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Antoine Girard, Mathieu Rouanne. Comment to: Relationship between the expression of PD-1/PD-L1 and 18 F-FDG uptake in bladder cancer. *European Journal of Nuclear Medicine and Molecular Imaging*, 2019, 46 (6), pp.1212-1213. 10.1007/s00259-019-04296-6 . hal-02127098

HAL Id: hal-02127098

<https://hal-univ-rennes1.archives-ouvertes.fr/hal-02127098>

Submitted on 13 May 2019

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Letter to the editor

Title: Comment to: Relationship between the expression of PD-1/PD-L1 and ¹⁸F-FDG uptake in bladder cancer

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Conflict of Interest or Financial Support: none

Ethical approval: This article does not describe any studies with human participants performed by any of the authors.

Key Words: ¹⁸F-FDG PET, Bladder Cancer, PD-1/PD-L1 expression, Immunohistochemistry, Immunotherapy

Word count: Text: 623 words

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Letter to the editor

Title: Comment to: Relationship between the expression of PD-1/PD-L1 and ^{18}F -FDG uptake in bladder cancer

Text:

Dear Sir,

The study by Chen, et al. addresses an important question regarding the predictive value of ^{18}F -FDG uptake to assess PD(L)-1 expression in bladder cancer [1]. Arguably, this innovative study is of major interest as multiple issues hamper the standardization of programmed cell death ligand-1 (PD-L1) scoring in tumour tissue. Recently, the Food and Drug Administration (FDA) has issued a drug safety notification warning against the use of frontline single-agent immune checkpoint inhibitors for patients with PD-L1-low expressing urothelial carcinoma. In August 2018, PD-L1 status has been incorporated into the labels for pembrolizumab and atezolizumab for existing frontline approvals in cisplatin-ineligible urothelial carcinoma. Therefore, this article shed light on a particularly relevant question from a clinical and scientific perspective but also raises technical issues regarding the method described.

First, the authors did not detail the staining protocol nor the antibody used to assess PD-L1 expression. Four PD-L1 assays have been FDA/ European Medicines Agency (EMA) approved in urothelial carcinoma, including the Dako 28-8, 22-c3 and the Vantana SP142 and SP263 monoclonal antibodies [2]. Divergent staining results have been reported in bladder cancer, leading to different detection rates of eligible patients for first-line treatment with immune checkpoint blockade [3]. The threshold $>1\%$ for PD-L1 positivity was clearly defined by Chen et al [1]. However, it remains unclear which PD-L1 stained cells have been evaluated in this study (tumour cells? immune cells? both of them?). The authors should have reported the PD-L1 score for each component of the tumour immune microenvironment. Indeed, the variability of PD-L1 expression across the type of stained cells may lead to variable relationship with ^{18}F -FDG uptake values.

Second, as immune checkpoint inhibitors are validated treatments in metastatic urothelial cancer, it would have been of interest to evaluate the correlation between ^{18}F -FDG uptake and PD-L1 expression in metastases rather than in the primary tumour. Indeed, the latter one is easily accessible by endoscopic transurethral resection. Five immune checkpoint inhibitors obtained accelerated approval by the FDA to treat metastatic bladder cancer [4], and molecular and functional imaging is increasingly recognized as a reliable tool for cancer staging [5, 6]. Burgess et al, recently reported a discordance of PD-L1 immune cells expression between primary and metastatic urothelial carcinoma lesions [7]. Thus, the correlation between ^{18}F -FDG uptake and PD-1/PD-L1 expression in distant metastases may have been more variable.

Third, non-muscle invasive bladder cancer (NMIBC) stands for more than 70% of bladder cancers at time of diagnosis. NMIBCs are often detected as an infracentimetric thickening of the bladder wall. Approximately one third (13/38) of tumours analysed by Chen, et al, were NMIBC disease [1]. In this situation, ^{18}F -FDG uptake for most of the tumours \leq pT1 may have been underestimated due to partial volume effect. This phenomenon is well known to underestimate the uptake intensity in small or thin structures (i.e. $< 8\text{-}12$ mm approximately depending on the full width at half maximum (FWHM) of the reconstructed image resolution) [8]. Tumour size (presumably in the great axis) was included in the multivariate analysis, but

1 the thickness was not. In addition, stromal lymphocyte infiltration is associated with tumour
2 invasion depth in pT1 bladder cancer, suggesting that the immune infiltration is different
3 between muscle-invasive and non-muscle invasive tumours [9]. Indeed, most of those tumours
4 \leq pT1 were PD-1 and PD-L1 negative, which may bias the results.

5
6 Arguably, predicting and monitoring the response to immune checkpoint inhibitors are
7 becoming key issues in the near future. Numerous phase III trials evaluating anti-PD(L)1
8 immune checkpoint inhibitors are currently ongoing in both localized and advanced bladder
9 cancer settings. Metabolic and molecular imaging will have a major impact on patients'
10 management in the new era of cancer immunotherapy. The study reported by Chen et al, takes
11 first step on a very promising way [1].
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