### New Platforms and Developments of Immunotherapy Biomarkers Beyond PD-L1 in Tissue

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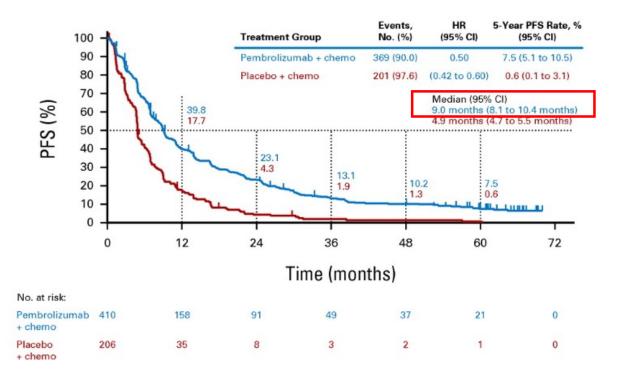


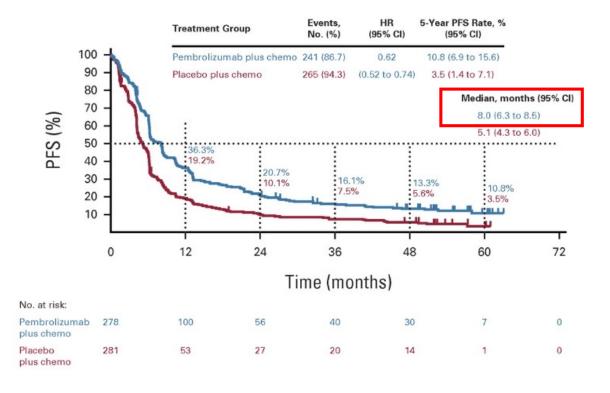


## Approximately 50% of patients with advanced non-small cell lung cancer will develop disease progression within 8 to 9 months

### **KEYNOTE-189**

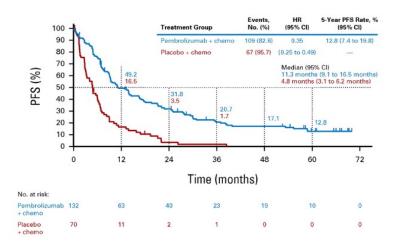
#### **KEYNOTE-407**



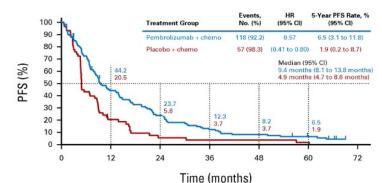


### How do we risk stratify these patients?

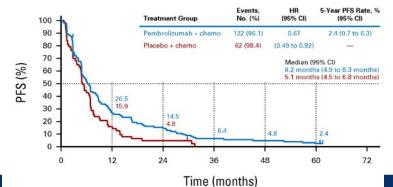
### **KEYNOTE-189**





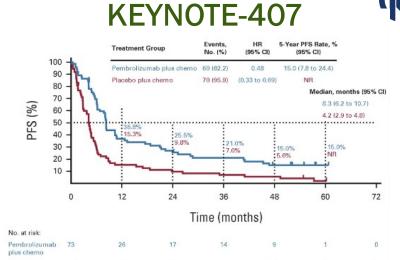


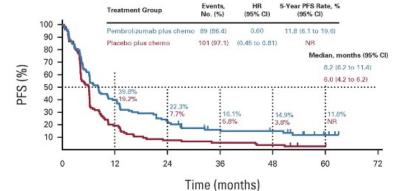




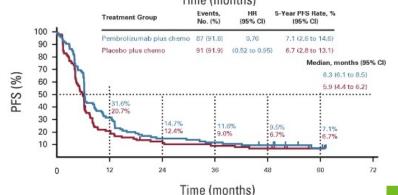


Garassino M. J Clin Oncol. 2023 Novello S. J Clin Oncol. 2023





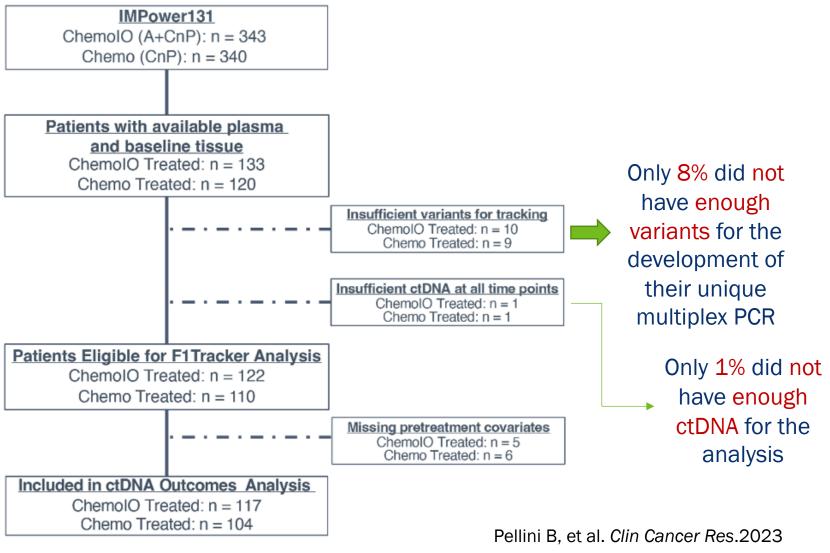
plus chemo



# Can ctDNA monitoring risk stratify advanced SCC prior to the start of maintenance atezolizumab in patients receiving carboplatin + nab-paclitaxel + atezolizumab (IMpower 131)?



Tumor-informed assay (FoundationOne Tracker)

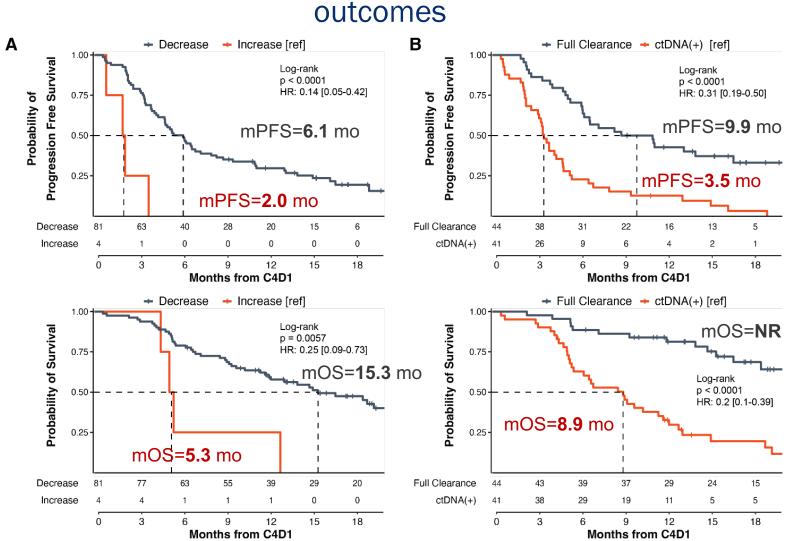


### ctDNA Monitoring on Atezolizumab + Carboplatin + Nab-Paclitaxel (A +CnP) identifies patients with SCC with higher risk for poorer



Analysis limited to the 85 patients with detectable ctDNA pretreatment

Decrease is defined as <u>any</u> decrease



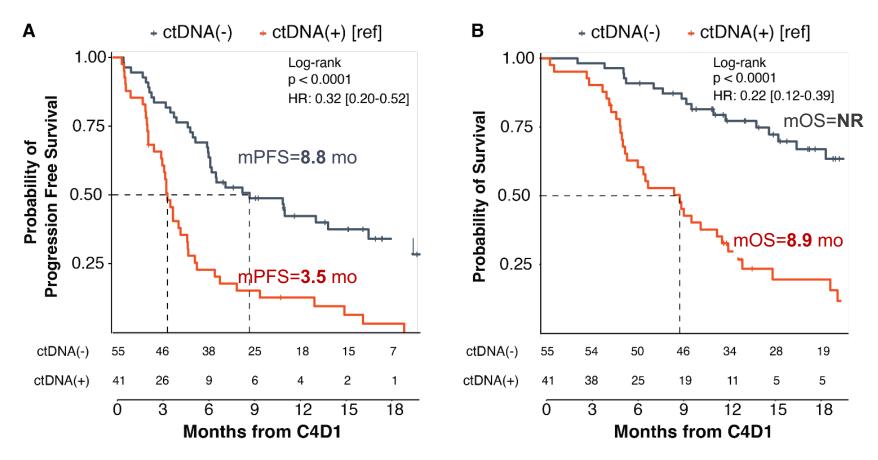
PFS and OS are landmarked from C4D1 (i.e., 9 weeks into induction therapy)

## ctDNA detection at C4D1 on Atezolizumab + Carboplatin + Nab-Paclitaxel (A +CnP) can risk stratify patients with SCC before maintenance therapy start



#### ctDNA + at C4D1

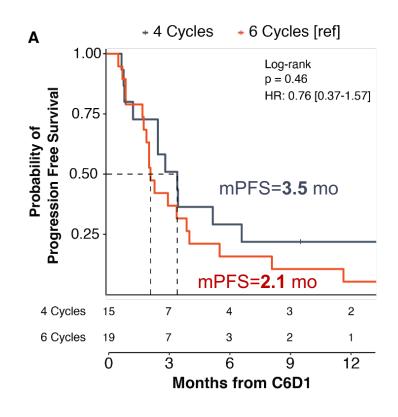
PFS and OS are landmarked from C4D1 (i.e., 9 weeks into induction therapy)

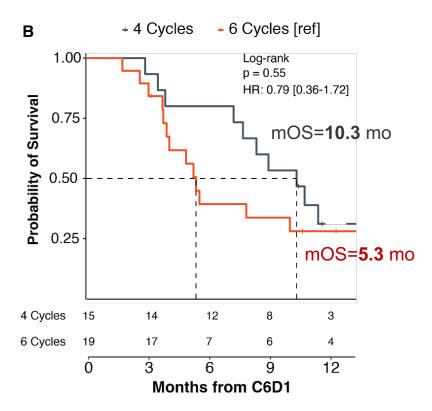


Analysis in 96 patients with available samples at C4D1 regardless of baseline sample availability

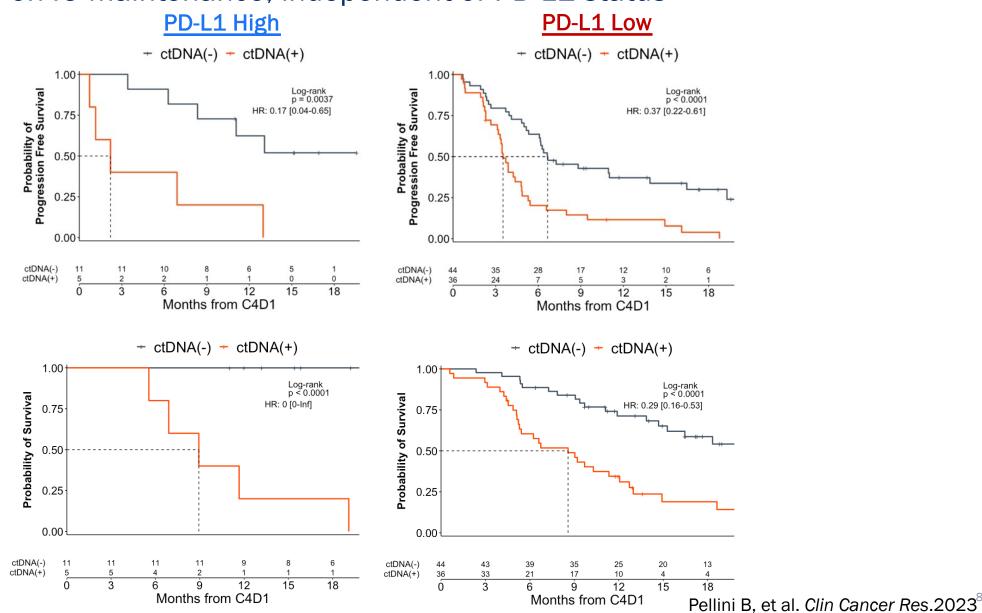
# Additional cycles of induction Atezolizumab + Carboplatin + Nab-Paclitaxel (A +CnP) are not associated with improved outcomes in patients with ctDNA + at C4D1





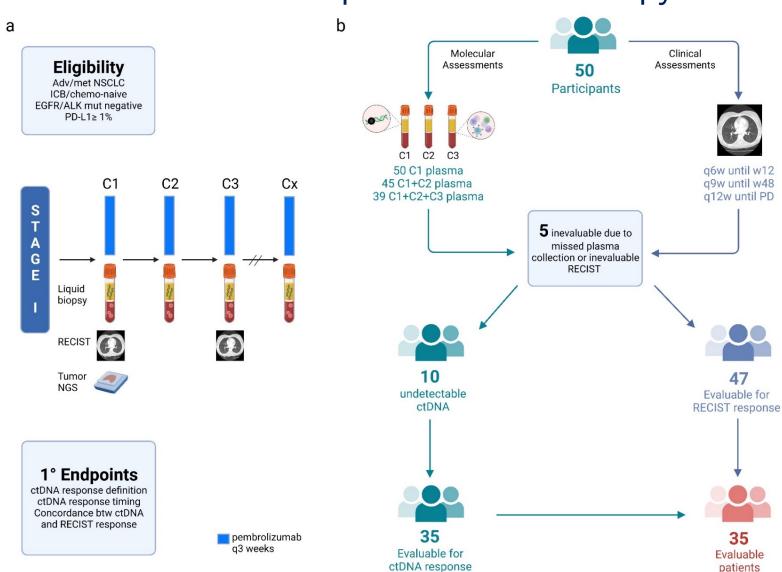


Full clearance of ctDNA at C4D1 on chemolO identifies patients with poorer outcomes on IO maintenance, independent of PD-L1 status



### BR.36: A ctDNA-directed phase II study of molecular response adaptive immunotherapy in NSCLC



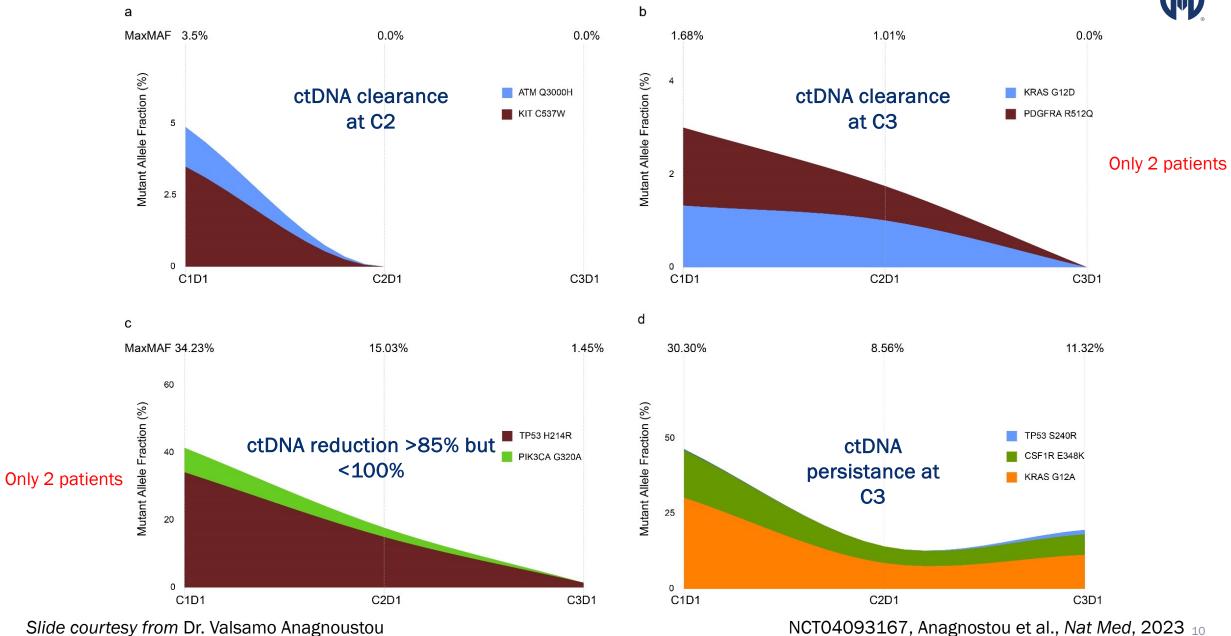


Tumor-uninformed assay (PGDx elio™ plasma resolve)

- NGS with 33-gene panel
  - **WBC-informed** 
    - TAT 1 week

20% not evaluable for molecular response due to ctDNA (-) at baseline

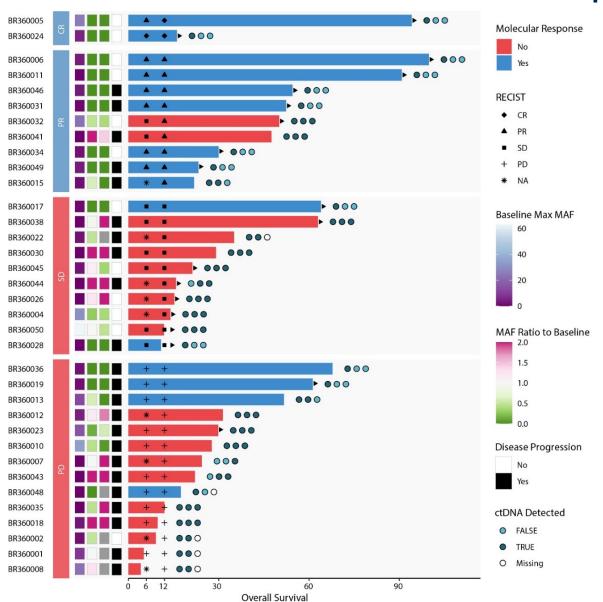
### Patterns of ctDNA kinetics



NCTO4093167, Anagnostou et al., Nat Med, 2023 10

### ctDNA-RECIST response concordance





BR.36 stage 1 met its primary endpoint.

The sensitivity of molecular response for RECIST best overall response was 82%, (90% CI: 52% - 97%), specificity was 75% (90% CI: 56.5% - 88.5%).

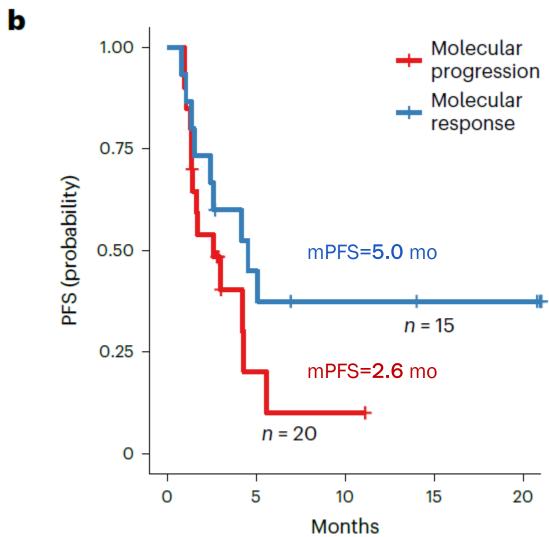
RECIST response (BOR)	Molecular response	
	mR	mPD
CR/PR	9 (82%)	2 (18%)
No RECIST response	6 (25%)	18 (75%)
iRECIST response	-	
iCR/iPR	10 (83%)	2 (17%)
No iRECIST response	5 (22%)	18 (78%)

(Weeks)

## ctDNA molecular response risk stratifies patients with mNSCLC with





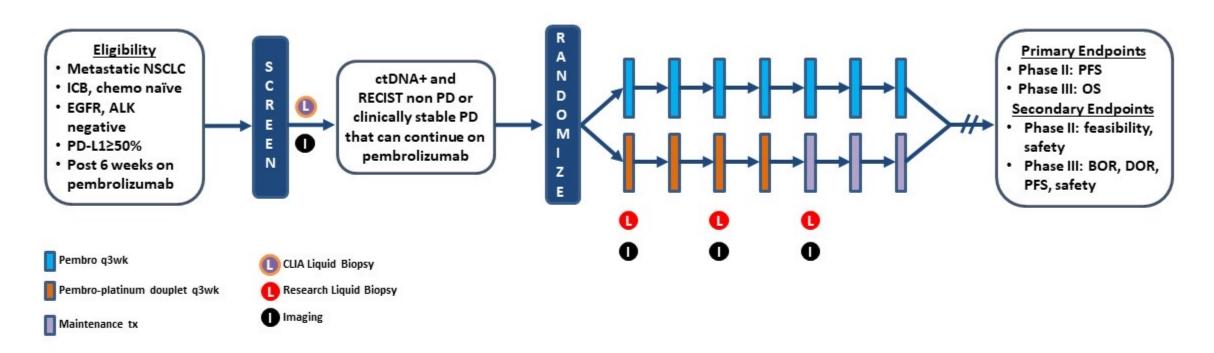


C 1.00 Molecular progression Molecular response 0.75 OS (probability) mOS= NR 0.50 n = 15mOS=7.2 mo 0.25 n = 200 5 15 20 25 10 Months

Median time to ctDNA response was 2.1 months



BR.36 stage 2-A Biomarker-Directed, Open Label, Multi-Center Phase II/III Study of Molecular Response Adaptive Immuno-Chemotherapy in Patients with NSCLC



### Conclusions



- CGP-informed ctDNA monitoring on chemolO in advanced NSCLC can inform durability of treatment benefit
- ctDNA detection during induction chemolo may offer an opportunity to identify patients at high-risk for disease progression and inform selection of novel personalized maintenance treatment strategies
- Tumor-uninformed (agnostic) ctDNA molecular response at 6-8 weeks into IO monotherapy in advanced PD-L1(+) NSCLC is can identify patients with improved clinical outcomes
- Treatment intensification with the addition of chemotherapy based on ctDNA MR for patients mNSCLC with PD-L1≥ 50% with is under investigation (NCT04093167)

### Questions?





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