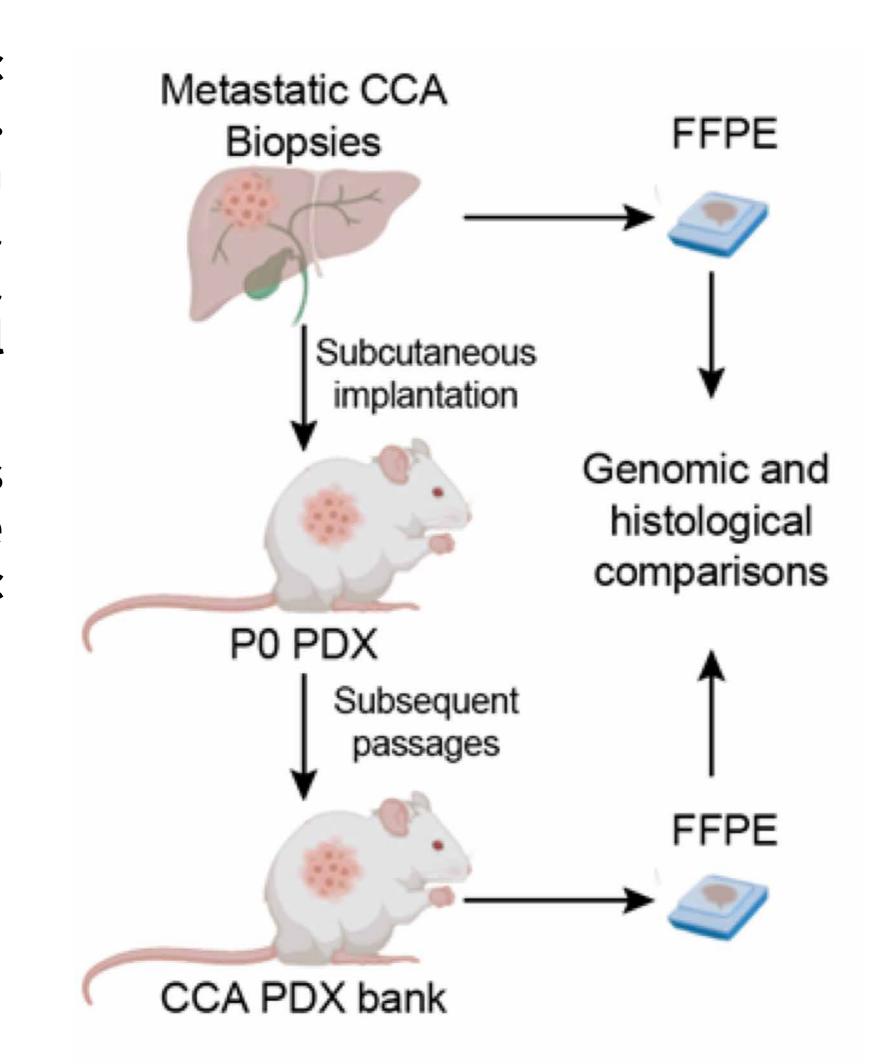
Aims

- **1.** Compare the clinical characteristics of IDH^{mt} vs IDH^{wt} CCA.
- 2. Evaluate the survival outcomes of IDH^{mt} patients treated with IDH inhibitors.

PART 2: Generation of a CCA PDX collection

Tumour pieces from metastatic lesions were implanted into NOD. CB-17-Prkdc scid/Rj mice. Upon growth of the engrafted tumours, a tumour piece was implanted into a new recipient mouse for the model perpetuation.

H&E staining, IHC of CCA markers (KRT19 and HepPar1), and NGS were used for histological and genomic characterizations.



RESULTS

Table 1: Baseline characteristics comparing IDH^{mt} with IDH^{wt} patients.

	IDH ^{mt} (N=77)	IDHwt (N=254)	P-value
Age [Median (range)]	59 (35-84)	62 (27-92)	<0.01
Gender (Female, %)	50 (64.9%)	113 (44.4%)	<0.01
Stage IV at diagnosis (N, %)	55 (71.4%)	144 (54.6%)	0.02
Surgery (N, %)	17 (22%)	113 (44.4%)	<0.01
Adjuvant therapy (N, %)	6 (7.79%)	65 (25.5%)	<0.01
Metastasic location (N, %)			0.9
No visceral	10 (12.9%)	37 (14.5%)	
Visceral	62 (80.5%)	204 (80.3%)	
No metastasis	5 (6.4%)	13 (5.1%)	
First-line regimen (N, %)			0.7
Single agent	6 (7.7%)	26 (10.2%)	
Gem-based combination	60 (77.9%)	200 (78.7%)	
Other treatment	6 (7.7%)	12 (4.6%)	
Best supportive care	5 (6.4%)	16 (6.3%)	
MSI status (N, %)	1 (1.3%)	4 (1.5%)	1
OS [median (95% CI)]	19.5 (16.2-24.1)	15.1 (13.6-18.2)	0.3*

MSI: microsatellite instability, OS: overall survival. *Log-rank p-value

Figure 1: Overall and progression-free survival of IDH^{mt} patients who received 2L IDH inhibitors.

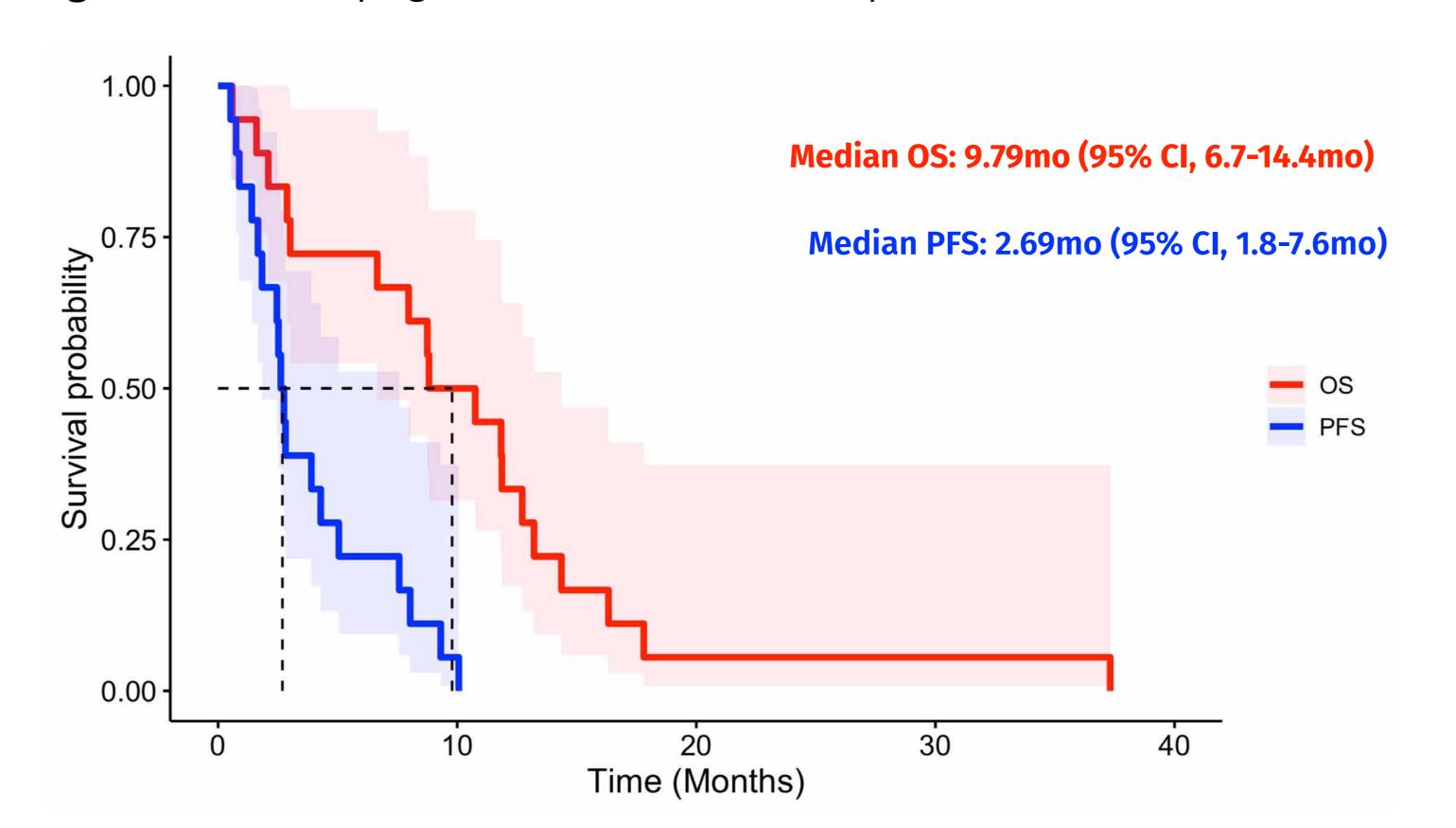
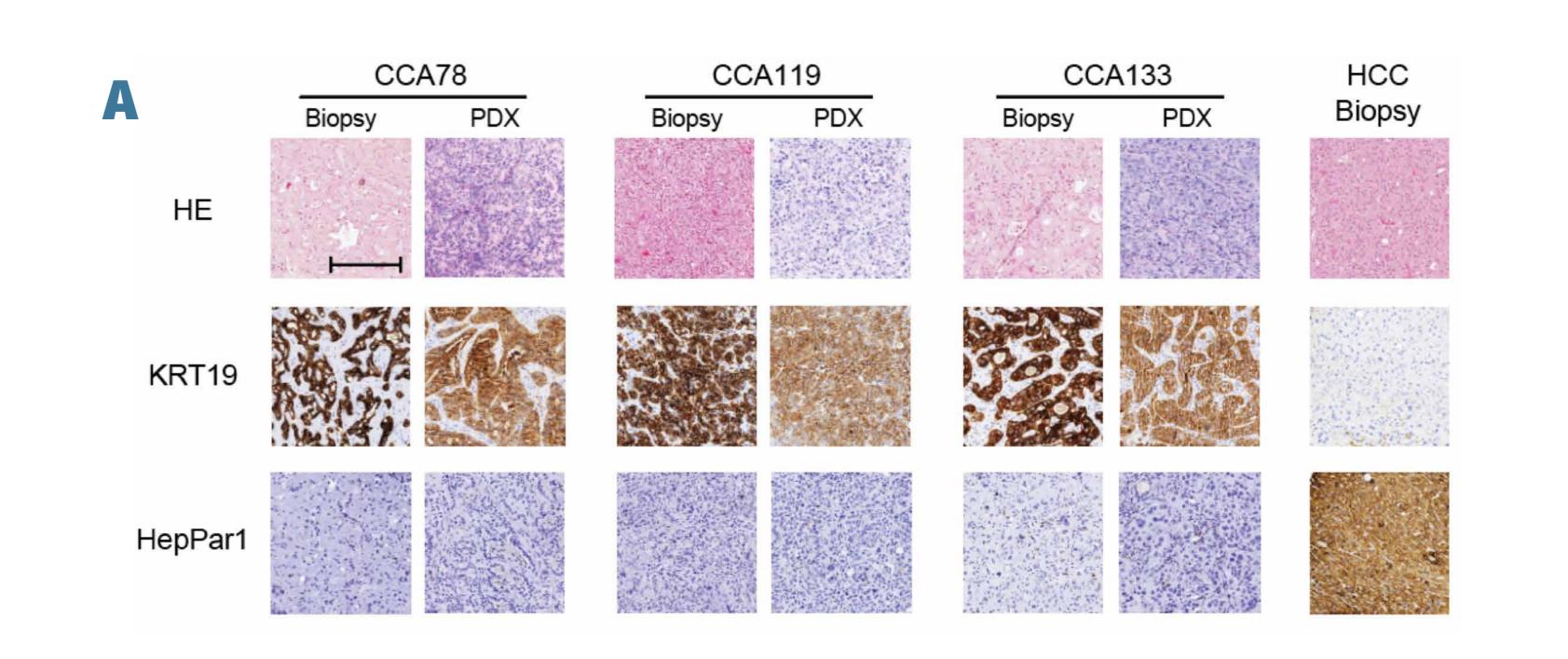


Figure 2: Histopathologic, immunohistochemistry (A) and mutational analysis confirmed that CCA_PDXs maintain the histologic and genomic features of the original biopsy specimens.



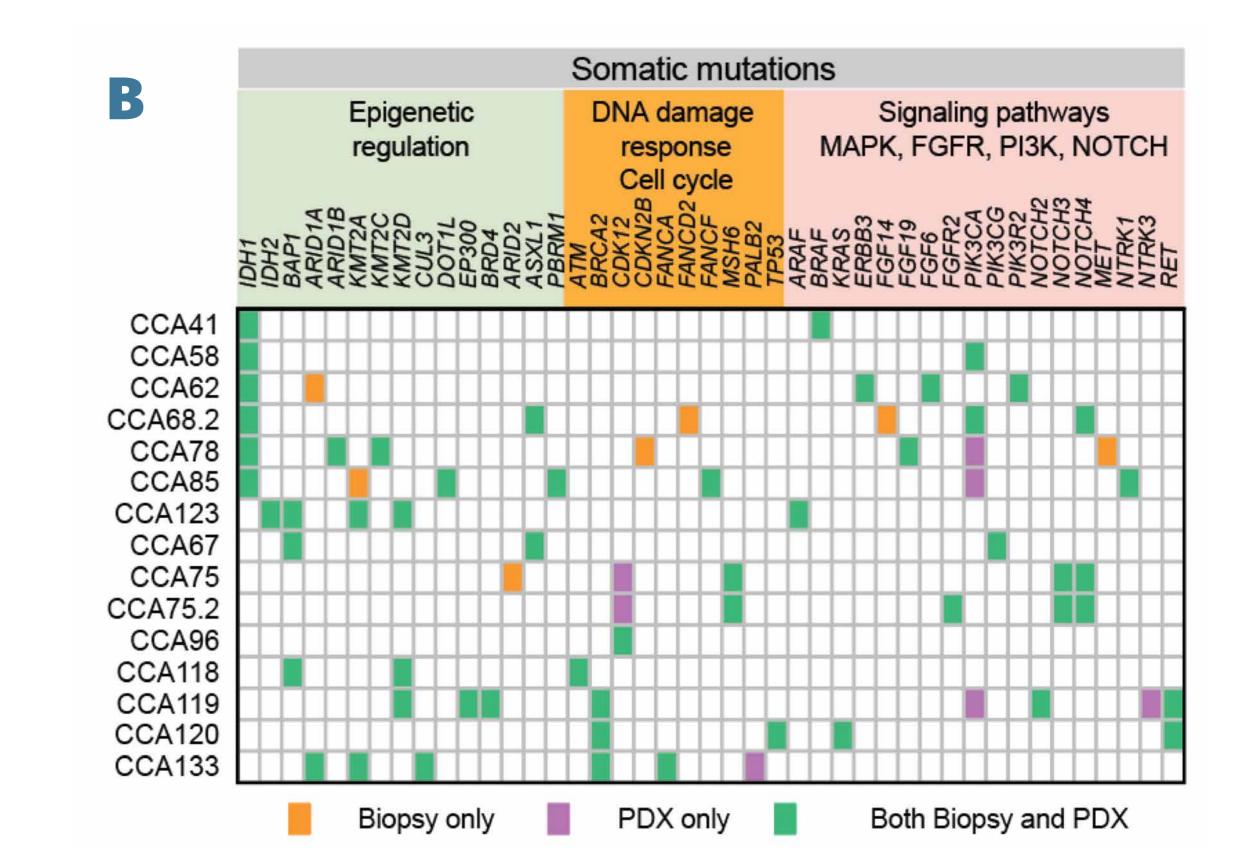
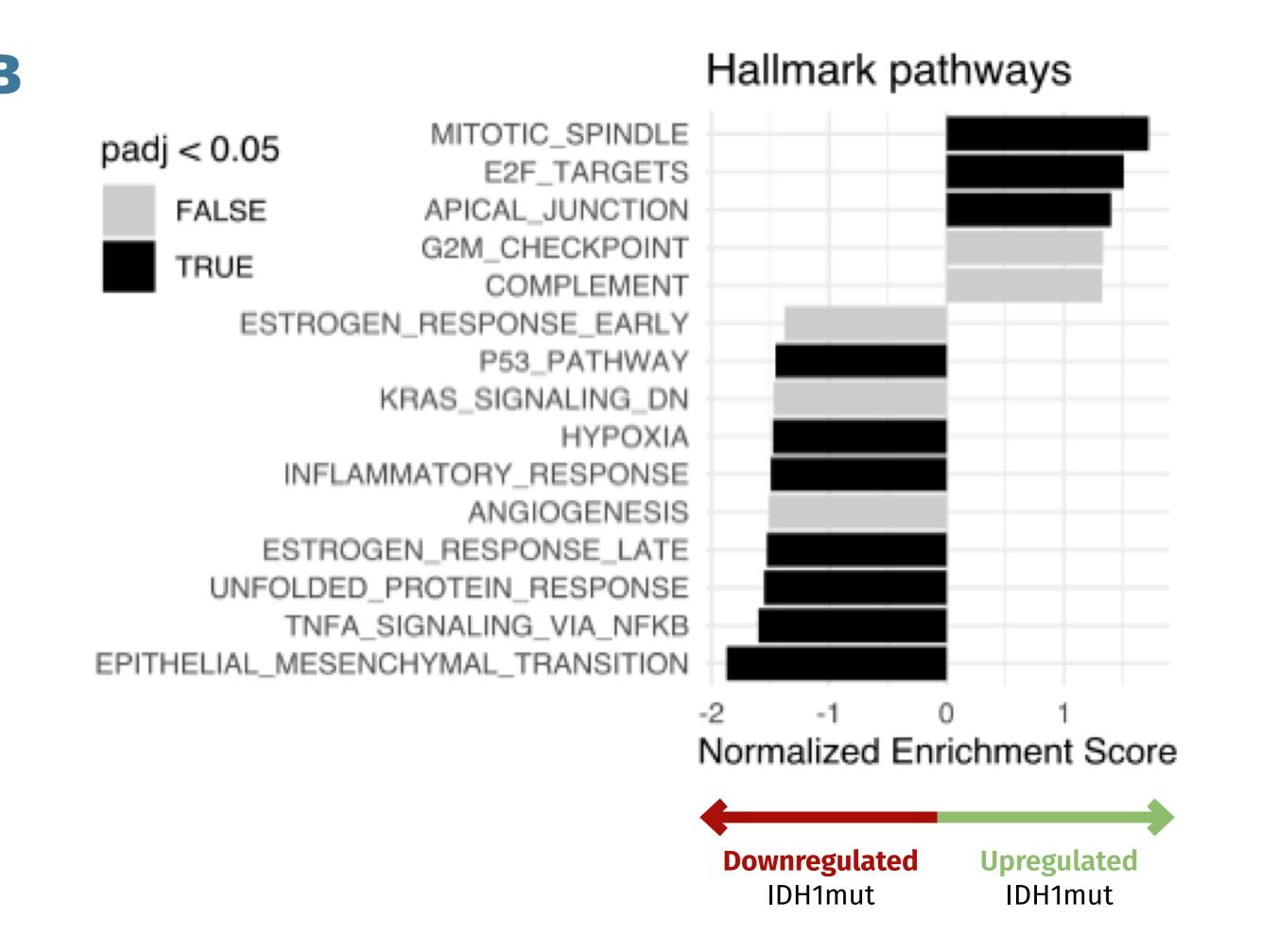


Figure 3: Differential gene expression analysis (A) and pathway enrichment (B) comparing IDH^{mt} with IDH^{mt} shows a downregulation of immune-related pathways in IDH^{mt} tumours.





CONCLUSIONS

- IDH^{mt} tumours are more frequent in young female patients, with more advanced disease at presentation. Patients treated with IDH inhibitors at our tertiary centre present similar survival outcomes to those reported in the phase III ClarIDHy trial.
- Genome-wide expression profiling of PDX derived from advanced CCA patients suggest the potential involvement of IDH1^{mt} in CCA tumour immune microenvironment alterations, suggesting a **potential synergistic effect of IDH inhibitors with immune modulating agents.**