Validation of PDACai v2.0 in predicting relative benefit from frontline FOLFIRINOX (FFX) and gemcitabine/nab-paclitaxel (GA) for patients (pts) with metastatic pancreatic cancer (mPDAC)







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PANCREATIC CANCER ACTION **NETWORK**

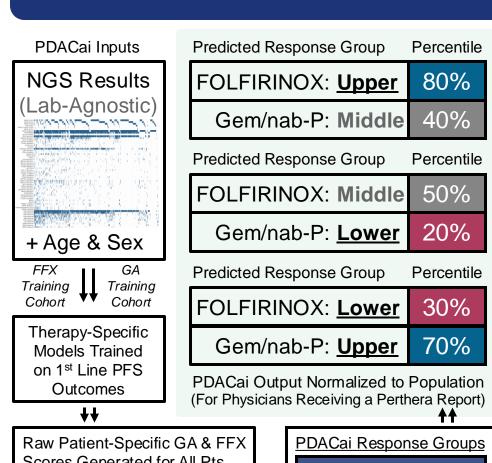
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Background

- Nearly 50% of pts with mPDAC never receive a 2nd line therapy for metastatic disease following frontline FFX or GA
- Genomic alterations in the DDR pathway² (e.g. BRCA1/2) are associated with longer progression-free survival (PFS) on platinum-containing regimens (e.g. FFX). PDACai uses the full NGS profile to predict benefit from GA and/or FFX in mPDAC.
- PDACai v2.0 redefines the mutational landscape in mPDAC by removing ATM mutations from the DDR pathway (top FFX feature) and adding new variant-level features (e.g. TP53 GOF).

PDACai Methods & Examples



Scores Generated for All Pts Jpper Tertile (67-100%) +FFX/GA Validation Cohorts ↓↓↓↓ PDACai Consensus Rankings Lower Tertile (0-33% (0-100th Percentile per Cohort)

Figure 1 - PDACai learned to predict outcomes based on NGS testing results from Training Cohorts where pts received either FFX & GA in the 1st line setting. PDACai predictions were independently analyzed for FFX & GA Validation Cohorts.

2nd Line Gem/nab-P (+/- New Trial Agent)

3rd Line Clinical Trial (see Perthera Report)

2nd Line Clinical Trial (see Perthera Report)

3rd Line Gemcitabine (+/- nab-Paclitaxel)

1st Line Gem/nab-P (+/- New Trial Agen

3rd Line Clinical Trial (see Perthera Report)

eligibility for nearby/accessible clinical trials

US pts whose physicians order a Perthera Report

may be eligible to participate in prospective efforts to

validate PDACai (frontline/adjuvant, non-interventional)

previous clinical/treatment history

PDACai predictions were analyzed

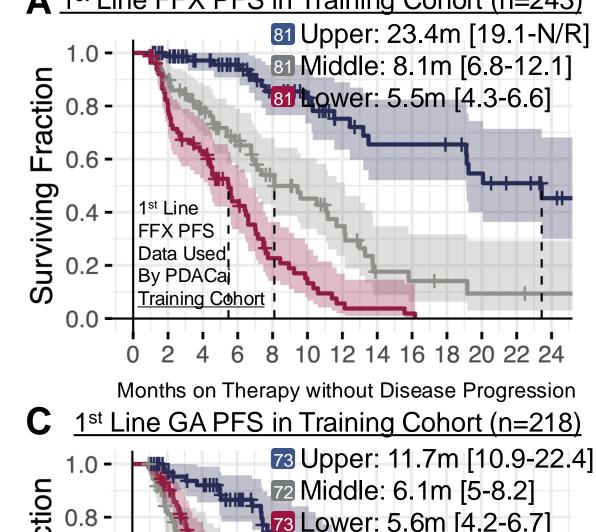
captured via the Perthera Registry.

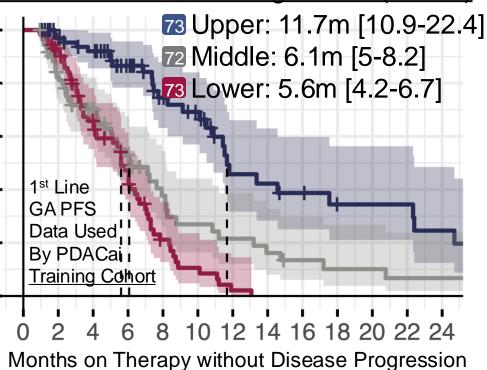
retrospectively for pts who received

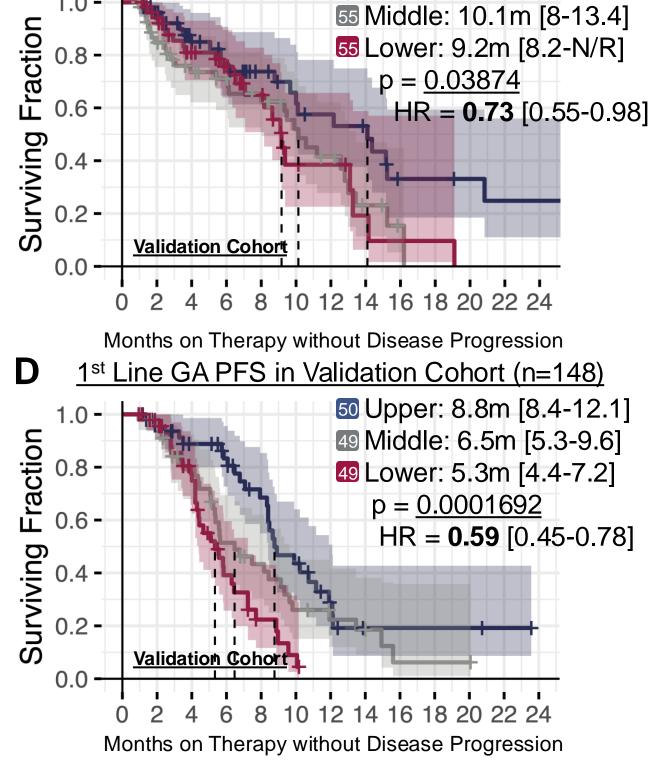
a Perthera Report. Outcomes were

NOTE: A Perthera Report's ranked therapy options

▲ 1st Line FFX PFS in Training Cohort (n=243)







B 1st Line FFX PFS in Validation Cohort (n=165)

55 Upper: 14.1m [10-39.6]

Figure 2 – KM curves of PFS on 1st line therapies from pts allocated to independent training (A,C) and validation (B,D) cohorts. Actual median PFS [plus 95% CI] in months were summarized in pts assigned to lower, middle, or upper thirds based on PDACai response group predictions (see Fig 1). The predictive utility of PDACai was confirmed in the independent validation cohorts (B,D) by comparing PFS across tertiles (see p-values and hazard ratios (HR) [plus 95% CI]).

1st Line PFS Stratified by Normalized PDACai Predictions

Advanced OS Correlates with PDACai Predictions

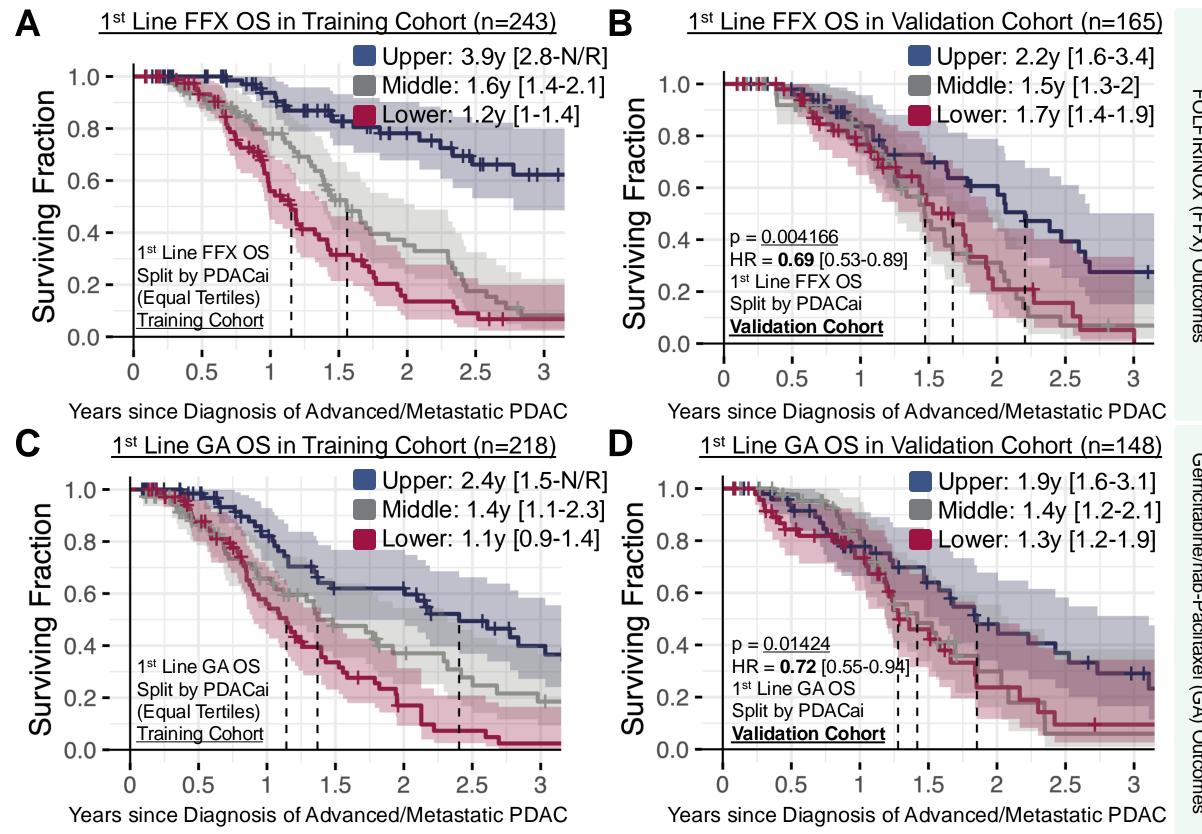


Figure 3 – KM curves of overall survival advanced diagnosis for treatment-specific PDACai predictions within the training (A,C) and validation (B,D) cohorts who received 1st line FFX (A,B) or GA (C,D).

#GI25 Poster #776

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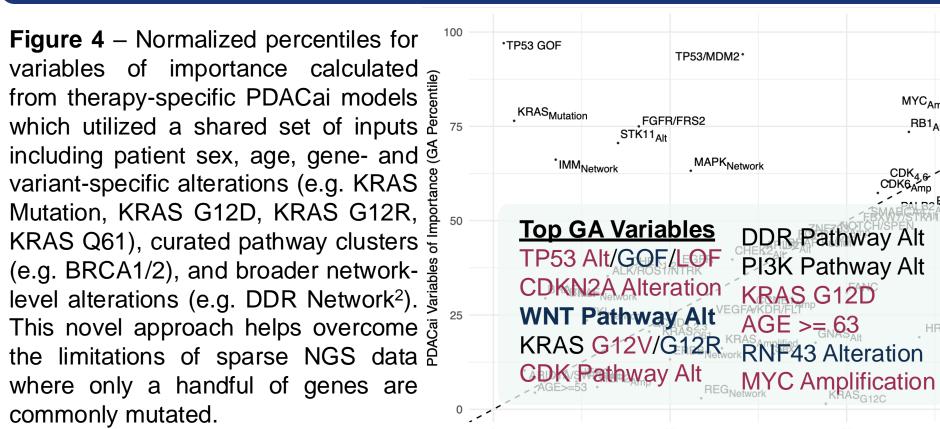
Conflicts of Interest

Perthera is a privately-held precision oncology company delivering multi-omic insights and real-world outcomes to support investigators, sponsors, and health systems

References

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- 2. Pishvaian et al. JCO Precision Oncology. Oct 2019; PMID: 35100730
- 3. Takebe et al. #GI24 DOI: <u>10.1200/JCO.2024.42.3 suppl.698</u>
- 4. Zohar et al. #GI24 DOI: 10.1200/JCO.2024.42.3_suppl.694
- 5. Hendifar et al. ASCO <u>10.1200/JCO.2020.38.15_suppl.464</u>

Treatment-Specific PDACai Variables of Importance



Top FFX Variables BRCA_{1,2} DDR Pathway Alt **WNT Pathway Alt BRCA1/2 Mutation** KRAS G12D/G12R **CDKN2A Alteration APC Mutation KRAS** Rare Variant **AKT2** Amplification **RNF43** Alteration AGE >= 63 **CDK Pathway Alt**

Conclusions

- Response to chemotherapy is heterogeneous and difficult to predict in pts with mPDAC
- PDACai v2.0 successfully predicted differences in PFS for both FFX and GA cohorts (better than v1.0)
- Guidance for 1st line decisions is now feasible with upfront NGS and PDACai v2.0 (faster than v1.0)
- Efforts to validate PDACai predictions for 2nd line variations of FFX (FOLFIRI vs FOLFOX) are ongoing
- Continuous prospective validation of AI/ML models that utilize lab-agnostic NGS results is warranted

PDACai predictions correlated with OS/PFS outcomes for FFX & GA cohorts. PDACai inputs routine NGS results and assigns Upper vs Middle vs Lower "Predicted Response Groups" from normalized FFX & GA scores (0-100%).

Recent Studies Explored Top NGS Features

<u>TP53 WT vs GOF vs LOF</u>⁴ & *More* (KRAS Variants⁵, BRCA/DDR⁶) - FFX Cohort: Longer OS for TP53 WT vs GOF/LOF

WNT Pathway Alterations (RNF43/GNAS/APC/CTNNB1)³ - GA Cohort: Shorter OS/PFS for TP53 LOF vs GOF/WT - Prognostic Associations Observed for FFX & GA Cohorts

Questions? eblais@perthera.com

BRCA1AIL PIK3CB/PIK3CG/PIK3R1 2

KRAS_{A146/K117/etc}

PDACai Variables of Importance (FFX Percentile)

DDR Pathway Alt

PI3K Pathway Alt

KRAS G12D

AGE >= 63