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## Failure to validate existing clinical prediction scale for response to PD-1 monotherapy in advanced melanoma in national cohort study

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## CORRESPONDENCE



## Clinical Studies

# Failure to validate existing clinical prediction scale for response to PD-1 monotherapy in advanced melanoma in national cohort study

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## INTRODUCTION

Treatment with targeted therapy and immune checkpoint inhibitors has significantly improved survival of patients with advanced melanoma. Unfortunately, a large proportion of patients are either primary non-responders or will eventually develop secondary resistance.

In 2017, Nosrati and colleagues published a prediction scale in the *British Journal of Cancer*, which included five clinical parameters that were associated with lower response to anti-PD-1 treatment; female sex (1 point), age <65 years (1 point), history of ipilimumab (anti-CTLA-4) treatment (2 points), elevated lactate dehydrogenase (LDH) (1 point), and the presence of liver metastasis (2 points) [1]. This study used a derivation cohort of 228 patients treated in California, and a validation cohort of 87 patients treated in Switzerland. The primary outcome measure was best tumour response to treatment evaluated using computed tomography at 12 and 16 weeks after the first administration of anti-PD-1 monotherapy, and every 12 weeks thereafter.

The aim of this correspondence is to validate the prediction scale, published by Nosrati and colleagues.

## PATIENTS AND METHODS

### Registry

Since 2013, all patients with advanced melanoma in the Netherlands are referred to 1 of the 14 expert hospitals and data are prospectively registered in the Dutch Melanoma Treatment Registry (DMTR).

Data are collected from patient files by trained data managers and approved by the treating physicians. In compliance with Dutch regulations, the DMTR was approved by a medical ethical committee (METC Leiden University Medical Center, 2013) and is not considered subject to the Medical Research Involving Human Subjects Act.

### Patients and data

We extracted data for all patients registered between July 2013 and July 2018. Patients without response evaluation scans  $\geq 10$  weeks after start of treatment ( $n = 284$ ), with missing data

on the clinical parameters included in the prediction scale ( $n = 134$ ), or with uveal melanoma ( $n = 17$ ) were excluded. Baseline characteristics at the start of anti-PD-1 monotherapy were collected, including serum LDH, age, sex, previous treatments and the presence of liver metastasis. Response was defined as complete response (CR) or partial response (PR), based on clinical judgement of the medical team.

## RESULTS

Between July 2013 and July 2018, 1292 patients started anti-PD-1 treatment and met inclusion criteria. Baseline characteristics are summarised in Table 1A, including differences between the derivation cohort of Nosrati et al. and our national cohort. Patients' sex was more equally distributed in our cohort. Furthermore, our cohort contained more patients with WHO performance score  $>0$ , fewer patients with elevated LDH levels, fewer BRAF wild type melanoma, and fewer patients who were previously treated with ipilimumab or targeted therapy.

Table 1B presents all clinical parameters that were found to be significantly associated with response to anti-PD-1 monotherapy in the univariate analysis by Nosrati et al. Both prior ipilimumab treatment (odds ratio (OR) = 0.73 95% confidence interval (CI); 0.56–0.96,  $P = 0.02$ ) and the presence of liver metastases (OR = 0.70 (95% CI 0.54–0.90),  $P = 0.006$ ) were also found to be significantly correlated with lack of response to treatment in our cohort.

Figure 1 shows the predictive value of the clinical prediction scale of 0–7 points of Nosrati et al. With an AUC of 0.55 ( $P = 0.001$ ), this scale did not predict response to anti-PD1 monotherapy in our cohort.

## DISCUSSION

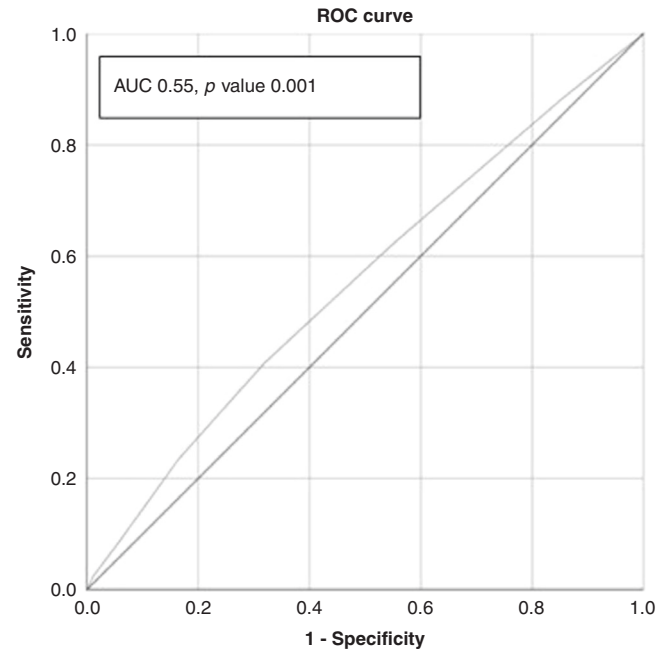
We could not confirm the predictive value of the clinical prediction scale of 0–7 points for response to anti-PD-1 monotherapy as published by Nosrati et al. A possible explanation could be the significantly higher ORR in the derivation (63.3%) cohort from Nosrati et al. compared to our cohort (49.8%), which could have led to an initial overestimation of the predictive value of their prediction scale. Additionally, our cohort differed from the group treated by Nosrati et al. when comparing the pre-treatment. More patients received prior targeted therapy in our cohort, while more patients received prior ipilimumab treatment in the group from Nosrati et al. Therefore, our cohort more closely resembles the current clinical setting where ipilimumab is less frequently given as a first line monotherapy for patients with advanced melanoma.

**Table 1.** Baseline characteristics and performance of prediction scale: (A) Comparison of baseline characteristics between validation cohort of Nosrati and colleagues and our cohort, using descriptive statistics. (B) Significance of predictive clinical parameters of Nosrati's univariate analysis in our cohort, calculated using logistic regression.

(A) Variable	Nosrati Number (%)	van der Kooij Number (%)	
Age, years			
Mean +/- SD	62.5 +/- 13.1	63.3 +/- 12.9	
Age <65 years	126 (55.3)	627 (48.5)	
Sex			
Male	148 (64.9)	771 (59.7)	
Female	80 (35.1)	521 (40.3)	
Primary site			
Cutaneous	200 (87.7)	1032 (79.9)	
Mucosal	13 (5.7)	43 (3.3)	
Acral		32 (2.5)	
Eye			
Unknown	15 (6.6)	185 (14.3)	
ECOG			
0	157 (68.9)	725 (56.1)	
1	65 (28.5)	419 (32.4)	
2	5 (2.2)	58 (4.5)	
3	1 (0.4)	6 (0.5)	
Unknown		84 (6.5)	
LDH			
Normal	150 (65.8)	939 (72.7)	
Elevated	78 (34.2)	353 (27.3)	
BRAF mutation			
Negative	162 (72.0)	619 (47.9)	
Positive	63 (28.0)	626 (48.5)	
Unknown	3 (1.3)	47 (3.6)	
Liver metastasis			
No	160 (70.2)	968 (74.9)	
Yes	68 (29.8)	324 (25.1)	
Lung metastasis			
No	94 (42.1)	595 (46.1)	
Yes	132 (57.9)	678 (52.5)	
Unknown		19 (1.4)	
Brain metastasis			
No	178 (78.1)	961 (74.4)	
Yes	50 (21.9)	294 (22.8)	
Unknown		37 (2.8)	
Prior ipilimuma			
No	<b>81 (35.5)</b>	<b>1021 (79.0)</b>	
Yes	147 (64.5)	271 (21.0)	
Prior targeted therapy			
No	<b>174 (76.3)</b>	<b>1144 (88.5)</b>	
Yes	54 (23.7)	148 (11.5)	
(B)	ORR (%)	OR (95% CI)	P value
Total cohort	49.8	NA	NA
Age ≥65 years	49.7	Ref.	Ref.
Age <65 years	50.3	1.03 (0.82–1.28)	0.82
Normal LDH	51.6	Ref.	Ref.

**Table 1.** continued

(B)	ORR (%)	OR (95% CI)	P value
Elevated LDH	45.7	0.79 (0.62–1.01)	0.06
Male sex	50.5	Ref.	Ref.
Female sex	49.3	0.96 (0.77–1.19)	0.69
No prior ipilimumab	51.6	Ref.	Ref.
Prior ipilimumab	43.9	0.73 (0.56–0.96)	0.02
No liver metastasis	52.2	Ref.	Ref.
Liver metastasis	43.3	0.70 (0.54–0.90)	0.006








**Fig. 1** Receiver operation characteristics (ROC) curve of the clinical prediction scale of 0–7 points of Nosrati et al. to predict response to anti-PD-1 monotherapy in our cohort. The Area Under Curve (AUC) of our cohort (blue line) is shown. The red line indicates random prediction.

Although the prediction scale could not be validated in our cohort, we did show that prior ipilimumab treatment and the presence of liver metastases was associated with a smaller response chance. This lack of response in the group of patients that has been pre-treated with ipilimumab could be due to the fact that patients who already progressed on prior immune checkpoint inhibition have a primary or acquired resistance to this type of treatment [2]. And therefore might also be less susceptible to a second line of immunotherapy.

In recent years, multiple meta-analyses have been published investigating the sex-dependent magnitude of benefit following treatment with immune checkpoint inhibition. The first study showed that men have more benefit from immune checkpoint inhibition, including anti-PD-1 [3], whereas the latter three showed no difference in efficacy and overall survival [4–6]. Our study supports the findings that sex on itself is not a predictor for response to anti-PD-1 treatment.

Failure to validate the prediction scale by Nosrati et al. indicates that response to anti-PD-1 monotherapy cannot only be predicted by clinical parameters, but is influenced by other factors. Examples currently being studied include tumour-intrinsic factors, immune cells and cytokines both in tumour tissue and blood [7, 8] and include more readily available blood parameters, such as LDH,

S100B, absolute leucocyte, lymphocyte, neutrophil counts and their ratios [9–11]. While further research on predictive models is encouraged, validation of these models in sufficiently large independent cohorts is of even more importance to test robustness and clinical applicability.

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## DATA AVAILABILITY

Study protocol: available from MKvdk (e-mail, m.k.van\_der\_kooij@lumc.nl). Statistical code: not available. Data set: can be applied for at <https://dica.nl/dmtr/onderzoek>.

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## AUTHOR CONTRIBUTIONS

Conceptualisation, MKvdk, OMD and EK; data curation, MKvdk; formal analysis, MKvdk; methodology, MKvdk, OMD and EK; resources, FWPJvdB, MJB-S, JWBdG, GAPH, DP, RsvR, KPMS, HMW, AAMvdV, GV, CUB, MWJMW, JBAGH, AJMvdE and EK; software, MJBA; supervision, OMD and EK; visualisation, MKvdk; writing—original draft, MKvdk; writing—review and editing, OMD, MJBA, FWPJvdB, MJB-S, JWBdG, GAPH, DP, RsvR, KPMS, HMW, AAMvdV, GV, CUB, MWJMW, JBAGH, AJMvdE and EK. All authors have read and agreed to the published version of the manuscript.

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## COMPETING INTERESTS

MJBS has served as an advisory board member for Bristol-Myers Squibb, Novartis, Merck and Pierre Fabre. AJMvdE has served as a speaker for Bristol-Myers Squibb and Novartis and an advisory board member for Bristol Myers Squibb, MSD oncology, Amgen, Roche, Novartis, Sanofi, Pfizer, Ipsen, Merck, Pierre Fabre and has received research grants not related to this paper from Sanofi, Roche, Bristol Myers Squibb, TEVA and Idera. CUB has served as an advisory board member for Bristol-Myers Squibb, MSD, Roche, Novartis, GlaxoSmithKline, AstraZeneca, Pfizer, Lilly, GenMab, and Pierre Fabre, and reports to have ownership interests in Uniti Cars, Neon Therapeutics, and Forty Seven, and received commercial grants from Novartis, Bristol-Myers Squibb, and NanoString. JWBdG is a paid consultant for Bristol Myers Squibb, MSD, Pierre Fabre, and Servier. GAPH is an unpaid consultant/advisory board member for Bristol Myers Squibb, MSD, Roche, and Novartis. DP has served as an advisory board member for Amgen, Bristol Myers Squibb and Pierre Fabre. AAMvdV is a paid consultant for Bristol Myers Squibb, MSD, Novartis, Roche, Pfizer, Eisai, Ipsen, Pierre Fabre, Sanofi, and Bayer. JH is a paid consultant for AIMM, Neon Therapeutics, Immunocore, Vaximm, and Neogene Therapeutics, and reports receiving commercial research grants from Bristol Myers Squibb, MSD, Novartis, and Neon Therapeutics. KPMS has served as a consultant and/or advisory board member for Bristol Myers Squibb, Novartis, MSD, Pierre Fabre and AbbVie and received honoraria/research support not related to this manuscript from Novartis, Roche and MSD. All paid to

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#### **ETHICS APPROVAL AND CONSENT TO PARTICIPATE**

The DMTR was approved by a medical ethical committee (METC Leiden University Medical Center, 2013) and is not considered subject to the Medical Research Involving Human Subjects Act. Patient consent was waived due to the fact that the

DMTR was not considered to be subject to the Medical Research Involving Human Subjects Act.

#### **ADDITIONAL INFORMATION**

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