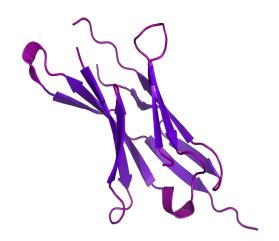


## Immune monitoring in cancer patients following treatment with anti-PD-1 antibody

NSCLC patients not responding to nivolumab show lowered frequency of co-stimulatory receptor-deficient CD8 T cells

#### PD-1, programmed cell death protein 1, aka CD279

- Cell surface receptor
- Expressed mainly on T cells and is induced by T cell activation
- Two ligands: PD-L1 and PD-L2
- Down-regulates T cell activation
- Acts as checkpoint to guard against autoimmunity



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# Cancer immunologists scoop medicine Nobel prize

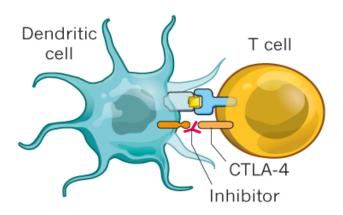
One of the hottest areas in cancer research, immunotherapy can dramatically extend lives

## IMMUNE BOOST

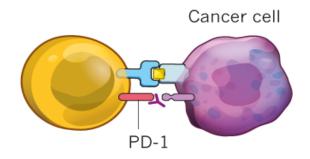
Several methods are showing promise in helping immune sentinels called T cells to attack cancer.

#### CHECKPOINT INHIBITOR DRUGS

'Checkpoint' proteins block T-cell activity. Inhibitor drugs can release the brakes on T cells at different stages.



The CTLA-4 checkpoint protein prevents dendritic cells from priming T cells to recognize tumours. Inhibitor drugs block the checkpoint.



The PD-1 checkpoint protein prevents T cells from attacking cancer cells. The inhibitor drug allows T cells to act.

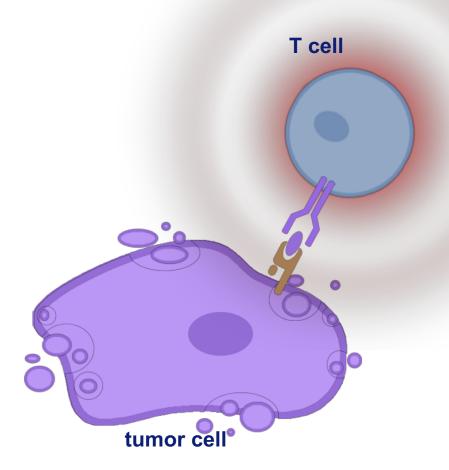
onature



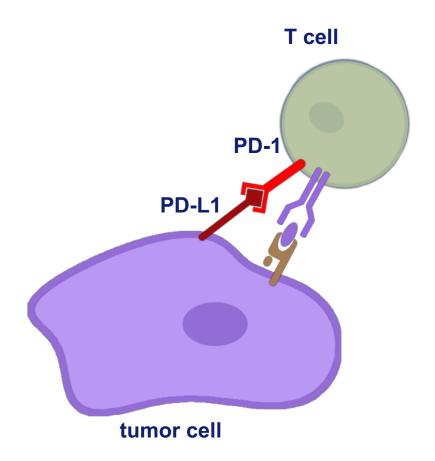


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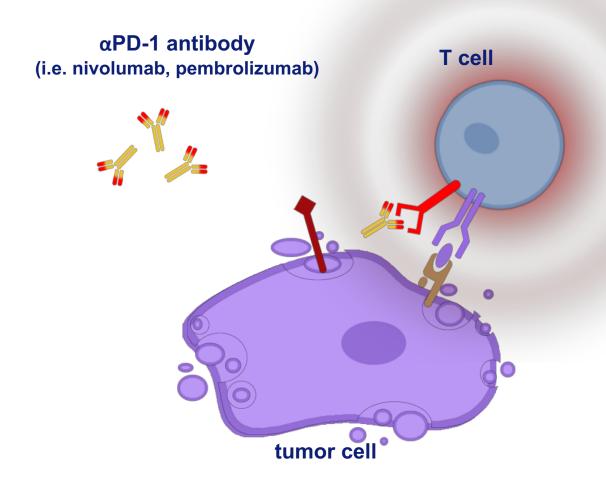
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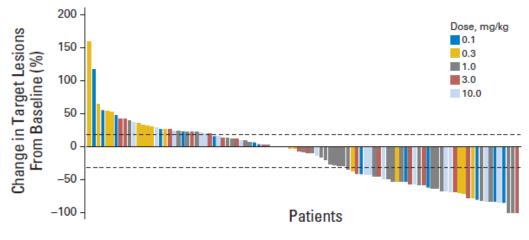


αPD-L1 antibody (i.e. Atezolizumab, Durvalumab, Avelumab)

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#### Nivolumab – first clinical data in advanced melanoma

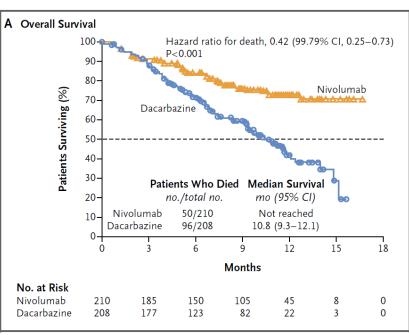
Survival, Durable Tumor Remission, and Long-Term Safety in Patients With Advanced Melanoma Receiving Nivolumab



Overall response rate after ipilimumab\*: 31%

Topalian et al. JCO, 2014

Melanoma without BRAF Mutation



Nivolumab in Previously Untreated

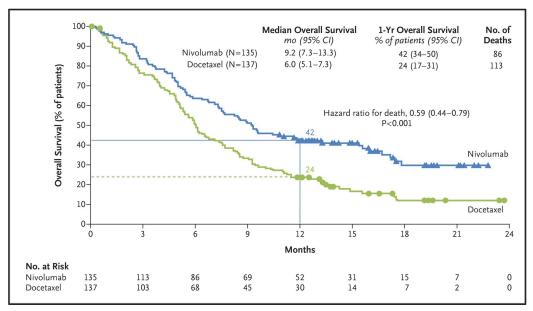
Overall response rate as frontline therapy 40%

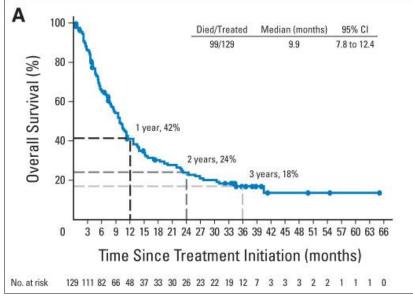
Robert et al. NEJM, 2015

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<sup>\*</sup> anti-CTLA4 mAb (BMS)

#### **Anti-PD-1 Immune Therapy in NSCLC – Promises and Challenges**





Brahmer, N Engl J Med, 2015

Gettinger, J Clin Oncol, 2015

- increased overall survival compared to standard treatment
- approved by FDA and EU in cancer types of multiple origins



## FDA and EU-approved Immune Checkpoint Inhibitors (ICI) Available in The Netherlands

Anti-CTLA4 Yervoy® Ipilimumab (Bristol-Myers Squibb)

Anti-PD-1 Opdivo® Nivolumab (Bristol-Myers Squibb)

**Keytruda**® **Pembrolizumab** (Merck)

Anti-PDL-1 Tecentriq® Atezolizumab (Roche)

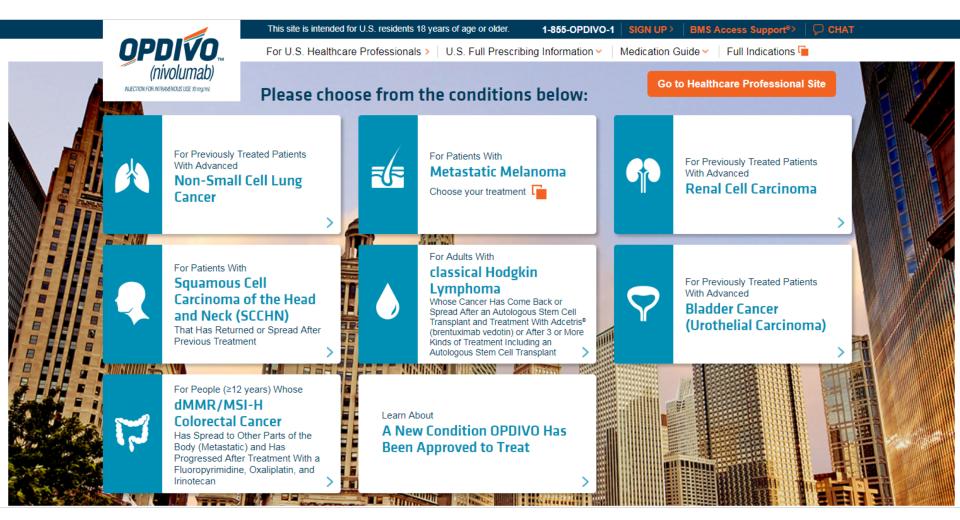
Imfinzi® Durvalumab (Astra Zeneca)

Bavencio® Avelumab (Merck Pfizer)

These drugs are increasingly gaining first-line indications

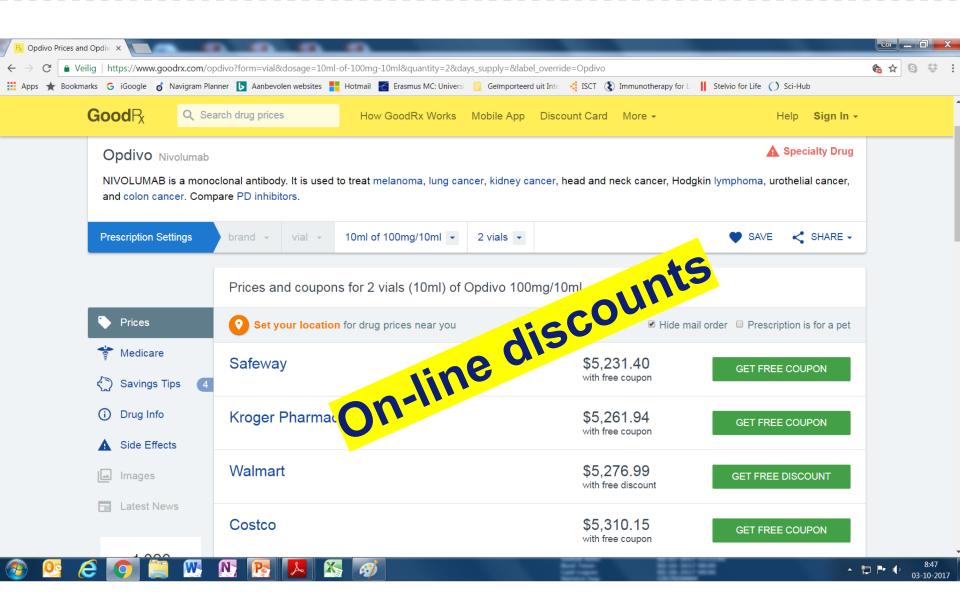


#### PD-1 inhibitor therapy has become big business



Bristol-Myers Squibb website





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#### **Current challenges of PD-1 treatment**

- ➤ Many patients do not demonstrate clinical benefit
- ➤ Many patients show clinical toxicities
- Costs of patient treatment are high

#### **Hypothesis**

Frequencies of immune cells and their subset distributions in blood predict response to PD-1 treatments and facilitate treatment stratification



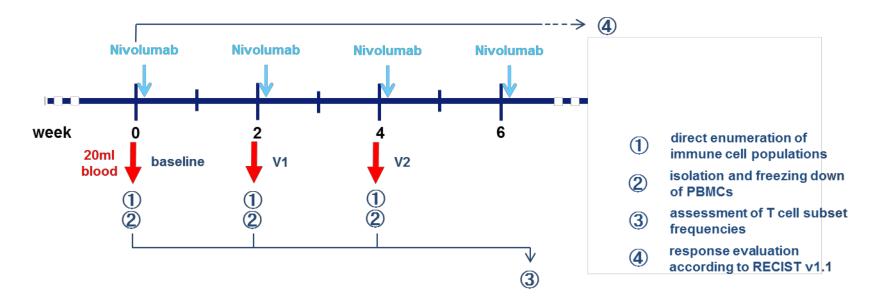
## Blood Parameters are Associated with Clinical Response to Immune Checkpoint Inhibitors

Marker	ICI therapy	Cancer	N	Study results	Reference
Lymphocyte count	Ipilimumab Ipilimumab	Melanoma Melanoma	51, 73 82, 40	$\geq$ 1000 per $\mu$ l at week 6 → ↑ OS ↑ At 2–8 weeks $vs$ baseline → ↑ response	(Delyon et al, 2013; Ku et al, 2010) (Bjoern et al, 2016; Martens et al, 2016b)
	Ipilimumab <mark>Nivolumab</mark>	Melanoma Melanoma	95 98	↑ At week 12 vs baseline $\rightarrow$ ↑ OS $\geqslant$ 1000 per $\mu$ l at week 3-6 $\rightarrow$ ↑ OS	(Simeone et al, 2014) (Nakamura et al, 2016)
Relative lymphocyte count	Ipilimumab Pembrolizumab	Melanoma Melanoma	209 <mark>616</mark>	↑ Baseline → ↑ OS ↑ Baseline → ↑ OS	(Martens et al, 2016a) (Weide et al, 2016)
Total leucocyte count	Ipilimumab	Melanoma	59	↓ Baseline → ↑ response	(Gebhardt et al, 2015)
Eosinophil count	Ipilimumab Ipilimumab Ipilimumab	Melanoma Melanoma Melanoma	209 59 73	↑ Baseline → ↑ OS     ↑ At week 3 vs baseline → ↑ response     ↑ At week 6 vs baseline → ↑ OS	(Martens et al, 2016a) (Gebhardt et al, 2015) (Delyon et al, 2013)
Relative eosinophil count	Pembrolizumab	Melanoma	616	↑ Baseline → ↑ OS	(Weide et al, 2016)
Neutrophil count	Ipilimumab Ipilimumab Nivolumab	Melanoma Melanoma Melanoma	59 720 <mark>98</mark>	↓ Baseline → ↑ response ↓ Baseline → ↑ PFS and OS < 4000 per $\mu$ l at week 3–6 → ↑ OS	(Gebhardt et al, 2015) (Ferrucci et al, 2016) (Nakamura et al, 2016)
Neutrophil/lymphocyte ratio	Ipilimumab Ipilimumab Nivolumab	Melanoma Melanoma NSCLC	58, 185 187 <mark>175</mark>	<ul> <li>↓ Baseline → ↑ OS</li> <li>↓ Baseline → ↑ PFS and OS</li> <li>↓ Baseline → ↑ OS</li> </ul>	(Khoja et al, 2016; Zaragoza et al, 2016) (Ferrucci et al, 2015) (Bagley et al, 2017)
Derived neutrophil/ lymphocyte ratio	Ipilimumab	Melanoma	720	↓ Baseline → ↑ PFS and OS	(Ferrucci et al, 2016)
Monocyte count	Ipilimumab	Melanoma	209	↓ Baseline → ↑ OS	(Martens et al, 2016a)



#### **MULTOMAB**

(prospective saMpling in intravenoUsLy Treated Oncology patients: Monoclonal AntiBodies )



#### **Collaborations:**

Translational Pharmacology (group Mathijssen)
Pulmonary Diseases (group Aerts)

#### Patient numbers per October 2018:

Melanoma: >150 NSCLC: >250

Also other tumor types

#### **Efforts:**

- Collection of patient blood pre- and post-treatment
- ➤ Processing/storage of whole blood, serum, PBMC, DNA/RNA
- Measurement of antibody levels in sera (pharmacokinetics)
- Measurement of immune cells in whole blood /PBMC (immune profiling)
- > Clinical patient evaluation (tumor burden, toxicity, response)

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#### Immune monitoring – methods applied

A) Blood processing

1. Immediate Analysis on Whole Blood (<24h)

2. Isolation and Cryopreservation of PBMC

3. Storage of Plasma

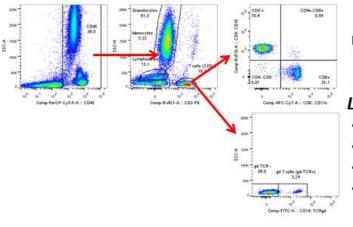
4. Storage of Whole blood DNA/RNA

B) Multiplex flow cytometry (12colors)

FACS Celesta

- > Enumeration of 18 immune cell subsets
- > Assess expression of T cell markers for:
  - Maturation
  - Co-stimulation
  - Co-inhibition
  - Chemokines
  - Total of 300 combinations

#### C) Absolute numbers and T cell markers



#### Lymphocytes:

- · B cells
- NK cells
- · T cells
- yδ T cells

#### Granulocytes:

- Eosinophils
- Immature neutrophils
- Mature neutrophiles
- PMN-MDSCs

#### Monocytes:

- Classical monocytes
  - M-MDSCs
- · Intermediate monocytes
- · Non-classical monocytes
- · Dendritic cells

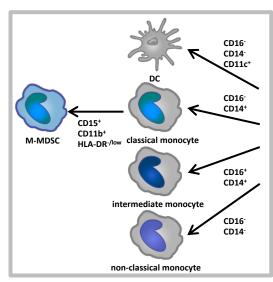
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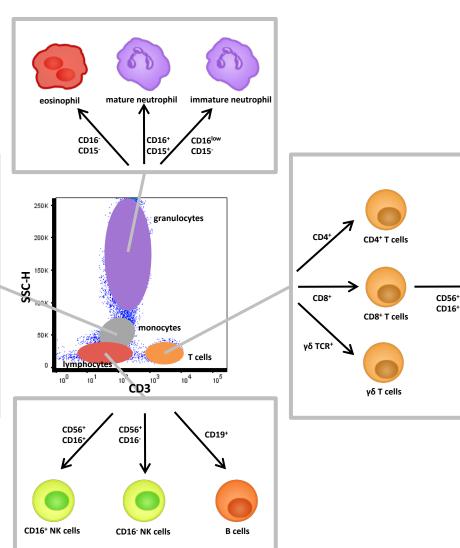
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#### Flow Cytometric Analysis of Blood

#### 12-color flow cytometry

enumeration of 18 immune cell populations







NK T cells

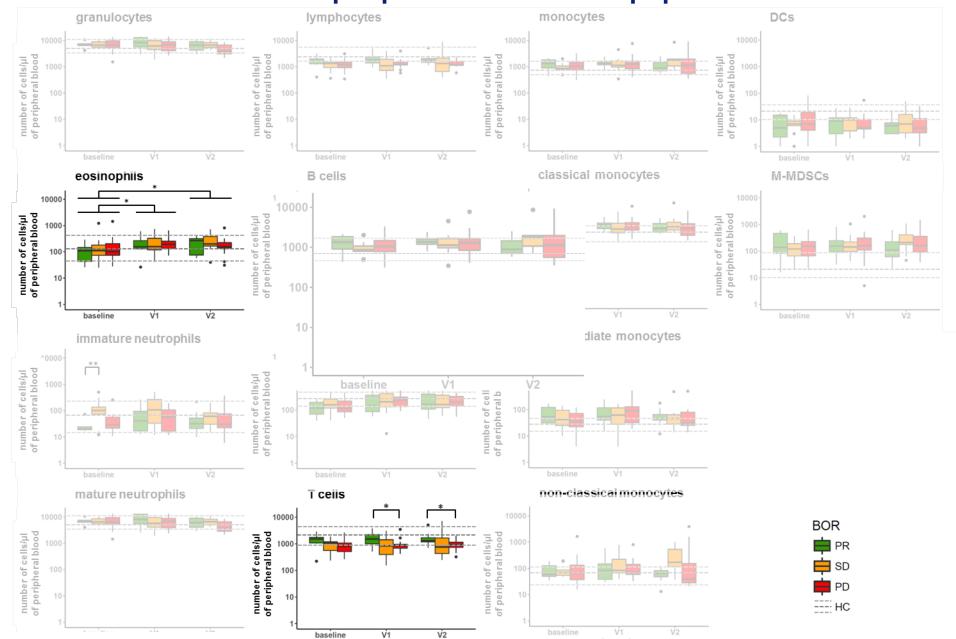
#### **Exploratory Cohort**

Tumor type:		NSCLC
Treatment*:		Nivolumab, Q2W, 3mg/kg
Median age in years (ra	inge):	65 (35-79)
Sex: - female - male		30 (42.3%) 41 (57.7%)
BOR: - progressive dis - stable disease ( - partial respons	SD)	32 (45.1%) 25 (35.2%) 14 (19.7%)
Median follow-up in da	ys (range):	242 (35-544)
WHO performance statu	ıs: 0 1 unknown	16 (22.5%) 37 (52.1%) 18 (25.4%)
Histology of primary lung tumor:	adenocarcinoma squamous cell carcinoma great cell carcinoma	48 (67.6%) 21 (29.6%) 2 (2.8%)

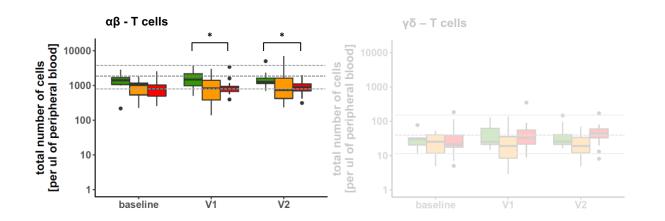
<sup>\*</sup> all patients received platinum-based pre-treatment

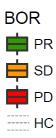


## Nivolumab treatment in general does not result in changed numbers of peripheral immune cell populations



#### Patients responding to nivolumab show high numbers of CD8 T cells





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#### Flow Cytometric Analysis of Blood

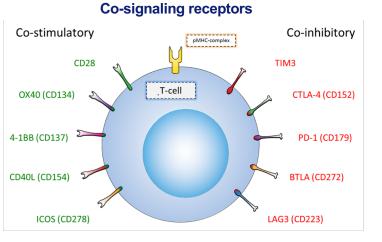
#### 12-color flow cytometry

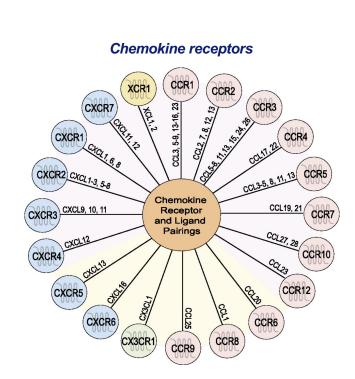
- enumeration of 18 immune cell populations
- assess expression of T cell markers for: maturation, co-stimulation, co-inhibition and chemokines (>300 combinations)

# granulocytes monocytes T cells

CD3

Naive T cell





T<sub>EM</sub> cell

T<sub>Eff</sub> cell

**Differentiation** 

 $T_{CM}$  cell

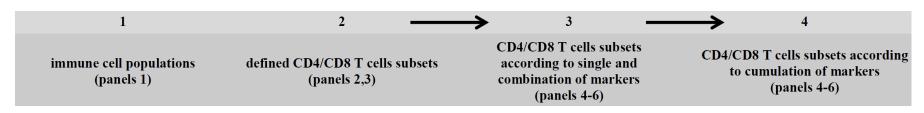
T<sub>SCM</sub> cell

#### Multiplex flow cytometry – panels 2-6

<b>2</b> <sup>b</sup>	T cell proliferation/ regulatory T cell markers	Ki67, CD25, FOXP3, PD-1
<b>3</b> <sup>b</sup>	T cell maturation markers	CCR7, CD45RA, CD95, CD69, CD27, CD103
<b>4</b> <sup>b</sup>	T cell co-inhibitory receptors	CD57, LAG3, BTLA, PD-1, TIM3
<b>5</b> <sup>b</sup>	T cell co-stimulatory receptors	CD28, OX40, 4-1BB, CD40L, ICOS
<b>6</b> <sup>b</sup>	T cell chemoattractant receptors	CXCR3, CXCR4, CCR1, CCR4, CCR5

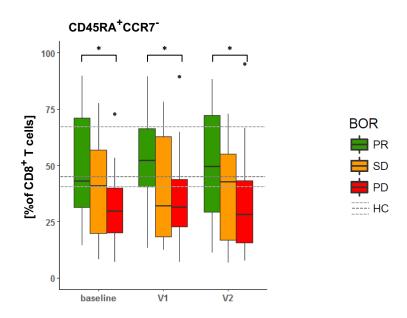
<sup>&</sup>lt;sup>b</sup> assessment of T cell subset frequencies in PBMC samples

#### Multiplex flow cytometry – analysis work scheme



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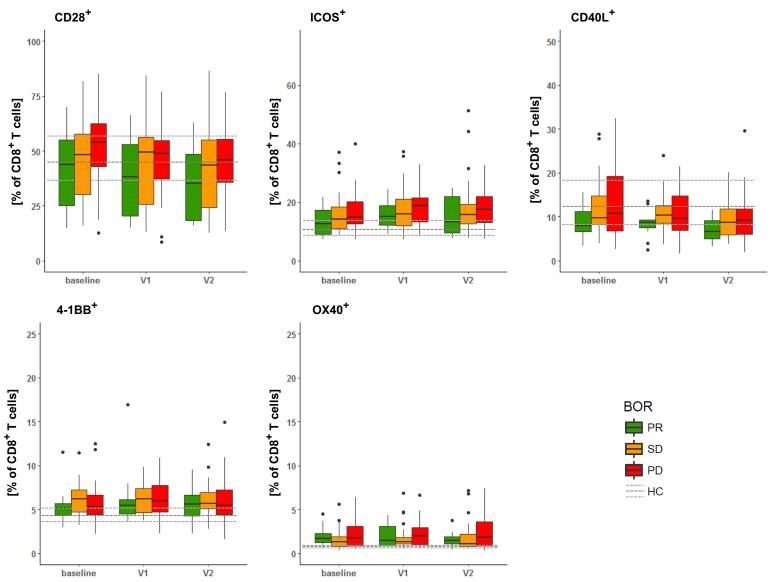
## Patients with PR show enhanced frequencies of CD8 T cells with CD45RA+CCR7- and CD95+CD69- phenotypes



These observations suggest a role for differentiated/tissue-egressed (possibly antigen experienced) CD8 T cells

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#### Individual co-signaling receptors do not predict response



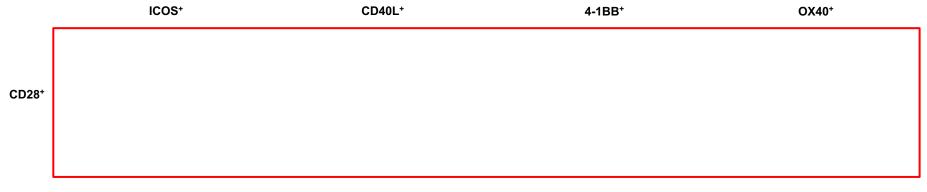
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OX40<sup>+</sup>

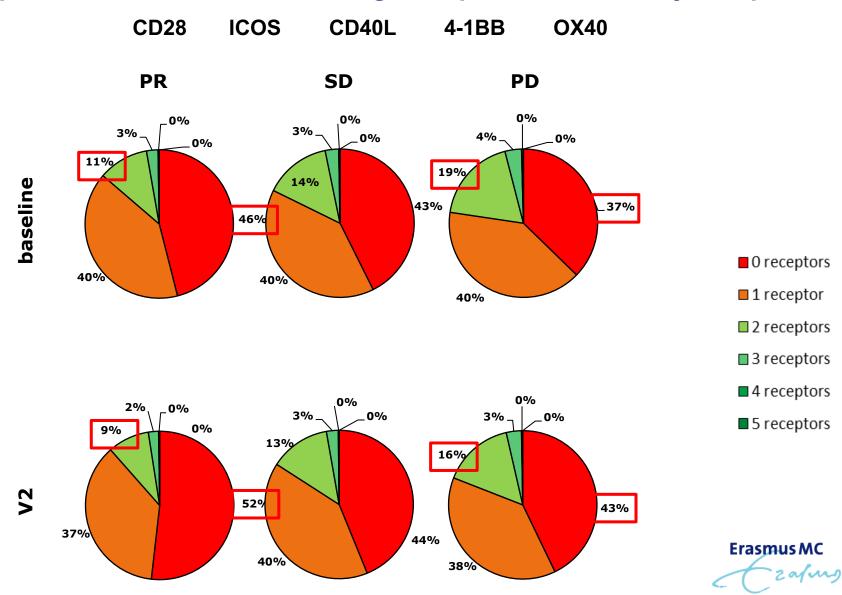
4-1BB+

CD40L+

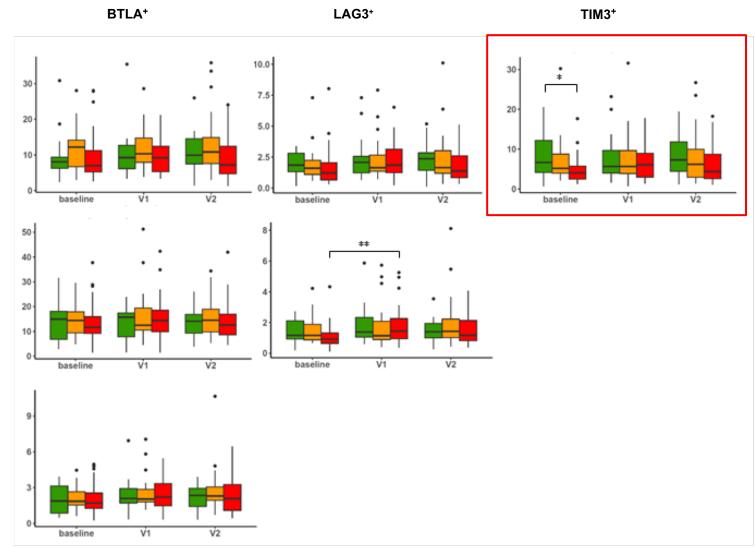
## Patients with PR display reduced frequencies of CD8 T cells co-expressing CD28 and CD40L, ICOS or ICOS



## Patients showing response to therapy are characterized by higher frequencies of CD8 T cells lacking multiple co-stimulatory receptors



## Patients with PR display enhanced frequencies of CD8 T cells co-expressing PD-1 and TIM-3



Cells double-positive for indicated receptors [% +CD8+ T cells]

PD-1+

TIM3+

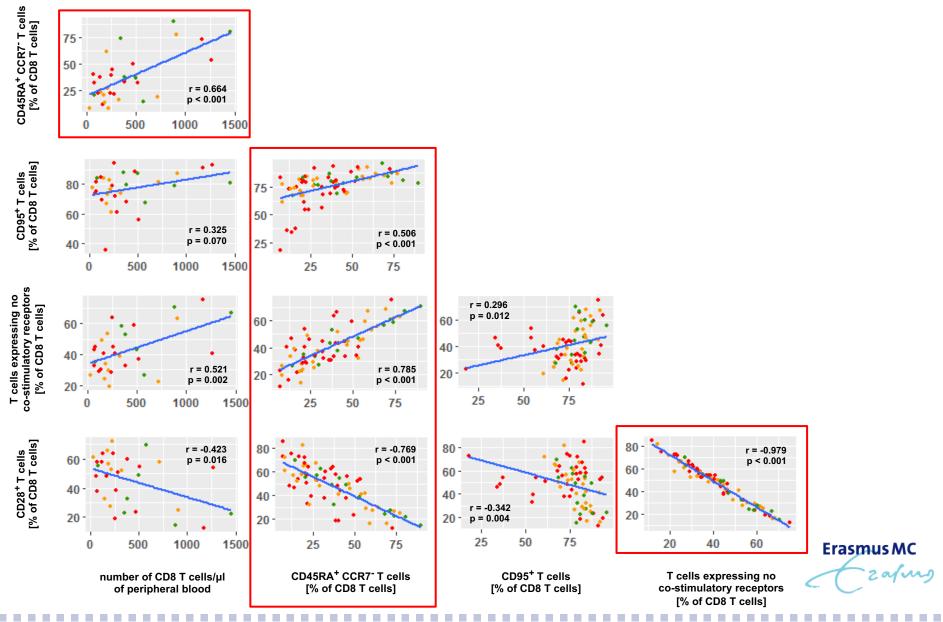
LAG3<sup>+</sup>

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Observations suggest a decrease function of CD8 T cells

## Number and differentiation phenotypes of CD8 T cells in PR patients are correlated to lack of co-stimulatory receptors



#### **Summary:**

#### NSCLC patients with partial response to nivolumab display at baseline:

- higher numbers of CD8<sup>+</sup> T cells
- enhanced frequency of terminally differentiated CD8<sup>+</sup> T cells (CD45RA<sup>+</sup>CCR7<sup>-</sup> and CD95<sup>+</sup>CD69<sup>-</sup>
- enhanced frequency of CD8<sup>+</sup> T cells devoid of co-stimulatory receptors (CD28<sup>-</sup> ICOS<sup>-</sup> CD40L<sup>-</sup> 4-1BB<sup>-</sup> OX40<sup>-</sup>)
- enhanced frequency of CD8<sup>+</sup> T cells co-expressing co-inhibiting receptors PD1 and TIM3
- → NSCLC patients responding to nivolumab have enhanced frequency of CD45RA+CCR7- CD8 T cells lacking co-stimulatory receptors
- → NSCLC patients responding to nivolumab appear to possess more antigen-experienced T cells

#### **Next steps:**

- Correlate findings with overall and progression free survival
- Investigate whether this profile is also found in other tumor types
- Correlate immune profiles with treatment-related toxicity
- Correlate immune profiles with nivolumab pharmacokinetics

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#### **Acknowledgements**

### Laboratory of Tumor Immunology, Department of Medical Oncology

- Cor Berrevoets
- Mandy van Brakel
- Merle van Geldrop
- Priscilla de Graaf
- Dora Hammerl
- Dian Kortleve
- Yarne Klaver
- Andre Kunert
- Cor Lamers
- Pim Mutsaers
- Astrid Oostvogels
- Chumud Phantunane
- Maud Rijnders
- Luc Veenman
- Bas Weening
- Rebecca Wijers
- Reno Debets













## Laboratory of Translational Pharmacology, Department of Medical Oncology

- Edwin Basak
- Daan Hurkmans
- Stijn Koolen
- Ron Matthijssen









#### **Department of Pulmonary Diseases**

- Joachim Aerts



#### **Department of Medical Oncology**

- Astrid van der Veldt
- Stefan Sleijfer



