



**PERSPECTIVAS
EM ONCOLOGIA VIII**



**13, 14 e 15 FEV 2020
SHERATON PORTO**

Biomarcadores em Imunoterapia

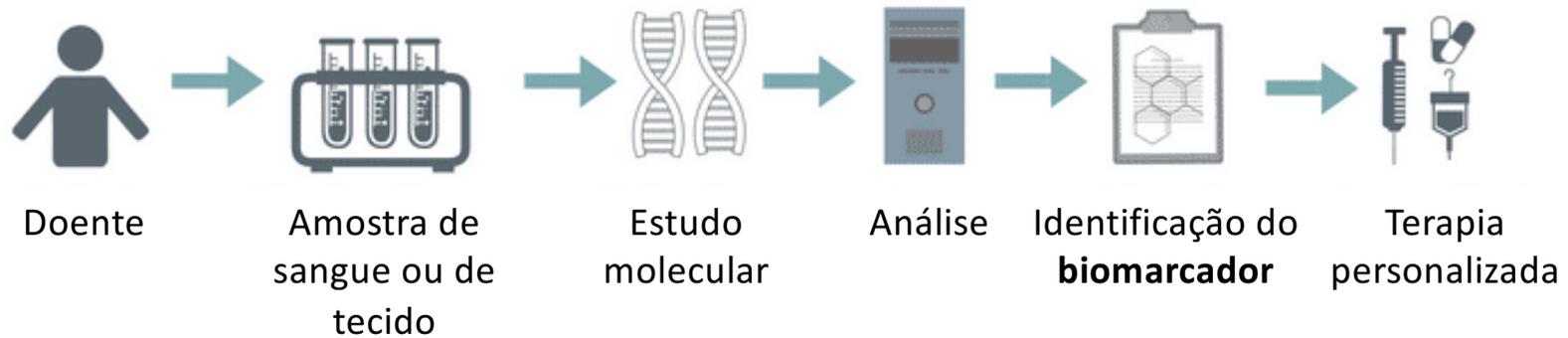
Porto, 13 de Fevereiro 2020



E-mail: hnovaisbastos@med.up.pt
Website: www.heldernovaisbastos.pt



A função do biomarcador



Medicina clássica

One size fits all?



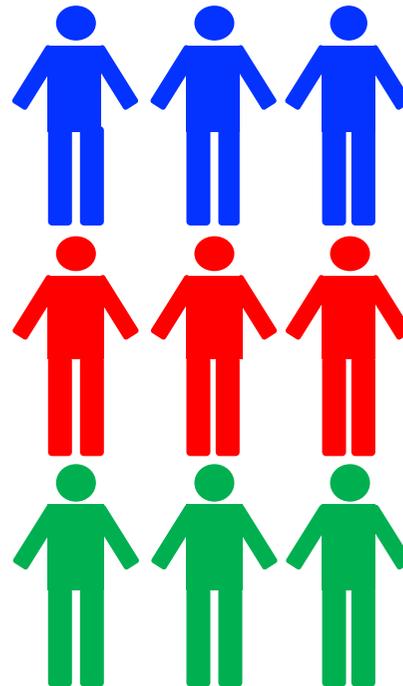
Estratificação

Doentes agrupados por

- perfis clínicos
- subtipos imagiológicos
- subtipos histológicos

Medicina moderna

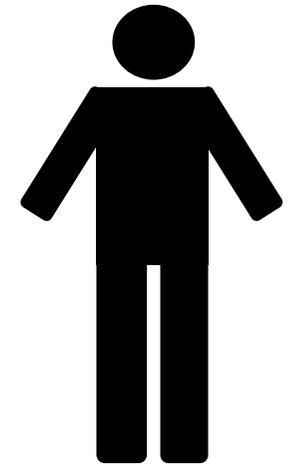
Categorização de doentes



Personalização

Doente individualizado

- perfil histológico
- perfil genético
- biomarcadores

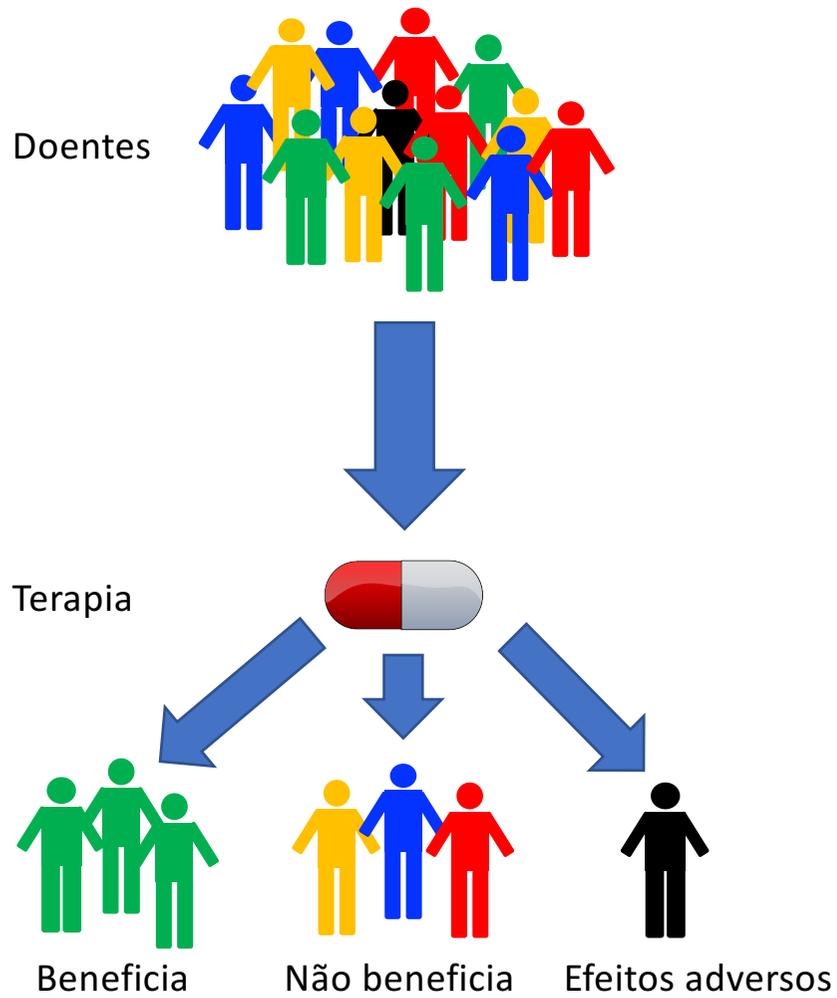


Medicina de futuro

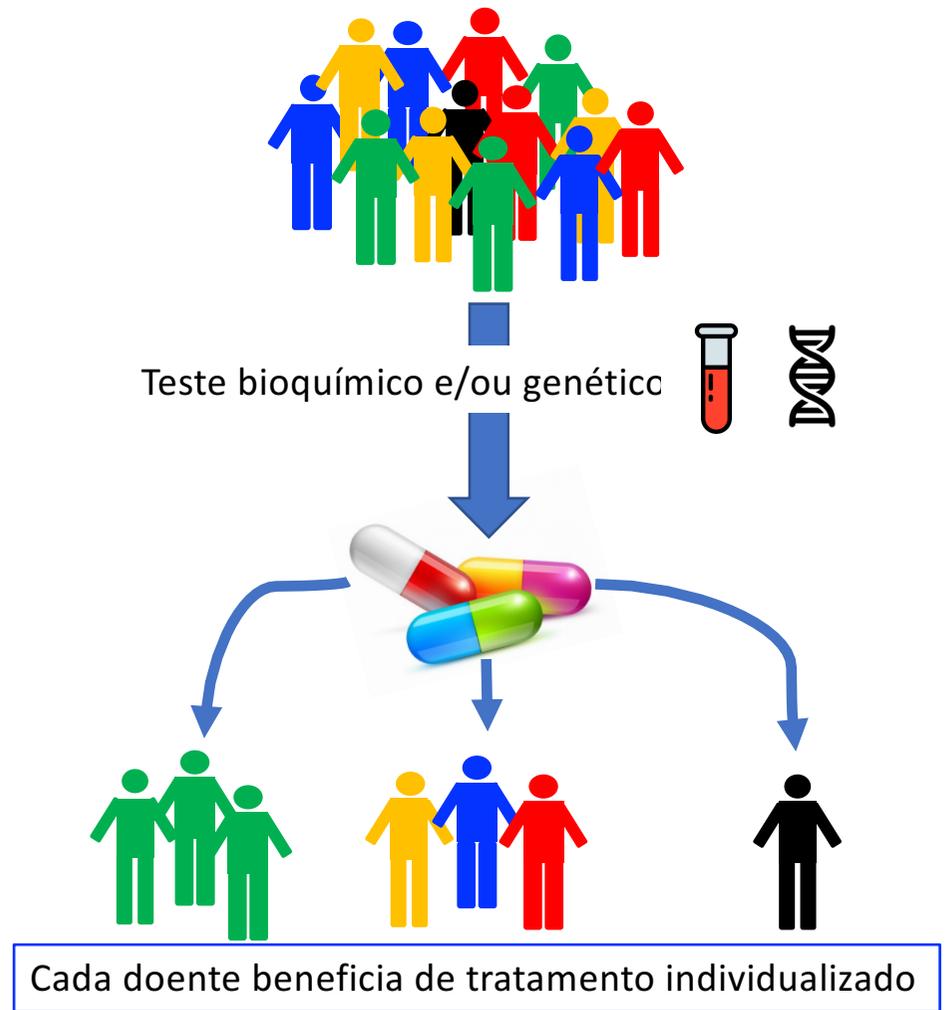
Medicina de precisão

Personalização da medicina no Cancro

Medicina convencional



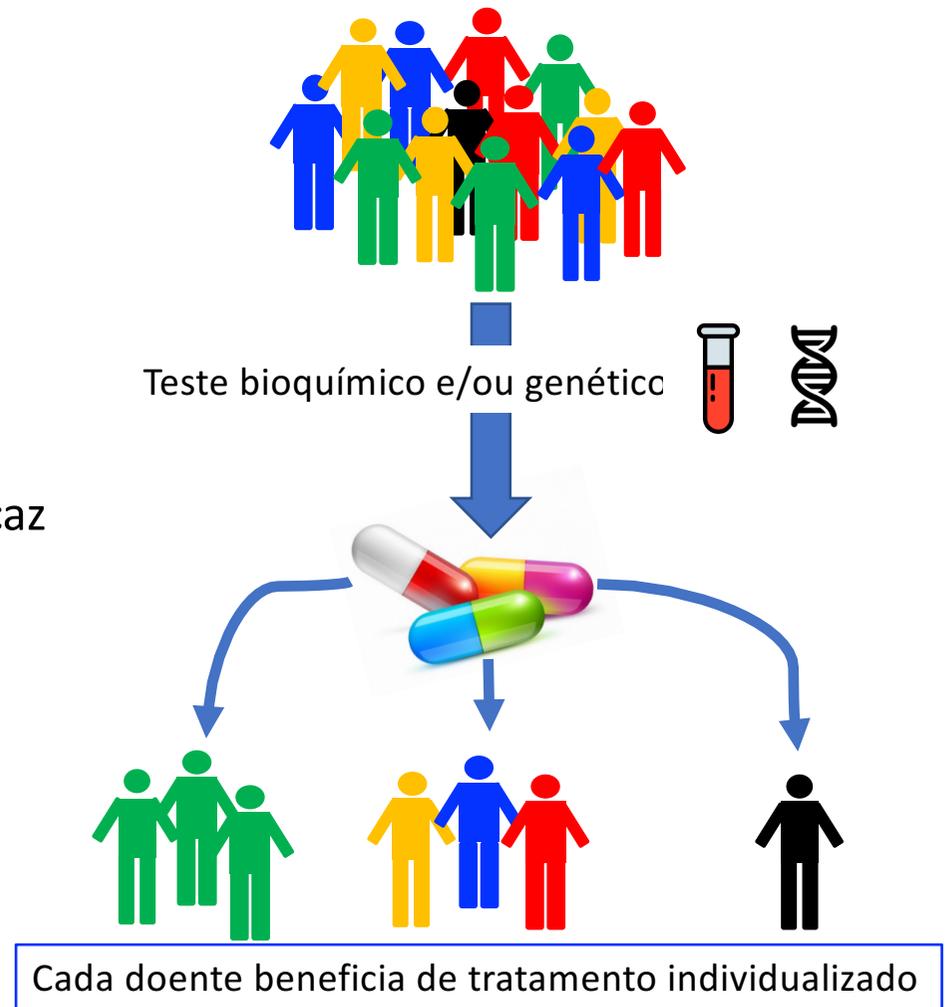
Medicina personalizada



Vantagens da medicina personalizada

- Melhor resultado clínico
- Menor frequência de efeitos adversos
- Previne exposição desnecessária a terapia ineficaz
- Tratamento mais custo-efectivo

Medicina personalizada



Expressão PD-L1

Teff (effector T-cell) gene signature

Tumor Mutational Burden

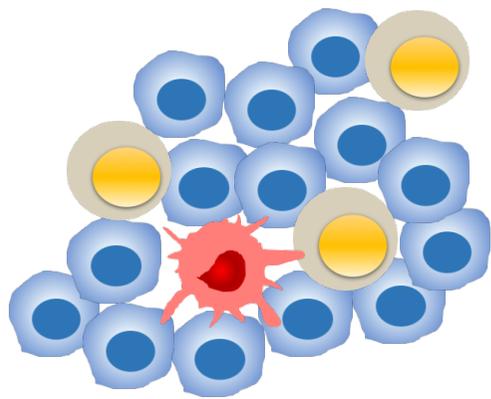
Perspectivas futuras

Expressão PD-L1

Teff (effector T-cell) gene signature

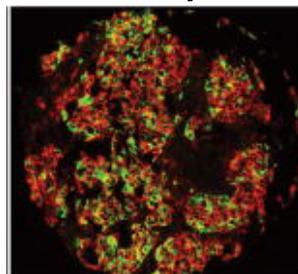
Tumor Mutational Burden

Perspectivas futuras



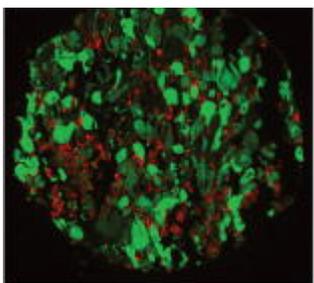
Microambiente tumoral

Cancro do pulmão

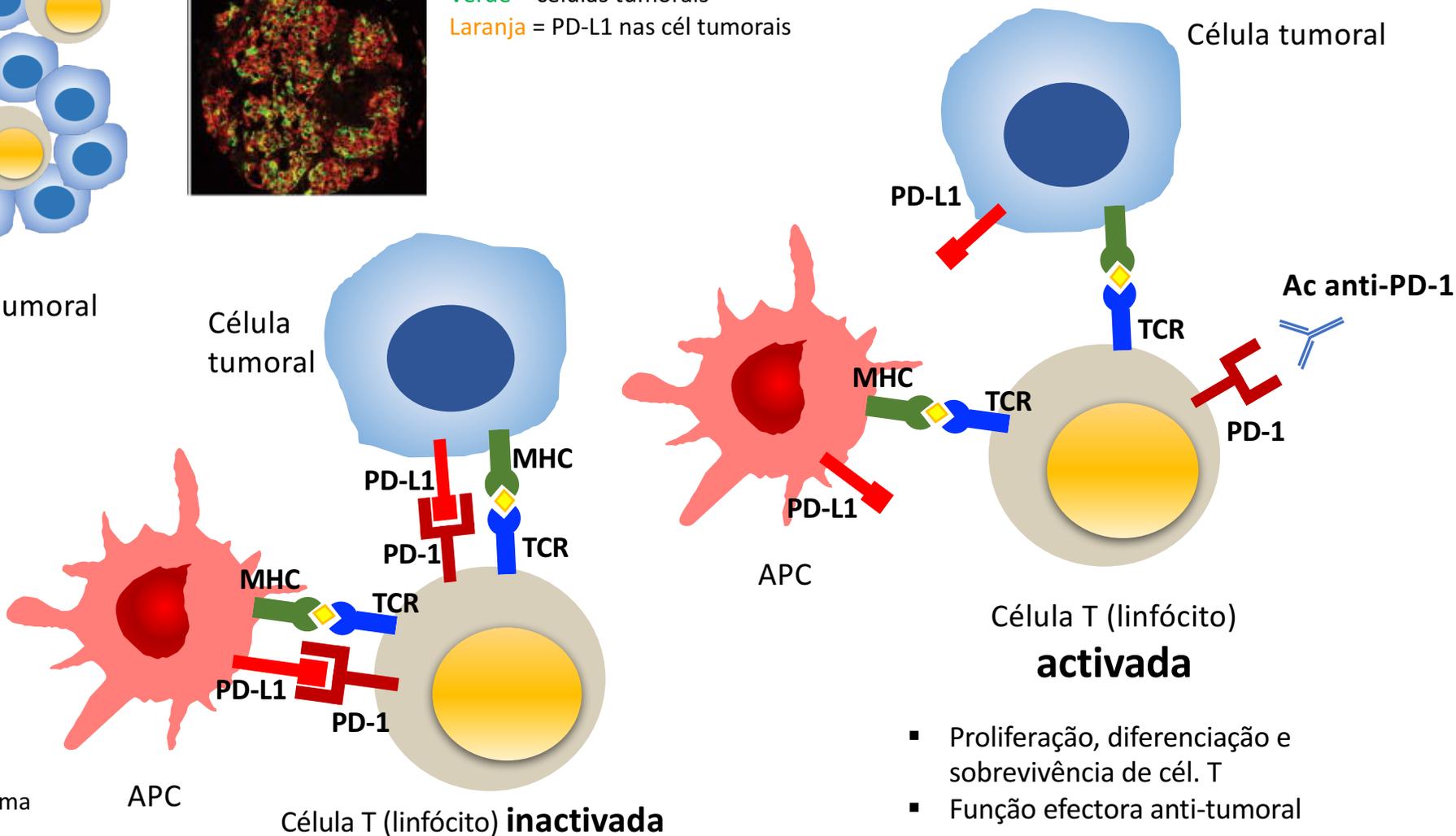


Verde = células tumorais
 Laranja = PD-L1 nas cél tumorais

Melanoma

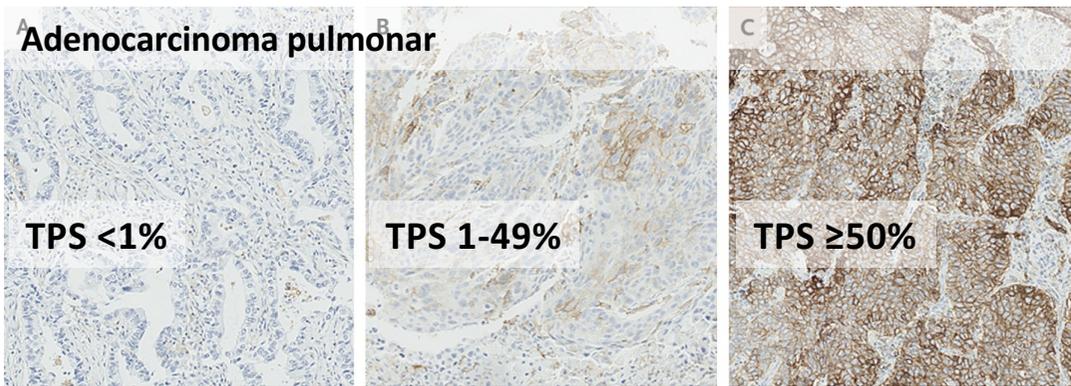


Verde = células tumorais
 Vermelho = PD-L1 no estroma



A importância do biomarcador revelada no KEYNOTE-001

Expressão PD-L1 testada através de ensaio de imuno-histoquímica pharmDx (PD-L1 IHC 22C3; Dako/Agilent)



$$\text{TPS}(\%) = \frac{\text{No. PD-L1-stained tumor cells}}{\text{Total No. of viable tumor cells}} \times 100$$

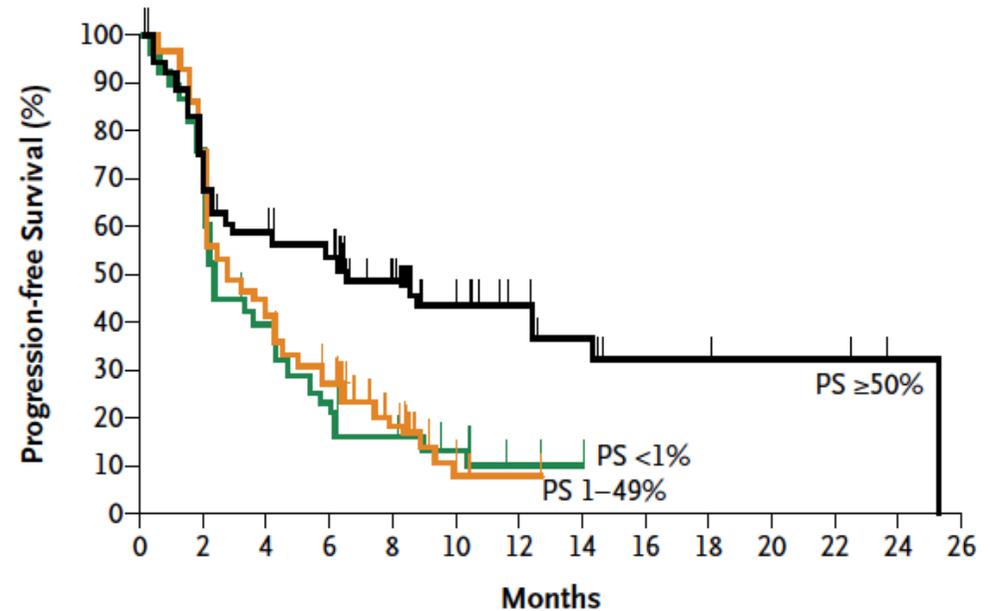


Pembrolizumab provado no CPCNP

- 2ª linha monoterapia: TPS ≥1 %
- 1ª linha monoterapia: TPS ≥50 %

EUROPEAN MEDICINES AGENCY
SCIENCE. MEDICINES. HEALTH.

A All Patients



No. at Risk

PS ≥50%	119	86	66	60	38	20	13	8	4	3	3	3	1	0
PS 1-49%	161	122	70	45	21	4	1	0	0	0	0	0	0	0
PS <1%	76	52	29	17	11	6	2	0	0	0	0	0	0	0

Garon EB et al. NEJM 2015

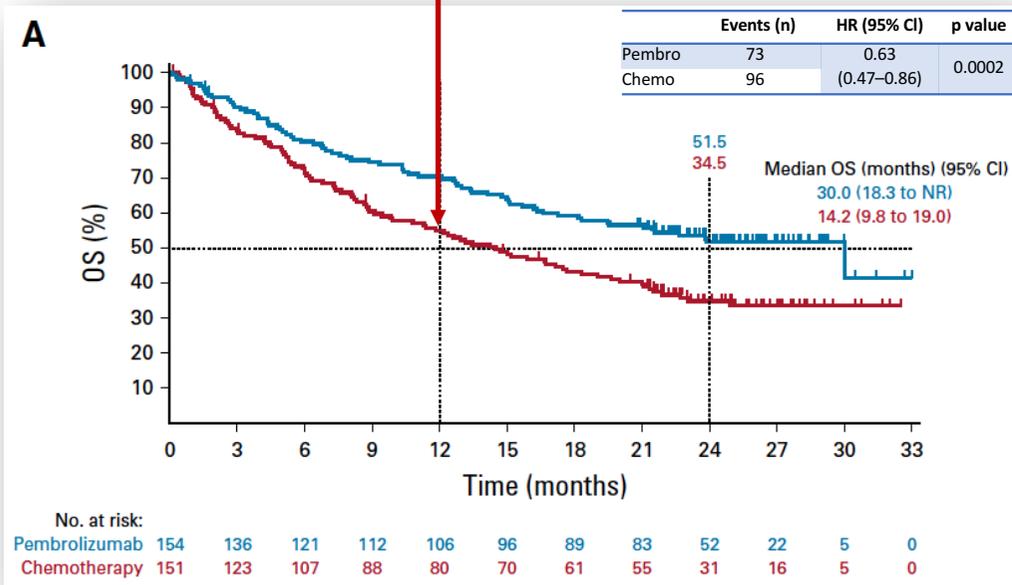
Impacto do PD-L1 como biomarcador de resposta a Pembrolizumab (1L)

Tratamento de doente com tumor PD-L1+ e sem alvo mutacional até em 2016:

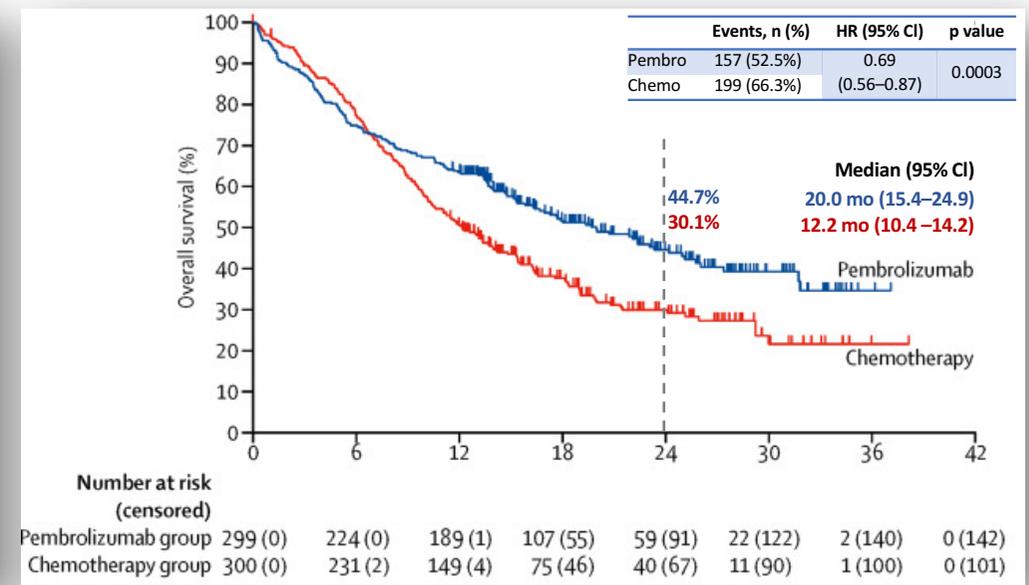


Morte

KEYNOTE-024

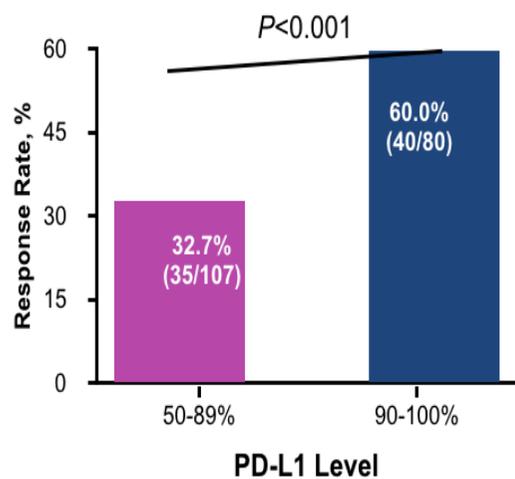


KEYNOTE-042



Pembrolizumab monoterapia 1L CPCNP: PD-L1 $\geq 90\%$ vs 50-89%

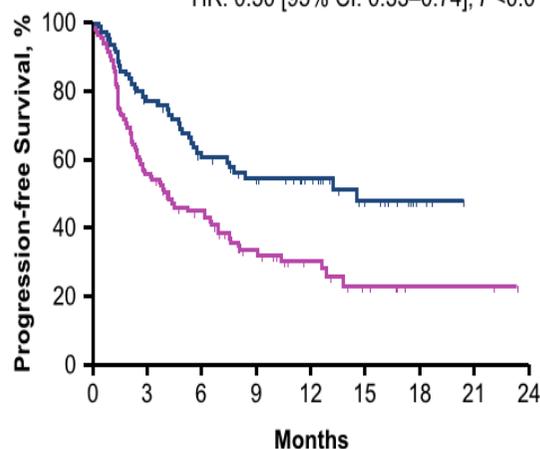
ORR



PFS

	N	mPFS (95% CI)
PD-L1 90-100%	80	14.5 mo (6.0-NR)
PD-L1 50-89%	107	4.1 mo (1.7-6.6)

HR: 0.50 [95% CI: 0.33-0.74], $P < 0.01$

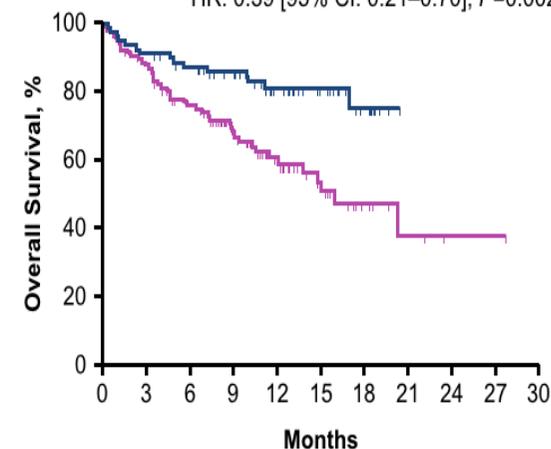


	No. at risk	0	3	6	9	12	15	18	21	24
PD-L1 90-100%	80	58	44	34	27	12	3	0	0	0
PD-L1 50-89%	107	59	42	25	13	6	2	2	0	0

OS

	N	mOS (95% CI)
PD-L1 90-100%	80	NR (NR-NR)
PD-L1 50-89%	107	15.9 mo (11.2-20.7)

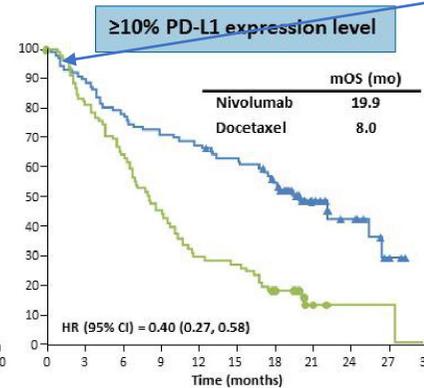
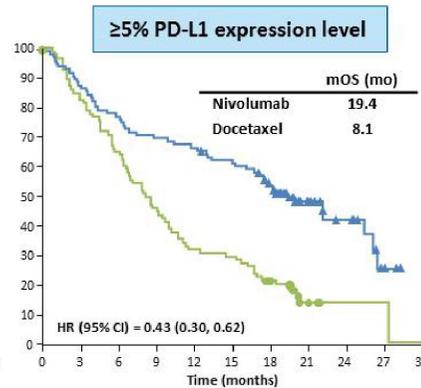
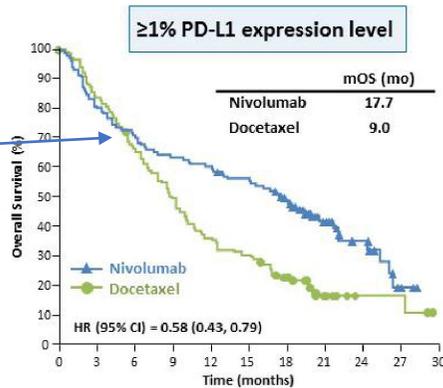
HR: 0.39 [95% CI: 0.21-0.70], $P = 0.002$



	No. at risk	0	3	6	9	12	15	18	21	24	27	30
PD-L1 90-100%	80	73	66	57	38	22	10	0	0	0	0	0
PD-L1 50-89%	107	92	75	51	33	18	8	4	1	1	0	0

Impacto da expressão PD-L1 na OS por Nivolumab (2L): CheckMate 057 trial

Efeito tardio do tratamento

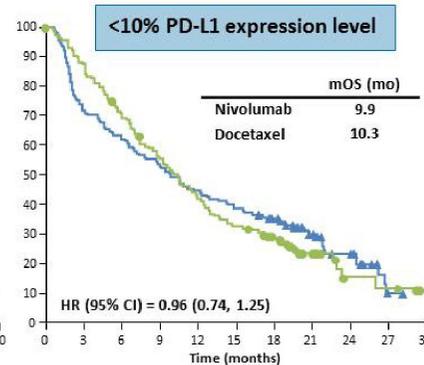
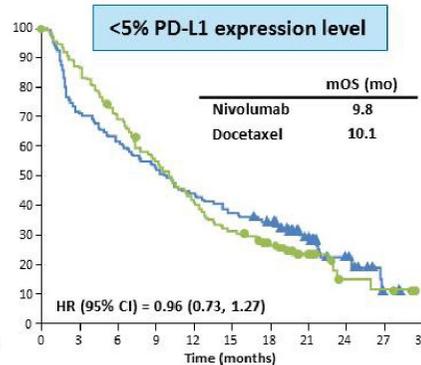
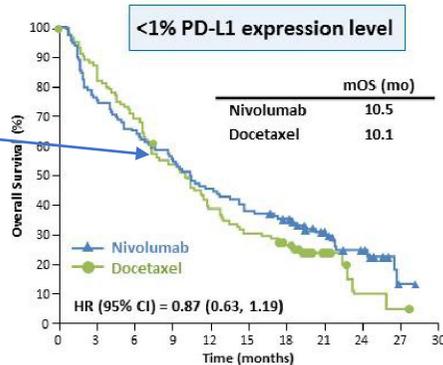


Efeito precoce do tratamento

Number of patients at risk

Nivolumab	123	99	87	78	74	68	56	24	13	3	0	95	83	73	66	63	58	47	20	12	3	0	86	77	67	61	58	53	44	19	11	3	0
Docetaxel	123	102	80	61	44	37	24	8	3	3	0	86	70	55	39	27	28	17	4	1	1	0	79	63	50	35	23	21	13	3	1	1	0

Efeito reverso do tratamento com o tempo

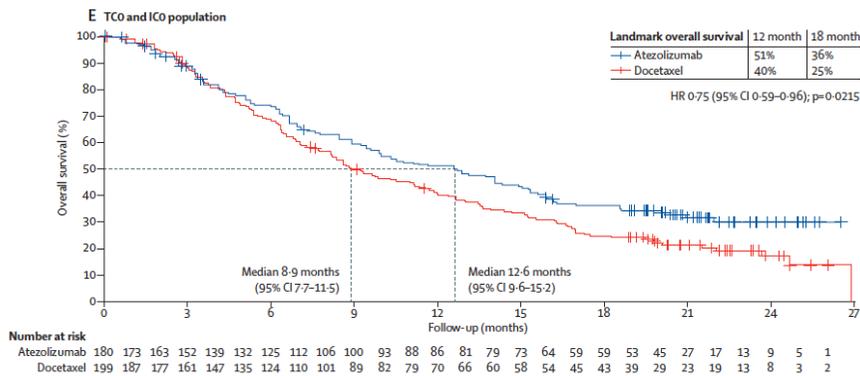


Number of patients at risk

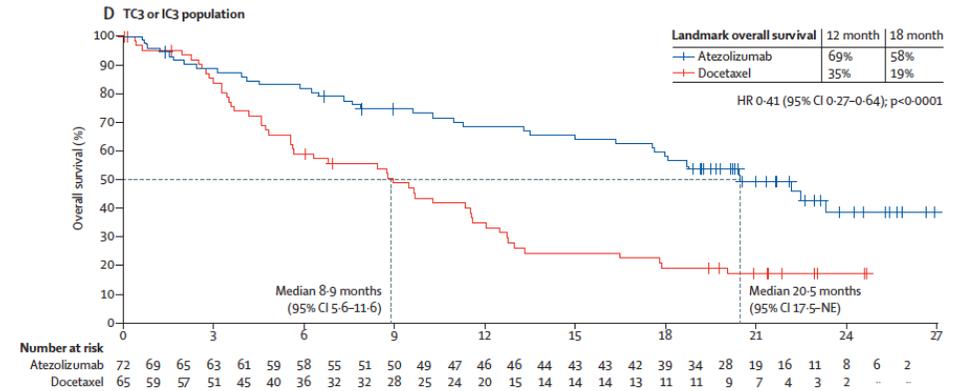
Nivolumab	108	82	70	61	49	41	36	23	13	1	0	136	98	84	73	60	51	45	27	14	1	0	145	104	90	78	65	56	48	28	15	1	0
Docetaxel	101	87	69	53	38	30	24	11	2	1	0	138	119	94	75	55	42	31	15	4	3	0	145	126	99	79	59	46	35	16	4	3	0

PD-L1 NÃO tem impacto na OS por Atezolizumab (2L): OAK trial

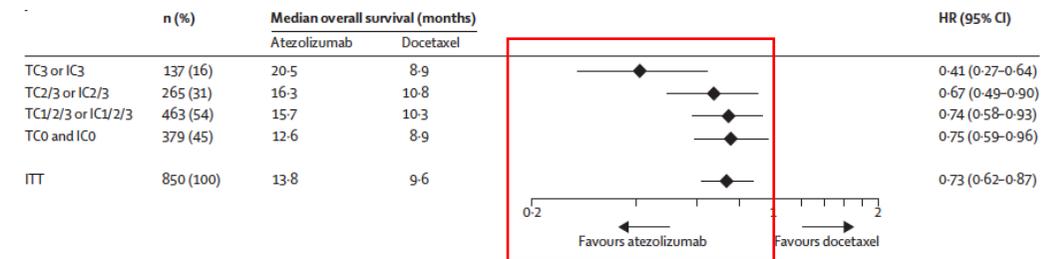
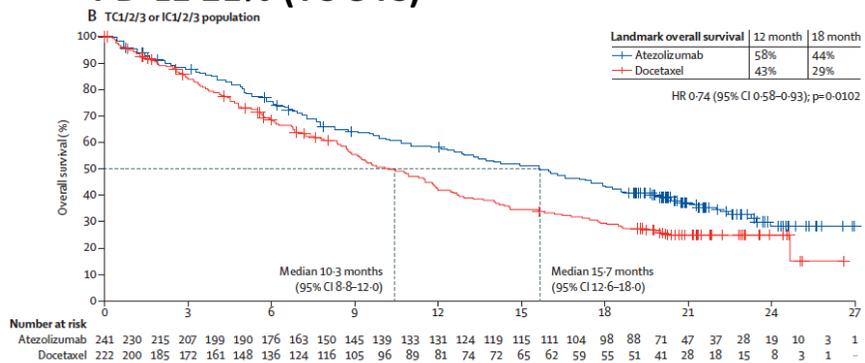
PD-L1 <1% (TC e IC)



PD-L1 ≥50% TC ou ≥10% IC



PD-L1 ≥1% (TC e IC)



TC: tumour cells; IC: tumour-infiltrating immune cells
 PD-L1 Antibody: VENTANA SP142 PD-L1 immunohistochemistry assay

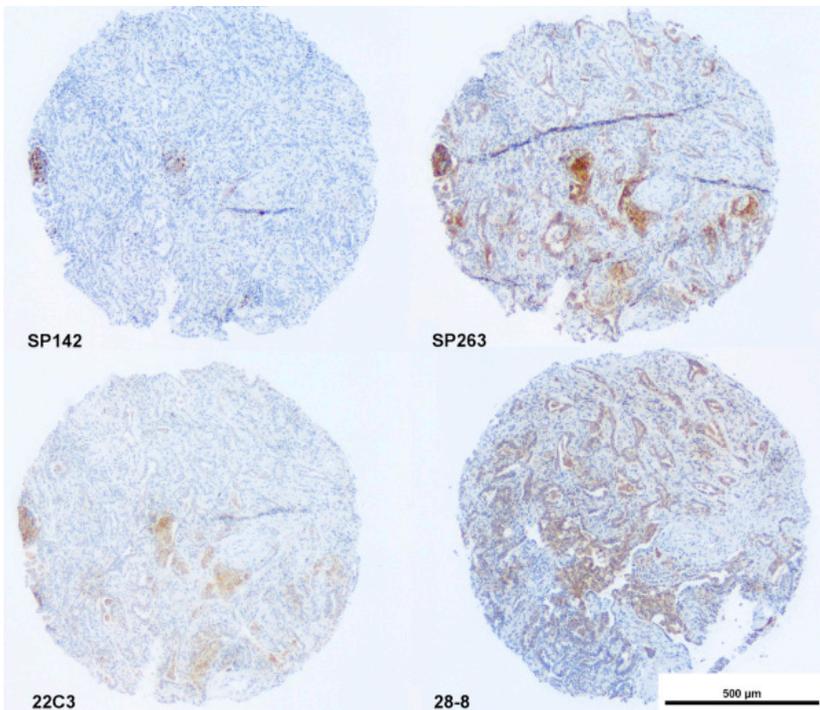
Desafios na utilização de PD-L1 como biomarcador

Variabilidade inter-ensaio

Clones de anticorpo usados nos estudos clínicos:

SP142 (Atezolizumab)

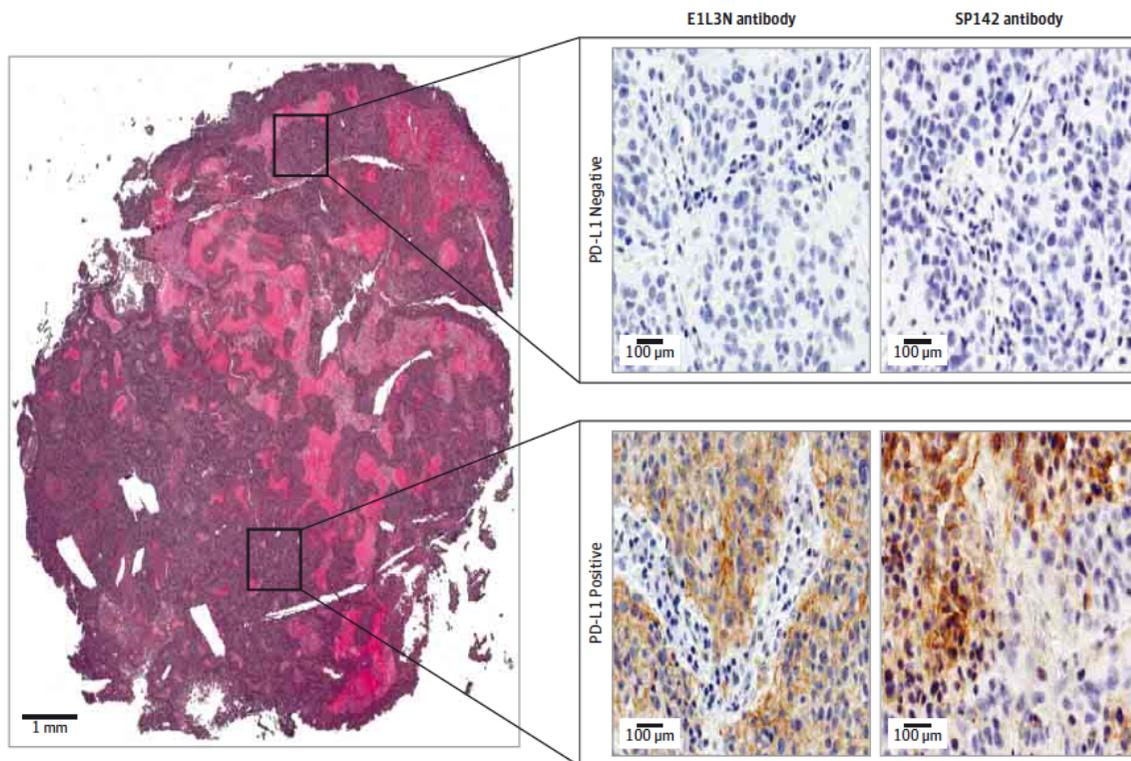
SP263 (Durvalumab)



22C3 (Pembrolizumab)

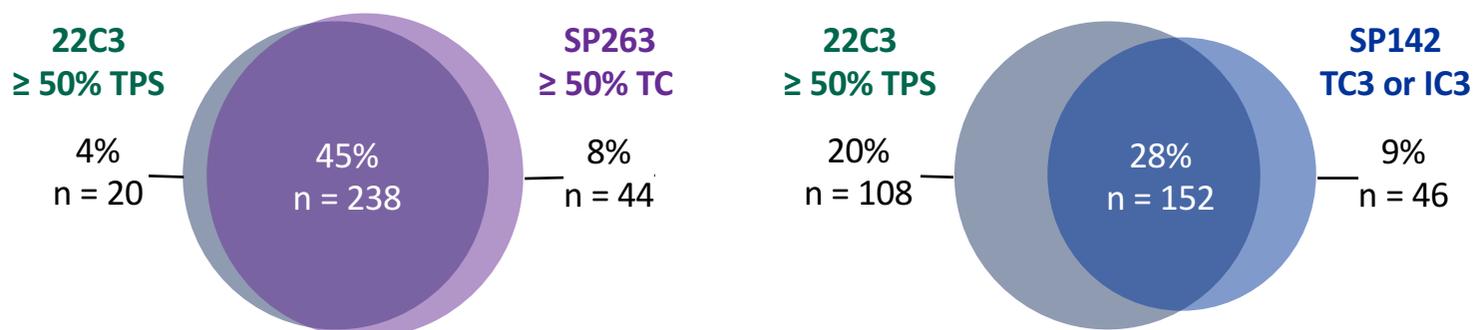
Clone 28-8 (Nivolumab)

Heterogeneidade de expressão no mesmo tumor



Hendry S et al. J Thorac Oncol. 2018
McLaughlin J et al. JAMA Oncol. 2016

Sobreposição entre subgrupos de expressão PD-L1 alta (IMpower110)



- **Elevada sobreposição entre 22C3 e SP263** observada no subgrupo de expressão $\geq 50\%$
- Uma grande proporção do subgrupo SP142 TC3 /IC3 abrangeu também o subgrupo 22C3 $\geq 50\%$ TPS.

Clones de anticorpo usados nos estudos clínicos:

22C3 (Pembrolizumab)

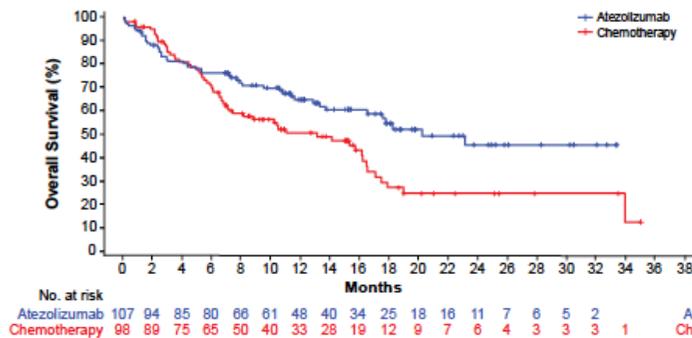
SP263 (Durvalumab)

SP142 (Atezolizumab)

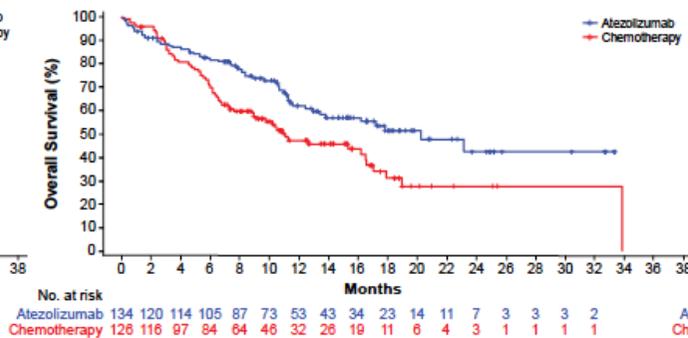
22C3 and SP263 overlap: BEP within the ITT-WT population, n = 530.
22C3 and SP142 overlap: BEP within the ITT-WT population, n = 534.

OS em subgrupos de expressão PD-L1 alta no IMpower110 (1L)

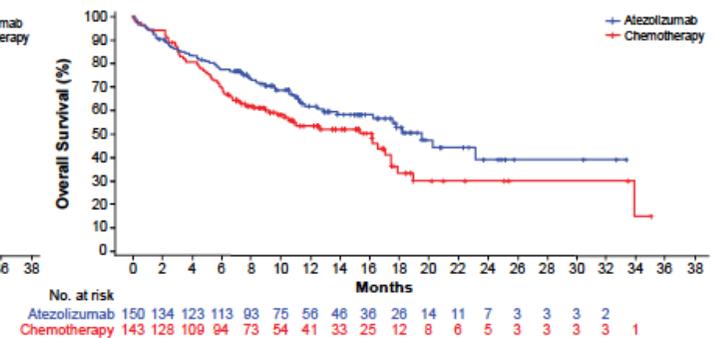
SP142 (TC3 or IC3-WT)^a



22C3 BEP-WT (TPS ≥ 50%)^a



SP263 BEP-WT (TC ≥ 50%)^a



	Atezo (n = 107)	Chemo (n = 98)
mOS, mo	20.2	13.1
HR ^b (95% CI)	0.59 (0.40, 0.89)	

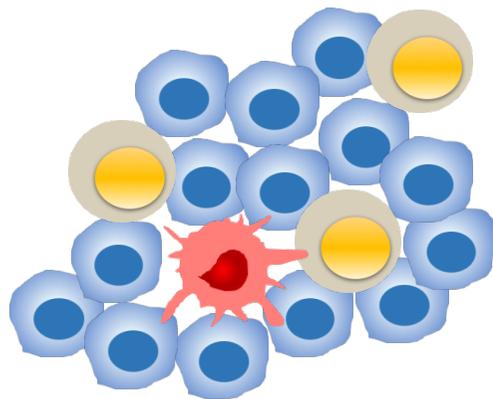
	Atezo (n = 134)	Chemo (n = 126)
mOS, mo	20.2	11.0
HR ^c (95% CI)	0.60 (0.41, 0.86)	

	Atezo (n = 150)	Chemo (n = 143)
mOS, mo	19.5	16.1
HR ^c (95% CI)	0.71 (0.50, 1.00)	

^a SP142 TC1/2/3 or IC1/2/3-WT (n = 554); 22C3 BEP-WT (n = 534); SP263 BEP-WT (n = 546). ^b Stratified. ^c Unstratified.

Clones de anticorpo usados nos estudos clínicos:
22C3 (Pembrolizumab)
SP263 (Durvalumab)
SP142 (Atezolizumab)

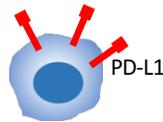
Quimioterapia
+
Inibidores PD-1/PD-L1



Microambiente tumoral



↑ Células apresentadoras de antígeno (APC)



↑ **Expressão de PD-L1 nas células tumorais**

↓ Células tumorais



↑ Células T CD8+

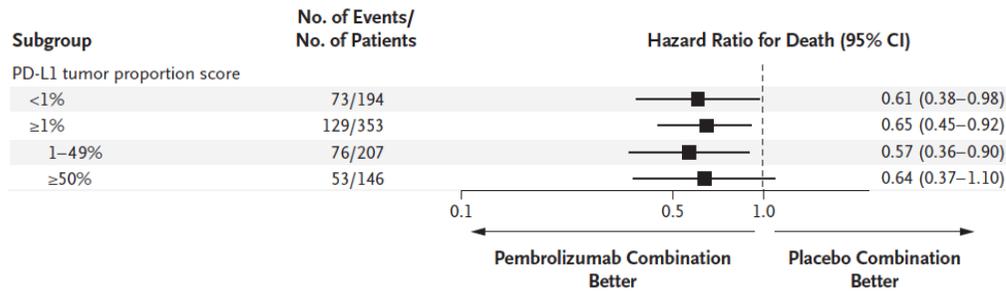
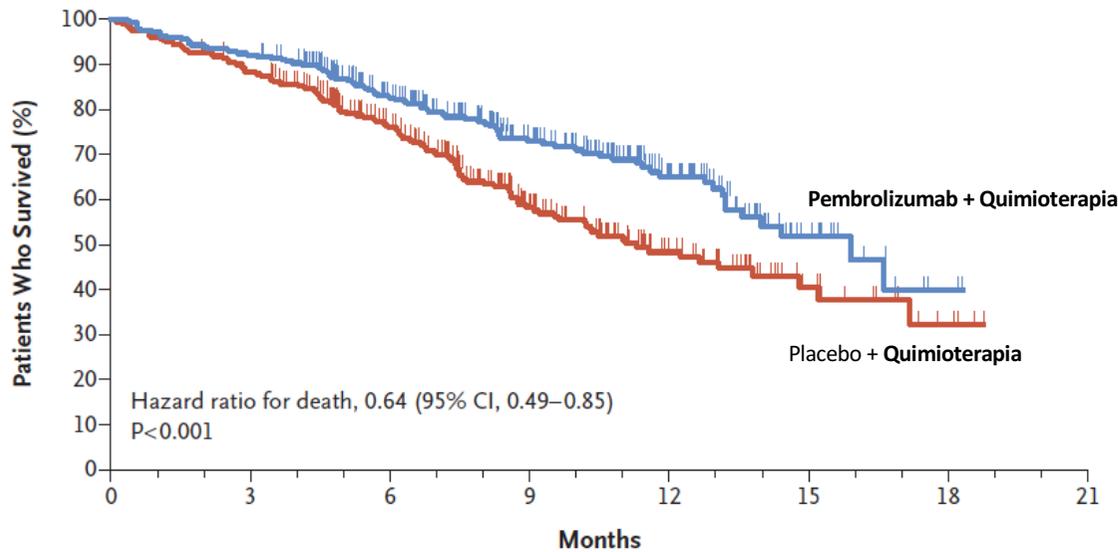
↓ Células Treg ou MDSCs



↑ Macrófagos

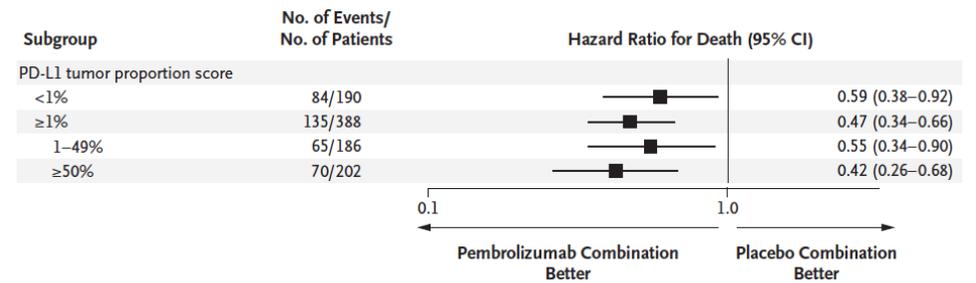
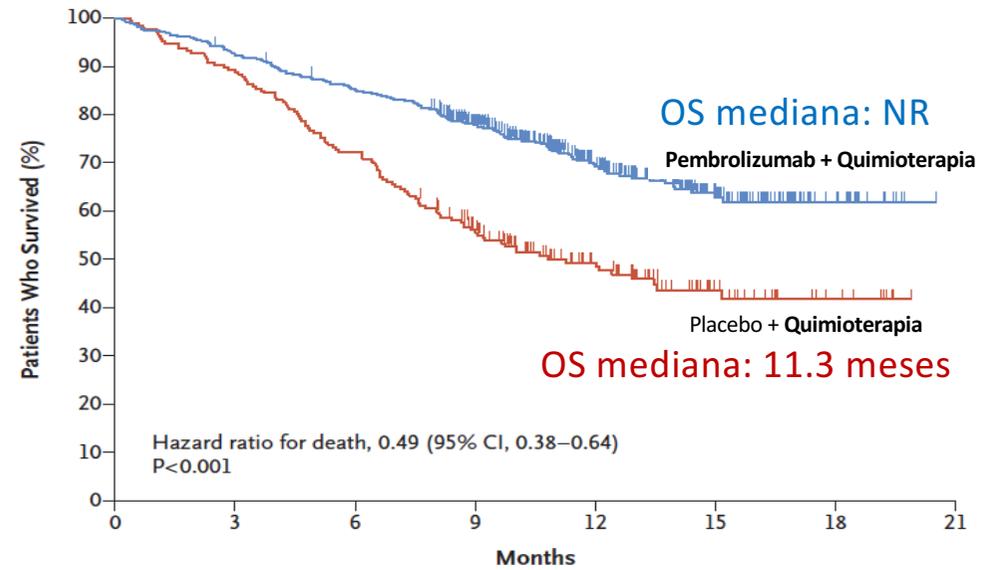
Combinações QT+IT

Squamous Non-Small-Cell Lung Cancer (KN-407)



Paz-Ares L et al, NEJM 2018

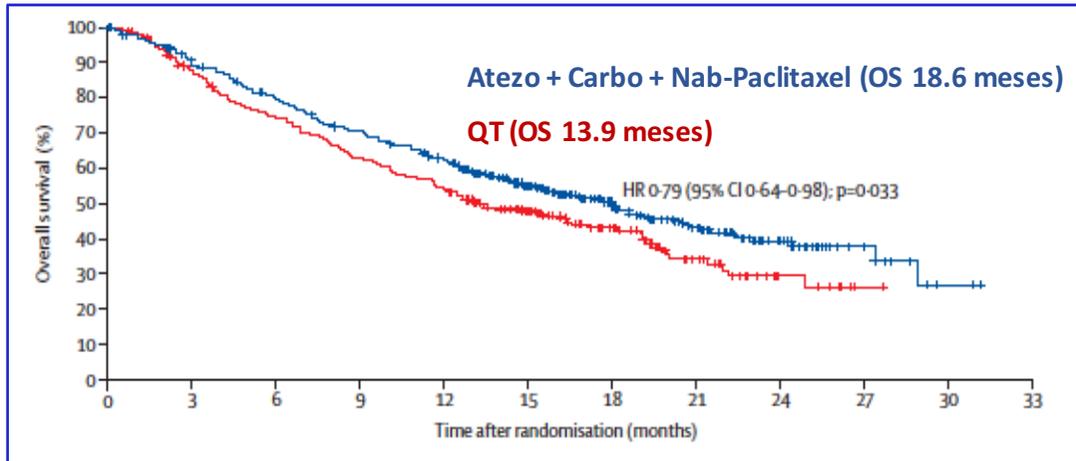
Nonsquamous NSCLC EGFR/ALK negative (KN-189)



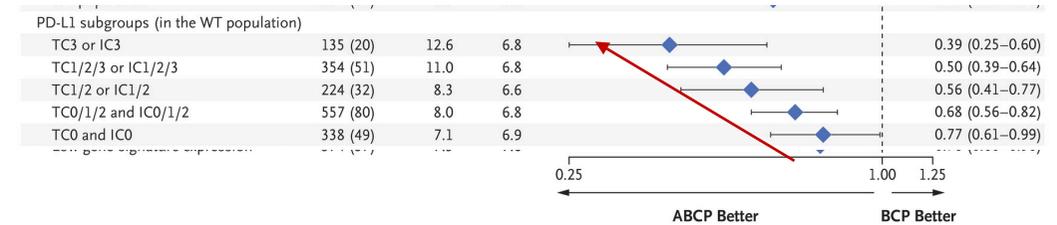
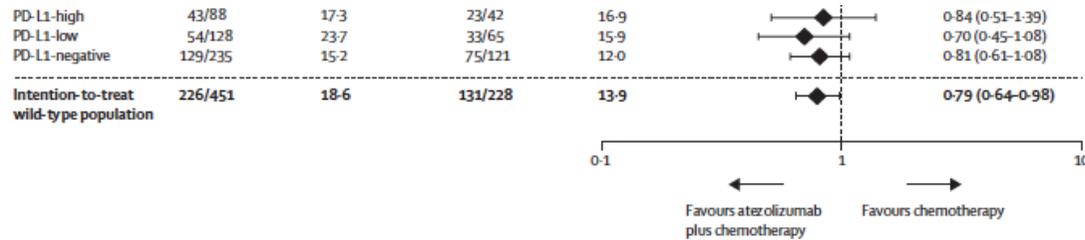
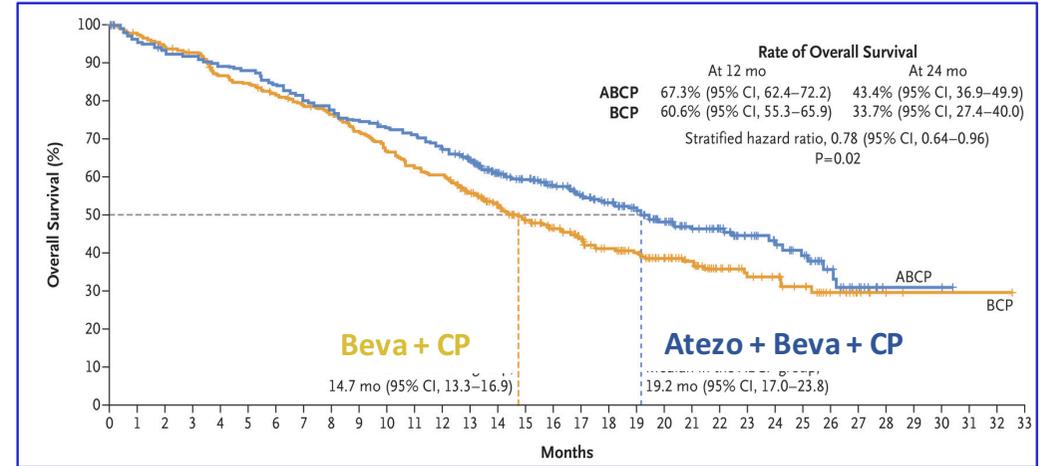
Gandhi L et al, NEJM 2018

Combinações QT+IT

IMpower 130



IMpower 150



West et al. Lancet Oncol 2019; Socinski et al. NEJM 2018

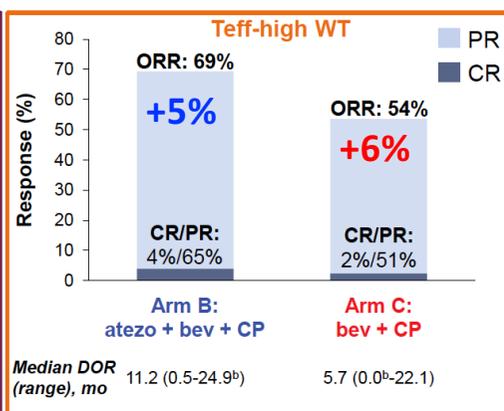
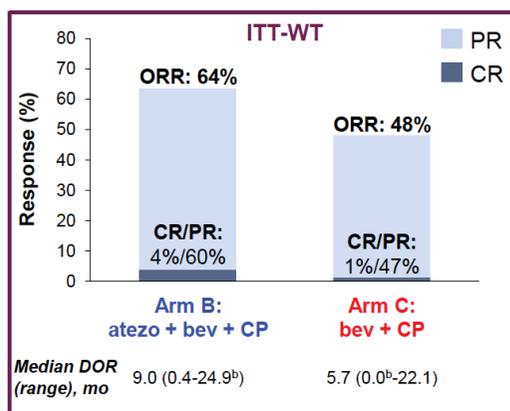
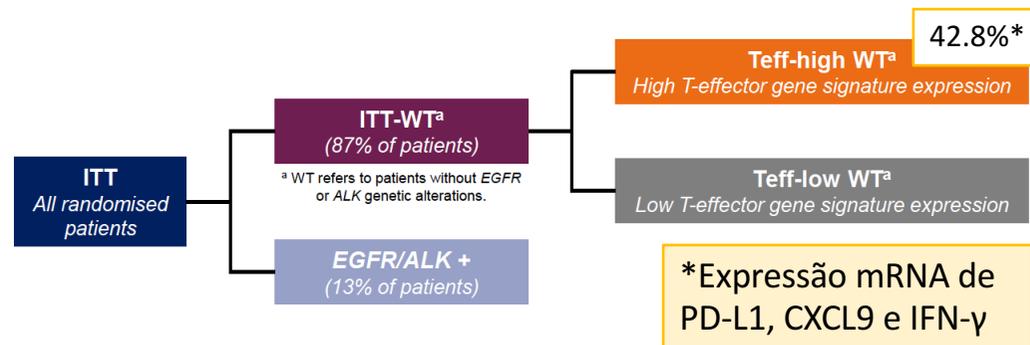
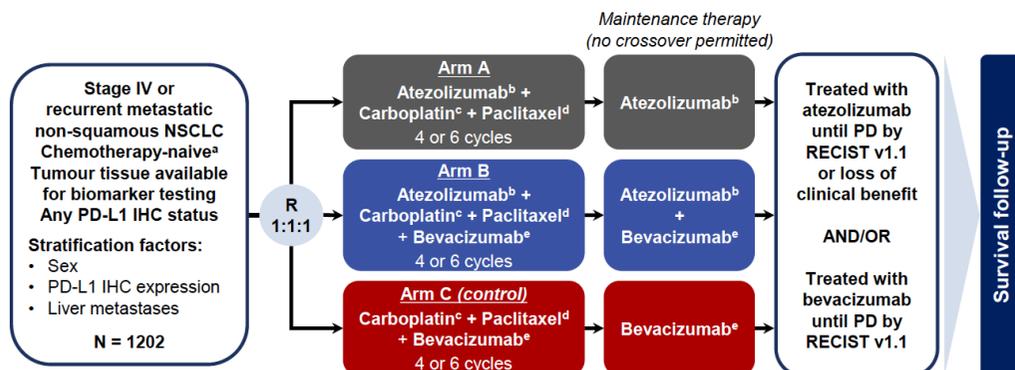
Expressão PD-L1

Teff (effector T-cell) gene signature

Tumor Mutational Burden

Perspectivas futuras

Teff (effector T-cell) gene signature - IMpower 150



B Hazard Ratios for Disease Progression or Death in Biomarker Subgroups

Population	No. of Patients (%)	Median Progression-free Survival (mo)		Hazard Ratio (95% CI)
		ABCP	BCP	
ITT population	800 (100)	8.3	6.8	0.61 (0.52-0.72)
Patients with EGFR or ALK genetic alterations	108 (14)	9.7	6.1	0.59 (0.37-0.94)
WT population	692 (87)	8.3	6.8	0.62 (0.52-0.74)
PD-L1 subgroups (in the WT population)				
TC3 or IC3	135 (20)	12.6	6.8	0.39 (0.25-0.60)
TC1/2/3 or IC1/2/3	354 (51)	11.0	6.8	0.50 (0.39-0.64)
TC1/2 or IC1/2	224 (32)	8.3	6.6	0.56 (0.41-0.77)
TC0/1/2 and IC0/1/2	557 (80)	8.0	6.8	0.68 (0.56-0.82)
TC0 and IC0	338 (49)	7.1	6.9	0.77 (0.61-0.99)
Teff subgroups (in the WT population)				
High gene-signature expression	284 (43)	11.3	6.8	0.51 (0.38-0.68)
Low gene-signature expression	374 (57)	7.3	7.0	0.76 (0.60-0.96)

0.25 1.00 1.25

ABCP Better BCP Better

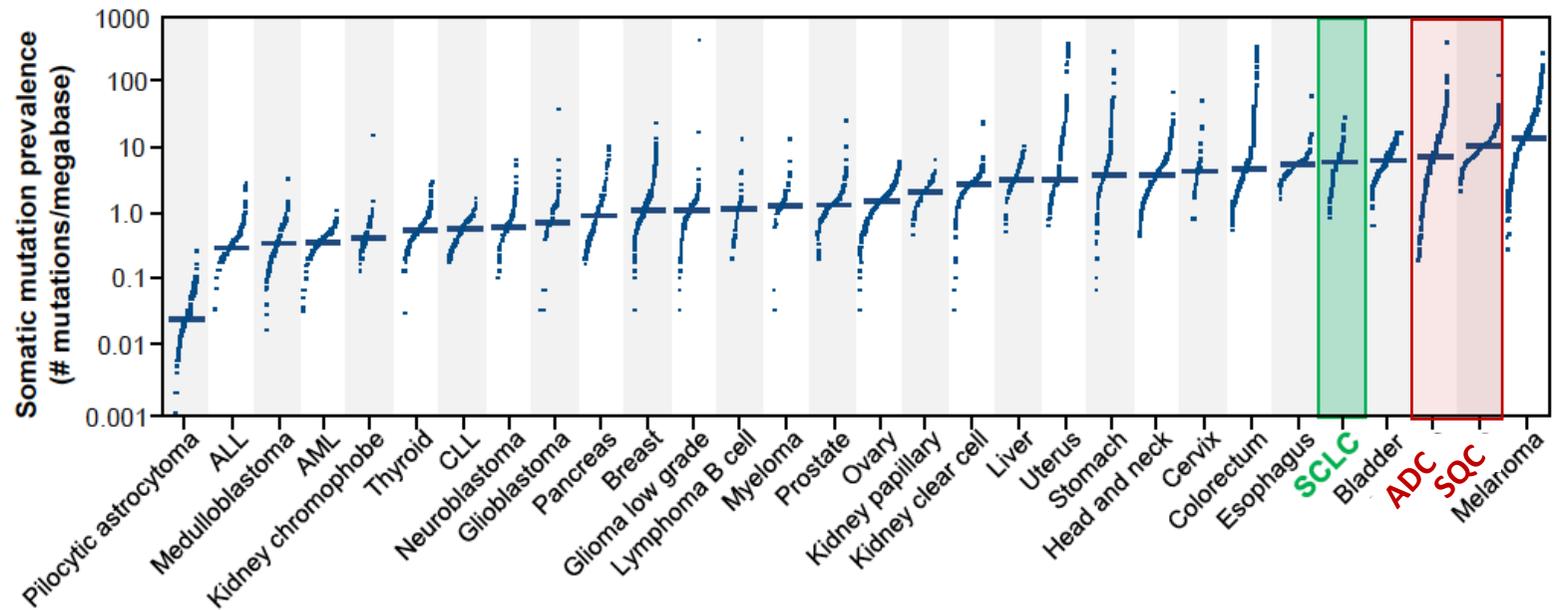
Expressão PD-L1

Teff (effector T-cell) gene signature

Tumor Mutational Burden

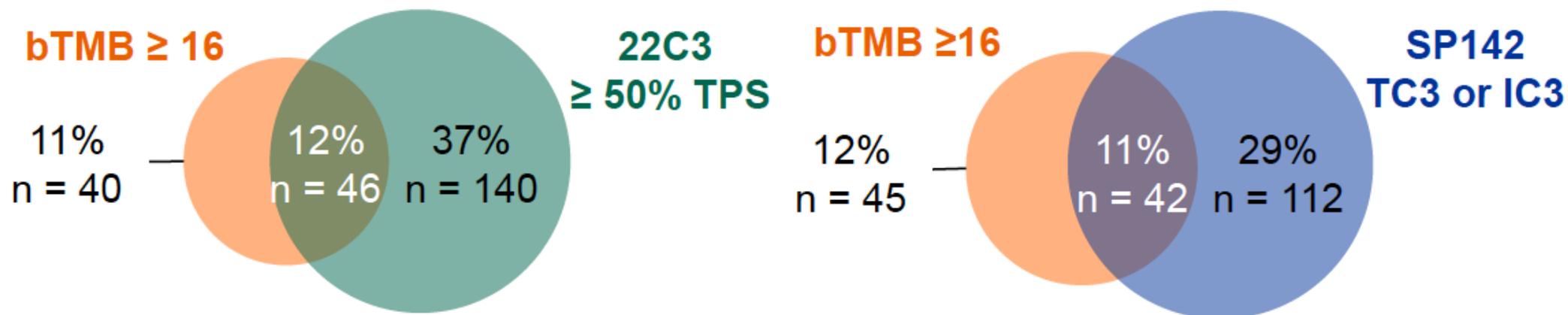
Perspectivas futuras

Tumor Mutational Burden



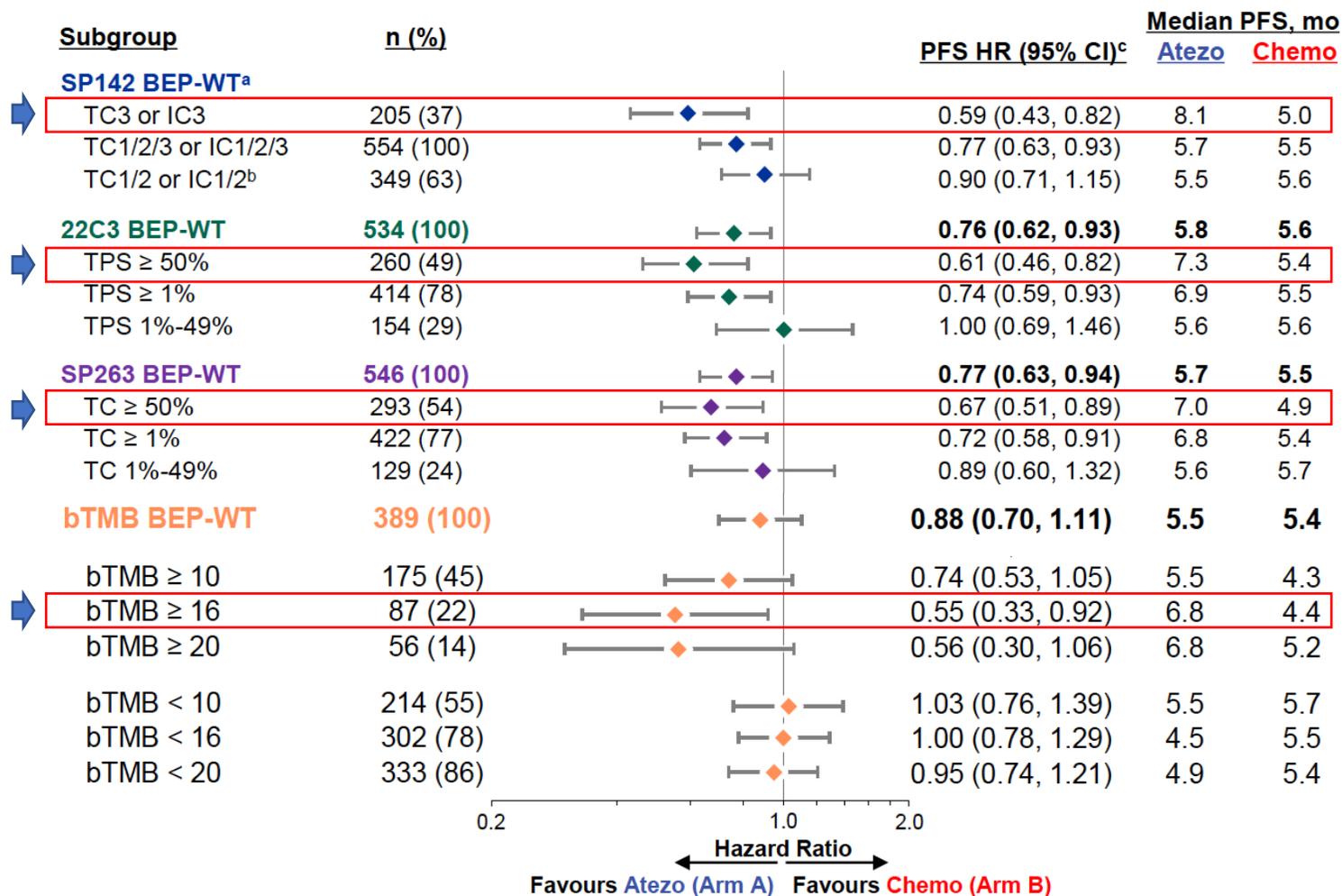
Alexandrov LB, et al. Nature 2013;500:415-421

PD-L1 (SP142/22C3) e TMB identificaram populações distintas no IMpower110



- No IMpower110, TMB foi avaliada usando *blood-based assay* da Foundation Medicine)
- bTMBscore de 16 é equivalente a 16 mutações/1.1 Mb, ou ≈ 14.5 mut/Mb

Benefício idêntico na OS e PFS com Atezo (1L) em PD-L1 alto ou TMB ≥16

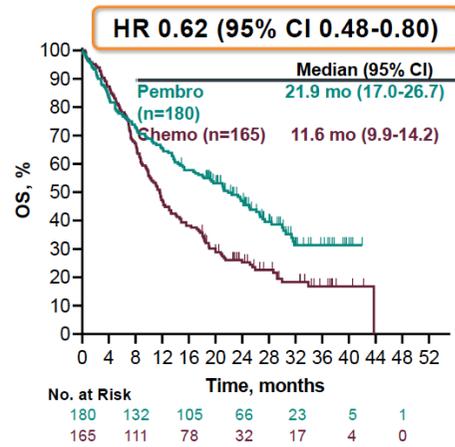
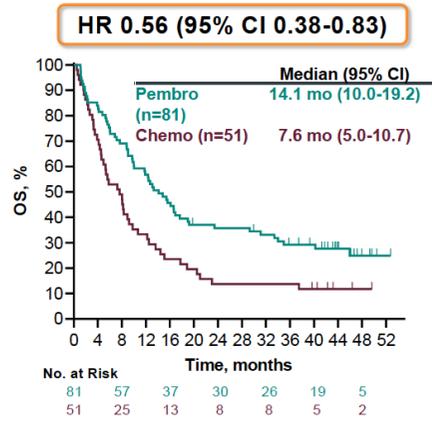


Benefício de Pembro (mono 1L) pode estar restrito a TMB ≥ 175 mut/exoma

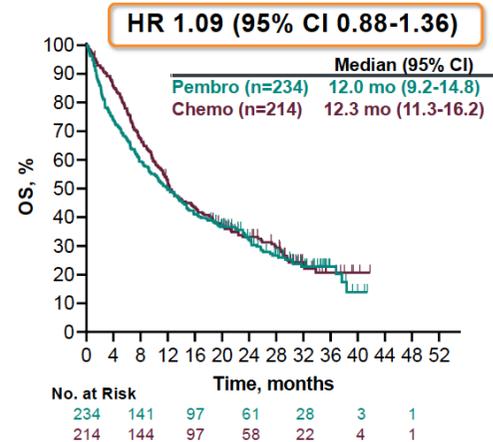
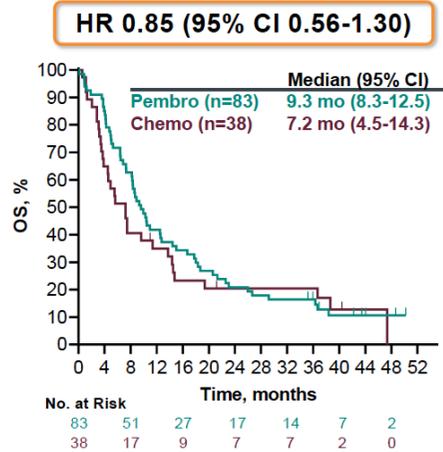
KEYNOTE-010 (PD-L1 $\geq 1\%$)

KEYNOTE-042 (PD-L1 $\geq 1\%$)

tTMB ≥ 175 mut/exome

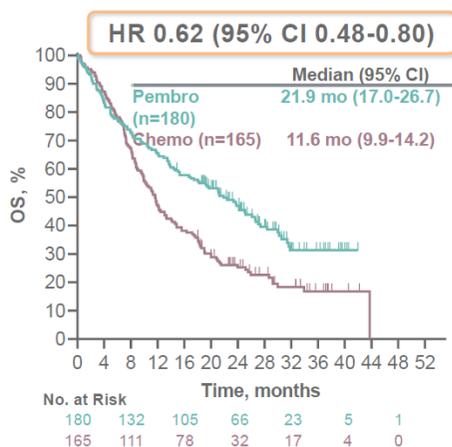
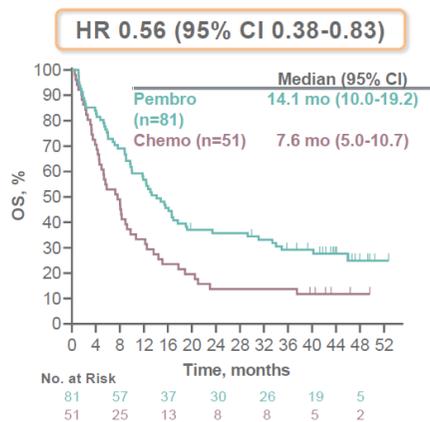


tTMB < 175 mut/exome

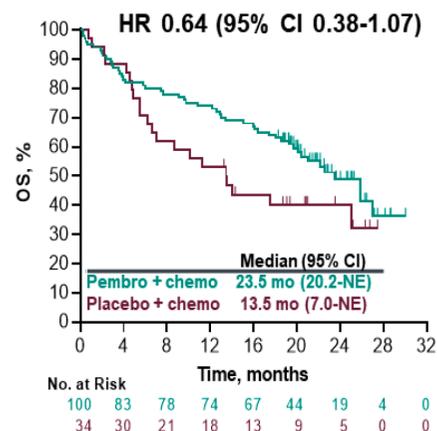


TMB pode não prever o resultado com QT+IT

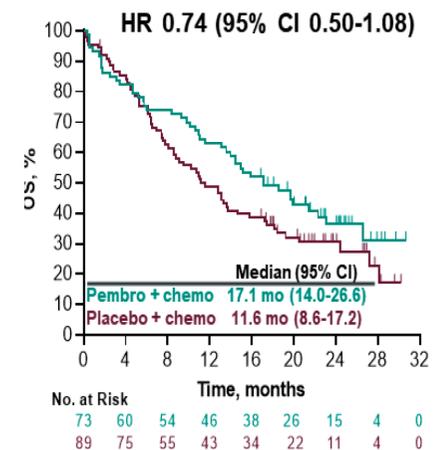
KEYNOTE-010 (PD-L1 $\geq 1\%$) KEYNOTE-042 (PD-L1 $\geq 1\%$)



KEYNOTE-189

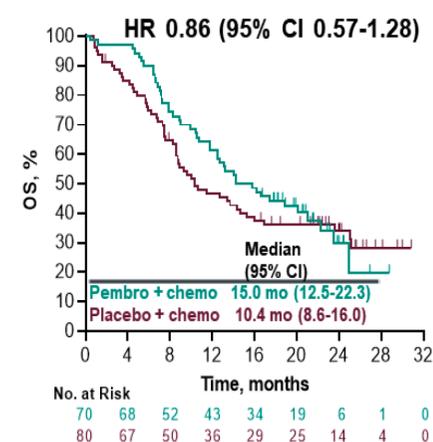
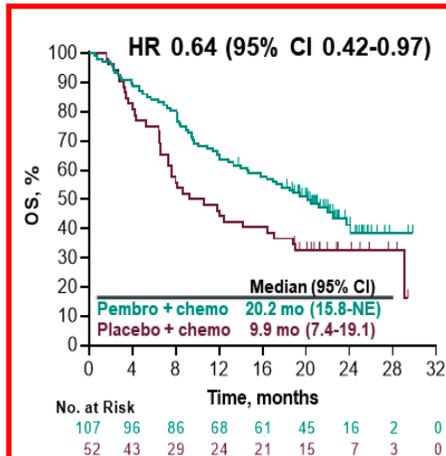
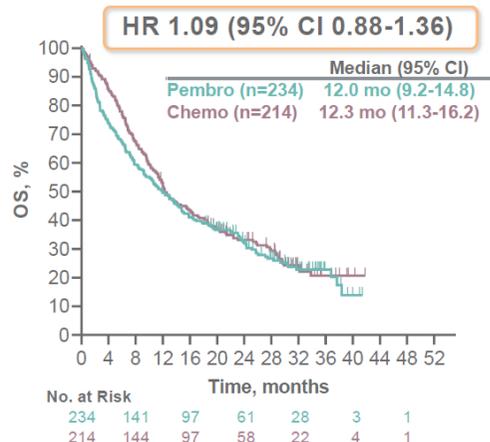
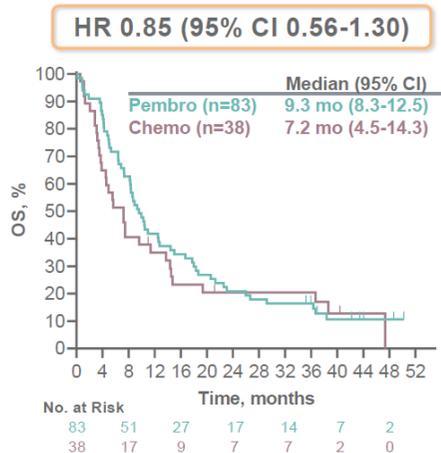


KEYNOTE-407

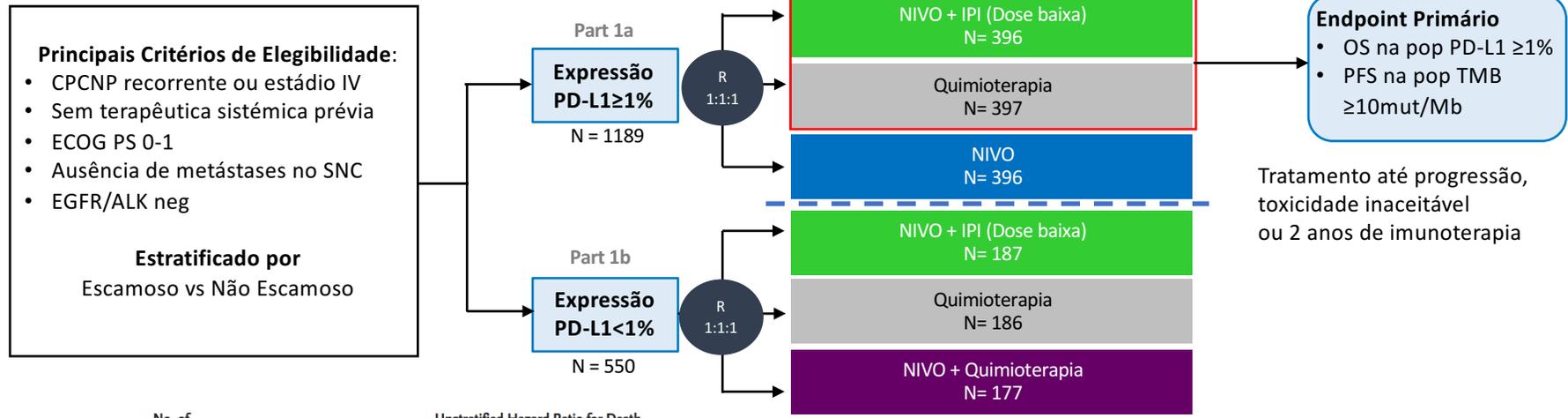


tTMB ≥ 175 mut/exome

tTMB < 175 mut/exome

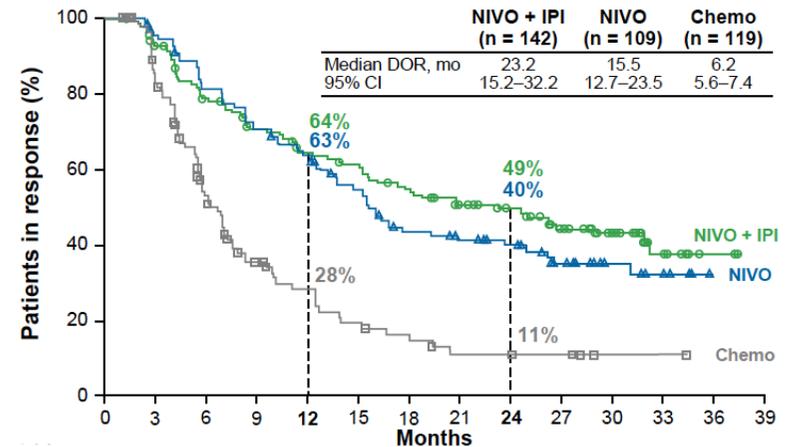


CheckMate 227: Nivolumab + Ipilimumab vs Quimioterapia (1L)



Subgroup	No. of Patients	Median Overall Survival Nivolumab + ipilimumab (N=583) months	Median Overall Survival Chemotherapy (N=583) months	Unstratified Hazard Ratio for Death (95% CI)
Randomized Groups				
PD-L1 expression level				
All patients	1166	17.1	13.9	0.73 (0.64–0.84)
<1%	373	17.2	12.2	0.62 (0.49–0.79)
≥1%	793	17.1	14.9	0.79 (0.65–0.96)
Additional Exploratory Subgroup Analyses				
PD-L1 expression level				
1–49%	396	15.1	15.1	0.94 (0.75–1.18)
≥50%	397	21.2	14.0	0.70 (0.55–0.90)
Tumor mutational burden				
Low, <10 mut/Mb	380	16.2	12.6	0.75 (0.59–0.94)
High, ≥10 mut/Mb	299	23.0	16.4	0.68 (0.51–0.91)
Tumor histologic type				
Squamous	236	14.8	9.2	0.69 (0.52–0.92)
Nonsquamous	557	19.4	17.2	0.85 (0.69–1.04)

DOR, meses (PD-L1 ≥ 1%)



Peters, S. et al Nivolumab + Low-Dose Ipilimumab Versus Platinum-Doublet Chemotherapy as First-Line Treatment for Advanced Non-Small Cell Lung Cancer: CheckMate 227 Part 1 Final Analysis, presented at ESMO 2019
 Hellmann et al. Nivolumab plus Ipilimumab in Advanced Non-Small-Cell Lung Cancer. N Engl J Med. 2019 Sep 28. doi: 10.1056/NEJMoa1910231

Expressão PD-L1

Teff (effector T-cell) gene signature

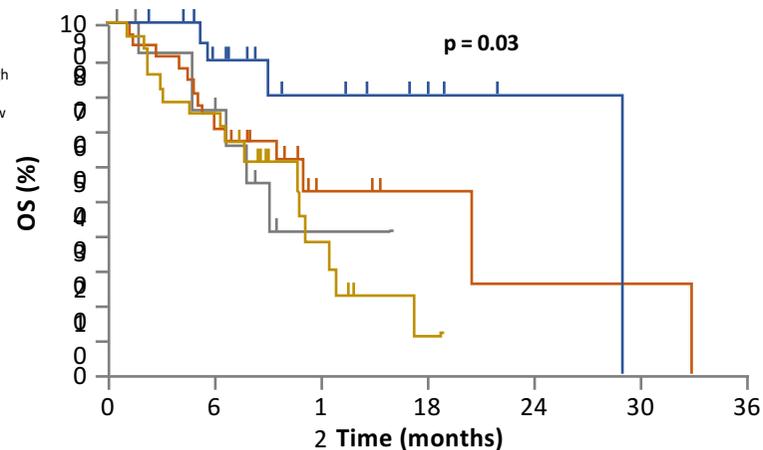
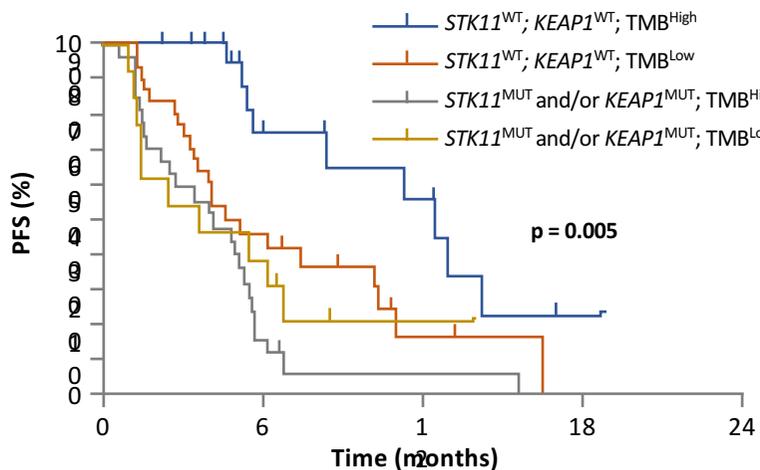
Tumor Mutational Burden

Perspectivas futuras

Integração de alterações genómicas *STK11* e *KEAP1* com TMB e outros biomarcadores: towards a composite panel?

Group	PFS, mo
<i>STK11</i> ^{WT} ; <i>KEAP1</i> ^{WT} ; TMB ^{High}	12.4
<i>STK11</i> ^{WT} ; <i>KEAP1</i> ^{WT} ; TMB ^{Low}	4.5
<i>STK11</i> ^{MUT} and/or <i>KEAP1</i> ^{MUT} ; TMB ^{High}	4.1
<i>STK11</i> ^{MUT} and/or <i>KEAP1</i> ^{MUT} ; TMB ^{Low}	3.6

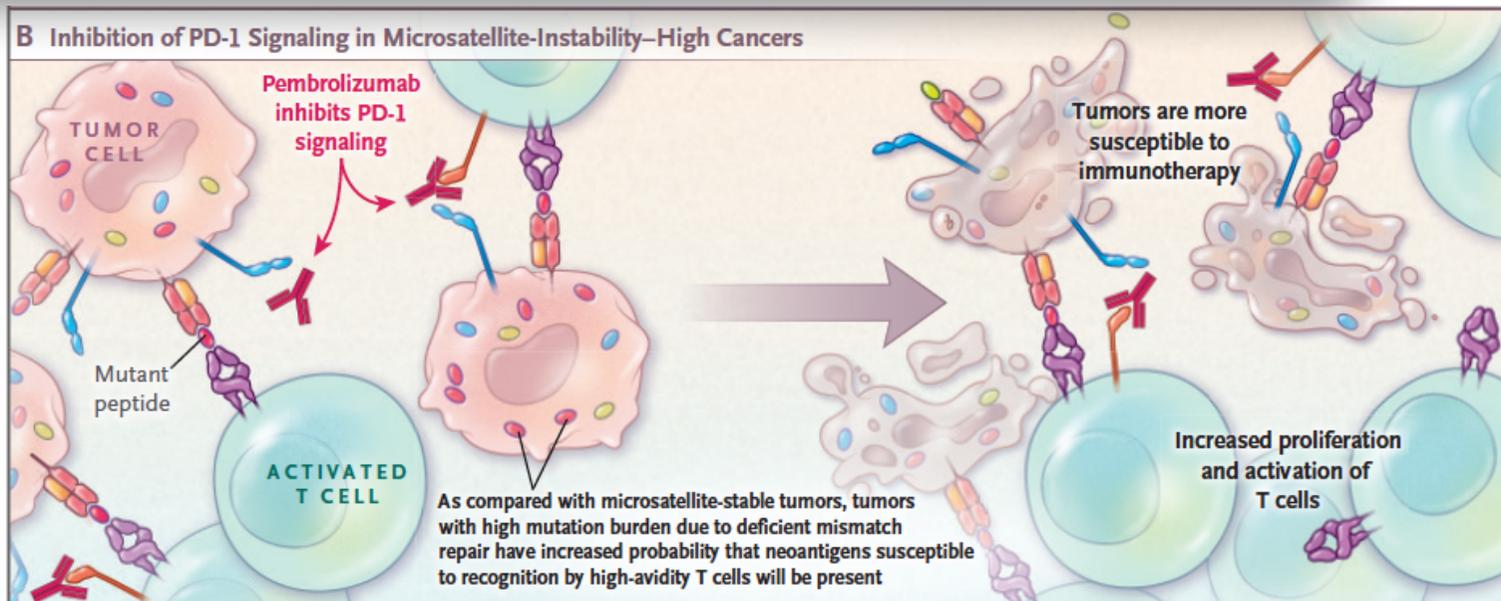
Group	OS, mo
<i>STK11</i> ^{WT} ; <i>KEAP1</i> ^{WT} ; TMB ^{High}	28.9
<i>STK11</i> ^{WT} ; <i>KEAP1</i> ^{WT} ; TMB ^{Low}	20.4
<i>STK11</i> ^{MUT} and/or <i>KEAP1</i> ^{MUT} ; TMB ^{High}	10.7
<i>STK11</i> ^{MUT} and/or <i>KEAP1</i> ^{MUT} ; TMB ^{Low}	9.1



dMMR/MSI-H predicts the efficacy of anti-PD-1/ PD-L1 immunotherapy

First FDA Approval Agnostic of Cancer Site — When a Biomarker Defines the Indication

Steven Lemery, M.D., M.H.S., Patricia Keegan, M.D., and Richard Pazdur, M.D.



N Engl J Med. 2017

Signaling Mechanism of PD-1 and PD-L1 and Inhibition of PD-1 Signaling in Microsatellite-Instability–High Cancers.

MHC1 denotes major histocompatibility complex 1.

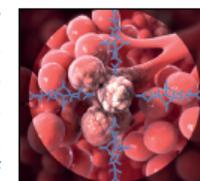
Lancet Respir Med 2018

Lung cancer immunotherapy biomarkers: refine not reject



Lung cancer is the leading cause of cancer-related mortality in the USA and UK, and for patients with metastatic non-small-cell lung cancer (NSCLC) 5-year survival is around 1%. The new era of immunotherapy drugs has given hope to millions, with impressive initial results for improving both progression-free and overall survival in these patients. The approval of immune checkpoint inhibitors has previously been tied to concentrations of the biomarker PD-L1, thought to predict treatment response. On April 16, 2018, two large

obtain a result, which can be 2 weeks or more. Another issue is the different assays and cutoffs available for PD-L1 testing that could lead to misclassification of PD-L1 status for some patients. These trials suggest that PD-L1 does still offer guidance as to who will respond better, and that for some indications PD-L1 status still plays a part. But if approval for use of pembrolizumab and other checkpoint inhibitors no longer always requires the determination of PD-L1 status, will physicians move away from testing? With an increase in the patient group eligible for



Artwork by iStockphoto Ltd/SE

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