Cleveland Clinic

Abstract 536 : Molecular Differences with Therapeutic Implications in Early-onset Compared to Average-onset Cholangiocarcinoma

MERIT AWARD THE ASCO FOUNDATION RECIPIENT Sanjay Goel^g, Tiago Biachi de Castria^h, Vaia Florouⁱ, Kanika G. Nair^{a,j}, Suneel D. Kamath^{a,j}, Alok A. Khorana^{a,j}



Thejus Jayakrishnan^a, Yasmine Baca^b, Joanne Xiu^b, Mehrie H Patel^a, Benjamin Weinberg^c, Emil Lou^d, Jashodeep Datta^e, Moh'd Kushman^f, Pat Gulhati^g, ^aTaussig Cancer Institute, Cleveland Clinic Foundation, ^bCaris Life Sciences, ^cGeorgetown University of Minnesota, ^eUniversity of Minnesota, eUniversity of Miami - Sylvester Comprehensive Cancer Center, d ^fWashington University Siteman Cancer Center, ^gCancer Institute of New Jersey, Rutgers University^hMoffitt Cancer Center, ^hHuntsman Cancer Institute, ⁱUniversity of Utah Health, ^jCase Comprehensive Cancer Center

Background

- Cholangiocarcinoma (CCA) has one of the fastest rising incidence rates (8% APR) among early-onset (eo) cancers.
- Real-world age-stratified multi-omic characterization of eoCCA could reveal molecular features with prognostic and therapeutic implications.
- We sought to compare the molecular characteristics of eoCCA with average-onset CCA (aoCCA) utilizing a realworld multi-omics dataset.

Methods

- The study comprised patients whose tumors underwent molecular analysis at Caris Life Sciences (Phoenix, AZ) using whole exome and whole transcriptome analyses.
- Patients were categorized by age as eoCCA defined as <50 years and aoCCA >50 years.
- values were adjusted for multiple testing and significant at Q<0.05 for molecular considered comparisons and Q<0.25 for Gene Set Enrichment Analysis (GSEA).
- Insurance claims data was used for survival comparison using Kaplan-Meier estimates.

Results

The baseline characteristics listed below:

Table 1. Baseline Characteristics		
Characteristic	eoCCA	aoCCA
	(n=453)	(n=5134)
Median Age (years)	44	68
Sex: Female	60.70%	54.60%
Sex: Male	39.30%	45.40%
Primary Tumor Site		
Intrahepatic	57%	53%
Extrahepatic	14%	16%
Gallbladder	21%	23%
Others	8%	8%

Author Contact : thayyit@ccf.org Acknowledgement: The Sandra and Stephen Hardis Endowment Fund Copies of this poster obtained through Quick Response (QR) Code are for personal use only and may not be reproduced without permission from ASCO[®] or the author of this poster X @thejusjay

Targetable Molecular Alterations

- (Figure 1).
- \bullet aoCCA, Q<0.001)



Conclusions

**q<0.05 Figure 1. Prevalence of Targetable Molecular Alterations Immune Oncology Relevant Markers • MSI-H tumors (4.1% eoCCA vs 2.4% aoCCA), and high- TMB (6.1% vs 5.1%) were numerically higher in eoCCA (Q=1). The Interferon Gamma - IFG Score (Fold P=0.06 Change FC: 1.1, Q=0.01) and T-Cell aoCCA eoCCA Inflammation Score TIS (FC:17.3, Q=0.03) Figure 3. Distribution were higher in aoCCA (Figure 2). of Inflamed Tumors The distribution of inflamed tumors were Enriched in aoCCA Enriched in aoCCA not significantly different, P=0.06 NES=-1.46 NES=-1.58 P=0.00 (Figure 3). P=0.00 On GSEA, angiogenesis was enriched in eoCCA (normalized enrichment score NES=1.51, Q=0.16) while IFG (NES= -1.58,

Q = 0.06) and inflammatory response (NES= -1.46, FDR-p=0.18) were enriched in aoCCA.

• Differences in the prevalences of targetable alterations were observed

FGFR2 fusion was significantly more prevalent in eoCCA (15.7% vs 5.9% in

eoCCA aoCCA



In the largest age-stratified analysis of molecular characteristics of CCA, we identified crucial differences, including higher prevalence of FGFR2 fusions and significant differences in immunotherapy-related markers.

Patients with eoCCA experienced better outcomes that was impacted by the FGFR2 fusion status.

Our findings, especially higher FGFR2 fusion prevalence in eoCCA, underscore the need for NGS testing



CONQUER CANCER

Survival Analysis

- Median OS was longer in eoCCA (16.5 vs 13.3 mo), HR 0.86, 95%CI 0.78-0.95, *P*=0.004.
- Among patients exposed to immunotherapy, median OS was longer in patients with eoCCA (19.2 vs 8.1 mo) - HR 0.54, 95%Cl 0.30-0.98, p=0.04 (Figure 5).
- No survival difference was identified in patients on chemotherapy (HR 0.91, 95%CI 0.76-1.08, P=0.28).



Figure 5. Survival Analysis Results (Left) and Time on Treatment (Right) for Patients Exposed to Immunotherany

Survival by FGFR2 Fusion Status

• Median OS was highest for patients with FGFR2 FusioneoCCA 21.7 vs aoCCA 18.6mo, followed by FGFR2 Fusion negative -eoCCA 15.0 vs aoCCA 12.2mo, *P*=0.002 (Figure 6).



Figure 7 Shows differentially expressed molecular alterations Figure 6. FGFR2 Fusion Status and Survival Outcomes (Q<0.05).





Figure 7. Volcano Plot of Molecular Differences in FGFR2 Fusion Positive vs Negative eoCCA (left) and aoCCA (right)

References:

- 1. Koh et al. Patterns in Cancer Incidence Among People Younger Than 50 Years in the US, 2010 to 2019. JAMA Network Open. 2023
- 2. Ogobuiro et al. Multiomic characterization to reveal a distinct molecular landscape in young-onset pancreatic cancer. JCO. 2022