

NOVEL DIGITAL IMAGE APPROACH OF MULTIPLEX IMMUNOFLUORESCENCE BASED PD-L1 EXPRESSION ENABLES THE STRATIFICATION OF ADVANCED NSCLC PATIENTS TREATED WITH DURVALUMAB

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Background Pathologist-based scoring of PD-L1 expression on tumor cells using IHC¹ has shown clinical utility in predicting favorable overall survival in advanced non-small cell lung cancer (NSCLC) patients treated with anti-PD-(L)1 therapies including durvalumab.²⁻³ Quantitative Continuous Scoring (QCS)⁴ enables the continuous measurement of the PD-L1 expression on single cells and the selection of the PD-L1 expression cutoff that best stratifies anti-PD-L1-treated patients with respect to prevalence and log-rank test p-value.⁵ We present here the extension of QCS to PD-L1 measured by multiplex immunofluorescence (mIF)⁶ to evaluate its ability to optimize patient stratification.

Methods Pre-treatment tumor samples from advanced NSCLC patients enrolled in durvalumab nonrandomized phase 1/2 trial (CP1108/NCT01693562)², were stained by mIF panel containing PD-L1.⁶ Similarly to IHC PD-L1 QCS, mIF PD-L1 QCS consists of two deep-learning models, first to segment epithelium regions and second to detect membrane, cytoplasm and nuclei of each epithelium cell, transferring for the second model annotations from IHC to mIF domain.⁷ The mIF images are normalized based on batch statistics prior to image analysis. PD-L1 expression is measured for each epithelium cell as the average of the PD-L1 signal in the segmented membrane. Cells with expression higher than an expression threshold (T_{PD-L1}) are considered positive. A slide is considered QCS-positive if it comprises a greater percentage of PD-L1 positive cells (QCS-score) than a cutoff value (CoV).

Results The QCS-scores are computed on 119 NSCLC patients treated with durvalumab. As a first proof of concept that QCS-scoring can replicate tumor proportion scoring (TPS), we optimize T_{PD-L1} as to maximize the correlation between QCS and TPS scores (figure 1). Second, we estimate for different combinations of (T_{PD-L1} , CoV) the log rank p-value associated with the stratification between patients with low and high QCS scores. A subregion of the parameter space was identified for which the stratification is significant ($p < 0.01$) with more than 50% prevalence in the positive subgroup (figure 2). The p-value is minimized ($p = 7.2 \cdot 10^{-5}$) for ($T_{PD-L1} = 37$, $CoV = 0.75\%$), yielding a median OS of 5.58 months and 13.44 months in the QCS negative and positive subgroups respectively, similar to those of IHC PD-L1 manual scoring with 25% cutoff.

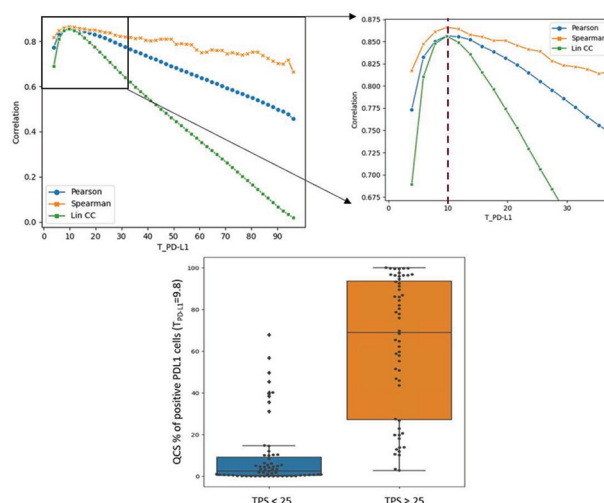
Conclusions The extension of QCS to mIF imaging provides opportunities to evaluate continuous PD-L1 expression of single tumor cells in relation to spatial distribution of other cells (e.g. PD1+ CD8+ T cells) and identify predictive biomarkers of tumor-immune cell interactions of anti-PD-(L)1 therapies.

Trial Registration CP1108/NCT01693562

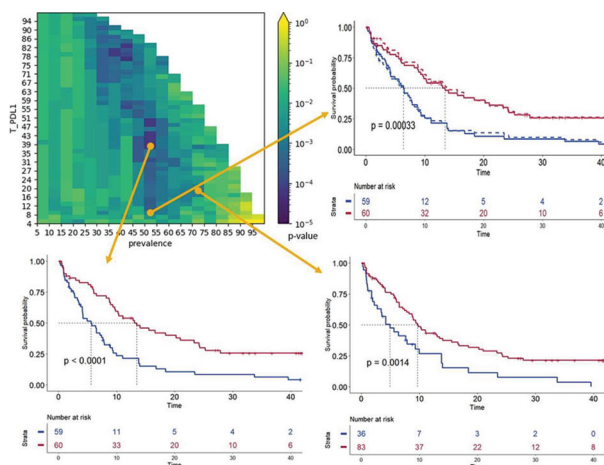
REFERENCES

1. Rebelatto M, et al. Development of a programmed cell death ligand-1 immunohistochemical assay validated for analysis of non-small cell lung cancer and head and neck squamous cell carcinoma. *Diagnostic Pathology* 2016 Oct 8;11(1):95.
2. Antonia S, et al. Clinical Activity, Tolerability, and Long-Term Follow-Up of Durvalumab in Patients With Advanced NSCLC *Journal of thoracic Oncology* 2019 Oct ; 14 (10):1794–1806

3. Rizvi NA, et al. Durvalumab With or Without Tremelimumab vs Standard Chemotherapy in First-line Treatment of Metastatic Non-Small Cell Lung Cancer. *JAMA Oncology* 2020;6(5) :661–674
4. Gustavson M, et al. Novel approach to HER2 quantification: digital pathology coupled with AI-based image and data analysis delivers objective and quantitative HER2 expression analysis for enrichment of responders to trastuzumab deruxtecan (T-DXd; DS-8201), specifically in HER2-low patients. *Cancer Res* 2021, 81 (4_Supplement): PD6–01.
5. Schmidt G, et al. Computational pathology delivers objective and quantitative PD-L1 expression analysis for enrichment of responders to durvalumab in non-small cell lung cancer (NSCLC). *J Immunother Cancer* 2021;9(Suppl 2):A1–A1054
6. Meinecke L, et al., Presence of TLS and combined high densities of PD-L1+ macrophages & CD8+ T cells predict long-term overall survival for patients with advanced NSCLC treated with durvalumab. *Cancer Res* 2022 82 (12_Supplement): 1235.
7. Brieu N, et al. Stain Isolation-based Guidance for Improved Stain Translation, Medical Imaging with Deep Learning (MIDL) 2022, <https://arxiv.org/abs/2207.00431>



Abstract 579 Figure 1 Correlation to pathologist-based TPS score Top: Lineplots of Pearson and Spearman correlations as well as of Lin correlation coefficient between the pathologist-based TPS scores and the QCS-based scores, computed for increasing expression threshold values (TPD-L1). The QCS shows maximum Pearson (0.856), Spearman (0.866) and Lin (0.856) correlations to the manual TPS score for $TPD-L1 = 10$. Bottom: QCS scores within the negative and positive patient subgroups as per pathologist assessment of IHC PD-L1 TPS.



Abstract 579 Figure 2 OS patient stratification Log rank p-values for OS stratification obtained by spanning the parameter space associated to the QCS, the higher CoV the lower the prevalence of the positive patient subgroup. Top right: Kaplan Meier (KM) curves obtained with manual IHC PD-L1 TPS score at 25% cutoff

(dashed line) and with median split for the QCS expression cut-off (TPD-L1=10) maximizing the correlation to TPS (full line). Bottom: KM curves of the QCS-based stratification as to minimize the p-value (TPD-L1=37) for a minimum prevalence of 50% (left) and as to maximize the prevalence (TPD-L1=18) (right).

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