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Patient and Tumor-Specific Multiplexed Circulating Tumor DNA Response to Multi-Modality Therapy for Esophageal and Rectal Cancer: Preliminary Results of a Pilot Study

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Background

 Non-operative management of locally-advanced esophageal and rectal cancer is becoming more prevalent

 Highly sensitive and specific serum biomarkers of treatment response are needed – ctDNA monitoring shows great promise



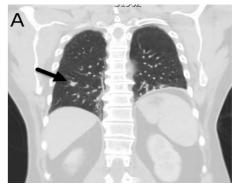
Design

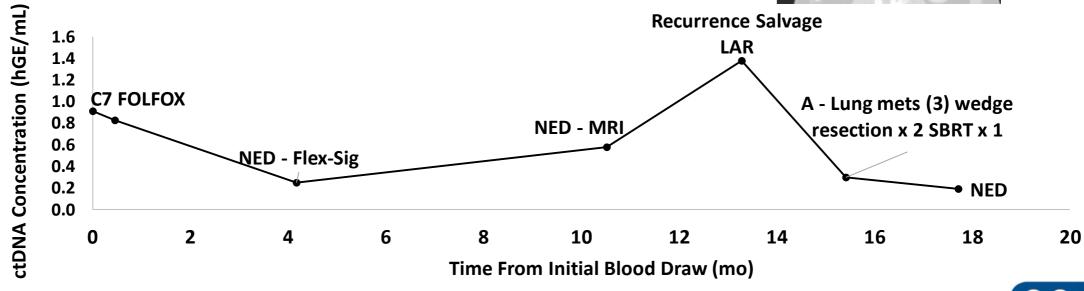
- Plasma specimens were collected at baseline, throughout neoadjuvant therapy, immediately prior to surgery and in follow up.
- Plasma samples underwent:
 - WES and library preparation
 - Somatic variants called using MuTect (1.1.4, Broad Institute)
 - SNV calls filtered to identify bait target sites
 - DIDA-Seq error-correction libraries prepared as per Butler et al., 2019



Persistent ctDNA precedes clinical recurrence with 8 month lead-time

cT2N1M0 rectal adenocarcinoma

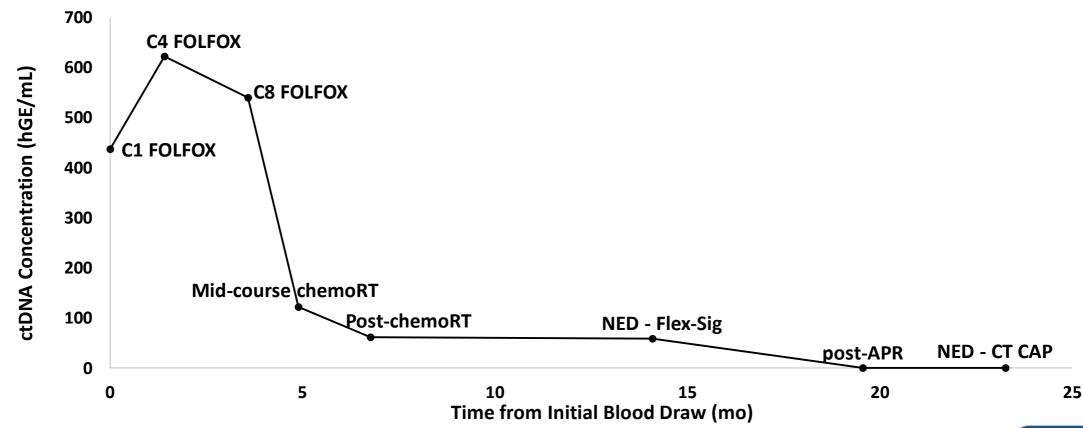






Persistent ctDNA precedes clinical recurrence with 5 month lead-time

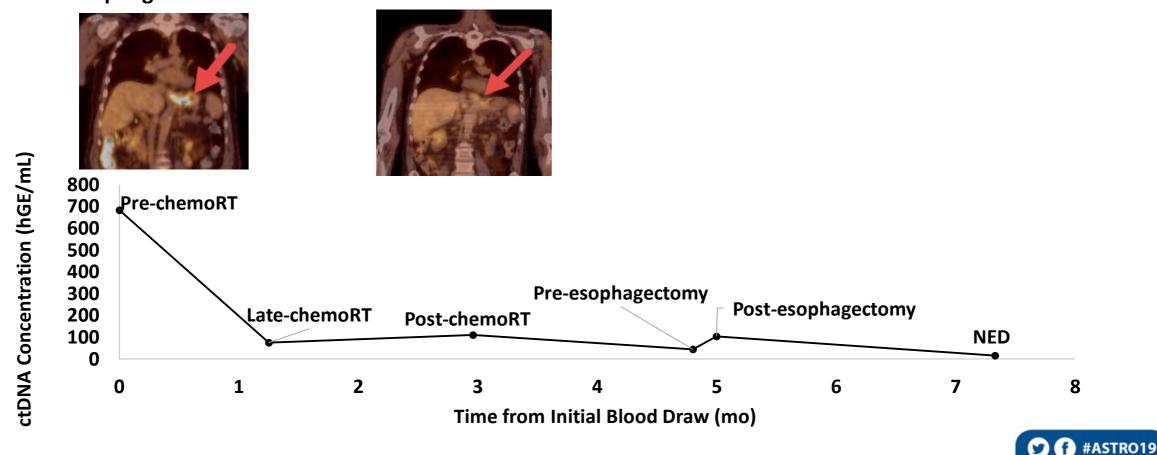






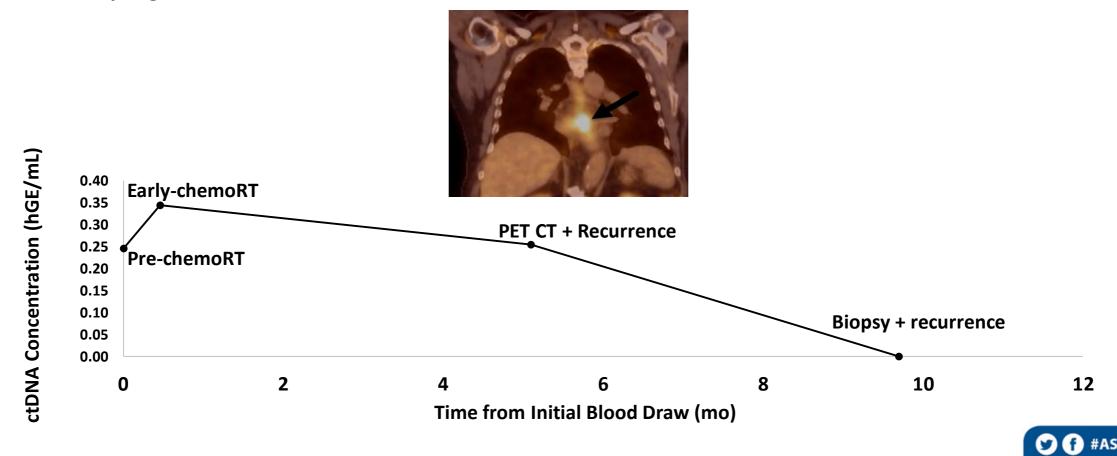
ctDNA declines with neoadjuvant chemoRT and after resection

cT3N1 esophageal adenocarcinoma



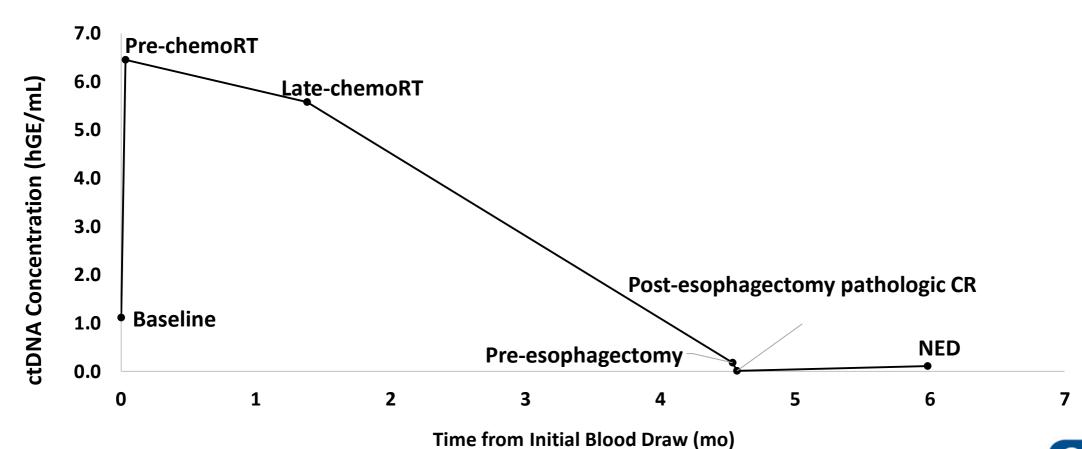
ctDNA undetectable at time of recurrence

cT3Nx esophageal adenocarcinoma



Absence of ctDNA associated with pCR

cT2N0 esophageal adenocarcinoma





Conclusions

- CtDNA associated with treatment response, confirming pCR and predicts local recurrences with significant clinical lead times.
- Tumor heterogeneity and high tumor mutational burden may reduce sensitivity.
- High sensitivity ctDNA detection could initiate early salvage surgery in non-operative patients and improve early detection of locally progressive or metastatic disease.



Questions

