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Endobronchial Ultrasound-Guided Transbronchial Needle Aspiration for PD-L1 Testing In Non-Small Cell Lung Cancer

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1 **Endobronchial Ultrasound-Guided Transbronchial Needle Aspiration for PD-L1 Testing In**
2 **Non-Small Cell Lung Cancer**

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24

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59 **Abbreviation List**

- 60
- 61 ALK: anaplastic lymphoma kinase gene
- 62 CNB: core needle biopsy
- 63 CT: computed tomography
- 64 EGFR: epidermal growth factor receptor gene
- 65 EBUS-TBNA: endobronchial ultrasound-guided transbronchial needle aspiration
- 66 EUS: endoscopic ultrasonography
- 67 FNA: fine needle aspiration
- 68 ICH: immunohistochemistry
- 69 OR: odds ratio
- 70 OS: overall survival
- 71 NOS: not otherwise specified
- 72 NSCLC: non-small cell lung cancer
- 73 PD: disease progression
- 74 PD-1: programmed death-1
- 75 PD-L1: programmed death-ligand 1
- 76 PR: partial response
- 77 TPS: tumor proportion score
- 78 VATS: video-assisted thoracoscopic surgery
- 79

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80 **Abstract**

81 *Rationale:* PD-L1 expression on cancer cells is a clinically important biomarker to select NSCLC
82 patients for treatment with PD-1/PD-L1 inhibitors. Clinical trials of immunotherapy in patients with
83 non-small cell lung cancer have required histology for PD-L1 testing, while in clinical practice
84 cytology samples are commonly acquired in patients with advanced disease.

85 *Objectives:* This study investigates sampling adequacy of endobronchial ultrasound-guided
86 transbronchial needle aspiration (EBUS-TBNA) for PD-L1 testing when compared to other
87 methods. Furthermore, the relationship between clinico-pathological characteristics and PD-L1
88 expression in the study population have been examined.

89 *Methods:* Five hundred seventy-seven NSCLC specimens were analysed from consecutive patients
90 with NSCLC across six centres in United Kingdom and one in the United States between January
91 2015 and December 2016.

92 *Main Results:* In the EBUS-TBNA group (189 specimens), the overall percentage of patients with
93 successful PD-L1 testing was 94.7%. There was no significant difference in sampling adequacy
94 with other methods of tissue acquisition. Older subjects had higher failure rates of PD-L1 testing
95 (OR= 1.06, p=0.008). In multivariate analysis, advanced N-stage (p=0.048) and presence of brain
96 metastasis (p<0.001) were associated with high PD-L1 expression.

97 *Conclusion:* This large multicenter study shows that EBUS-TBNA provides samples adequate for
98 PD-L1 testing and that advanced N stage and the presence of brain metastasis are associated with
99 high PD-L1 expression.

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101

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103 Introduction

104

105 Immune checkpoint inhibitors have demonstrated significant clinical utility in patients with
106 advanced non-small cell lung cancer (NSCLC), and several anti PD-1 and anti PD-L1 monoclonal
107 antibodies have been approved as first or second-line therapies ¹⁻⁵. These agents interfere with both
108 costimulatory and co-inhibitory pathways regulating the antigen specific T-cell response ⁶. PD-1 is
109 a cell-receptor involved in programmed cell death. The PD-1 receptor binds to the ligands PD-L1
110 and PD-L2 and results in downregulation of anti-tumor cytolytic T-cell activity, inducing T cell
111 exhaustion and immune tolerance.

112 The correlation between PD-L1 immunohistochemistry (IHC) expression, measured by the
113 proportion of cancer cells positively staining for PD-L1, and the overall response to anti-PD-1 or
114 anti-PD-L1 agents has been demonstrated in clinical trials. In the landmark KEYNOTE-024 trial¹,
115 Pembrolizumab, an anti-PD-1 agent, resulted in better progression-free survival and overall survival
116 compared to standard chemotherapy in patients with a tumor proportion score of 50% or greater.
117 Therefore, PD-L1 IHC expression is currently used to select patients with advanced lung cancer
118 who may benefit from first line immunotherapy alone. In this study however, core biopsies of tumor
119 were mandated for trial entry¹.

120 Endobronchial ultrasound-guided transbronchial needle aspiration (EBUS-TBNA) is a minimally
121 invasive technique, proven to be effective in obtaining cytology samples suitable for the molecular
122 characterization of NSCLC ⁷. However, despite its routine use in clinical practice, patients
123 undergoing tissue acquisition by EBUS-TBNA alone were excluded from immunotherapy trials ⁸.

124 We therefore conducted a large, pragmatic, multi-center study to examine whether samples
125 obtained by EBUS-TBNA were suitable for PD-L1 assessment and selection of patients for immune
126 checkpoint inhibition. We compared the diagnostic yield of different methods including cytology
127 samples, small biopsies and lung resections. We also systematically collected patient and procedure
128 characteristics to define factors that predicted a reliable PD-L1 result and PD-L1 expression.

129

130 **Methods**131 *Study design*

132 This study included consecutive patients with known or suspected NSCLC undergoing tissue
133 acquisition procedures between January 2015 and December 2016 across six centers in the United
134 Kingdom (University College London Hospital, University Hospital Birmingham, Lancashire
135 Teaching Hospital, Nottingham University Hospitals, University of South Manchester and
136 Papworth Hospital, Cambridge) and one center in the United States (Johns Hopkins University).
137 The specimens were obtained by EBUS-TBNA, percutaneous fine needle aspiration (FNA),
138 percutaneous core needle biopsy (CNB), medical thoracoscopy, video-assisted thoracoscopic
139 surgery (VATS) or open thoracotomy. Samples were analyzed and interpreted according to local
140 protocols and there was no centralized reporting. Genotyping was performed in all non-squamous
141 NSCLC or in other subtypes according to clinical judgment. Non-squamous NSCLC samples were
142 prioritized for mutation testing of the epidermal growth factor receptor gene (EGFR),
143 rearrangement of the anaplastic lymphoma kinase gene (ALK) and ROS-1 re-arrangement where
144 necessary. Samples were subsequently evaluated for PD-L1 expression.

145

146

147 *EBUS-TBNA samples*

148 EBUS-TBNA was performed with a dedicated linear echo-endoscope as previously described⁹. The
149 procedure at John Hopkins and University College London Hospital were carried out under general
150 anesthesia in 100% (14/14) and (33/49) 67% of cases respectively. All the other cases were done in
151 the outpatient setting with patients given moderate sedation with midazolam and fentanyl. In brief,
152 under direct ultrasound guidance, the lymph node was aspirated using either a 19, 21, 22 or 25-
153 gauge needle. The site and number of lymph nodes punctured were at the operator's discretion.
154 Four passes per lymph node were routinely performed in all cases. If these passes did not visually

155 return adequate material, at least 2 more passes from the same lymph node were additionally
156 performed. A suction syringe was applied to the needle during lymph node aspiration. On-site
157 evaluation of samples was not routinely employed. The samples obtained at EBUS-TBNA were
158 expelled from the needle using the stylet and placed into liquid fixative for cell-block processing.
159 The specimen was centrifuged to form a pellet, suspended in agar, fixed in neutral buffered
160 formalin or alcohol-based fixative, and processed as a cell block from which a single hematoxylin
161 and eosin (H&E)-stained section was cut. Further sections were cut and used for IHC staining as
162 required.

163

164 *PD-L1 assessment*

165 All the centers involved in this study used the Dako PD-L1 IHC 22C3 pharmDx
166 immunohistochemical assay (Dako, Glostrup, Denmark). This assay uses a monoclonal antibody
167 (humanized IgG4) that recognize the extracellular domain of PD-L1 to assess PD-L1 expression in
168 formalin-fixed, paraffin embedded (FFPE) tissue. The IHC staining procedure was performed on a
169 Dako Autostainer Link 48 platform with a validated staining protocol. PD-L1 expression was
170 evaluated by tumor proportion score (TPS), which is defined as the percentage of viable tumor cells
171 with at least partial membrane staining relative of all viable tumor cells in the examined section. All
172 other stained cells, such as tumor-associated immune cells, normal/non-neoplastic cells, and
173 necrotic cells, were excluded from evaluation. A minimum of 100 viable tumor cells were required
174 to consider the specimen adequate. The scoring was interpreted as: no PD-L1 expression
175 (TPS<1%); low PD-L1 expression (TPS 1-49%); and high PD-L1 expression (TPS \geq 50%), in line
176 with current clinical practice and immunotherapy licensing. Ethical approval was not required given
177 the observational nature of the study. All data were prospectively recorded in each center, though
178 the study design is retrospective as reported previously⁷. Treatment strategies were fully disclosed
179 to the patients and were discussed in multidisciplinary team meetings.

180

181 **ENDPOINTS AND STATISTICAL ANALYSIS**

182 The primary aim of the study was to evaluate the diagnostic performance of PD-L1 testing in
183 specimens obtained by EBUS-TBNA in patients with NSCLC compared to other methods.
184 Secondary endpoints were to define clinico-pathological characteristics associated with a reliable
185 PD-L1 result and also to define clinical features associated with PD-L1 high expression.
186 Associations between baseline characteristics and a successful PD-L1 test were assessed using chi-
187 square tests, chi-square trend tests, and t-tests as appropriate. Baseline variables considered were
188 age, performance status, smoking status, TNM stage, presence of brain metastasis, pathological
189 tumor differentiation, actionable mutations and sampling method. Individual factors associated with
190 PD-L1 level (none, low or high), and with high PD-L1 were assessed using ordinal regression and
191 logistic regression respectively. Predictors of high PD-L1 level were further investigated through a
192 multi-variable model generated using forward selection and backward elimination processes,
193 assessing all variables with a p-value<0.25 on univariate analysis. All statistical calculations were
194 performed using STATA version 16 (StataCorp, College Station, TX).

195

196 **RESULTS**

197 *Study population*

198 Five hundred seventy-seven NSCLC specimens were analyzed from consecutive patients with
199 NSCLC. Three hundred eighteen subjects (55%) were male and the median age of the study
200 population was 68 years (range, 31-96 years). Tissue acquisition techniques included 189 (33%)
201 EBUS or EUS, 72 (12%) endobronchial biopsy, 167 (29%) CT-guided procedures, 124 (21%)
202 surgical excisions or resections, 6 (1%) pleural biopsy and 19 (3%) other site specimens.
203 Demographic and baseline characteristics are summarized in Table 1. Three hundred seventy-eight
204 patients (66%) had a final diagnosis of Adenocarcinoma, 151 (26%) Squamous Cell Carcinoma
205 while 48 (8%) received other diagnoses (Adenosquamous, not otherwise specified (NOS), Large
206 Cell Carcinoma, Other). The presence of EGFR mutations was reported in forty-one patients (7%),

207 ALK rearrangement in seven patients (1%) and ROS-1 rearrangement in only one case (<1%). For
208 EBUS-TBNA, 22-gauge needle was used in 78% of cases, 21-gauge needle in 20%, 19 and 25-
209 gauge needle in 1%. Lymph node stations sampled by EBUS-TBNA were reported in
210 Supplementary Table 1. Seven patients (3.7%) who underwent EBUS-TBNA had complications,
211 none of which resulted in early interruption of the procedure. In particular, significant bleeding
212 determined by the operator was documented in six cases (3.2%) while one patient (0.5%)
213 experienced desaturation with early recovery after the procedure. No patients required inpatient
214 admission after the procedure. The complication rates for endobronchial and transbronchial forceps
215 biopsies, CT-guided biopsies and surgery were 4.1%, 9.0% and 11.5%, respectively.

216

217 *PD-L1 assessment*

218 PD-L1 assessment was reported to be feasible in the majority of cases (Table 2). The overall rate of
219 assessment failure was 5% (29 patients). EBUS-TBNA provided adequate sampling for reliable PD-
220 L1 testing in 95% (179/189) of patients. PD-L1 testing was feasible in 155/167 CT guided biopsies
221 (93%), 70/72 endobronchial biopsies (97%), 123/124 surgical specimens (99.2%) and 6/6 pleural
222 biopsies (100%).

223 Failure rate among patients diagnosed with “other” methods was 21% (4/19), resulting in a
224 statistically significant difference when compared to the rest of the study population. Successful
225 PD-L1 assessment rates were similar across the different centers (range, 88.8%-100%). In the
226 EBUS-TBNA group, no differences were observed in yield between 21 and 22-gauge needle
227 ($p=0.39$). Older age was the only predictor of failure of PD-L1 assessment ($OR= 1.06$, $p=0.008$) in
228 univariate analysis. The likelihood of a successful PD-L1 assay did not vary according to study site,
229 gender, ethnicity, smoking status, pack years, performance status, T-stage, N-stage, M-stage,
230 histological subtype, actionable mutations, biopsy type (original vs re-biopsy), presence of brain
231 metastases or receipt of prior radiotherapy. Although the non-squamous samples also underwent
232 analysis for EGFR mutations and ALK and ROS-1 rearrangement, specifically no differences were

233 observed in PD-L1 assessment failure rate between adenocarcinoma and squamous cell carcinoma
234 (p=0.825).

235

236

237 *Predictors of PD-L1 expression*

238 PD-L1 tumor proportion staining was negative (<1%) in 234 patients (42.7%), low (1-49%) in 159
239 patients (29.0%), and high ($\geq 50\%$) in 155 patients (28.3%). PD-L1 expression was not influenced
240 by the tissue sampling method (Table 3). However, we found that PD-L1 high expression was
241 associated with the presence of brain metastasis (p= 0.009). In the model, dividing the study
242 population into high expression (TPS $\geq 50\%$) versus no or low expression (TPS < 50%) we found
243 that high PD-L1 expression was associated with advanced N-stage (p=0.024), M1 stage (p=0.031),
244 Adenocarcinoma subtype (p=0.023) and presence of brain metastasis (p<0.001). The final
245 multivariate model showed that higher N-stage (p=0.048) and the presence of brain metastasis
246 (p<0.001) were independently associated with high PD-L1 expression.

247

248 *Response to immunotherapy*

249 Fifty-six patients received immune checkpoint inhibitors (44 Pembrolizumab, 10 Nivolumab, 1
250 Atezolizumab, 1 Durvalumab). Table 4 demonstrates the response to immune checkpoint inhibitors
251 according to the line of treatment. 25 (44.6%) patients had disease progression, 20 (35.7%) patients
252 had stable disease, while 11 (19.6%) patients achieved a partial response. All patients with a partial
253 response were observed to have high PD-L1 expression. Disease response was not associated with
254 mode of tissue sampling.

255

256 **DISCUSSION**

257 *PD-L1 expression as predictive biomarker in NSCLC*

258 In the management of advanced NSCLC, molecular subtyping and PD-L1 status assessment have
259 become critical in selecting the most appropriate treatment¹⁰. Recently, several anti PD-1 and anti
260 PD-L1 agents have been approved by the FDA and EMA for patients with metastatic NSCLC who
261 do not harbor an EGFR mutation or ALK rearrangement both in the first and second line settings.
262 Pembrolizumab has received approval for first-line monotherapy for patients with tumor in which at
263 least 50% of cells express PD-L1 or in second-line treatment for patients with tumor whose at least
264 1% cells express PD-L1 on cell surface¹¹⁻¹⁴. However, these trials specifically excluded patients
265 with tissue acquired by EBUS-TBNA despite at least 1/3 of patients having this procedure in
266 clinical practice. In 2016, the Papanicolaou Society of Cytopathology recommended against the use
267 of cytology samples for PD-L1 IHC testing due to insufficient data¹⁵. Similarly, the Pulmonary
268 Pathology Society¹⁶ highlighted the lack of validation for cytology preparation in PD-L1 testing,
269 though for many patients with advanced NSCLC they are often the only specimens available. PD-
270 L1 analysis is now also required in Europe for patients with stage III disease to receive
271 immunotherapy after concurrent chemoradiotherapy¹⁷. For these patients, EBUS-TBNA provides an
272 important dual purpose of providing a tissue diagnosis as well as accurately mapping malignant
273 intra-thoracic lymph nodes. In this study, we show that EBUS-TBNA provides samples suitable for
274 PD-L1 testing and that response rates to immunotherapy do not depend upon modality of tissue
275 acquisition.

276 Several limitations of assessing PD-L1 expression are recognized. These include the tumor spatial
277 heterogeneity among different sections of the same sample or at different sites coupled with the
278 dynamic changes in PD-L1 expression over time¹⁸. However, at this time it represents the only
279 biomarker approved by the regulatory agencies for first line immunotherapy in NSCLC, while
280 others (microsatellite instability, tumor mutation burden, tumor microenvironment, gut
281 microbiome) are currently under investigation or in late stage of development.

282

283 *PD-L1 quantification in EBUS-TBNA*

284 Few studies have investigated the feasibility of PD-L1 assessment by EBUS-FNA¹⁹. Stoy et al²⁰
285 examined the PD-L1 quantification in cytology specimens and they showed successful assessment
286 in 90.9% (20/22) of patients. This study included sixteen EBUS-TBNA, four endobronchial fine
287 needle aspirations and two bronchoscopic-FNA of peripheral nodules; two unsuccessful EBUS tests
288 were because the cell block had <100 cells. They also found a good concordance in two patients
289 who had same site both cytology and histology samples. In another single-center retrospective study
290 collecting 188 patients with lung cancer, Heyman et al.²¹ found that cytology specimens were
291 adequate for PD-L1 quantification in 90% of patients, while small biopsy and surgical resection
292 completed assessment rates were 96% and 99%, respectively. Interestingly, only 25 of 214
293 specimens (11.7%) were from EBUS-TBNA, while 36.0% of samples were from surgical resection
294 which is not commonly performed in patients who are currently candidates for immune checkpoint
295 inhibitors. Similar results are described in a larger study which included 252 EBUS-TBNA samples
296 and compared cytology, small biopsies and surgical resections. The authors reported 92% sample
297 adequacy for PD-L1 testing for cytology or small biopsy specimens²². In this study the fixation
298 process (formalin only versus methanol/alcohol only versus both) did not influence the PD-L1
299 staining. Very recently, Biswas et al, using the PD-L1 22C3 pharmDx assay, confirmed that EBUS-
300 TBNA was able to allow the PD-L1 quantification in 86% of cases²³. These studies reflect our
301 findings in 566 patients in whom the rate of failure of PD-L1 testing was 4.8% in the EBUS group.
302 Other studies have investigated the concordance in PD-L1 expression between EBUS-TBNA and
303 other samples²⁴. Sakakibara et al.²⁵ found a good correlation between EBUS samples and surgical
304 samples in both primary ($r=0.75$; $p=0.08$, $n=6$) and metastatic site ($r=0.93$; $p:0.02$, $n=5$); However,
305 the IHC antibody used (EPR1161, Abcam, Cambridge, Massachusetts) was not one of the approved
306 companion assays developed with immune checkpoint inhibitors.

307 An important finding from our study is that older age was associated with a higher chance of a
308 failed PD-L1 assessment. EBUS-TBNA has previously been shown to have an excellent safety
309 profile coupled with an excellent yield for malignancy in older subjects^{26,27}. Our data suggest that

310 specimen quality may be inferior in the older patient, perhaps reflecting the challenges of obtaining
311 sufficient diagnostic material in this important group of patients.

312

313 *Clinico-pathological features of PD-L1 expression*

314 In this large multicenter study, we report the novel findings that PD-L1 expression is associated
315 with higher N-stage and the presence of brain metastasis. Previous studies have shown conflicting
316 results. Shimoji et al²⁸ reported that in 220 patients undergoing surgical resection, PD-L1
317 expression was correlated with younger age, smoking habit and solid pattern in adenocarcinoma
318 subjects, while multivariate analysis however revealed that only the solid adenocarcinoma subtype
319 was an independent predictor of PD-L1 expression. This study however was limited by fact that
320 only patients with early stage disease were included and all samples were from surgical resections.
321 In another article²⁹ using the E1L3N assay in 297 patients, the authors found that PD-L1 expression
322 on tumor cells was higher in men ($p < 0.0001$), older ($p = 0.0321$), smokers ($p < 0.0001$), high
323 histologic grade ($p = 0.0012$) and squamous cell histotype ($p = 0.0412$) patients. More recently, a
324 larger retrospective cohort study of 2402 surgically resected stage I-III NSCLC patients found that
325 PD-L1 positivity was more frequent in never smokers, higher disease stages and larger tumors³⁰. In
326 this study, PD-L1 expression in adenocarcinoma patients was associated with better clinical
327 outcomes (OS, time to relapse and relapse-free survival), though these data are heterogeneous
328 among the previous published papers³¹⁻³³.

329 Our study confirms findings of a recent metanalysis³⁴ showing that PD-L1 expression was
330 increased in patients with lymph node metastasis (OR = 1.34, 95% CI: 1.19–1.50, $P < 0.001$) and
331 TNM stage (OR = 1.45, 95% CI: 1.18–1.78; $P < 0.001$) but also for the first time that PD-L1 strong
332 positive patients were more likely to have brain metastases. These data are, of great interest as the
333 immune checkpoint inhibitors clinical trials excluded patients with presence of untreated or
334 unstable brain metastasis^{1-4,35} and the management of patients with NSCLC and brain metastases is
335 evolving³⁶.

336

337 *Study limitations*

338 To our knowledge, this is the largest study to assess the feasibility of PD-L1 testing using different
339 modes of tissue acquisition and provides multi-center real world data. However, there are several
340 limitations. First, PD-L1 expression, was evaluated by local pathology units only without any
341 control of inter-observer variability. However, the same approved assay was used in each center. No
342 specific assessment of concordance between EBUS specimens and surgical lymph node sampling
343 was planned in this study. This would have required surgical sampling of intra-thoracic lymph
344 nodes which is currently not standard practice in patients with advanced disease for whom
345 immunotherapy is currently licensed. The study included patients biopsied before the routine
346 approval of checkpoint inhibitors and many of the subjects who received immunotherapy were
347 within clinical trials. Thus, the small number of patients treated with immune checkpoint antibodies
348 did not allow any further consideration of factors that may predict response to immunotherapy.

349

350 **Conclusions**

351 EBUS-TBNA represents an important investigation for tissue acquisition in patients with lung
352 cancer as well as for lymph node staging. In this multicenter study, we have demonstrated that
353 EBUS-TBNA allows adequate sampling for testing PD-L1 in a broad population of NSCLC
354 patients. We have also reported that patients with advanced N-stage and brain metastasis are more
355 likely to express high levels of PD-L1. Finally, these data provide evidence that EBUS-TBNA
356 samples are suitable for complete molecular profiling, including PD-L1 testing, to allow decisions
357 regarding treatments and clinical trial eligibility to be made.

358

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366

367 Authors Contribution

368 Each author contributed to the study as follows:

369 Conception and design of the work (NN, SMJ, DRB, RB, IW, ADL, KK); Data acquisition (FP, AdB,
370 UM, ME, SJ, LY) ; Statistical Analysis (MN); Drafting the work or revising it critically (FP, MN, ADL,
371 IW, MM, ME, RB, SMJ, KK, AnB, LY, NN). NN is the guarantor of the paper, taking responsibility for
372 the integrity of the work as a whole, from inception to published article.

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TABLES

Table 1. Baseline characteristics.

		All patients N=577
Age	Median	68
	IQR	61 – 74
	Range	31 – 96
Sex	Male	318 (55%)
	Female	259 (45%)
Ethnicity	Caucasian	429 (88%)
	Other	57 (12%)
	Missing	91
Smoking status	Current	142 (27%)
	Former	312 (59%)
	Never	77 (15%)
	Missing	46
Pack years (those with smoking status data only)	0	77 (15%)
	<20	60 (11%)
	20+	298 (56%)
	Missing (but >0)	96 (18%)
Performance status	0	122 (25%)
	1	262 (53%)
	2	67 (14%)
	3	31 (6%)
	4	9 (2%)
	Missing	86
T-stage	1	85 (16%)
	2	168 (32%)
	3	125 (24%)
	4	151 (29%)
	Missing	48
N-stage	0	121 (23%)
	1	63 (12%)
	2	202 (38%)
	3	142 (27%)
	Missing	49
M-stage	0	218 (42%)
	1	307 (58%)
	Missing	52

		All patients N=577
Histology	Adenocarcinoma	378 (66%)
	Squamous	151 (26%)
	Other	48 (8%)
Sampling method	EBUS/EUS	189 (33%)
	Endobronchial biopsy	72 (12%)
	CT guided biopsy	167 (29%)
	Surgical	124 (21%)
	Pleural	6 (1%)
	Other	19 (3%)
Actionable mutation	ALK	7 (1%)
	EGFR	41 (7%)
	HER-2	2 (<1%)
	ROS1	1 (<1%)
	None	526 (91%)
Brain metastases	No	381 (84%)
	Yes	70 (16%)
	Missing	127
Received radiotherapy	No	310 (58%)
	Yes	221 (42%)
	Missing	46

Table 2. PDL1 assessment success rate.

		N=577
Overall	Overall	548/577 (95%)
Age group*	<60	3/130 (2%)
	60-69	8/187 (4%)
	70-79	13/203 (6%)
	80+	5/57 (9%)

NB. No difference according to sex, ethnicity (caucasian vs others), smoking status, pack years, performance status, T-stage, N-stage, M-stage, Histology, EGFR, biopsy type (original vs re-biopsy), presence of brain metastases, receipt of radiotherapy. Sampling method is non-significant if the "other" group is excluded. Failure rate among the "other" group is 21% (4/19), significantly higher than the other methods. *Odds ratio for age as a continuous variable is 1.06 (p-value=0.008).

Table 2. PDL1 assessment success rate.

		Fail	P-value
Overall	Overall	548/577 (95%)	n/a
Age	<60	3/130 (2%)	0.008*
	60-69	8/187 (4%)	
	70-79	13/203 (6%)	
	80+	5/57 (9%)	
Sex	Male	19/318 (6%)	0.248

		Fail	P-value
	Female	10/259 (4%)	
Ethnicity	Caucasian	0/57 (0%)	0.061
	Other	25/429 (6%)	
Smoking status	Current	3/77 (4%)	0.604 ^s
	Former	16/312 (5%)	
	Never	8/142 (6%)	
Pack years (those with smoking status data only)	0	3/77 (4%)	0.511 ^s
	<20	6/60 (10%)	
	20+	13/298 (4%)	
Performance status	0	4/122 (3%)	0.086 ^s
	1	15/262 (6%)	
	2	5/67 (7%)	
	3	3/31 (10%)	
	4	1/9 (11%)	
T-stage	1	5/85 (6%)	0.949 ^s
	2	7/168 (4%)	
	3	8/125 (6%)	
	4	7/151 (5%)	
N-stage	0	6/121 (5%)	0.254 ^s
	1	2/63 (3%)	
	2	7/202 (3%)	
	3	11/142 (8%)	
M-stage	0	13/218 (6%)	0.276
	1	12/307 (4%)	
Histology	Adenocarcinoma	23/378 (6%)	0.253
	Squamous	4/151 (3%)	
	Other	2/48 (4%)	
Sampling method	EBUS/EUS	10/189 (5%)	0.098 ^s
	Endobronchial biopsy	2/72 (3%)	
	CT guided biopsy	12/167 (7%)	
	Surgical	1/124 (1%)	
	Pleural	0/6 (0%)	
	Other	4/19 (21%)	
Actionable mutation	Any	3/51 (6%)	0.769
	None	26/526 (5%)	

		Fail	P-value
EGFR mutation	No	26/536 (5%)	0.486
	Yes	3/41 (7%)	
Brain metastases	No	22/381 (6%)	0.129
	Yes	1/70 (1%)	
Received radiotherapy	No	20/310 (6%)	0.089
	Yes	7/221 (3%)	

^s For p-value calculation, the “other” group is excluded.

*P-value calculated treating factor as a continuous variable.

^sP-value calculated using test for trend.

Table 3. Association with strong PDL1 expression

Factor		None/weak	Strong	P-value
Age	Median	68	67	0.249
	IQR	61-74	59-73	
Sex	Male	215 (72%)	84 (28%)	0.913
	Female	178 (71%)	71 (29%)	
Ethnicity	Caucasian	294 (73%)	110 (27%)	0.140
	Non-Caucasian	36 (63%)	21 (37%)	
Smoking status	Current	93 (69%)	41 (31%)	0.653
	Former	217 (73%)	79 (27%)	
	Never	55 (74%)	19 (26%)	
Pack years	0	55 (74%)	19 (26%)	0.586
	<20	41 (76%)	13 (24%)	
	20+	200 (70%)	85 (30%)	
Performance status	0	87 (74%)	31 (26%)	0.656
	1	176 (71%)	71 (29%)	
	2	42 (68%)	20 (32%)	
	3	19 (68%)	9 (32%)	
	4	4 (50%)	4 (50%)	
T-stage	1	58 (73%)	22 (28%)	0.942
	2	118 (73%)	43 (27%)	
	3	84 (72%)	33 (28%)	
	4	101 (70%)	43 (30%)	
N-stage	0	90 (78%)	25 (22%)	0.024
	1	49 (80%)	12 (20%)	

Factor		None/weak	Strong	P-value
	2	137 (70%)	58 (30%)	
	3	83 (63%)	48 (37%)	
M-stage	0	157 (77%)	48 (23%)	0.031
	1	200 (68%)	95 (32%)	
Histology	Adenocarcinoma	241 (68%)	114 (32%)	0.023
	Squamous	116 (79%)	31 (21%)	
	Other	36 (78%)	10 (22%)	
EGFR mutation	No	366 (72%)	144 (28%)	0.925
	Yes	27 (71%)	11 (29%)	
Any mutation	No	359 (72%)	141 (28%)	0.887
	Yes	34 (71%)	14 (29%)	
Re-biopsy	No	275 (72%)	107 (28%)	0.910
	Yes	75 (71%)	30 (29%)	
Sampling method	EBUS/EUS	120 (67%)	59 (33%)	0.073
	Endobronchial biopsy	49 (70%)	21 (30%)	
	CT guided biopsy	118 (76%)	37 (24%)	
	Surgical	95 (77%)	28 (23%)	
	Pleural	4 (67%)	2 (33%)	
	Other	7 (47%)	8 (53%)	
Brain metastases	No	266 (74%)	93 (26%)	<0.001
	Yes	35 (51%)	34 (49%)	
Received radiotherapy	No	215 (74%)	75 (26%)	0.118
	Yes	145 (68%)	69 (32%)	

Table 4. Response to immunotherapy

Immunotherapy treatment line	Response	All patients (56)
1 st line	PR	7 (28%)
	Stable	10 (40%)
	PD	8 (32%)
2 nd line	PR	2 (8.3%)
	Stable	7 (29.2%)
	PD	15 (62.5%)
3 rd or more	PR	1 (14.3%)
	Stable	1 (14.3%)
	PD	5 (71.4%)