

Immunotherapy: the quest for a biomarker

Prof Keith M Kerr

Department of Pathology,

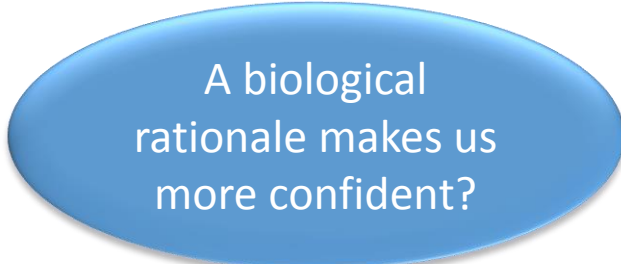
Aberdeen University School, Aberdeen Royal Infirmary, UK

Disclosures

- I have acted as consultant for Roche Genentech, Astra Zeneca, Pfizer, Eli Lilly, Novartis, Boehringer Ingelheim, Clovis, Bristol Myers Squibb, Merck Sharp Dohme
- I have received honoraria for speaker bureau from Roche Genentech, Astra Zeneca, Pfizer, Eli Lilly, Novartis, Boehringer Ingelheim, Bristol Myers Squibb

Biomarkers

- Biological features which are associated with disease behaviour
- Predictive biomarkers 'predict' the likely outcome from a therapy
- The ideal biomarker: always correct
 - Easy and practical to measure
 - Present or absent
 - Stable and functionally unique
 - 100% predictive
- Usually biologically related to the system being examined
 - The drug target
 - A co-factor of the drug target
 - A factor negating drug effect

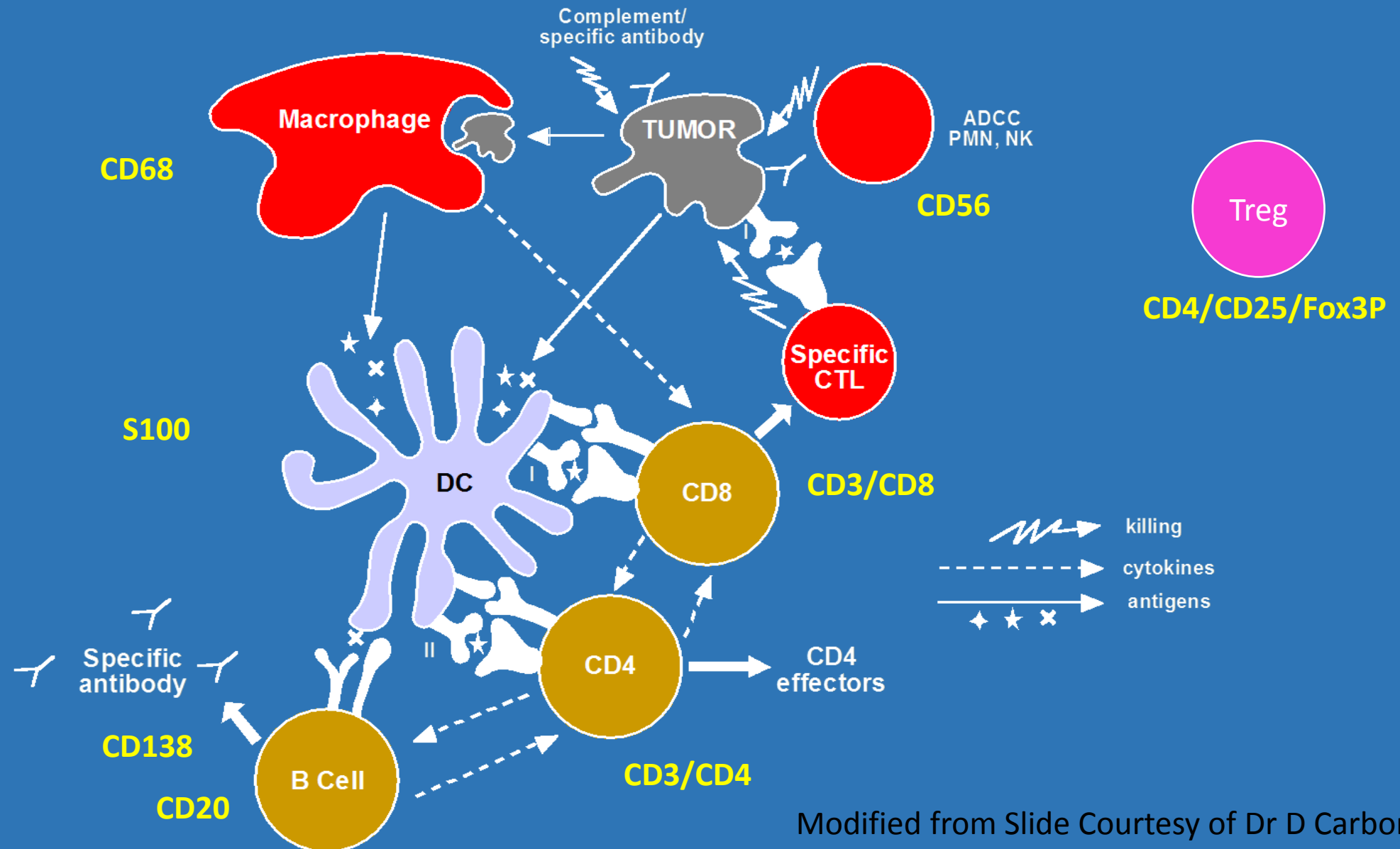


A biological rationale makes us more confident?

Biomarkers in lung cancer (so far.....)

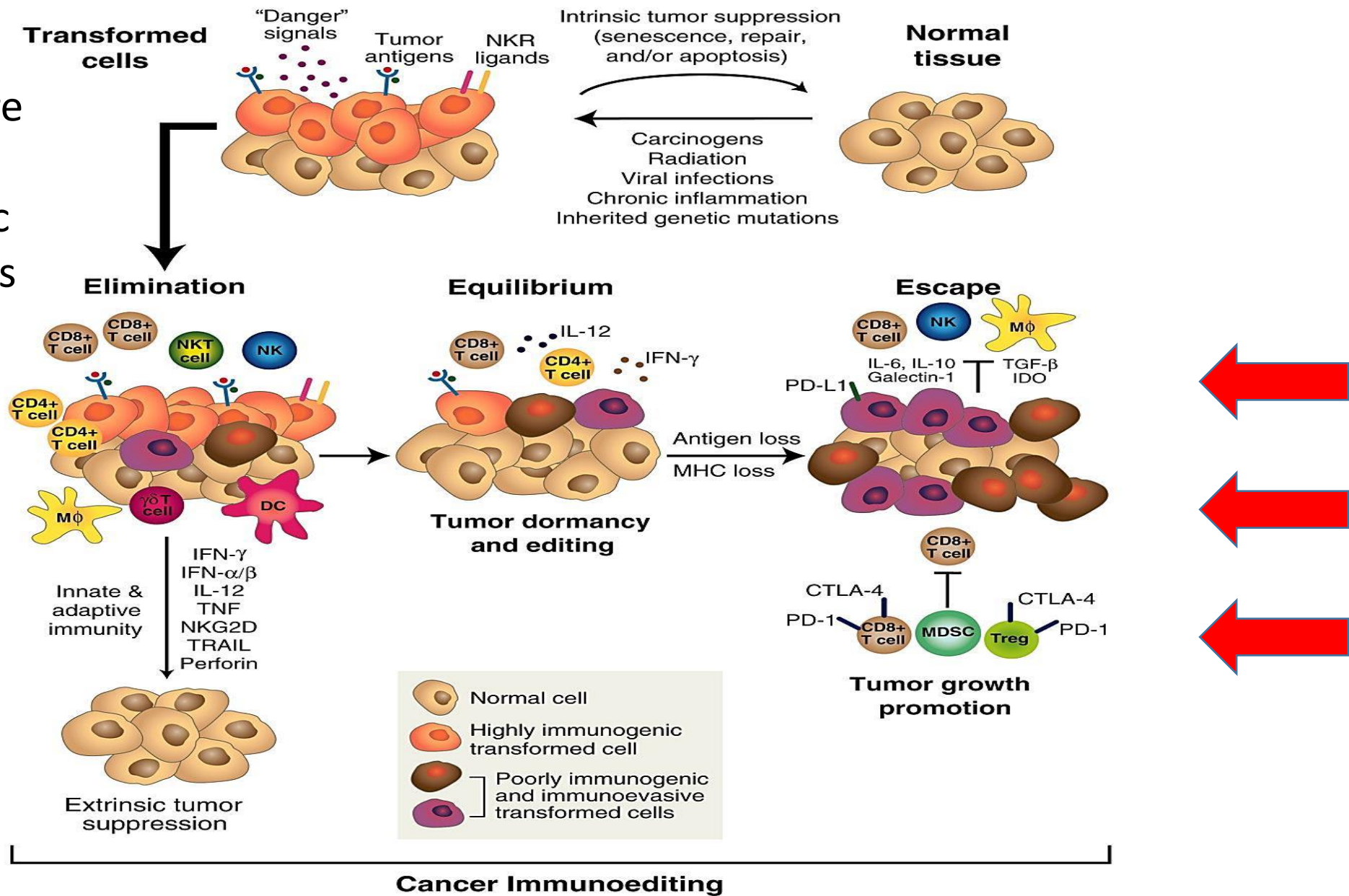
- Our experience is with the 'low hanging fruit'
- Addictive oncogenes
 - Main driver of tumour
 - Mutation or translocation: relatively easy to measure
 - 'Present or Absent' at least in terms of current thinking
- And yet.....
 - Best response rates are 60-70%
 - Testing is not 'fool-proof'

Tumour immune response are VERY complex and involve many factors

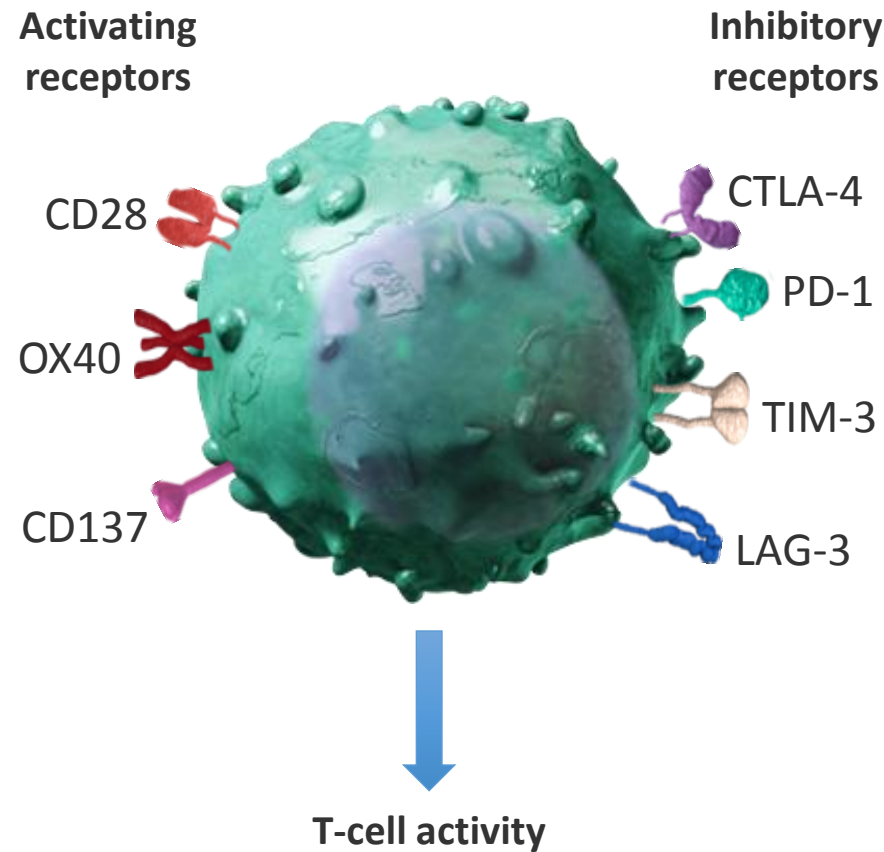


Modified from Slide Courtesy of Dr D Carbone, MD

Lung cancers are amongst the MOST Antigenic of solid tumours

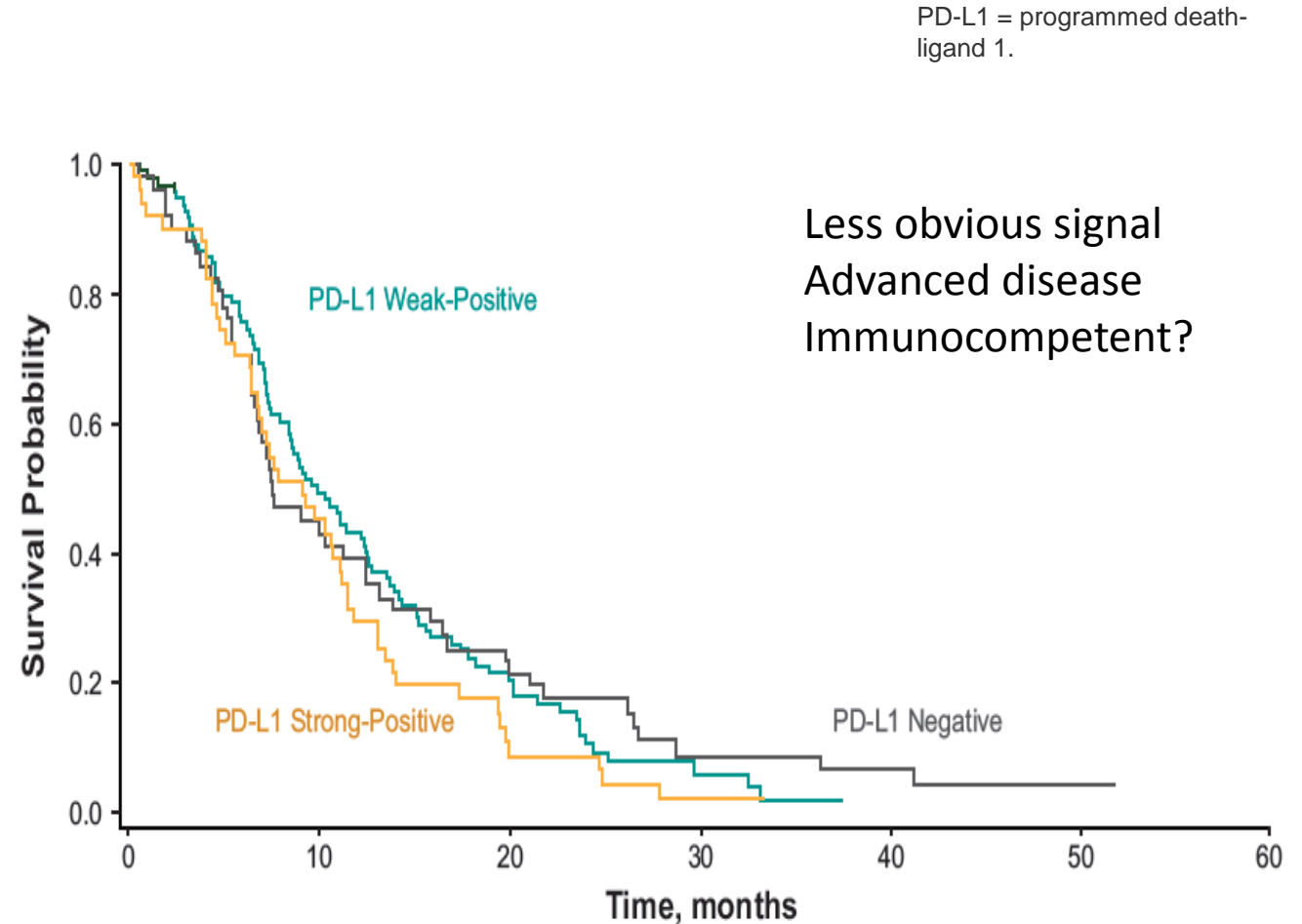
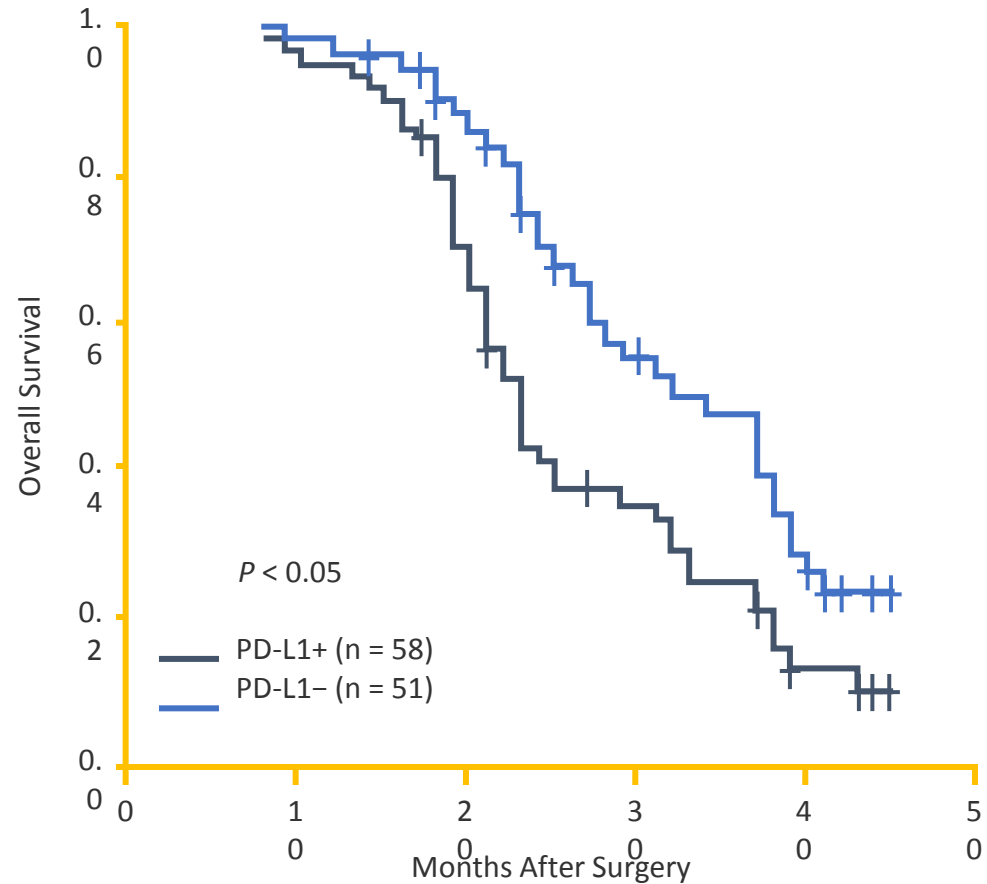


Tumours may alter T cell activation through immune checkpoint signaling



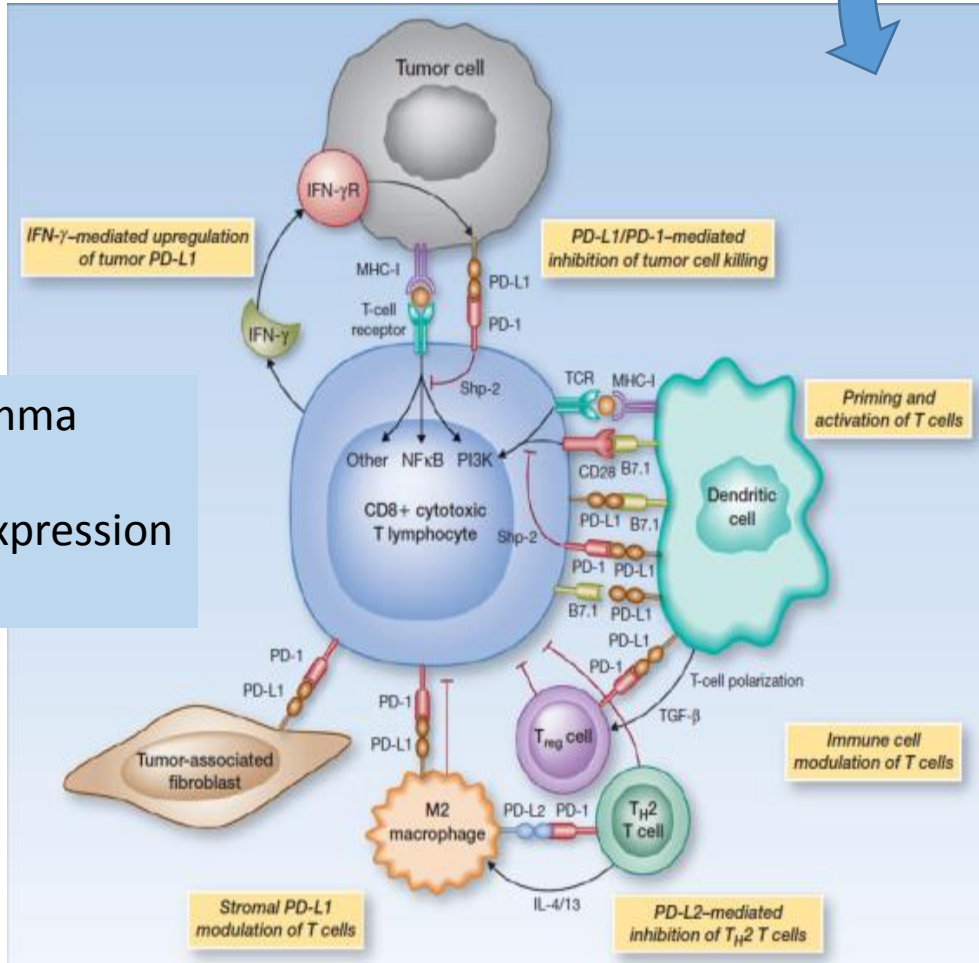
- These inhibitory checkpoints probably help host (normal) tissues avoid autoimmune responses
- Tumours can dysregulate checkpoints and activating pathways, and consequently inhibit the immune response
- Targeting checkpoints can reactivate an immune response

The inhibitory (checkpoint) molecule PD-L1 is associated with poor prognosis in patients with resected NSCLC^{1,2}



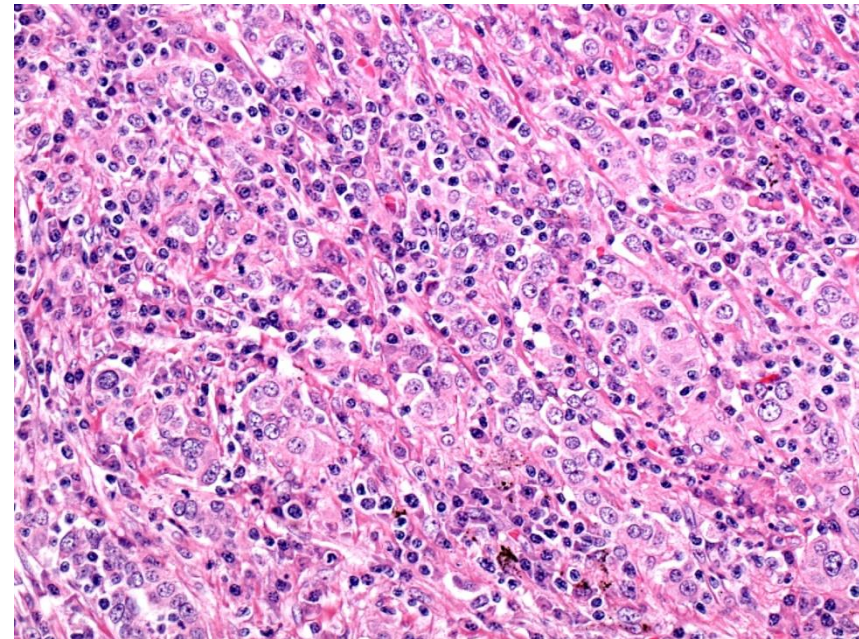
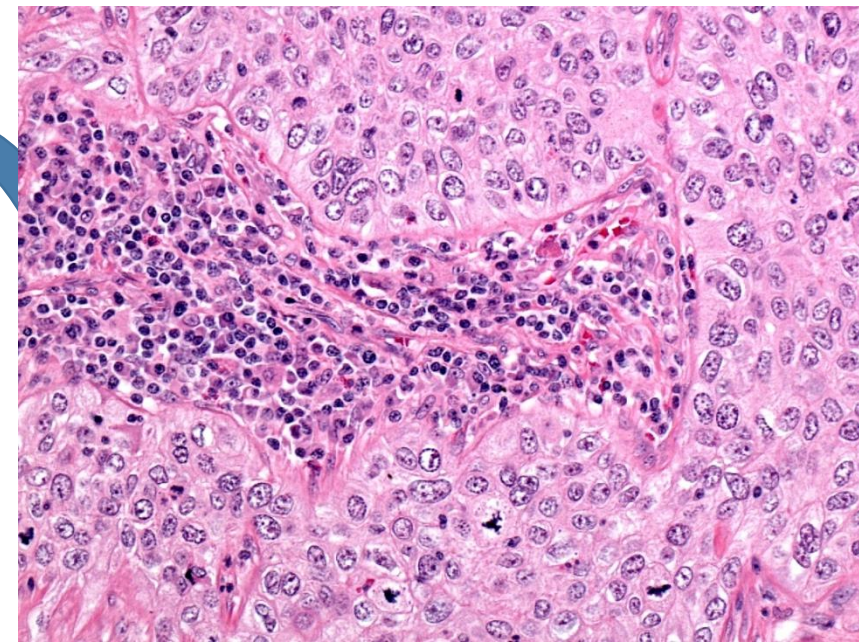
High levels of PD1 or PDL1 protein expression (IHC) may inhibit Immune response

Interferon gamma upregulates Tumour cell expression of PD-L1



Chen, et al. Clin Cancer Res 2012

Block PD1 or PDL1
Immune damage to tumour



Headline studies: Outcomes and PD-L1 expression

Agent	Study	Study Design	Treatment Line	Histology	PD-L1 Positive Definition	PD-L1 positive, %	ORR, % (n/N)		Median PFS, months		Median OS, months	
							Positive	Negative	Positive	Negative	Positive	Negative
Nivolumab	CheckMate 063 [Rizvi 2015]	Phase 2, single arm	≥3rd	Squamous	≥5% in ≥100 cells	33	24 (6/25)	14 (7/51)	NA	NA	NA	NA
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Nivolumab + ipilimumab	CA209-012 [Antonia SJ ASCO 2014]	Phase 1, multi-cohort	1st	Any	≥5% in ≥100 cells	42	19 (3/16)	14 (3/22)	3.3	3.1	NR	NR
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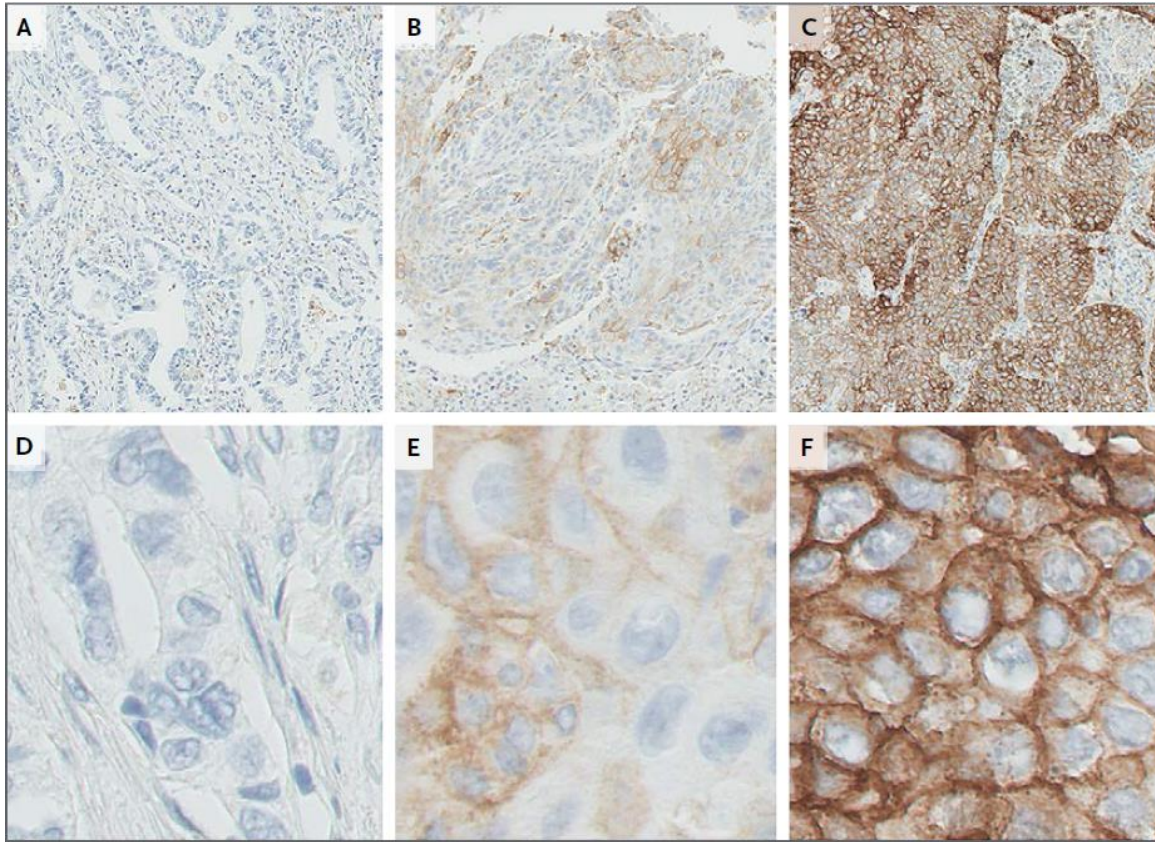
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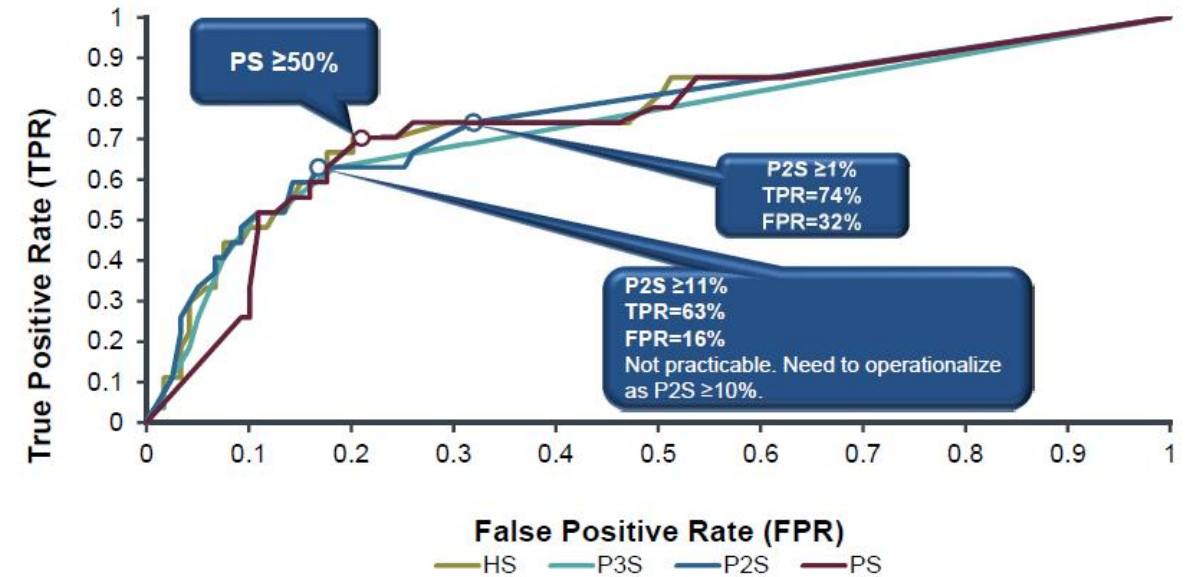
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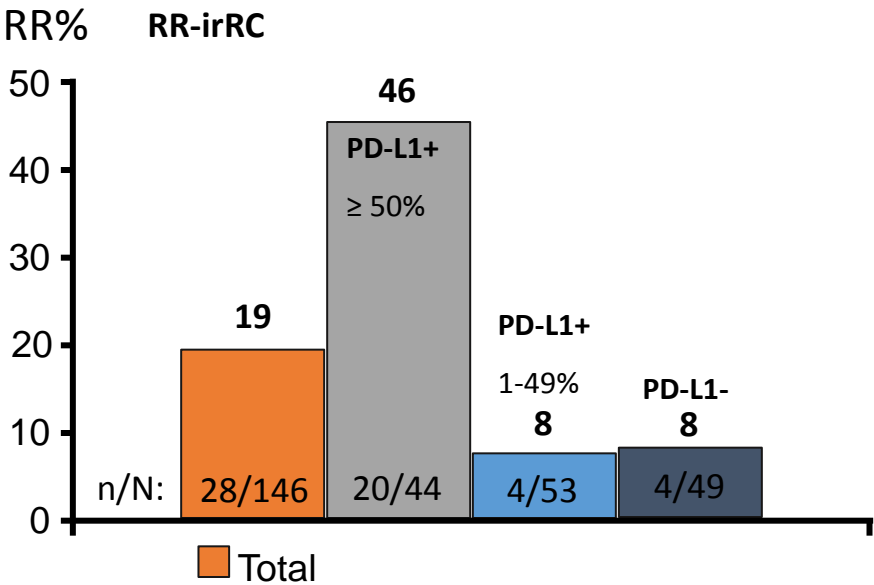
22C3 assay and Pembrolizumab



Comparison of Scoring Methods and Cutoffs Using ROC Analysis with Unconfirmed irRC



Pembrolizumab in NSCLC: OS Per Proportional Scores (TPS)

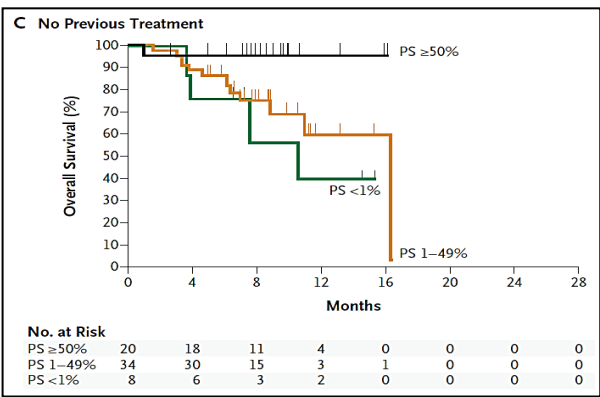
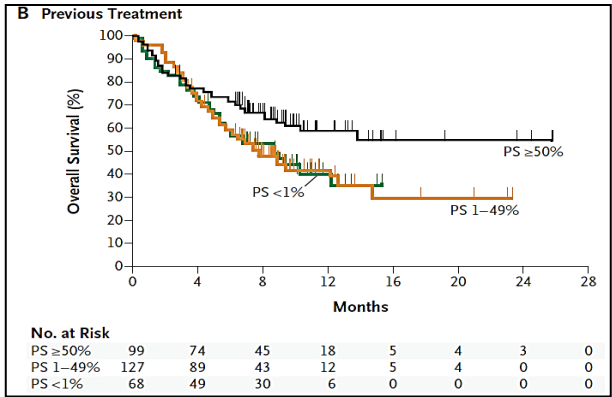
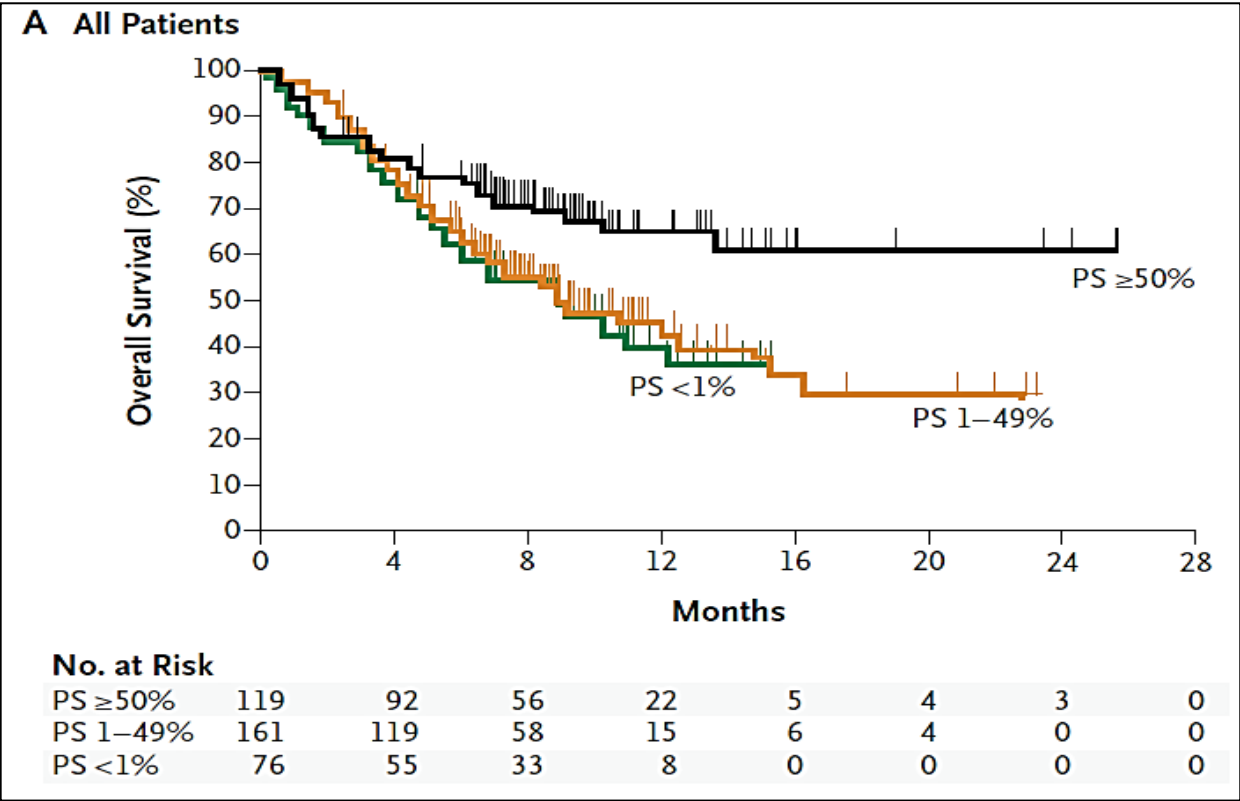


Gandhi L, et al. AACR 2014. Abstract CT105.

$\geq 50\%$ IHC cut off Tumour cell expression

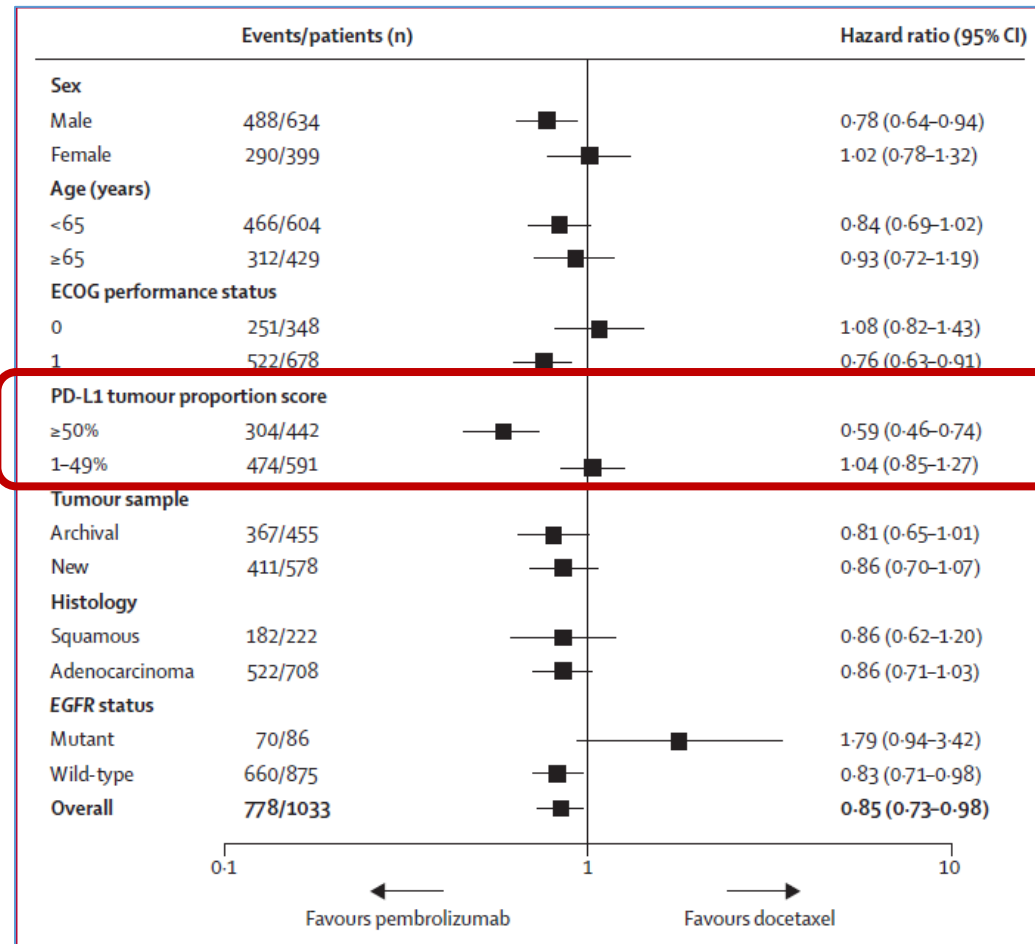
22C3 clone IHC based assay

Expression in TILs added no predictive value

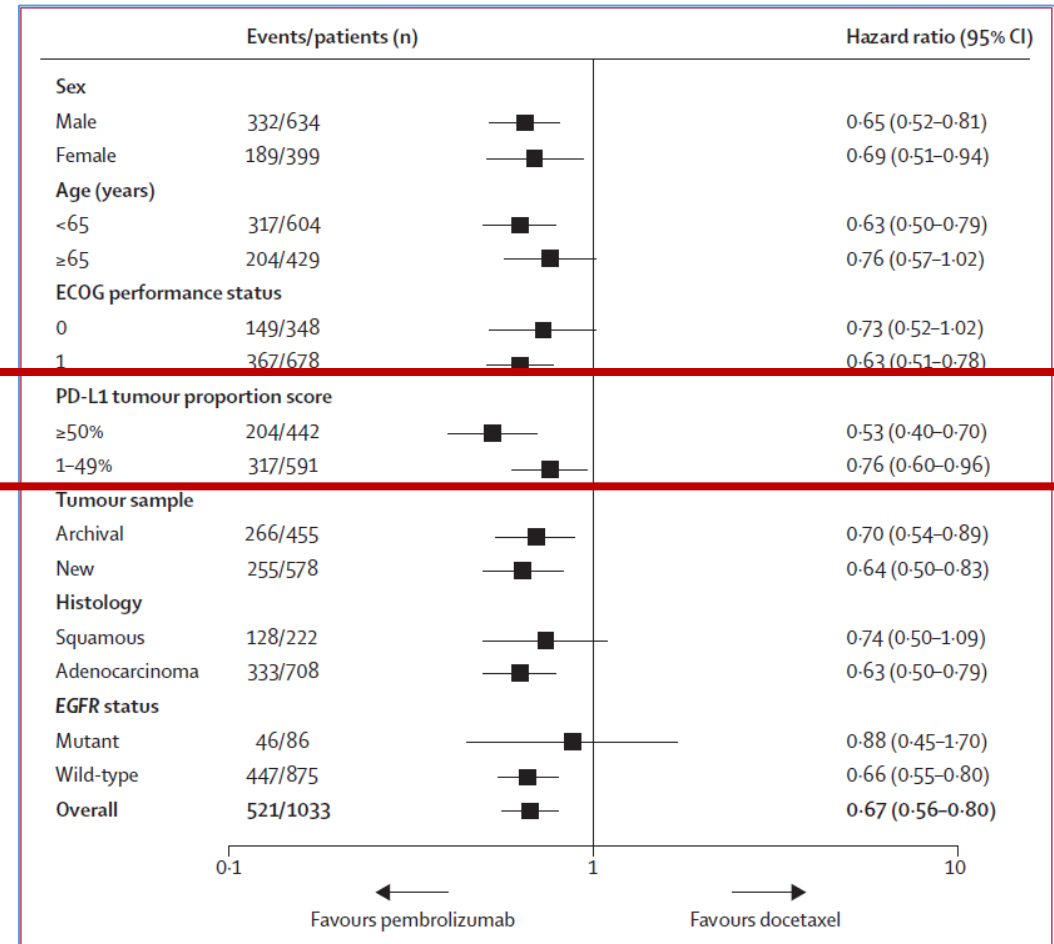


Garon EB et al NEJM 2015

Keynote 010: Pembrolizumab Phase2/3 trial

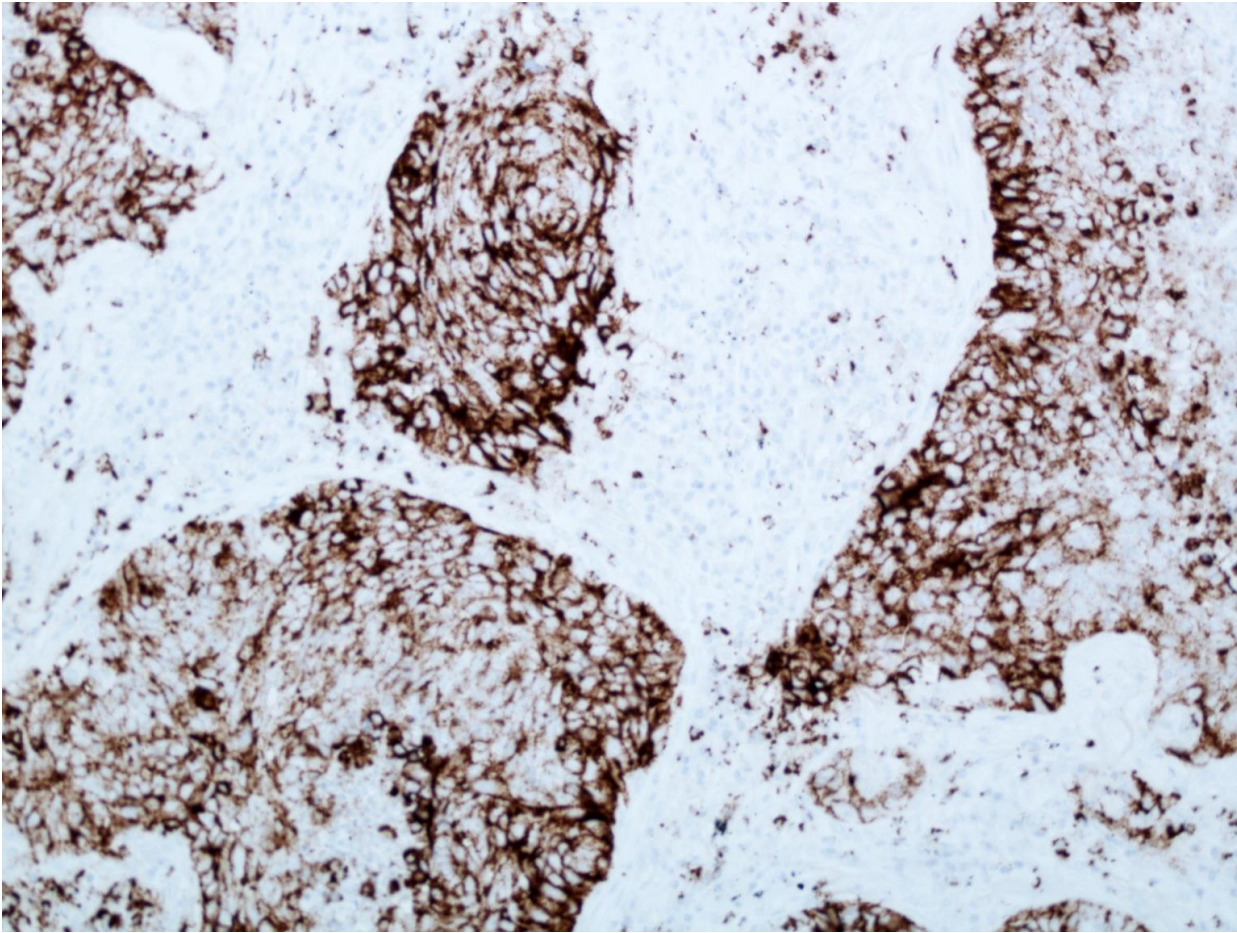


Progression Free Survival



Overall Survival

Atezolizumab & anti-PD-L1 SP142 clone based assay



Tumour cells: % TC positive

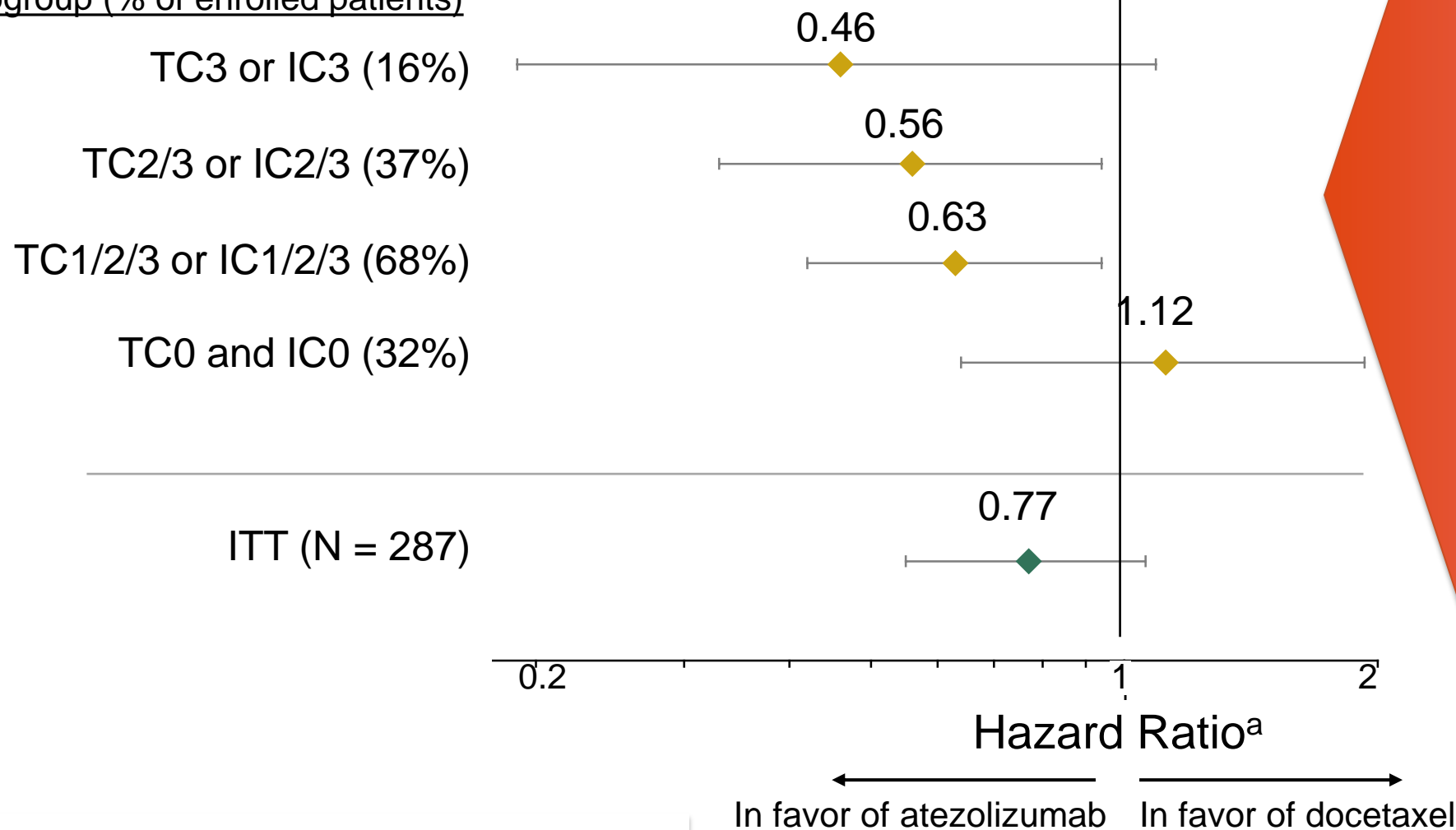


Immune cells: % area of tumour infiltrated

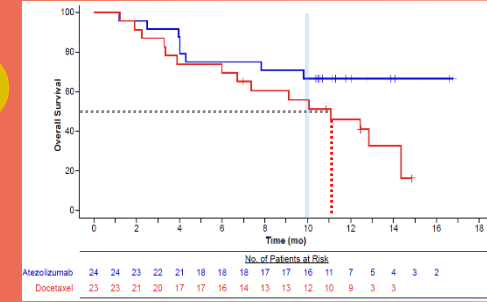
POPLAR: PD-L1 Expression Subgroups

Interim OS Atezolizumab >1L

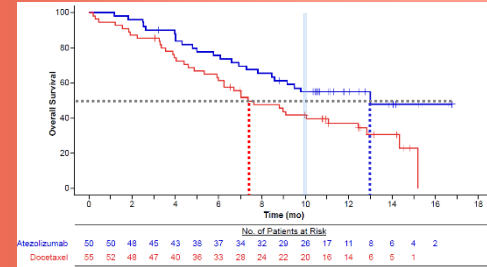
Subgroup (% of enrolled patients)



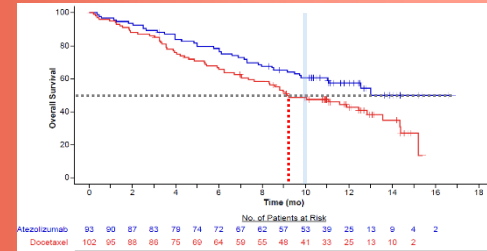
TC3 IC3



TC2/3 IC2/3



TC1/2/3 IC1/2/3



Spira A et al, ASCO June 2015

Spira A. et al., atezolizumab (MPDL3280A)

PRESENTED AT:

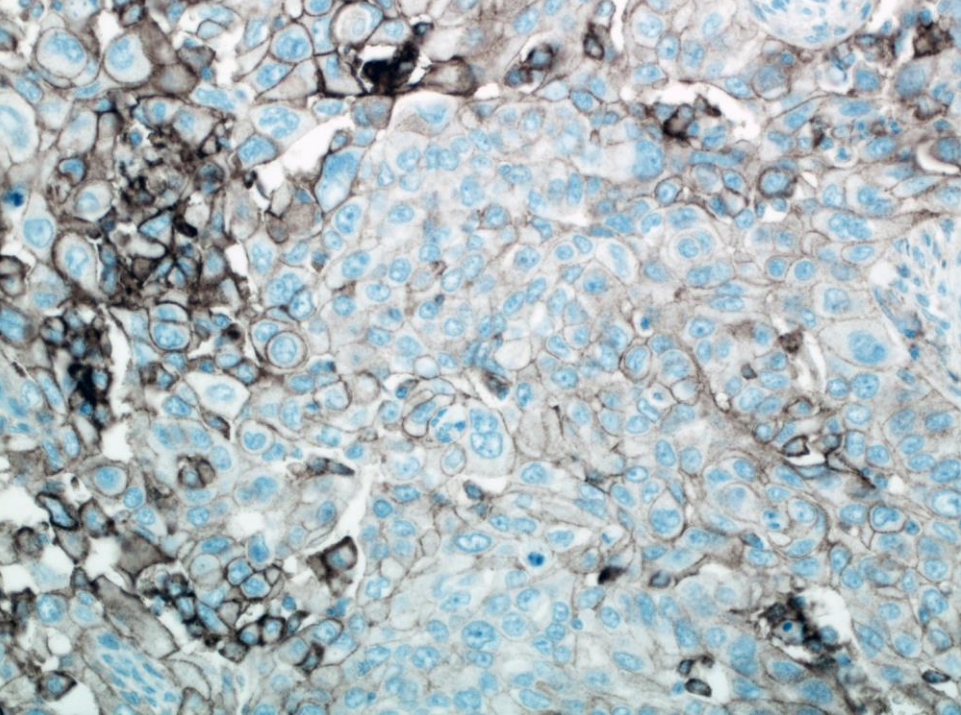
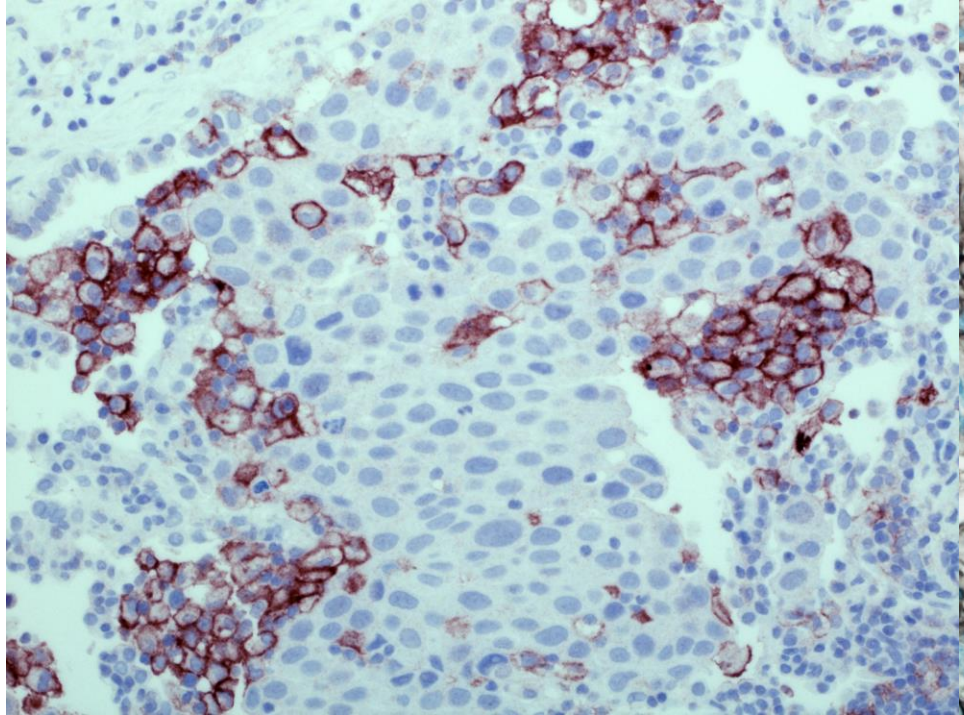
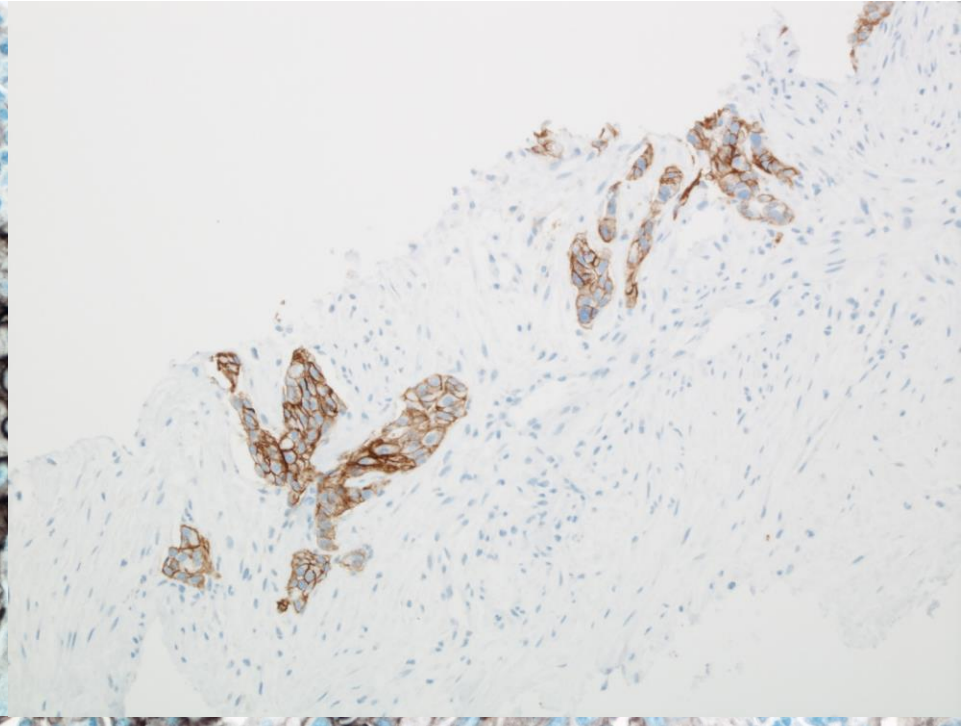
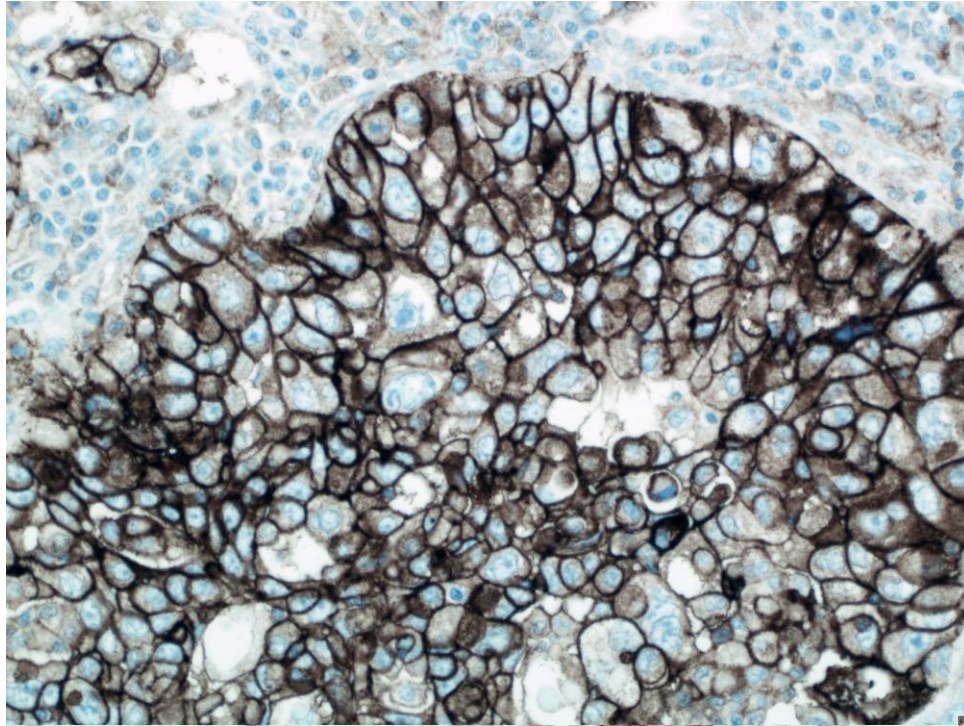
ASCO Annual '15 Meeting

Nivolumab

28-8 clone assay

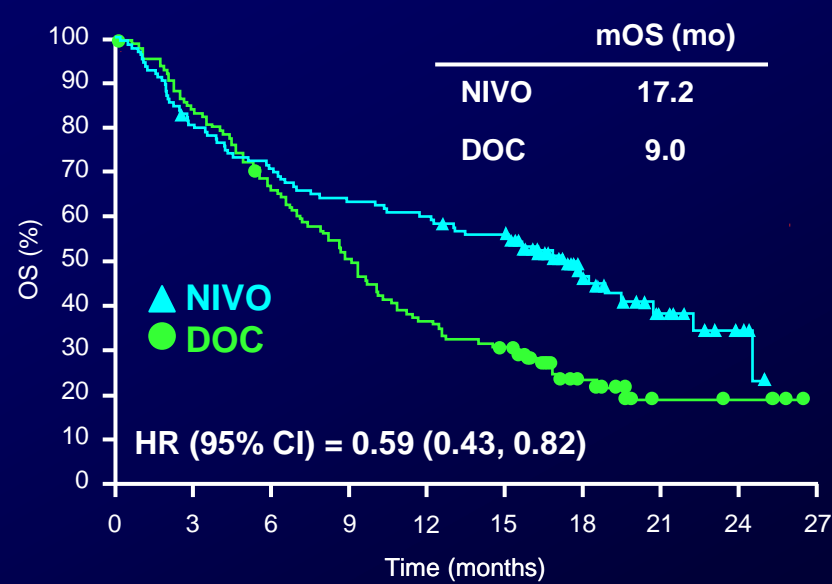
Complimentary or
Companion diagnostic
Assay?

1% threshold is
'positive'

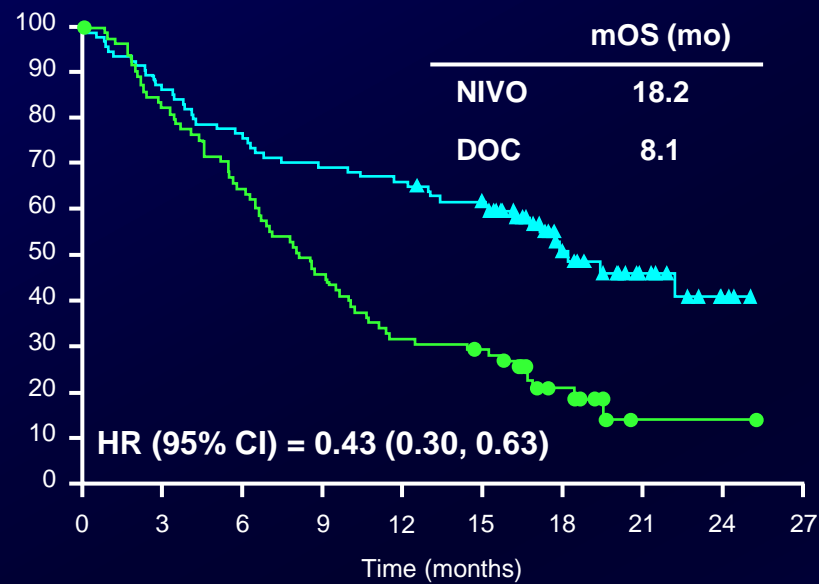


CheckMate 057: Non-Squamous - OS by PD-L1 Expression

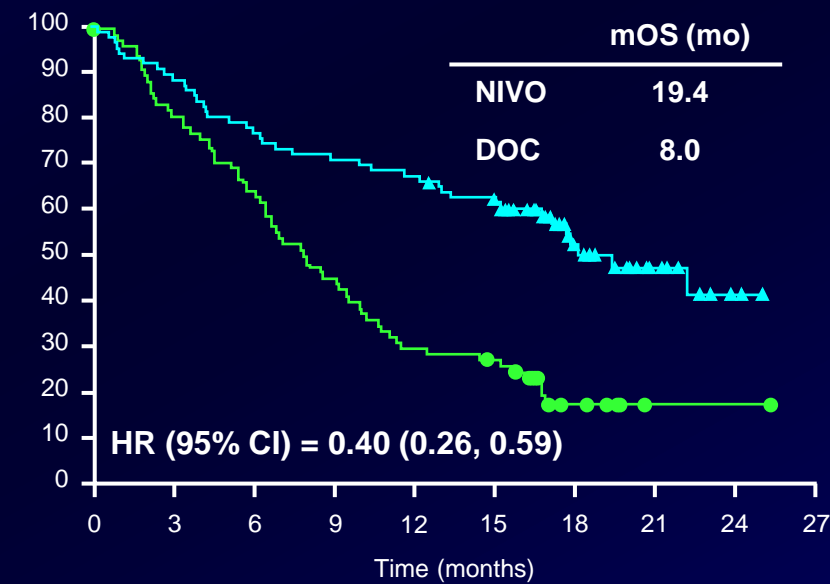
≥1% PD-L1 expression level



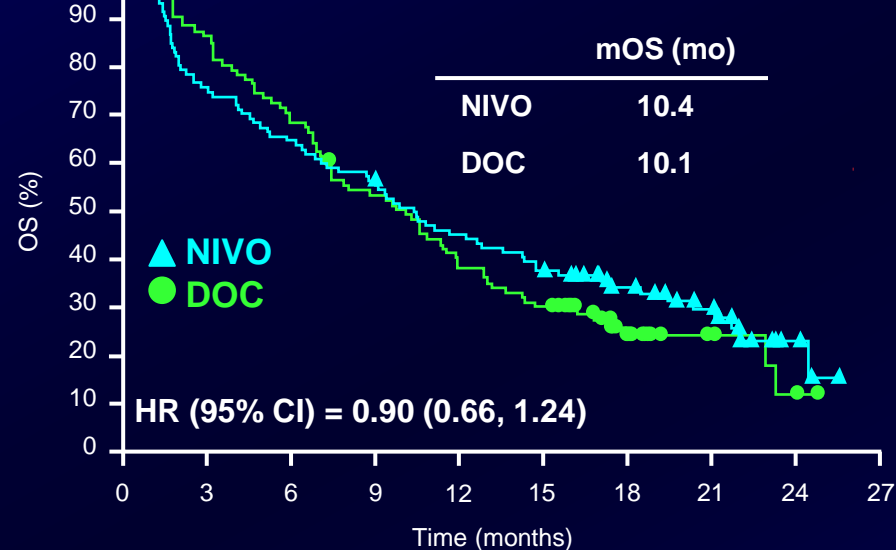
≥5% PD-L1 expression level



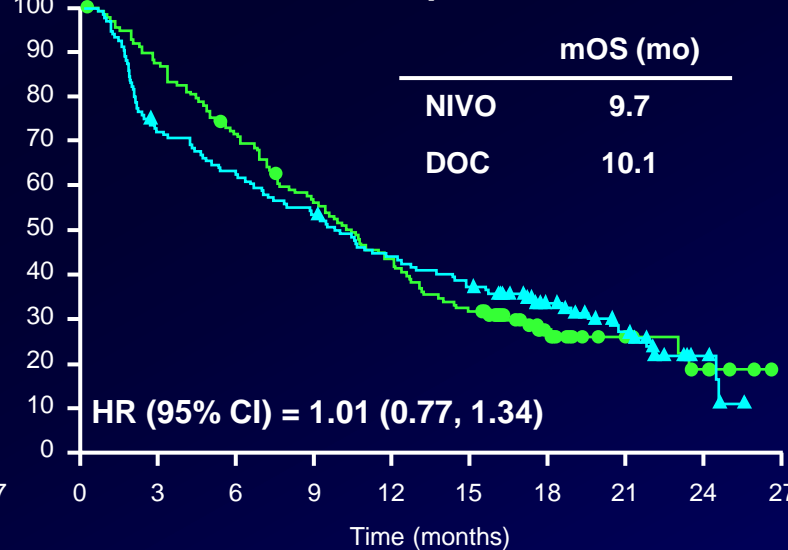
≥10% PD-L1 expression level



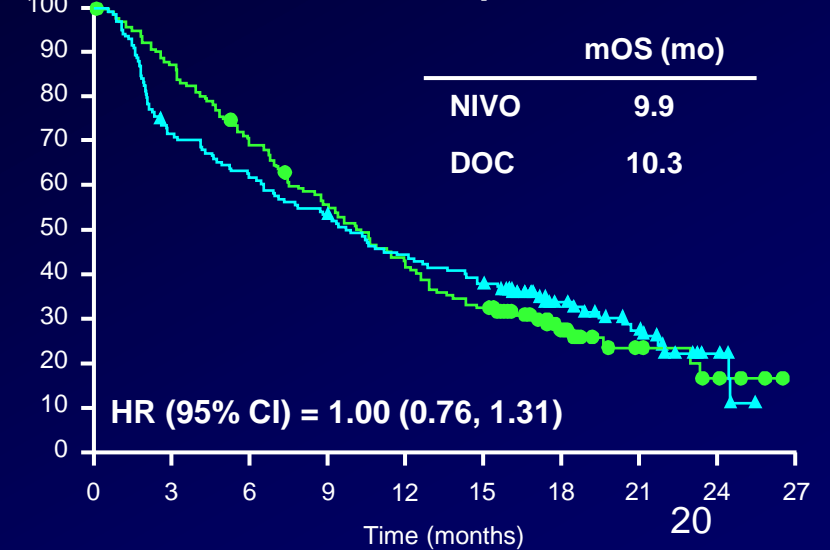
<1% PD-L1 expression level



<5% PD-L1 expression level



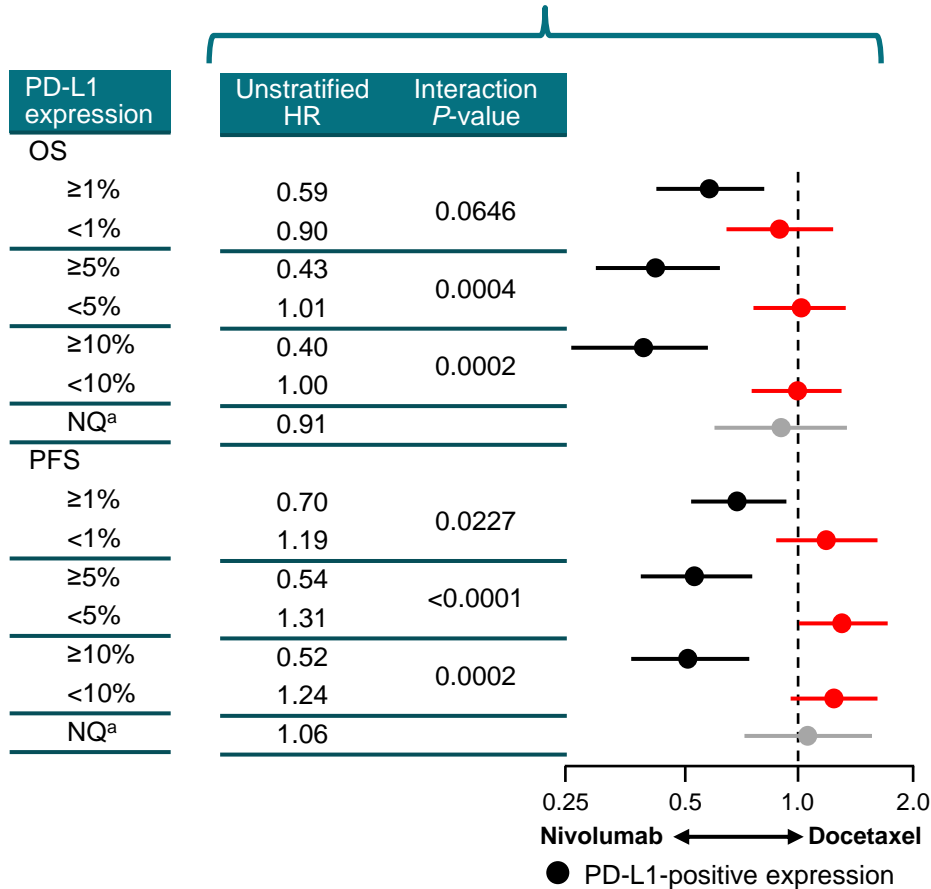
<10% PD-L1 expression level



PD-L1 Expression and Outcome in Nivolumab Phase 3 Trials

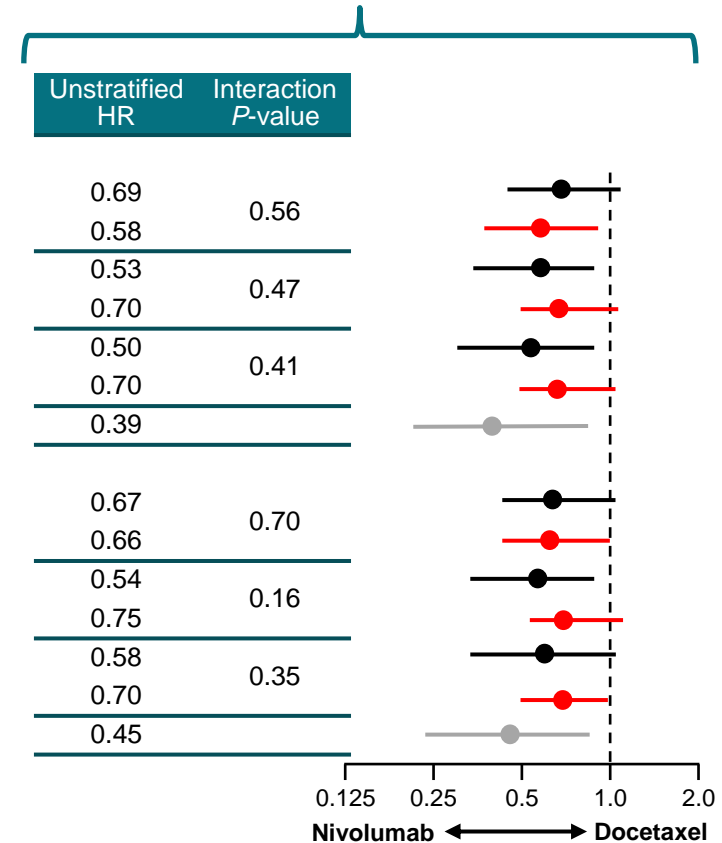
Non-squamous NSCLC (CheckMate 057)³

PREDICTIVE



Squamous NSCLC (CheckMate 017)^{1,2}

NOT PREDICTIVE

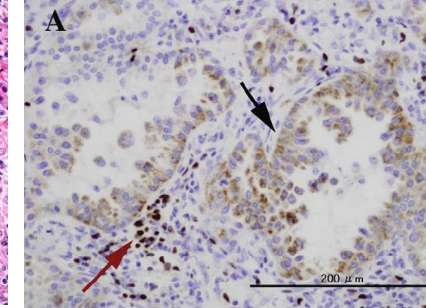
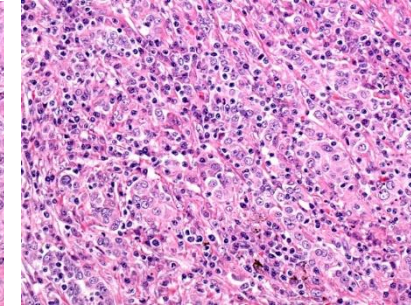
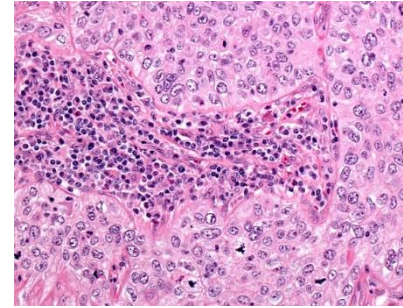


Alternative Potential Biomarkers for Response?

- **Immune gene signatures**
- **Immune cells**
 - Overall infiltrate
 - Specific cell types
- **Other Immune checkpoints**
 - PD-L2, IDO, etc
- **Mutational Burden**

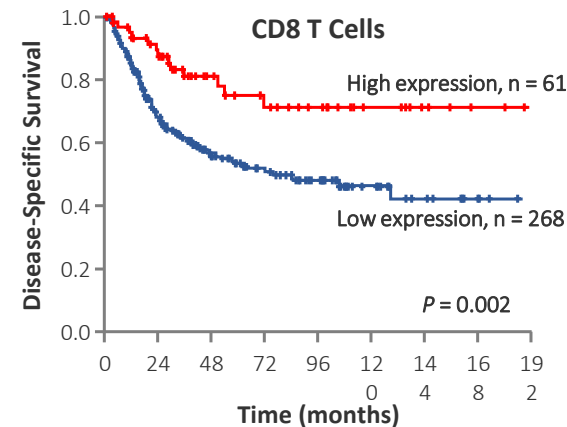
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 - PD-L2, IDO, etc
- Mutational Burden

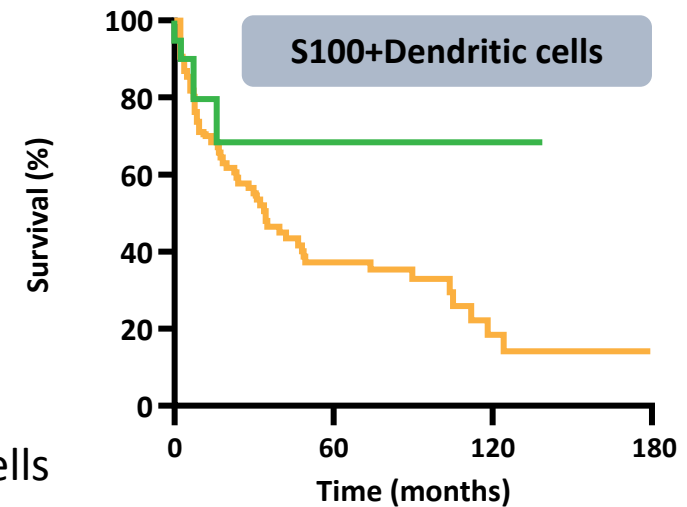


Where is the infiltrate?

FoxP3+ Tregs



Or CD4+ or CD56+ lymphoid cells



Alternative Potential Biomarkers for Response?

- **Immune gene signatures**
- **Immune cells**
 - Overall infiltrate
 - Specific cell types
- **Other Immune regulators**
 - PD-L2, IDO, LAG3,
 - Interferon gamma
- **Mutational Burden**

Anti-PD-L1 therapy – Durvalumab

PD-L1 protein AND
interferon gamma mRNA expression

Higher RR (46%) in combined expression versus
Interferon gamma (33%) or PL-L1 (27%) alone

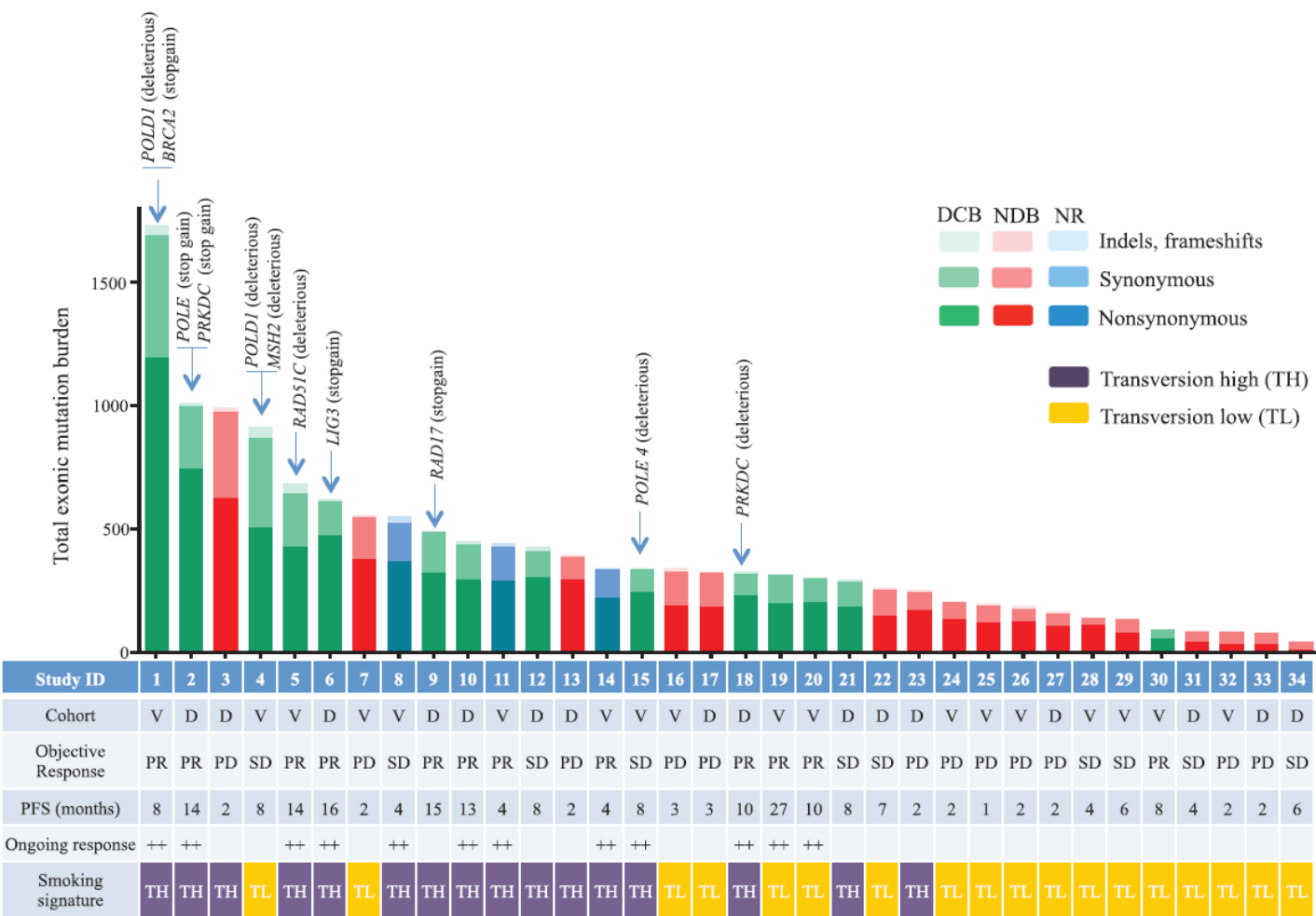
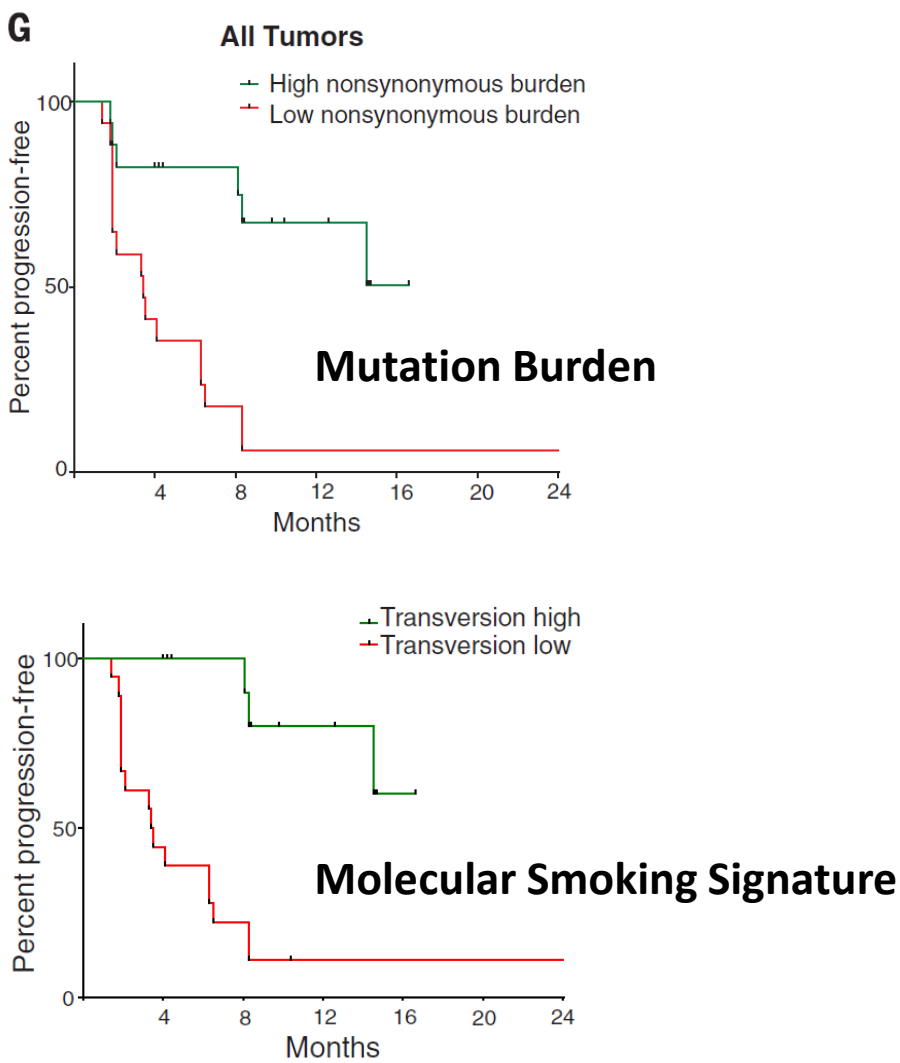
Higgs et al, 15LBA, ECCO 2015

Anti-PD-1 therapy – Pembrolizumab

PD-L1 & PD-L2 IHC better than either alone

Crowley et al. LBA, ECCO 2015

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

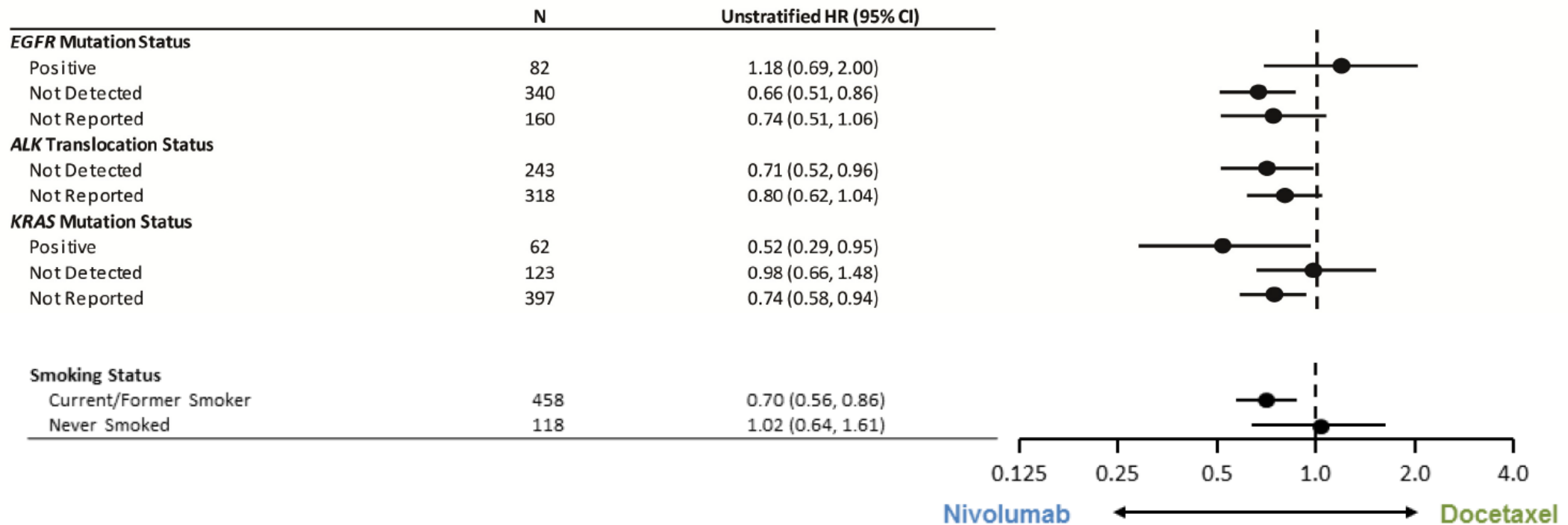


Other factors related to genomic instability

- Polymerase E (POLE) mutations
- Mismatch repair genes (MMR)
- Microsatellite Instability (MSI)

- Smoking signatures in mutations
- Smoking

Smoking and Mutational status: Checkmate 057 – Nivolumab – non-squamous tumours



Biomarker 'positivity': present or absent?

Your tumour is 'negative'

Addictive oncogenic mutation or fusion gene is ABSENT

You will not benefit from therapy

Your tumour is 'positive'

Addictive oncogenic mutation or fusion gene is PRESENT

You will benefit from therapy

Biomarker 'positivity': present, absent or graduated?

Your tumour is 'negative'
Addictive oncogenic mutation or fusion gene is ABSENT
You will not benefit from therapy

Your tumour is 'positive'
Addictive oncogenic mutation or fusion gene is PRESENT
You will benefit from therapy

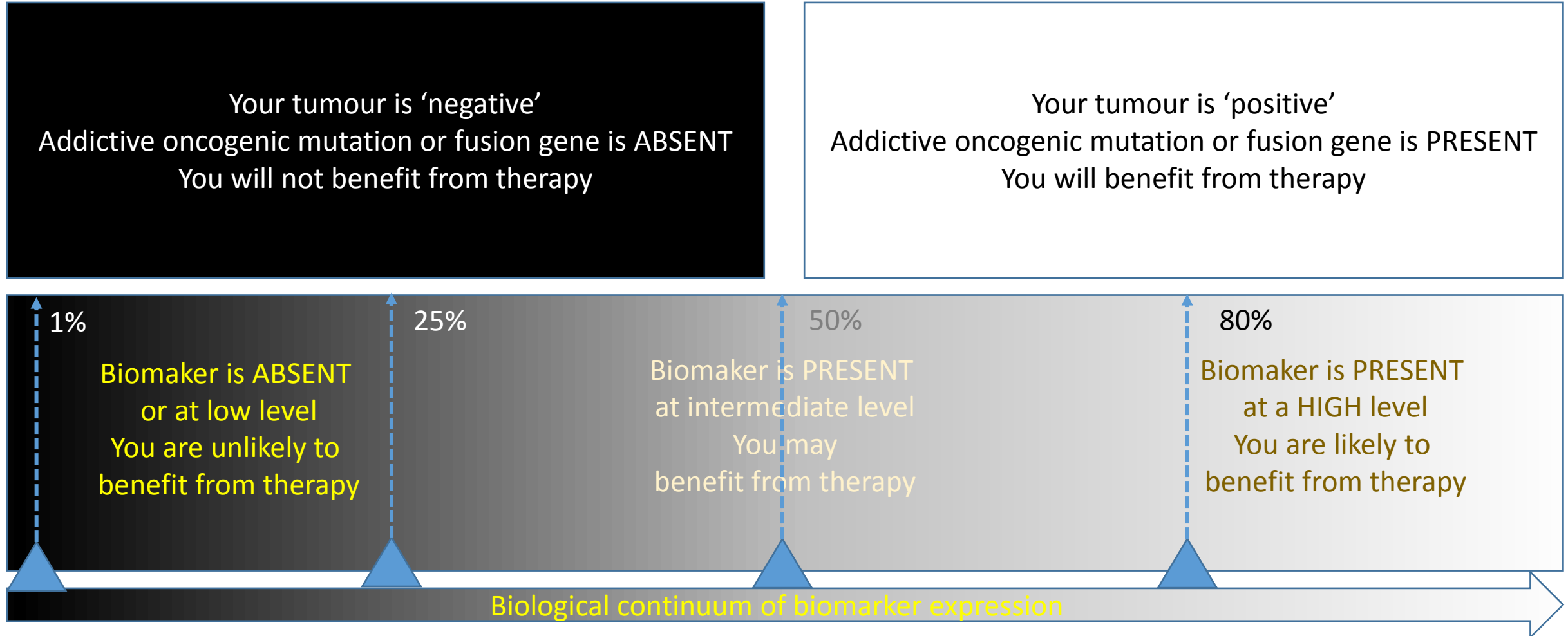
Biomarker is ABSENT
or at low level
You are unlikely to
benefit from therapy

Biomarker is PRESENT
at intermediate level
You may
benefit from therapy

Biomarker is PRESENT
at a HIGH level
You are likely to
benefit from therapy

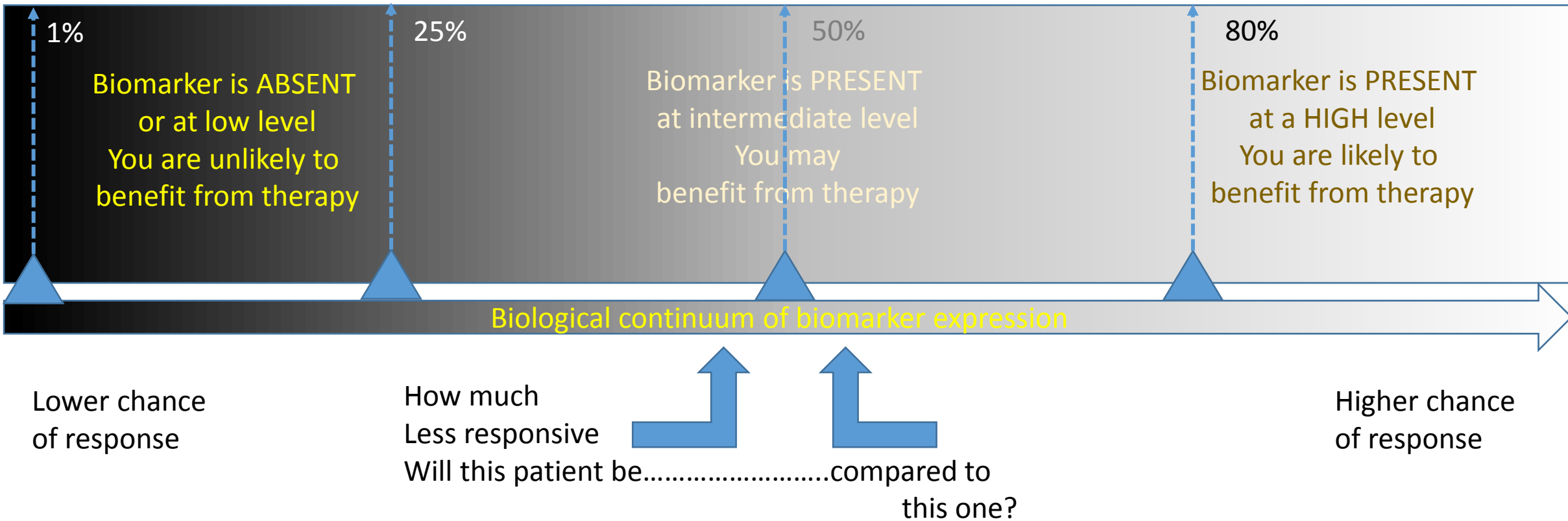
Biological continuum of biomarker expression

Biomarker 'positivity': present, absent or graduated?



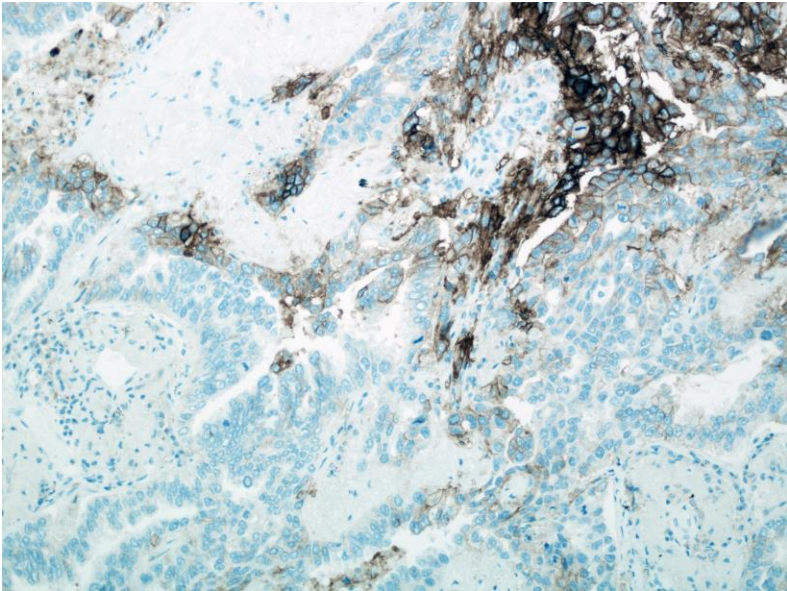
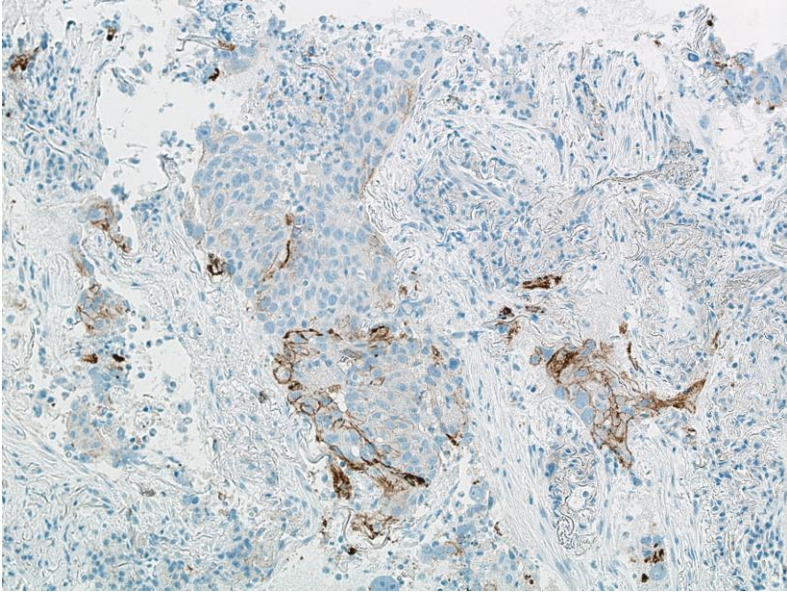
How do we define 'positive'? Where do we set the cut-off value?

Clinical efficacy versus PD-L1 positivity.....



Differential effects depend upon the Dose-response relationship

Heterogeneity and PD-L1



- Expression is dynamic
 - Expression is heterogeneous
 - Sampling 'error' must occur
 - Greater impact at lower thresholds
-
- Part of the reason why the biomarker appears worse than it is
 - Trials 'control' for heterogeneity to some extent

PD-L1 immunohistochemistry as a biomarker

- **Is the drug target a 'singular' factor in the target system?**
- **Is the biomarker present or absent?**
- **Is the biomarker stable and functionally unique?**
- **Is the biomarker easily measured?**
- **Is the biomarker 100% predictive?**

- **But it is all we currently have and what chance anything else is better?**

It depends how you look at this.
What 'odds' of success are you willing to accept?



PD-L1 as a predictive immune biomarker: assays, sample collection and analysis in NSCLC studies

		Pembrolizumab Merck	Nivolumab Bristol-Myers Squibb	Atezolizumab Roche/Genentech	Durvalumab AstraZeneca	Avelumab Pfizer/Merck Serono
Sample Source and Collection	PD-L1 Assay	<ul style="list-style-type: none"> Prototype or clinical trial IHC assay (22C3 Ab) 	<ul style="list-style-type: none"> Dako automated IHC assay (28-8 Ab) 	<ul style="list-style-type: none"> Central laboratory IHC assay Ventana PD-L1 (SP142) 	<ul style="list-style-type: none"> Ventana automated IHC (BenchMark ULTRA using Ventana PD-L1 (SP263) clone) 	<ul style="list-style-type: none"> Dako assay Clone not known
		<ul style="list-style-type: none"> Surface expression of PD-L1 on tumour specimen 	<ul style="list-style-type: none"> Surface expression of PD-L1 on tumour cells 	<ul style="list-style-type: none"> Surface expression of PD-L1 on TILs or tumour cells 	<ul style="list-style-type: none"> Surface expression of PD-L1 on tumour cells 	<ul style="list-style-type: none"> Surface expression of PD-L1 on tumour cells
		<ul style="list-style-type: none"> Ph I: Fresh or archival tissue 	<ul style="list-style-type: none"> Archival or fresh tissue 	<ul style="list-style-type: none"> Archival or fresh tissue 	<ul style="list-style-type: none"> Unknown 	<ul style="list-style-type: none"> Unknown
	Definition of Positivity [†]	<p>IHC Staining:</p> <ul style="list-style-type: none"> Strong vs weak expression PD-L1 expression required for NSCLC for enrollment <ul style="list-style-type: none"> Note that one arm of KEYNOTE 001 trial requires PD-L1⁺ tumours <p>Tumour PD-L1 expression:</p> <ul style="list-style-type: none"> ≥50% PD-L1⁺ cut-off: 32% (41/129) 1–49% PD-L1⁺ cut-off: 36% (46/129) 	<p>IHC Staining:</p> <ul style="list-style-type: none"> Strong vs weak expression Patients not restricted by PD-L1 status in 2nd- & 3rd-line Ph III 1st-line trial in PD-L1⁺ <p>Tumour PD-L1 expression:</p> <ul style="list-style-type: none"> 1% PD-L1 + cut off 5% PD-L1⁺ cut-off: 59% (10/17) 5% PD-L1⁺ cut-off: 49% (33/68) 10% PD-L1 + cut off 	<p>IHC Staining Intensity (TC: 0, 1, 2, 3):</p> <ul style="list-style-type: none"> IHC 3 (≥50% PD-L1⁺) IHC 2,3 (≥5% PD-L1⁺) IHC 1,2,3 (≥1% PD-L1⁺) IHC 0,1,2,3 (all patients with evaluable status)^{6,7} PD-L1 expression required for NSCLC for enrolment in Ph II trials <p>IC: TIL PD-L1 expression:</p> <ul style="list-style-type: none"> IHC 3 (≥10% PD-L1⁺): 11% (6/53) PD-L1 low (IHC 1, 0): 62% (33/53) 	<p>IHC Staining Intensity:</p> <ul style="list-style-type: none"> Not presented to date <p>Tumour PD-L1 expression:</p> <ul style="list-style-type: none"> PD-L1 + cut off 25% PD-L1⁺: 34% (20/58) PD-L1⁻: 50% (29/58) 	<p>IHC Staining Intensity:</p> <ul style="list-style-type: none"> Not presented to date <p>Tumour PD-L1 expression (all doses):</p> <ul style="list-style-type: none"> PD-L1 + cut off 1% PD-L1⁺: 34% (20/58) PD-L1⁻: 50% (29/58)

PD-L1 IHC: what chance one test?

- Is all anti-PD-L1 IHC the same?
- Are all the Companion Diagnostics the same?
- **Can we use any IHC for any drug?**
 - No evidence to support this practice
 - One IHC – multiple scoring definitions
 - Implications for how PD-L1 IHC would be reported by pathologists
 - % cells positive
 - Indicate different thresholds?
 - Mention actual drugs in report?
- **How far is it safe to deviate from trial-validated practice?**

Immunotherapy and Biomarkers

- Biologically rational therapeutic approach
- Biomarkers based upon a putative understanding of
 - Likelihood of antigenicity and therefore an immune response
 - Evidence of an immune response
 - Evidence of an inhibitory mechanism
 - Evidence of the specific target – PD-L1
- PD-L1 is a realistic biomarker
 - Nature of this biomarker presents issues
 - Complex environment with multiple drugs and assays