

Development of a novel RNA-based immune checkpoint inhibitor response signature (ICI-PRS) that avoids standard-of-care (SOC) prognostic signal for use in patients with urothelial cancer (UC) Shepherd JH, Mayhew GM, Guo J, Beebe KD, Eisner JR, Milburn MV

BACKGROUND

- Immune checkpoint inhibitors (ICIs) show durable responses in a portion of urothelial carcinoma patients¹, making predictive biomarkers beneficial
- Elevated PD-L1 or total mutation burden (TMB) are associated with ICI response¹, but also with response to chemotherapy^{2,3} in bladder cancer
- Confirmatory trials of ICIs within PD-L1 biomarker positive patients have failed, leading to withdrawal of atezolizumab and durvalumab in bladder cancer
- Better biomarkers, that distinguish ICI-response from non-ICI standard-of-care (SOC) are needed

METHODS

- RNA expression data from the IMvigor210 trial¹ (n=298) and from the Cancer Genome Atlas (TCGA) bladder cancer cohort⁴ (n=405) were combined, with 2/3 used to identify treatment-agnostic and ICI-specific prognostic genes (training set)
- Candidate ICI-specific genes were developed into an ICI-specific survival signature (sigi162) and applied to the withheld 1/3 of IMvigor210 and TCGA (test set)
- A novel composite ICI predictive response signature (ICI-PRS), featuring sigi162 and other selected signatures reflective of both tumor and immune microenvironment was developed in the same training sets and tested in IMvigor210 and TCGA test sets and two independent ICI-treated bladder cancer datasets: UNC-106 (BACI; n=84 receiving ICI>3 weeks)⁵ and BCAN UC-GENOME (BCAN; n=105)⁶

Figure 1: Prognostic Value of Sigi162 Signature (A) and TMB (B) in ICI-treated IMvigor210 and SOC-treated TCGA BLCA



- Sigi162 signature was trained to be prognostic for ICI-treated patients (e.g., IMvigor210, green lines) but not non-ICI treated patients (e.g., TCGA BLCA, blue lines). Note separation of green vs blue lines, with blue lines near 1 for Hazard Ratio of Cox Proportional Hazard models of overall survival when data is stratified at given quantile in Figure 1A
- TMB is prognostic in both ICI- and non-ICI treated patients. Note overlapping green and blue lines in Figure 1B
- Sigi162 signature was not clearly prognostic for ICI treatment in IMvigor210 test set (solid green line in Figure 1A), thus necessitating improvement with the addition of other features

C)

D)

A)

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Figure 2: ICI Predictive Response Signature (ICI-PRS) Combines Figure 4: PD-L1 and TMB are Complimentary to ICI-PRS Signature Sigi162 with Tumor Cell and Immune Microenvironment Features IMviaor210 test set IMvigor210 test set Α B) PD-L1 IC IHC **ICI-PRS** ICI-specific survival sig. (sigi162) tat3 acitivatio Ratio 0.50 E2F/RB cell cycle regulation (E2F/RB cell cycle regulation 0.25 Q cell activation 1.5 IC0 IC1 IC2+ 0.5 T cell activation 2 PD-L1 IC IHC T cell activation 3 B) Immune checkpoint signaling Response mmune activation Response cell chemokine sianalina CR 0.025 0.000 0.025 0.050 Tumor cell T in CR/PR 0.50 SD mmune ce Feature stratification quantile in SD/PD IMvigor210 **TCGA BLCA** IMvigor210 log2 TMB +1 IC Level T-test, p = 1.2e-10 ICI-PRS in Iraining Sets AUC = 0.805 **IMvigor test dataset** IMvigor test dataset LR p.val model LR p.val Model OS: IC OS ~ TMB 0.25 (base model) (base model) p < 0.0001 p = 0.31OS: IC + ICI-PRS OS ~ TMB + ICI-PRS 0.00714 0.00911 low vs. high split by median value PD-L1 IHC levels in peritumoral immune cells (IC) have been associated with clinical response and survival in ICI-treated patients (IMvigor210 test set shown in Figure 4A) IMvigor210 IMvigor210 **TCGA BLCA** T-test, p = 0.0029 in ets AUC = 0.710 ICI-PRS appears to improve both PD-L1 IHC and TMB predictive strength in ς Υ IMvigor210 training (data not shown) and testing data (Fig. 4B-C). LR = likelihood ratio. ICI-PR: esting PD-L1 and/or TMB may be complimentary to ICI-PRS use p = 0.0038low vs. high split by median value SUMMARY AND CONCLUSIONS CR=complete response; PR=partial response;

SD=stable disease; PD=progressive disease

A new ICI-PRS, incorporating sigi162 with select RNA features reflective of tumor cells and the immune microenvironment (Figure 2A), was trained in the IMvigor210 training set, where it was associated with response and survival for ICI-treated patients, but was still not associated with survival in non-ICI treated TCGA BLCA training (Figure 2C)

In the separate testing dataset, the ICI-PRS signature was significantly associated with response and survival for ICI-treated patients only (e.g., IMvigor210) but not in non-ICI treated patients (e.g., TCGA BLCA) (Figure 2D)

Figure 3: ICI-PRS Signature Status is Associated with Response and Survival in Independent Cohorts of ICI-treated BLCA Patients



When applied to independent cohorts of patients treated with ICI therapy (BACI (A) and BCAN (B)), ICI-PRS status is associated with both clinical response and overall survival

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• Herein we describe the development of a novel ICI-PRS and initial demonstration that it provides ICI-treatment specific prognostic value for bladder cancer patients

ICI-PRS status was associated with extended survival in patients treated with ICI but not those treated with non-ICI SOC therapy

These results support the further evaluation of the ICI-PRS and potential development as a diagnostic test for selecting UC patients most likely to benefit from ICI therapy

References

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