

## **SCLC BIOMARKERS**

### Carl M. Gay, M.D., Ph.D. University of Texas MD Anderson Cancer Center

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## A comprehensive list of SCLC biomarkers recommended by international guidelines







#### Horn et al, NEJM 2018



Speaker: Carl M. Gay, M.D., Ph.D. MDACC

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### **IMpower133**: Phase 3 – EP + atezolizumab with

#### atezolizumab maintenance in ED- SCLC

#### (NCT02763579)

(100102100013)				Atezolizumab	Placebo			
	Carboplatin + Etoposide + Atezolizumab	Sex						
		Male	261 (65)	12.3	10.9		1	0.74 (0.54–1.02)
-	Carbo (AUC 5) D1+ Etop 100mg/m2 D1-3	Female	142 (35)	12.5	9.5	·		0.65 (0.42–1.00)
	4 Atore 1200mg D1 c2W x 4 evelop	Age	217 (54)	10.1	11.5		1	0.02 (0.64, 1.22)
	+ Alezo 1200mg DT q3W X 4 cycles	   	217 (54)	12.1	11.5		_	0.92(0.64 - 1.32)
		ECOC score	100 (40)	12.5	9.0		1	0.55 (0.56-0.77)
	Followed by	0	140 (35)	16.6	12.4		_	0.79 (0.49-1.27)
Histologically documented	Atezo 1200mg D1 g3W until PD or	1	263 (65)	11.4	9.3		1	0.68 (0.50-0.93)
ED-SCI C	unacceptable toxicity	Brain metastases						(,
LD-00E0		Yes	35 (9)	8.5	9.7		•	→ 1.07 (0.47–2.43)
No prior treatment     R		No	368 (91)	12.6	10.4	<b>-</b>	i	0.68 (0.52–0.89)
• ECOG PS 0 – 1		Liver metastases	1 (0 (27)	0.2	7.0		(	0.01 (0.55, 1.00)
No autoimmune disease		Yes	149 (37)	9.3	/.8		-	0.81 (0.55 - 1.20)
	Carboniatin + Etoposide + Placebo	Tumor mutational burde	204 (00) m	10.0				0.84 (0.43=0.90)
• No active CNS disease		<10 mutations/Mb	139 (34)	11.8	9.2	<b>_</b>	4	0.70 (0.45-1.07)
		≥10 mutations/Mb	212 (53)	14.6	11.2		i	0.68 (0.47-0.97)
	Carbo (AUC 5) D1+ Etop 100mg/m2 D1-3	<16 mutations/Mb	271 (67)	12.5	9.9			0.71 (0.52–0.98)
	+ Placebo q3W x 4 cycles	≥16 mutations/Mb	80 (20)	17.8	11.9	·	-	0.63 (0.35-1.15)
Co-primary endpoints: OS, PFS		Intention-to-treat	403 (100)	12.3	10.3			0.70 (0.54–0.91)
	Followed by	population			0,1	1	0	2.5
Secondary endpoints: ORR, DOR,	Placebo D1 d3W until PD or unacceptable				4			
6-month PFS, QoL	toxicity				Ate	zolizumab Better	Placebo Better	
(N=500)	toxicity							
(14-500)								

Subgroup

C Overall Survival According to Baseline Characteristics

#### Higher TMB did not predict benefit with addition of ICI in IMpower133

No. of Patients (%) Median Overall Survival (mo) Hazard Ratio for Death (95% CI)

#### Horn et al, NEJM 2018





## **CASPIAN**: Phase 3 – EP + durvalumab with durvalumab maintenance in ES-SCLC



Durvalumab plus platinum–etoposide versus platinum–etoposide in first-line treatment of extensive-stage small-cell lung cancer (CASPIAN): a randomised, controlled, open-label, phase 3 trial

Luis Paz-Ares, Mikhail Dvorkin, Yuanbin Chen, Niels Reinmuth, Katsuyuki Hotta, Dmytro Trukhin, Galina Statsenko, Maximilian J Hochmair, Mustafa Özgüroğlu, Jun Ho Ji, Oleksandr Voitko, Artem Poltoratskiy, Santiago Ponce, Francesco Verderame, Libor Havel, Igor Bondarenko, Andrzej Kazarnowicz, György Losonczy, Nikolay V Conev, Jon Armstrong, Natalie Byrne, Norah Shire, Haiyi Jiang, Jonathan W Goldman, for the CASPIAN investigators\*



#### Paz-Ares et al, Lancet 2019





## **CASPIAN**: Phase 3 – EP + durvalumab with durvalumab maintenance in ES-SCLC



#### ESMO 2019 update:

- 277 with evaluable samples with Ventana PD-L1 (SP263)
- 5% and 22% of pts with expression ≥1% in tumor and immune cells, respectively
- PD-L1 expression as continuous variable had no impact on OS, PFS, or ORR

Similarly, PD-L1 expression was infrequent and had no bearing on outcomes with addition of ICI in CASPIAN.

Paz-Ares et al, Lancet 2019; Paz-Ares et al. ESMO 2019



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## **Personalization in SCLC may look different**

### ARTICLE

doi:10.1038

## Comprehensive genomic profiles of small cell lung cancer



Ubiquitous bi-allelic loss of tumor suppressor genes *TP53* and *RB1*Absence of actionable oncogenic drivers

#### George et al, Nature 2015



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### **Transcriptional diversity underlies immune** phenotypic diversity Antigen presentation IFNy signature

subtype

Cibersortx absolute abundance

1.0

0.5

0.0

NS







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subtype

HLA-A

HLA-B

B2M

HLA-C

HLA-DPA1

HLA-DPB1



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CXCL9

PSMB10

IDO1

LAG3

TIGIT



# Transcriptional subtype predicts differential benefit of the addition of immunotherapy (IMP133)





Gay et al. Cancer Cell, 2021



# Transcriptional subtype predicts differential benefit of the addition of immunotherapy (CASPIAN)



Gay et al. Cancer Cell, 2021





## What about the un-inflamed subtypes?

Could we make them all inflamed? Maybe...



LSD1 inhibition directs neuroendocrine subtypes (e.g. SCLC-A or –N) toward more inflamed states.

#### Hiatt et al. Clin Cancer Res, 2022





## What about the un-inflamed subtypes?

Could we make them all inflamed? Maybe...







# If subtype, or any biomarker, is needed in SCLC, how might we do that practically?

- Perform biomarker analysis in parallel with treatment.
- Plasma-based approaches.





## If subtype, or any biomarker, is needed in SCLC, how might we do that practically?

• Perform biomarker analysis in parallel with treatment.

<u>S1929</u>: Phase II Study of Maintenance Atezolizumab Versus Atezolizumab in Combination with Talazoparib in Patients with SLFN11 Positive Extensive Stage Small Cell Lung Cancer (ES-SCLC)



SWOG1929 successfully integrated an IHC for SLFN11 – a predictive biomarker for PARP inhibitor sensitivity in SCLC – into frontline maintenance by performing assay *during* initial therapy.







## If subtype, or any biomarker, is needed in SCLC, how might we do that practically?

• Plasma-based approaches.

- Transcriptional subtype, as well as other expression-based features, are epigenetically encoded, including in ctDNA, and can be rapidly and longitudinally collected.
- Longitudinal collections highlight the limitations of archival tissue analysis.



#### Heeke et al. Cancer Cell 2024



## **SCLC Biomarkers: Summary**

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- Genomic biomarkers are out, for now.
- Even established expression-based biomarkers (e.g. PD-L1) are not predictive.
- SCLC transcriptional subtypes delineate tumor diversity.
  - SCLC-I patients experience greatest benefit from ICI.
  - Other subtypes may benefit from intensification or novel immune-based strategies (e.g. BiTEs).
- Aggressive natural history requires creative strategies to assess biomarkers without delaying care.
- Transcriptional biomarkers may evolve with treatment assess longitudinally?
- Should surface protein targets be integrated into biomarker plans?
  - DLL3 vs. B7H3 vs. SEZ6 vs. TROP2, etc.

