



Society for Immunotherapy of Cancer

Advances in Cancer Immunotherapy™

New Directions and Immunotherapy Resistance: How to Treat Patients Progressing on Therapy

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#LearnACI

Disclosures

- Consultant for: Genentech, Bristol-Myers Squibb, Merck, AstraZeneca, Maverick, Blueprint Medicine, Syndax, Ariad, Nektar, ArcherDX, Mirati, NextCure, Novartis, EMD Serono
- Research funding (to institute): AstraZeneca, Lilly, Genentech, Bristol-Myers Squibb
- I will be discussing non-FDA approved indications during my presentation.

NSCLC: Advances in precision genomic medicine

Actionable genomic mutation by NGS?



Treat with targeted therapy



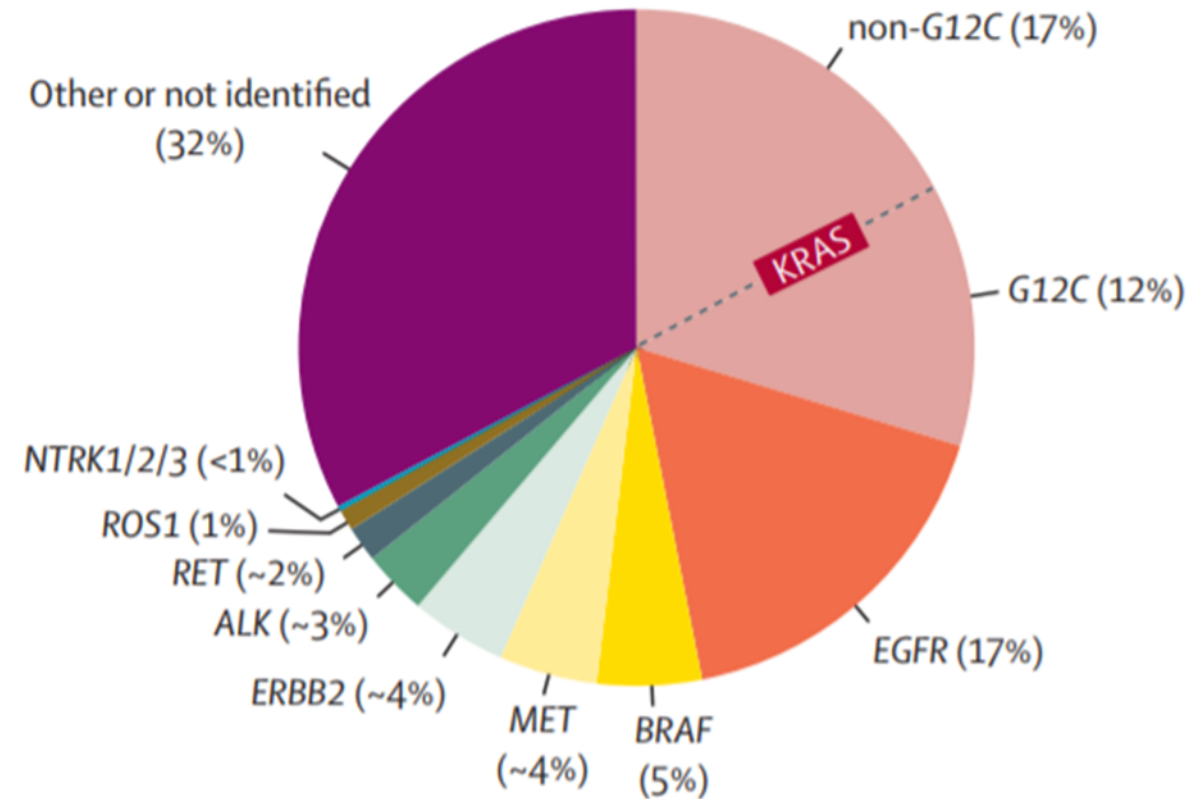
Upon development of acquired resistance

Repeat biopsy & ctDNA analysis



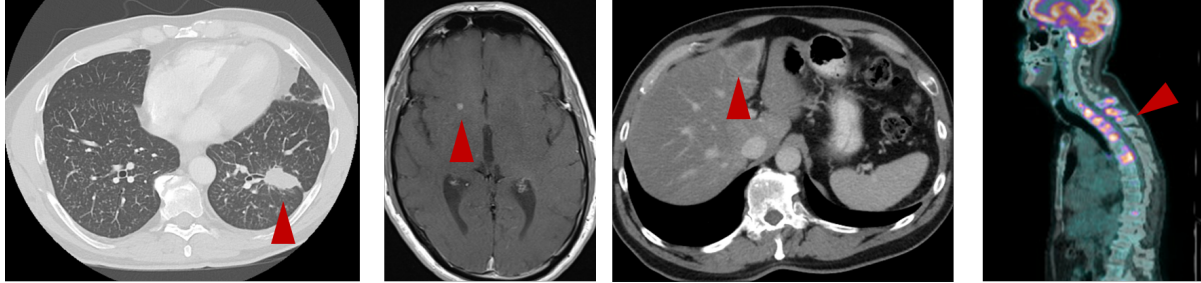
Treat resistance mutations if possible

C Oncogenic mutations in NSCLC

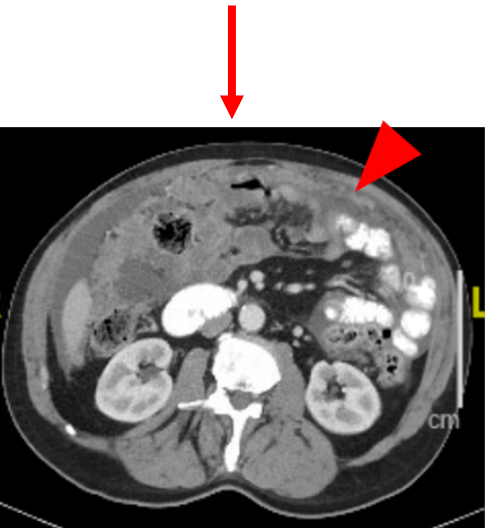


NSCLC: Advances in precision genomic medicine

62yoM with stage IV NSCLC & EGFR exon 19 deletion

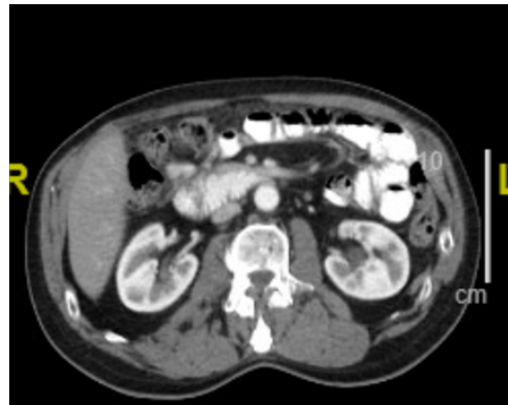


Responded to osimertinib (EGFR TKI), developed resistance at 20 months, new peritoneal carcinomatosis and ascites



**Acquired
RET Fusion**

Added
selpercatinib
(RET TKI)
to osimertinib



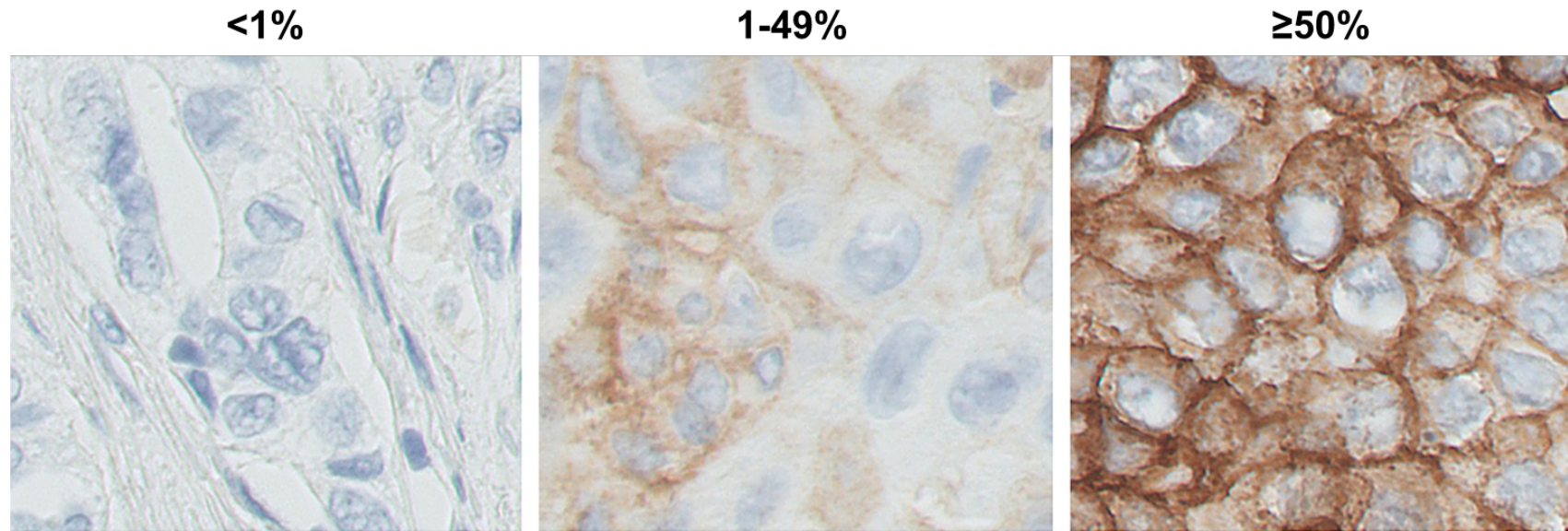
Acquired
Resistance



**EGFR C797S +
RET G810S**

Chemotherapy

Biomarkers for Immunotherapy Selection



PD-L1 Tumor Proportion Score (TPS)

Developing Precision Medicine for Immunotherapy

If no targetable mutation, check PD-L1

PD-L1 TPS $\geq 50\%$

PD-L1 TPS $< 50\%$

**Pembrolizumab
Monotherapy**

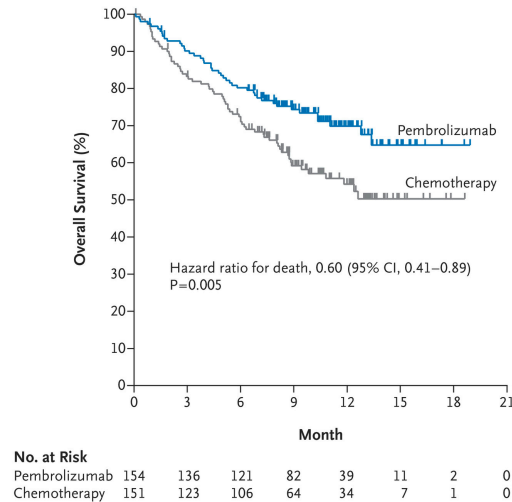
**Pembrolizumab
+ Chemotherapy**

*Upon development of
resistance*

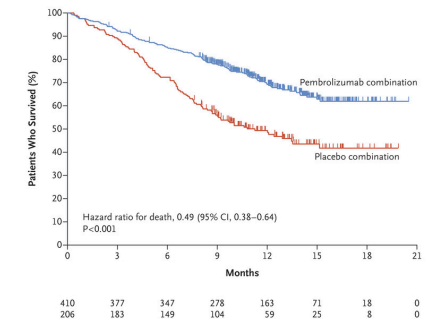
?

**Limited understanding
of resistance**

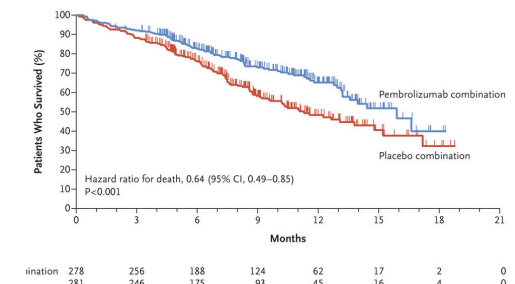
KEYNOTE 024
Pembro vs Chemo
in NSCLC
with PD-L1 TPS $\geq 50\%$



KEYNOTE 189
Chemo +/- Pembro
in Nonsquamous NSCLC
with PD-L1 TPS 0-100%



KEYNOTE 407
Chemo +/- Pembro
in Squamous NSCLC with
PD-L1 TPS 0-100%



What are the known mechanisms of primary and acquired resistance to immunotherapy in lung cancer?



Biomarkers of Immunotherapy Efficacy

- Pathologic Factors
 - PD-L1 expression
- Genomic Factors
 - Tumor mutational burden, specific genomic subtypes
- Circulating Factors
 - Neutrophil-lymphocyte ratio (NLR)
- Immunologic Factors
 - Gene expression signatures
 - Immune cell subsets

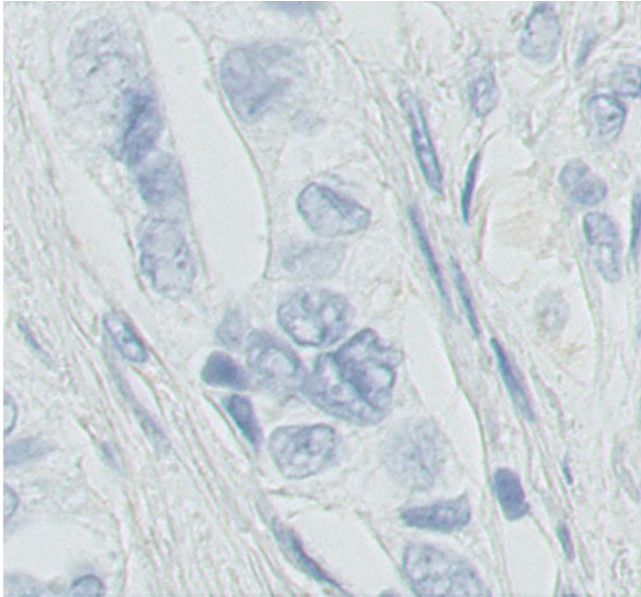
PD-L1 Expression

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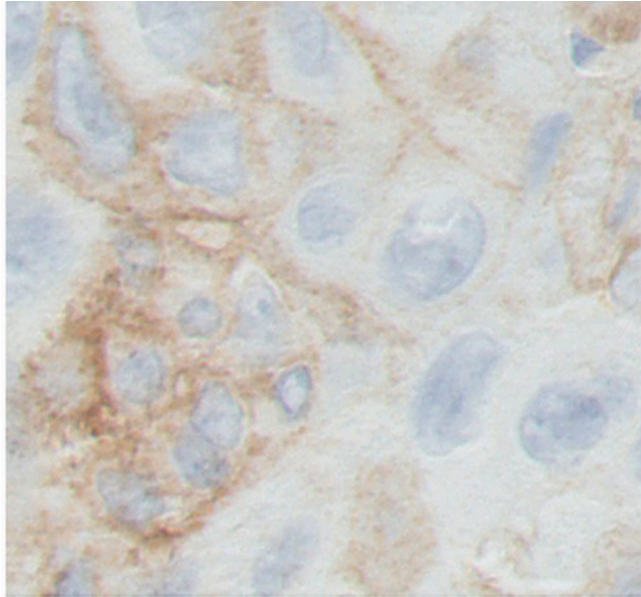
PD-L1 as a predictive biomarker

PD-L1 Tumor Proportion Score (TPS)

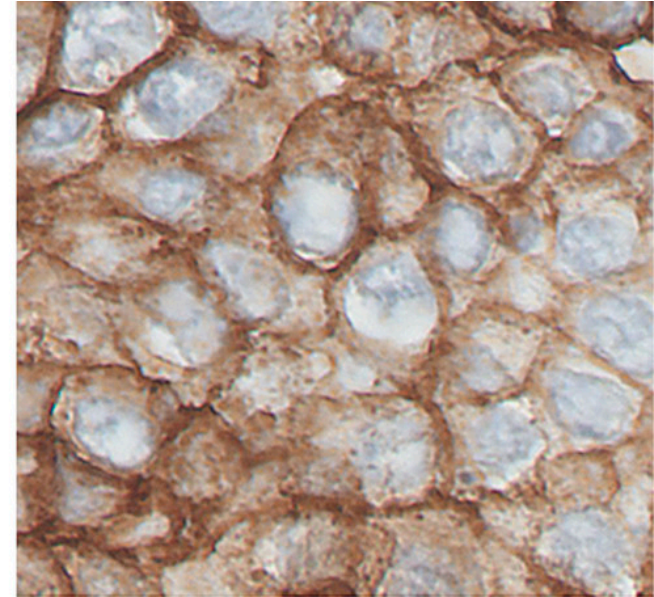
<1%



1-49%

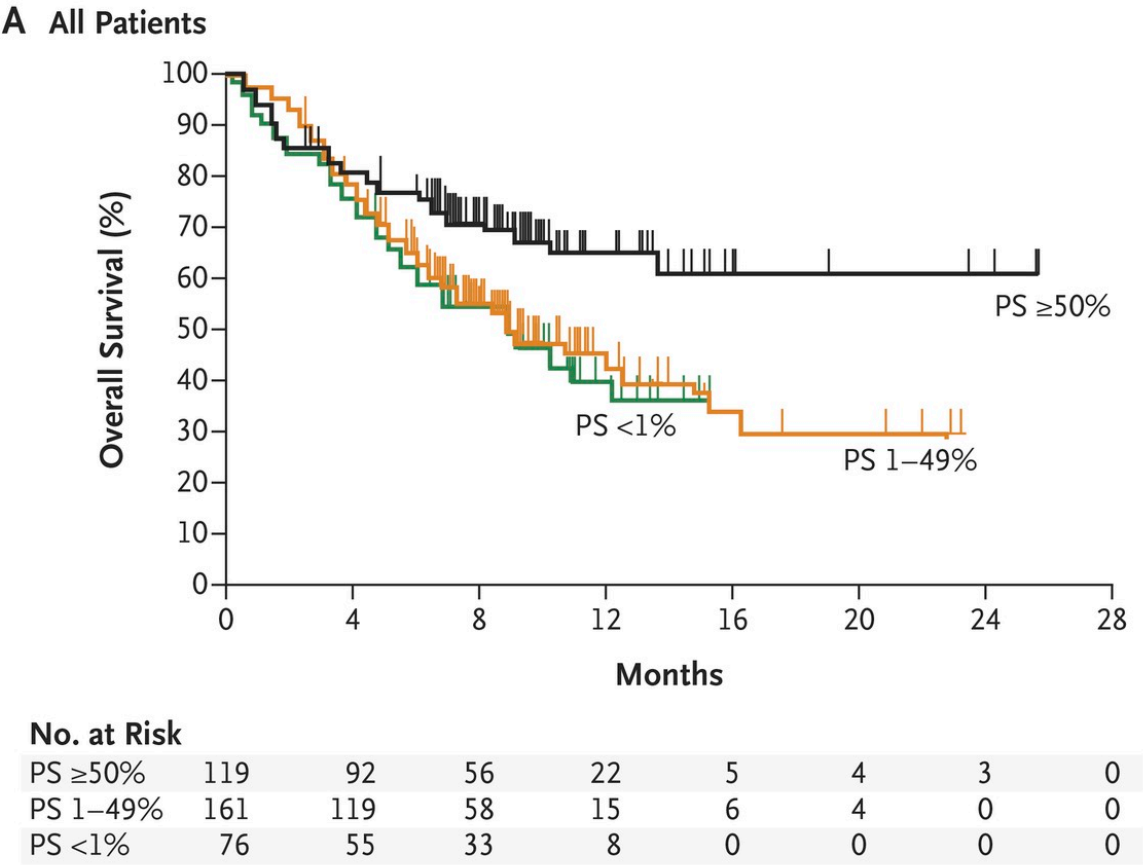


≥50%



PD-L1 as a predictive biomarker

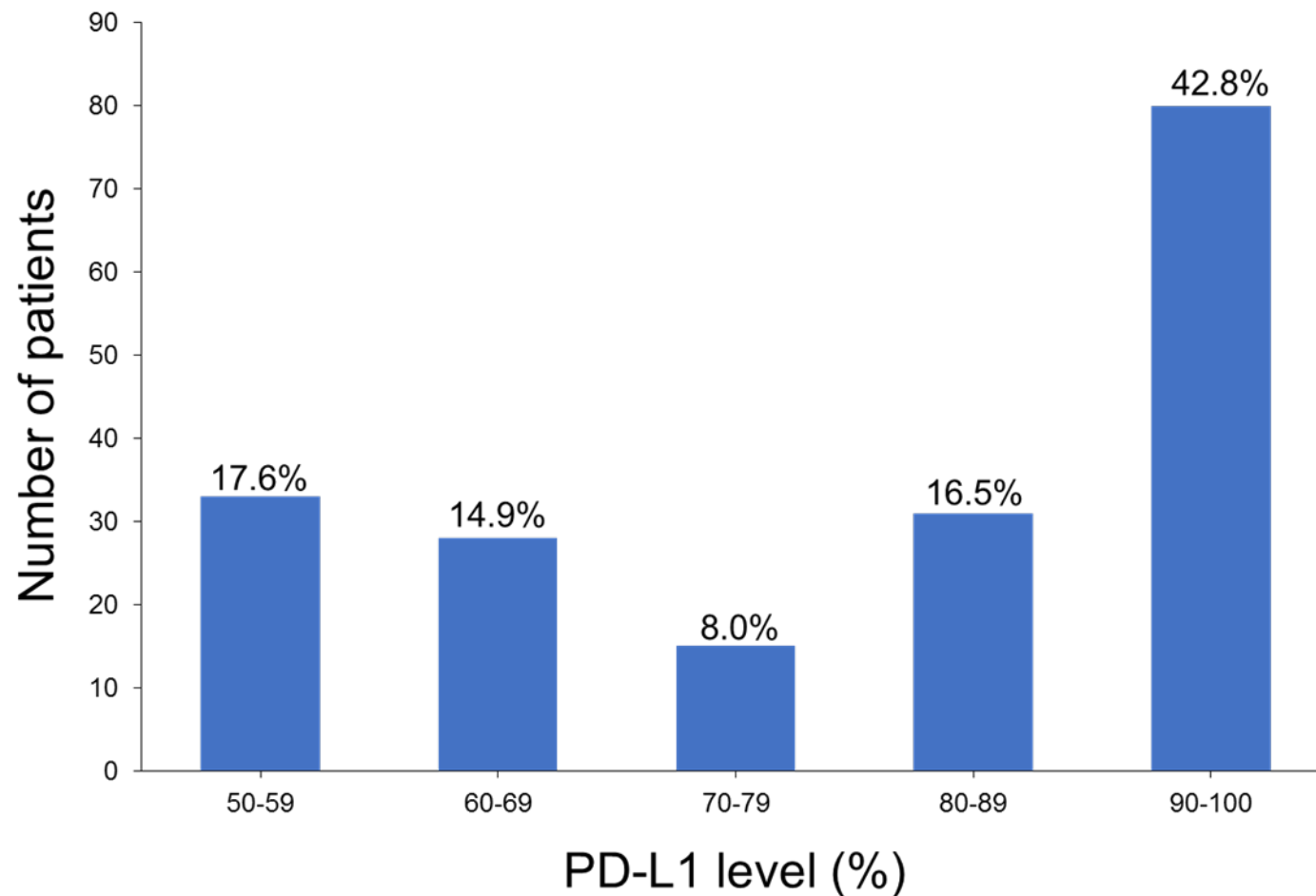
KEYNOTE 001



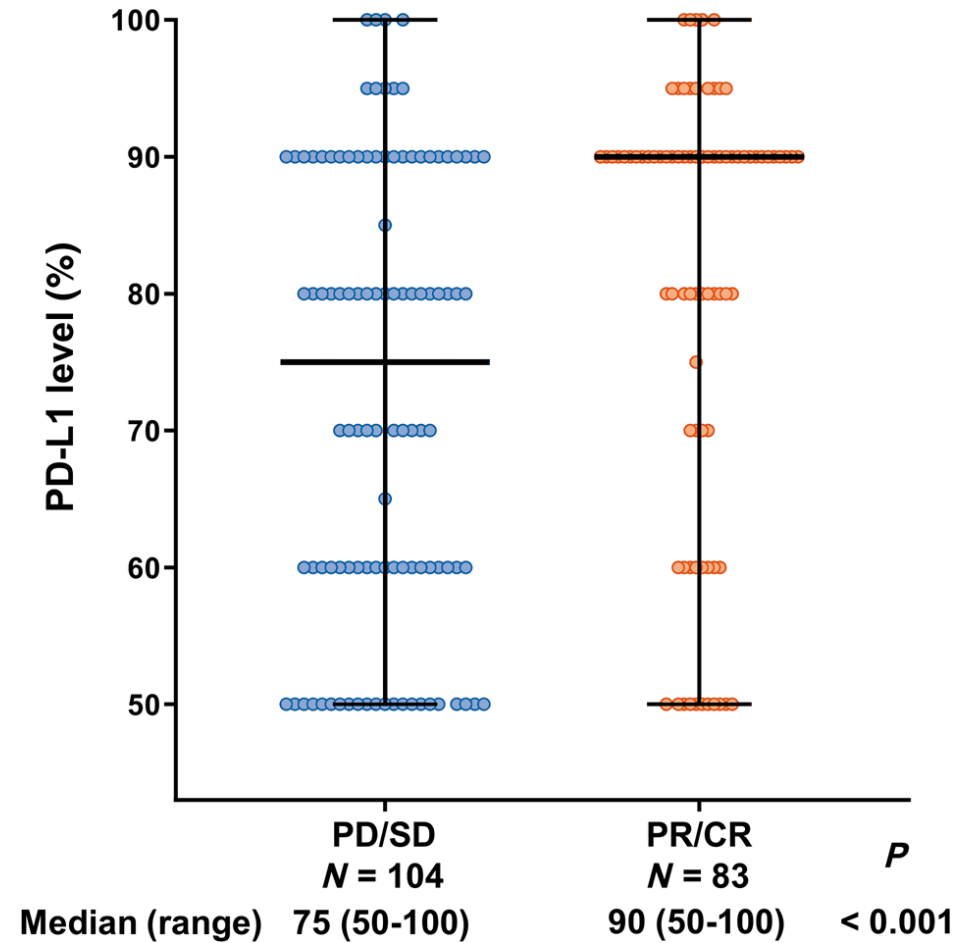
PD-L1 TPS $\geq 50\%$: Is higher even better?

Retrospective analysis:

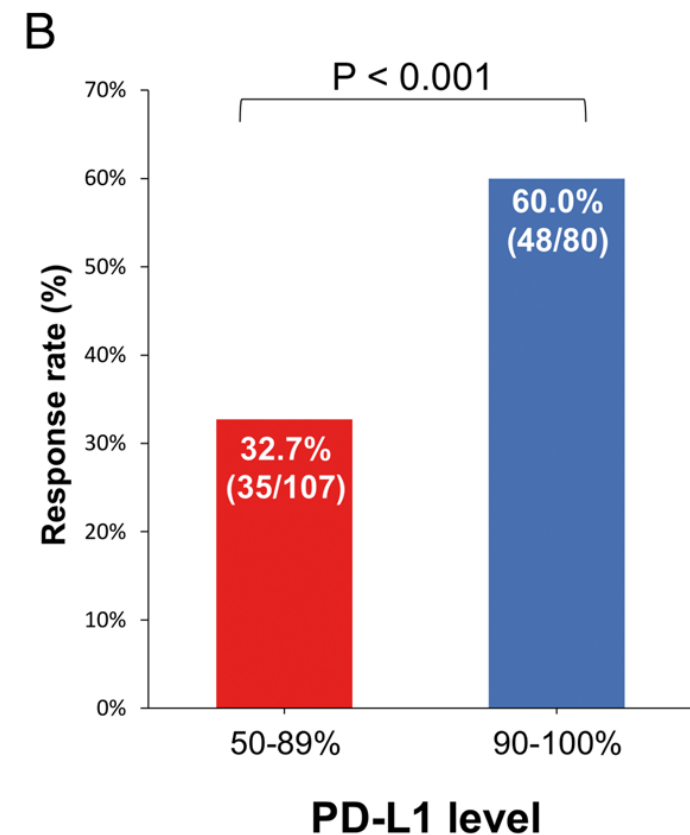
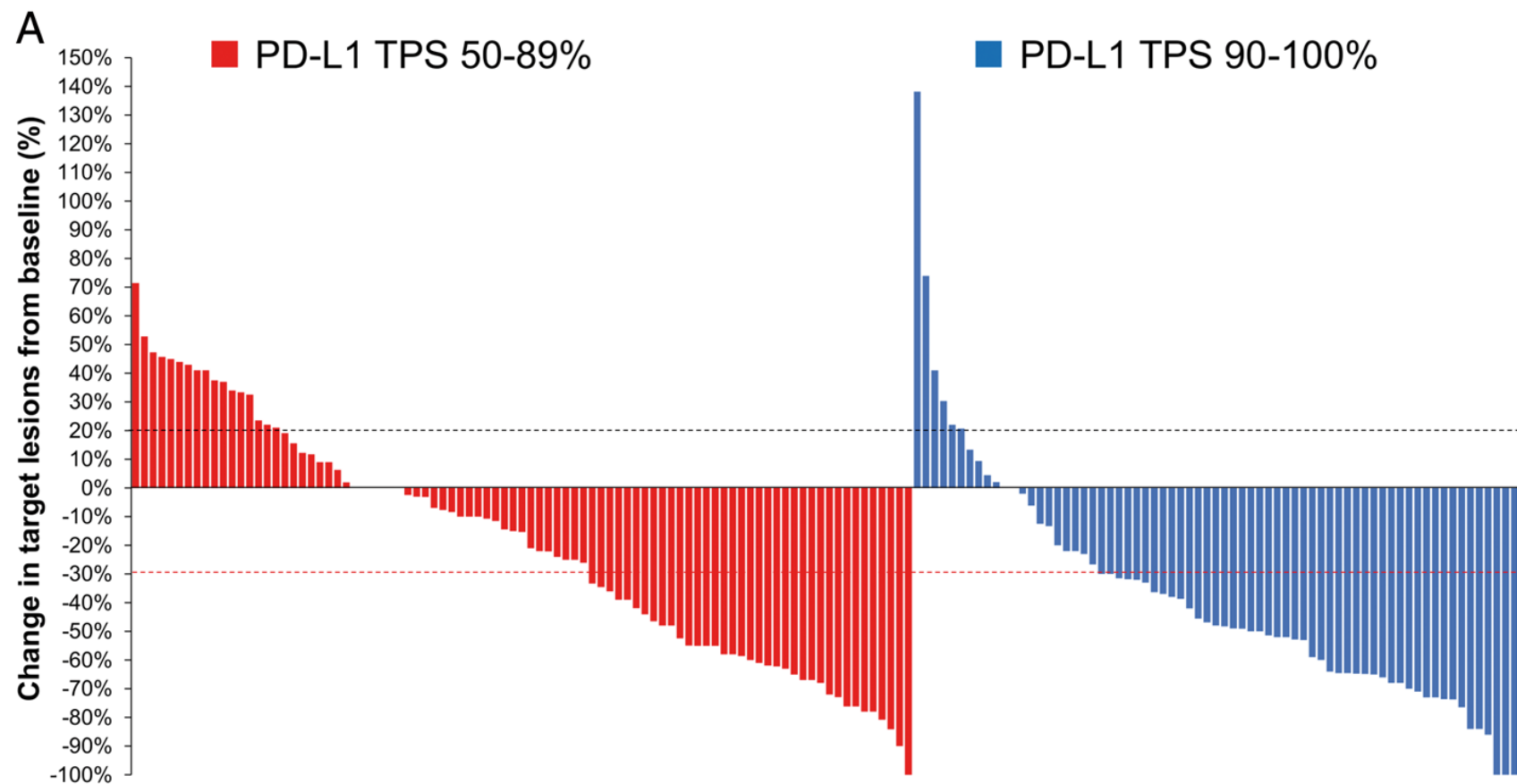
- 4 academic institutions
- 187 patients with NSCLC
- 1st-line commercial pembrolizumab
- PD-L1 TPS $\geq 50\%$
- EGFR/ALK excluded



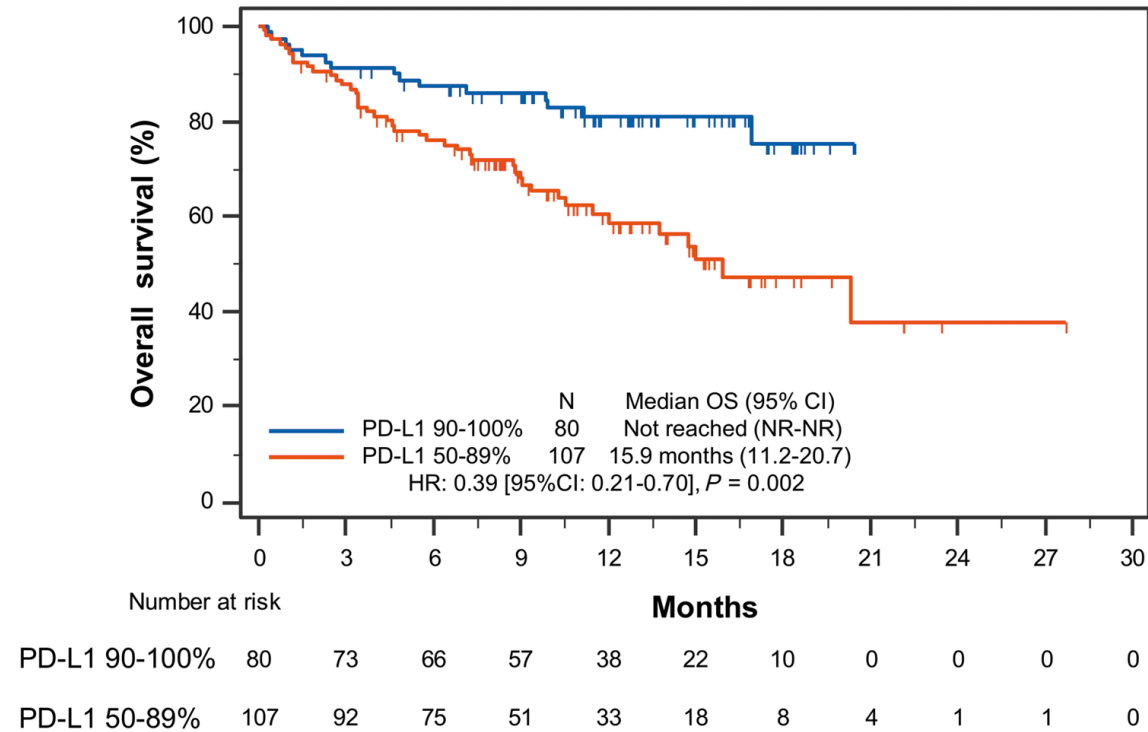
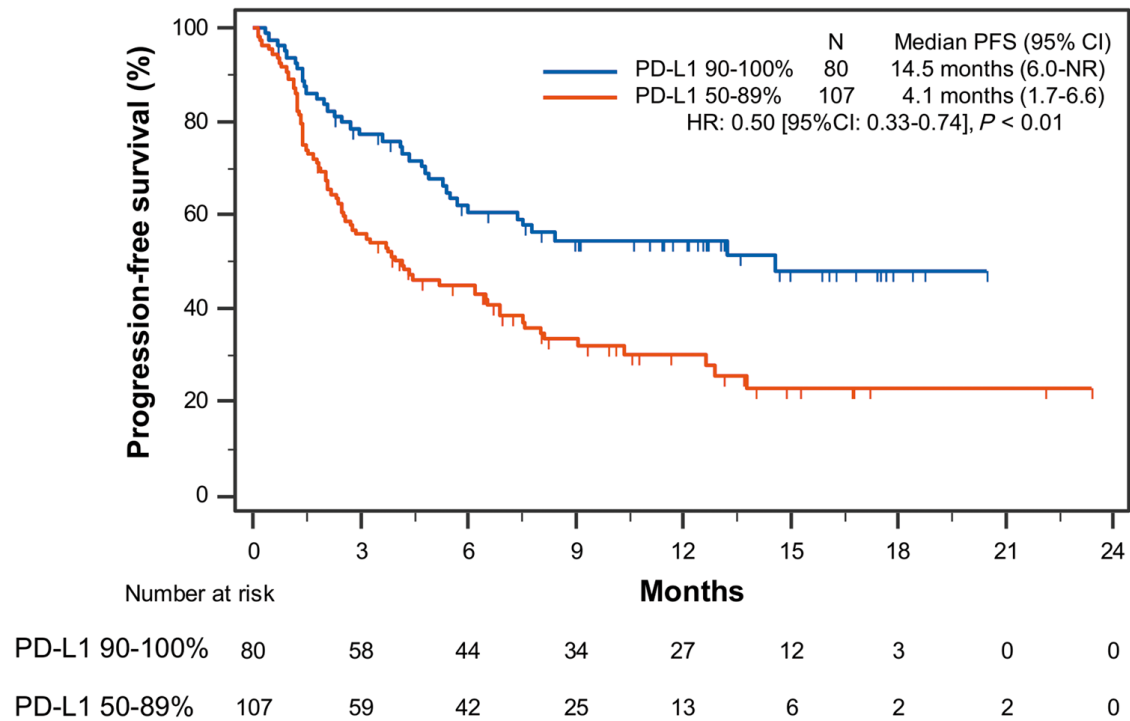
PD-L1 TPS in responders vs nonresponders



PD-L1 TPS 50-89% vs $\geq 90\%$



PD-L1 TPS 50-89% vs ≥90%



Cemiplimab in Very High PD-L1 Expressing NSCLC

EMPOWER-Lung 1

	PD-L1 ≥90%	PD-L1 >60 to <90%	PD-L1 ≥50 to ≤60%	PD-L1 <50% or unknown
Number of patients	98 vs 94	89 vs 90	96 vs 96	73 vs 74
Overall survival				
Median, months (95% CI)	NR (17.3-NE) vs 15.1 (11.1-NE)	22.1 (17.9-NE) vs 12.0 (9.6-19.2)	21.9 (13.2-NE) vs 14.0 (9.4-19.3)	16.5 (11.6-NE) vs 15.2 (10.2-NE)
Hazard ratio (95% CI)	0.46 (0.25-0.85)	0.47 (0.27-0.80)	0.77 (0.49-1.23)	1.082 (0.68-1.72)
Progression-free survival				
Median, months (95% CI)	15.3 (10.4-18.7) vs 5.9 (4.3-6.2)	6.2 (4.2-8.4) vs 4.2 (4.1-5.7)	4.3 (2.8-6.3) vs 6.2 (5.0-6.2)	4.1 (2.6-6.1) vs 5.0 (4.2-6.2)
Hazard ratio (95% CI)	0.28 (0.17-0.46)	0.55 (0.38-0.80)	0.79 (0.56-1.12)	0.82 (0.56-1.18)
Tumour response				
Objective response rate, % (95% CI)	46 (36-56) vs 18 (11-27)	39 (29-50) vs 20 (12-30)	32 (23-43) vs 23 (15-33)	26 (17-38) vs 22 (13-33)
Data are median (95% CI), hazard ratio (95% CI), and objective response rate % (95% CI). NE=not evaluable. NR=not reached. PD-L1=programmed cell death ligand 1.				
Table 3: Correlation of survival and objective response with baseline PD-L1 proportion scores for cemiplimab versus chemotherapy				

Tumor Mutational Burden

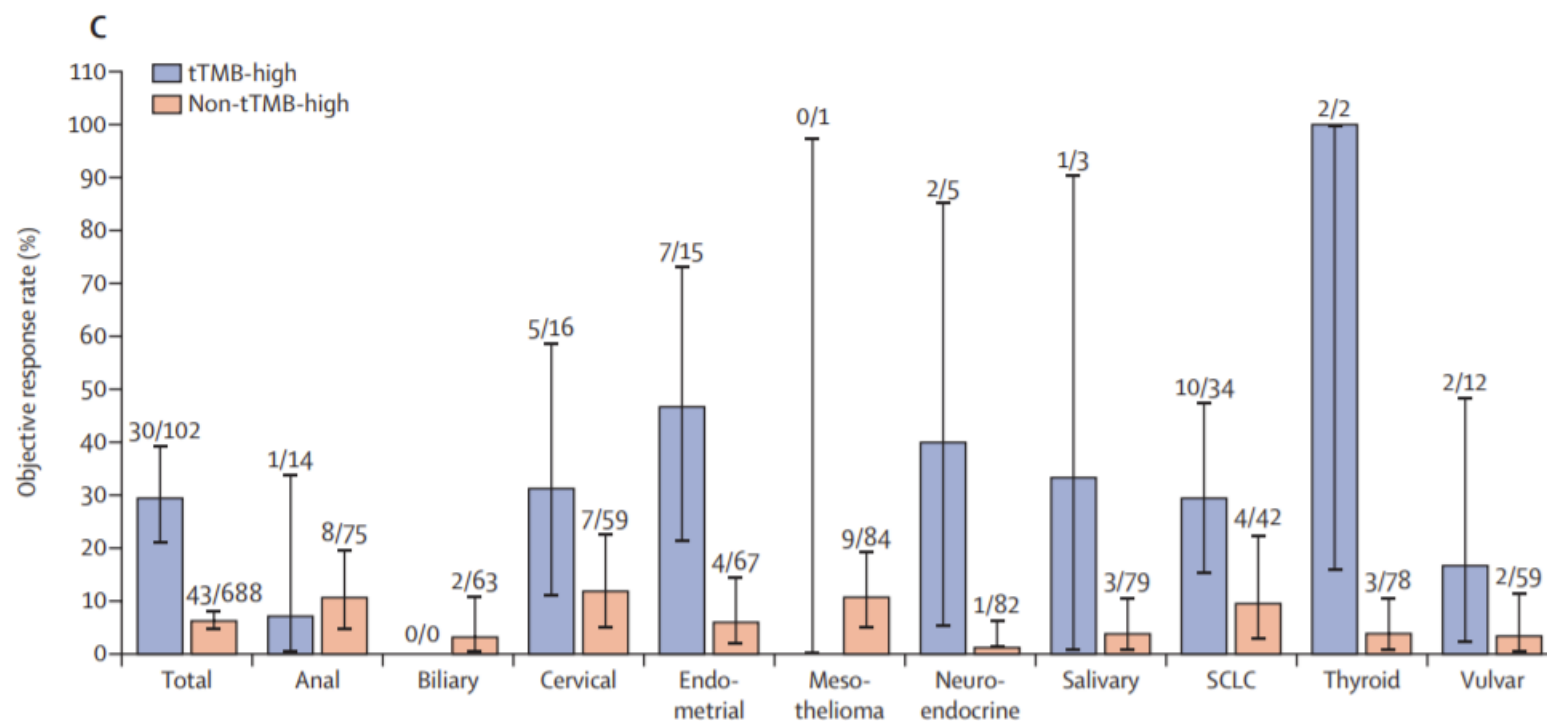
A horizontal decorative bar spanning the width of the slide, positioned below the title. It consists of three segments: a light blue segment on the left, a dark blue segment in the middle, and an orange segment on the right. The segments are separated by thin white lines, and the dark blue segment has a slight diagonal cut on its right side.

KEYNOTE-158: Pembrolizumab in TMB-high Cancers

	tTMB-high (n=102)	tTMB-high (excluding MSI-H; n=81)*	Non-tTMB- high (n=688)
Best response			
Complete response	4 (4%)	3 (4%)	11 (2%)
Partial response	26 (25%)	20 (25%)	32 (5%)
Stable disease	14 (14%)	11 (14%)	227 (33%)
Non-complete response or non-progressive disease†	0	0	3 (<1%)
Progressive disease	48 (47%)	38 (47%)	349 (51%)
Not evaluable‡	1 (1%)	1 (1%)	13 (2%)
Not assessed§	9 (9%)	8 (10%)	53 (8%)
Objective response rate	29% (21–39)	28% (19–40)	6% (5–8)

Data are n (%) or % (95% CI). MSI-H=high microsatellite instability. RECIST=Response Evaluation Criteria in Solid Tumors. tTMB-high=high tissue tumour mutational burden. *Excludes 14 patients who were MSI-high and seven additional patients who had missing MSI status. †Patients without measurable disease per central review at baseline who did not have a complete response or progressive disease. ‡Patients who did not have a post-baseline imaging assessment evaluable for response. §Patients who did not have post-baseline imaging.

Table 2: Objective response (per RECIST version 1.1), assessed by independent central review in the efficacy population

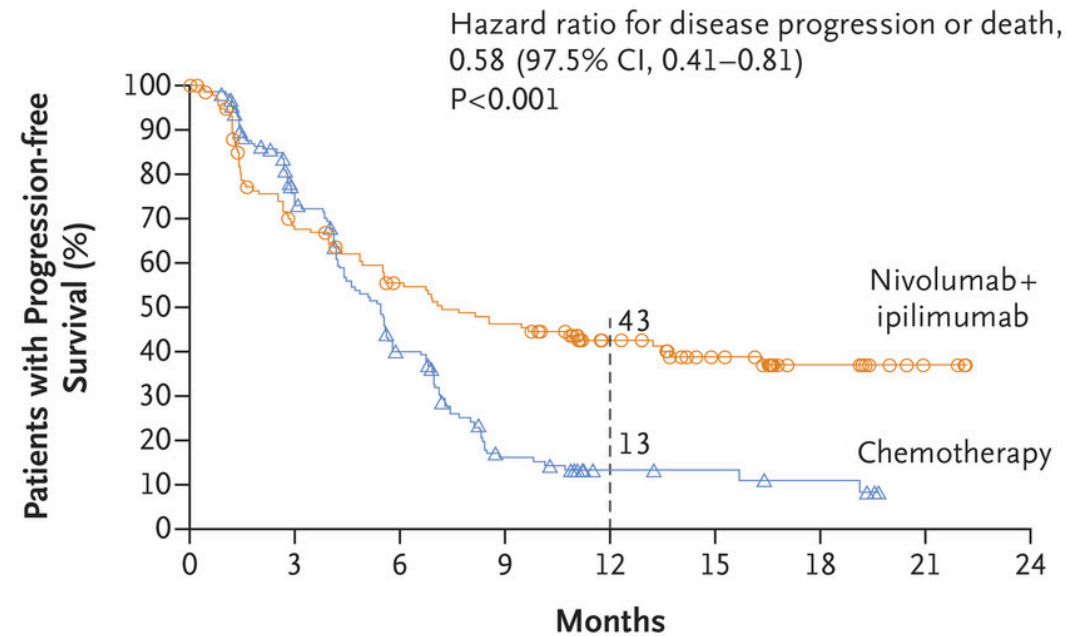


Tumor mutational burden (TMB)

CheckMate 227

High TMB: ≥ 10 mutations/megabase

A Progression-free Survival



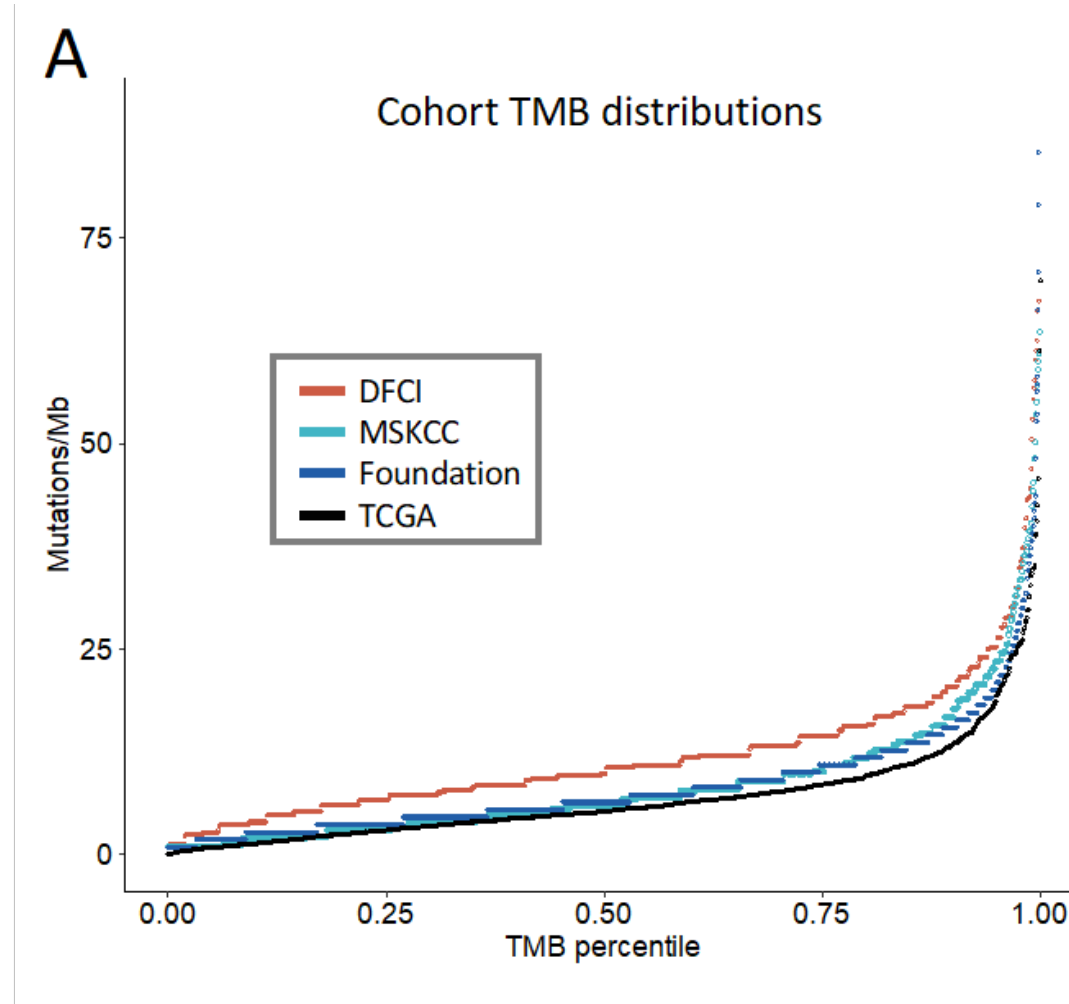
No. at Risk

Nivolumab + ipilimumab	139	85	66	55	36	24	11	3	0
Chemotherapy	160	103	51	17	7	6	4	0	0

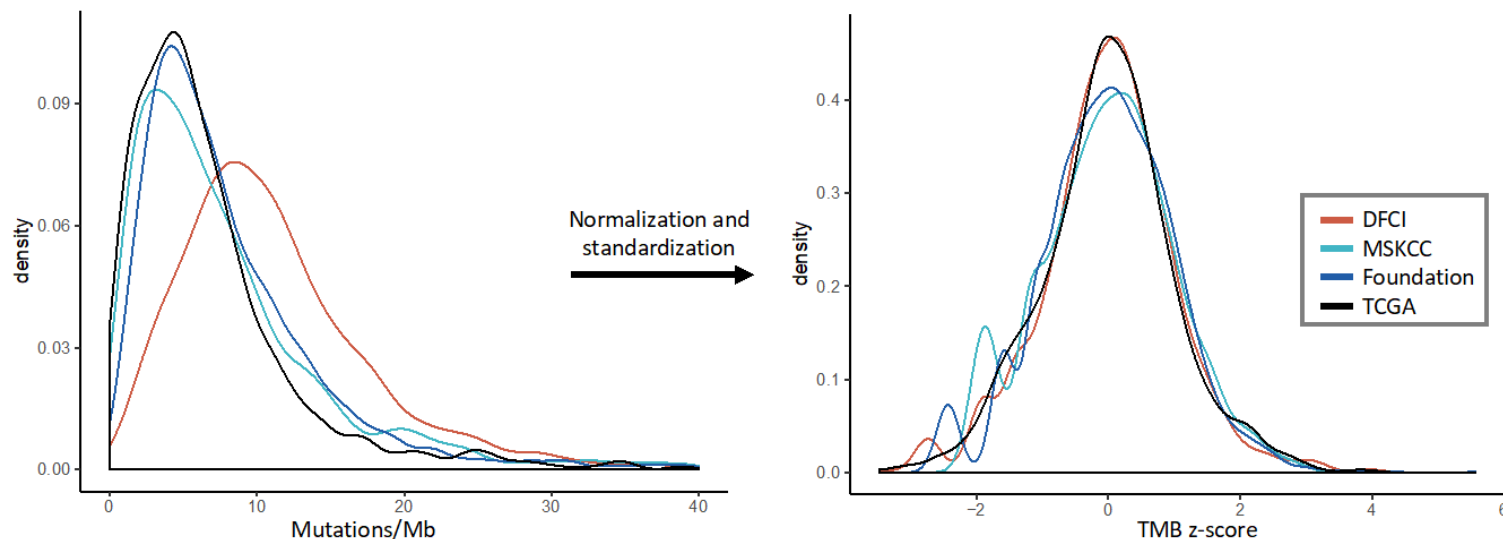
Tumor mutational burden (TMB)

- Challenges with TMB
 - Still not used routinely for treatment selection in NSCLC
 - Limited access to published trial data
 - Cohort sizes at any individual institution are relatively small
 - Differences across platforms make it difficult to compare and combine data

Tumor mutational burden (TMB)

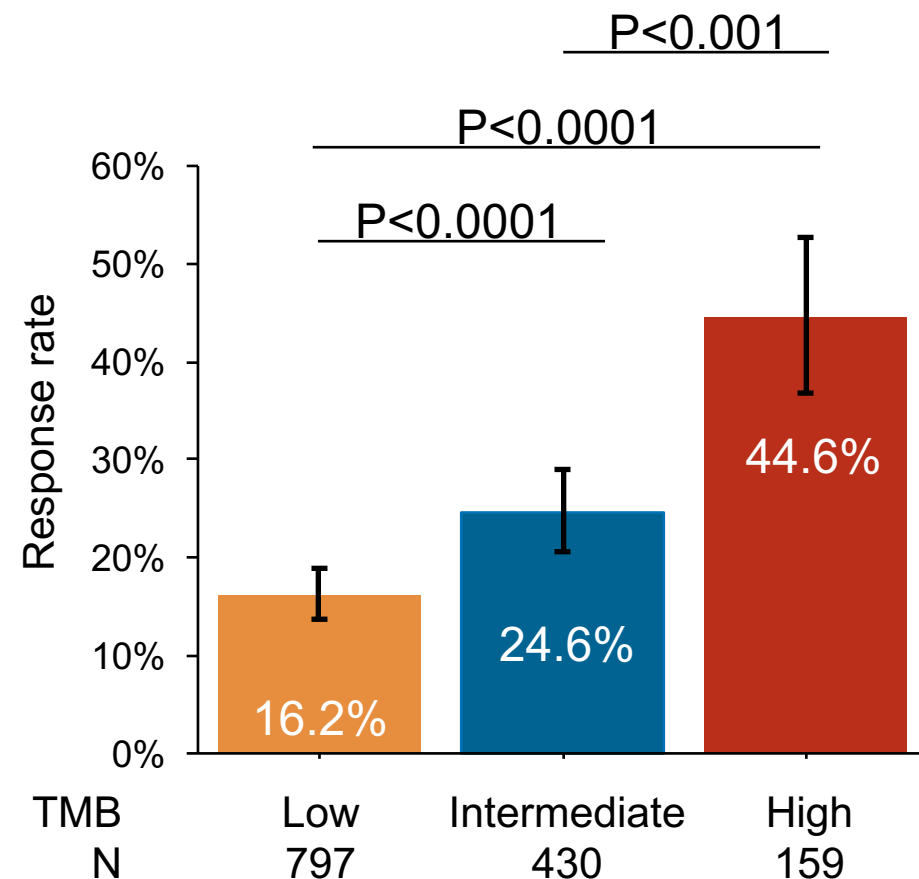
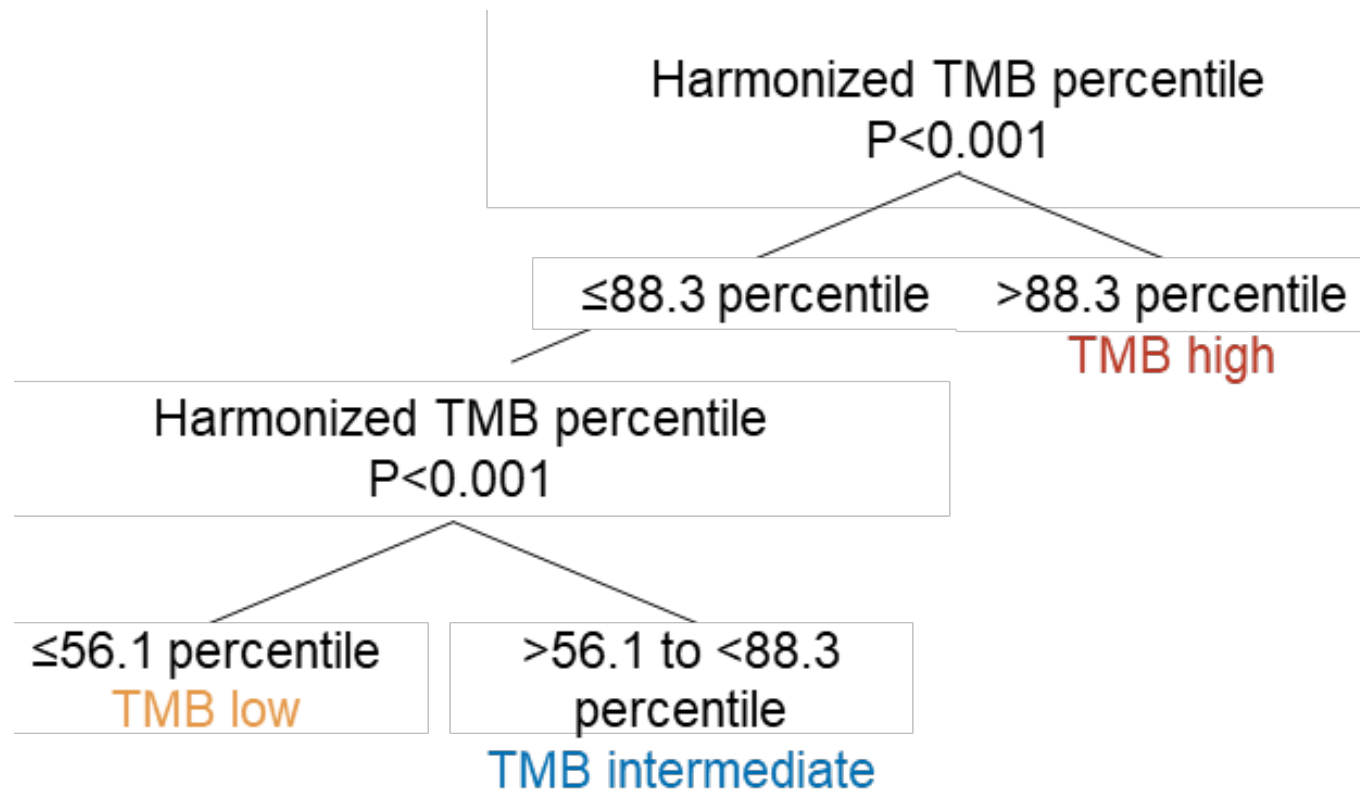


TMB Harmonization

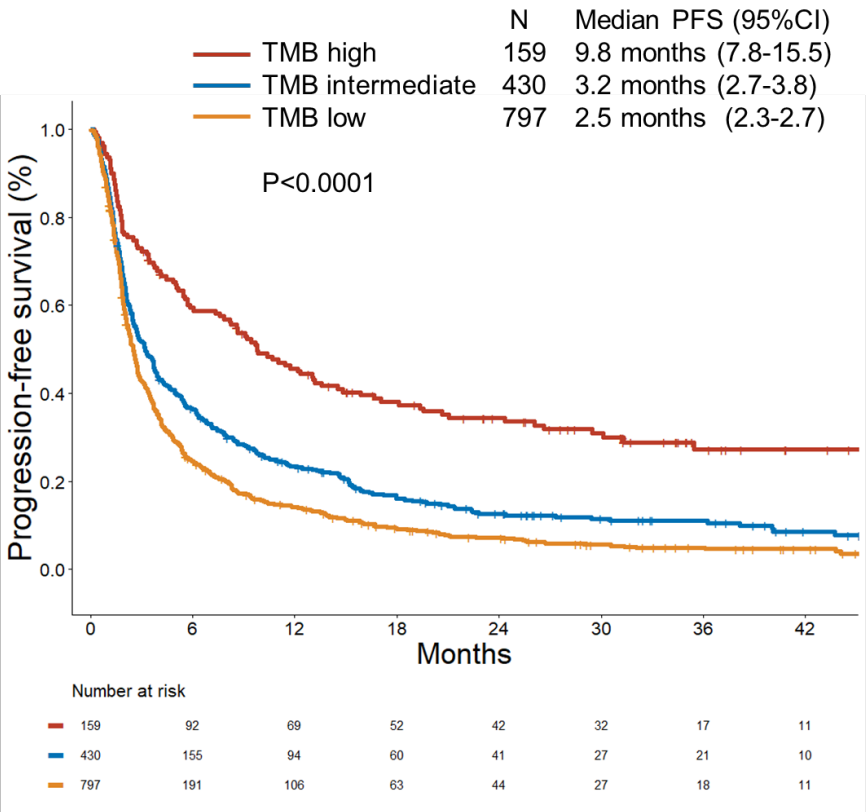


Percentile	TMB z-score	DFCI TMB	MSKCC TMB	Foundation TMB	TCGA TMB (mutation count)
10th	-1.04	4.81	2.27	2.83	1.84 (55)
20th	-0.47	7.22	3.89	4.45	3.35 (101)
30th	-0.24	8.42	4.78	5.30	4.18 (125)
40th	0.00	9.87	5.90	6.36	5.25 (158)
50th	0.17	11.07	6.89	7.27	6.10 (183)
60th	0.45	13.24	8.76	8.97	7.58 (228)
70th	0.70	15.47	10.82	10.80	9.41 (282)
80th	0.95	18.05	13.34	13.00	11.31 (339)
90th	1.38	23.49	19.10	17.90	15.43 (463)

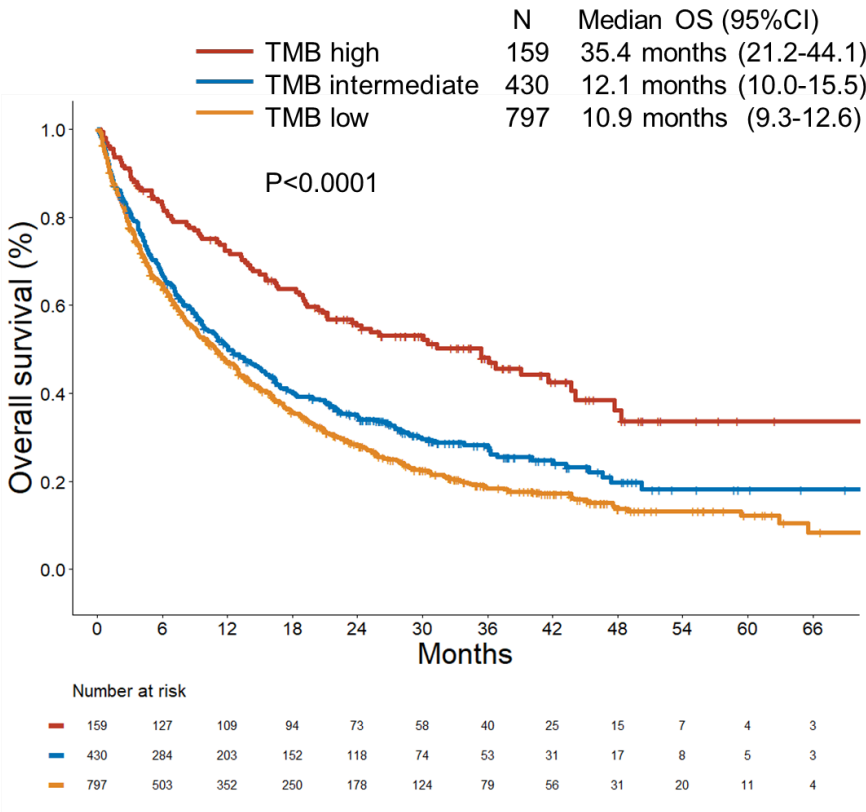
TMB Groupings & PD-1 Response in NSCLC



TMB Groupings & PD-1 Response in NSCLC



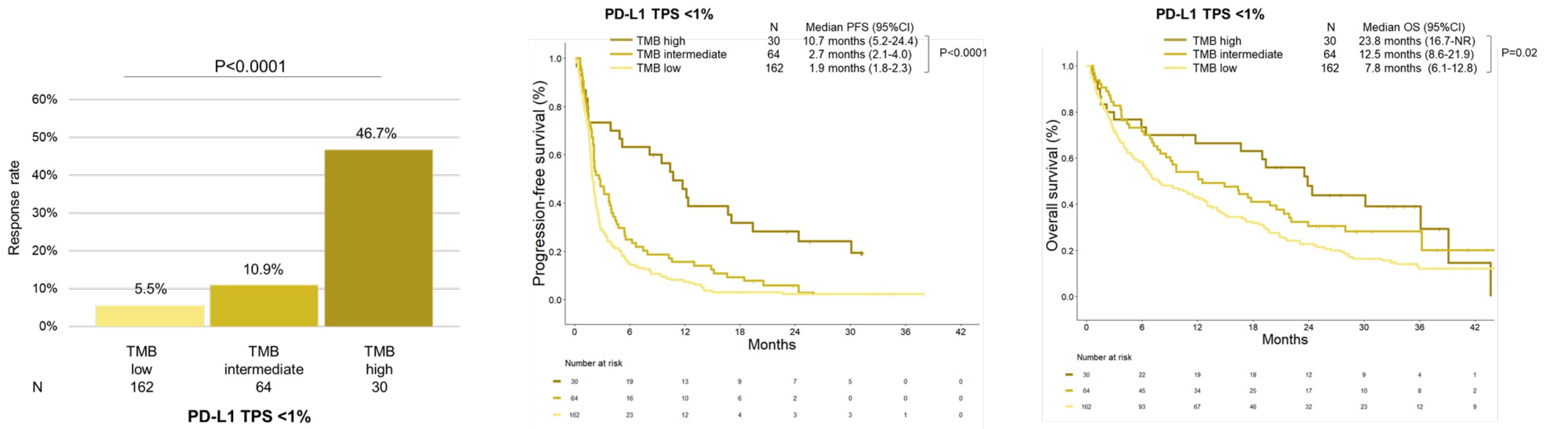
	PFS HR [95%CI], P
TMB high vs TMB low	0.39 [0.32-0.48], P<0.0001
TMB high vs TMB intermediate	0.57 [0.42-0.65], P<0.0001
TMB intermediate vs TMB low	0.76 [0.67-0.86], P<0.0001



	OS HR [95%CI], P
TMB high vs TMB low	0.46 [0.37-0.58], P<0.0001
TMB high vs TMB intermediate	0.56 [0.43-0.71], P<0.0001
TMB intermediate vs TMB low	0.84 [0.73-0.96], P=0.01

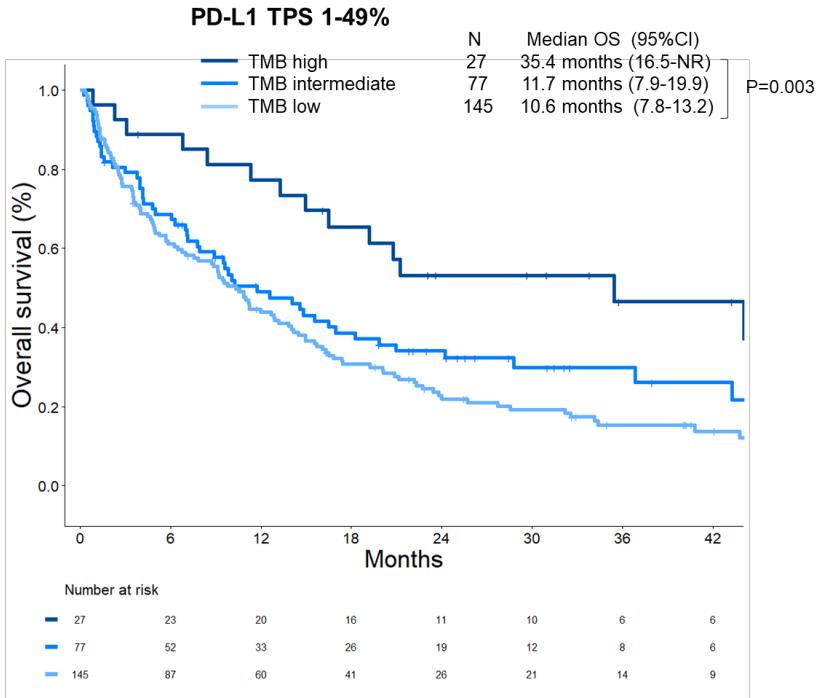
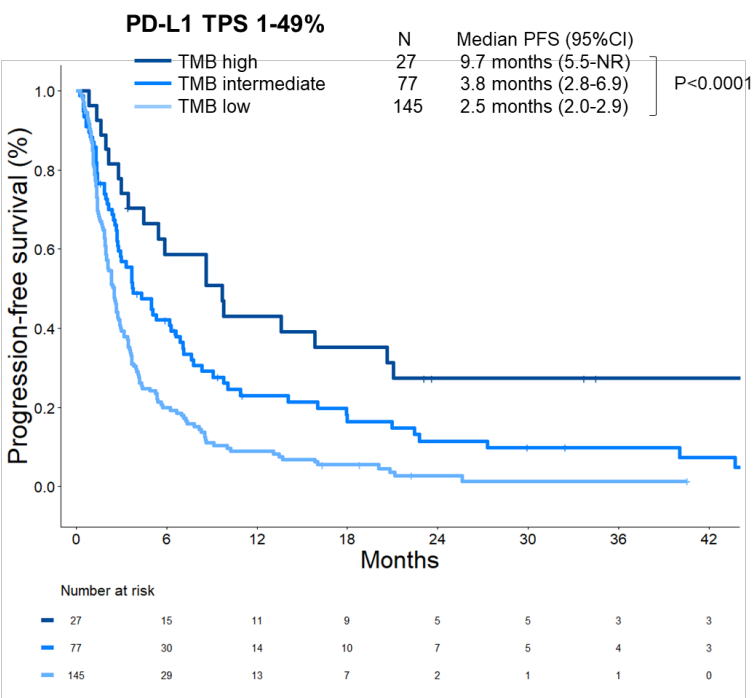
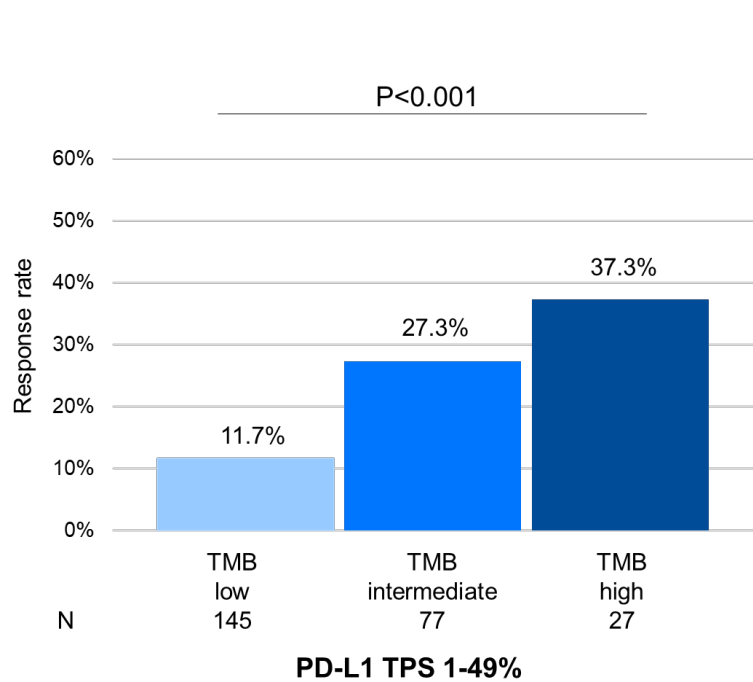
TMB Groupings & PD-1 Response in NSCLC

PD-L1 Negative (TPS <1%) NSCLC



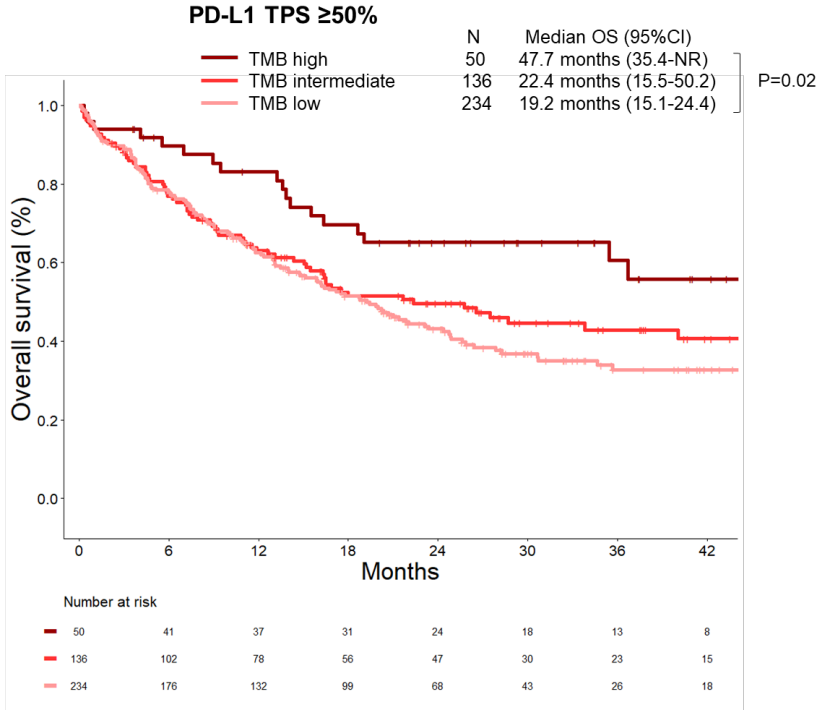
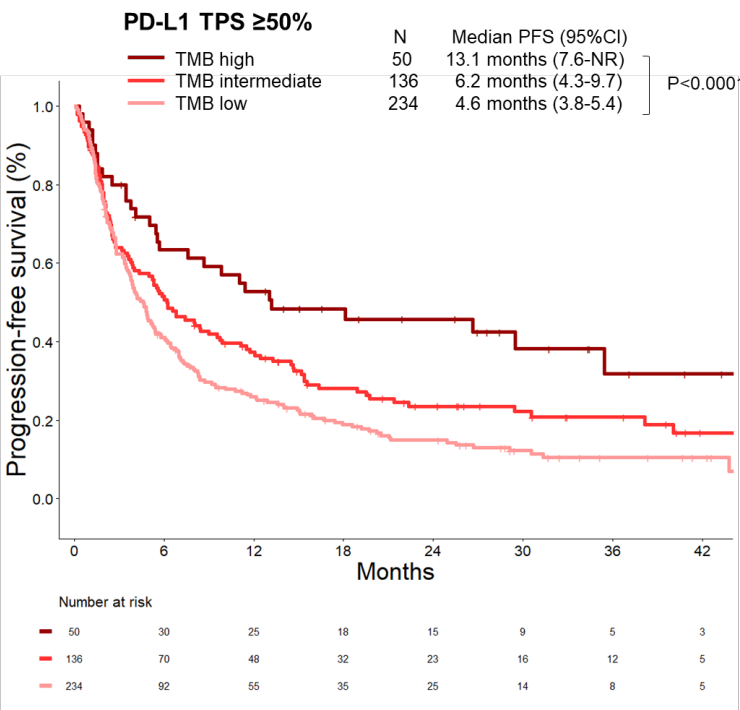
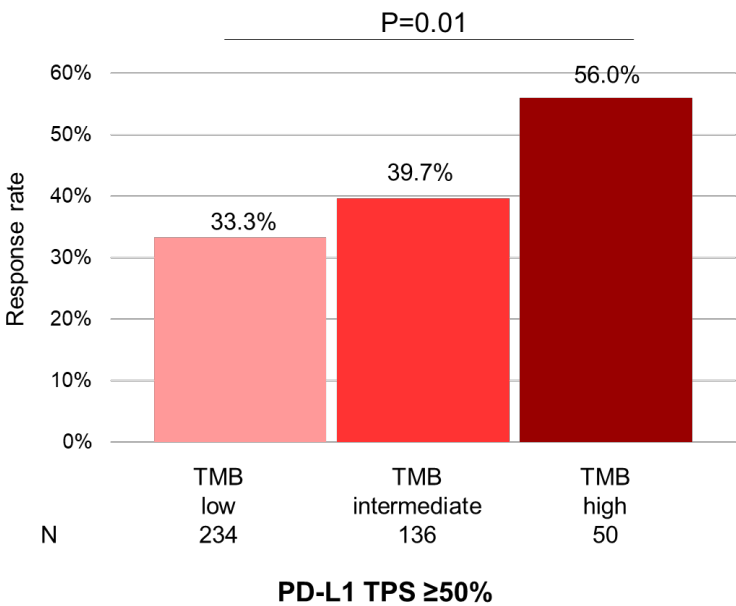
TMB Groupings & PD-1 Response in NSCLC

PD-L1 Low (TPS 1-49%) NSCLC

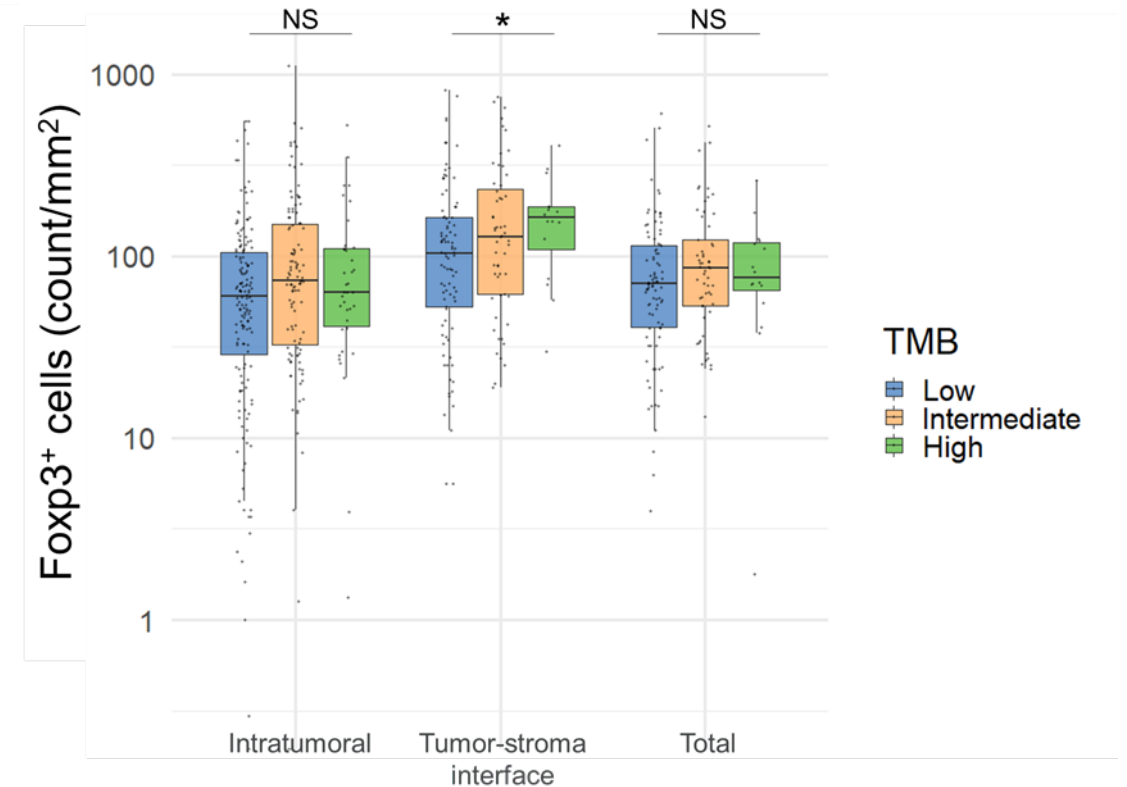
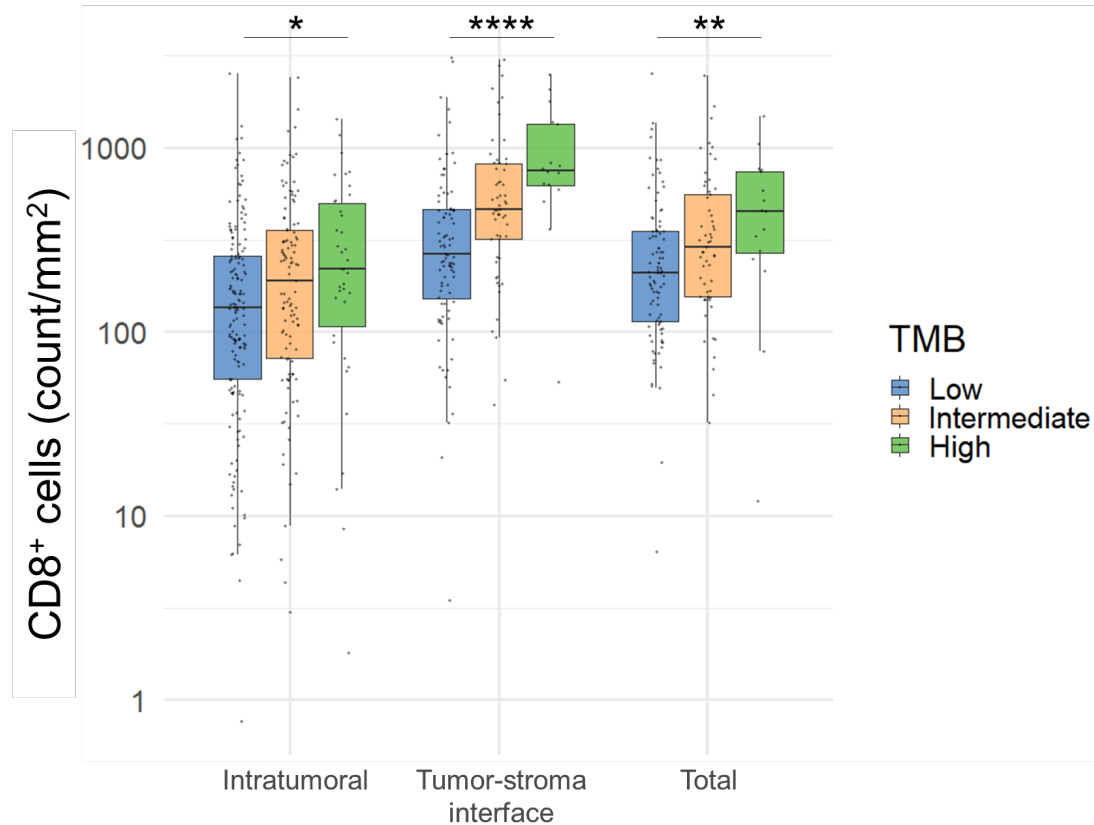


TMB Groupings & PD-1 Response in NSCLC

PD-L1 High (TPS ≥50%) NSCLC



TMB Groupings & Have Distinct Cell Populations

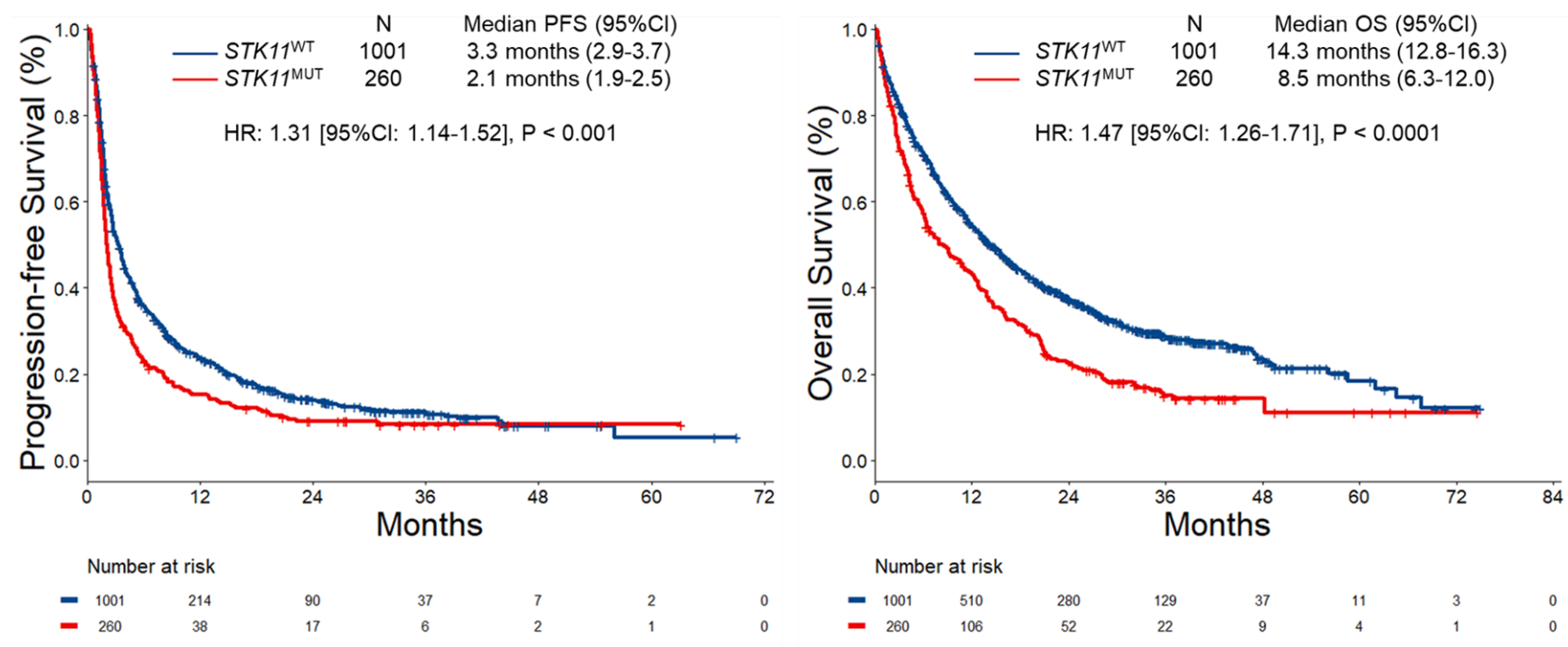


Other Genomic Factors Impacting Immunotherapy Efficacy



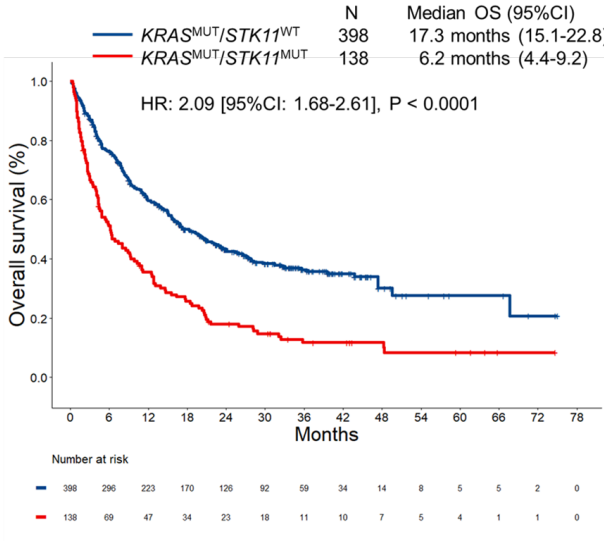
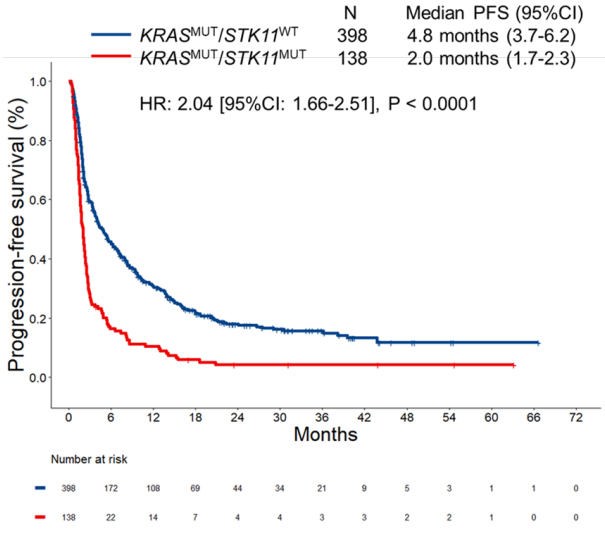
Impact of STK11/LKB1 mutations

Impact of STK11 mutation on immunotherapy efficacy in all comers with NSCLC

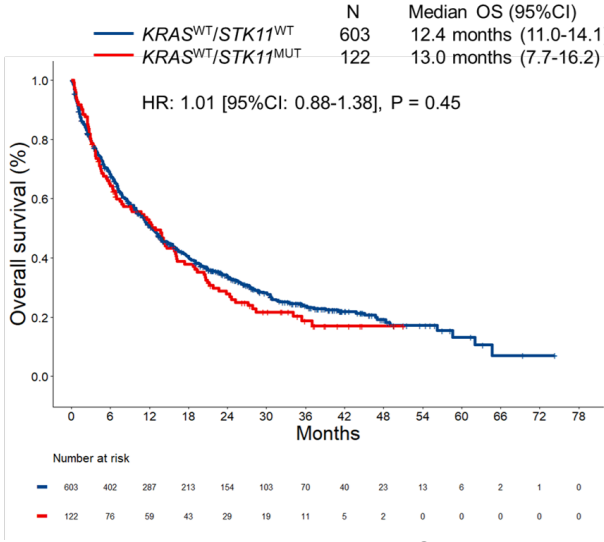
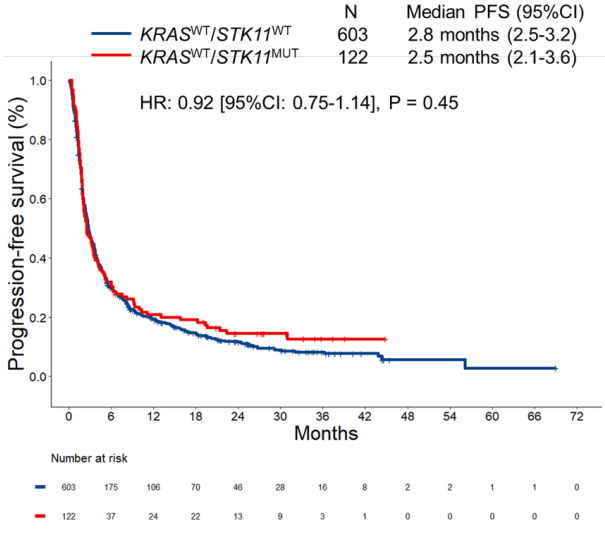


Impact of STK11/LKB1 mutations

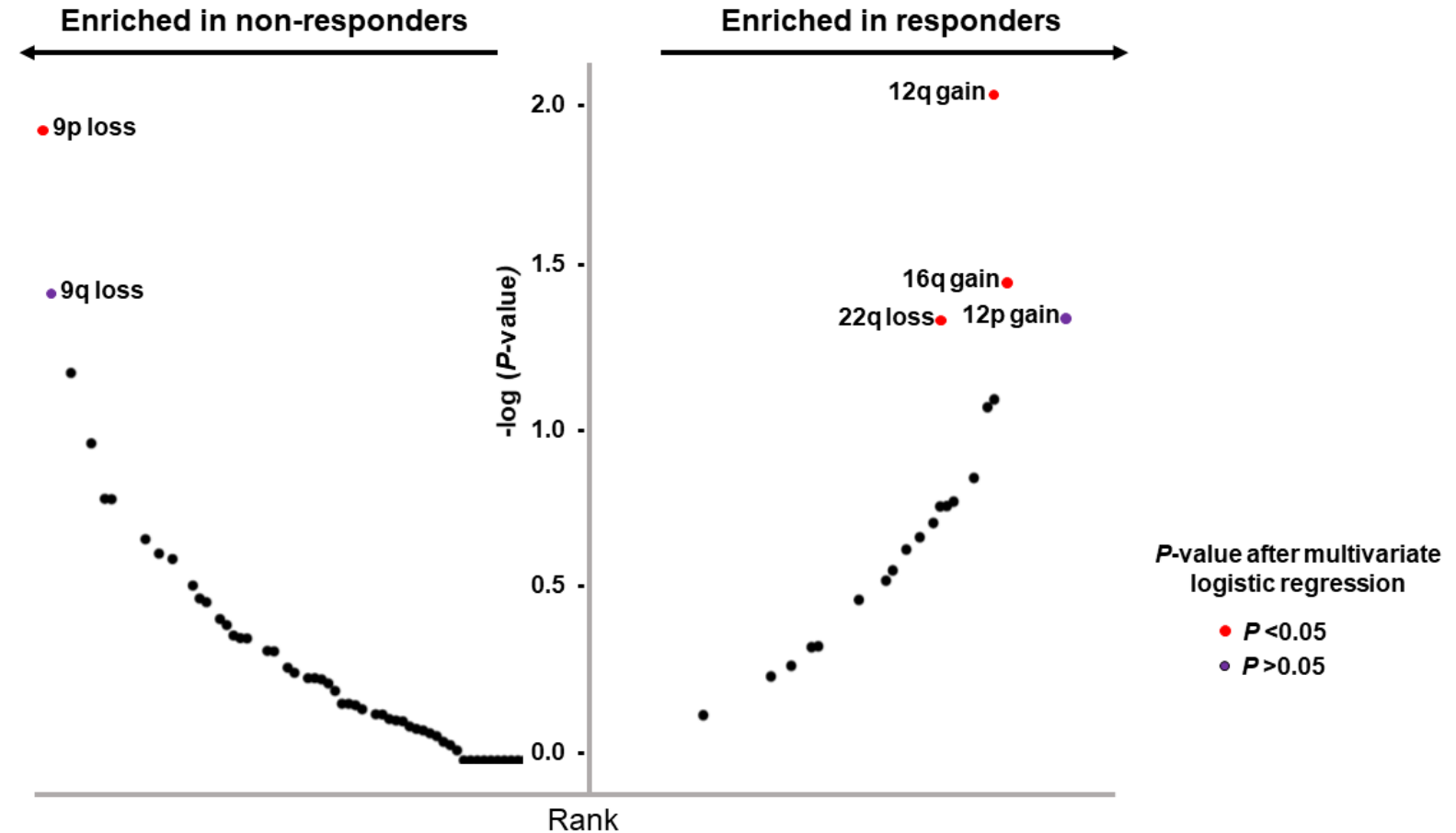
KRAS mutant NSCLC



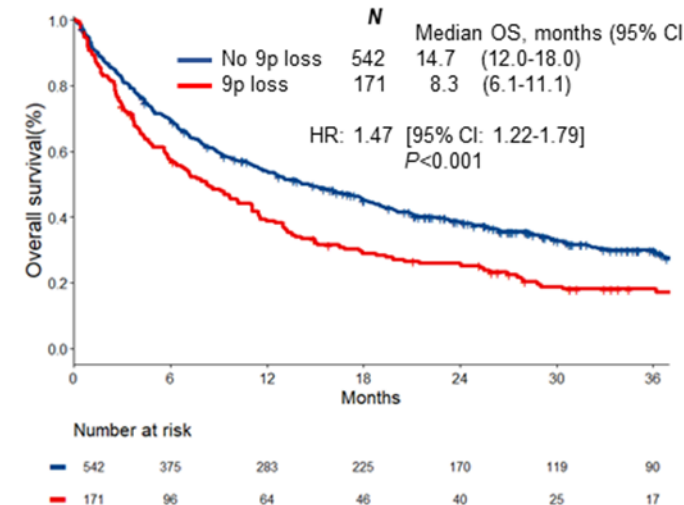
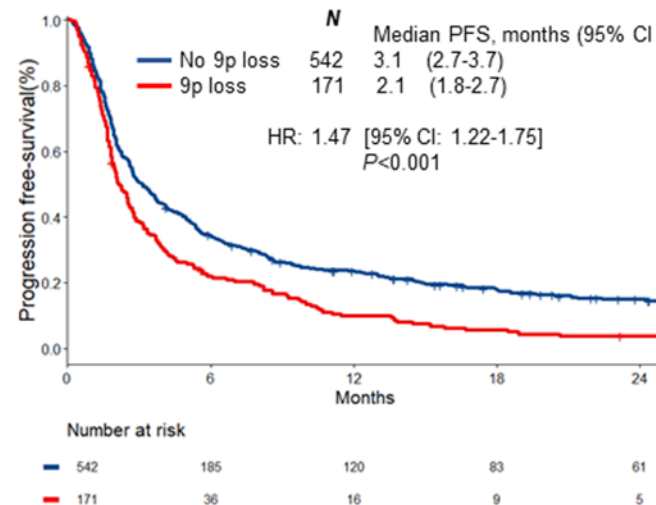
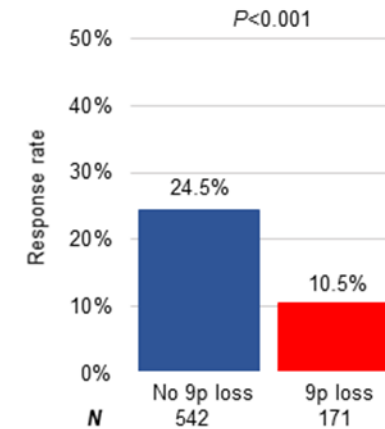
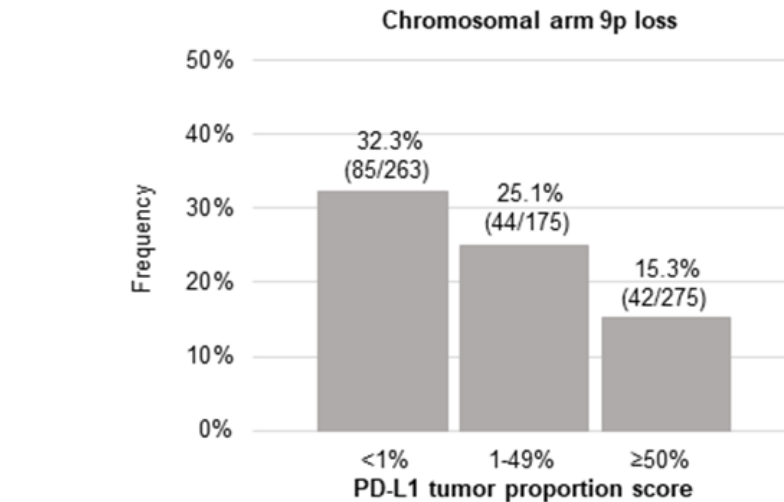
KRAS wild-type NSCLC



Aneuploidy and Immunotherapy Efficacy



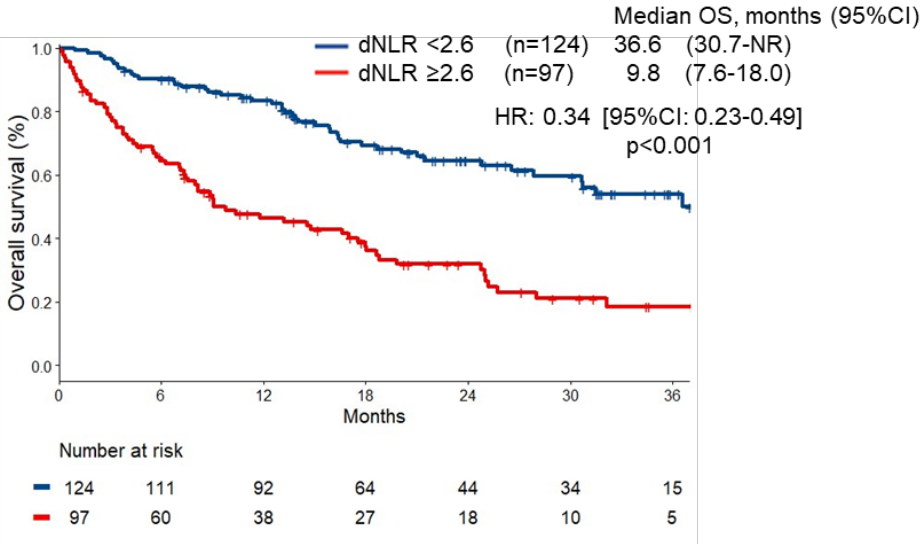
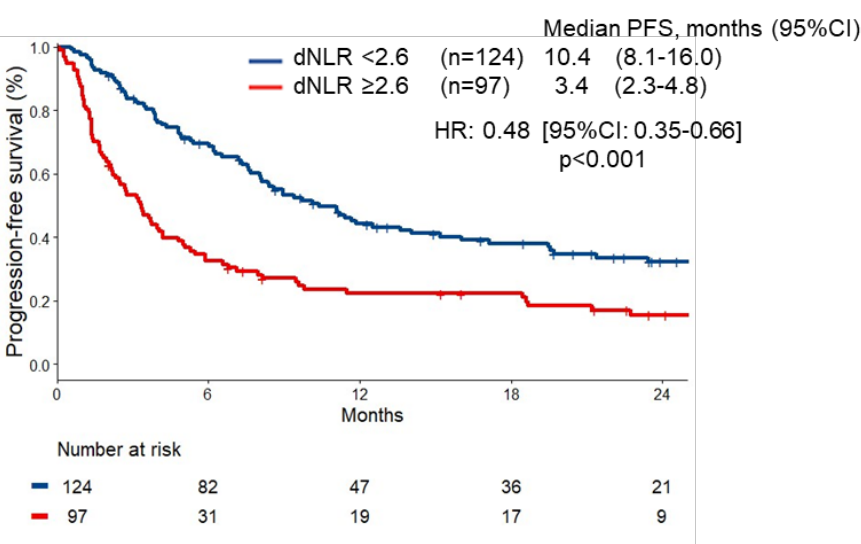
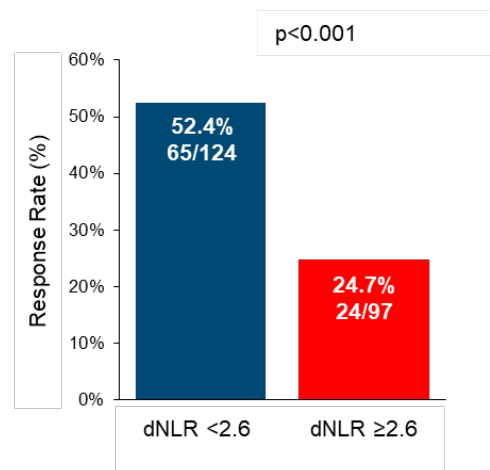
Impact of 9p Loss on Immunotherapy Efficacy



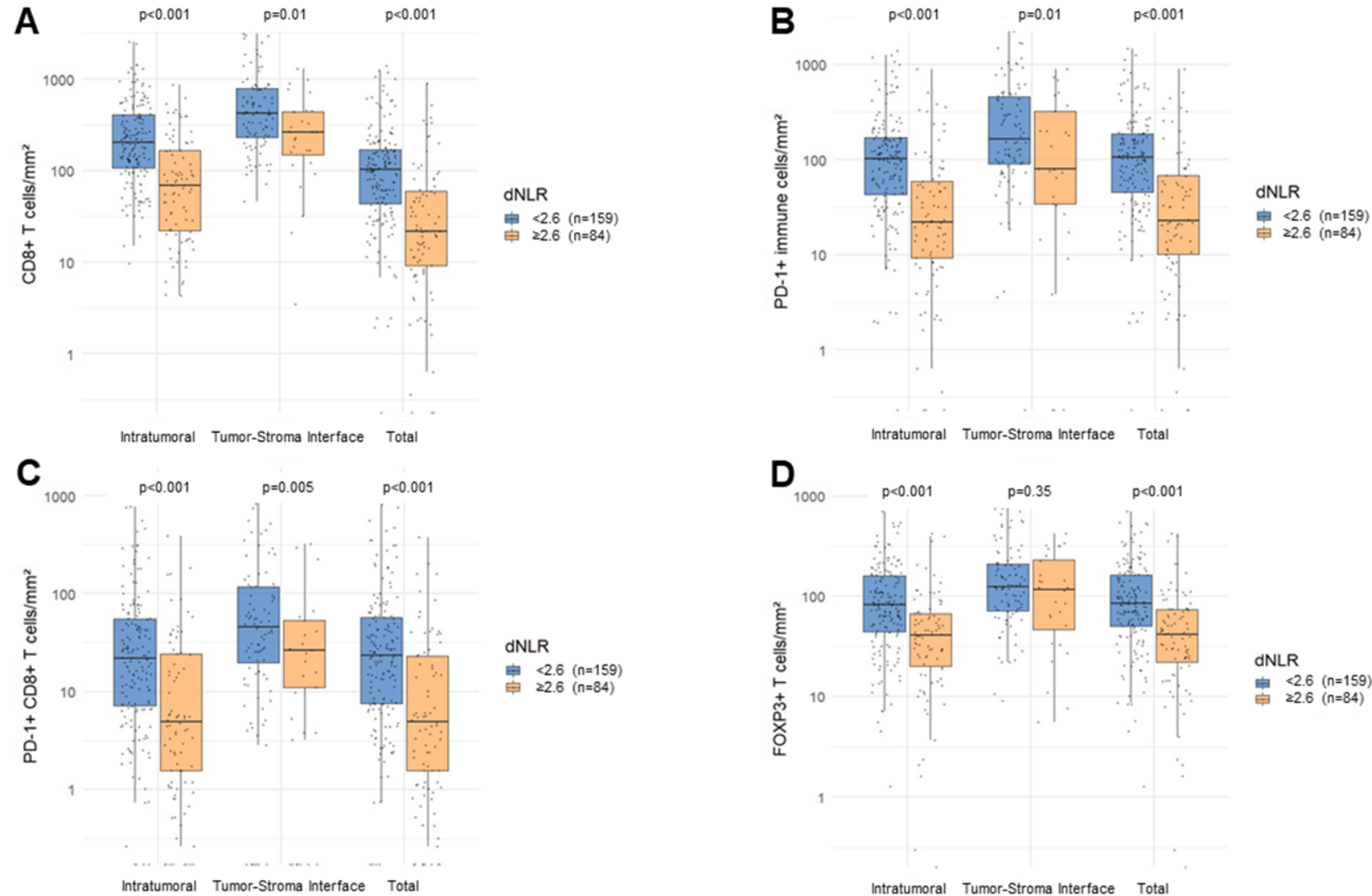
Circulating Factors

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Neutrophil-Lymphocyte Ratio (dNLR) & PD-1 Efficacy



High peripheral blood dNLR correlates with low tumor immune cell infiltration

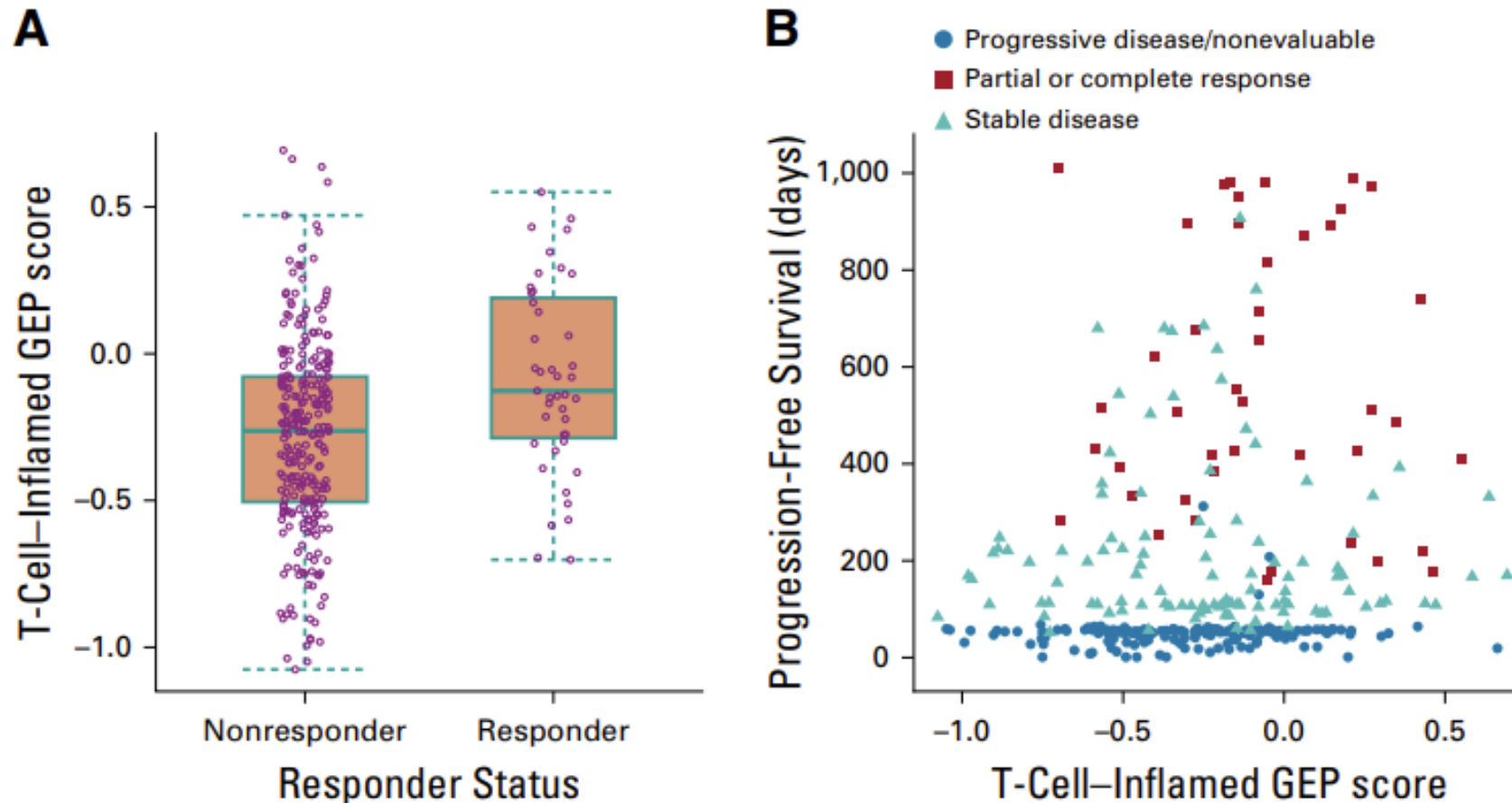


Immune Microenvironment



Gene Expression Profile (GEP) & PD-1 Efficacy

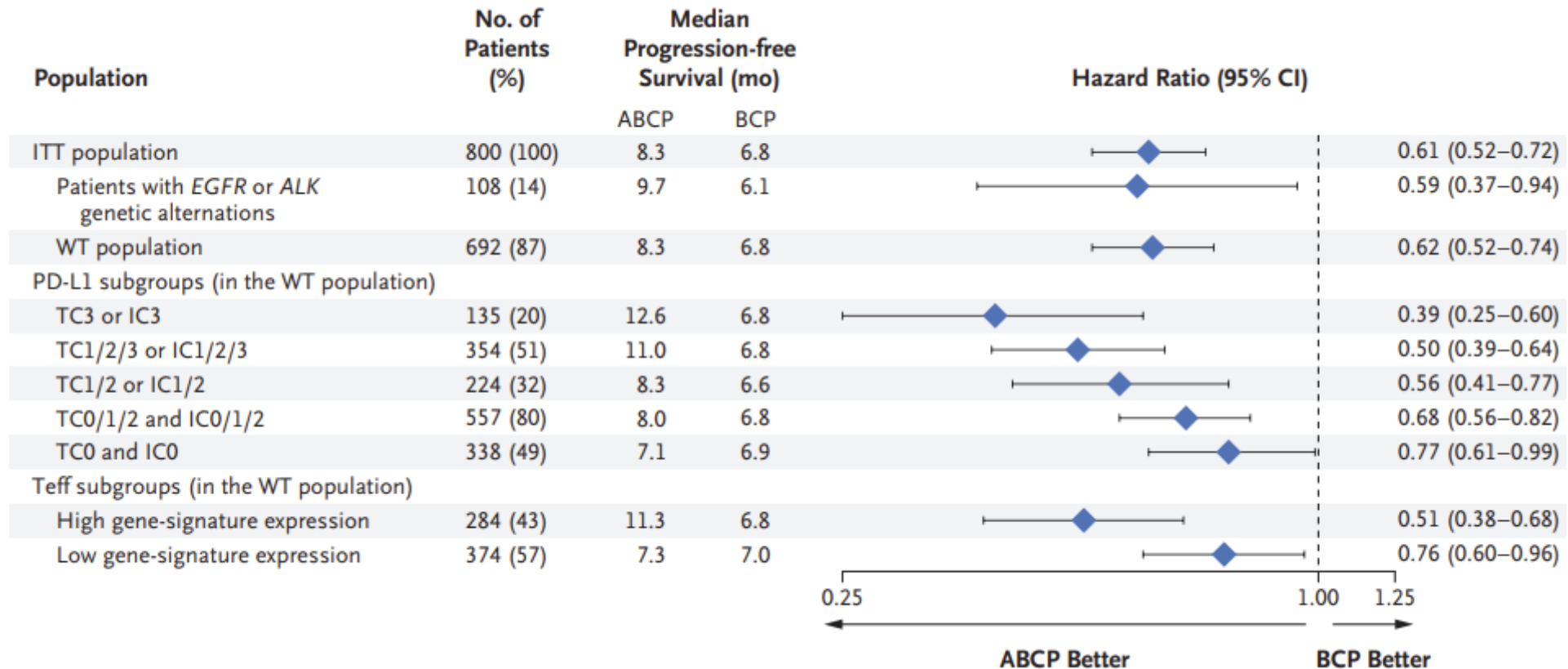
KEYNOTE-028 (Across 20 Cancers)



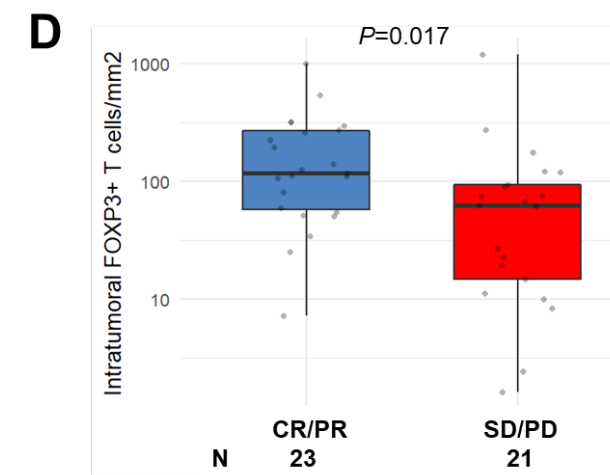
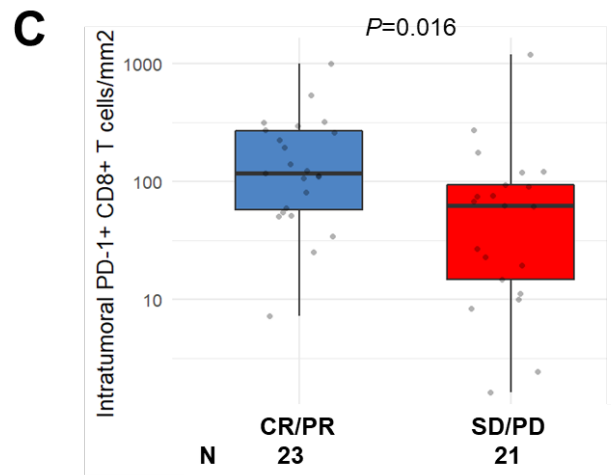
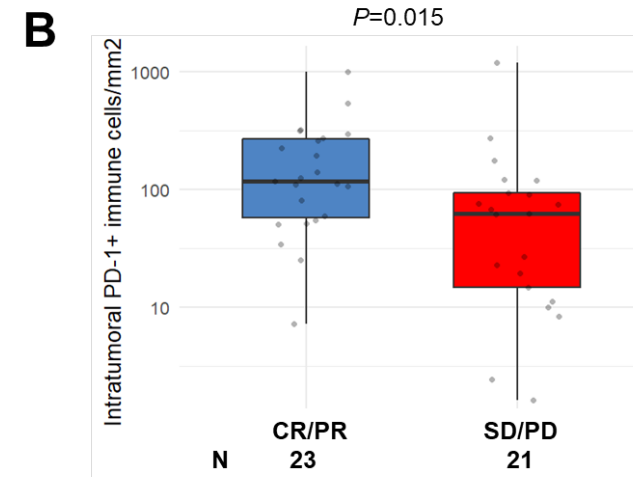
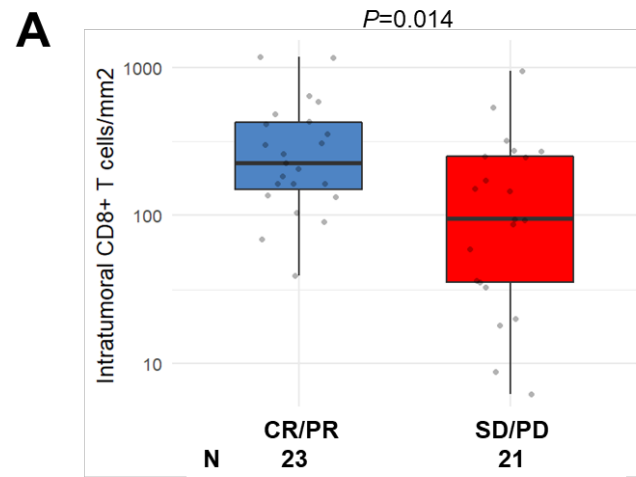
Effector T-cell (Teff) Gene Signature

IMpower150

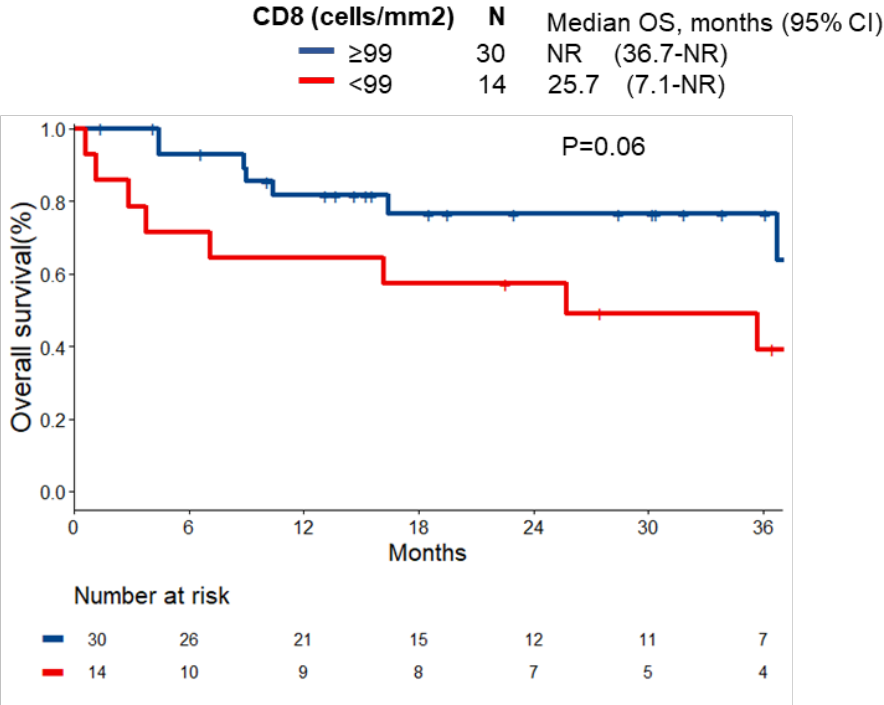
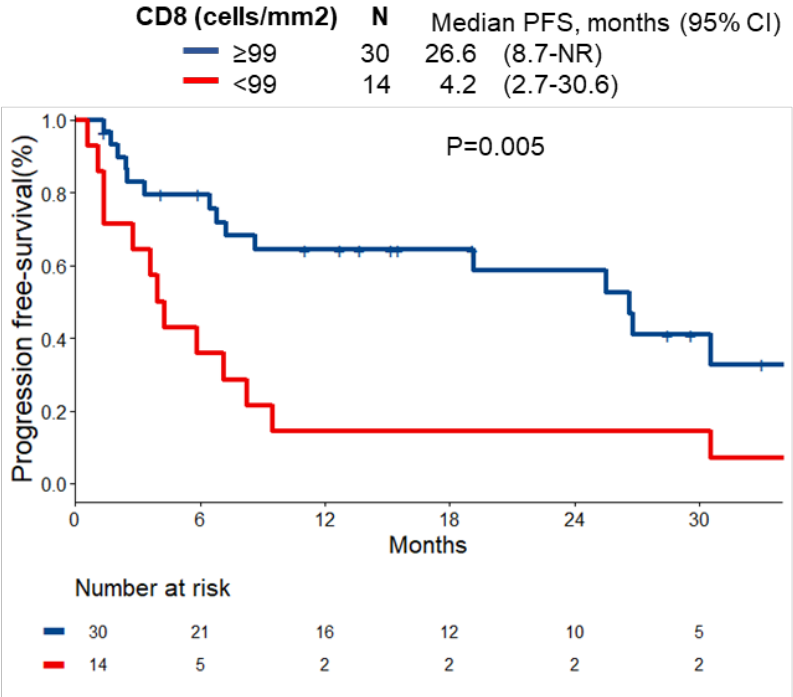
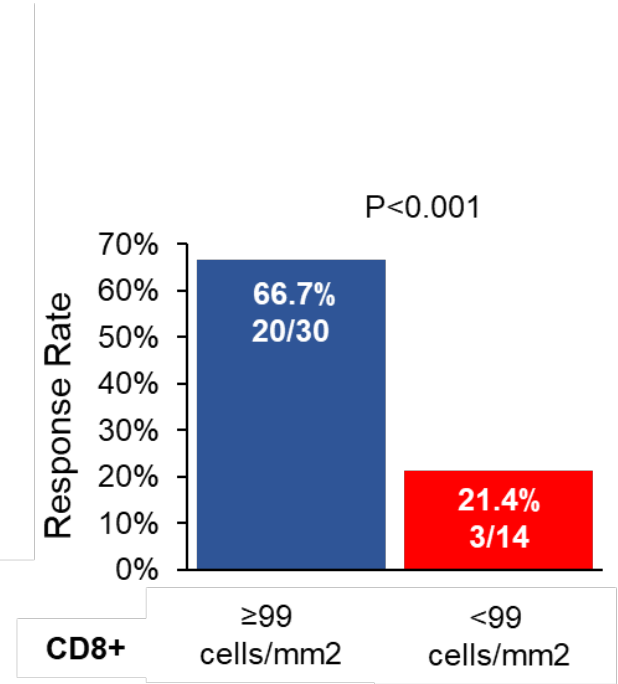
B Hazard Ratios for Disease Progression or Death in Biomarker Subgroups



Immunoprofile & 1st Line Pembrolizumab Response



Immunoprofile & 1st Line Pembrolizumab Response

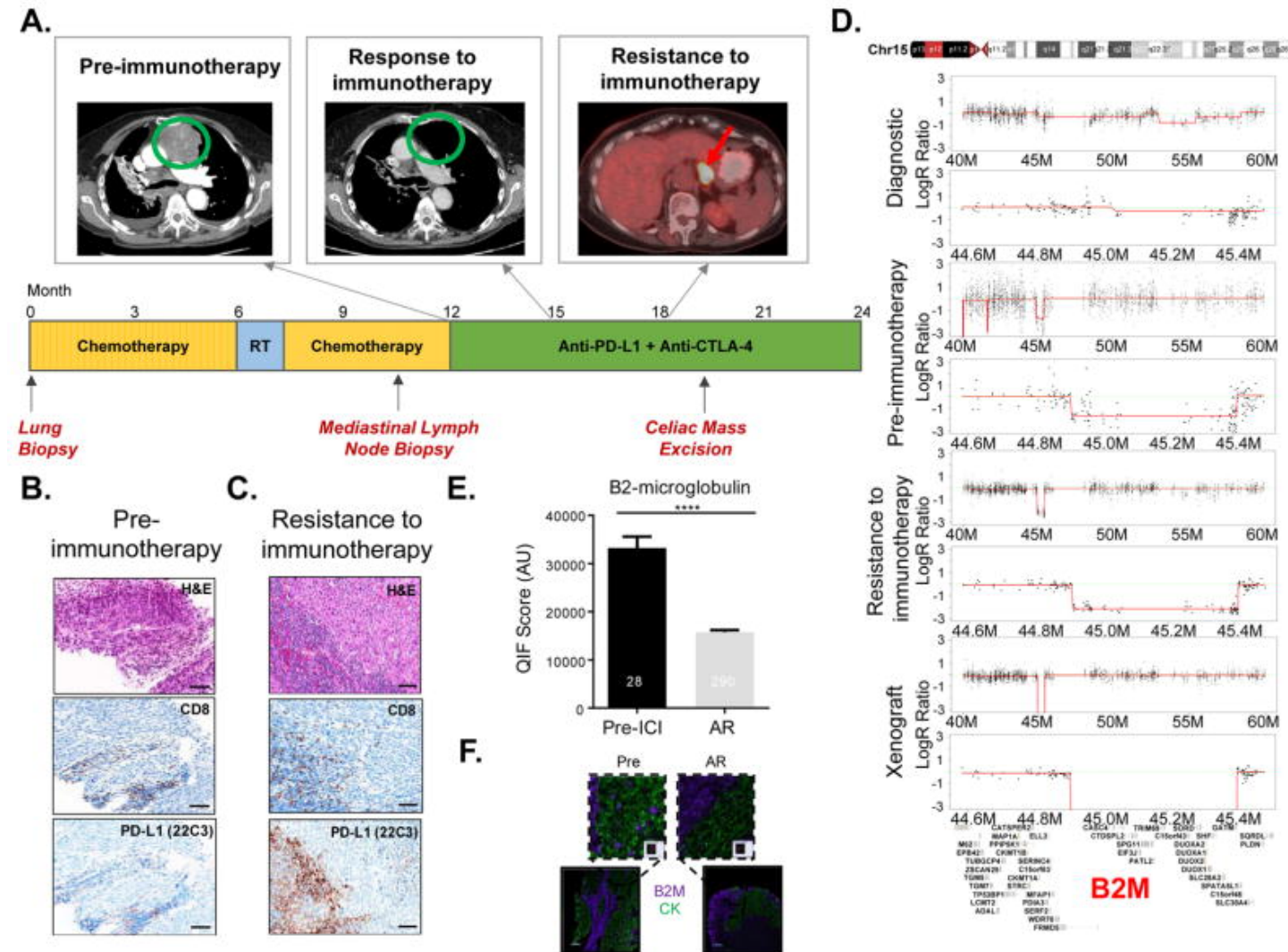


What mediates acquired resistance to immunotherapy in lung cancer?

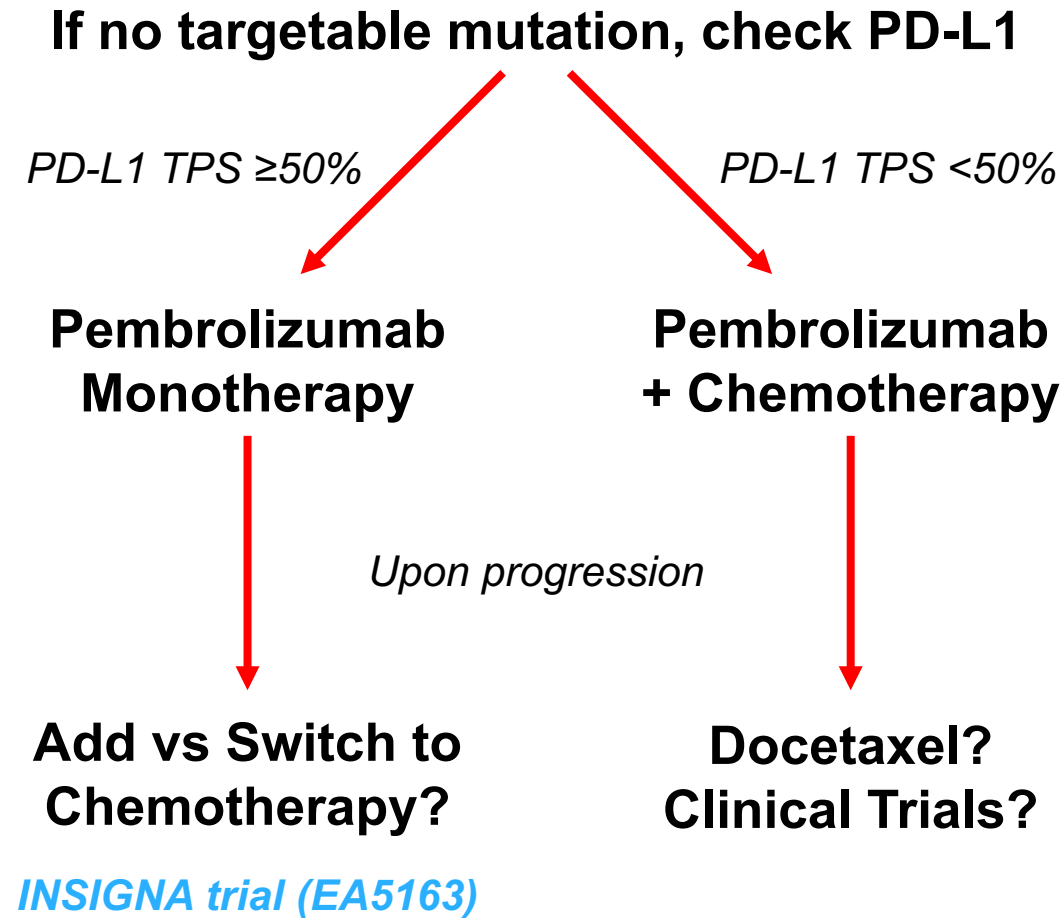


Impaired HLA Class I Antigen Presentation

14 ICI-resistant lung cancer samples



Management of Resistance to Immunotherapy



Conclusions



- Several clinical, pathologic, genomic, and immunologic factors impact primary response and resistance to immune checkpoint inhibitors in lung cancer
- New strategies need to be considered for integrating multiple, continuous biomarkers (e.g. PD-L1 expression, TMB, TILs) to predict response and resistance to immunotherapy
- Mechanisms of primary and acquired resistance to PD-1 inhibitors are still not well understood
- Additional diagnostics and therapeutics are necessary to make precision immunotherapy a reality in clinic to prevent and overcome treatment resistance

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