

# **New Therapies for Squamous Cell and Small Cell Lung Carcinomas**

**Alberto Chiappori, MD**

**Senior Member**

**Department of Thoracic Oncology**

**Moffitt Cancer Center**

**11th Annual Winter Cancer Symposium**

**La Concha Renaissance Hotel**

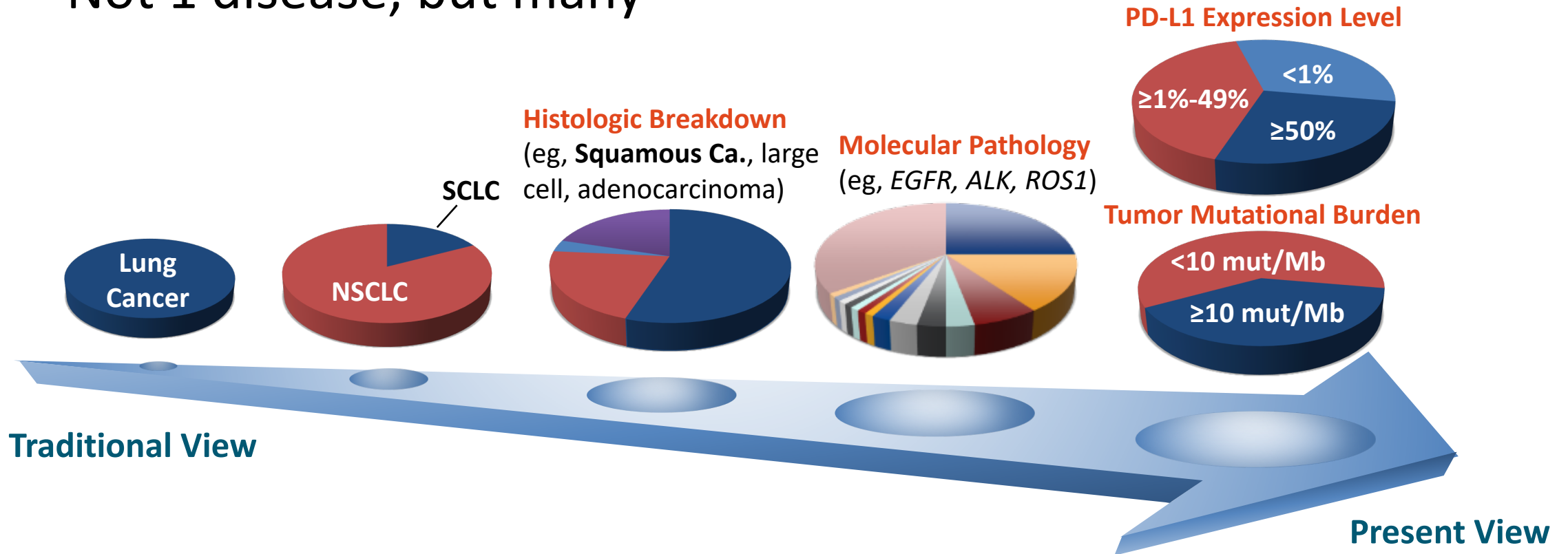
**San Juan, Puerto Rico**

**March 4-6, 2022**

# Evolution of Therapy in Lung Cancer

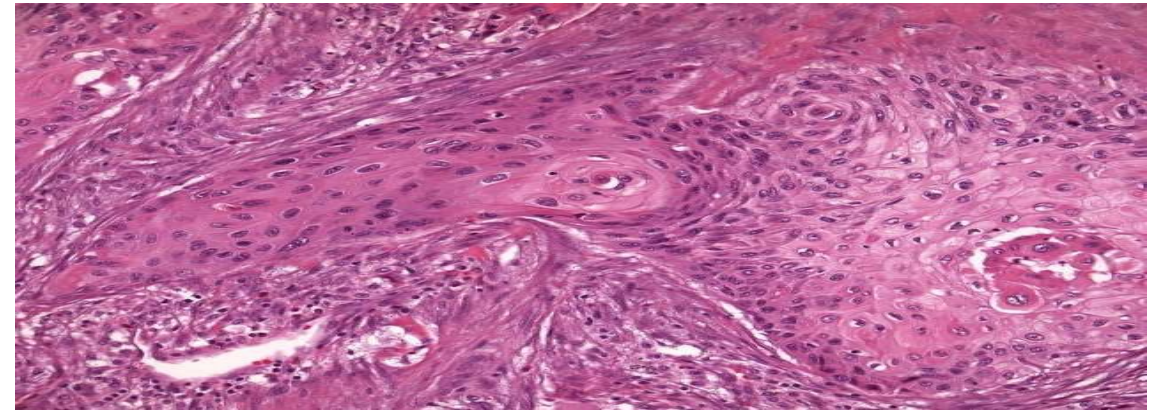
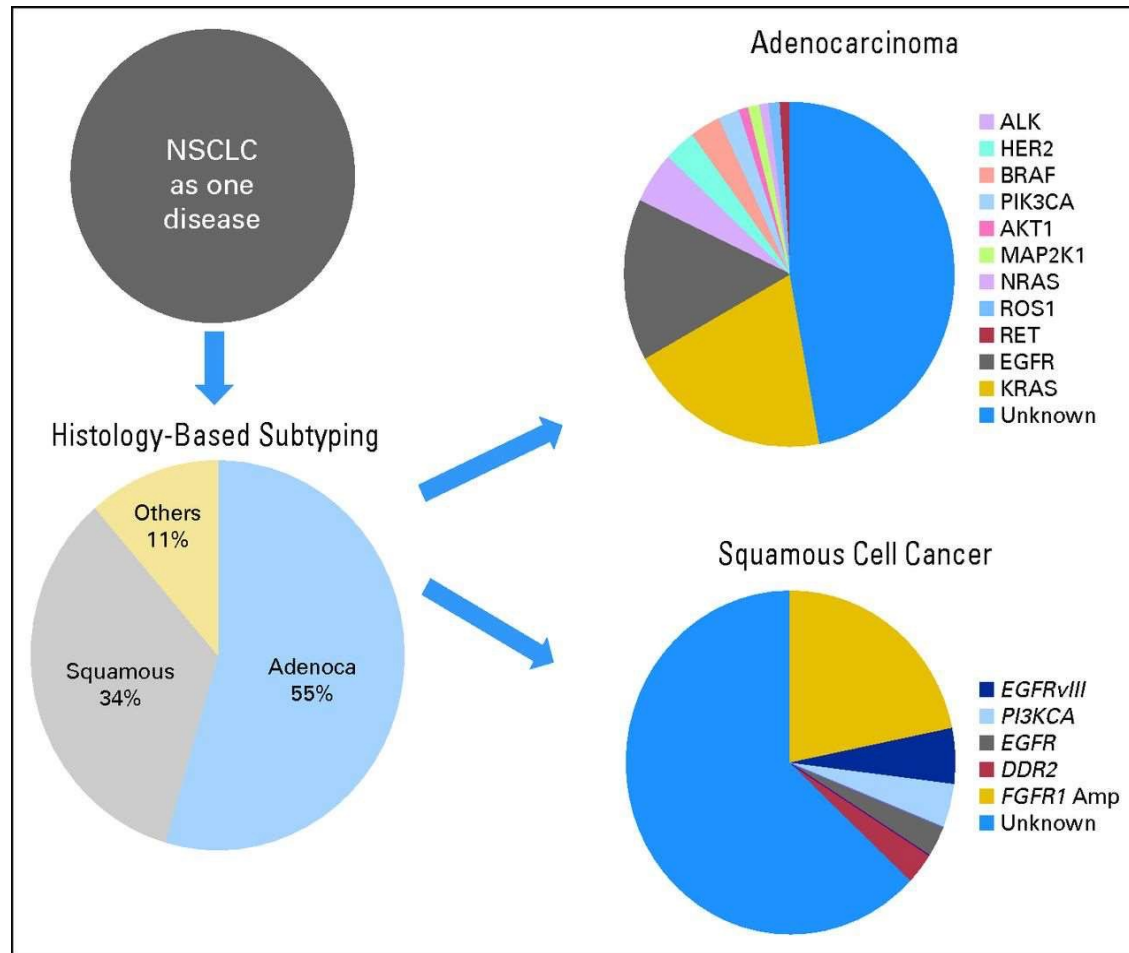
## NSCLC – Adenocarcinoma vs. Squamous Carcinoma

- Not 1 disease, but many



# Evolution of Therapy in Lung Cancer

## Squamous Carcinoma – Molecular Targets

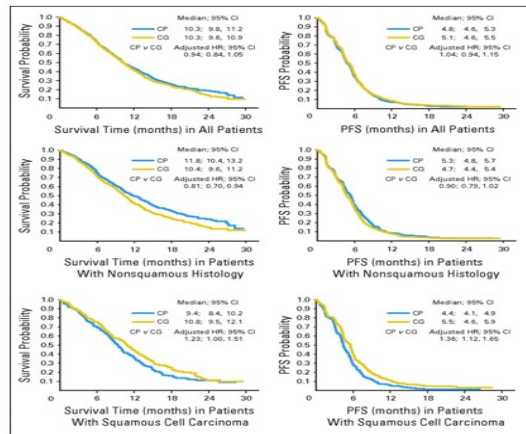


- 20-30% of NSCLC
- Strongly associated with cigarette smoking
- Historically was treated like lung adenocarcinomas
- Despite advances in the personalized treatment of adenocarcinoma, effective targeted therapy for squamous cell has remained elusive
- Squamous cell lung cancer lacks druggable targets
- Most substantial impact in treatment has come from histology agnostic approaches

# Squamous Cell Carcinoma

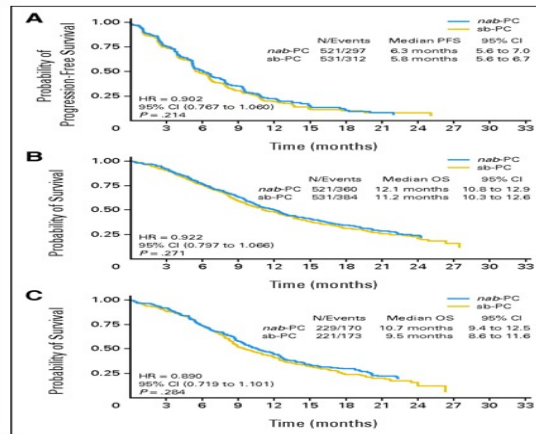
## Treatment Evolution: Chemotherapy, moAbs, TKIs

### Chemotherapy: Pemetrexed and nab-Paclitaxel



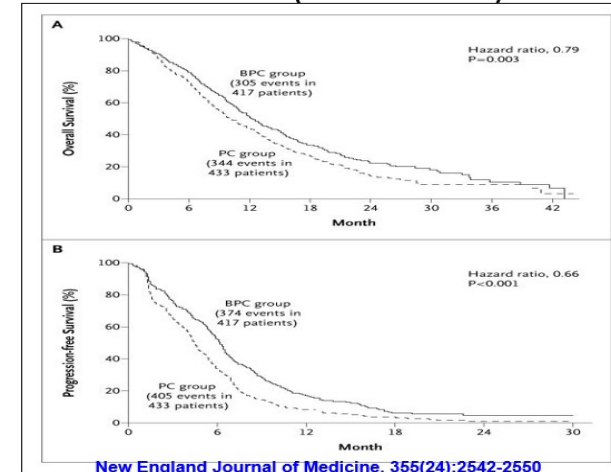
Giorgio Vittorio Scagliotti et al. JCO 2008;26:3543-3551

### Squamous



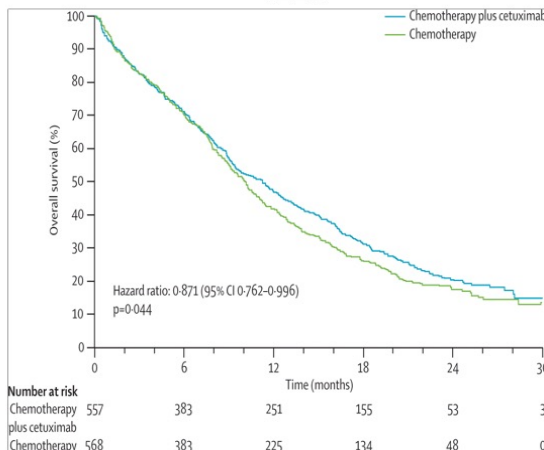
Mark A. Socinski et al. JCO 2012;30:2055-2062

### Biologics (anti-VEGFR): Bevacizumab and Ramucirumab



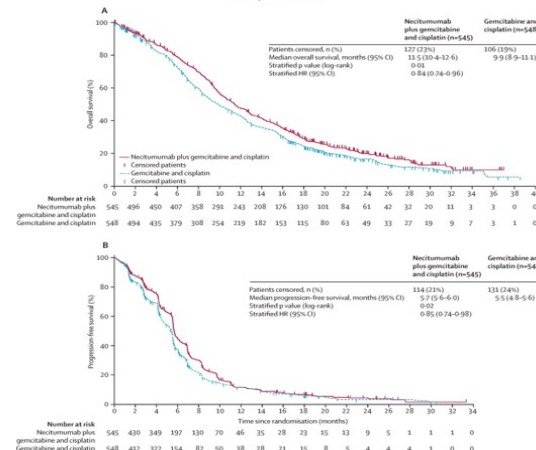
New England Journal of Medicine. 355(24):2542-2550

### Biologics (anti-EGFR): Cetuximab and Necitumumab



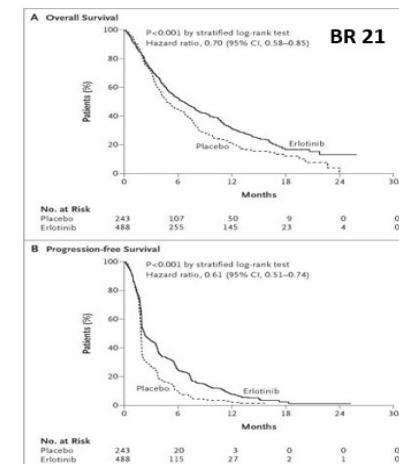
Lancet 2009; 373: 1525-1531

### SQUIRE

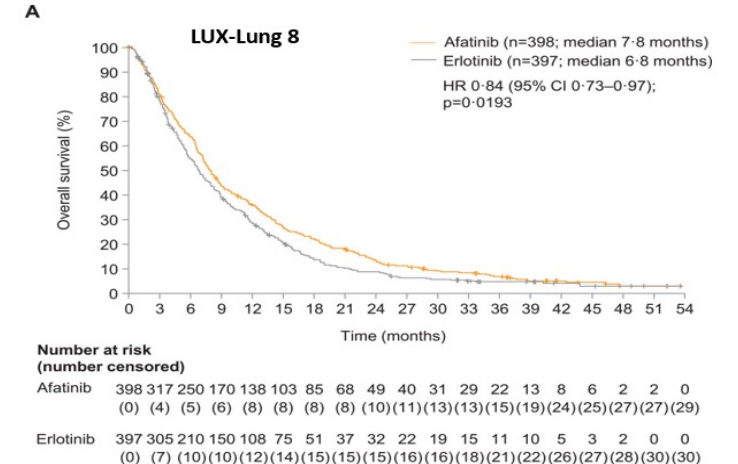


Lancet Oncology 2015, 16(7): 763-774

### TKI: Erlotinib and Afatinib



N Engl J Med 2005 Jul 14;353(2):123-32

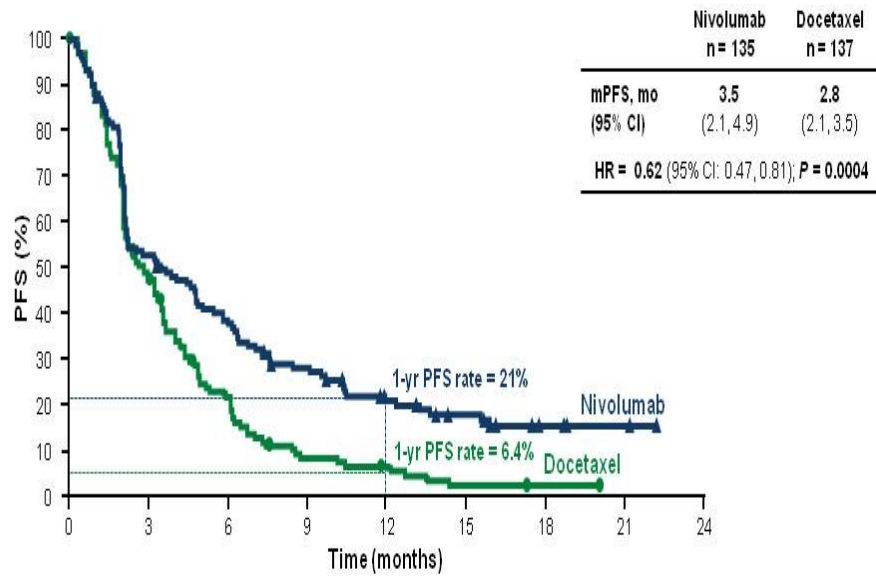


EClinicalMedicine. 2021 Jul; 37: 100940.

# Nivolumab

## Phase III Squamous NSCLC – CM-017

### Progression-Free Survival



#### Number of Patients at Risk

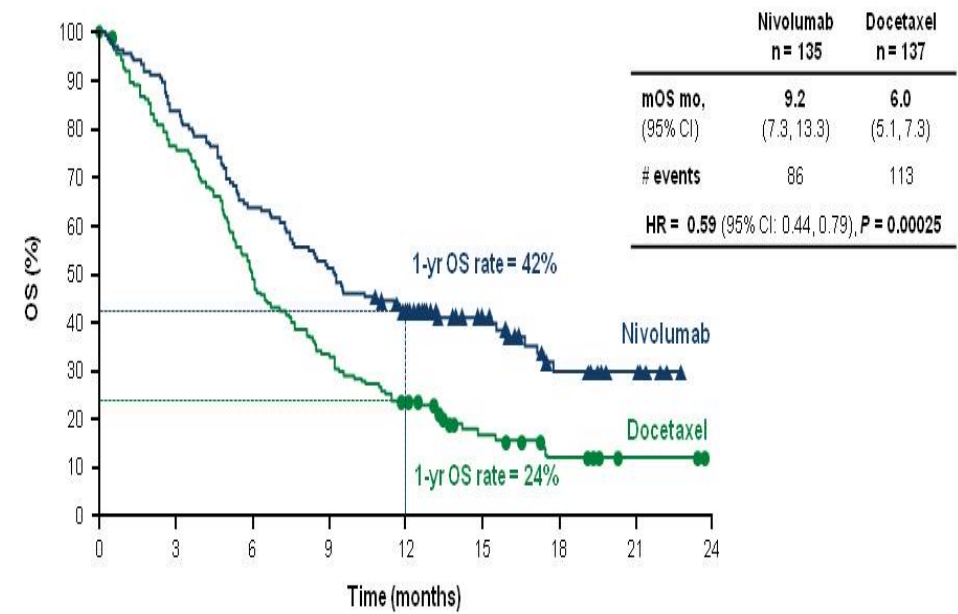
	0	3	6	9	12	15	18	21	24
Nivolumab	135	68	48	33	21	15	6	2	0
Docetaxel	137	62	26	9	6	2	1	0	0

PFS per investigator.

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PRESENTED AT: ASCO Annual '15 Meeting

### Overall Survival



#### Number of Patients at Risk

	0	3	6	9	12	15	18	21	24
Nivolumab	135	113	86	69	52	31	15	7	0
Docetaxel	137	103	68	45	30	14	7	2	0

Symbols represent censored observations

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PRESENTED AT: ASCO Annual '15 Meeting

# Single-Agent ICI for Advanced NSCLC

## High PD-L1 Expression

### KEYNOTE-024

#### Open-label, randomized phase III study

Stratified by ECOG PS (0 vs 1), histology (squamous vs nonsquamous), and enrollment site

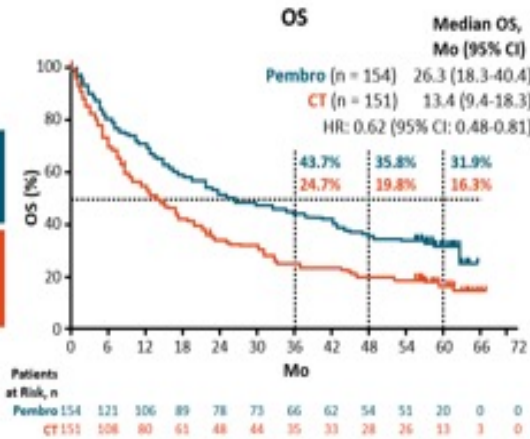
Chemotherapy-naïve, stage IV NSCLC; PD-L1 selected (TPS ≥50%\*); ECOG PS 0/1; no EGFR/ALK mutations; no untreated CNS mets or autoimmune disease requiring tx (N = 305)

**Pembrolizumab 200 mg IV Q3W for up to 35 cycles (n = 154)**

**Platinum-Doublet CT (histology based) for 4-6 cycles (n = 151)**

\*Staining of ≥50% tumor cells by 22C3 companion diagnostic IHC assay.

Reck. JCO. 2021;21:2339.



### IMpower110

#### Randomized phase III study

Stratified by sex, ECOG PS, histology, and tumor PD-L1 status

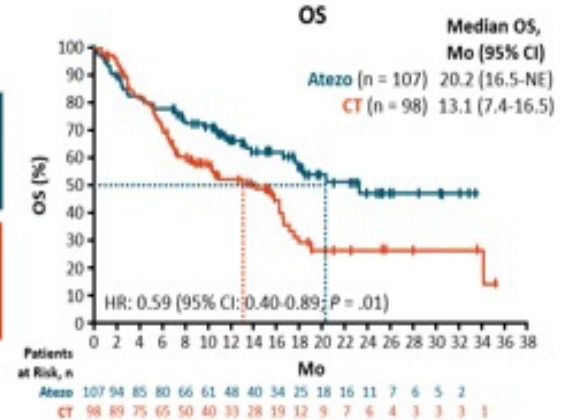
Untreated LA or metastatic NSCLC of any histology; PD-L1 ≥50% on TC or 10% on IC\*;  
no sensitizing EGFR or ALK alterations; ECOG PS 0/1; no untreated or unstable CNS mets or pneumonitis requiring tx (N = 205\*)

**Atezolizumab 1200 mg Q3W until PD (n = 107)**

**Platinum-Doublet CT (histology based) for 4-6 cycles (n = 98)**

\*Staining of ≥50% tumor cells (TC) or ≥10% tumor-infiltrating immune cells (IC) using SP142 complementary diagnostic IHC assay.  
\*572 patients with PD-L1 ≥1% on TC or IC enrolled overall.

Herbst. NEJM. 2020;383:1328.



### EMPOWER-Lung 1

#### Randomized phase III study

Stratified by histology (squamous vs nonsquamous) and region (Europe, Asia, vs ROW)

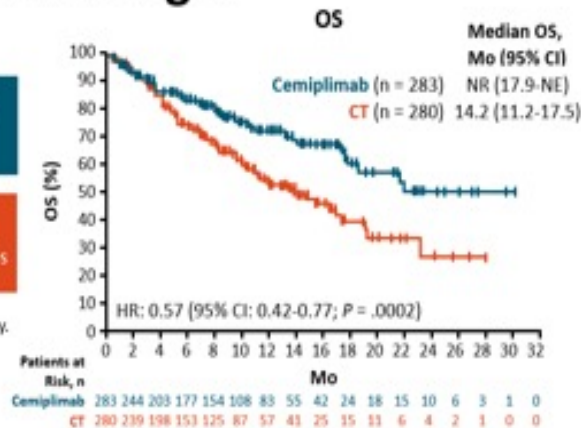
Treatment naïve advanced NSCLC, PD-L1 ≥50%\*;  
no EGFR/ALK/ROS1 mutations, ECOG PS 0/1 (N = 563\*)

**Cemiplimab 350 mg Q3W Until PD or 108 wk**

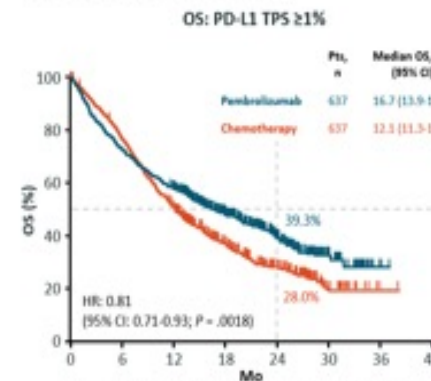
**Platinum-Doublet CT (histology based) for 4-6 cycles**

\*Staining of ≥50% tumor cells using 22C3 companion diagnostic IHC assay.  
\*710 patients enrolled in overall ITT population.

Sezer. Lancet. 2021;397:592.

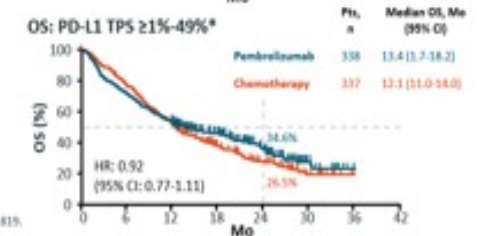
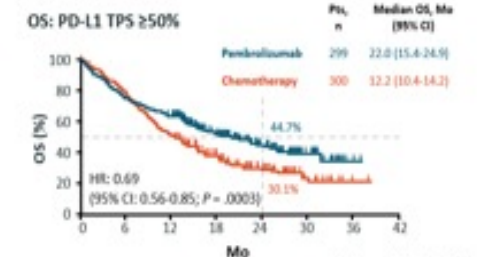


### KEYNOTE-042



\*Exploratory analysis; no alpha allocated to this comparison.

McK. Lancet. 2019;393:815.



# Chemo-IO Combinations for Advanced NSCLC

## Histology based

Trial	Comparison	Selection	ORR, %	PFS HR	OS HR
KEYNOTE-189 <sup>1,2</sup>	Pembro or placebo + carbo/pem	PD-L1 unselected; nonsq	48.3 vs 19.9	0.49	0.56
IMpower130 <sup>3</sup>	Atezo + carbo/nab-pac vs CT alone	PD-L1 unselected; nonsq	49.2 vs 31.9	0.64	0.79
IMpower150 <sup>4,5</sup>	Atezo + carbo/pac + bev vs CT + bev	PD-L1 unselected; nonsq*	63.5 vs 48.0	0.62	0.80
KEYNOTE-407 <sup>6-8</sup>	Pembro or placebo + carbo/pac or nab-pac	PD-L1 unselected; sq	62.6 vs 38.8	0.59	0.71
IMpower131 <sup>9</sup>	Atezo + carbo/nab-pac vs CT alone	PD-L1 unselected; sq		0.71	0.96

\*WT population (excludes patients with *EGFR* or *ALK* alterations).

### ■ FDA approvals

- No *EGFR* or *ALK* alterations
- PD-L1 agnostic; OS benefit observed in all subgroups

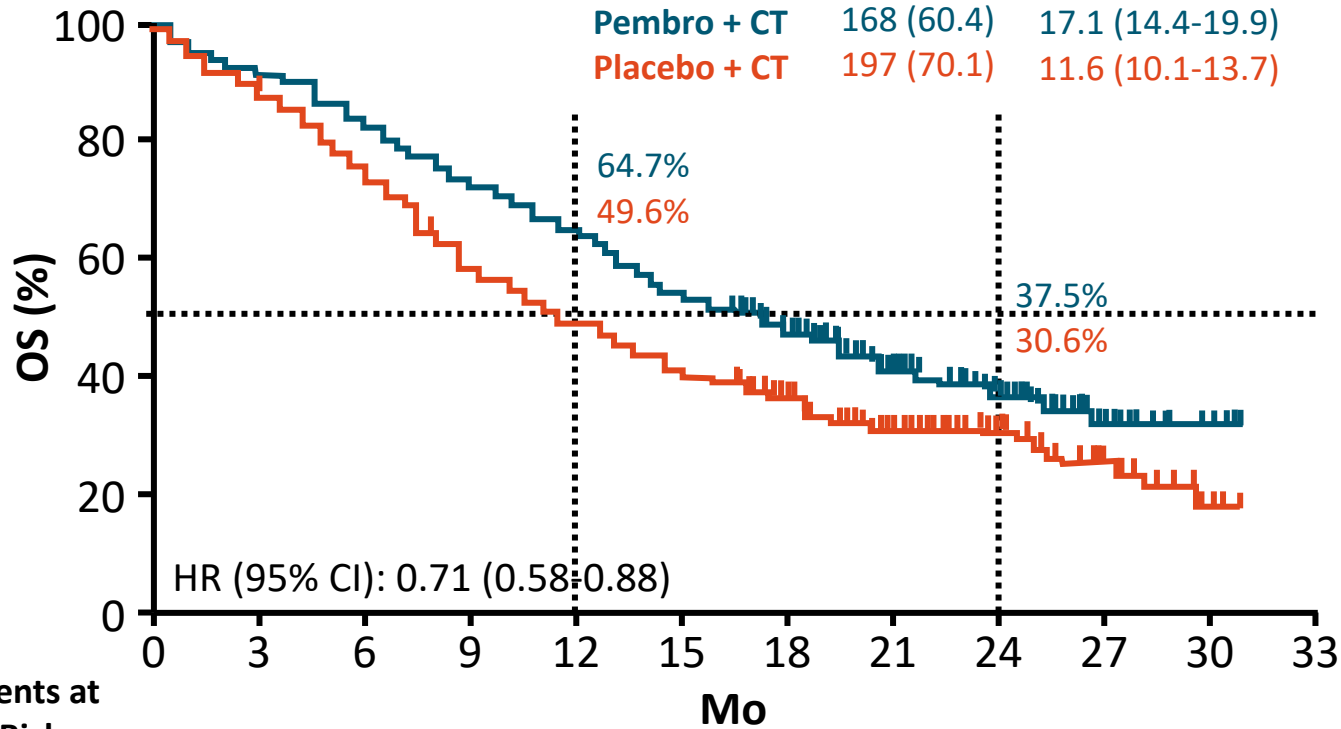
1. Gandhi. NEJM. 2018;378:2078. 2. Rodríguez-Abreu. Ann Oncol. 2021;32:881. 3. West. Lancet Oncol. 2019;20:924. 4. Socinski. NEJM. 2018;378:2288. 5. Socinski. AACR 2020. Abstr CT216. 6. Paz-Ares. NEJM. 2018;379:2040. 7. Paz-Ares. J Thorac Oncol. 2020;15:1657. 8. Robinson. ELCC 2021. Abstr 970. 9. Presented By Robert Jotte at 2018 ASCO Annual Meeting

# KEYNOTE-407:

## Pembrolizumab + Chemotherapy

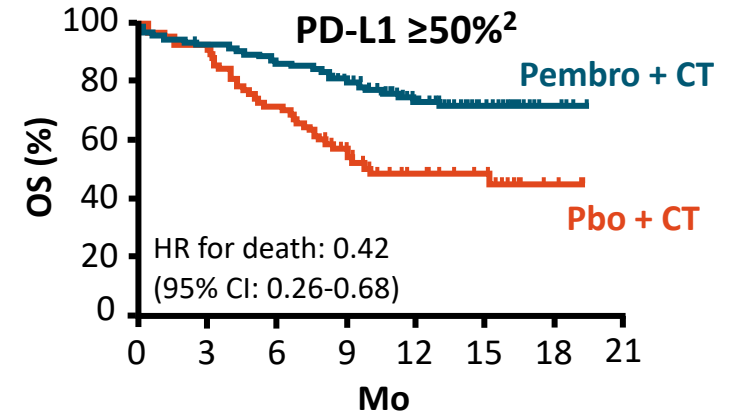
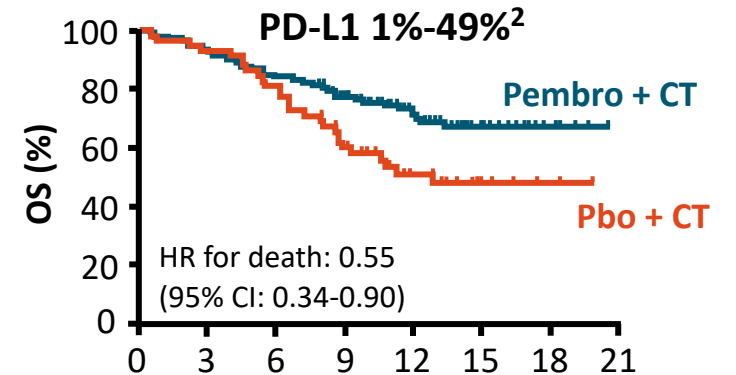
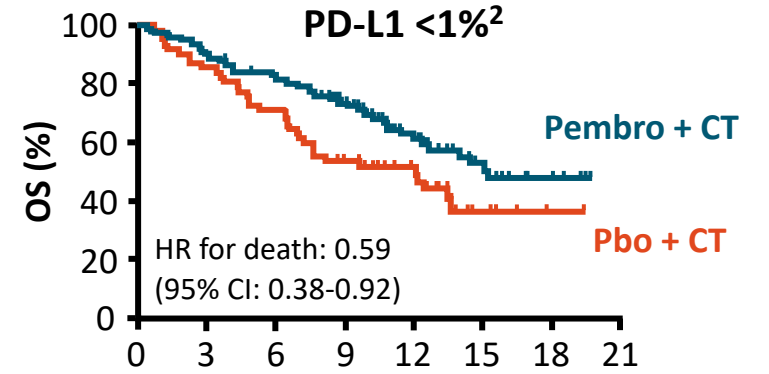
### OS in ITT Population<sup>1</sup>

	Events, n (%)	Median OS, Mo (95% CI)
<b>Pembro + CT</b>	168 (60.4)	17.1 (14.4-19.9)
<b>Placebo + CT</b>	197 (70.1)	11.6 (10.1-13.7)



Patients at Risk, n

<b>Pembro + CT</b>	278	256	232	203	180	150	119	80	46	14	4	0
<b>Pbo + CT</b>	281	245	210	163	137	113	91	61	36	16	3	0

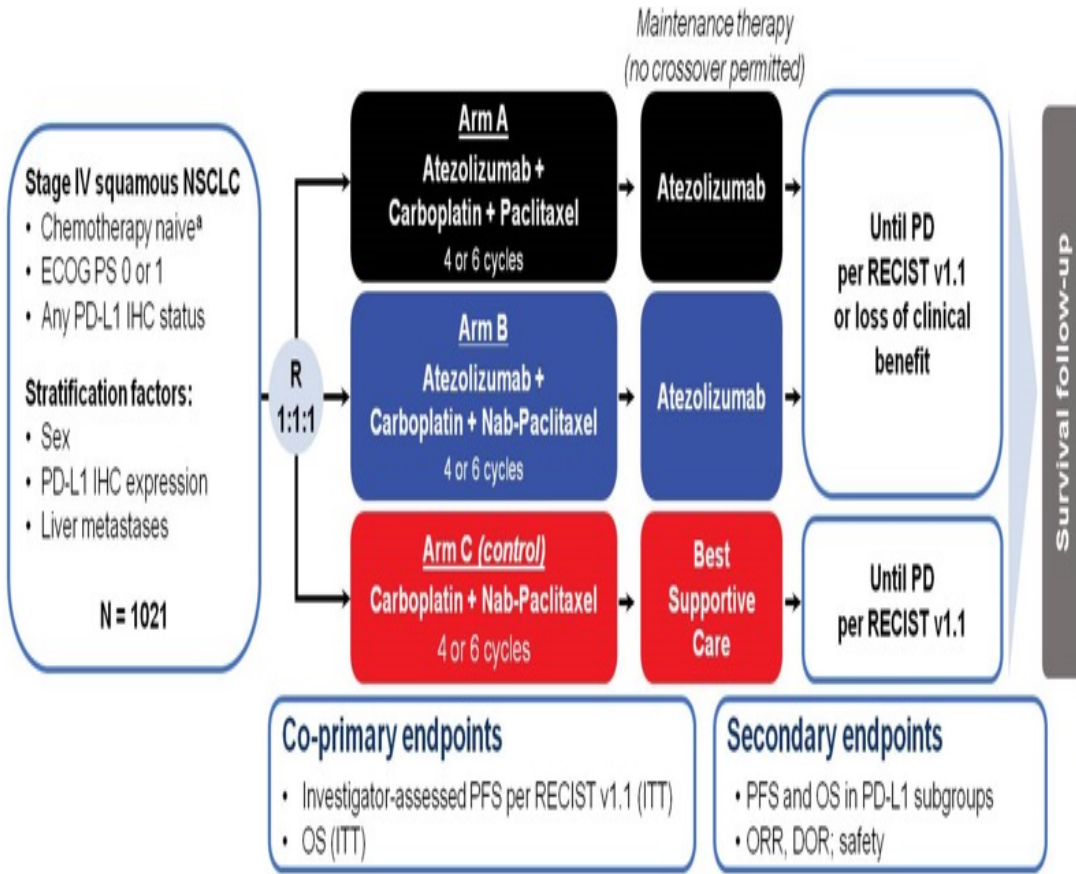


1. Paz-Ares. J Thorac Oncol. 2020;15:1657. 2. Gandhi. NEJM. 2018;378:2078.

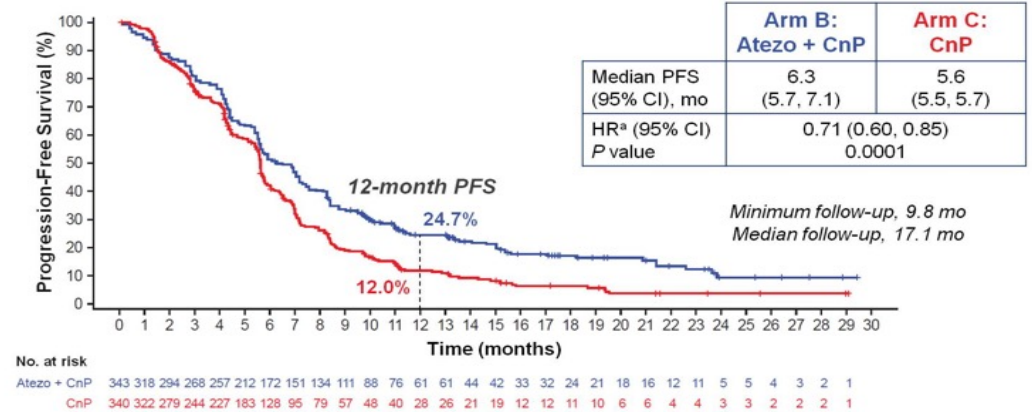


# IMpower 131

## Atezolizumab + Chemotherapy

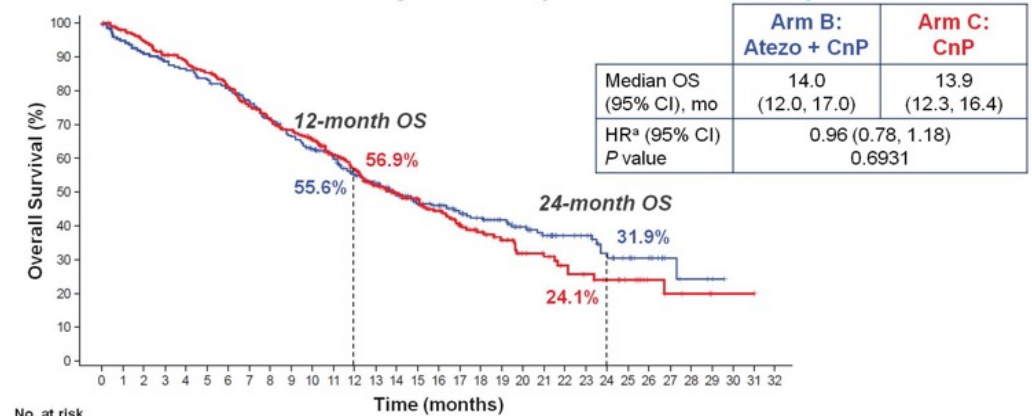


### INV-Assessed PFS in the ITT Population (Arm B vs Arm C)



Data cutoff: January 22, 2018.  
INV, Investigator. <sup>a</sup> Stratified HR.

### First Interim OS in the ITT Population (Arm B vs Arm C)



Data cutoff: January 22, 2018.  
<sup>a</sup> Stratified HR.

Atezolizumab 1200 mg IV q3w; carboplatin AUC 6 IV q3w; nab-paclitaxel 100 mg/m<sup>2</sup> IV qw; paclitaxel 200 mg/m<sup>2</sup> IV q3w.

<sup>a</sup> Patients with a sensitising EGFR mutation or ALK translocation must have disease progression or intolerance to treatment with ≥ 1 approved targeted therapies. Testing for EGFR mutation or ALK translocation was not mandatory.

<sup>b</sup> PD-L1 expression was evaluated using the VENTANA SP142 IHC assay.

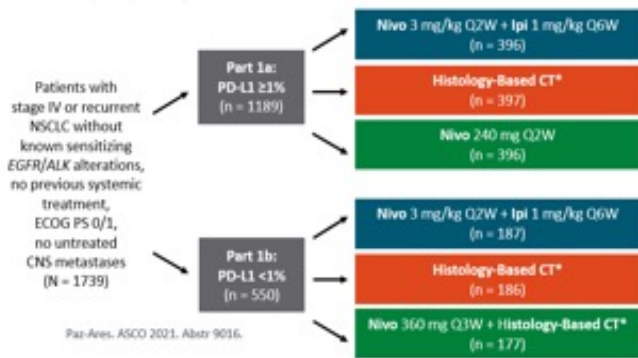
# ICI Combinations

## With and without Chemotherapy

### CheckMate 227

#### First-line Nivolumab + Ipilimumab vs Chemotherapy

- Randomized, open-label, multipart phase III trial  
Stratified by histology (squamous vs nonsquamous)



Up to 2 yr for immunotherapy

- Dual primary endpoints for nivo + ipi vs CT

- PFS in high TMB ( $\geq 10$  mut/Mb) population
- OS in PD-L1  $\geq 1\%$  population

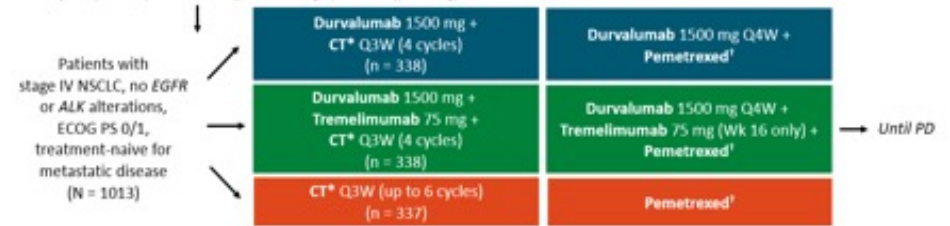
\*Nonsquamous: pemetrexed + cisplatin or carboplatin Q3W for 54 cycles with optional maintenance (pemetrexed after CT, or nivo + pemetrexed after nivo + pemetrexed)  
Squamous: gemcitabine + cisplatin or carboplatin Q3W for 54 cycles.

Pao-Ann, ASCO 2021, Abstr 9016.

### POSEIDON

#### First-line Durvalumab ± Tremelimumab + Chemotherapy in NSCLC

- Open-label, multicenter, randomized phase III trial  
Stratified by PD-L1 ( $\geq 50\%$  vs  $<50\%$ ), disease stage (IVA vs IVB), histology



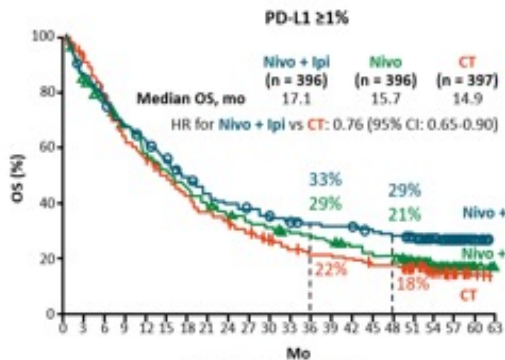
\*Gem + carbo or cis (squamous), pemetrexed + carbo or cis (nonsquamous), or nab-pac + carboplatin (either histology). Maintenance pemetrexed only given to patients with nonsquamous NSCLC who received first-line pemetrexed.

- Primary endpoints: PFS by BICR, OS (D + CT vs CT)
- Key secondary endpoints: PFS by BICR, OS, OS in pts with bTMB  $\geq 20$  mut/Mb (D + T + CT vs CT)
- Other secondary endpoints: ORR, DoR, BOR by BICR; 12-mo PFS; HRQoL; safety/tolerability

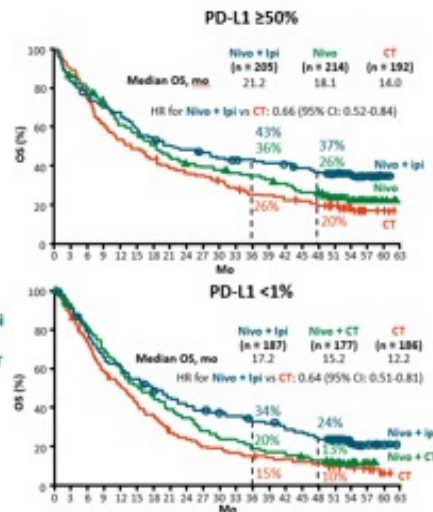
Johnson, WJCL 2021, Abstr P102.01.

### CheckMate 227

#### 4-Yr OS by PD-L1 Expression

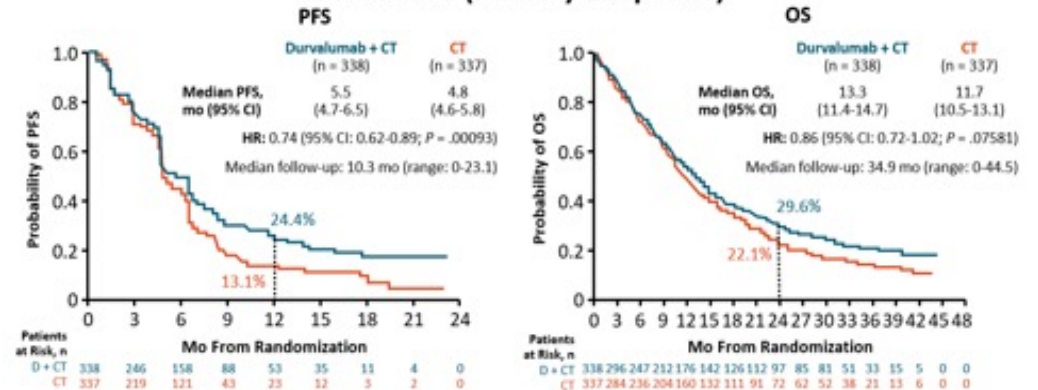


Pao-Ann, ASCO 2021, Abstr 9016.



### POSEIDON

#### OS and PFS (Primary Endpoints)



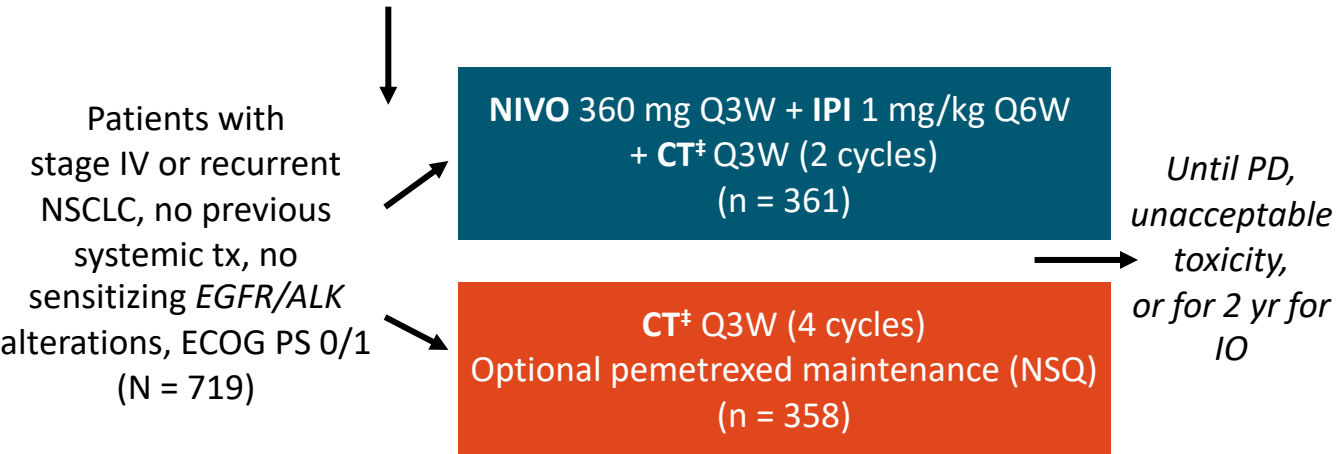
Johnson, WJCL 2021, Abstr P102.01.

# CheckMate 9LA

## Nivolumab/Ipilimumab + Chemotherapy in Advanced NSCLC

- Randomized, open-label, phase III study

Stratified by PD-L1 expression\* ( $\geq 1\%$  vs  $< 1\%$ ), sex, and histology (squamous vs nonsquamous)

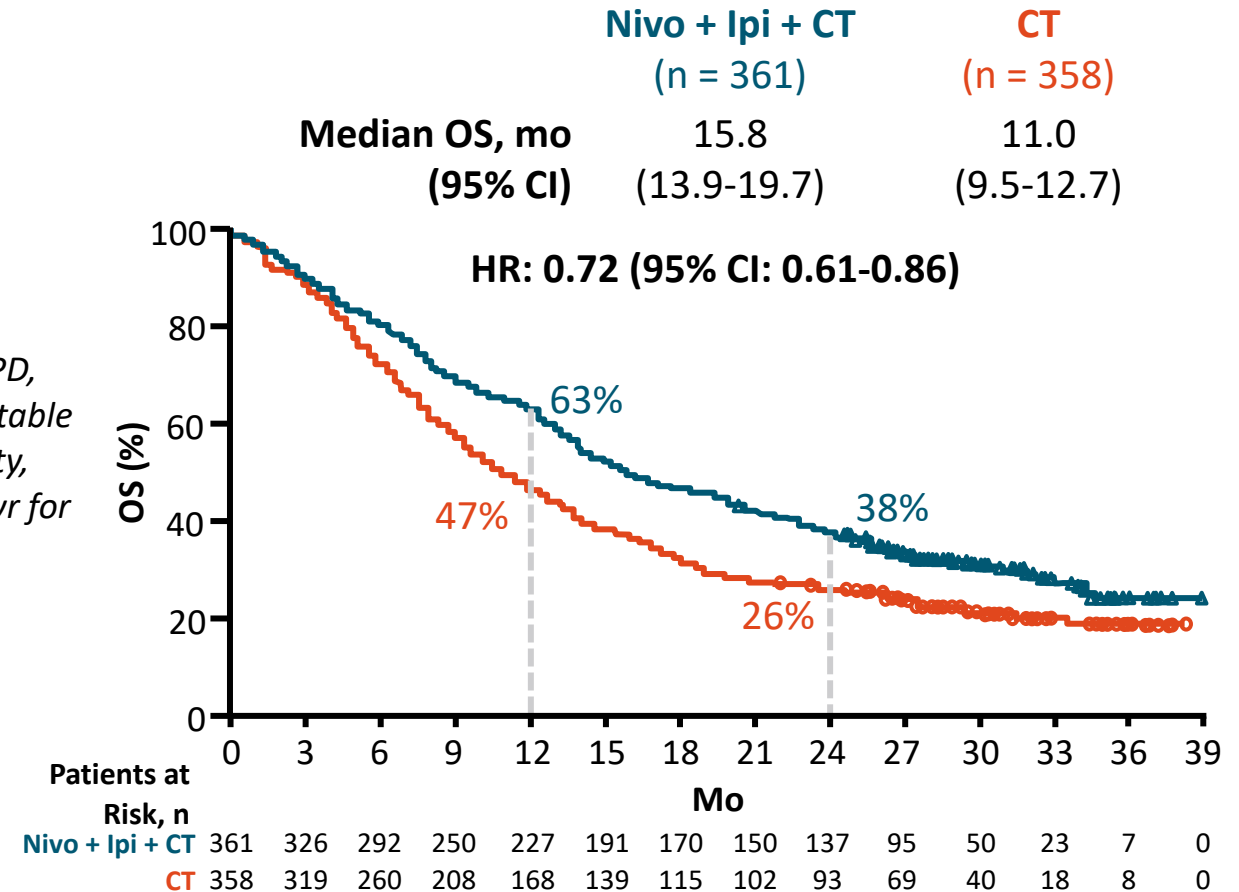


\*PD-L1 assessed by 28-8 IHC assay. <sup>†</sup>Patients unevaluable for PD-L1 were stratified to PD-L1  $< 1\%$  and capped to 10% of all randomized patients.

<sup>‡</sup>NSQ: platinum + pemetrexed; SQ: carboplatin + nab-paclitaxel.

- Regimen approved by FDA in May 2020

- DoR: 11.3 mo with nivolumab/ipilimumab + CT vs 5.6 mo with CT alone



# Immunotherapy Options for Advanced NSCLC

## High PD-L1 Expression Across Histologies

Parameter	KEYNOTE-024: Pembrolizumab (n = 154) <sup>1</sup>	IMpower110: Atezolizumab (n = 107) <sup>2</sup>	EMpower-Lung 1: Cemiplimab (n = 283) <sup>3</sup>	CheckMate 227: Nivo/Ipi (n = 205) <sup>4</sup>	CheckMate 9LA: Nivo/Ipi + CT (n = 76) <sup>5</sup>
PD-L1+ definition	TPS ≥50%*	TC3 or IC3 <sup>†</sup>	TPS ≥50%*	TPS ≥50% <sup>‡</sup>	TPS ≥50% <sup>‡</sup>
ORR, %	46.1	40.2	39.0	45.4	50.0
Median DoR, mo	29.1	38.9	16.7	31.8	26.0
Median PFS, mo	7.7 (HR: 0.50)	8.2 (HR: 0.59)	8.2 (HR: 0.54)	6.7 (HR: 0.60)	7.5 (HR: 0.59)
Median OS, mo	26.3 (HR: 0.62)	20.2 (HR: 0.76)	NR (HR: 0.57)	21.2 (HR: 0.66)	18.9 (0.67)

\*By PD-L1 22C3 IHC assay.

<sup>†</sup>Staining of ≥50% tumor cells (TC3) or ≥10% tumor-infiltrating immune cells (IC3) by PD-L1 SP142 IHC assay.

<sup>‡</sup>PD-L1 28-8 IHC assay.

1. Reck. JCO. 2021;39:2339. 2. Jassem. J Thorac Oncol. 2021;[Epub]. 3. Sezer. Lancet. 2021;397:592. 4. Paz-Arez. ASCO 2021. Abstr 9016. 5. Reck. ASCO 2021. Abstr 9000.

# Immunotherapy Options for Advanced NSCLC

## PD-L1 Expression Negative Across Histologies

Outcome Across Histologies	CheckMate 227: Nivo/Ipi (n = 187) <sup>1</sup>	CheckMate 9LA: Nivo/Ipi + CT (n = 135) <sup>2</sup>
ORR, %	27.3	31.1
Median DoR, mo	18.0	17.5
Median PFS, mo (HR)	5.1 (0.74)	5.8 (0.68)
Median OS, mo (HR)	17.2 (0.64)	17.7 (0.67)

OS for Nonsquamous	CheckMate 227: Nivo/Ipi <sup>1</sup>	CheckMate 9LA: Nivo/Ipi + CT <sup>2</sup>	KEYNOTE-189: Pembro + CT <sup>3</sup>
Median OS, mo (HR)	17.5 (0.69)	Not reported (0.75)	17.2 (0.51)
2-yr OS, %	25	38	39

OS for Squamous	CheckMate 227: Nivo/Ipi <sup>1</sup>	CheckMate 9LA: Nivo/Ipi + CT <sup>2</sup>	KEYNOTE-407: Pembro + CT <sup>4</sup>
Median OS, mo (HR)	15.9 (0.53)	Not reported (0.48)	15.0 (0.79)
2-Yr OS, %	22	33	30

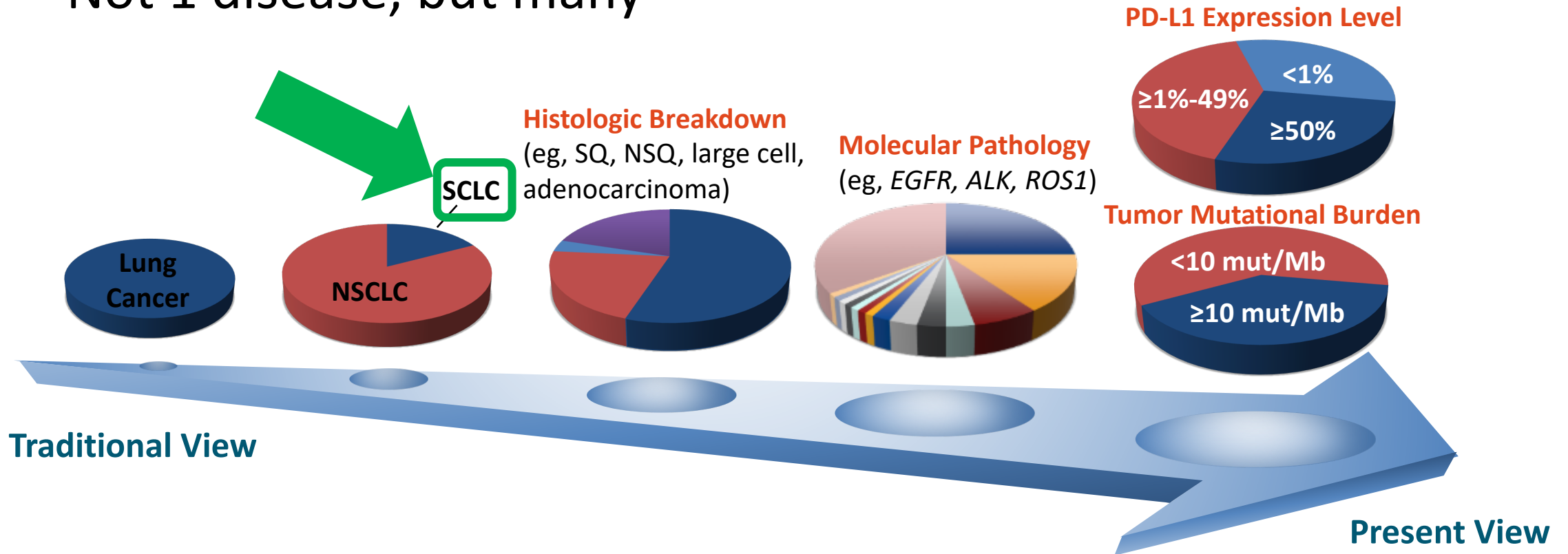
1. Paz-Ares. ASCO 2021. Abstr 9016. 2. Reck. ASCO 2021. Abstr 9000.

3. Rodríguez-Abreu. Ann Oncol. 2021;32:881. 4. Paz-Ares. J Thorac Oncol. 2020;15:1657.

# Evolution of Therapy in Lung Cancer

## Small Cell Lung Cancer (SCLC)

- Not 1 disease, but many

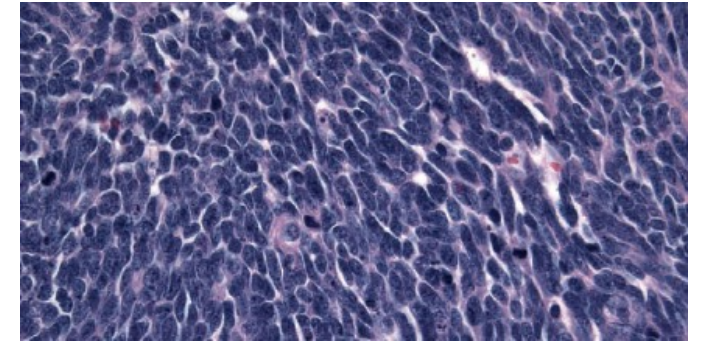


# SCLC Pathology

## Spectrum of Neuroendocrine Carcinomas (NEC)

- SCLC presents as malignant, epithelial, high-grade, neuroendocrine tumors<sup>[1,2]</sup>
  - Markers of epithelial origin
  - Neuroendocrine and neural differentiation markers: synaptophysin, chromogranin A, CD56
- SCLC falls along spectrum of WHO classification of neuroendocrine lung tumors<sup>[2,3]</sup>
- Potential therapeutic Implications

HPF View of SCLC Tumor<sup>[1]</sup>



WHO Classification <sup>[2,3]</sup>	Mitoses/ 10 HPF	Necrosis	Cytologic Features
Typical carcinoid	< 2	None	--
Atypical carcinoid	2-10	Generally punctate	--
Small-cell carcinoma	> 10	Generally abundant	Small size, scant cytoplasm, finely granular chromatin, faint nucleoli
Large-cell neuroendocrine carcinoma	> 10	Generally abundant	Cytologic features opposite SCLC

# SCLC

## Staging

The VALSG system classifies SCLC as either limited or extensive:<sup>2</sup>

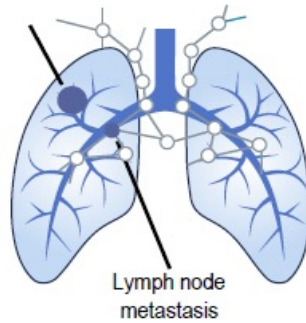
### Limited stage

- Disease confined to one hemithorax
- ± Regional lymph node metastases
- ± Pleural effusions independent of cytology

Often confined to an area that can be encompassed within a radiation port

Corresponds to stages I–III  
(T any, N any, M0)

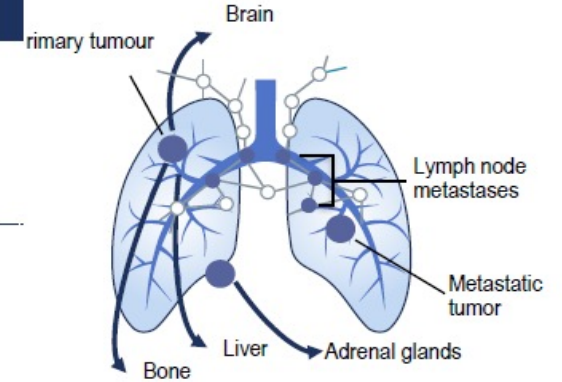
Primary tumour



### Extensive stage

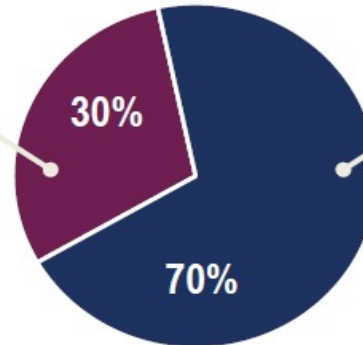
Any disease at sites beyond the definition of limited disease

Corresponds to stage IV  
(T any, N any, M1a/b)<sup>a</sup>



SCLC is predominantly diagnosed in advanced stages<sup>3</sup>

Limited stage



Extensive stage

- <sup>a</sup>Or T3–4 owing to multiple lung nodules that are too extensive or have tumoural/nodal volume that is too large to be encompassed in a tolerable radiation plan<sup>4</sup>
- SCLC, small-cell lung cancer; TNM, tumour, node, metastasis; VALSG, Veterans Administration Lung Study Group
- 1. Farago AF, et al. *Transl Lung Cancer Res* 2018;7:69–79; 2. Stahel RA, et al. *Lung Cancer* 1989;5:119–126; 3. National Cancer Institute. Small Cell Lung Cancer Treatment (PDQ®) – Health Professional Version. Available at: <https://www.cancer.gov/types/lung/hp/small-cell-lung-treatment-pdq> (Accessed November 2020);
- 4. National Comprehensive Cancer Network, Inc. NCCN Clinical Practice Guidelines in Oncology (NCCN Guidelines) for Small Cell Lung Cancer version 2.2020. Available at: [https://www.nccn.org/professionals/physician\\_gls/pdf/sclc\\_blocks.pdf](https://www.nccn.org/professionals/physician_gls/pdf/sclc_blocks.pdf) (Accessed November 2020)

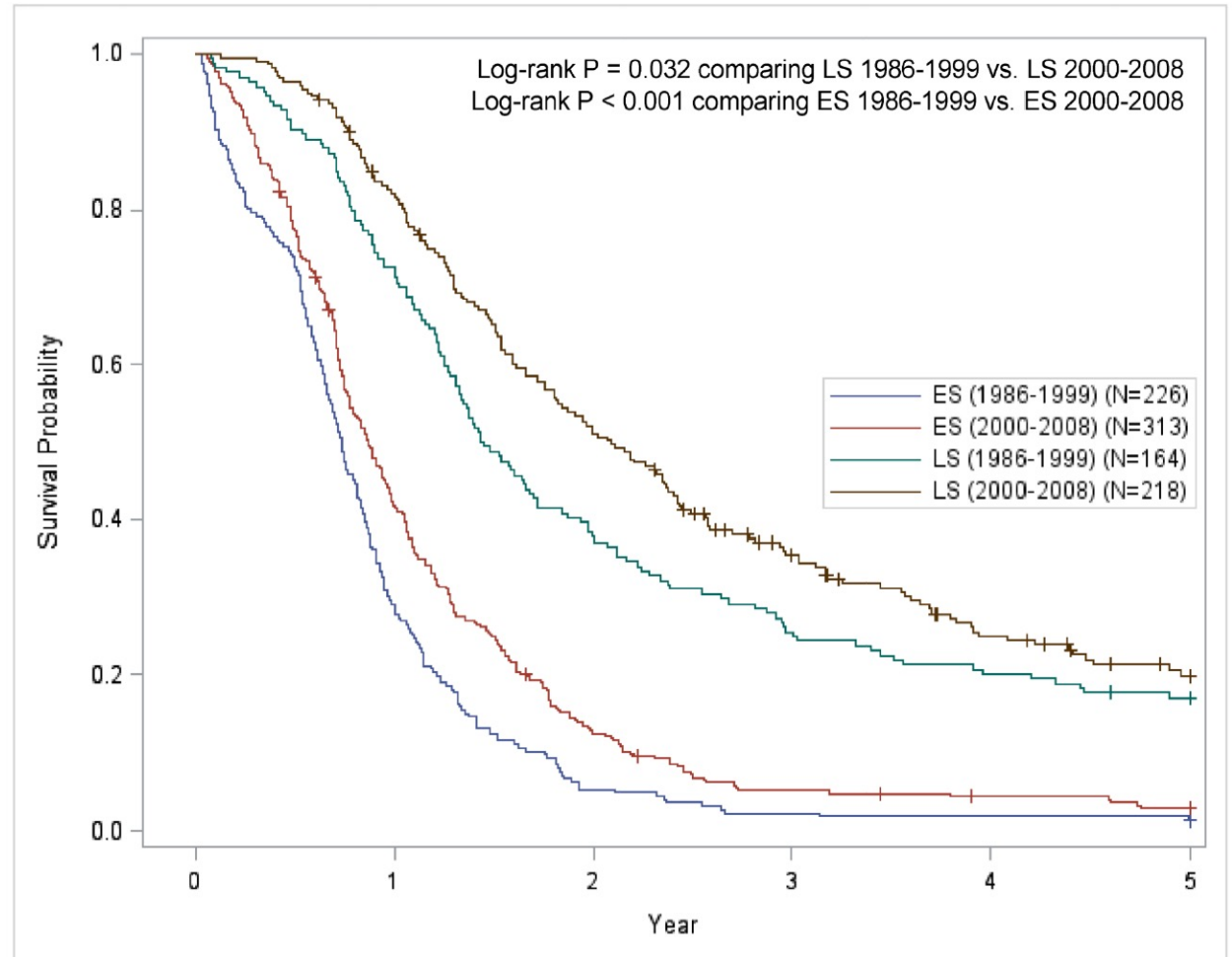


# SCLC

## Prognosis

Median survival time and 1-year, 2.5-year, and 5-year survival rate for each time-period<sup>3</sup>

	1986 to 1999 (N = 410)	2000 to 2008 (N = 593) <sup>3</sup>	P-value <sup>2</sup>
<b>Median survival time, months (95% CI)<sup>1</sup></b>			
Overall	11.3 (10.5 – 12.7)	15.2 (13.6 – 16.6)	
By stage			
Limited	17.3 (15.7 – 20.6)	25.1 (21.1 – 28.8)	
Extensive	8.8 (7.9 – 9.8)	10.4 (9.2 – 11.6)	
Not classifiable	16.1 (10.0 – 38.2)	26.1 (20.0 – 32.9)	
<b>1-year survival rate, %</b>			
Overall	48.1	60.1	< 0.001
By stage			
Limited	72.6	82.0	0.022
Extensive	28.8	41.8	0.001
Not classifiable	65.0	81.8	0.172
<b>2.5-year survival rate, %</b>			
Overall	15.9	22.6	< 0.001
By stage			
Limited	31.1	40.7	0.009
Extensive	3.6	6.8	< 0.001
Not classifiable	30.0	44.6	0.162
<b>5-year survival rate, %</b>			
Overall	8.3	11.1	< 0.001
By stage			
Limited	17.1	19.9	0.032
Extensive	1.3	2.8	< 0.001
Not classifiable	15.0	26.1	0.178



# Extensive-Stage SCLC

## First-line Chemotherapy

- SoC: cisplatin/carboplatin and etoposide for 4-6 cycles<sup>[1]</sup>
  - Response rates: ~ 50% to 75%
  - 2-yr OS: < 5%
  - Median OS: 9-11 mos
- Cisplatin vs. Carboplatin
  - Meta-analysis (N = 663)<sup>[2]</sup>
  - No differences in OS, PFS, ORR
- Irinotecan
  - superior to etoposide in Japan
  - Results not replicated in two US studies
- Pemetrexed
  - inferior to etoposide
- Inevitably, all patients progress: Recurrent Disease!

### Historic Management of SCLC

- Improving systemic therapy
  - Alternating “non-cross resistant” regimens
  - Maintenance
  - Dose escalation
  - Dose-intense regimens
  - Hematopoietic GFS
  - Two vs. more drugs
  - Four vs. six (more) cycles
  - Cisplatin vs. Carboplatin
  - Combinations with new drugs

## Extensive-Stage SCLC: “First-line” Radiation – PCI, Consolidation XRT

- PCI is guideline-recommended for pts in CR or PR; however, its use remains controversial
  - 2007: reduction in risk of brain metastases, improvement in median OS and 1-yr OS<sup>[1]</sup>
  - 2017: no improvement in OS with PCI vs observation<sup>[2]</sup>
- Thoracic RT following PCI provides OS benefit at 2 yrs<sup>[3]</sup>
  - 498 pts with response to first-line chemotherapy received PCI followed by randomization to TRT or no further therapy
    - 1-yr OS: 33% vs 28% (NS)
    - 2-yr: OS: 13% vs 3% ( $P = .001$ )

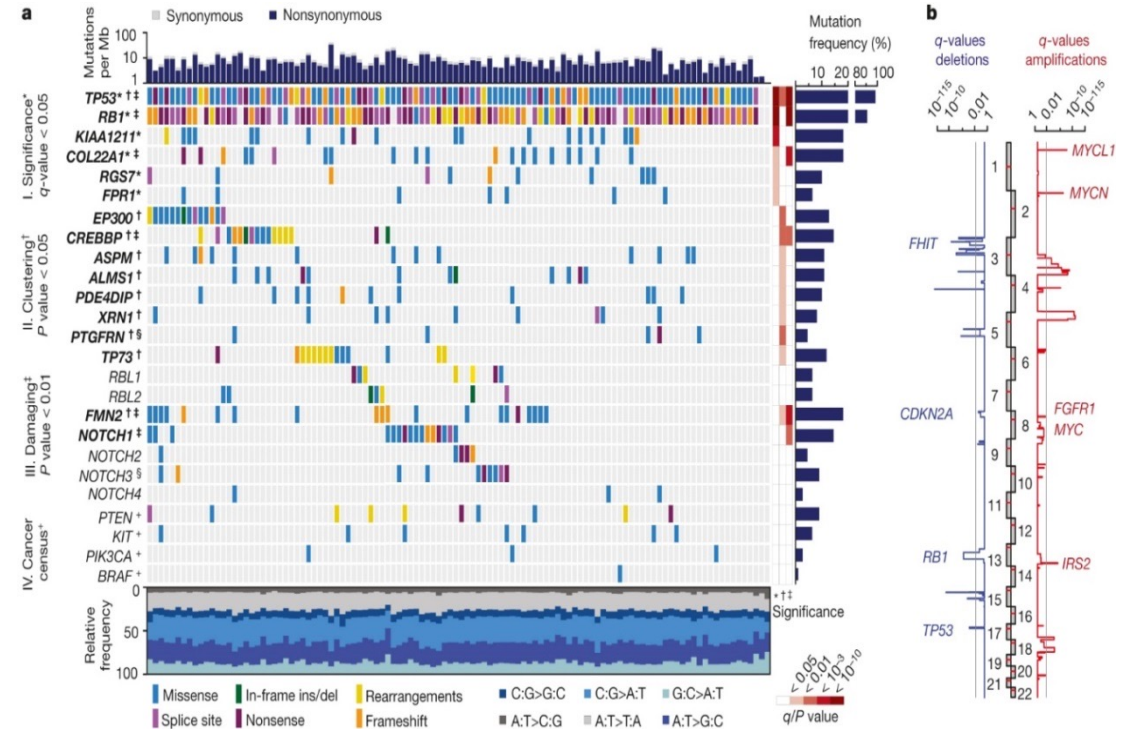
**REMAINS  
CONTROVERSIAL**

**NOT WIDELY ADOPTED**

# Genetic Alterations in SCLC

## No Clear Targetable Oncogenic Driver

- Nearly all tumors have loss or inactivation of *TP53* and *RB1*<sup>[1-4]</sup>
- *MYC* family member amplification is common<sup>[1-4]</sup>
  - *MYC-L1* > *N-MYC* > *C-MYC*<sup>[1]</sup>
  - Recurrent *RFL-MYCL1* fusions have been described<sup>[4]</sup>
- Additional alterations include:
  - *FGFR1* amplification (6%)<sup>[1]</sup>
  - *SOX2* amplification (27%)<sup>[4]</sup>
  - Recurrent point mutations in chromatin modifiers: *CREBBP*, *EP300*, *MLL* (~ 10-20%)<sup>[1,2]</sup>
  - Inactivating mutations in *NOTCH* family genes (25%)<sup>[2]</sup>
  - *EZH2*, regulator of chromatin remodeling implicated in acquired resistance<sup>[3]</sup>



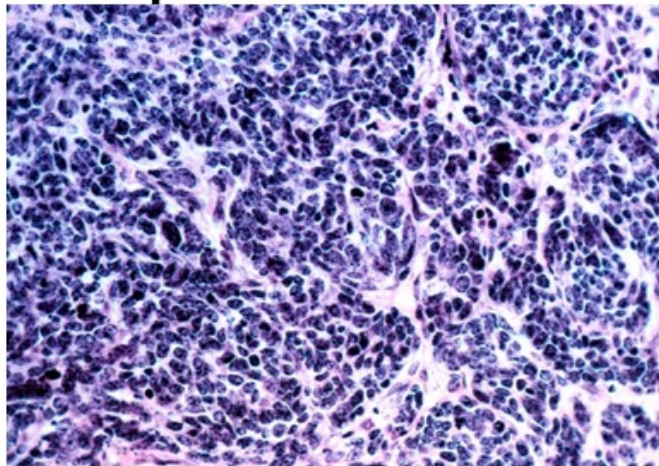
# Genetic Alterations in SCLC

## No Clear Targetable Oncogenic Driver

### Replication Stress, DDR and Genomic Instability

#### SCLC Tumor Profile:

- Mutation or loss of both TP53 and RB1
- High expression of ATM, ATR, CHK1, and CHK2, CDC25A, B, & C (relative to NSCLC)
- MYC amplification



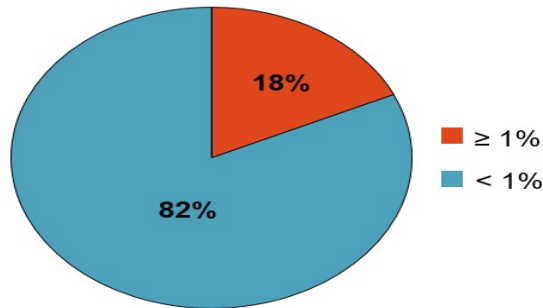
Net Effect

- Rapidly dividing tumor under immense replicative stress
- Dependence on robust DNA damage response (DDR) to maintain survival
- High level of genomic instability

# Immunotherapy and SCLC

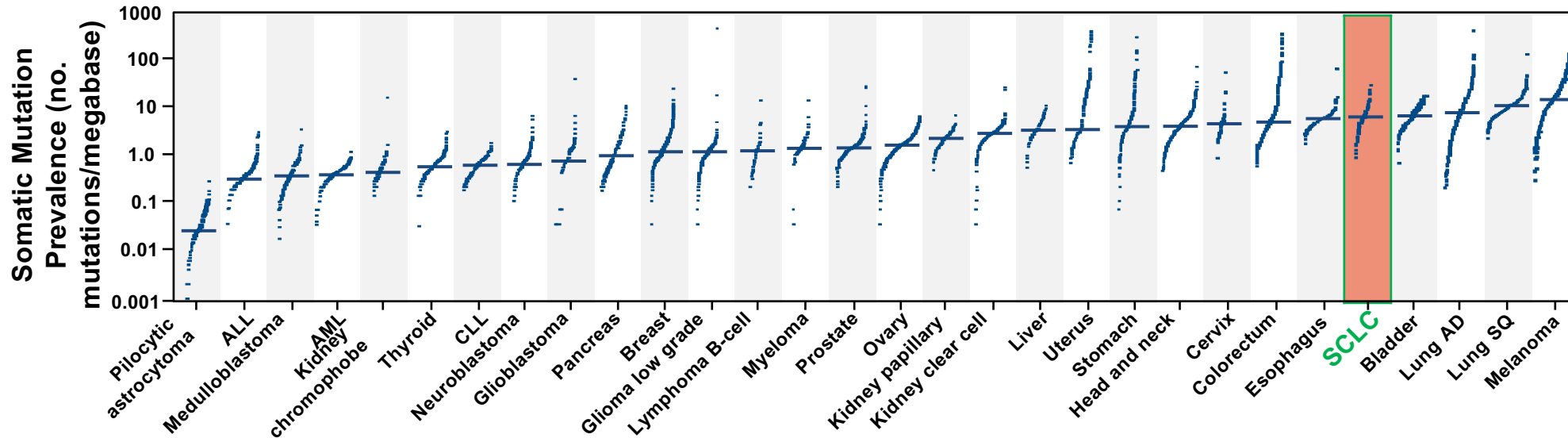
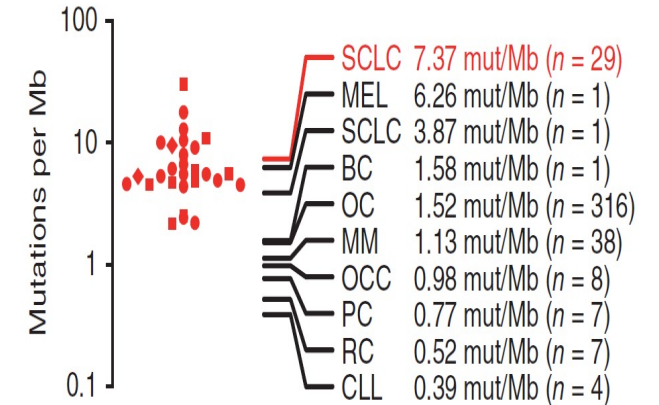
## PD-L1 expression and TMB

Tumor PD-L1 Expression in Nonrandomized Cohort (n = 159)\*



\*Pts with evaluable PD-L1 expression.

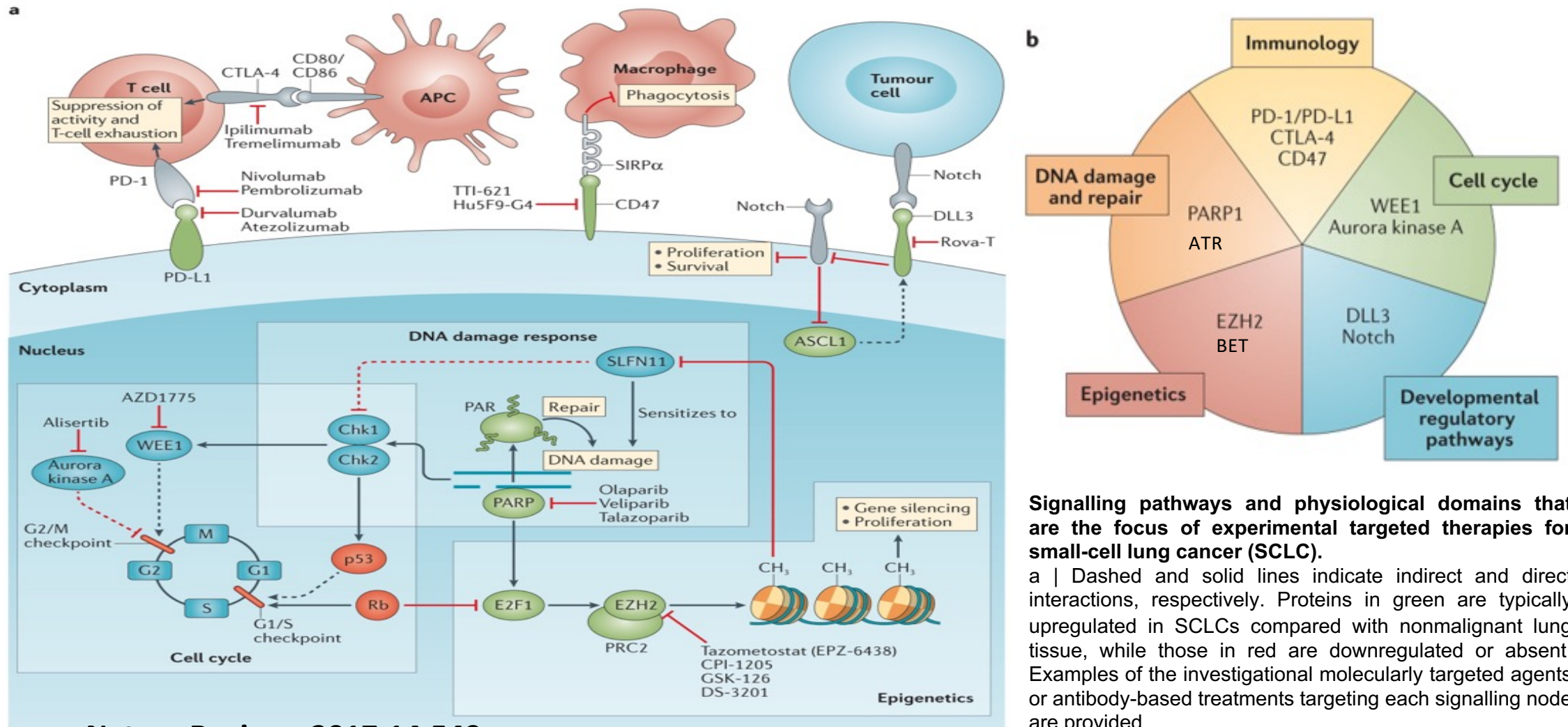
PD-L1 Expression	ORR, % (n/N)	
	Nivolumab (n = 98)	Nivolumab + Ipilimumab (n = 61)
< 1%	14 (9/64)	32 (10/31)
≥ 1%	9 (1/11)	10 (1/10)



**High mutational burden seen in SCLC**

# SCLC Systemic Therapy

## Current Areas of Advances

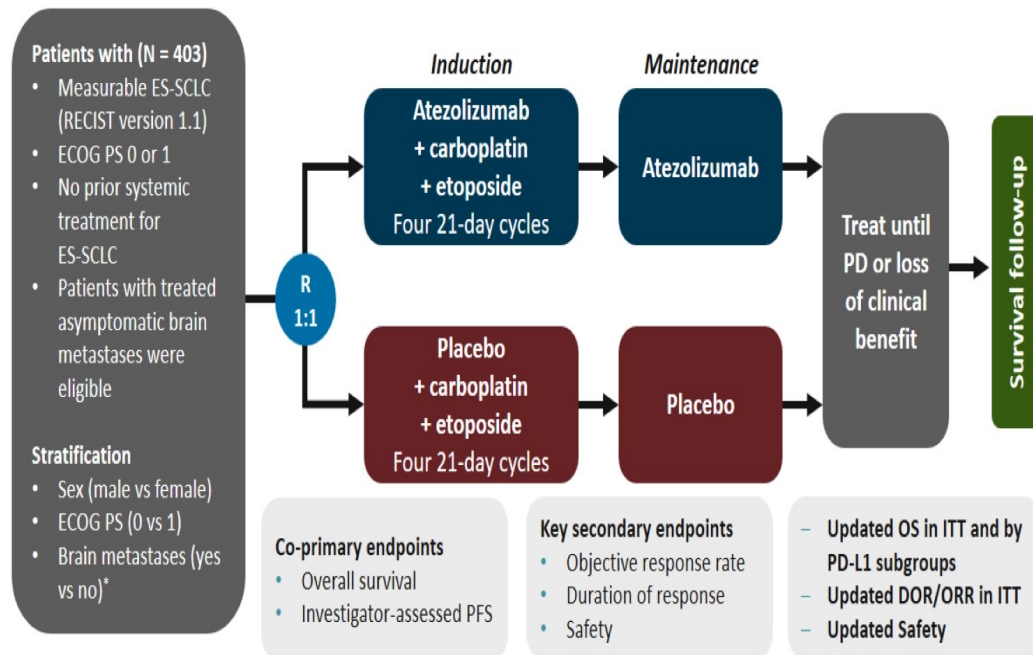


# ES-SCLC Immunotherapy

## New First Line Standard

### First-Line Treatment: IMpower133

#### Study Design



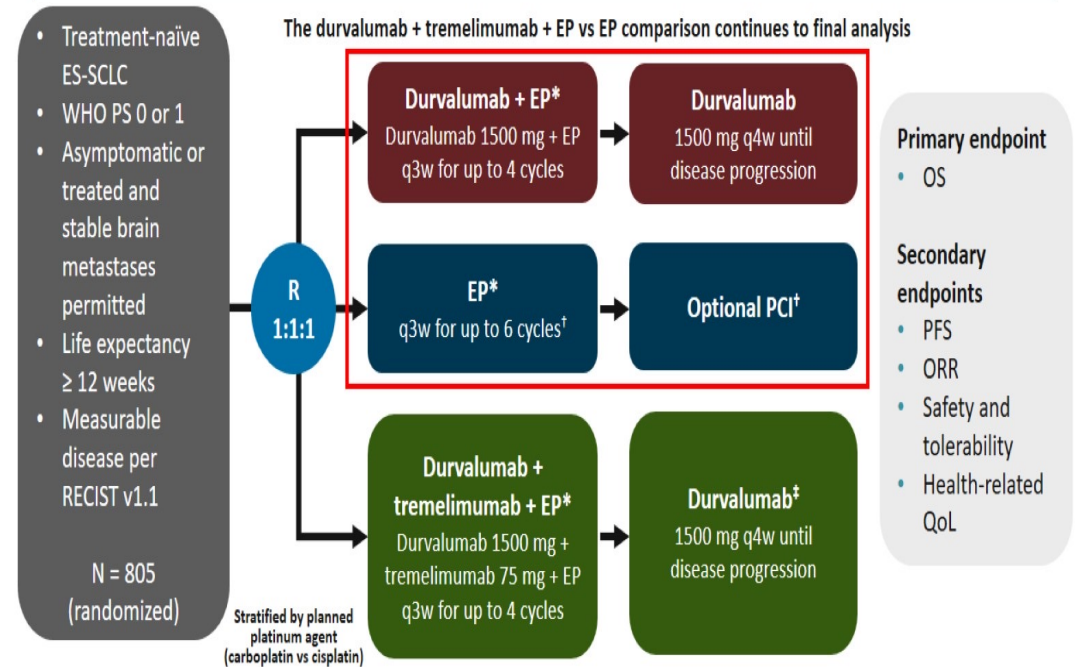
Note: Atezolizumab, 1200 mg IV, Day 1; Carboplatin, AUC 5 mg/mL/min IV, Day 1; Etoposide, 100 mg/m<sup>2</sup> IV, Days 1-3.

\*Only patients with treated brain metastases were eligible.

Horn L, et al. *N Engl J Med.* 2018;379:2220-2229; Reck M, et al. ESMO 2019. Presentation 17360.

### First-Line Treatment: CASPIAN

#### Study Design



\*EP consists of etoposide 80-100 mg/m<sup>2</sup> with either carboplatin AUC 5-6 or cisplatin 75-80 mg/m<sup>2</sup>; †Patients could receive an additional 2 cycles of EP (up to 6 cycles total) and PCI at the investigator's discretion; ‡Patients received an additional dose of tremelimumab post-EP.

Paz-Ares L, et al. *Lancet.* 2019;394:1929-1939; Paz-Ares L, et al. WCLC 2019. Presentation PLO2.11.

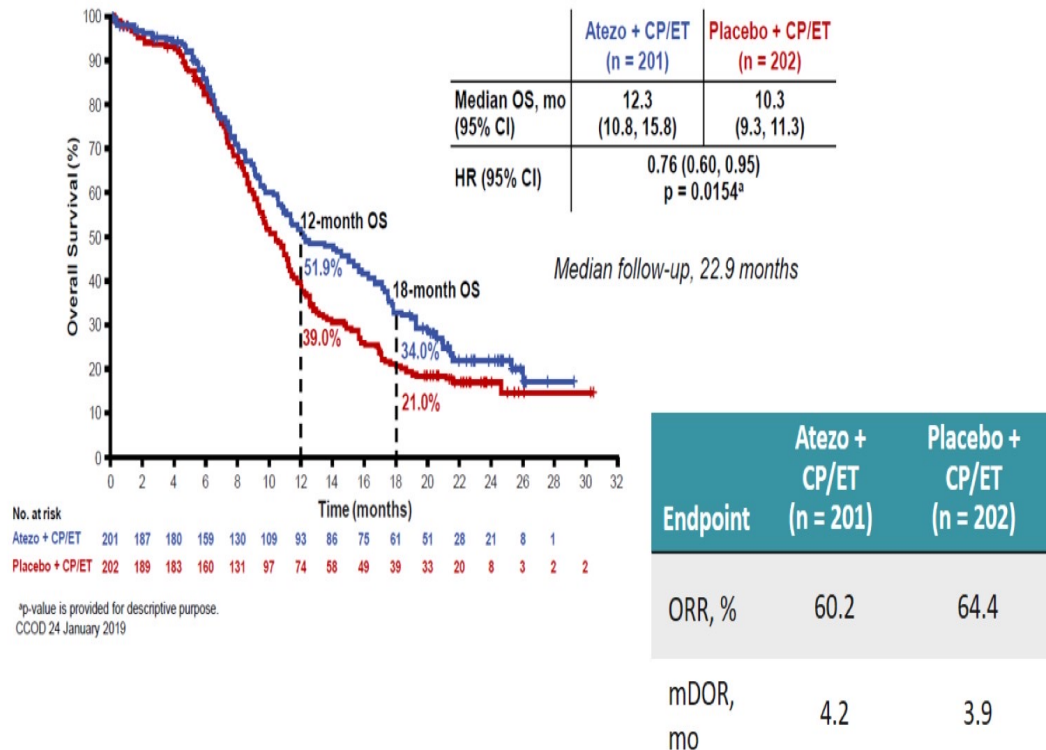


# ES-SCLC Chemo-Immunotherapy

## New First Line Standard

### First-Line Treatment: IMpower133

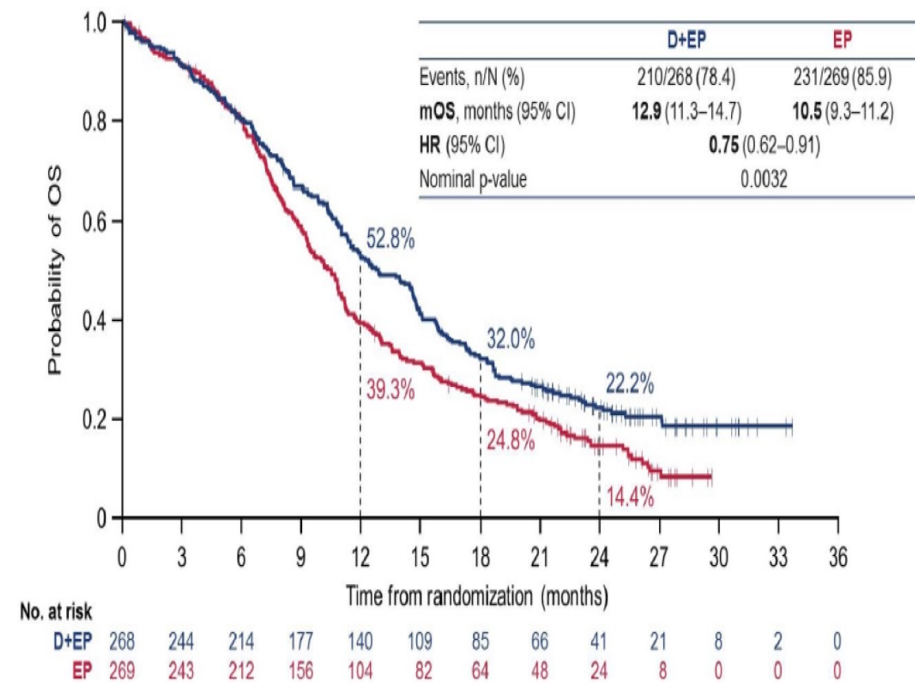
Updated Results



Reck M, et al. ESMO 2019. Presentation 17360.

### First-Line Treatment: CASPIAN

Updated OS



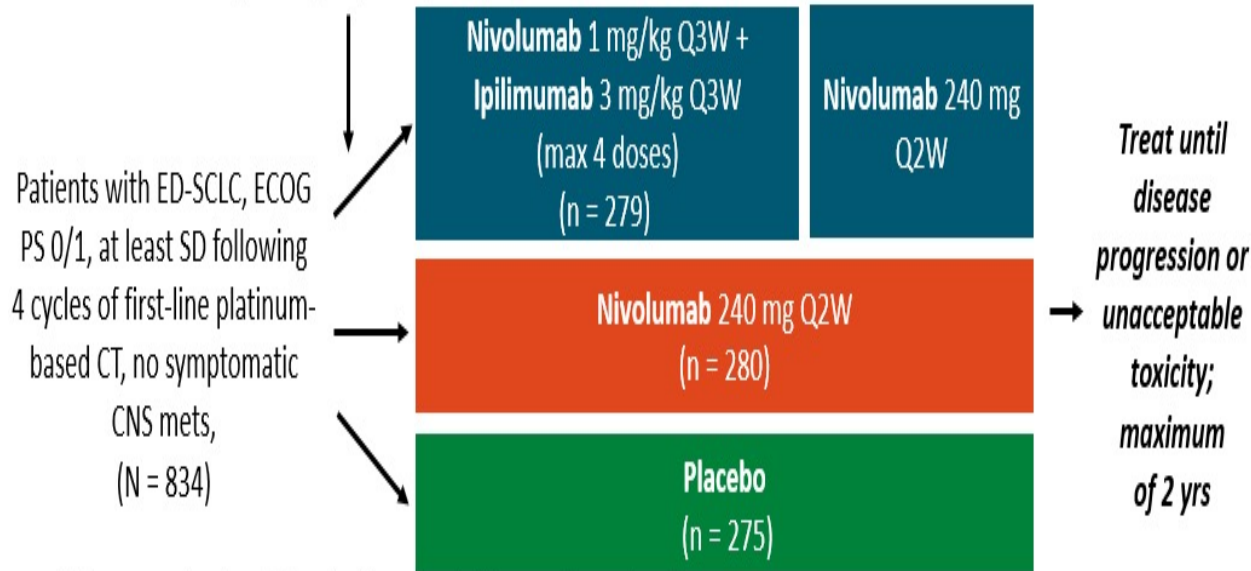
Paz-Ares L, et al. ASCO® 2020. Presentation 9002.

# Maintenance Ipi-Nivo

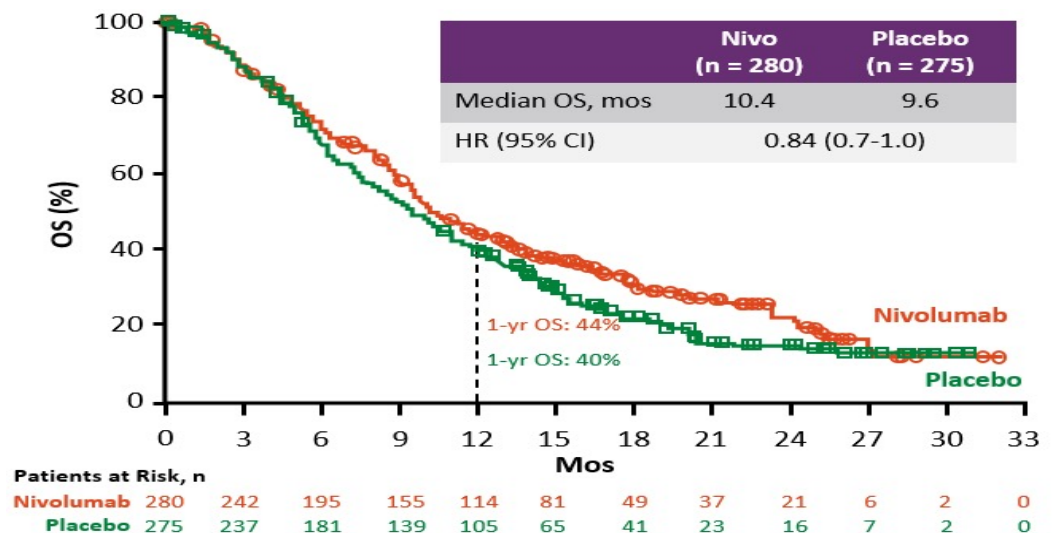
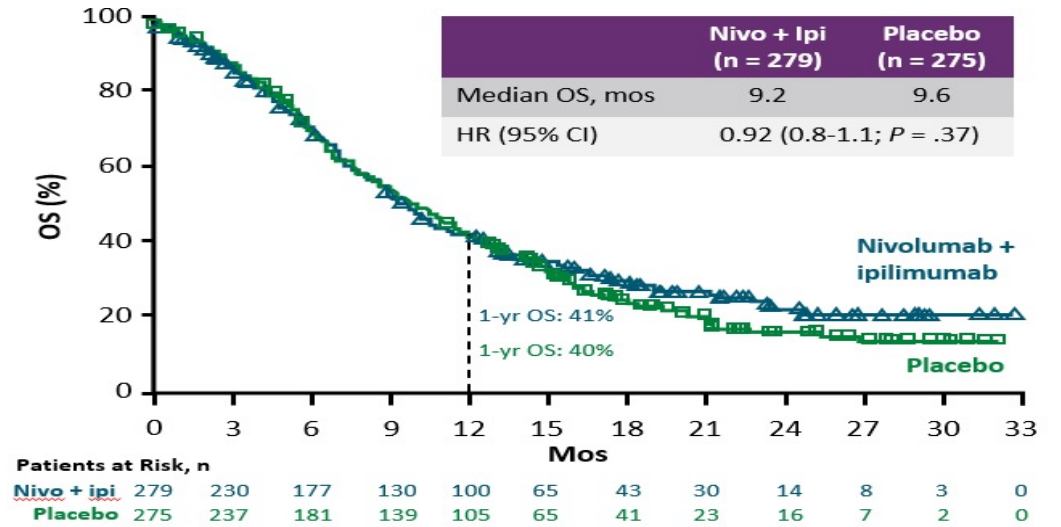
## CheckMate 451

- Randomized, double-blind phase III trial

Stratified by ECOG PS (0 vs 1), previous PCI (yes vs no), sex (male vs female)



- Primary endpoint: OS—nivolumab + ipilimumab vs placebo
- Secondary endpoints: OS—nivolumab vs placebo; PFS—nivolumab + ipilimumab and nivolumab vs placebo
- Exploratory endpoints: ORR, DoR, safety and tolerability



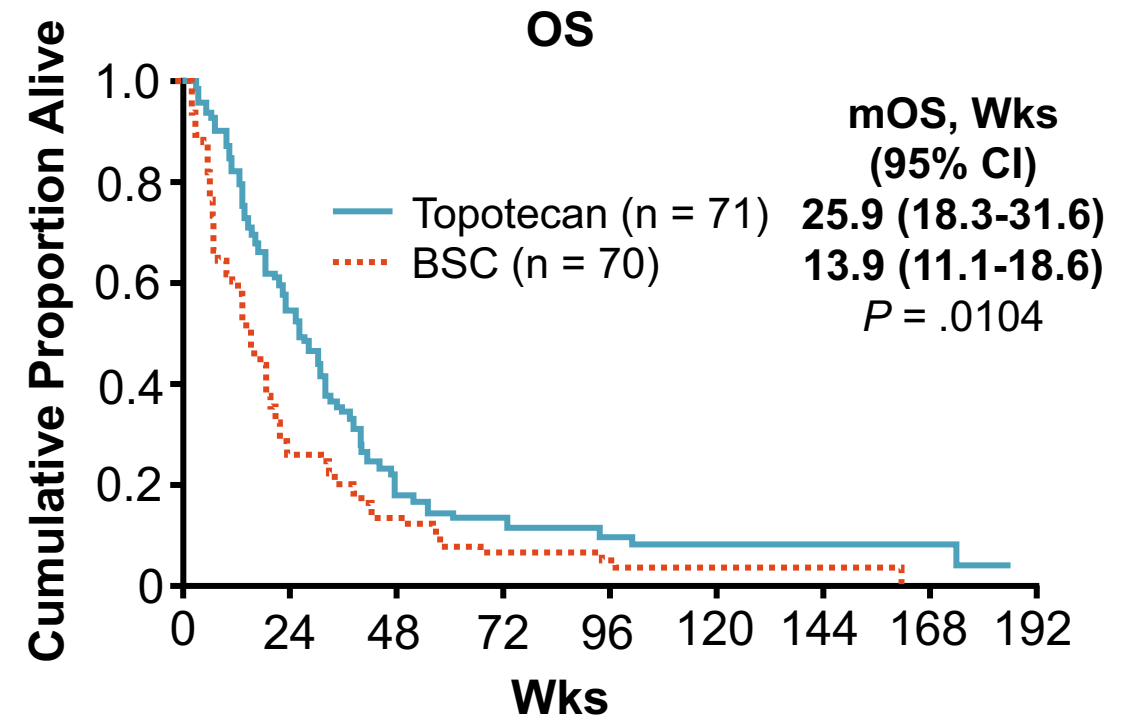
# Management of Relapsed SCLC

## Second-Line Topotecan

- Single-arm phase II study of second-line IV topotecan (N = 92)<sup>[1]</sup>
  - Overall median OS: 5.4 mos

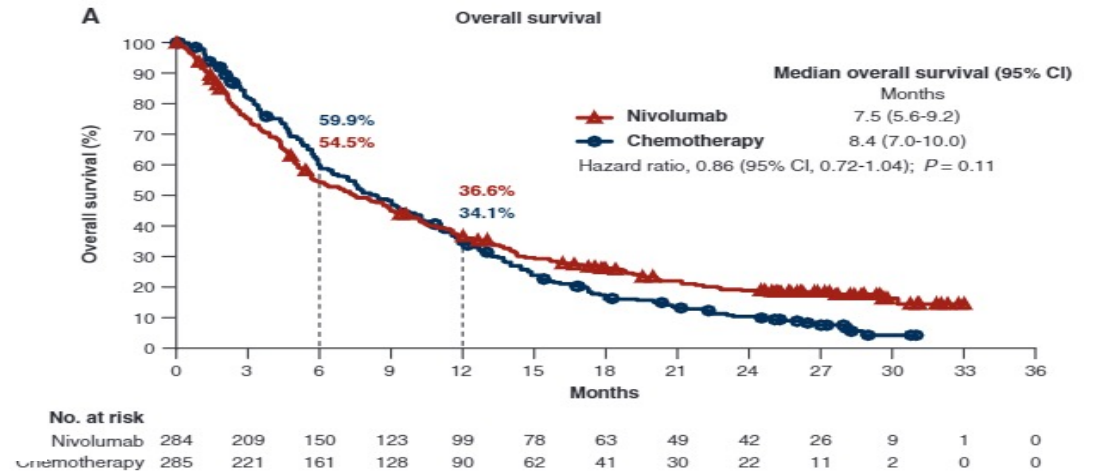
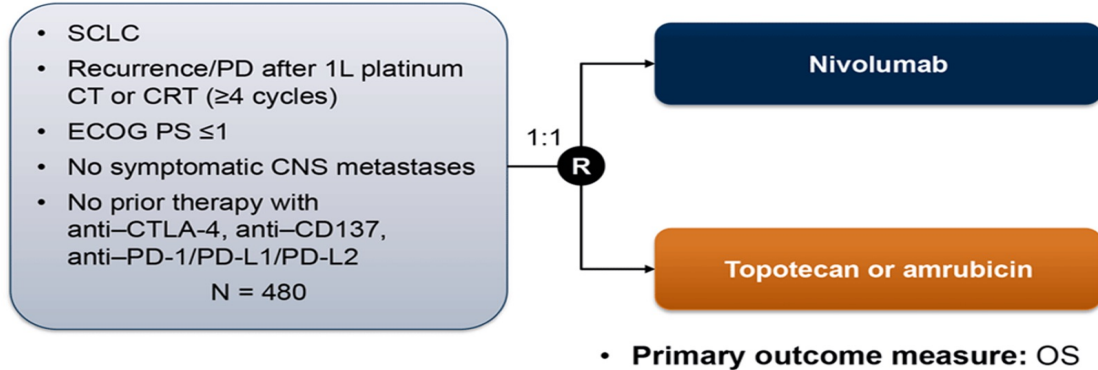
Outcome	Refractory (n = 47)	Sensitive (n = 45)
ORR, % (95% CI)	6.4 (1.3-17.6)	37.8 (23.8-53.5)
Median OS, mos	4.7	6.9

- Randomized phase III trial of BSC ± oral topotecan in pts with relapsed SCLC (N = 141)<sup>[2]</sup>



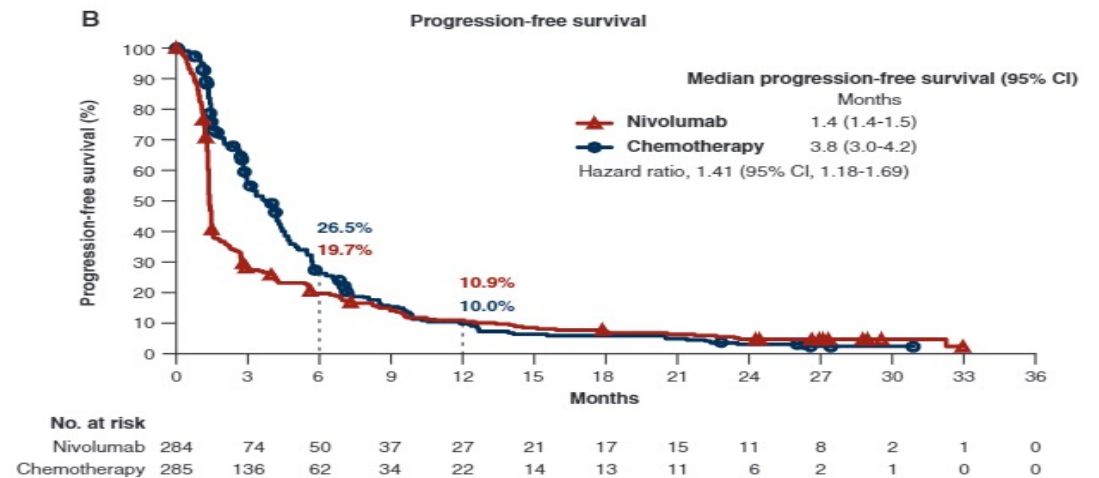
# CheckMate 331

## Nivolumab vs Topotecan/Amrubicin in Relapsed SCLC



1. Reck M et al. European Society for Medical Oncology 2018 Congress (ESMO 2018). Abstract 489.

Table 3. Summary of tumor response		
	Nivolumab (n = 284)	Chemotherapy (n = 285)
<b>Objective response<sup>a</sup></b>		
Patients with response, n	39	47
% of patients (95% CI)	13.7 (10.0-18.3)	16.5 (12.4-21.3)
Estimated odds ratio (95% CI)	0.80 (0.50-1.27)	
<b>Duration of objective response, months<sup>b</sup></b>		
Median (95% CI)	8.3 (7.0-12.6)	4.5 (4.1-5.8)
Range	0.0+ to 31.7+	1.6-23.9
<b>Best overall response, n (%)</b>		
Complete response	1 (0.4)	1 (0.4)
Partial response	38 (13.4)	46 (16.1)
Stable disease	58 (20.4)	116 (40.7)
Progressive disease	150 (52.8)	67 (23.5)
Could not be determined	37 (13.0)	55 (19.3)



# Relapsed SCLC

## Lurbinectidin (Phase II Basket Trial)

### Lurbinectidin<sup>[a,b]</sup>

- Synthetic analog of trabectedin used to treat soft-tissue sarcoma
- Selective inhibitor of oncogenic transcription
  - Covalently binds CG-rich sequences mainly located near promoters; inhibits RNA Pol II associated to DNA and leads to its specific degradation
- May also influence the TM via processes, including suppression of immune cells (eg, TAMs)

### Antitumor Activity

	Overall (n=105)
ORR, % (95% CI)	35.2 (26.2-45.2)
Best response	n (%)
- PR (confirmed)	37 (35.2) #
- SD	35 (33.3)
- PD	28 (26.7)
- NE* (non- evaluable)	5 (4.8)
Disease Control Rate,% (95% CI)	68.6 (58.8-77.3)

# 5 of 8 patients who failed prior immunotherapy had confirmed response

\* Treatment discontinuation without any tumor assessment performed

	Resistant CTFI < 90 days (n=45)	Sensitive CTFI ≥ 90 days (n=60)
ORR, % (95% CI)	22.2 (11.2-37.1)	45.0 (32.1-58.4)
Best response (confirmed)	n (%)	n (%)
- PR	10 (22.2) #	27 (45.0) #
- SD	13 (28.9)	22 (36.7)
- PD	18 (40.0)	10 (16.7)
- NE* (non- evaluable)	4 (8.9)	1 (1.7)
Disease Control Rate), % (95% CI)	51.1 (35.8-66.3)	81.7 (69.6-90.5)

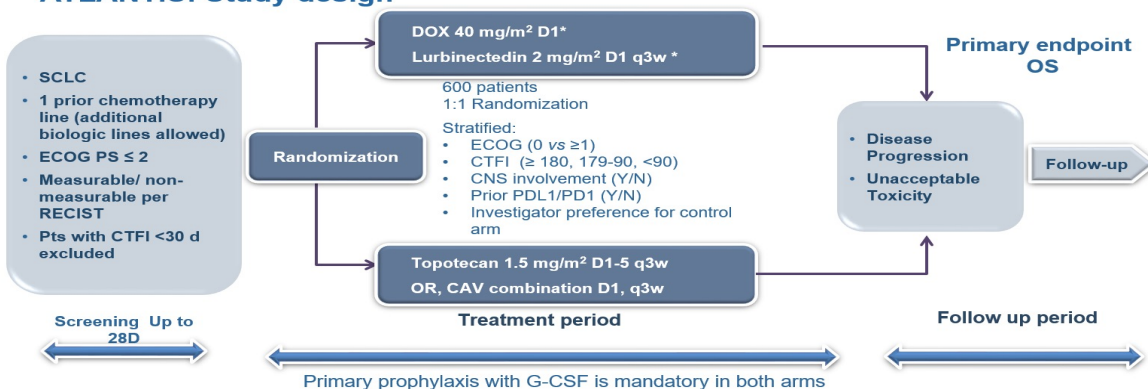
# 3 of 5 patients with resistant disease and 2 of 3 patients with sensitive disease who failed prior immunotherapy had confirmed response

\* Treatment discontinuation without any tumor assessment performed

# Phase III ATLANTIS Trial

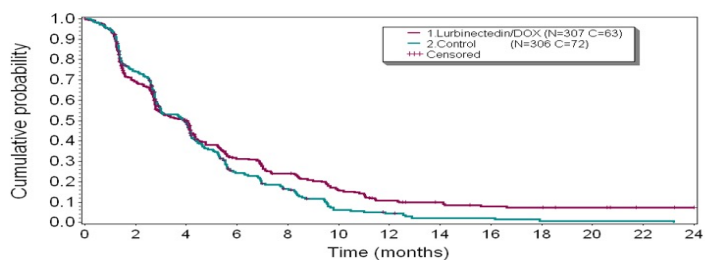
## OS and PFS

### ATLANTIS: Study design



\* Maximum 10 cycles, Lurbinectedin to be continued at 3.2 mg/m<sup>2</sup> D1 q3w

### PFS by IRC:

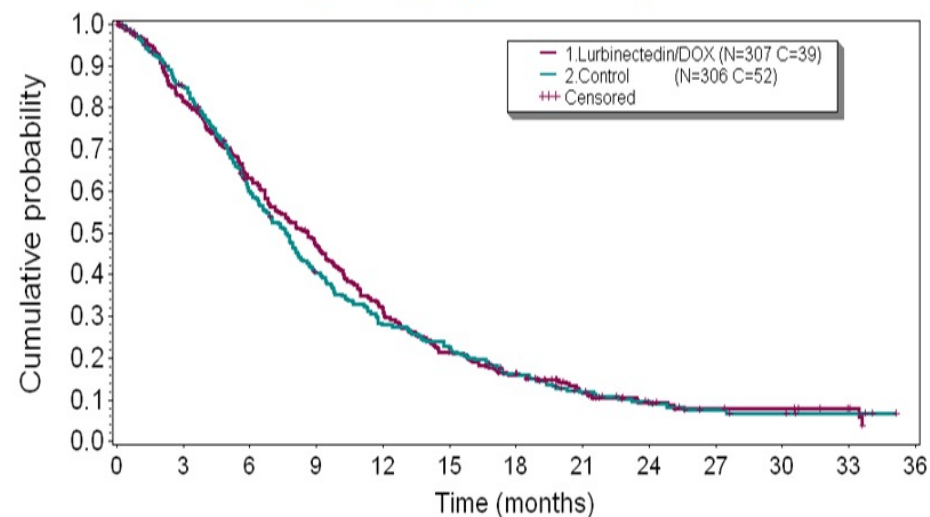


Number of patients at risk

1. Lurbinectedin+DOX	307	198	134	72	52	34	21	16	12	11	9	6	5
2. Control	306	196	119	60	32	11	7	3	3	1	1	1	1

	Lurbinectedin+DOX (N=307)	Control (N=306)	Parameter	p-value
Events, n (%)	244 (79.5)	234 (76.5)		
Censored, n (%)	63 (20.5)	72 (23.5)		
Median PFS (95% CI), months	4.0 (2.8, 4.2)	4.0 (3.0, 4.1)	HR: 0.831 (0.693, 0.996)	0.0437
Mean PFS, months	5.9	4.6		
PFS (%) at 6 months (95% CI)	31.3 (25.8, 36.9)	24.4 (19.1, 30.1)		0.0851
PFS (%) at 12 months (95% CI)	10.8 (7.1, 15.3)	4.4 (2.1, 8.1)		0.0129

### Overall Survival (ITT population)



Number of patients at risk

1. Lurbinectedin+DOX	307	247	188	138	91	62	43	25	14	10	9	5
2. Control	306	244	168	111	77	62	42	24	15	8	6	4

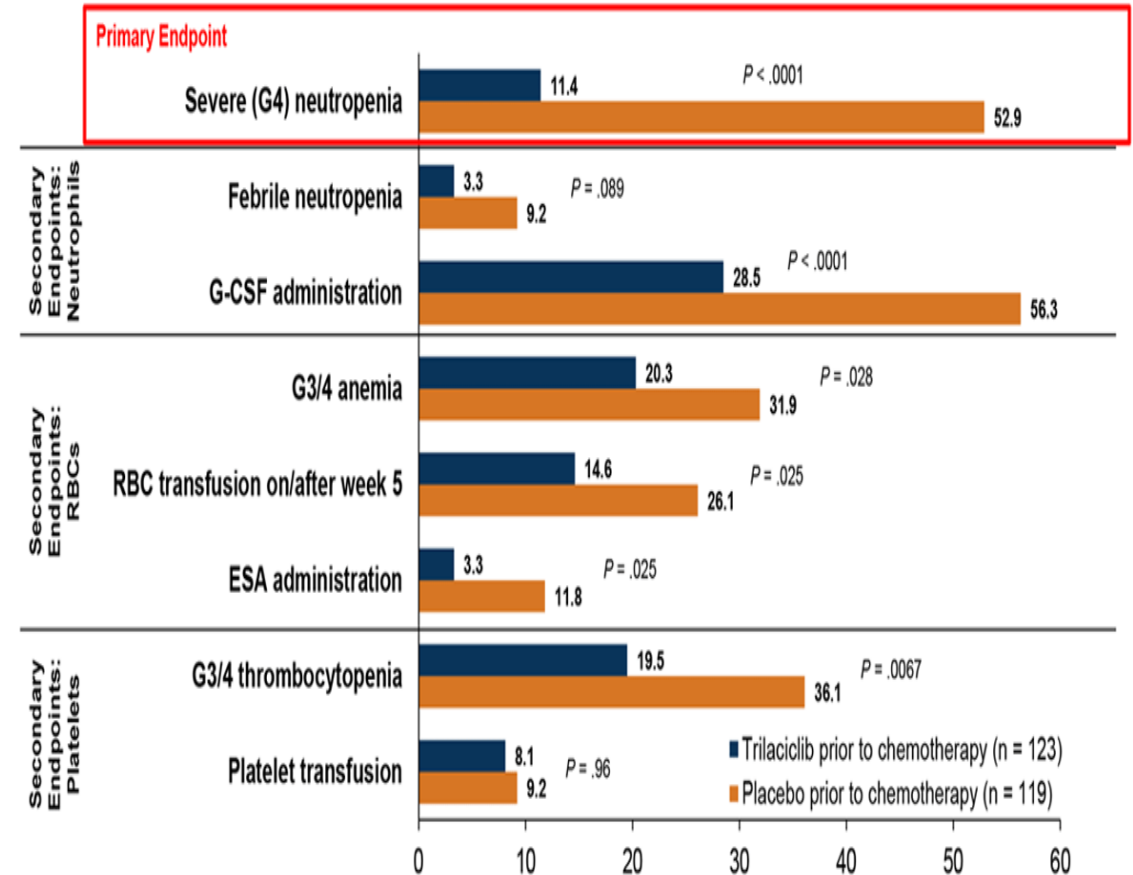
	Lurbinectedin+DOX (N=307)	Control (N=306)	Parameter	p-value
Events, n (%)	268 (87.3)	254 (83.0)		
Censored, n (%)	39 (12.7)	52 (17.0)		
Median OS (95% CI), months	8.6 (7.1, 9.4)	7.6 (6.6, 8.2)	HR: 0.967 (0.815, 1.148)	0.7032
Mean OS, months	10.6	9.9		

# Trilaciclib

## Randomized Trials – Pooled data

### Trilaciclib, a First-in-Class Myelopreservation Agent, Proactively Reduces Risks Associated with Myelosuppressive Chemotherapy

Study	Patient Population	Treatment Schedule
G1T28-05 (NCT03041311) <sup>1</sup>	Newly diagnosed (first-line) ES-SCLC	Trilaciclib 240 mg/m <sup>2</sup> IV QD prior to chemotherapy on days 1-3 of each 21-day E/P/A IV cycle for up to four cycles, followed by atezolizumab monotherapy (without trilaciclib) Q21D  Placebo IV QD prior to chemotherapy on days 1-3 of each 21-day E/P/A IV cycle for up to four cycles, followed by atezolizumab monotherapy (without placebo) Q21D
G1T28-02 (NCT02499770) <sup>2</sup>	Newly diagnosed (first-line) ES-SCLC	Trilaciclib 240 mg/m <sup>2</sup> IV QD prior to chemotherapy on days 1-3 of each 21-day E/P IV cycle  Placebo IV QD prior to chemotherapy on days 1-3 of each 21-day E/P IV cycle
G1T28-03 (NCT02514447) <sup>3</sup>	Previously treated (second-/third-line) ES-SCLC	Trilaciclib 240 mg/m <sup>2</sup> IV QD prior to topotecan 1.5 mg/m <sup>2</sup> IV QD on days 1-5 of each 21-day cycle  Placebo IV QD prior to topotecan 1.5 mg/m <sup>2</sup> IV QD on days 1-5 of each 21-day cycle

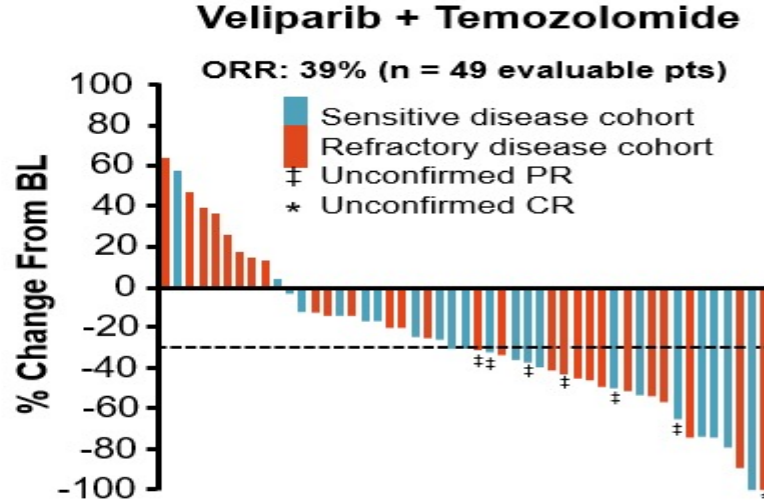
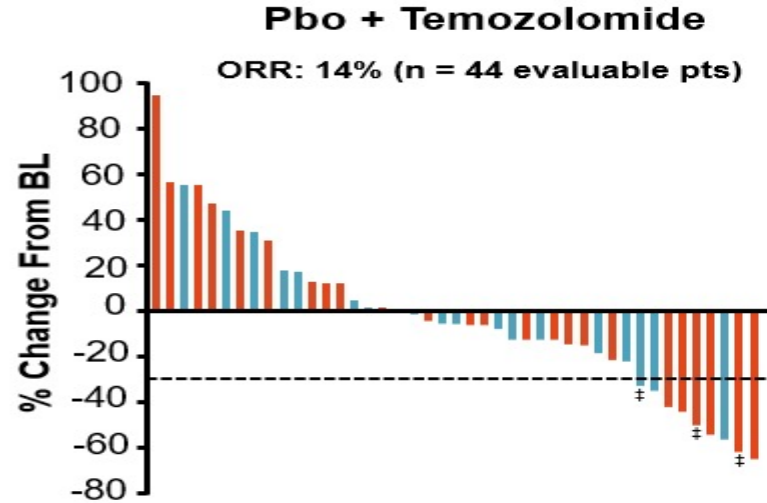


1. <https://clinicaltrials.gov/ct2/show/NCT03041311>. 2. <https://clinicaltrials.gov/ct2/show/NCT02499770>. 3. <https://clinicaltrials.gov/ct2/show/NCT02514447>.

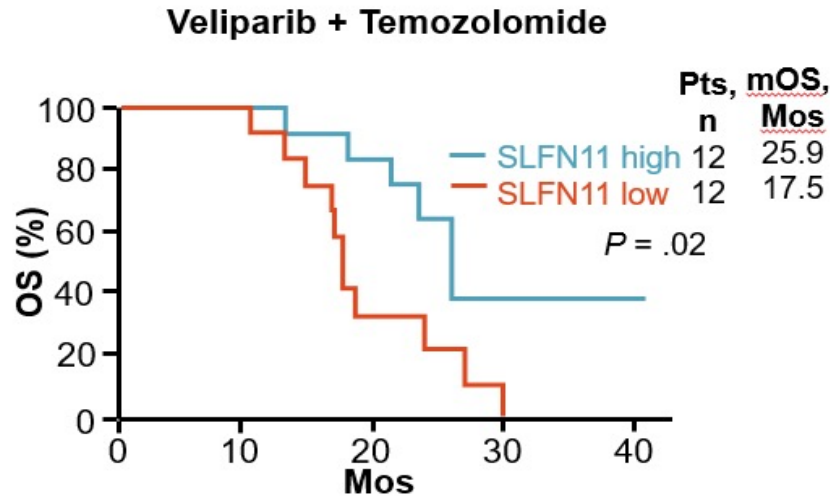
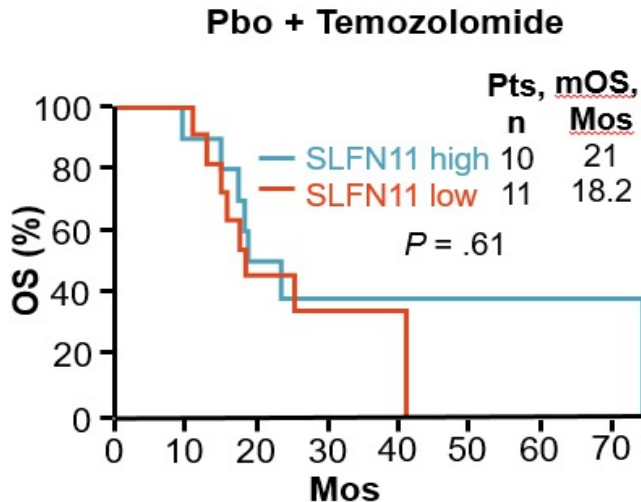
1. Weiss J et al. *Clin Lung Cancer*. 2021 Mar 26 [Epub ahead of print].

# Relapsed SCLC

## Veliparib +/- Temozolomide: ORR and OS in Pts With SLFN11 Expression



- Significantly better ORR with veliparib ( $P = .016$ )
- No significant difference in 4-mo PFS, mPFS, or mOS between arms
- Greater incidence of hematologic toxicities with veliparib combination

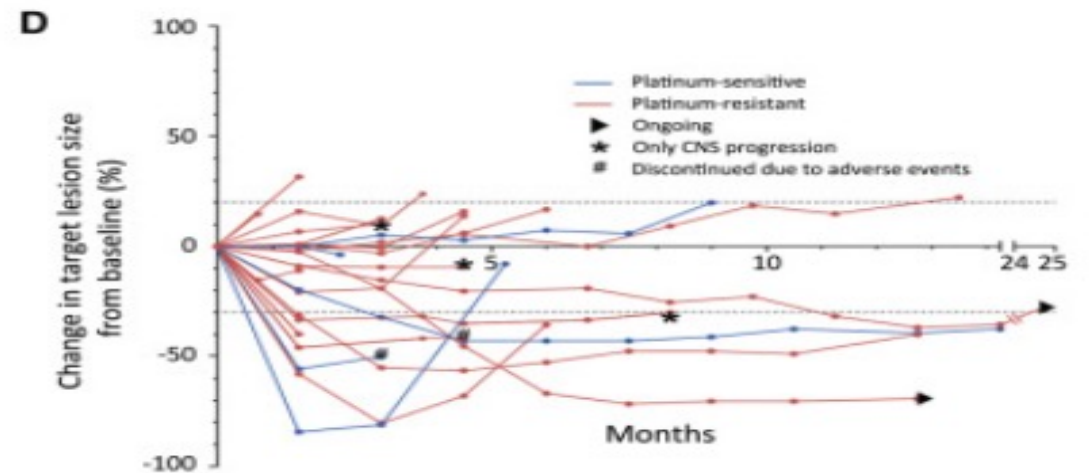
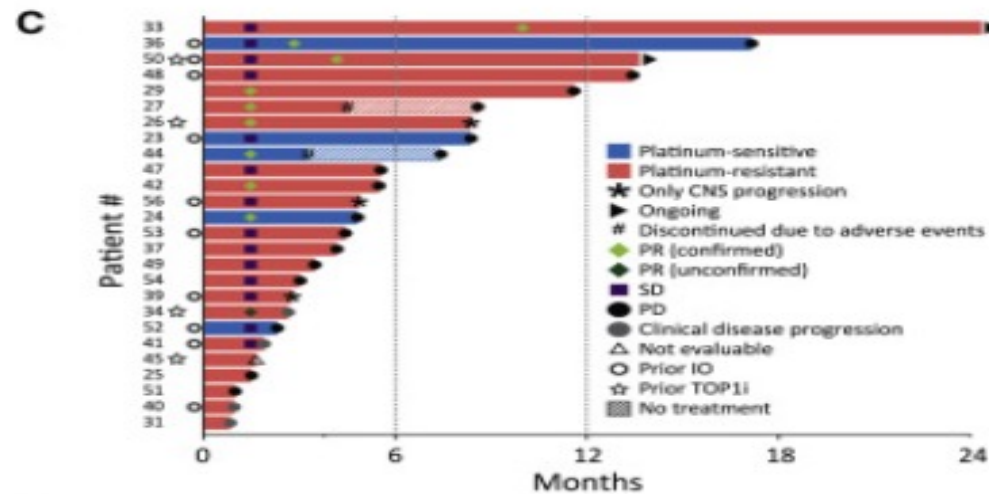
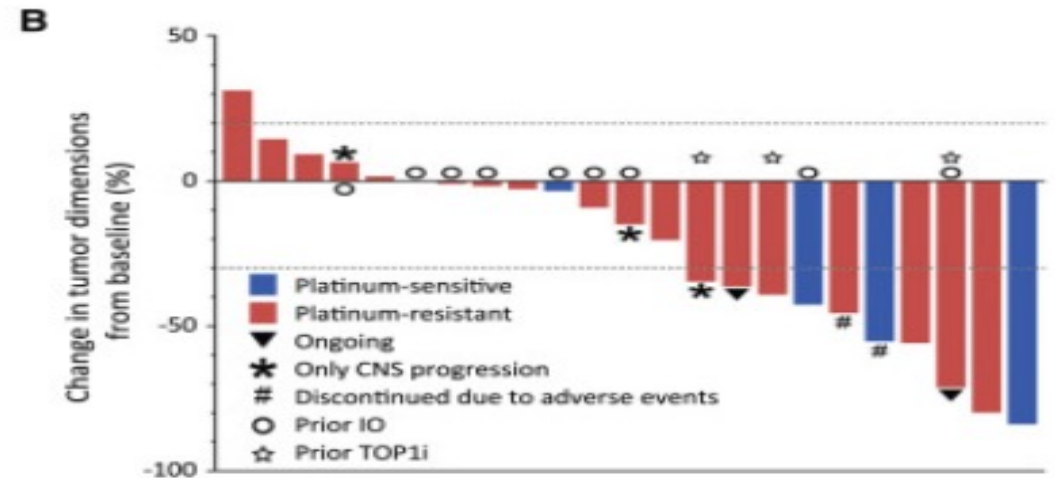
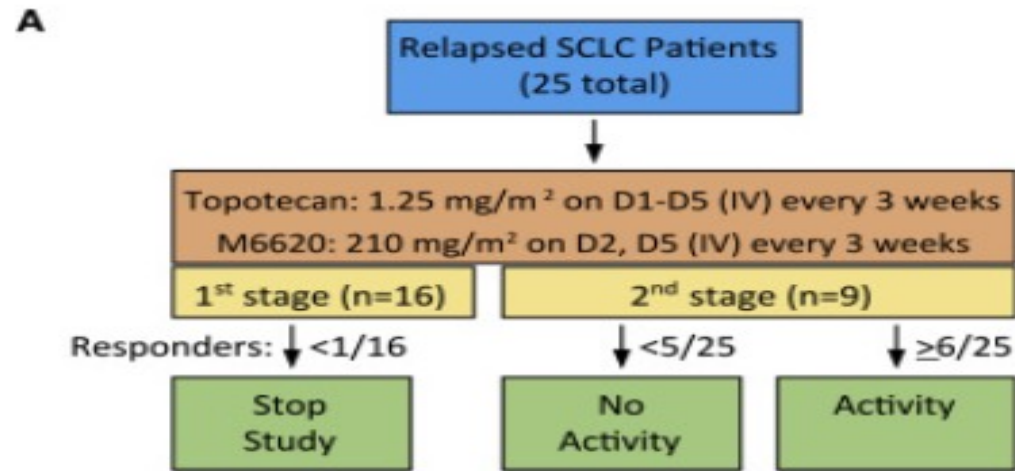


- SLFN11, a DDR protein, is aberrantly expressed in SCLC
- Veliparib: Trend toward better OS with higher tumor SLFN11 expression



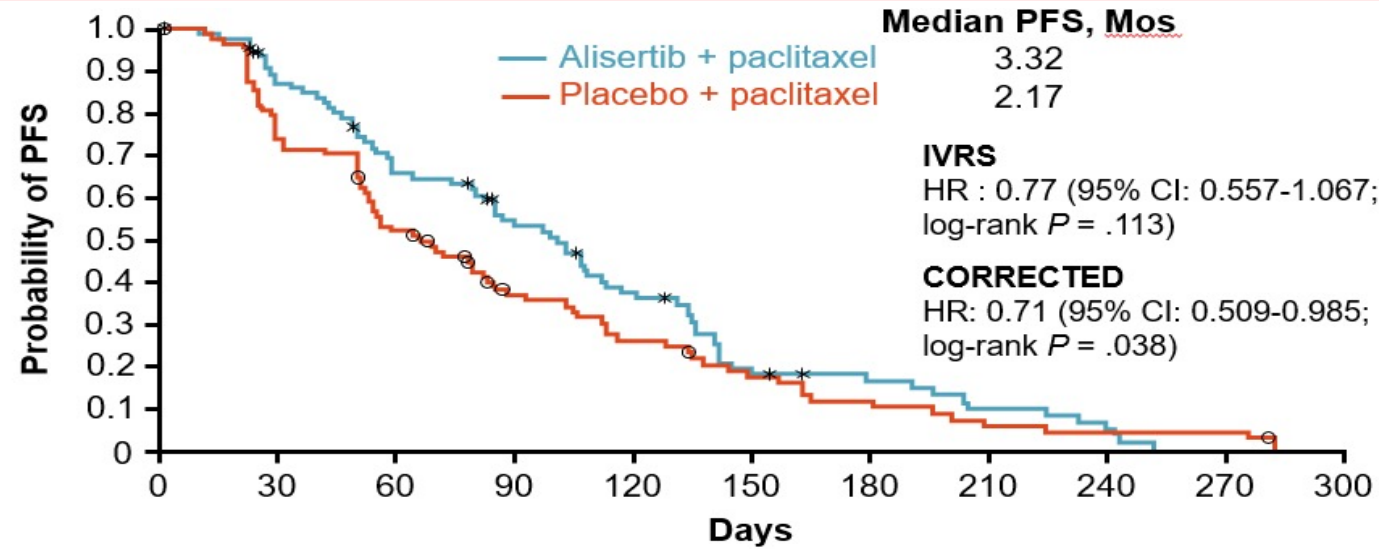
# Relapsed SCLC

## ATR inhibitor (M6620) – Study Design and ORR



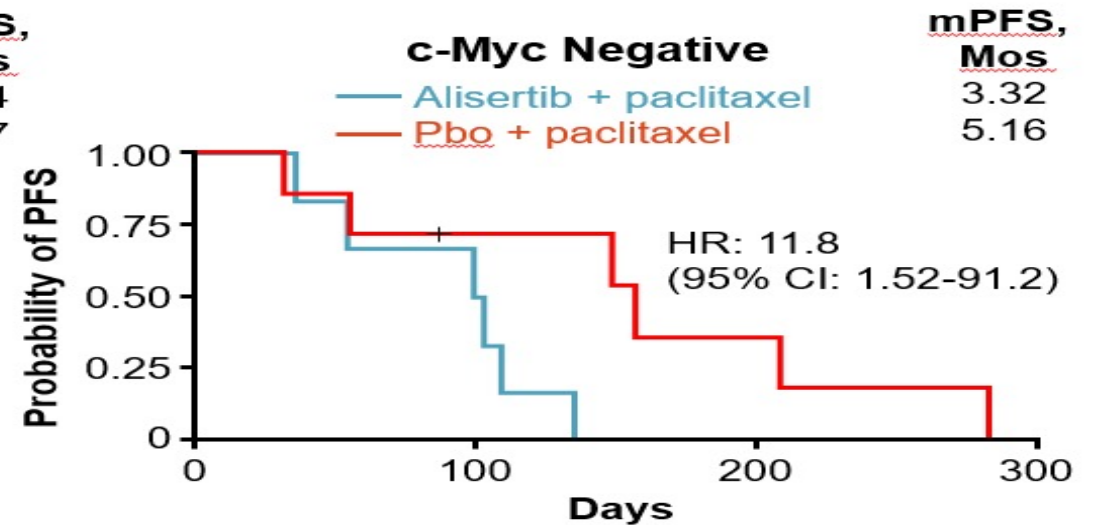
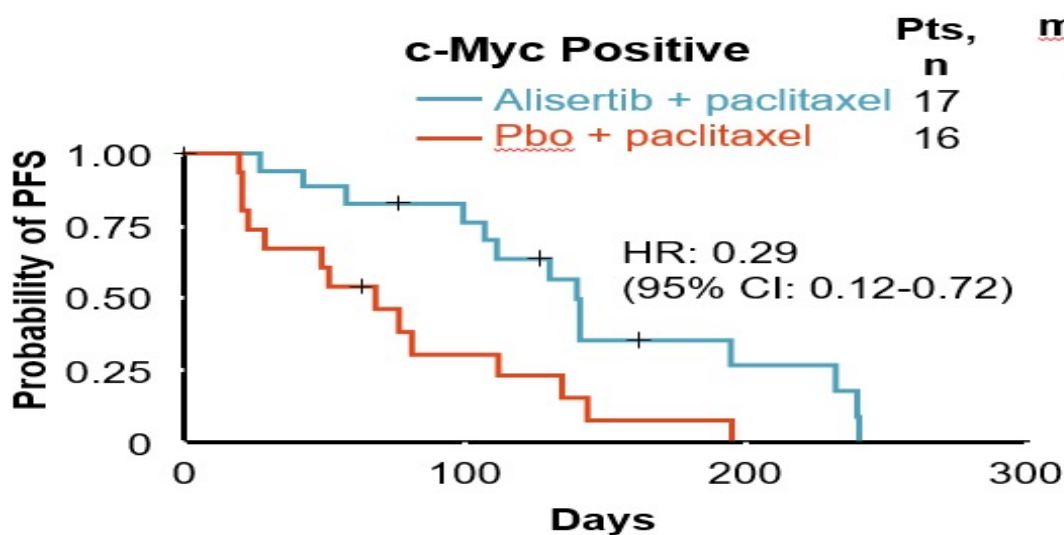
# Relapsed SCLC

## Alisertib + Paclitaxel vs Paclitaxel - PFS (ITT) and by c-Myc Expression



Archived tumor tissue available for 44 pts;  
c-Myc +ive vs c-Myc -ive:  
 $P_{\text{binary}} = .0006$

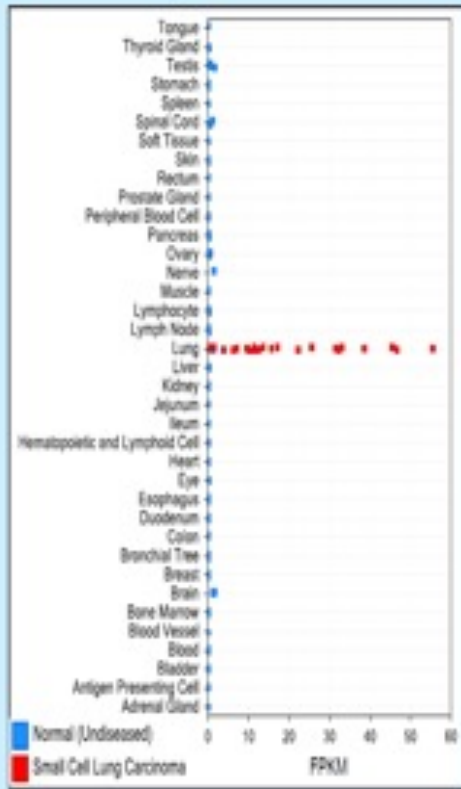
Owonikoko TK, et al. WCLC 2016. Abstract OA05.05.



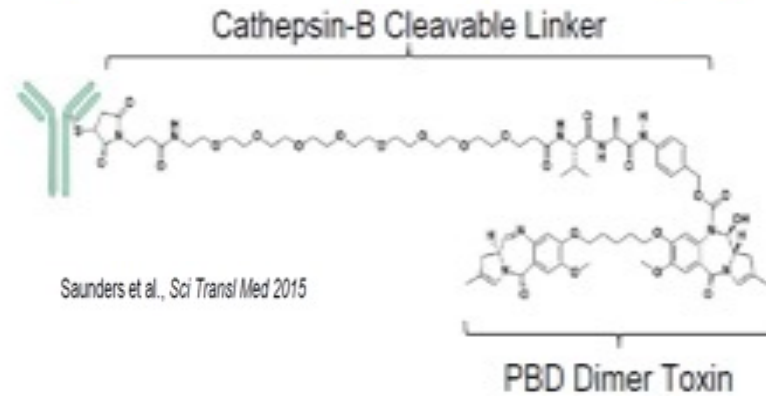
# DLL3 - NOTCH Ligand

## Antibody Drug Conjugate (ADC): Rova-T

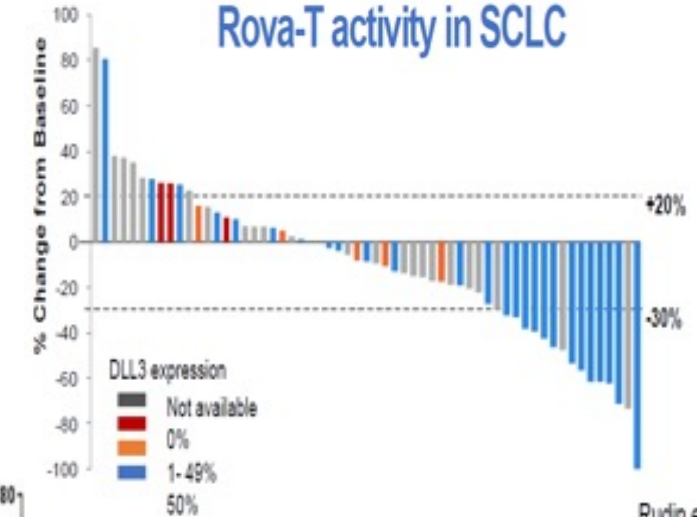
### RNA Seq: Tumor vs Normal



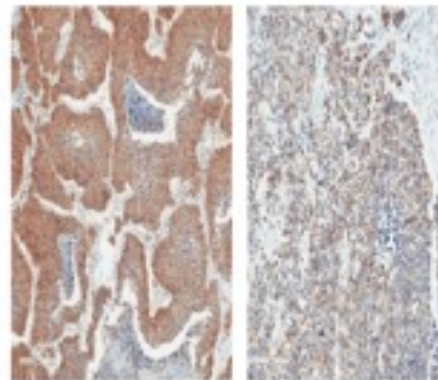
### Rovalpituzumab tesirine (Rova-T™): an ADC targeting DLL3



### Rova-T activity in SCLC



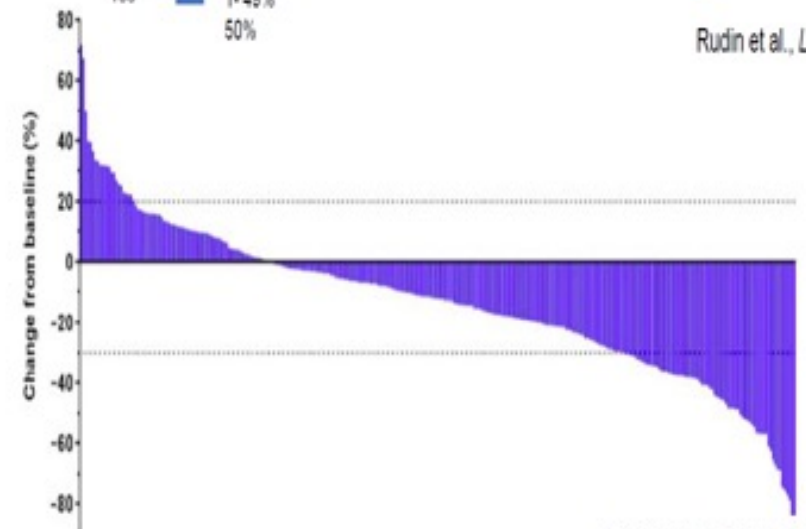
### SCLC Tumors



### Normal Lung



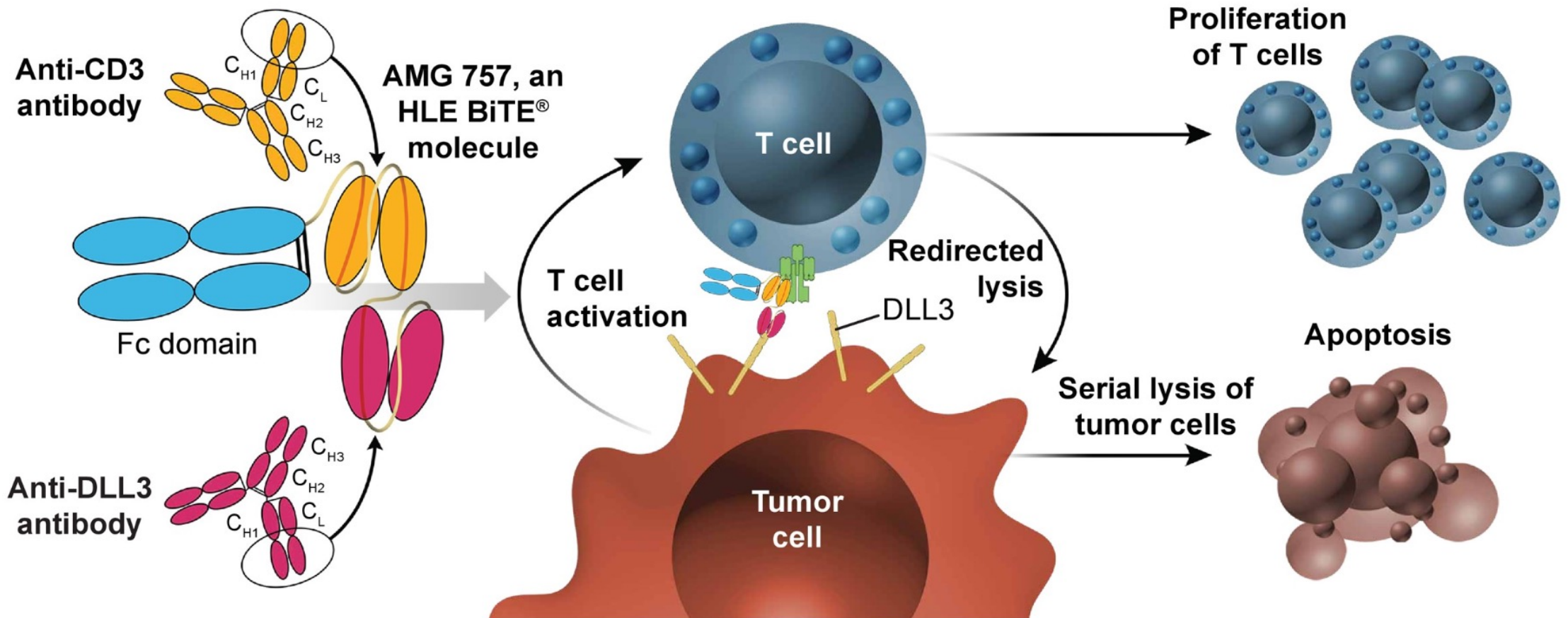
DELTA-LIKE LIGAND 3 (DLL3) IS A HIGHLY SPECIFIC TARGET IN SCLC



# AMG 757

## Half-life Extended DLL3-Directed Bispecific Antibody (BiTE)

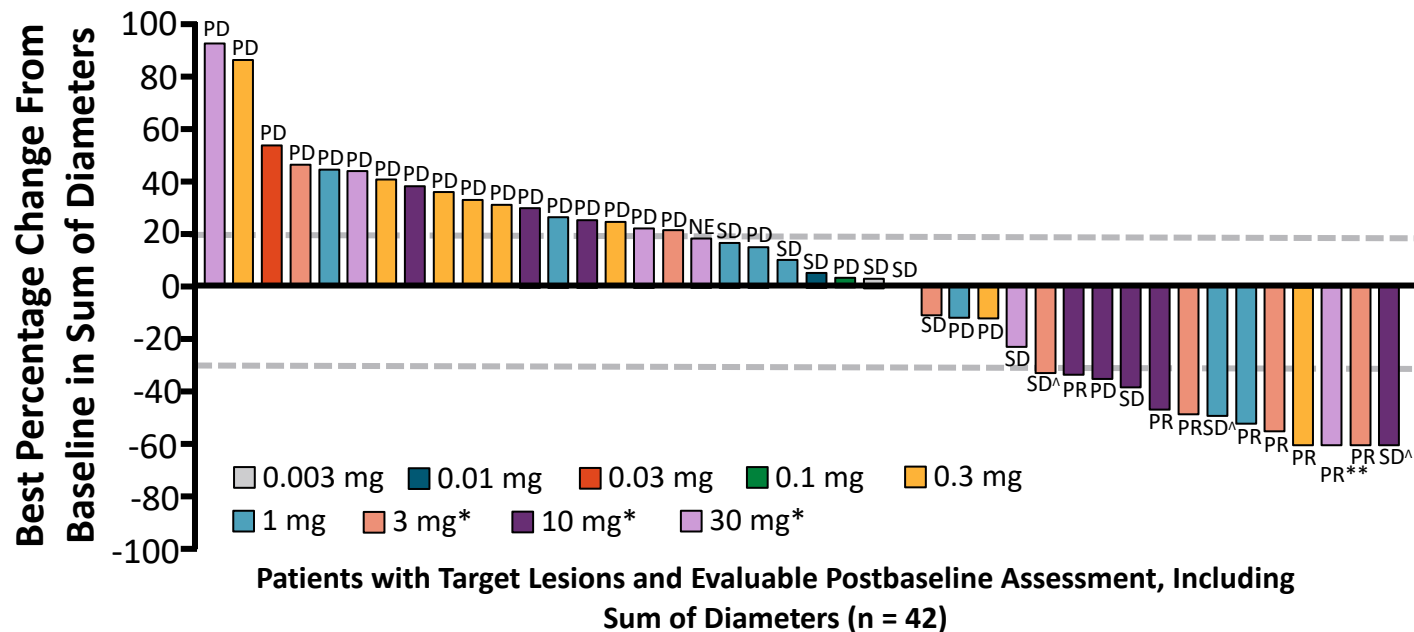
Figure 2. AMG 757 Is a Half-life Extended BiTE® Immuno-oncology Therapy



C<sub>H</sub>, heavy chain constant domain; C<sub>L</sub>, light chain constant domain; HLE BiTE®, half-life extended bispecific T-cell engager; CD, cluster of differentiation; DLL3, delta-like ligand 3; Fc, crystallizable fragment.

# AMG 757 in Relapsed SCLC

## Antitumor Activity



PR\*\*, unconfirmed PR; SD^, initial PR not confirmed on subsequent scan; NE, PD in post-baseline scan and went off study without confirmation scan. \*Step dosing.

†Includes those treated with ≥ 1 dose of AMG 757 and with follow-up ≥ 8 wks.

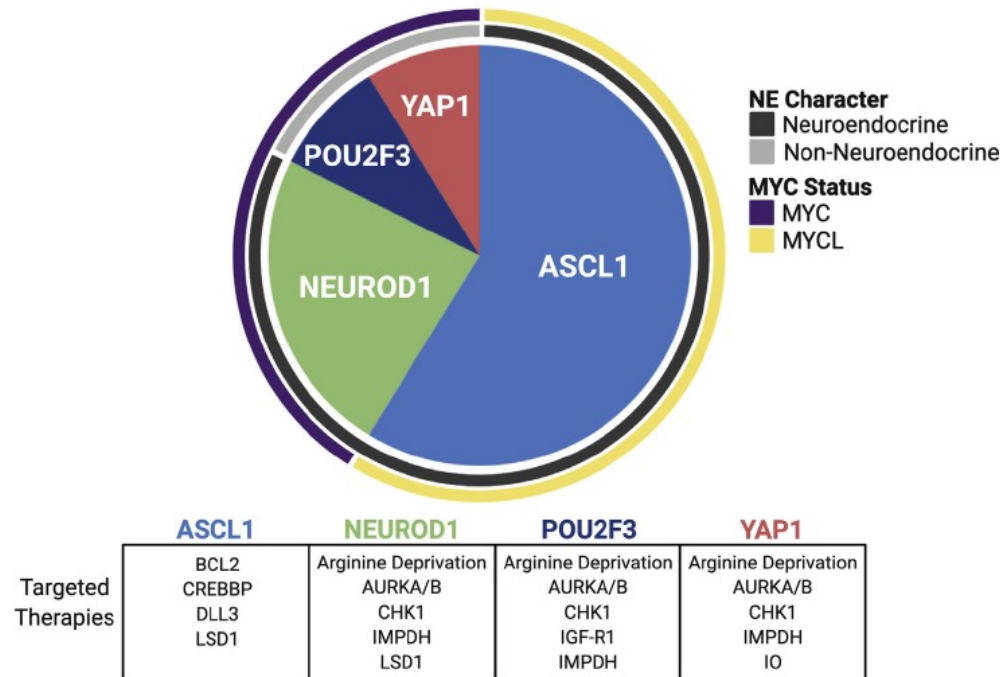
‡At target dose of 30 mg.

Response per mRECIST v1.1	Patients (n = 51 <sup>†</sup> )
Confirmed PR, n (%)	7 (14)
Confirmed PR by target dose, n/N (%)	
▪ 0.3 mg	1/12 (8)
▪ 1.0 mg	1/8 (13)
▪ 3.0 mg	3/9 (33)
▪ 10.0 mg	2/10 (20)
Unconfirmed PR, n (%)	1 (2) <sup>‡</sup>
SD, n (%)	11 (22)
DCR, %	37

- Antitumor activity observed with AMG 757 during dose exploration

# Molecular subtypes of SCLC

## Potential Therapeutic Implications



Data on molecular heterogeneity of SCLC holds promise for biomarker driven personalized therapeutic approaches

Preclinical studies suggest distinct therapeutic vulnerabilities in the novel marker-defined subtypes of SCLC

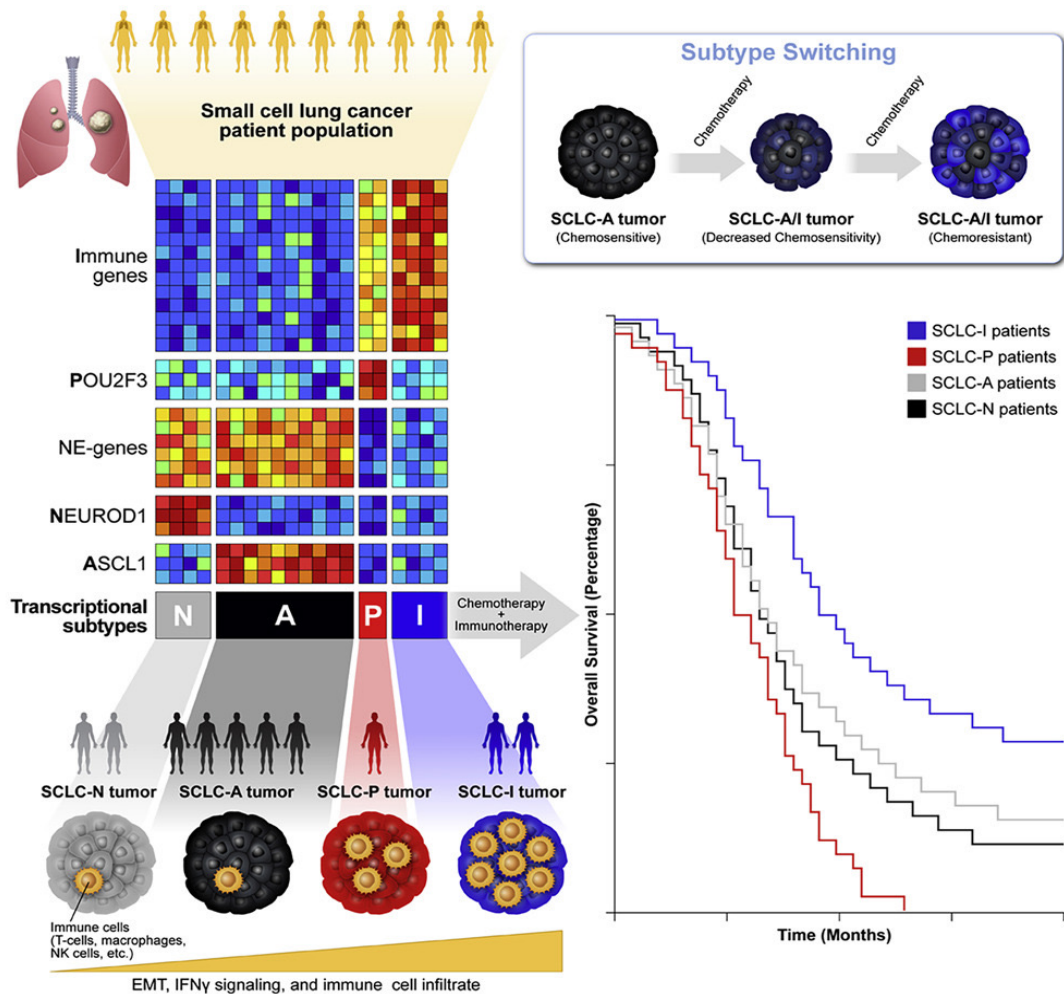
Clinical studies

- Retrospective
- Not prospective yet

**Figure 3.** Diagram of the relative abundance, MYC status, and NE character of the four molecular subtypes of SCLC, each identified by their key transcriptional regulator. These subtypes may exhibit distinct targetable vulnerabilities, which are represented in the table beneath the pie chart. Proportions of each subtype are as follows: ASCL1 (0.70, 95% CI: 0.60-0.79), NEUROD1 (0.11, 95% CI: 0.06-0.20), YAP1 (0.02, 95% CI: 0.01-0.09), POU2F3 (0.16, 95% CI: 0.10-0.26). ASCL1, achaete-scute homolog 1; AURKA/B, Aurora kinase A/B; BCL2, B-cell lymphoma 2; CREBBP, CREB-binding protein; CHK1, checkpoint kinase 1; DLL3, delta-like ligand 3; IMPDH, inosine-5' monophosphate dehydrogenase; IGF-R1, insulin-like growth factor 1 receptor; IO, immuno-oncology; LSD1, lysine-specific histone demethylase 1; NE, neuroendocrine; NEUROD1, neurogenic differentiation factor 1; POU2F3, POU class 2 homeobox 3; YAP1, yes-associated protein 1.

# Molecular subtypes of SCLC

## Potential Biomarkers Implications



- Four subtypes with unique molecular features and therapeutic vulnerabilities
  - Differential expression of ASCL1, NEUROD1, and POU2F3 defines SCLC subtypes
  - An inflamed, mesenchymal, subtype (SCLC-I) has low expression of ASCL1, NEUROD1, and POU2F3
- SCLC-I experiences greatest benefit from the addition of anti-PD-L1 to chemotherapy
- Subtypes with specific genomic alterations
  - All subtypes with similar mutational landscape (including TP53 and RB1)
- Phenotype (genomic, transcriptomic, proteomic)
  - SCLC-A: highly NE, epithelial, TTF-1+
  - SCLC-N: highly NE, largely TTF-1-
  - SCLC-P: non-NE, low EMT
  - SCLC-I: non-NE, high EMT
- Intratumoral subtype switching accompanies acquired resistance to platinum chemotherapy

# Notch Signaling in SCLC

## Determinant of response to ICI

### Notch signaling and efficacy of PD-1/PD-L1 blockade in relapsed small cell lung cancer

- Immunogenomic profiling of relapsed SCLC tumors treated with Immune Checkpoint Blockade (ICB).
- Tumors deriving clinical benefit from ICB exhibited cytotoxic T-cell infiltration, high expression of antigen processing and presentation machinery (APM) genes, and low neuroendocrine (NE) differentiation.
- Notch signaling, (correlates positively with low NE differentiation), most significantly predicts clinical benefit to ICB.
- Activation of Notch signaling (overexpression of NOTCH1 intracellular domain) in a (high) NE human SCLC cell line induces transition to a low NE phenotype
  - marked by increased expression/upregulation of APM genes
- Mechanistic link between Notch activation, low NE differentiation and increased intrinsic tumor immunity.

Roper, et al. Nat Commun. 2021 Jun 23;12(1):3880.

**Table 2 Notch signaling gene set is the most significant predictor of clinical benefit to immune checkpoint blockade across relapsed SCLC cohorts.**

Variable	Estimate <sup>a</sup>	t value <sup>a</sup>	p value <sup>a</sup>	FDR <sup>b</sup>
Hallmark Notch signaling	0.25	4.31	$9.8 \times 10^{-4}$	$5.9 \times 10^{-4}$
Immune signature	0.13	2.06	0.047	0.14
NE score	-0.07	-1.82	0.08	0.16
MYC expression	-0.04	-0.83	0.41	0.62
EZH2 expression	-0.03	-0.56	0.58	0.62

Outcome dependent variable = clinical benefit to immune checkpoint blockade.

<sup>a</sup>Estimates, t and p values calculated using multivariable logistic regression.

<sup>b</sup>False discovery rate was calculated using the Benjamini-Hochberg procedure.

**Table 1 Association between transcriptional subtypes and clinical benefit to immune checkpoint blockade across relapsed SCLC cohorts.**

Transcriptional subtype	Clinical benefit (# of tumors)	No clinical benefit (# of tumors)	p value <sup>a</sup>
ASCL1	2	22	0.11
NEUROD1	3	9	0.40
POU2F3	1	0	0.17
YAP1	2	8	0.78

<sup>a</sup>Statistical significance calculated using the two-tailed chi-squared test.