

I° CONVEGNO REGIONALE SIFO

Meeting di primavera

“IL FARMACISTA CLINICO  
E I NUOVI MODELLI DI CURA”



TAORMINA (ME),  
11-12-13 MAGGIO 2017

ID ECM: 313-188877



IV SESSIONE - IMMUNONCOLOGIA

## I biomarcatori nell'immunoncologia: L'importanza della selezione del paziente

*Hector Soto Parra MD  
Oncologia Medica*



# I biomarcatori nell'immunoncologia: Perché è importante la selezione del paziente?

- **Migliorare il risultato terapeutico** **-Efficacia-**
  - pochi pz/grandi benefici vs molti pz/pochi vantaggi
- **Contenere i costi** **-Farmacoeconomia-**
- **Controllo adeguatezza prescrittiva** **-Ente regolatorio-**
- **Disegno di nuovi studi** **-Ricerca-**

# Tumor and Immune Biomarkers Being Evaluated to Predict Better Outcomes to Immuno-Oncology Therapy

## Tumor Antigens

- Biomarkers indicative of hypermutation & neo-antigens may predict response to IO treatment

### Examples:

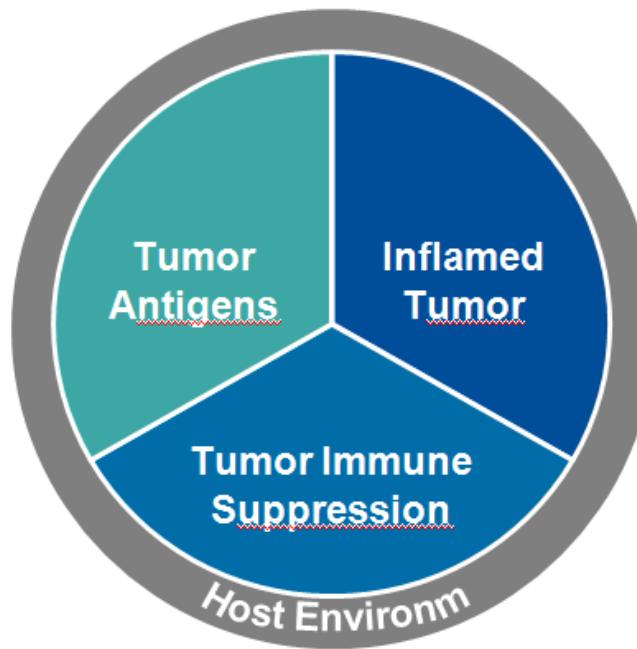
- *TMB, MSI-High, Neo-Antigens*

## Tumor Immune Suppression

- Biomarkers that identify tumor immune system evasion beyond PD-1/CTLA-4 to inform new IO targets and rational combinations

### Examples:

- *Tregs, MDSCs, IDO, LAG-3*



## Inflamed Tumor Microenvironment

- Biomarkers (intra- or peri-tumoral) indicative of an inflamed phenotype may predict response to IO treatment

### Examples:

- *PD-L1, Inflammatory Signatures*

## Host Environment

- Biomarkers which characterize the host environment, beyond tumor microenvironment, may predict response to IO treatment

### Examples:

- *Microbiome, Germline Genetics*

IDO = indoleamine-2,3 dioxygenase; LAG-3 = lymphocyte activation gene-3; MDSCs = myeloid-derived suppressor cells; MSI-High = microsatellite instability high; TMB = tumor mutational burden. Adapted from Blank C.U. et al., *Science* 2016;352:658–660.

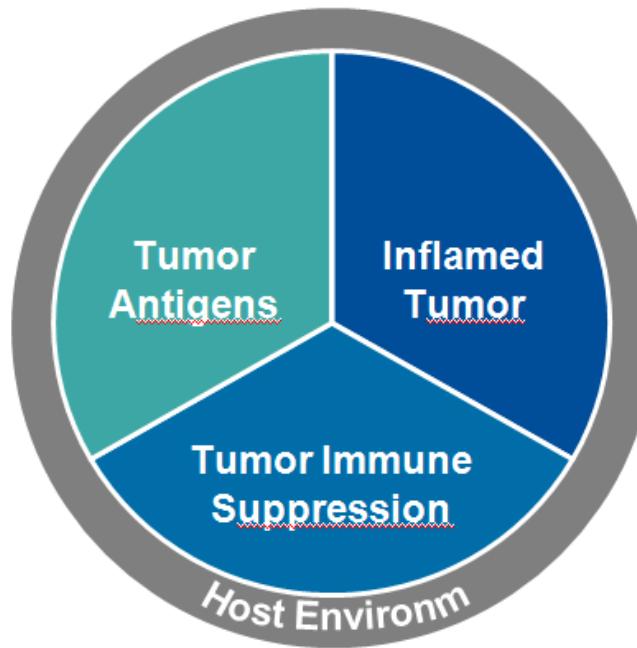
# Tumor and Immune Biomarkers Being Evaluated to Predict Better Outcomes to Immuno-Oncology Therapy

## Tumor Antigens

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### Examples:

- *TMB, MSI-High, Neo-Antigens*



## Inflamed Tumor Microenvironment

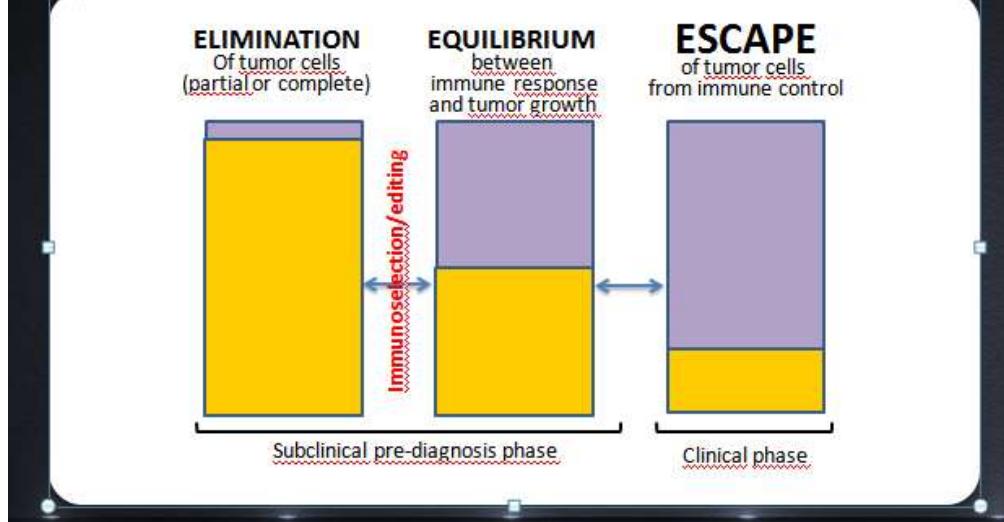
- Biomarkers (intra- or peri-tumoral) indicative of an inflamed phenotype may predict response to IO treatment

### Examples:

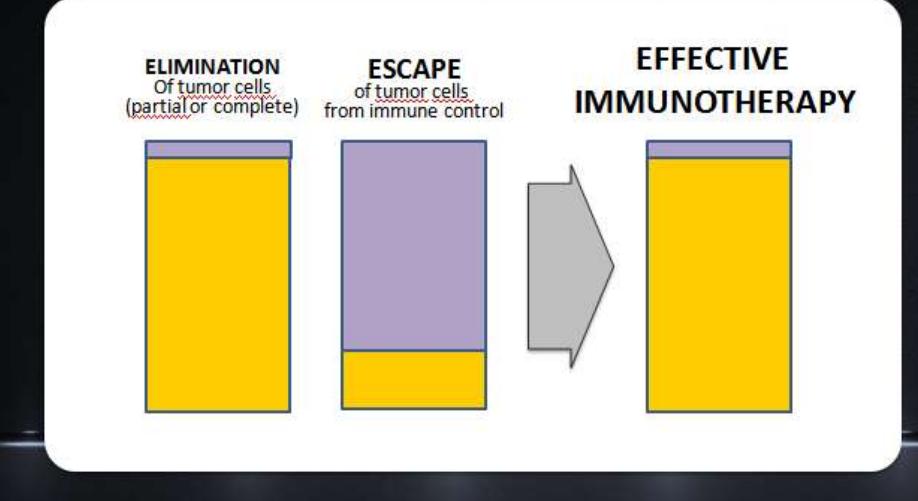
- *PD-L1, Inflammatory Signatures*

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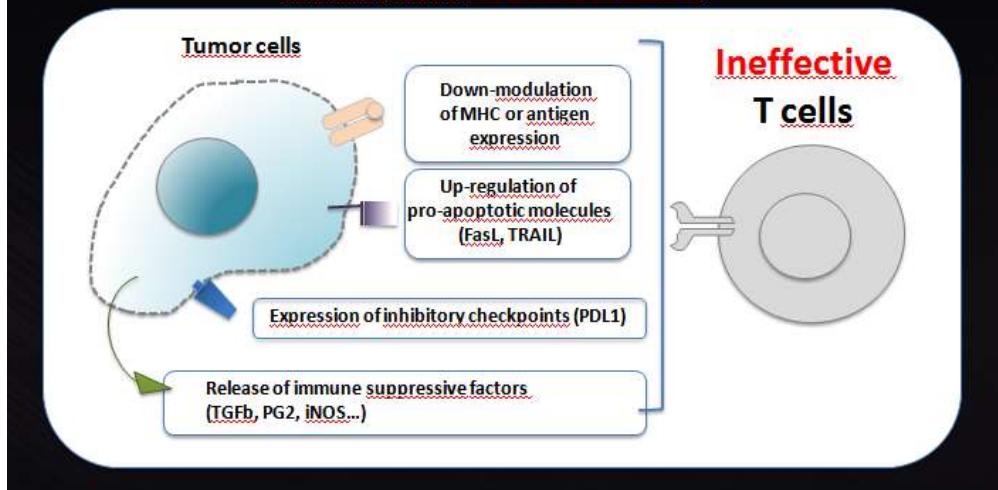
# Tumor immunity: a dynamic interaction



## CANCER IMMUNOTHERAPY: tilt the balance to immune tumor control

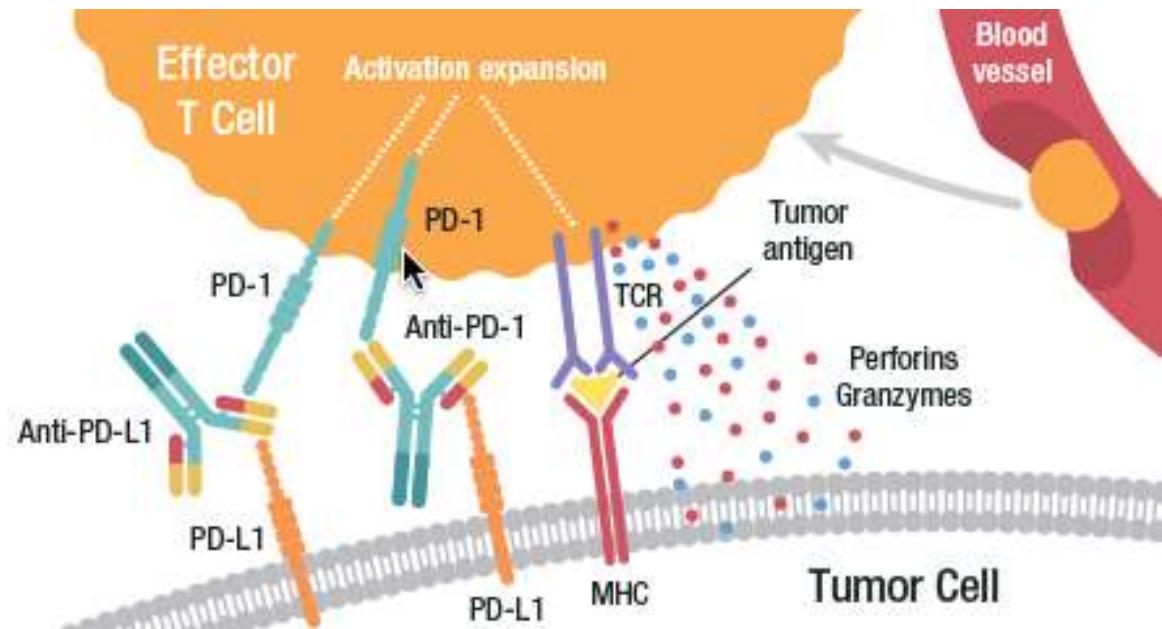


## Tumor immune escape mechanisms: tumor cells counterattack



**Anti-PD1**  
Nivolumab  
Pembrolizumab

**Anti-PDL1**  
Atezolizumab  
Durvalumab  
Avelumab



# Solid Tumors

## Anti-PD1/PDL1 FDA Approved

### Anti-PD1

**Nivolumab:** Melanoma, NSCLC, Renal, Bladder, H&N

**Pembrolizumab:** Melanoma, **NSCLC\***, H&N

### Anti-PDL1

**Avelumab:** Merkel cell carcinoma

**Atezolizumab:** NSCLC, Urothelial

**Durvalumab:** Urothelial

\* non-small cell lung cancer (NSCLC) whose tumors express PD-L1 as determined by an FDA-approved test.

# Randomised Trials of Anti PD-1 / anti-PD-L1 Agents

Study	Line	Agents	PD-L1	Result	HR
CheckMate 026	1 <sup>st</sup>	Nivo vs Chemo	≥1%	No difference in PFS, OS	1.15, 1.02
KEYNOTE-024	1 <sup>st</sup>	Pembro vs Chemo	>50%	Improved OS	0.60
CheckMate 017	2 <sup>nd</sup>	Nivo vs Docetaxel	Unselected	Improved OS	0.59
CheckMate 057	2 <sup>nd*</sup>	Nivol vs Docetaxel	Unselected	Improved OS	0.73
KEYNOTE-010	2 <sup>nd*</sup>	Pembro vs Docetaxel	>1%	Improved OS (2, 10 mg/kg)	0.61, 0.71
POPLAR	2 <sup>nd</sup> or 3 <sup>rd</sup>	Atezo vs Docetaxel	Unselected	Improved OS	0.73
OAK	2 <sup>nd</sup> or 3 <sup>rd</sup>	Atezo vs Docetaxel	Unselected	Improved OS	0.73

\*3<sup>rd</sup> line for EGFR or ALK +ve tumours

Borghaei et al. N Engl J Med 2015; 373: 1627-39. Brahmer et al. N Engl J Med 2015; 373: 123 – 35.

Herbst et al. Lancet 2016; 387: 1540 – 50. Fehrenbacher et al. Lancet 2016; 387: 1837 – 46.

Barlesi et al. ESMO 2016 LBA44. Socinski et al. ESMO 2016

Reck et al. N Engl J Med 2016; 375: 1823 - 33

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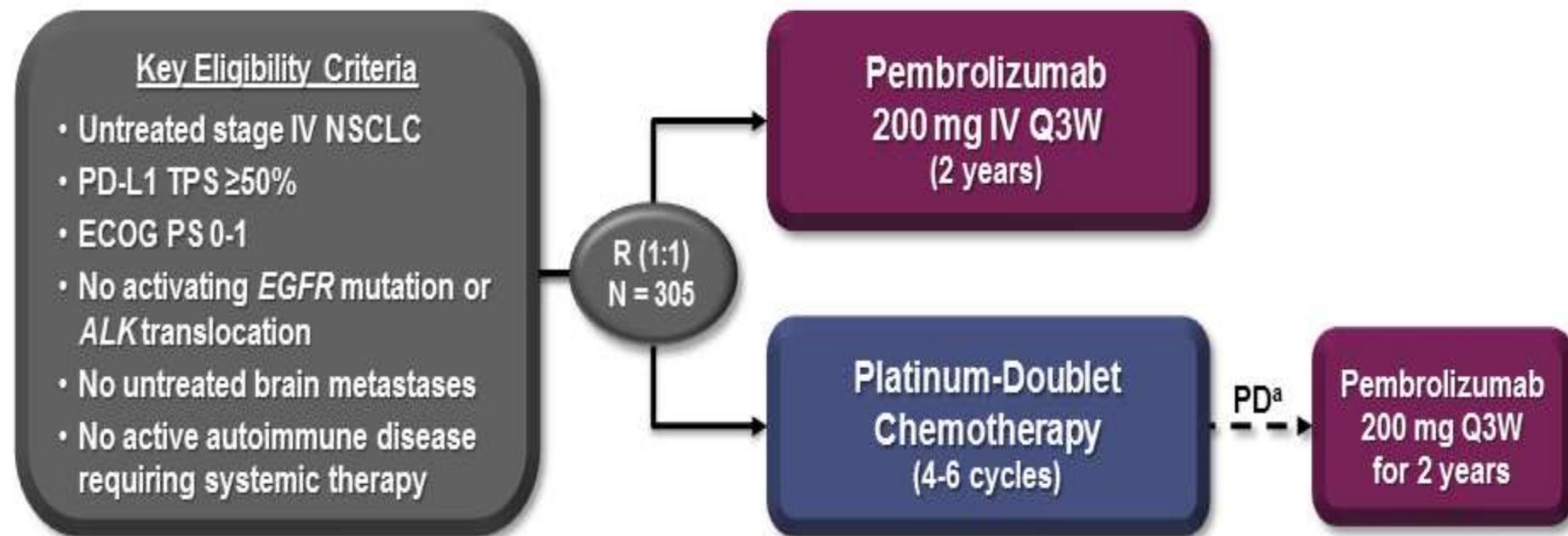
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Barlesi et al. ESMO 2016 LBA44. Socinski et al. ESMO 2016

Reck et al. N Engl J Med 2016; 375: 1823 - 33

# KEYNOTE-024 Study Design (NCT02142738)



## Key End Points

Primary: PFS (RECIST v1.1 per blinded, independent central review)

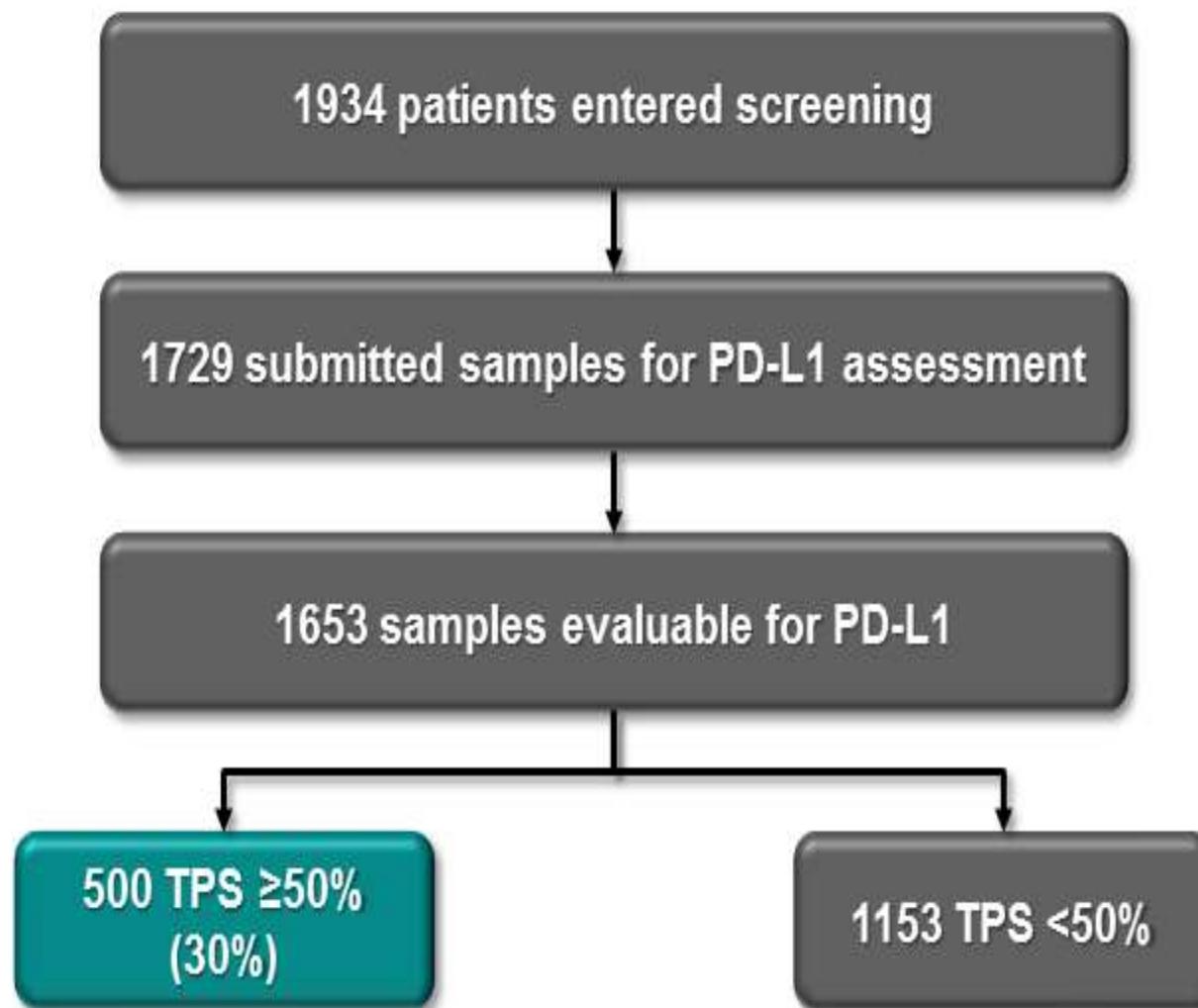
Secondary: OS, ORR, safety

Exploratory: DOR

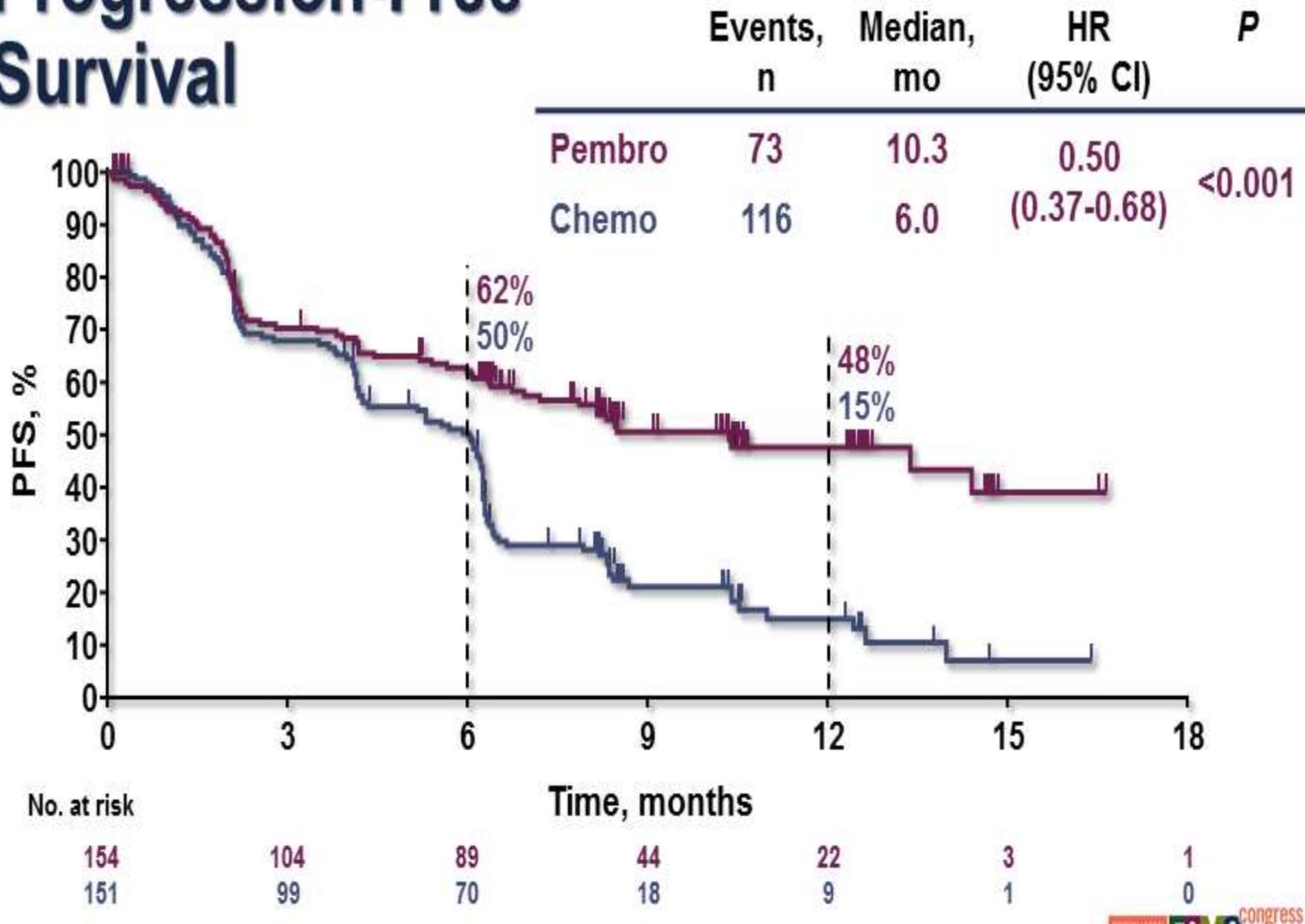
Median duration follow up: 11.2 months

<sup>a</sup>To be eligible for crossover, progressive disease (PD) had to be confirmed by blinded, independent central radiology review and all safety criteria had to be met.

# PD-L1 Screening

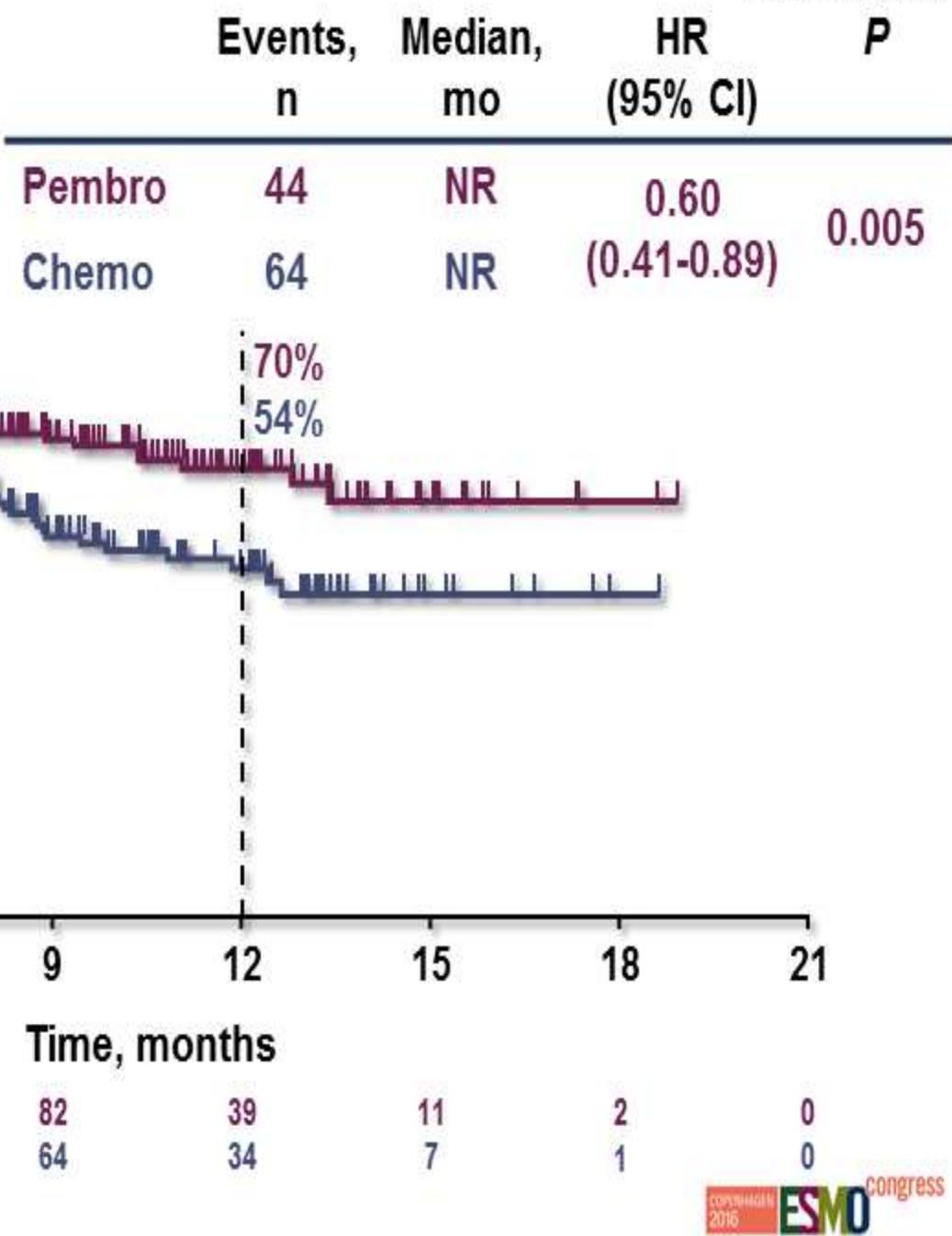


# Progression-Free Survival

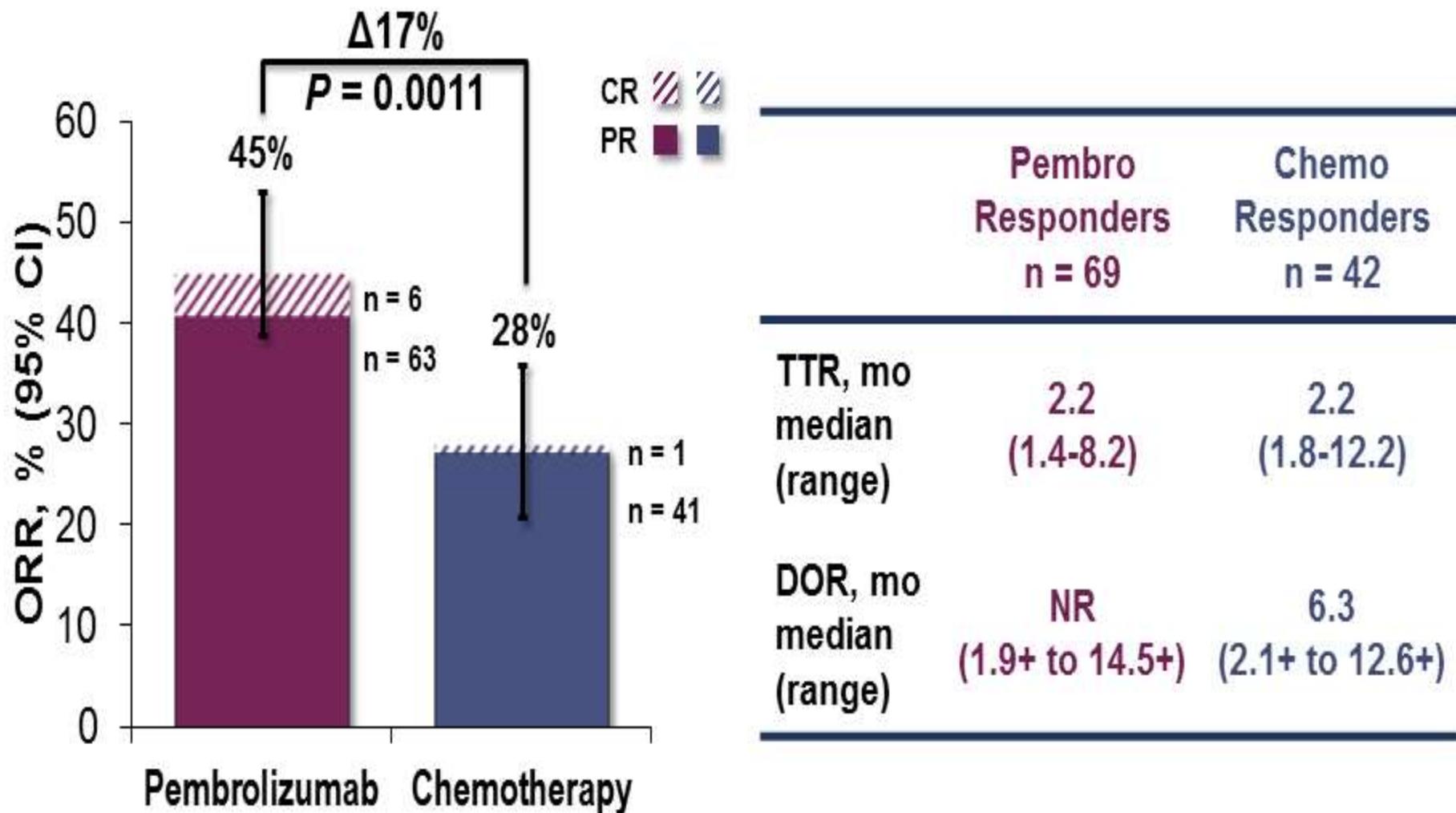


Assessed per RECIST v1.1 by blinded, independent central review.  
Data cut-off: May 9, 2016.

# Overall Survival

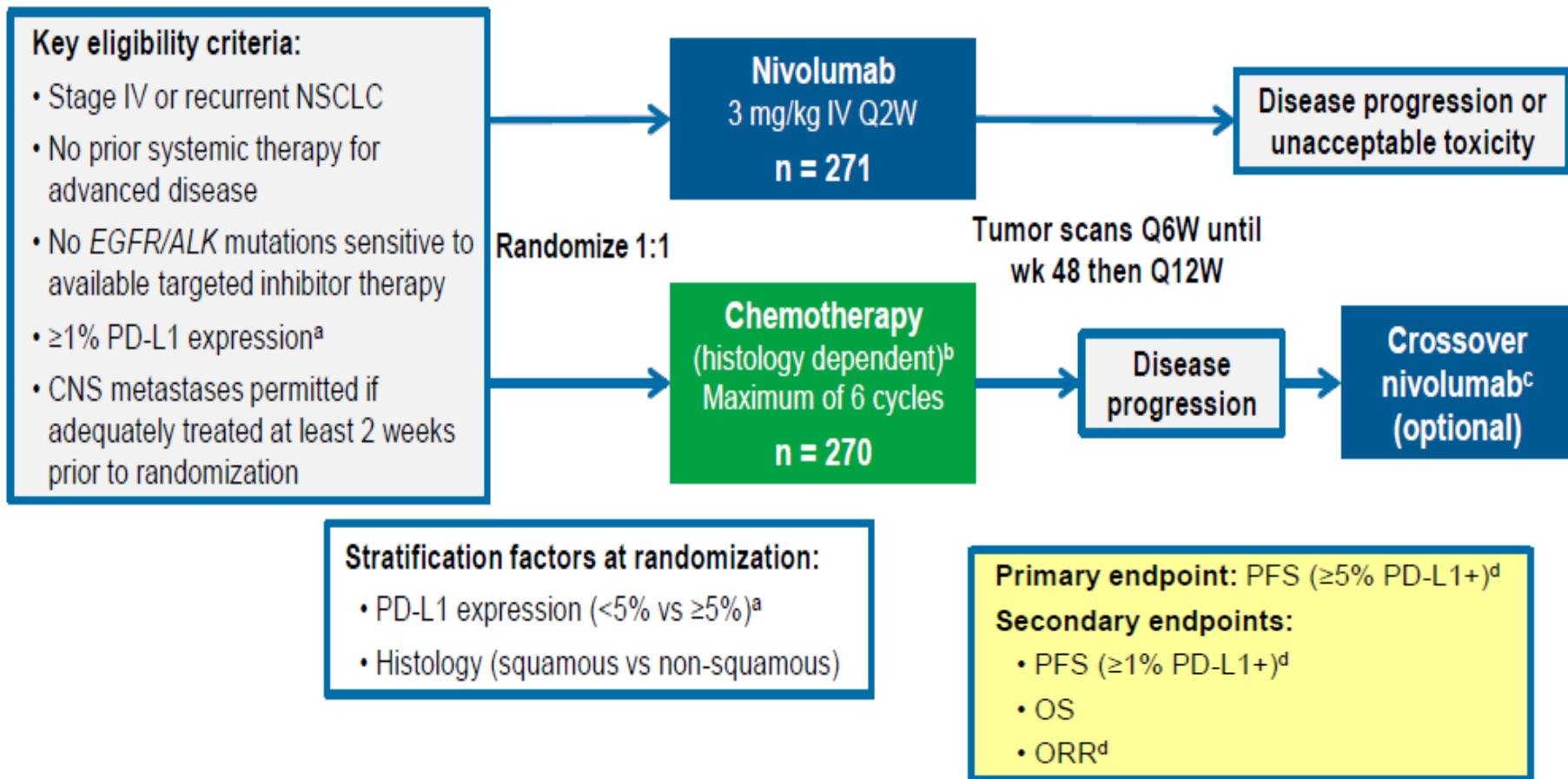


# Confirmed Objective Response Rate



Assessed per RECIST v1.1 by blinded, independent central review.  
Data cut-off: May 9, 2016.

# Phase 3 CheckMate 026 Study Design: Nivolumab vs Chemotherapy in First-line NSCLC



<sup>a</sup>Dako 28-8 validated; archival tumor samples obtained ≤6 months before enrollment were permitted; PD-L1 testing was centralized

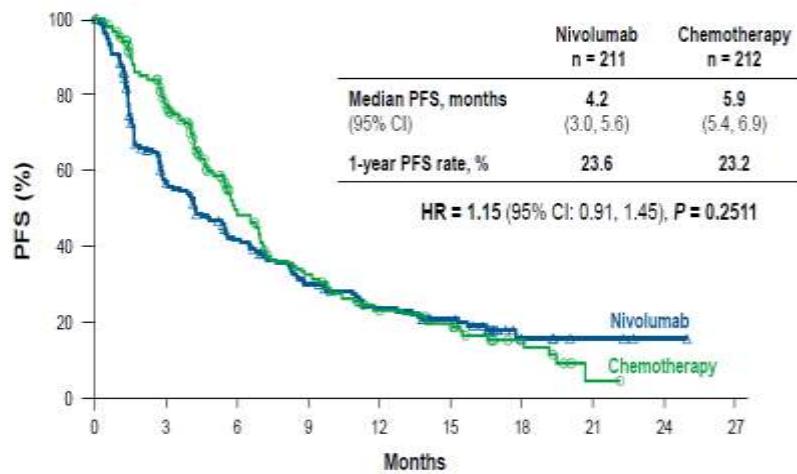
<sup>b</sup>Squamous: gemcitabine 1250 mg/m<sup>2</sup> + cisplatin 75 mg/m<sup>2</sup>; gemcitabine 1000 mg/m<sup>2</sup> + carboplatin AUC 5; paclitaxel 200 mg/m<sup>2</sup> + carboplatin AUC 6;  
Non-squamous: pemetrexed 500 mg/m<sup>2</sup> + cisplatin 75 mg/m<sup>2</sup>; pemetrexed 500 mg/m<sup>2</sup> + carboplatin AUC 6; option for pemetrexed maintenance therapy

<sup>c</sup>Permitted if crossover eligibility criteria met, including progression confirmed by independent radiology review

# PFS, OS & ORR

## Primary Endpoint (PFS per IRRC in $\geq 5\%$ PD-L1+)

CheckMate 026: Nivolumab vs Chemotherapy in First-line NSCLC



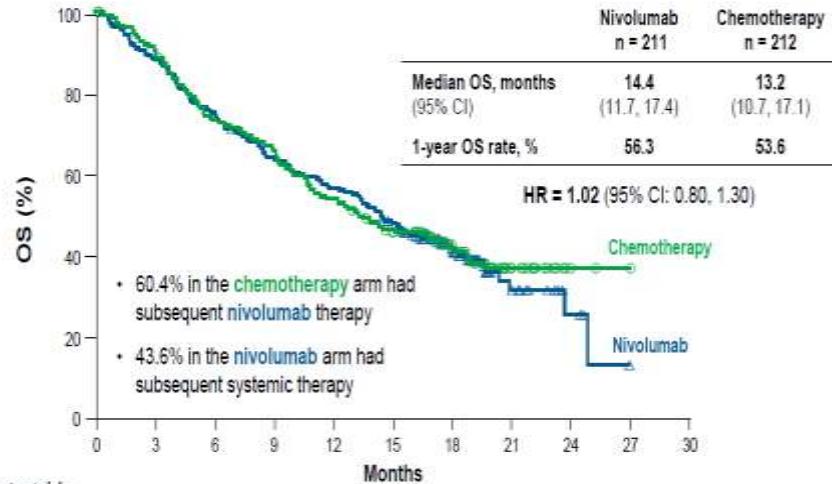
No. of patients at risk:

Nivolumab	211	104	71	49	35	24	6	3	1	0
Chemotherapy	212	144	74	47	28	21	8	1	0	0

All randomized patients ( $\geq 1\%$  PD-L1+): HR = 1.17 (95% CI: 0.95, 1.43)

## OS ( $\geq 5\%$ PD-L1+)

CheckMate 026: Nivolumab vs Chemotherapy in First-line NSCLC



No. of patients at risk:

Nivolumab	211	186	156	133	118	98	49	14	4	0	0
Chemotherapy	212	186	153	137	112	91	50	15	3	1	0

All randomized patients ( $\geq 1\%$  PD-L1+): HR = 1.07 (95% CI: 0.86, 1.33)

**Nivolumab  
(n = 211)**

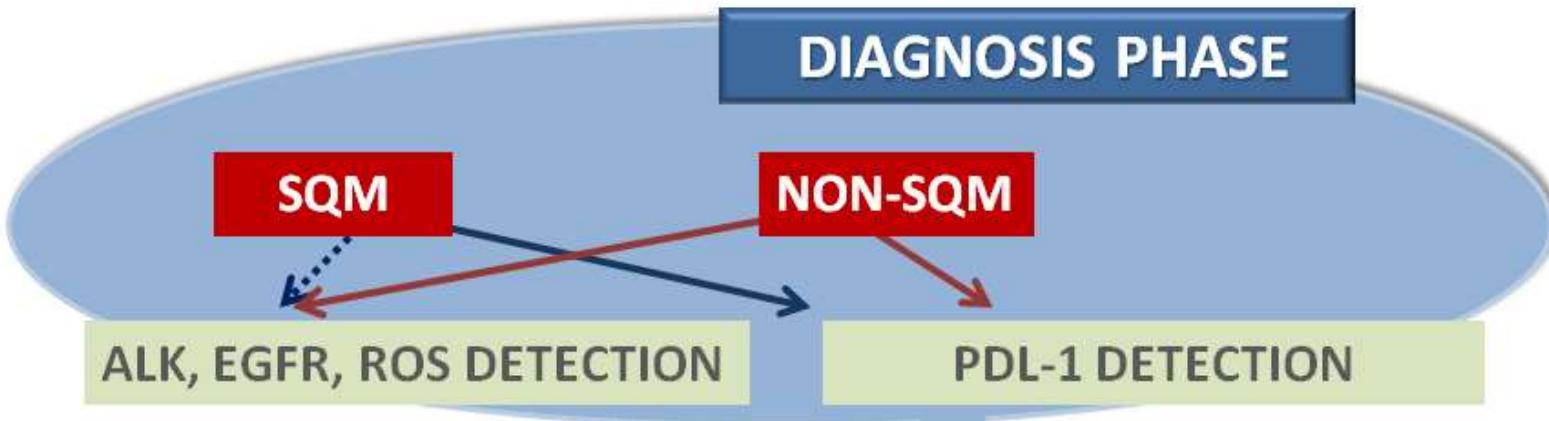
**Chemotherapy  
(n = 212)**

**ORR, % (95% CI)**

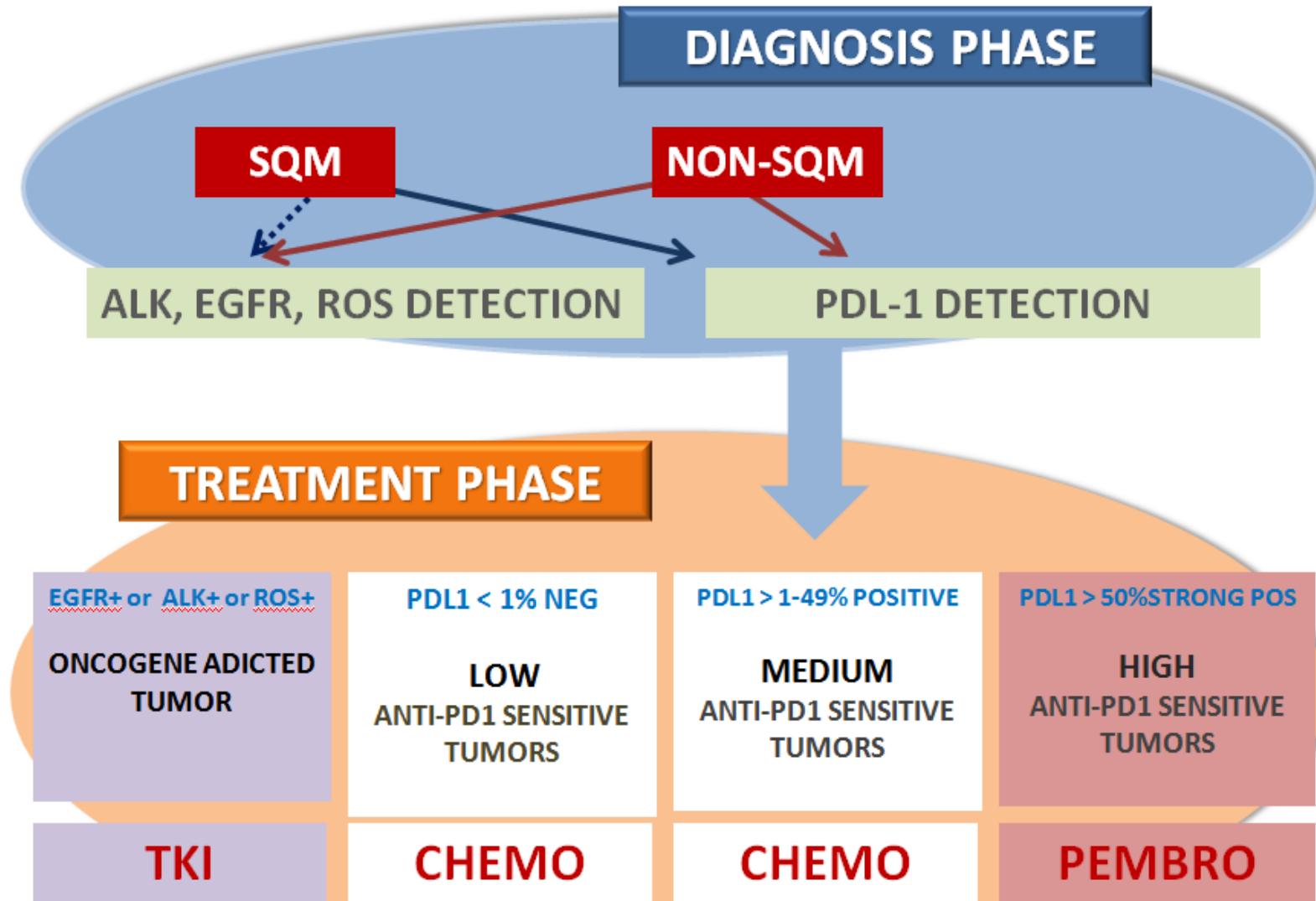
26.1 (20.3, 32.5)

33.5 (27.2, 40.3)

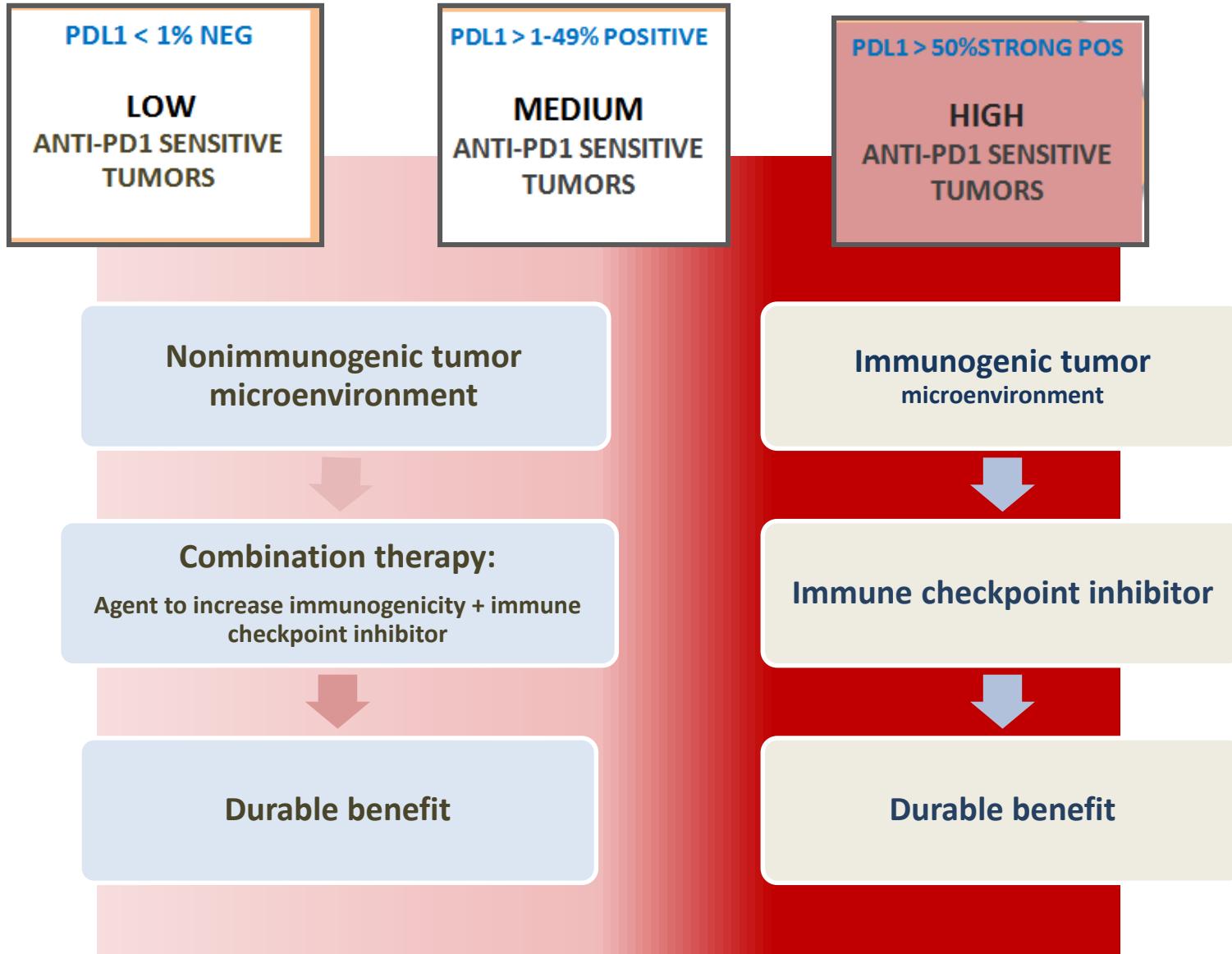
# 1° line NSCLC DX/TX Algorithm



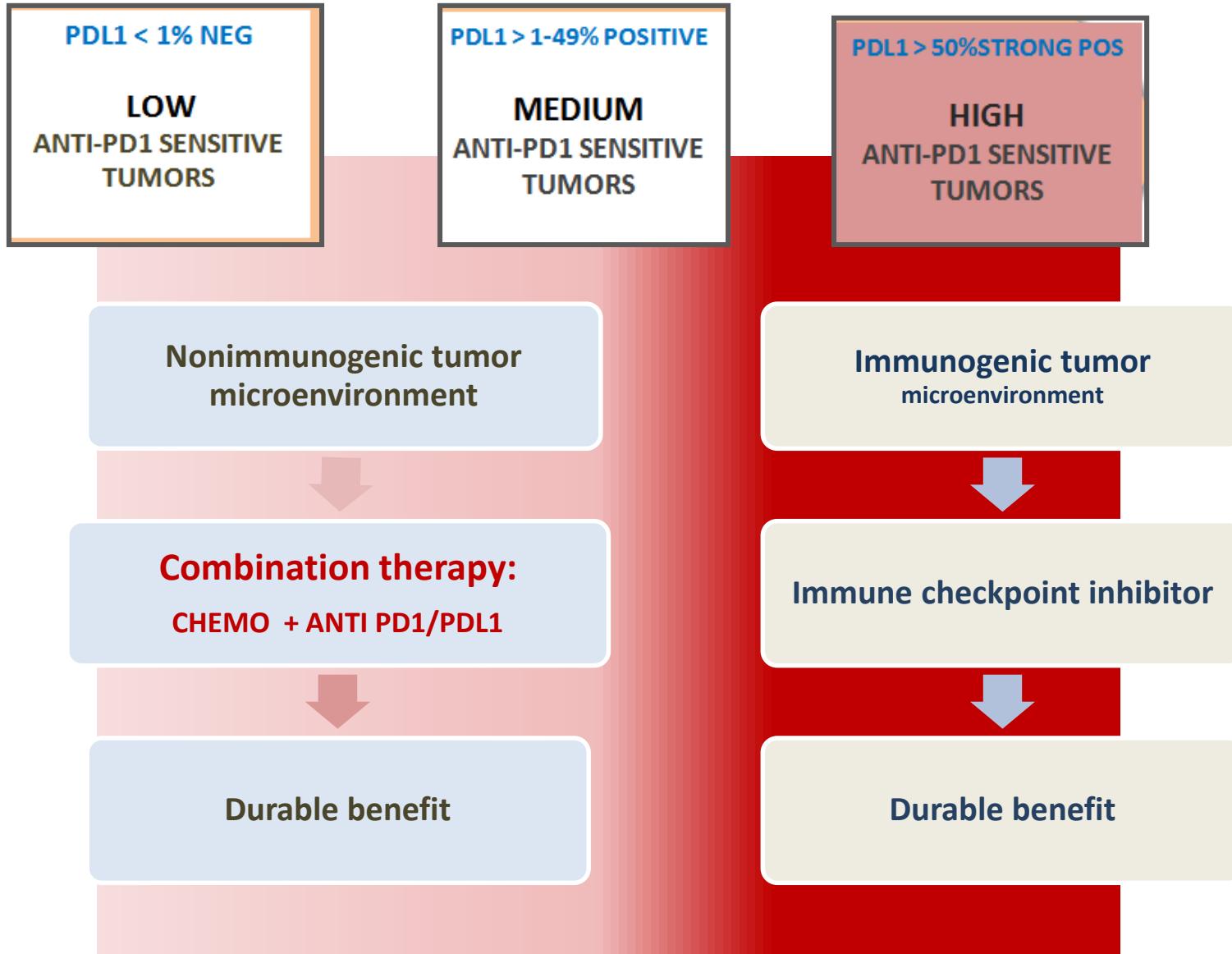
# 1° line NSCLC DX/TX Algorithm



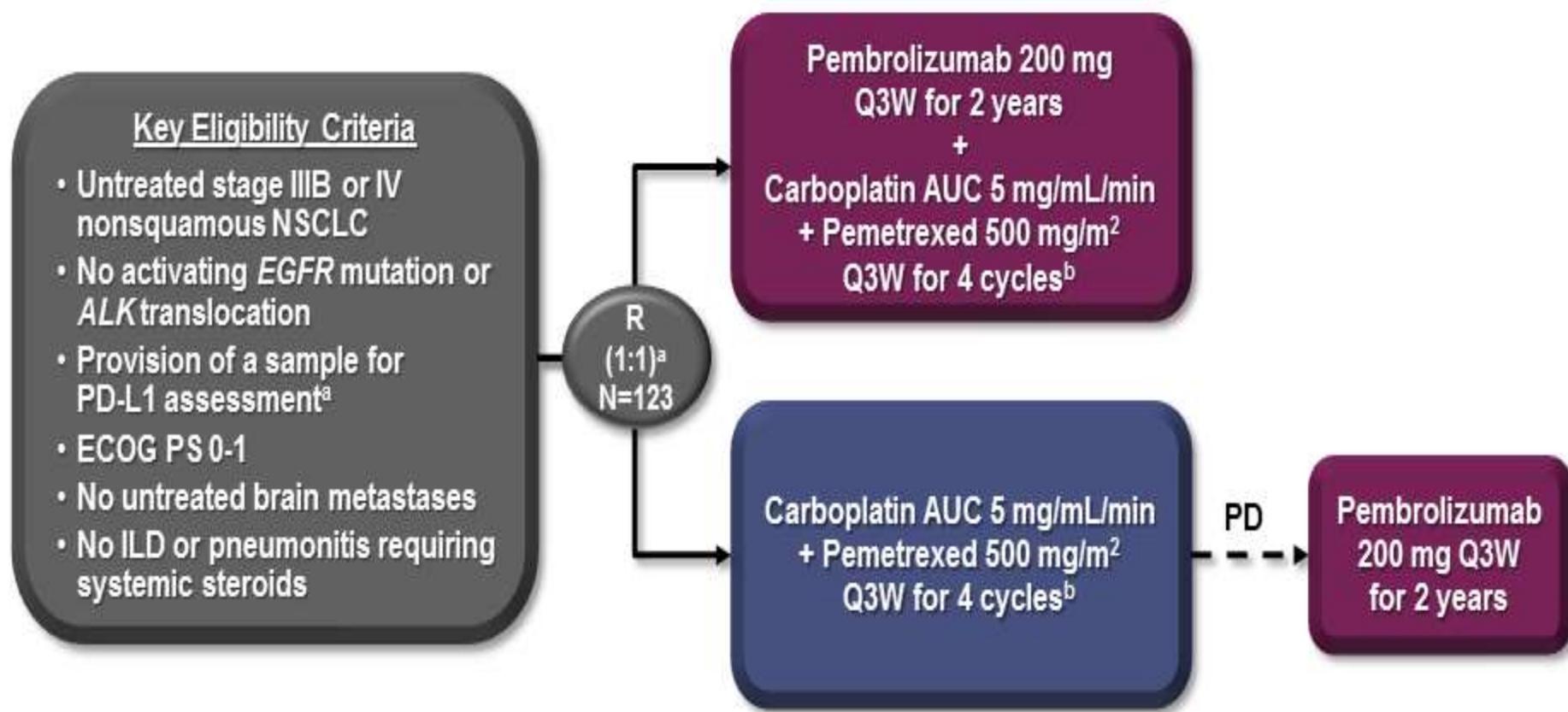
# 1° line NSCLC IO CLINICAL RESEARCH



# 1° line NSCLC IO CLINICAL RESEARCH



# KEYNOTE-021 Cohort G



## End Points

Primary: ORR (RECIST v1.1 per blinded, independent central review)

Key secondary: PFS

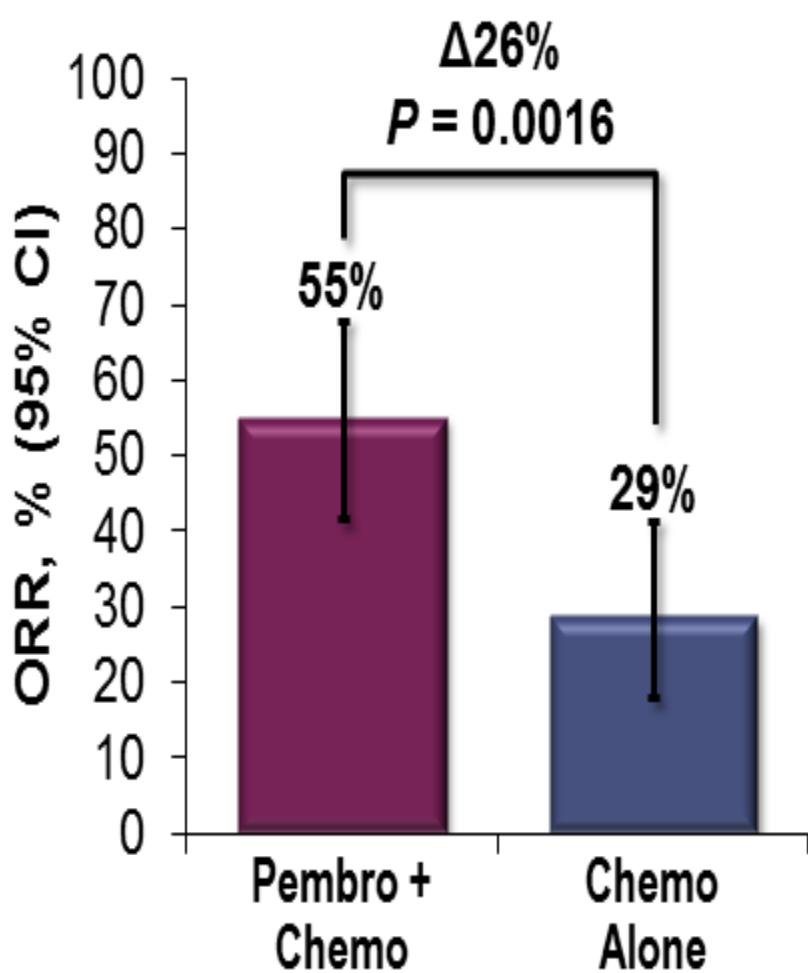
Other secondary: OS, safety, relationship between antitumor activity and PD-L1 TPS

PD=progressive disease.

<sup>a</sup>Randomization was stratified by PD-L1 TPS <1% vs ≥1%.

<sup>b</sup>Indefinite maintenance therapy with pemetrexed 500 mg/m<sup>2</sup> Q3W permitted.

# Confirmed Objective Response Rate (RECIST v1.1 by Blinded, Independent Central Review)



	Pembro + Chemo Responders n = 33	Chemo Alone Responders n = 18
TTR, mo median (range)	1.5 (1.2-12.3)	2.7 (1.1-4.7)
DOR, mo median (range)	NR (1.4+ - 13.0+)	NR (1.4+ - 15.2+)
Ongoing response, <sup>a</sup> n (%)	29 (88)	14 (78)

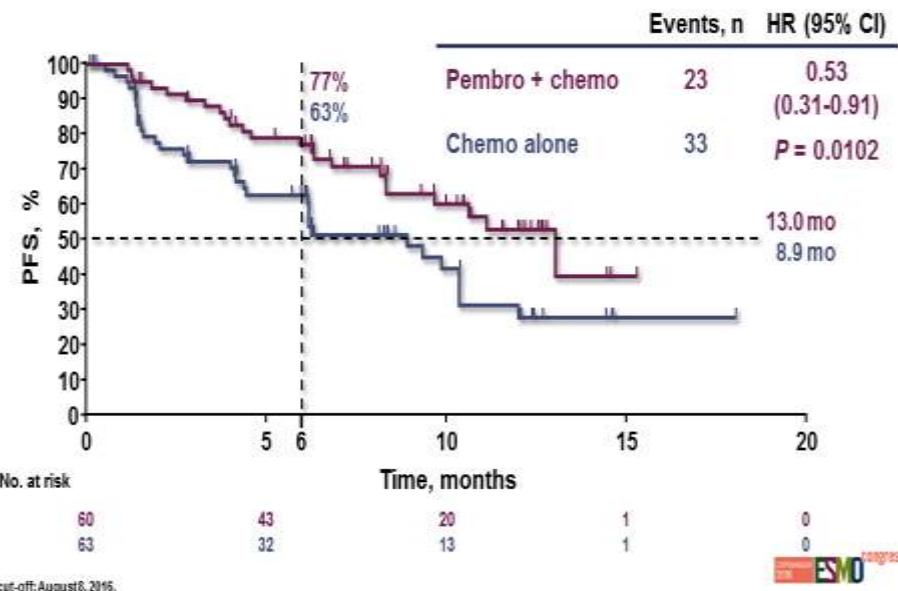
DOR = duration of response; TTR = time to response.

<sup>a</sup>Alive without subsequent disease progression.

# PFS&OS

## Progression-Free Survival (RECIST v1.1 by Blinded, Independent Central Review)

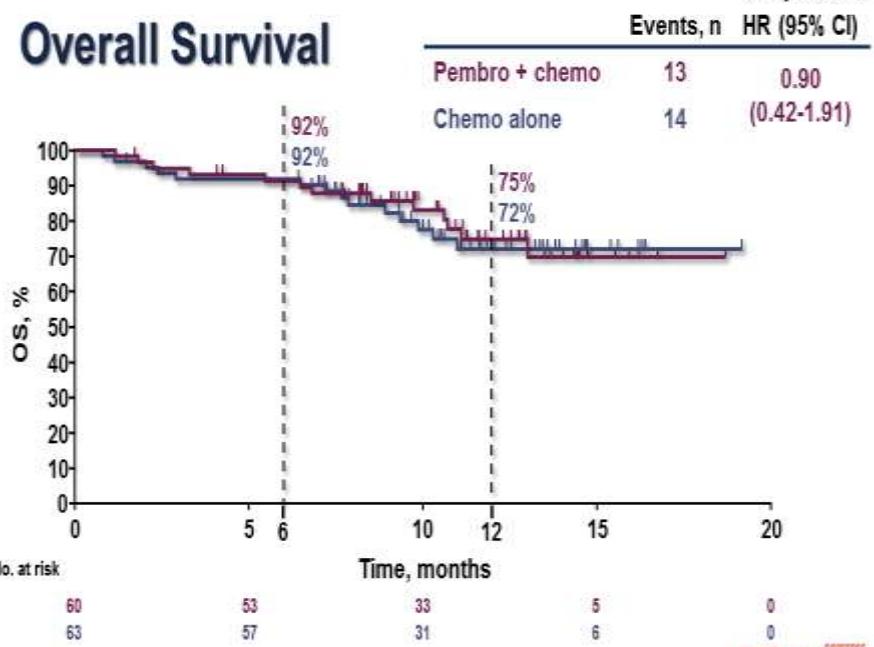
CJ Langer ESMO 2016



CJ Langer ESMO 2016.

## Overall Survival

CJ Langer ESMO 2016



# First-Line Nivolumab Monotherapy and Nivolumab Plus Ipilimumab in Patients With Advanced NSCLC: Long-Term Outcomes From CheckMate 012

Scott N. Gettinger,<sup>1</sup> Naiyer Rizvi,<sup>2</sup> Laura Q. Chow,<sup>3</sup> Hossein Borghaei,<sup>4</sup> Julie Brahmer,<sup>5</sup> Frances A. Shepherd,<sup>6</sup> Neal E. Ready,<sup>7</sup> David E. Gerber,<sup>8</sup> Scott J. Antonia,<sup>9</sup> Jonathan W. Goldman,<sup>10</sup> Rosalyn Juergens,<sup>11</sup> William J. Geese,<sup>12</sup> Tina C. Young,<sup>12</sup> Xuemei Li,<sup>12</sup> Matthew D. Hellmann<sup>2</sup>

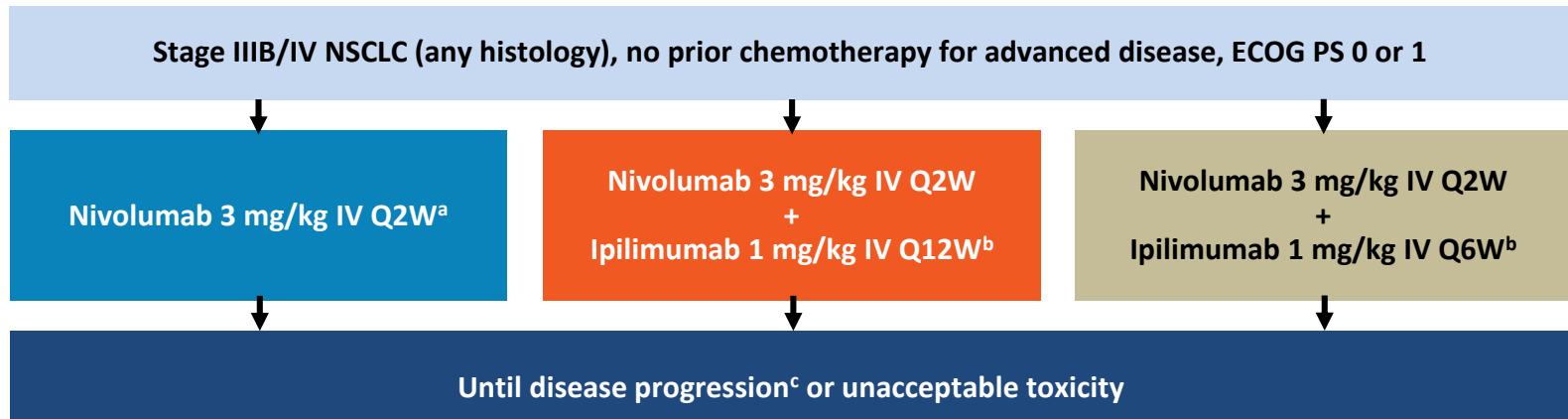
<sup>1</sup>Yale Cancer Center, New Haven, CT, USA; <sup>2</sup>Memorial Sloan Kettering Cancer Center, New York, NY, USA; <sup>3</sup>University of Washington, Seattle, WA, USA;

<sup>4</sup>Fox Chase Cancer Center, Philadelphia, PA, USA; <sup>5</sup>The Sidney Kimmel Comprehensive Cancer Center at Johns Hopkins, Baltimore, MD, USA; <sup>6</sup>Princess Margaret Cancer Centre, Toronto, Canada; <sup>7</sup>Duke University Medical Center, Durham, NC, USA; <sup>8</sup>University of Texas Southwestern Medical Center, Dallas, TX, USA;

<sup>9</sup>H. Lee Moffitt Cancer Center and Research Institute, Tampa, FL, USA; <sup>10</sup>University of California, Los Angeles, Los Angeles, CA, USA;

<sup>11</sup>Juravinski Cancer Centre, McMaster University, Hamilton, ON, Canada; <sup>12</sup>Bristol-Myers Squibb, Princeton, NJ, USA

# Phase 1 CheckMate 012 Study Design: First-Line Nivolumab ± Ipilimumab in NSCLC



**Primary endpoint:** safety and tolerability

**Secondary endpoints:** ORR (RECIST v1.1) and PFS rate at 24 weeks assessed by investigators

**Exploratory endpoints:** OS, efficacy by PD-L1 expression

- Updated data<sup>d</sup> presented here are based on median follow-up durations of 22 months (monotherapy) and 16 months (combination cohorts)
  - Overall additional follow-up relative to previous reports: monotherapy, +~18 months;<sup>1</sup> combination cohorts, +6 months<sup>2</sup>

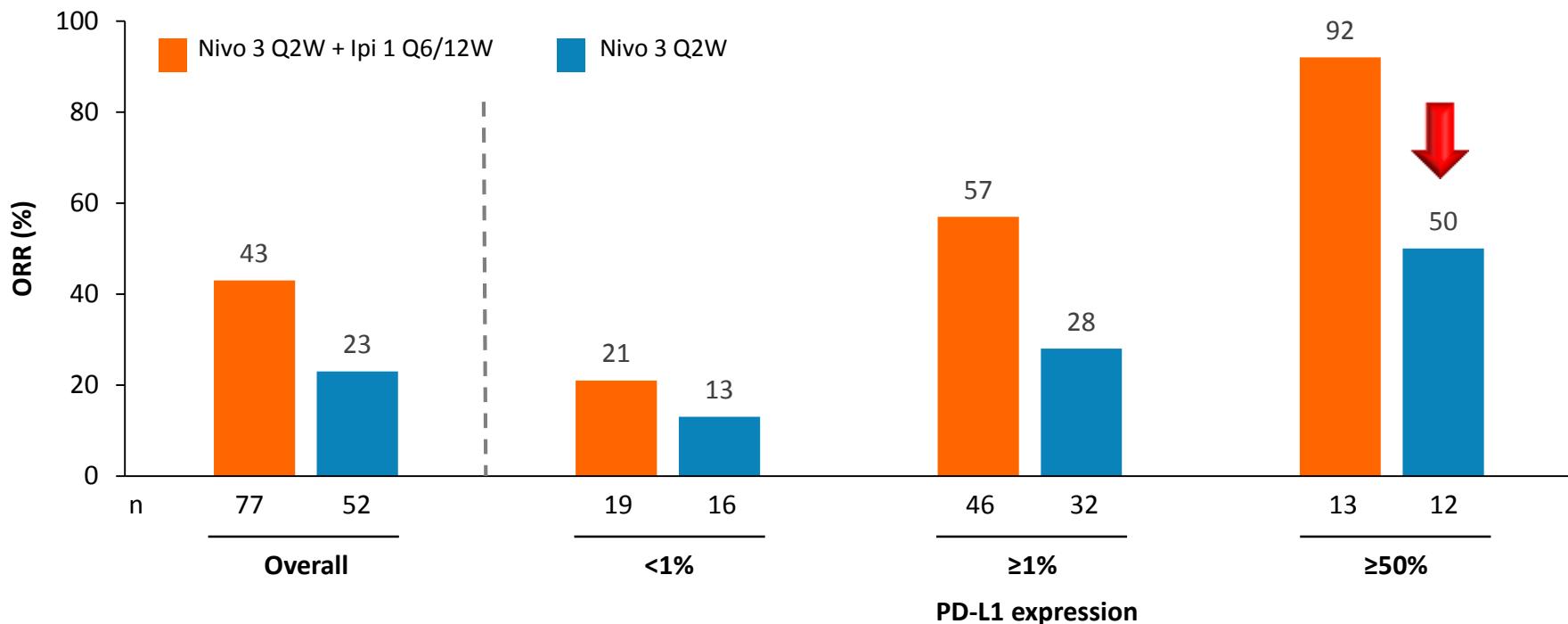
ClinicalTrials.gov number NCT01454102; <sup>a</sup>Treatment allocation not randomized; <sup>b</sup>Treatment allocation randomized; earlier cohorts evaluated other dosing schedules/regimens<sup>2</sup> <sup>c</sup>Patients tolerating study treatment permitted to continue treatment beyond RECIST v1.1-defined progression if considered to be deriving clinical benefit

<sup>d</sup>Based on a September 2016 database lock

1. Gettinger S, et al. *J Clin Oncol* 2016;34:2980–2987; 2. Hellmann MD, et al. *Lancet Oncol* 2016 Dec 5. [Epub ahead of print].

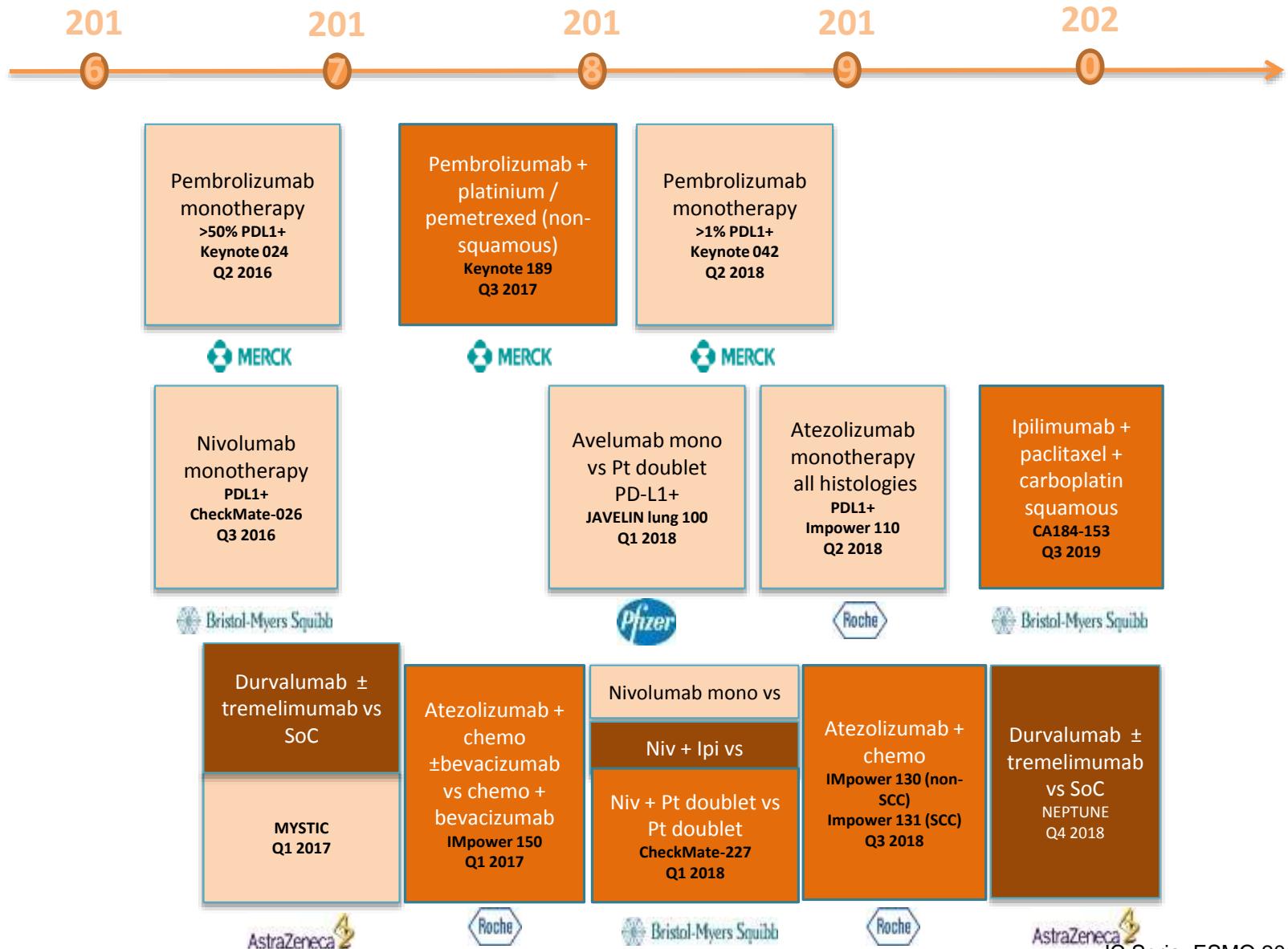
# Nivolumab ± Ipilimumab ORR by Tumor PD-L1 Expression

CheckMate 012: First-Line Nivolumab ± Ipilimumab in NSCLC



- 5 CRs (10%) were achieved in the nivolumab monotherapy cohort (1 in a patient with tumor PD-L1 expression <1%)
- 6 CRs (8%) were achieved in the nivolumab + ipilimumab cohorts<sup>a</sup> (3 in patients with tumor PD-L1 expression <1%)

# Upcoming randomized immunotherapy trials in 1<sup>st</sup> line NSCLC and projected read-out timelines



# Come cambia l'algoritmo di trattamento nella pratica clinica del NSCLC



## Immunoterapia: Cosa si ha fato imparare?

**1° LINEA**  
**2° LINEA**

# Randomised Trials of Anti PD-1 / anti-PD-L1 Agents

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CheckMate 057 <b>(Non-sqm)</b>	2 <sup>nd*</sup>	Nivol vs Docetaxel	Unselected	Improved OS	0.73
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Reck et al. N Engl J Med 2016; 375: 1823 - 33

# Summary of phase III studies of immunotherapy in previously treated patients

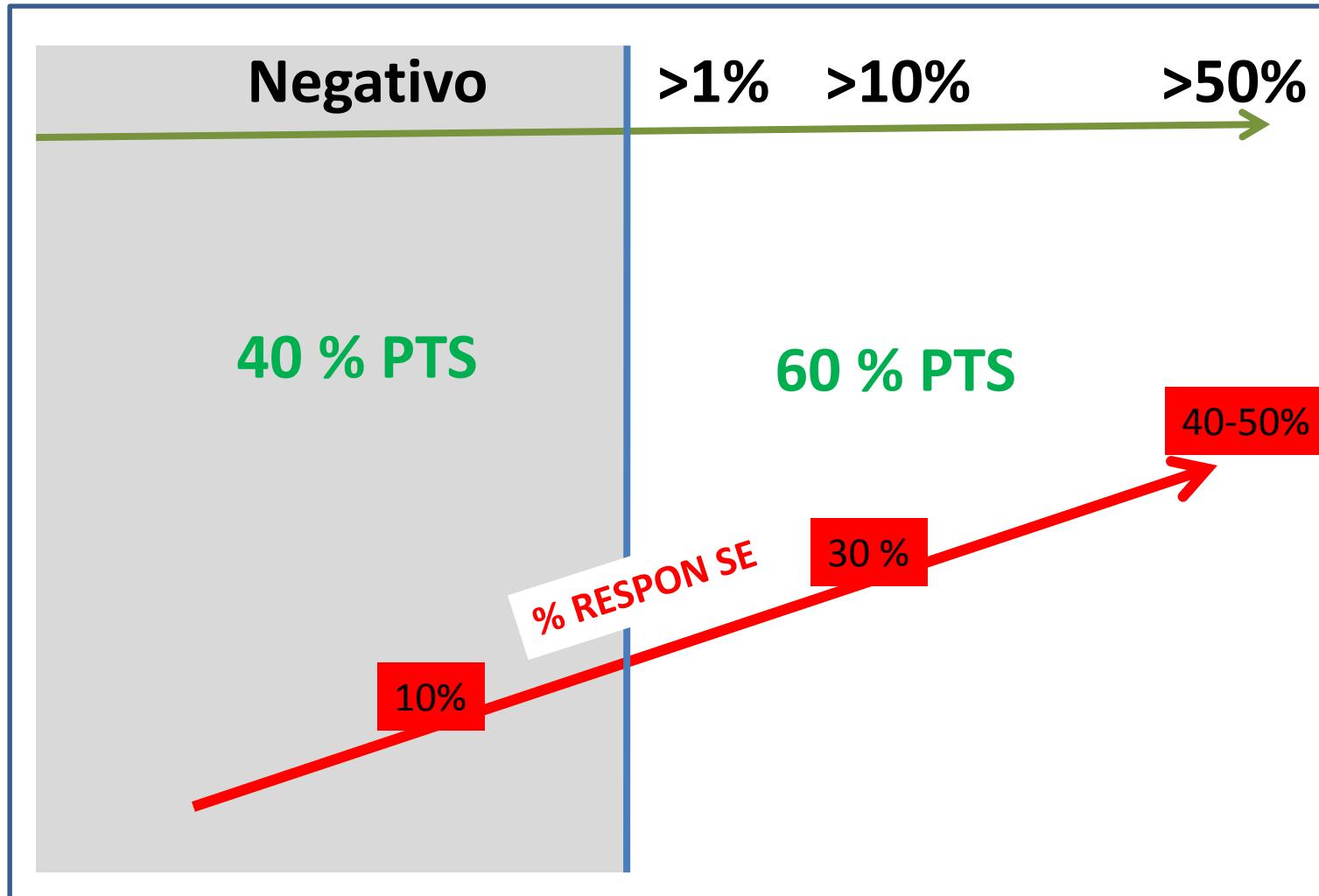
	CheckMate 017 <sup>1</sup> Nivolumab vs docetaxel	CheckMate 057 <sup>1</sup> Nivolumab vs docetaxel	KEYNOTE-010 <sup>2</sup> Pembrolizumab (2mg/kg or 10mg/kg) vs docetaxel	OAK <sup>3</sup> Atezolizumab vs docetaxel
<b>Phase of study</b>	III	III	II/III	III
<b>PD-L1 selected</b>	No	No	Yes (TPS* ≥1%)	No
<b>Study size, n</b>	272 (135 vs 137)	582 (292 vs 290)	1,033 (344 vs 346 vs 343)	1,225 (425 vs 425)*
<b>Histology</b>	Squamous	Non-squamous	All-comers	All-comers
<b>Line of therapy, %</b>				
2L	100	88	69	75
3L	0	11	20	25
>3L	0	<1	9	0
Other/unknown	0	0	<1	0
<b>Subsequent CIT (immunotherapy arm vs chemo arm), %</b>	<1 vs 2	1 vs 2	0.6 vs 1.7 vs 13.1	4.5 vs 17.2
<b>Crossover from chemo arm to study immunotherapy, %</b>	4	6	Not permitted	Not permitted
<b>Median OS, months</b>	9.2 vs 6.0	12.2 vs 9.5	10.4 vs 12.7 vs 8.5 2mg/kg: 0.71 (p=0.0008) 10mg/kg: 0.61 (p<0.0001)	13.8 vs 9.6 0.73 (p=0.0003)
<b>HR vs docetaxel (p value)</b>	0.62 (p=0.0004)	0.75 (p<0.001)		

\*850 in primary population  
NR = not reached

1. Borghaei, et al. ASCO 2016  
2. Herbst, et al. Lancet 2015; 3. Barlesi, et al. ESMO 2016

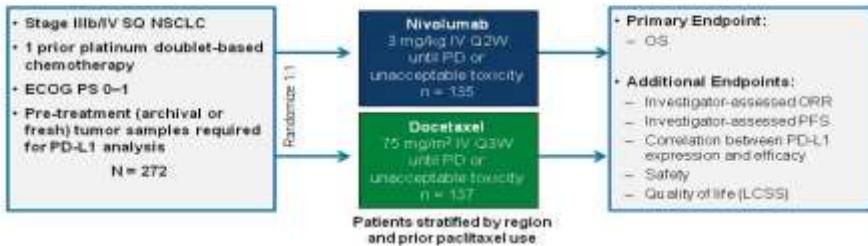
# 2°-Line NSCLC

## Anti PD-1 / anti-PD-L1 Agents , PD-L1 expression



# Second line – Nivolumab Checkmate 017 (SQM)

## CheckMate 017 (NCT01642004) - Study Design

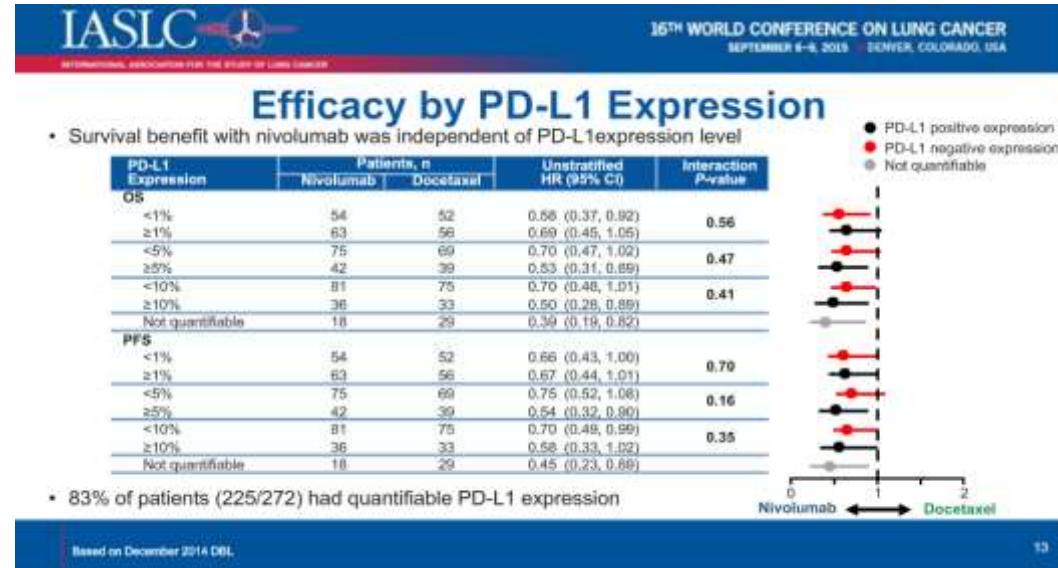


- One pre-planned interim analysis for OS
- At time of DBL (December 15, 2014), 199 deaths were reported (86% of deaths required for final analysis)
- The boundary for declaring superiority for OS at the pre-planned interim analysis was  $P < 0.03$

LCSS = Lung cancer symptom scale

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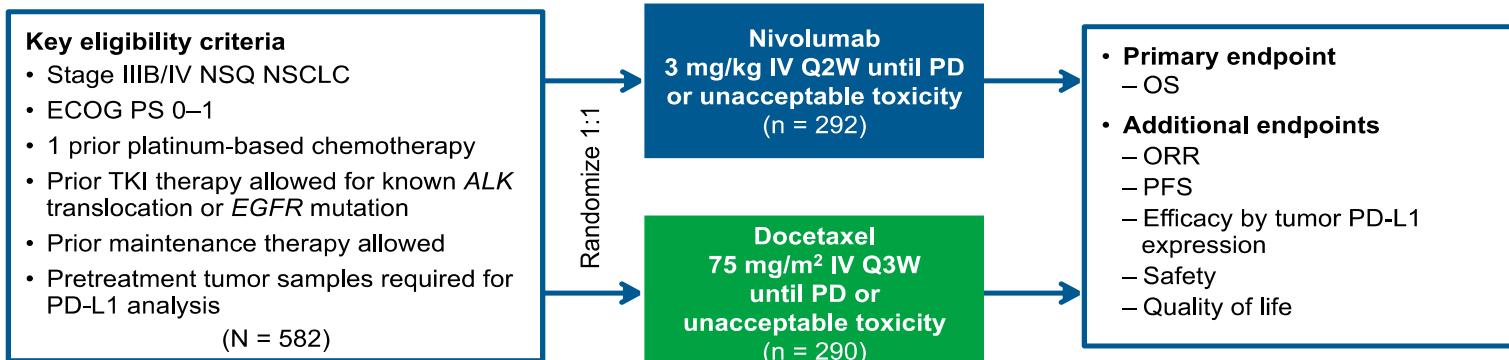
PRESENTED AT ASCO Annual Meeting



# Second line – Nivolumab

## Checkmate 057

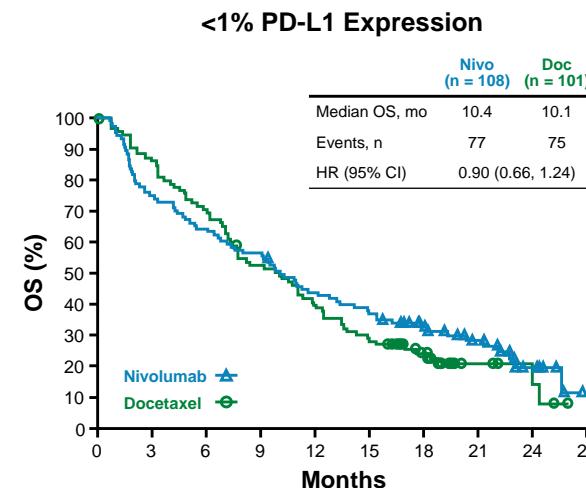
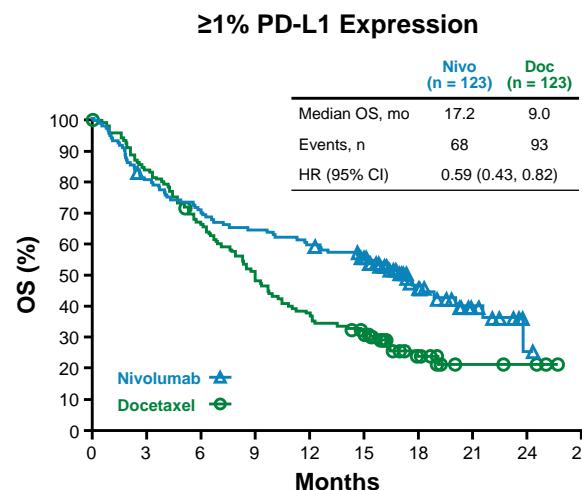
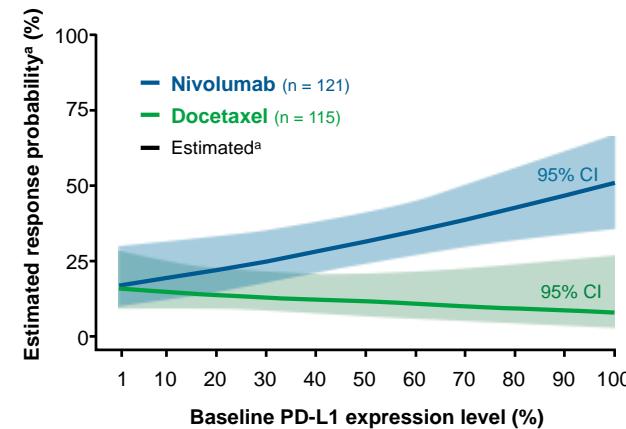
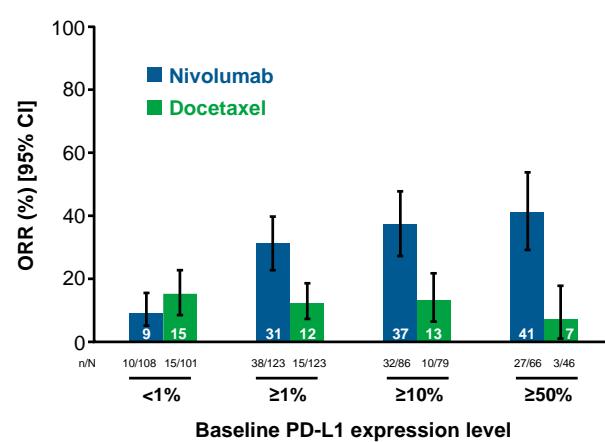
### Phase 3 CheckMate 057 Study Design: Nivolumab vs Docetaxel in Previously Treated NSQ NSCLC



Randomization stratified by prior maintenance therapy and line of therapy (second-line vs third-line)

# Second line – Nivolumab Checkmate 057

## PD-L1 Expression Continuum and Response Probability CheckMate 057: Nivolumab vs Docetaxel in Previously Treated NSQ NSCLC

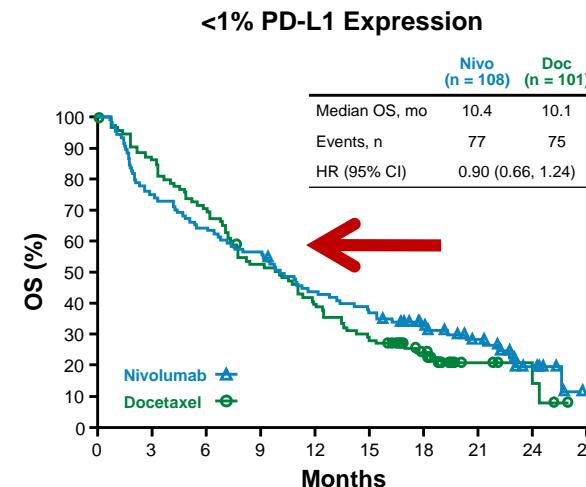
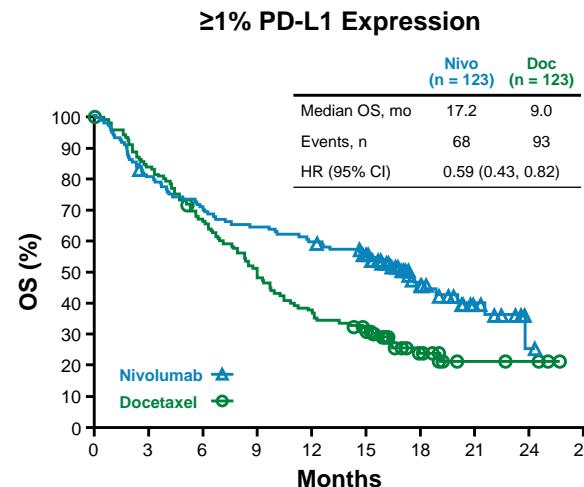
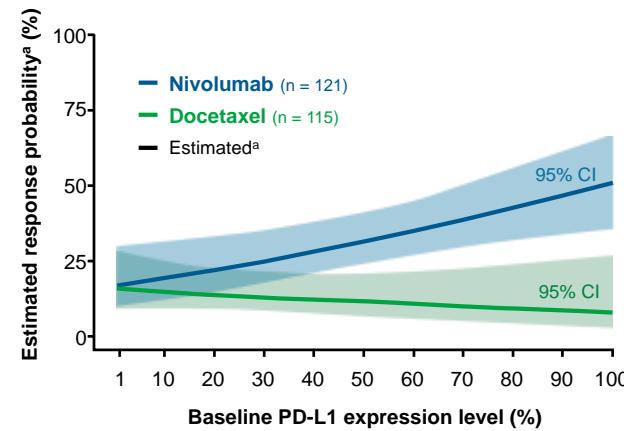
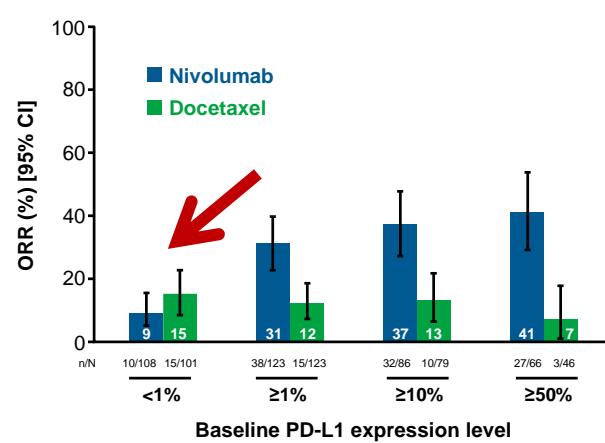


# Second line – Nivolumab

## Checkmate 057

### PD-L1 Expression Continuum and Response Probability

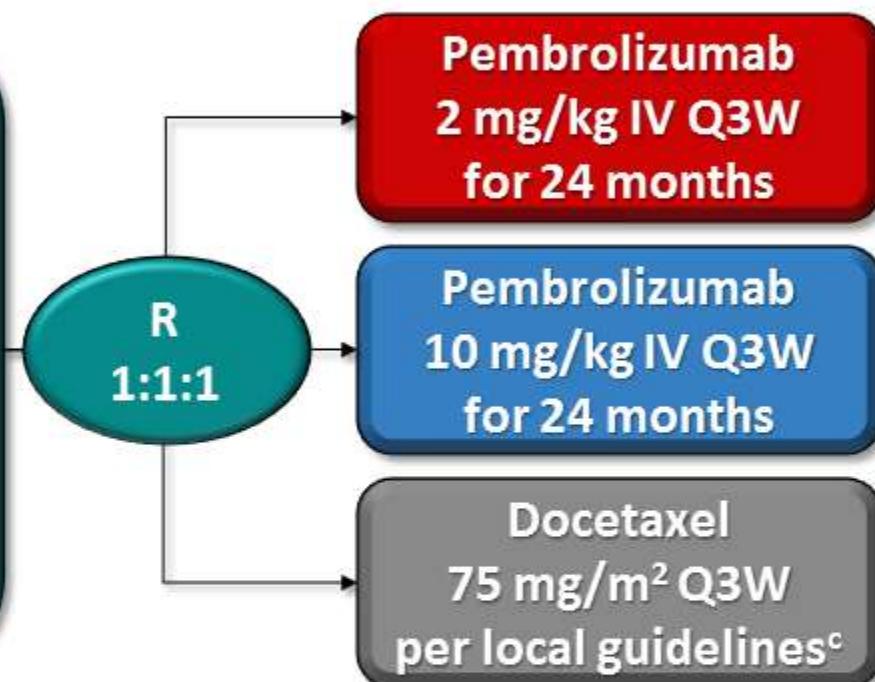
CheckMate 057: Nivolumab vs Docetaxel in Previously Treated NSQ NSCLC



# KEYNOTE-010 Study Design

## Patients

- Advanced NSCLC
- Confirmed PD after ≥1 line of chemotherapy<sup>a</sup>
- No active brain metastases
- ECOG PS 0-1
- PD-L1 TPS ≥1%
- No serious autoimmune disease
- No ILD or pneumonitis requiring systemic steroids



## Stratification factors:

- ECOG PS (0 vs 1)
- Region (East Asia vs non-East Asia)
- PD-L1 status<sup>b</sup> (TPS ≥50% vs 1%-49%)

## End points in the TPS ≥50% stratum and TPS ≥1% population

- Primary: PFS and OS
- Secondary: ORR, duration of response, safety

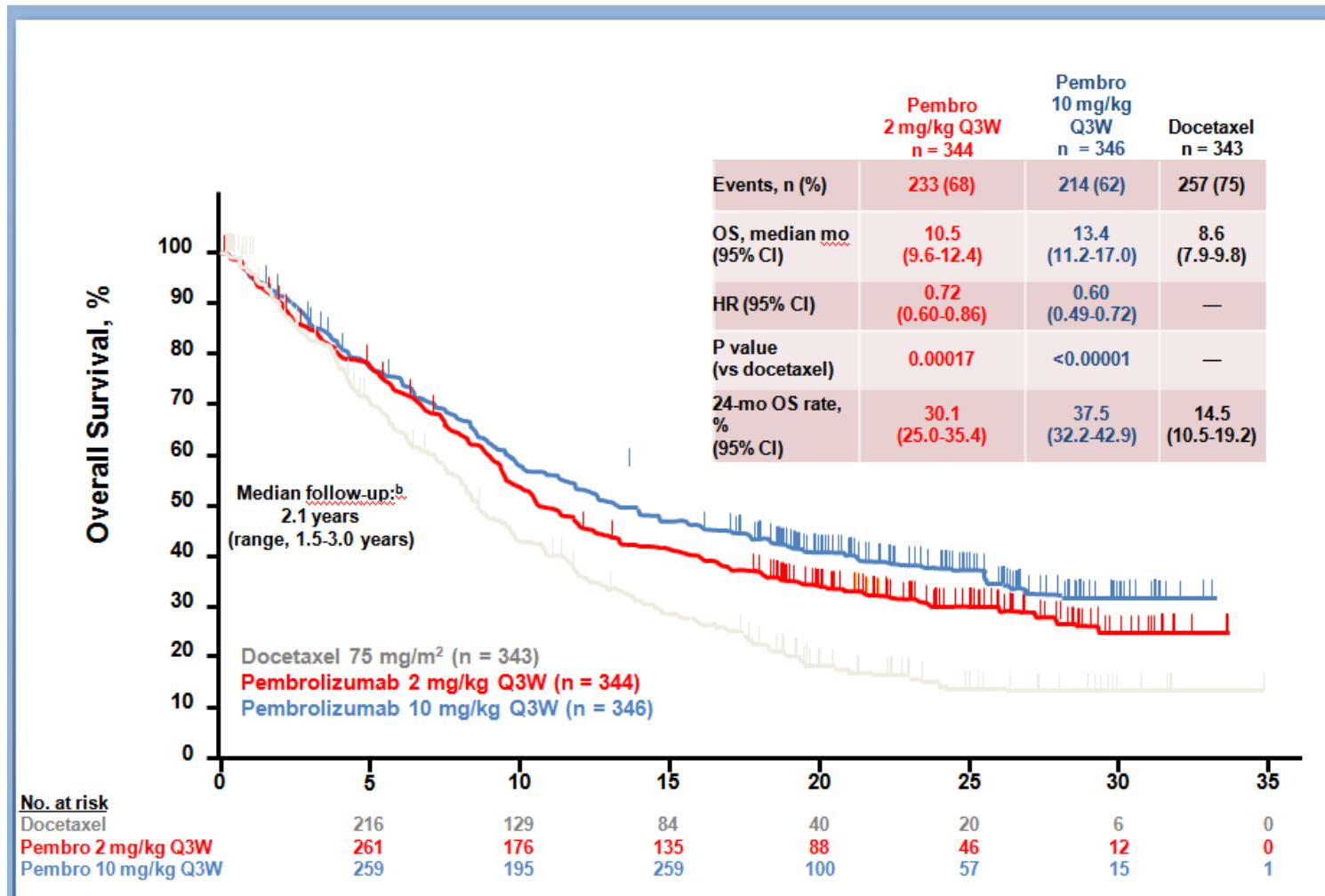
ClinicalTrials.gov, NCT01905657.

<sup>a</sup>Prior therapy must have included ≥2 cycles of platinum-doublet chemotherapy. An appropriate tyrosine kinase inhibitor was required for patients whose tumors had an EGFR sensitizing mutation or an ALK translocation.

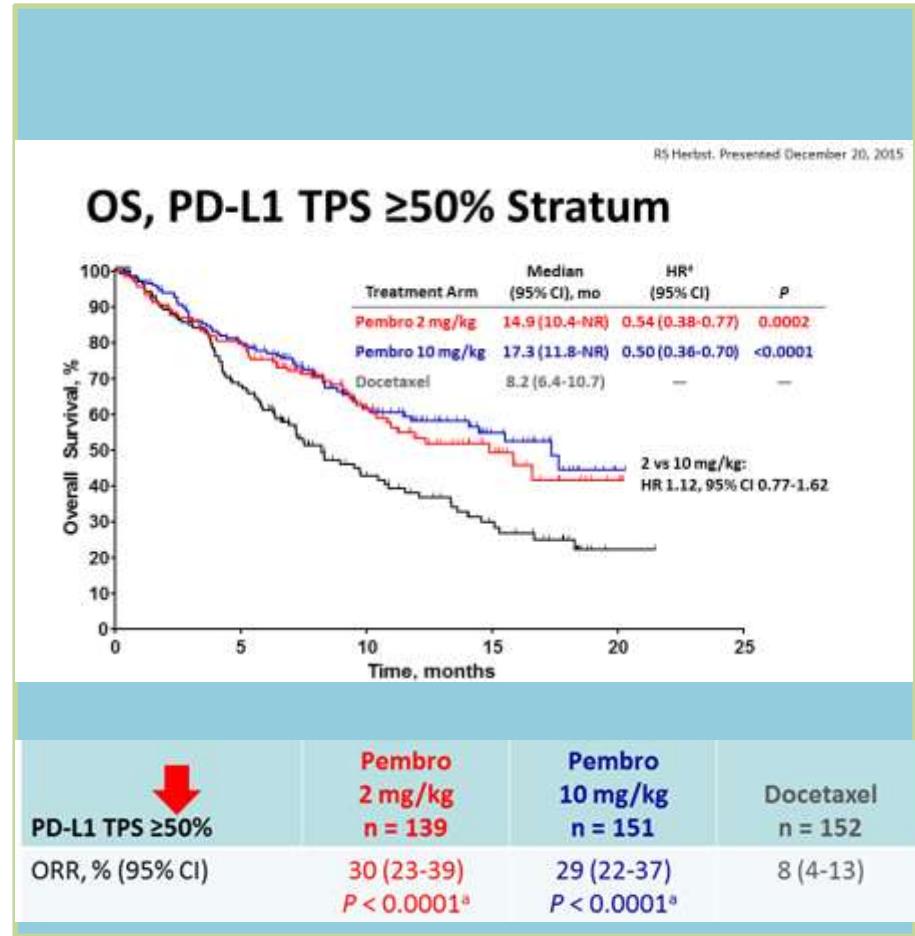
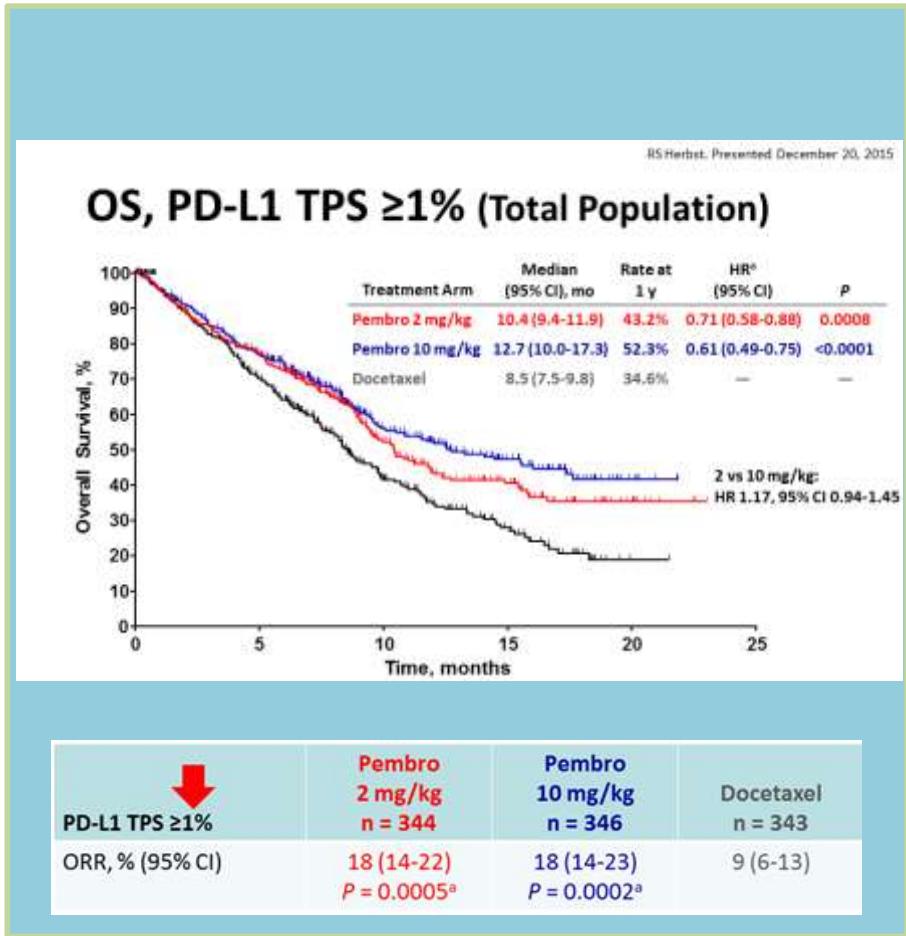
<sup>b</sup>Added after 441 patients enrolled based on results from KEYNOTE-001 (Garon EB et al. *N Engl J Med.* 2015;372:2018-28).

<sup>c</sup>Patients received the maximum number of cycles permitted by the local regulatory authority.

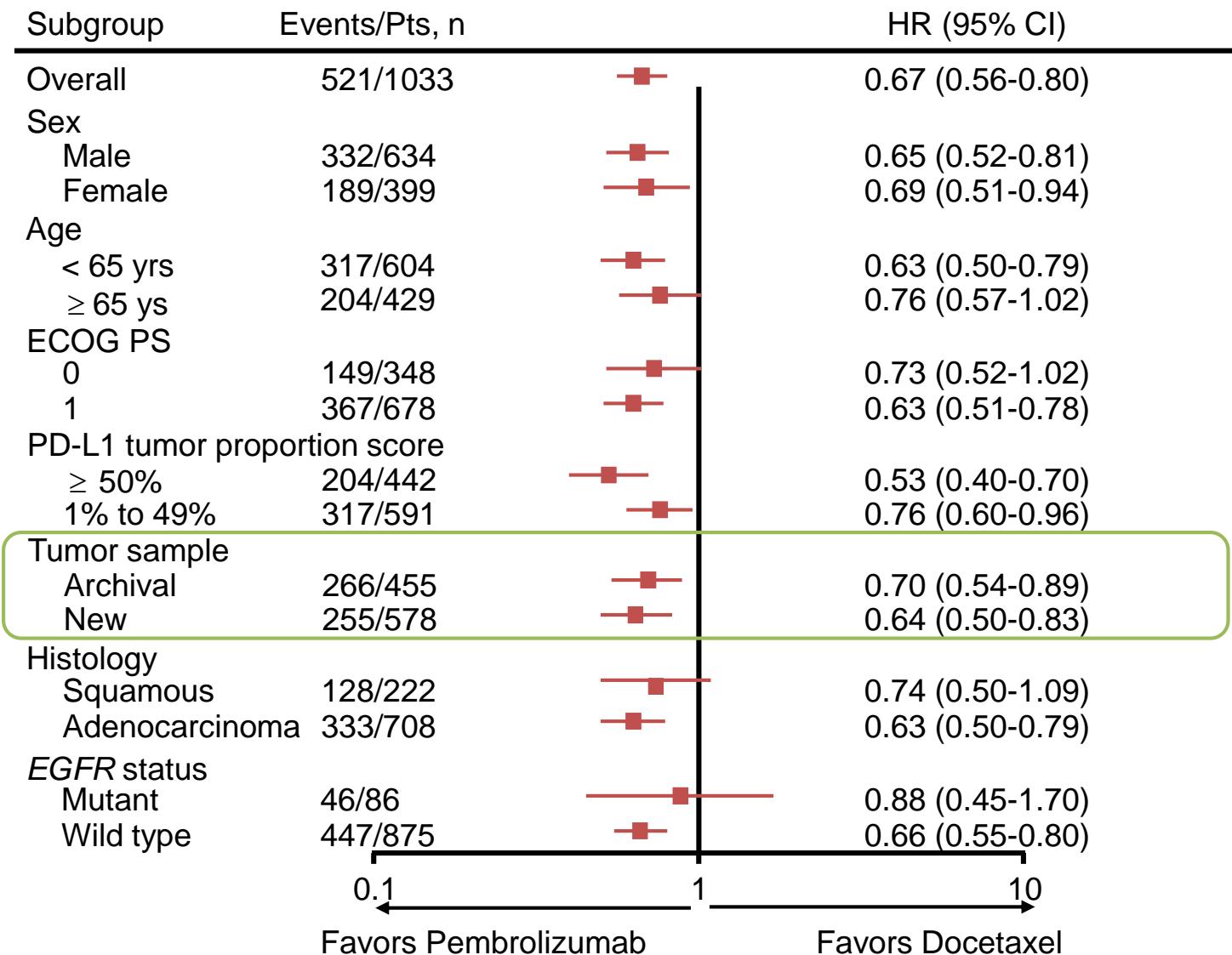
# OS KEYNOTE-010 STUDY



# Efficacy to PEMBROLIZUMAB according to PDL1 expresion (KN-010)



# KEYNOTE-010: OS by Pt Subgroup

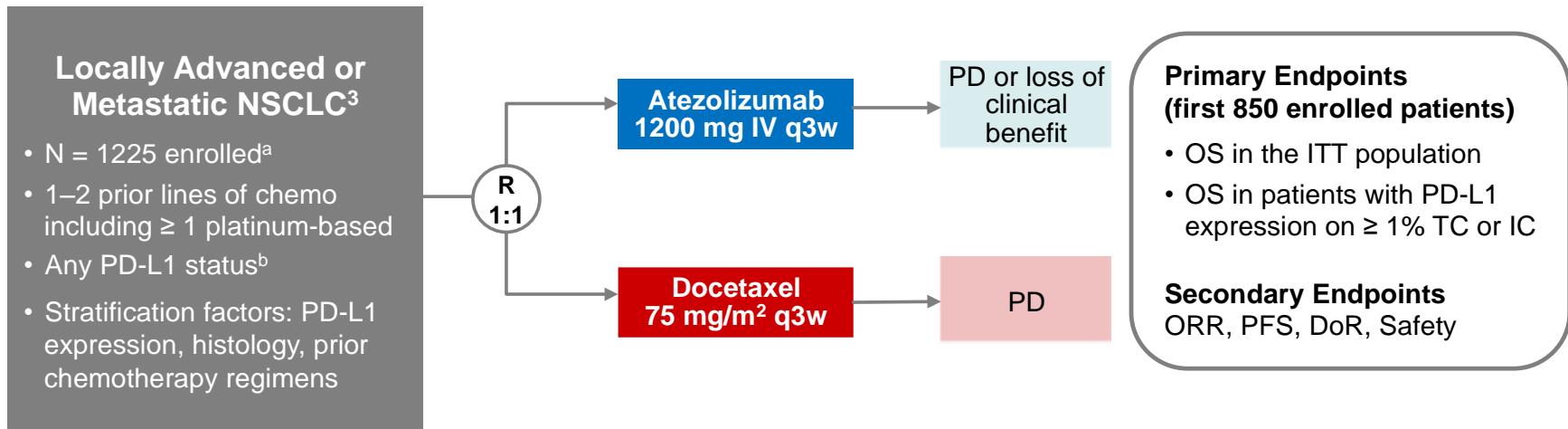


# Second line – Atezo Subgroups OAK PhIII

## Phase III OAK study design

Atezolizumab (anti–PD-L1) is an engineered mAb that inhibits the PD-L1/PD-1 and PD-L1/B7.1 interactions to restore anti-tumor T-cell activity and enhance T-cell priming<sup>1,2</sup>

### OAK study design



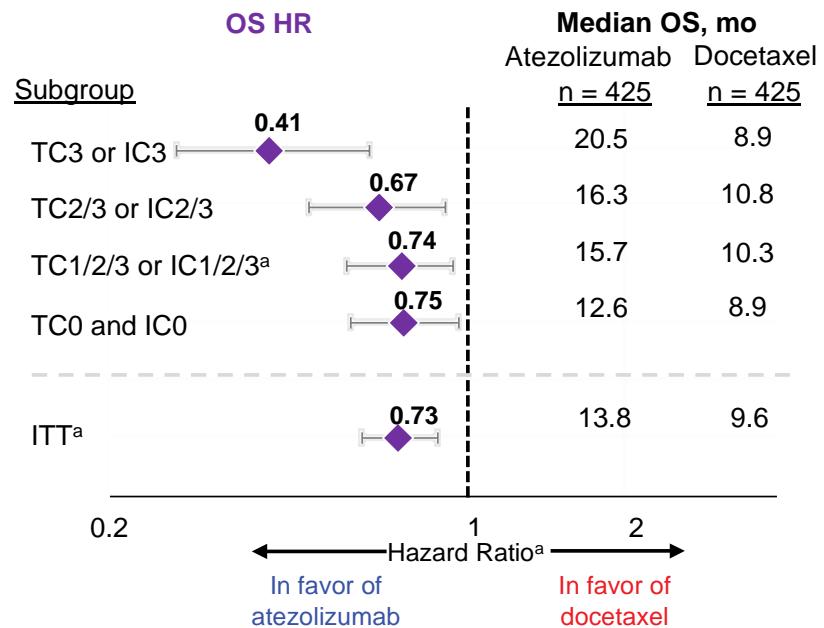
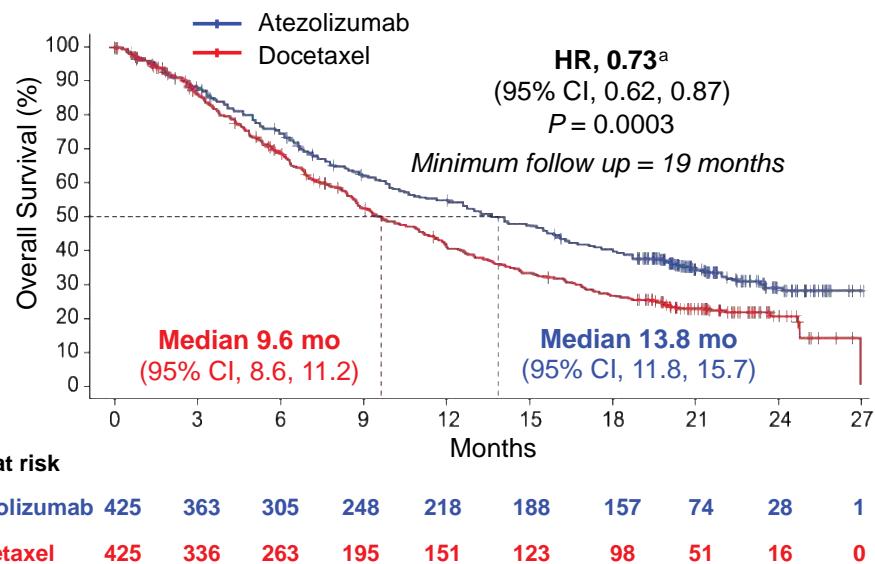
<sup>a</sup>A prespecified analysis of the first 850 patients provided sufficient power to test the co-primary endpoints of OS in the ITT and TC1/2/3 or IC1/2/3 subgroup (≥ 1% PD-L1 expression).

<sup>b</sup>PD-L1 expression assessed with VENTANA SP142 IHC assay

1. Herbst *Nature* 2014. 2. Chen *Immunity* 2013. 3. Barlesi et al. ESMO 2016 LBA44

# Second line – Atezo Subgroups OAK PhIII

## Overall survival, ITT (n = 850) and PD-L1 subgroups



<sup>a</sup>Stratified HR for ITT and TC1/2/3 or IC1/2/3. Unstratified HR for other subgroups.

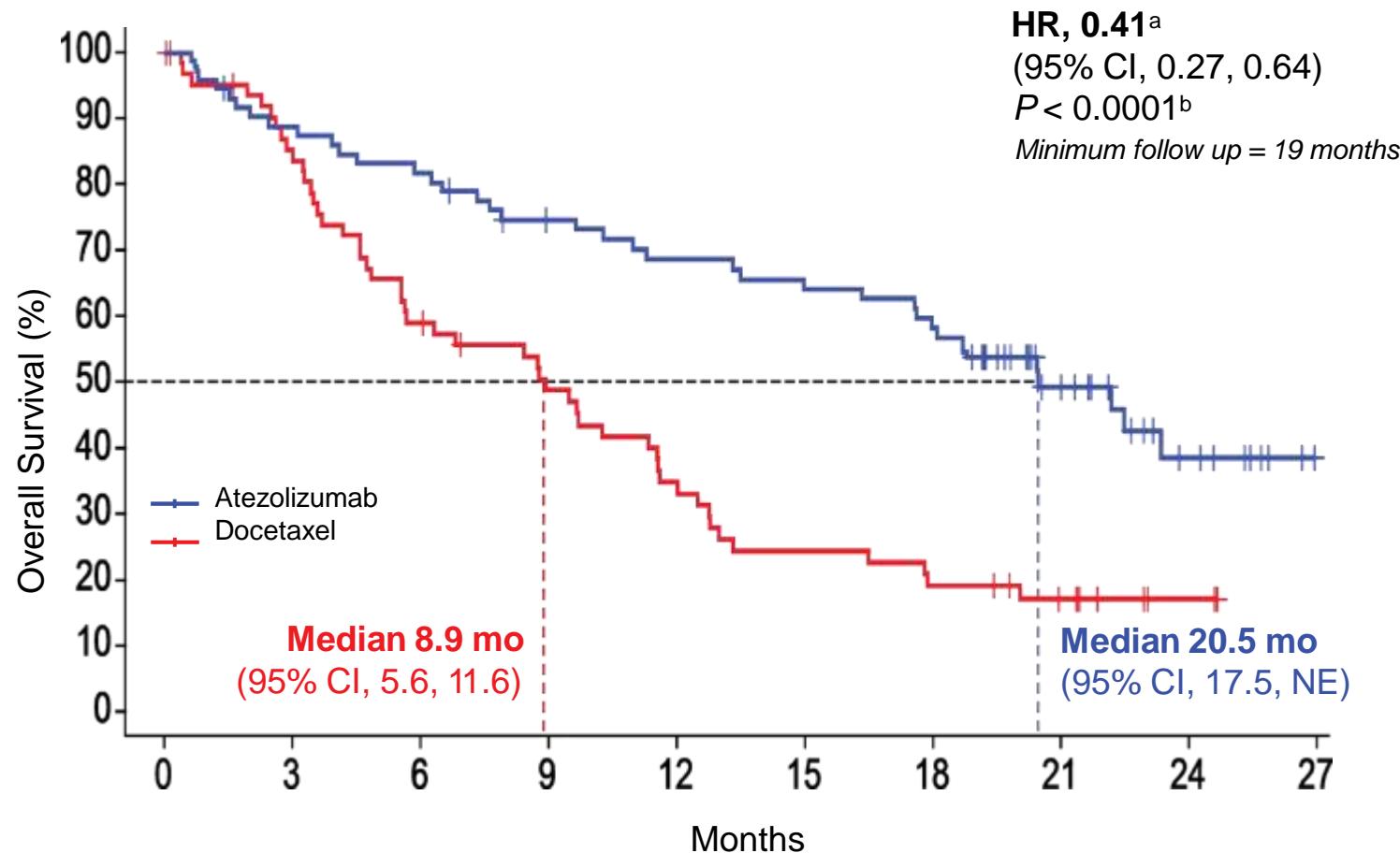
TC, tumor cells; IC, tumor-infiltrating immune cells; OS, overall survival.

Barlesi et al. ESMO 2016 LBA44

Gadgeel et al., WCLC 2016

# OS, PD-L1 EXPRESSION ON $\geq 50\%$ TC OR $\geq 10\%$ IC

## TC3 OR IC3; 16% OF PATIENTS



No. at Risk

Atezolizumab	72	69	65	63	61	59	58	55	51	50	49	47	46	46	44	43	43	42	39	34	28	19	16	11	8	6	2
Docetaxel	65	59	57	51	45	40	36	32	32	28	25	24	20	15	14	14	14	13	11	9	7	4	3	2			

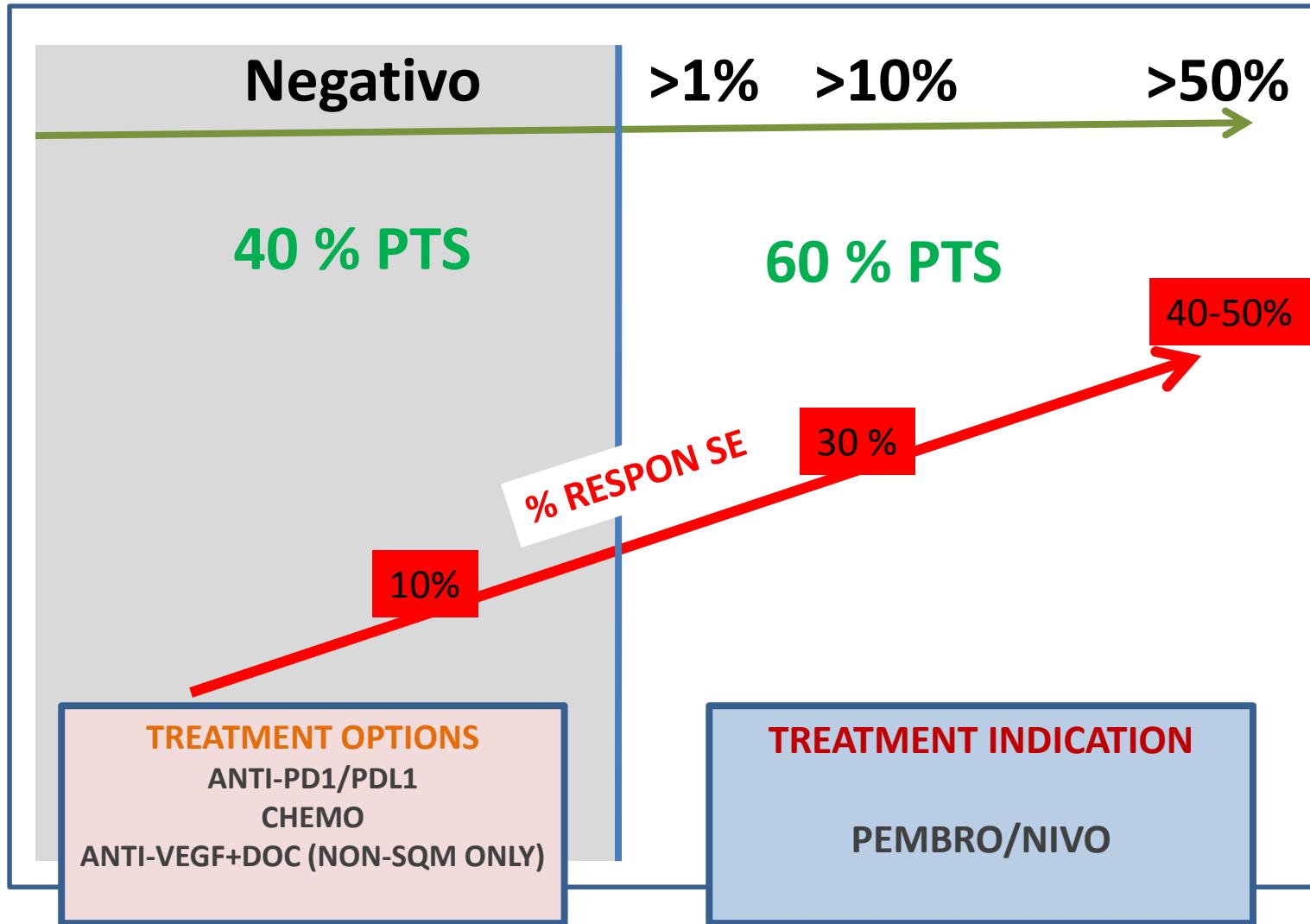
<sup>a</sup>Unstratified HR.

<sup>b</sup>P values for descriptive purpose only.

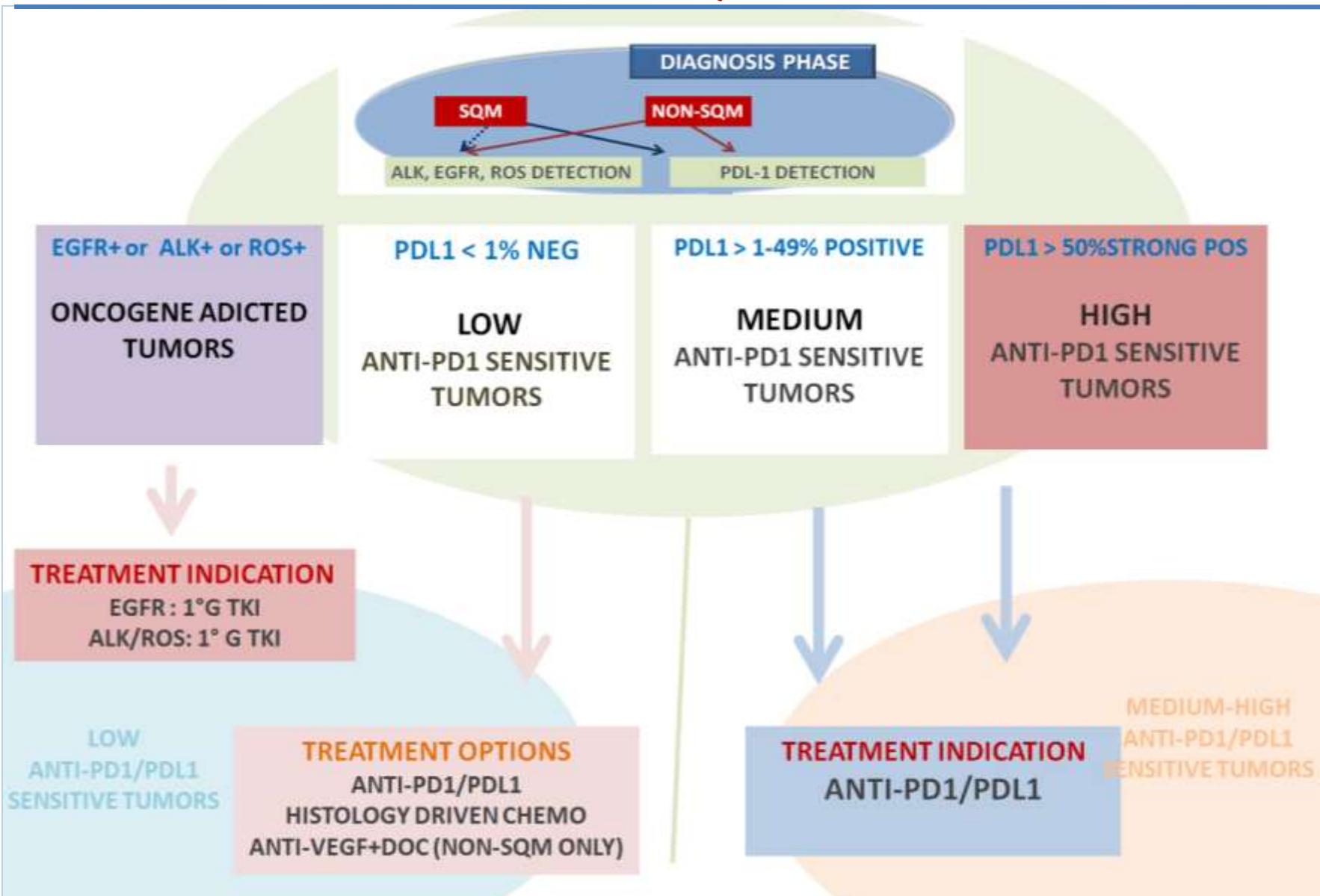
TC, tumor cells; IC, tumor-infiltrating immune cells; OS, overall survival.

# 2°-Line NSCLC (Non-Sqm)

## Anti PD-1 / anti-PD-L1 Agents , PD-L1 expression



# 2°-line NSCLC DX/TX algorithm: “1° Schenario” Non-SQM



# I biomarcatori nell'immunoncologia: L'importanza della selezione del paziente

Melanoma

Rene

Vescica

Testa e collo

PDL1 ?

# I biomarcatori nell'immunoncologia: L'importanza della selezione del paziente

Melanoma

Rene

Vescica

Testa e collo

PENSARE ?  


# PD-1 Inhibitors: Approved Indications in Advanced Melanoma

- **Nivolumab**

- Single agent (3 mg/kg q2w) for **unresectable or metastatic** melanoma with or without a *BRAF* V600 mutation
- In combination with ipilimumab for **unresectable or metastatic** melanoma
  - Nivolumab 1 mg/kg + ipilimumab 3 mg/kg q3w x 4 then nivolumab 3 mg/kg q2w

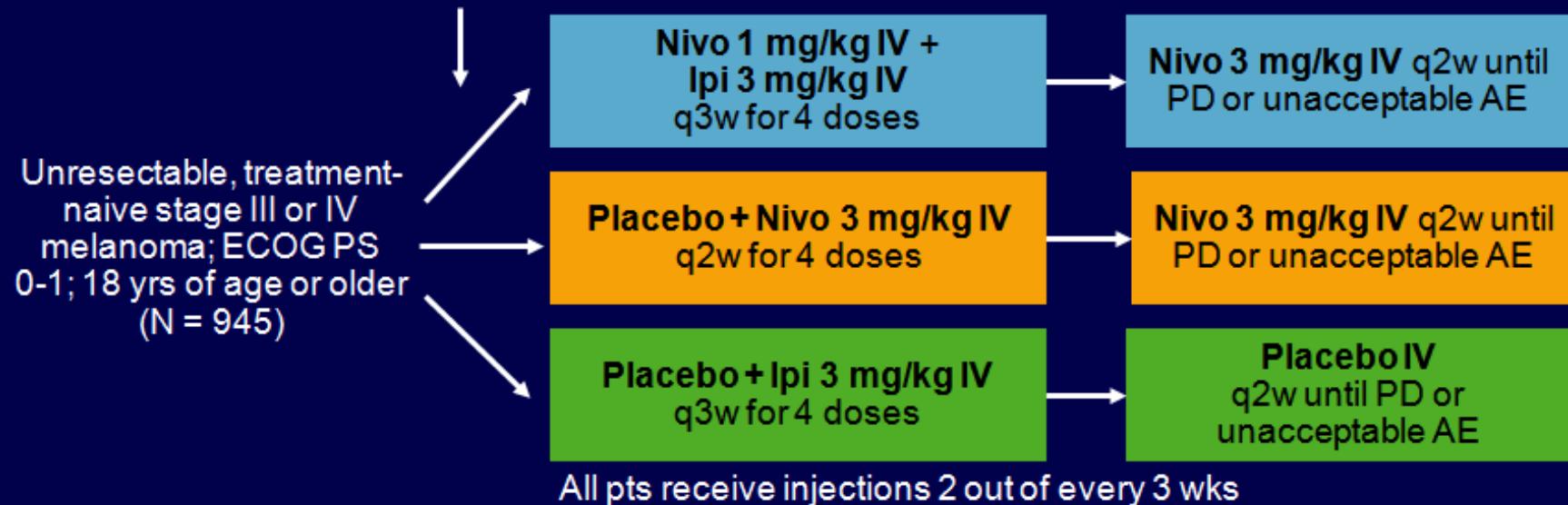
- **Pembrolizumab**

- Single agent (2 mg/kg q3w) for **unresectable or metastatic** melanoma

# Nivo + Ipi vs Nivo vs Ipi for First-line Treatment of Melanoma (Checkmate 067)

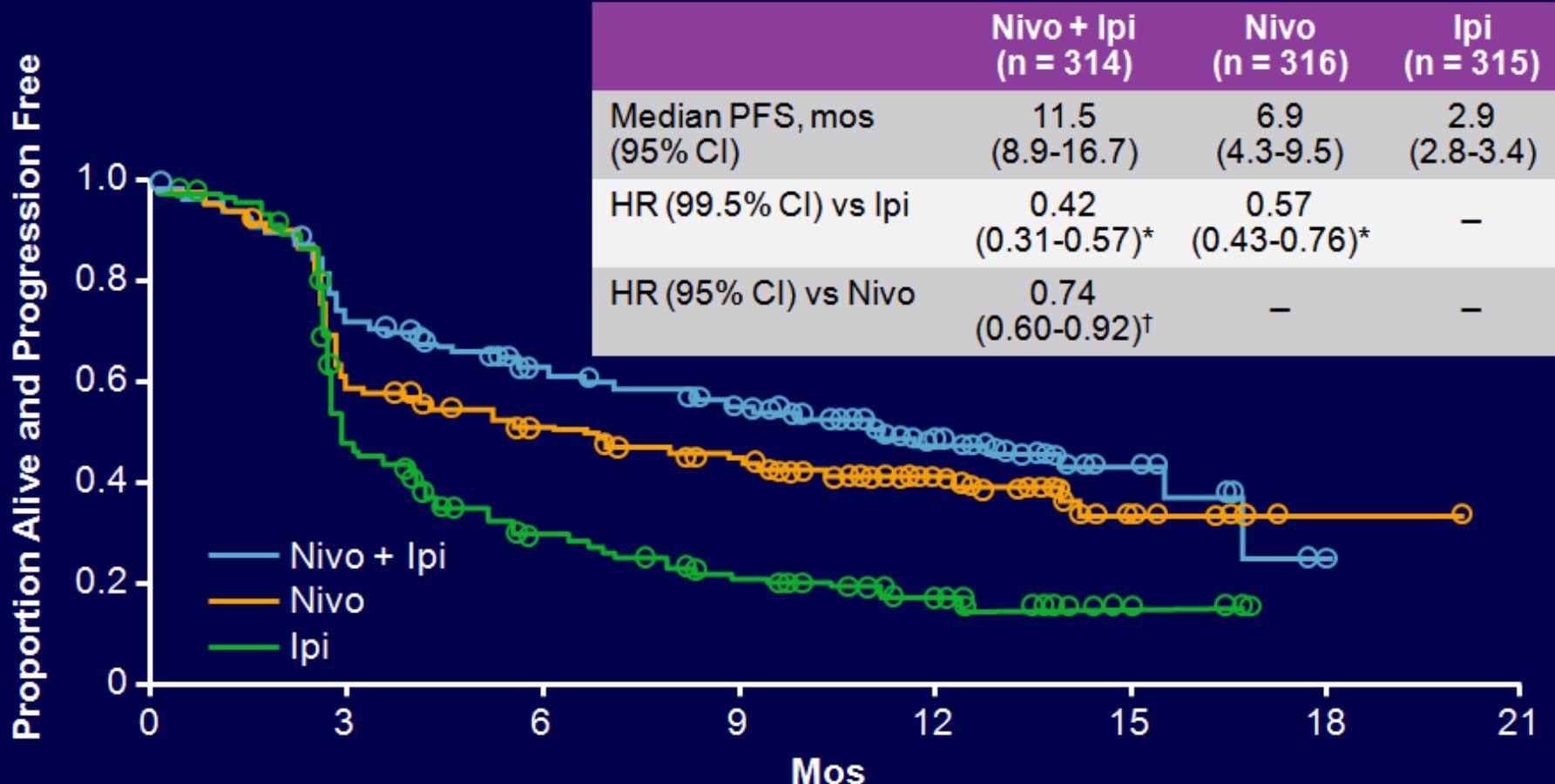
A randomized, double-blind phase III study

Stratified by tumor PD-L1 status (positive vs negative/indeterminate), BRAF mutation status (V600 mutation positive vs wild type), and AJCC metastasis stage (M0, M1a, or M1b vs M1c)



- Primary endpoint: OS, PFS
- Secondary endpoint: ORR, OS by PD-L1, safety

# CheckMate 067: Improved PFS With Nivo + Ipi or Nivo Alone vs Ipi Alone



\*Stratified log-rank  $P < .00001$  vs Ipi.

†Exploratory endpoint. Study not powered to detect a statistical difference between Nivo + Ipi and Nivo.

# CheckMate 067: Nivo + Ipi Provides Most Benefit for PD-L1lo, Similar to Nivo for PD-L1hi

## PD-L1 ≥ 5%\* (hi)

Median  
PFS, Mos HR

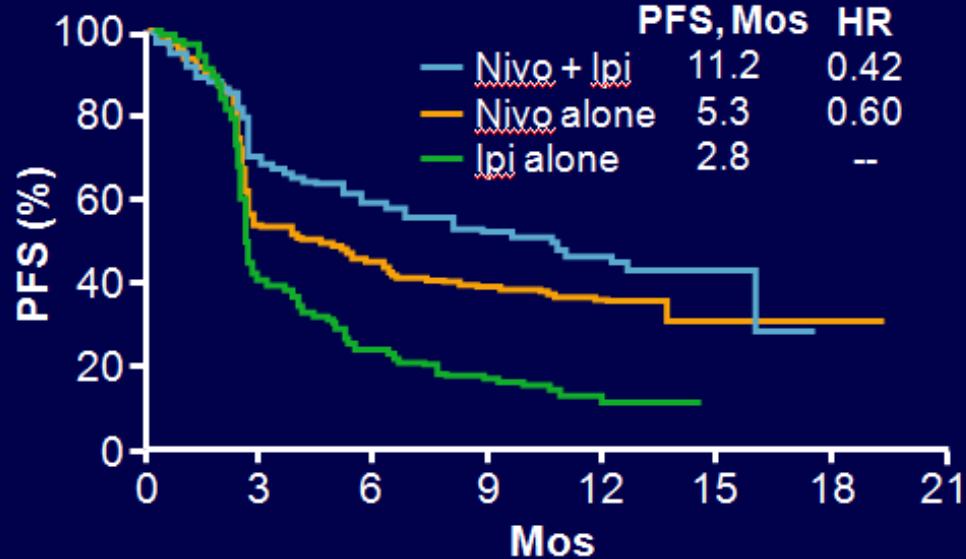
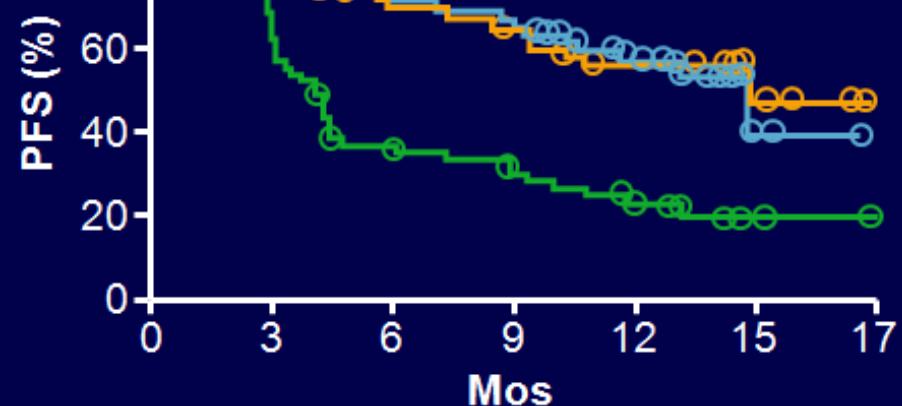
Nivo + Ipi	14.0	0.40
Nivo alone	14.0	0.40
Ipi alone	3.9	--

## PD-L1 < 5%\* (lo)

Median

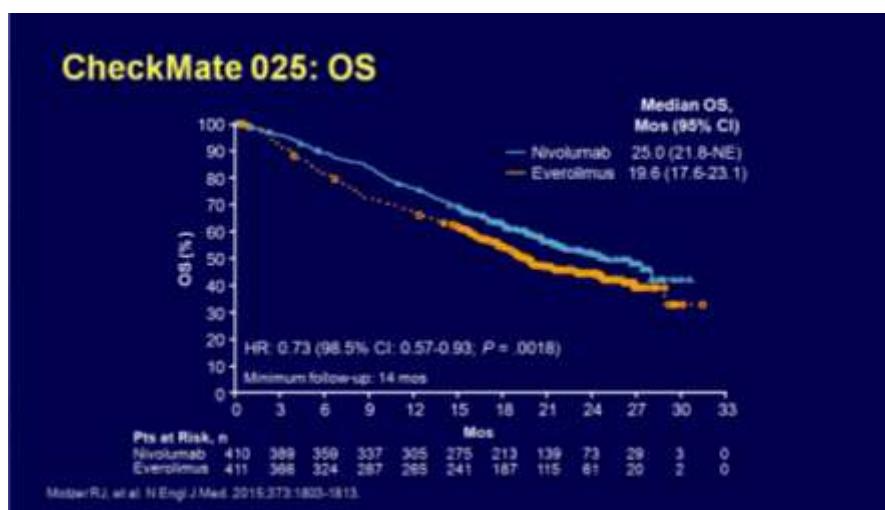
PFS, Mos HR

Nivo + Ipi	11.2	0.42
Nivo alone	5.3	0.60
Ipi alone	2.8	--

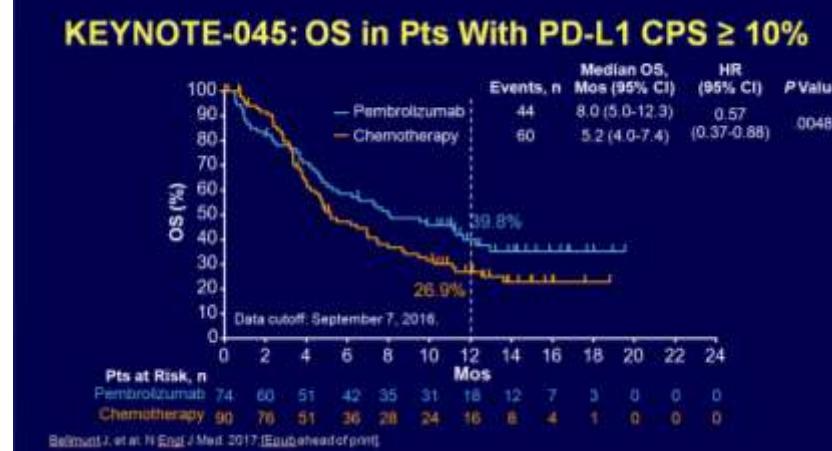
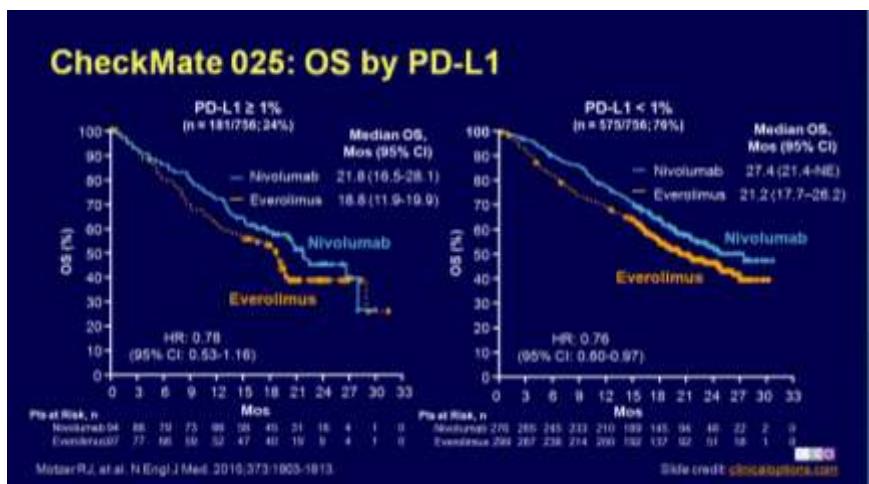
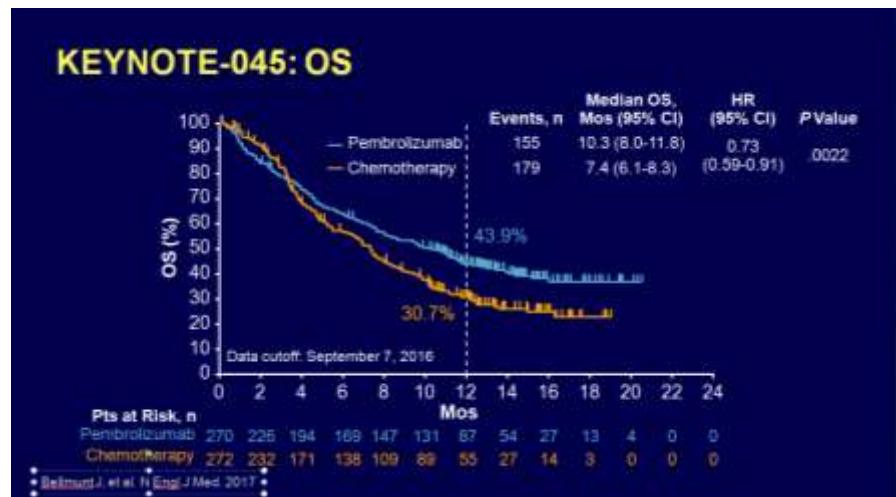


\*Pervalueated PD-L1 IHC assay based on PD-L1 staining of tumor cells in a section of at least 100 evaluable tumor cells.

## RENE – 2° linea



## VESCICA – 2° linea



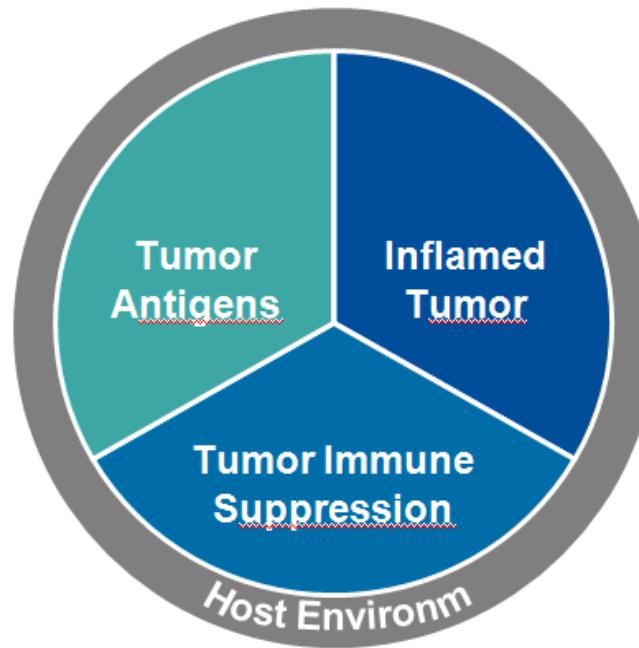
# Tumor and Immune Biomarkers Being Evaluated to Predict Better Outcomes to Immuno-Oncology Therapy

## Tumor Antigens

- Biomarkers indicative of hypermutation & neo-antigens may predict response to IO treatment

*Examples:*

- *TMB, MSI-High, Neo-Antigens*



## Inflamed Tumor Microenvironment

- Biomarkers (intra- or peri-tumoral) indicative of an inflamed phenotype may predict response to IO treatment

*Examples:*

- *PD-L1, Inflammatory Signatures*

IDO = indoleamine-2,3 dioxygenase; LAG-3 = lymphocyte activation gene-3; MDSCs = myeloid-derived suppressor cells;  
MSI-High = microsatellite instability high; TMB = tumor mutational burden. Adapted from Blank C.U. et al., *Science* 2016;352:658–660.

ORIGINAL ARTICLE

# PD-1 Blockade in Tumors with Mismatch-Repair Deficiency

D.T. Le, J.N. Uram, H. Wang, B.R. Bartlett, H. Kemberling, A.D. Eyring, A.D. Skora, B.S. Luber, N.S. Azad, D. Laheru, B. Biedrzycki, R.C. Donehower, A. Zaheer, G.A. Fisher, T.S. Crocenzi, J.J. Lee, S.M. Duffy, R.M. Goldberg, A. de la Chapelle, M. Koshiji, F. Bhajjee, T. Huebner, R.H. Hruban, L.D. Wood, N. Cuka, D.M. Pardoll, N. Papadopoulos, K.W. Kinzler, S. Zhou, T.C. Cornish, J.M. Taube, R.A. Anders, J.R. Eshleman, B. Vogelstein, and L.A. Diaz, Jr.

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## ABSTRACT

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### BACKGROUND

Somatic mutations have the potential to encode “non-self” immunogenic antigens. We hypothesized that tumors with a large number of somatic mutations due to mismatch-repair defects may be susceptible to immune checkpoint blockade.

### METHODS

We conducted a phase 2 study to evaluate the clinical activity of pembrolizumab, an anti-programmed death 1 immune checkpoint inhibitor, in 41 patients with progressive metastatic carcinoma with or without mismatch-repair deficiency. Pembrolizumab was administered intravenously at a dose of 10 mg per kilogram of body weight every 14 days in patients with mismatch repair-deficient colorectal cancers, patients with mismatch repair-proficient colorectal cancers, and patients with mismatch repair-deficient cancers that were not colorectal. The coprimary end points were the immune-related objective response rate and the 20-week immune-related progression-free survival rate.

The authors’ full names, academic degrees, and affiliations are listed in the Appendix. Address reprint requests to Dr. Diaz at the Johns Hopkins Sidney Kimmel Comprehensive Cancer Center, 1650 Orleans St., Rm. 590, Baltimore, MD 21287, or at ldiaz1@jhmi.edu.

This article was published on May 30, 2015, at NEJM.org.

N Engl J Med 2015;372:2509-20.

DOI: 10.1056/NEJMoa1500596

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# Pembrolizumab in Mismatch Repair–Deficient CRC: Overview

- **Mismatch repair defects lead to MSI-H**
  - **MSI-H** associated with hereditary nonpolyposis colorectal carcinoma and present in 15% of CRCs across all stages
  - **MSI-H** tumors harbor genetic mutations that are potentially recognizable by immune system; **abundant expression of neoantigens on tumor cells creates inflamed microenvironment with high expression of immune checkpoints, such as PD-1**
- **PD-1 blockade has antitumor effects in MSI-H tumors**
- Pembrolizumab, an immune checkpoint inhibitor of PD-1, evaluated in pts with treatment-refractory, progressive, metastatic cancer by mismatch repair status<sup>[1]</sup>
  - Received breakthrough therapy designation in 11/2015 for tx of MSI-H CRC
- Current report summarizes phase I data on PD-1 blockade with pembrolizumab in pts with mismatch repair–deficient CRC<sup>[2]</sup>

1. Le DT, et al. N Engl J Med. 2015;372:2509-2520.

2. Le DT, et al. ASCO 2016. Abstract 103.

# Pembrolizumab in Mismatch Repair–Deficient CRC: Study Design

- Eligibility for cohorts A and B:
  - Histologically confirmed metastatic or locally advanced CRC, with or without mismatch repair deficiency
  - Mismatch repair deficiency defined as: deficiency in MLH1, MSH2, MSH6 or PMS2 by IHC, or microsatellite instability in  $\geq 2$  loci by PCR
  - Measurable disease
  - $\geq 2$  previous cancer therapy regimens
  - ECOG PS  $\leq 1$
  - No previous anti–PD-1/PD-L1/PD-L2, anti-CD137, anti–OX-40, anti–CTLA-4
- Pembrolizumab 10 mg/kg every 2 wks
- Current report: updated data from cohort A

Cohort A (n = 28)  
MMRD CRC

Cohort B (n = 25)  
MMRP CRC

Cohort C (n = 30)  
MMRD non-CRC

# Pembrolizumab in Mismatch Repair–Deficient CRC: Pt Population

Characteristics	MMRD CRC (n = 28)	MMRP CRC (n = 25)
Median age, yrs (range)	49 (26-75)	62 (32-79)
Male, n (%)	15 (54)	16 (64)
ECOG PS, n (%)		
▪ 0	5 (18)	7 (28)
▪ 1	23 (82)	18 (72)
No. of previous treatments, median	3	4
Liver metastases, n (%)	14 (50)	15 (60)
Lynch syndrome, n (%)		
▪ Yes	15 (54)	0 (0)
▪ No	2 (7)	25 (100)
▪ Unknown	11 (39)	0 (0)

# Pembrolizumab in Mismatch Repair–Deficient CRC: Efficacy

Outcome	MMRD CRC (n = 28)	MMRP CRC (n = 25)
Median follow-up, mos	9.3	6
<b>ORR, % (95% CI)</b>	<b>57 (39-73)</b>	<b>0 (0-13)</b>
Response, %		
▪ CR	11	0
▪ PR	46	0
▪ SD (Wk 12)	32	16
▪ PD	4	44
▪ NE (no 12-wk scan)	7	40
Disease control rate, % (95% CI)	89 (73-96)	16 (6-35)
Median PFS, mos	NR	2.3
Median OS, mos	NR	5.98

# Tumor Mutation Burden As a Predictive Biomarker for Immuno-Oncology Therapies

The NEW ENGLAND JOURNAL of MEDICINE

ORIGINAL ARTICLE

## Genetic Basis for Clinical Response to CTLA-4 Blockade in Melanoma

Alexandra Snyder, M.D., Vladimir Makarov, M.D., Taha Merghoub, Ph.D.,  
Jianda Yuan, M.D., Ph.D., Jesse M. Zaretsky, B.S., Alexis Desrichard, Ph.D.,  
Logan A. Walsh, Ph.D., Michael A. Postow, M.D., Phillip Wong, Ph.D.,  
Teresa S. Ho, B.S., Travis J. Hollmann, M.D., Ph.D., Cameron Bruggeman, M.A.,  
Kasthuri Kannan, Ph.D., Yanyun Li, M.D., Ph.D., Ceyhan Elipenahli, B.S.,  
Cailian Liu, M.D., Christopher T. Harbison, Ph.D., Lisu Wang, M.D.,  
Antoni Ribas, M.D., Ph.D., Jedd D. Wolchok, M.D., Ph.D.,  
and Timothy A. Chan, M.D., Ph.D.

RESEARCH | REPORTS

CANCER IMMUNOLOGY

## Mutational landscape determines sensitivity to PD-1 blockade in non-small cell lung cancer

Naiyer A. Rizvi,<sup>1,2,\*†</sup> Matthew D. Hellmann,<sup>1,2,\*</sup> Alexandra Snyder,<sup>1,2,3,\*</sup> Pia Kvistborg,<sup>4</sup>  
Vladimir Makarov,<sup>3</sup> Jonathan J. Havel,<sup>3</sup> William Lee,<sup>5</sup> Jianda Yuan,<sup>6</sup> Phillip Wong,<sup>6</sup>  
Teresa S. Ho,<sup>6</sup> Martin L. Miller,<sup>7</sup> Natasha Rekhtman,<sup>8</sup> Andre L. Moreira,<sup>8</sup>  
Fawzia Ibrahim,<sup>1</sup> Cameron Bruggeman,<sup>9</sup> Billel Gasmi,<sup>10</sup> Roberta Zappasodi,<sup>10</sup>  
Yuka Maeda,<sup>10</sup> Chris Sander,<sup>7</sup> Edward B. Garon,<sup>11</sup> Taha Merghoub,<sup>1,10</sup>  
Jedd D. Wolchok,<sup>1,2,10</sup> Ton N. Schumacher,<sup>4</sup> Timothy A. Chan<sup>2,3,5,†</sup>

1. Snyder A, et al. *N Engl J Med* 2014;371:2189–2199

2. Rizvi NA, et al. *Science* 2015;348:124–128

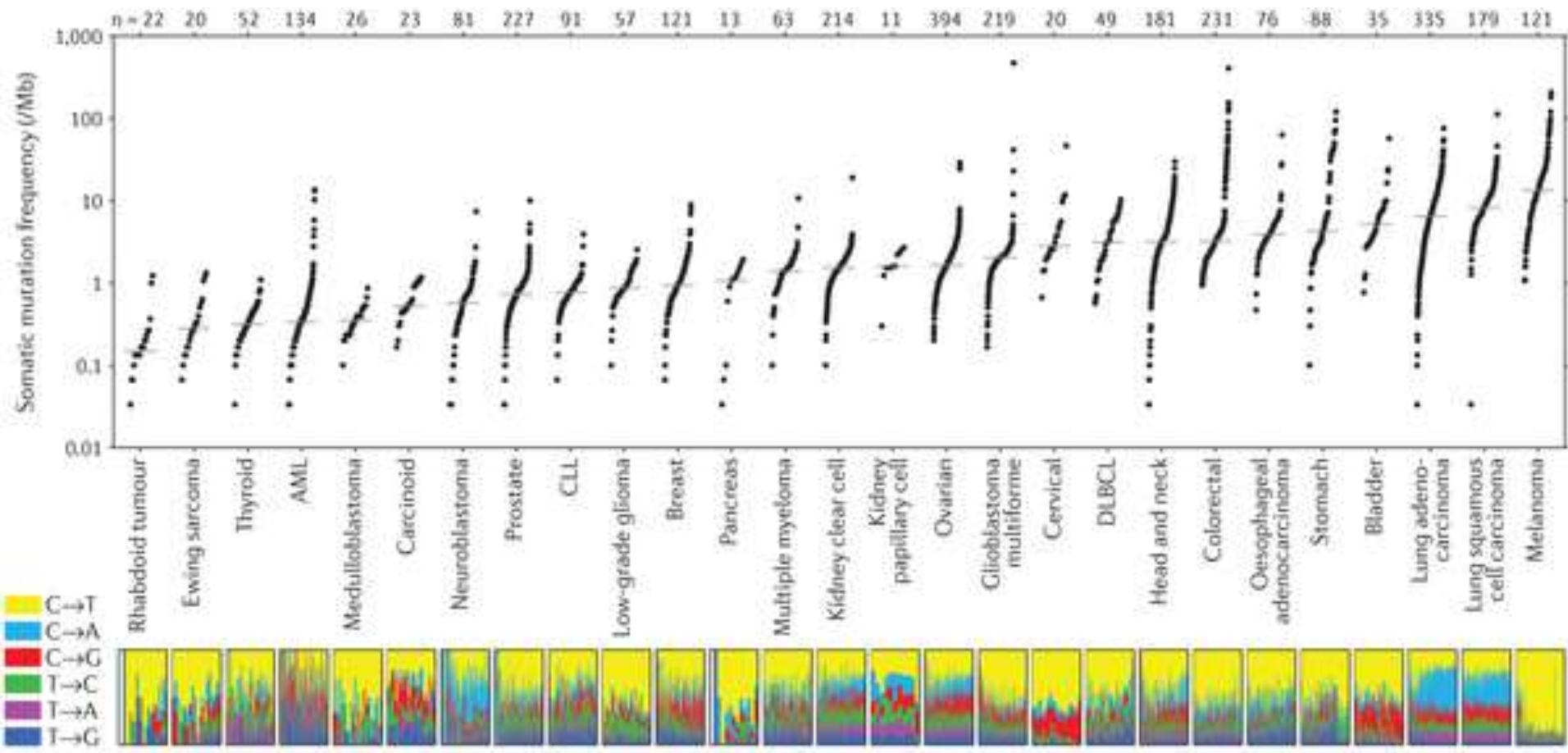
3. Van Allen EM, et al. *Science* 2015;350:207–211

4. Rosenberg JE, et al. *Lancet* 2016;387:1909–1920

5. Hugo W, et al. *Cell* 2016;165:35–44

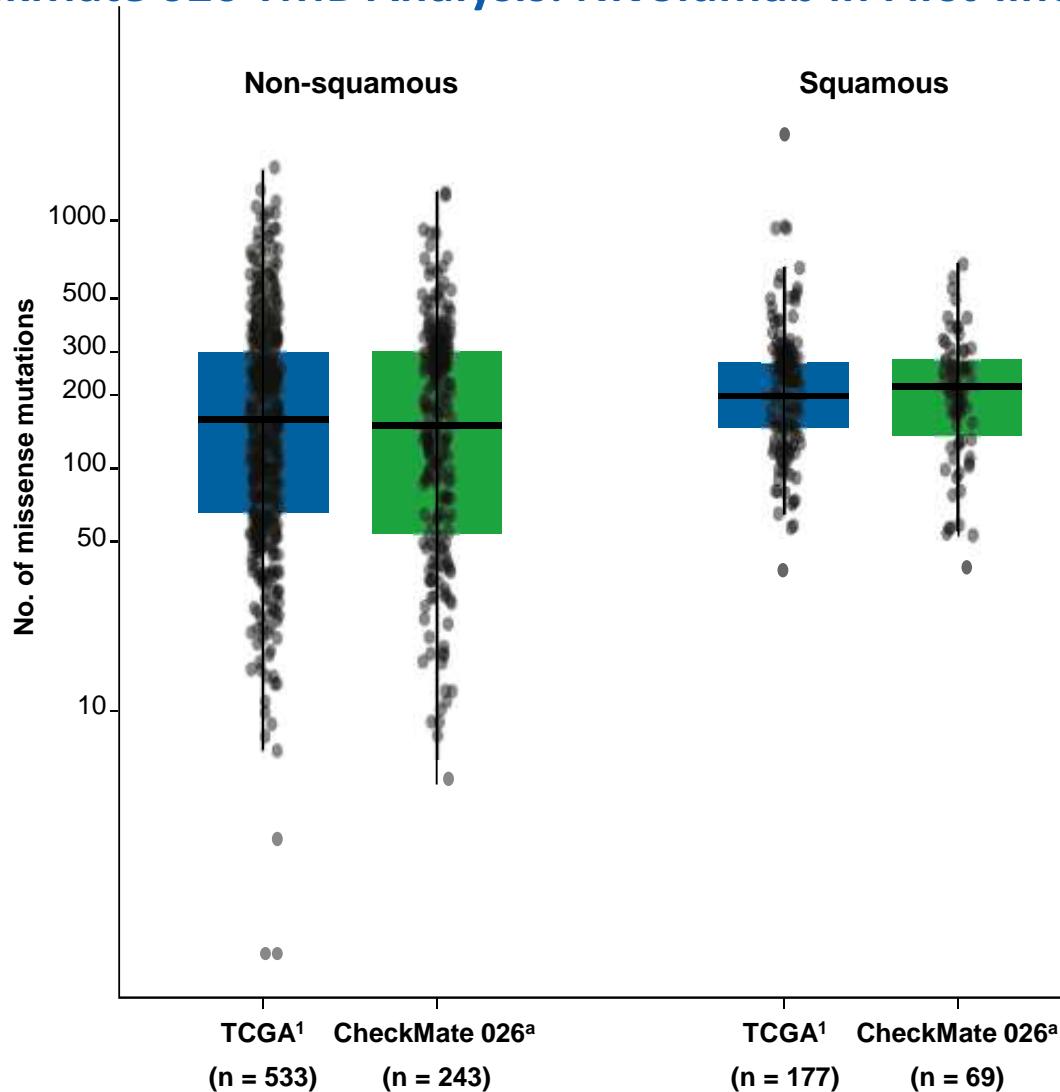
6. Hellmann M. Presented at the 14th International Congress on Targeted Anticancer Therapies; March 21–23, 2016; Washington, DC, USA. Oral O2.2

7. Kowanetz M, et al. Presented at the 2016 IASLC 17th World Conference; December 4–7, 2016; Vienna, Austria. Oral OA20.01



# TMB in The Cancer Genome Atlas<sup>1</sup> and CheckMate 026 Samples<sup>a</sup>

## CheckMate 026 TMB Analysis: Nivolumab in First-line NSCLC

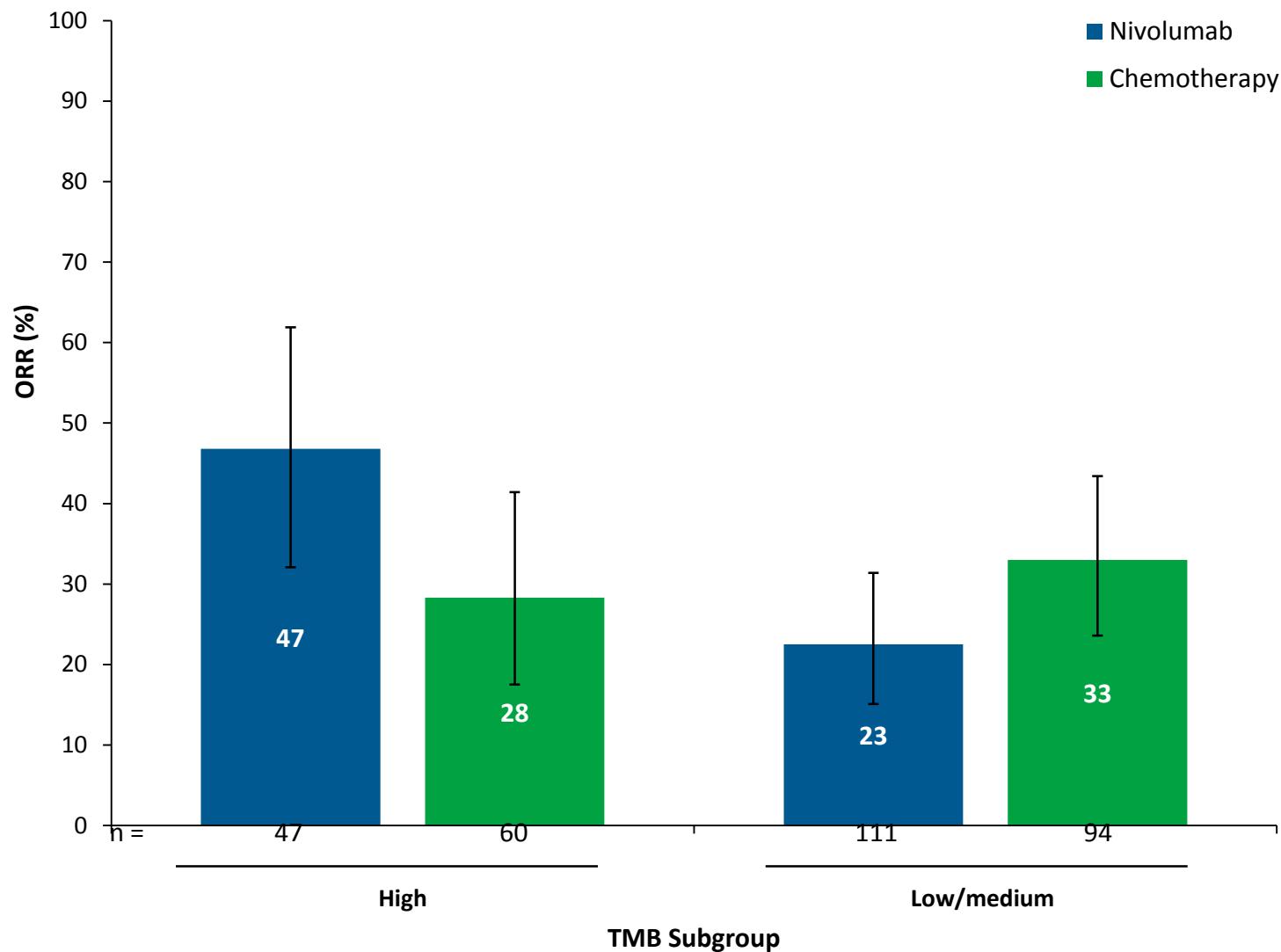


<sup>a</sup>Samples were from whole exome sequencing

1. Broad Institute TCGA Genome Data Analysis Center (2015): Firehose stddata\_\_2015\_02\_04 run. Broad Institute of MIT and Harvard. doi:10.7908/C19P30S6

# ORR by Tumor Mutation Burden Subgroup

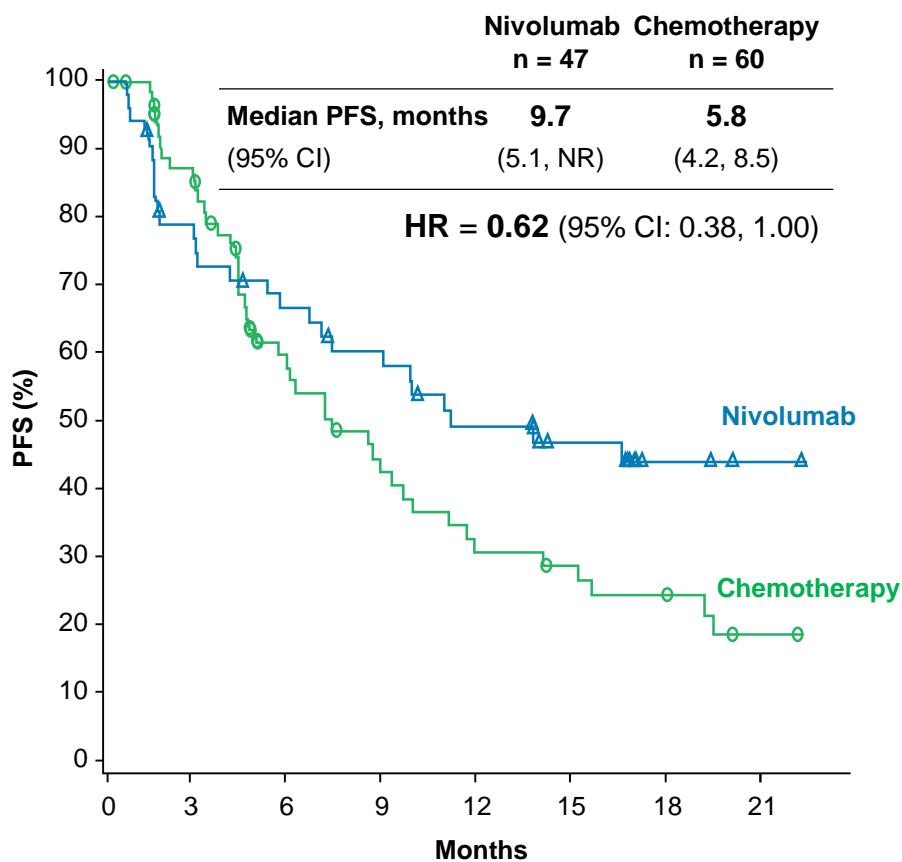
## CheckMate 026 TMB Analysis: Nivolumab in First-line NSCLC



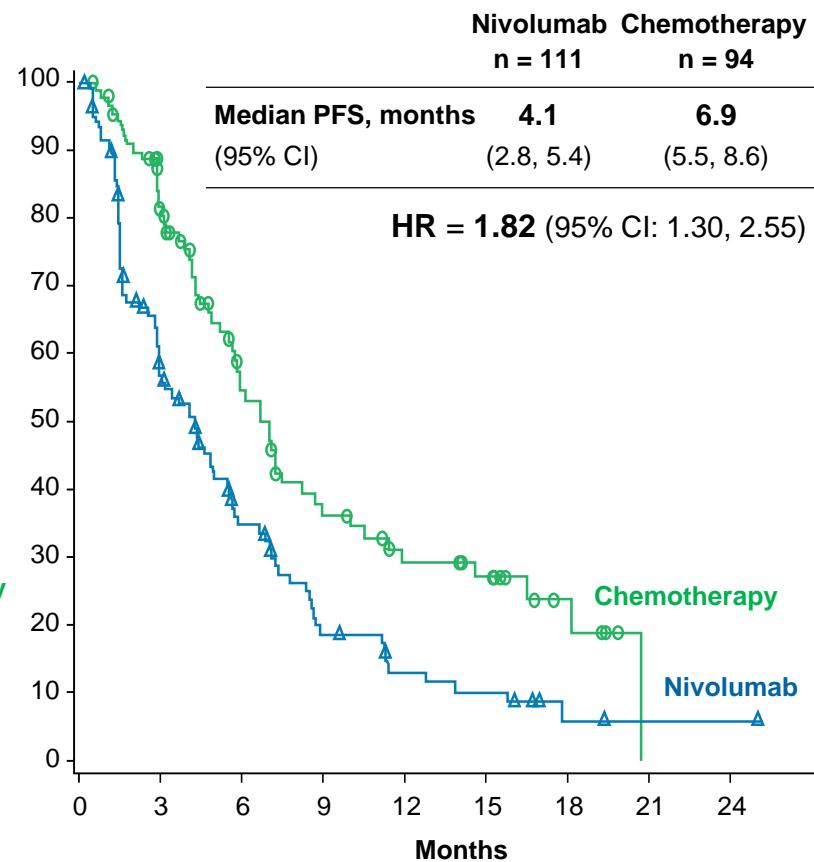
# PFS by Tumor Mutation Burden Subgroup

## CheckMate 026 TMB Analysis: Nivolumab in First-line NSCLC

### High TMB



### Low/medium TMB





A.O.U. Policlinico – Vittorio Emanuele

# The Role of ERCC-1 Polymorphisms as Predictive Biomarker of Response to Nivolumab in Advanced NSCLC

## Abstract ID: 6154

Medical Oncology Unit

A.O.U. Policlinico Vittorio Emanuele Catania, Italy

Vienna, 4 – 7 December 2016



# Methods: Study design

- Stage IIIB/IV NSCLCs
- Previous platinum-based therapy
- $\geq 18$  years
- PS (ECOG) 0 – 1
- Archival tissue availability
- No autoimmune diseases

Evaluation of single nucleotide polymorphisms (SNPs) by pyrosequencing analysis on tumor DNA:  
- **rs11615**  
- **rs3212986**  
- **rs2298881**

- Until PD or unacceptable toxicity
- Treatment beyond progression was allowed in presence of clinical benefit

Nivolumab 3 mg/kg q2w

- Responses were assessed according to RECIST 1.1 criteria → Radiological assessments were performed every 6 weeks.
- PFS was calculated from the beginning of the treatment until PD or patient's death.

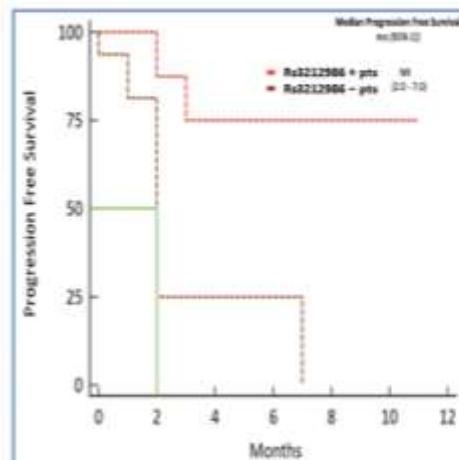


## Results: Objective Response Rate

	All patients	Rs3212986 + patients	Rs3212986 - patients
Type of response	(N = 24)	N= 8	N= 16
Complete Response — no. (%)	1 (4)	1 (12.5)	0
Partial Response — no. (%)	5 (21)	4 (50)	1 (6)
Stable Disease — no. (%)	5 (21)	2 (25)	3 (19)
Progressive Disease — no. (%)	13 (54)	1 (12.5)	12 (75)
ORR (95% CI) — %	25 (10-47)	62.5 (25-92)	6 (0-30)
Disease control rate (95% CI) — %	46 (25-67)	87.5 (47-100)	25 (7-52)

ORR is higher in Rs3212986 + pts than Rs3212986 - pts  
(62.5% VS. 6% p= 0.006)

## Results: Progression Free Survival



PFS is higher in Rs3212986 + pts than  
Rs3212986 - pts  
NR vs. 2.0 months  
(HR = 0.21, 95% CI: 0.07 to 0.58, P < 0.004)



# Conclusioni

- La determinazione del PDL1 ha un valore limitato per la selezione dei pazienti nei tumori solidi
- Nella ricerca biomedica è prioritario individuare nuovi biomarkers per questa tipologia di farmaci
- È necessaria una maggiore conoscenza dei meccanismi di risposta ai farmaci immunoterapici

I° CONVEGNO REGIONALE SIFO

Meeting di primavera

“IL FARMACISTA CLINICO  
E I NUOVI MODELLI DI CURA”



TAORMINA (ME),  
11-12-13 MAGGIO 2017

ID ECM: 313-188877



IV SESSIONE - IMMUNONCOLOGIA

## I biomarcatori nell'immunoncologia: L'importanza della selezione del paziente

*Hector Soto Parra MD  
Oncologia Medica*

