



Prolonged immune checkpoint inhibitor response of a patient with a RET rearranged non-small cell lung cancer and high tumour mutational burden: case report and review of the literature

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Abstract: Patients with oncogenic driven metastatic non-small cell lung cancer (NSCLC) represent a special subgroup of metastatic NSCLC, as for several oncogenic drivers, tyrosine kinase inhibitors (TKI, either registered or within a trial) are available. One of the more challenging drivers to target is the rearranged during transfection (RET) fusion, initially found in thyroid cancers, but also present in 1–2% of NSCLC patients. Initially, RET was targeted with multi-kinase inhibitors, with fairly disappointing results (generally no long-term responses and high rates of adverse effects). Selective TKIs are currently being investigated, and results are promising. Limited data is available regarding immune checkpoint inhibitor (ICI) efficacy in NSCLC patients with a RET rearrangement. We present a case of a male patient with stage IV NSCLC with a RET gene rearrangement, with low programmed death-ligand 1 (PD-L1) expression, but high tumour mutational burden (TMB), responding to second line nivolumab. To put these data into perspective, we also review the literature regarding treatment of NSCLC patients with a RET gene rearrangement with either TKIs or ICI.

Keywords: Non-small cell lung cancer (NSCLC); RET rearrangement; immune checkpoint inhibitors (ICIs); selective tyrosine kinase inhibitors (selective TKIs)

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Introduction

Immune checkpoint inhibitors (ICI) have become the new standard of care (SoC) in the treatment of metastatic non-small cell lung cancer (NSCLC) patients. First, in second line and beyond, the programmed death-1 (anti-PD-1) antibodies nivolumab (registration regardless of PD-L1 status) and pembrolizumab (for PD-L1 positive patients) became available, followed by the anti-PD-L1 antibody atezolizumab (registration regardless of PD-L1) (1). Recently, ICI have also become SoC in the first line treatment of patients having a metastatic NSCLC without an activating epidermal growth factor receptor (*EGFR*) mutation or anaplastic lymphoma kinase (*ALK*)

rearrangement. Pembrolizumab monotherapy is registered as first-line therapy for patients with a PD-L1 tumour proportion score (TPS) $\geq 50\%$ (Europe) or $\geq 1\%$ (USA) (1-5). Combinations of platinum-based chemotherapy and anti-PD-(L)1 inhibitors (pembrolizumab or atezolizumab, the latter combined with bevacizumab) have become SoC treatment regardless of PD-L1 status.

Data is limited regarding ICI efficacy in those with targetable oncogenic-driven advanced NSCLC. Markers generally associated with ICI benefit are PD-L1 expression and tumour mutational burden (TMB). However, PD-L1 is often constitutively expressed in tumours with an oncogenic driver, and correlation with response is often poor (5,6). In general, TMB is low in non-smoking related

oncogenic drivers, such as *EGFR* and *ALK* (6,7). Most ICI efficacy data come from patients with *EGFR* mutated NSCLC and to a lesser extent from patients with mutations in more rare driver genes (e.g., *ALK*, ROS proto-oncogene 1 (*ROS1*), rearranged during transfection (*RET*) fusion, ERBB2 receptor tyrosine kinase 2 [*ERBB2*, also known as *HER2*), BRAF serine/threonine kinase proto-oncogene (*BRAF*) and MET proto-oncogene (*MET*)]. For these rare driver genes, mainly retrospective data exist (6). In general, in NSCLC of non-smokers harbouring oncogenic driver mutations, responses are low and long-term outcome is poor (8). Based on a subgroup analysis of the IMpower 150 trial (n=80 tumours with sensitizing *EGFR* mutations, n=34 with *ALK* rearrangements) it is stated in the European Society for Medical Oncology (ESMO) metastatic NSCLC guideline that *EGFR* mutated and *ALK* rearranged tumour patients can be treated with a combination of chemotherapy and ICI (platinum doublet/bevacizumab/atezolizumab) after exhaustion of all available TKI's (1,9). BRAF V600 mutated patients can be treated with ICI (with or without chemotherapy) upon progression on dabrafenib/trametinib. For the rarer oncogenic drivers, no recommendation regarding ICI has been made (1).

The preferred first line treatment for a patient with NSCLC containing a targetable oncogenic driver mutation (*EGFR*, *ALK*, *ROS1*, *BRAF*) is a (registered) tyrosine kinase inhibitor (TKI). For other driver gene mutations, TKIs are only available in a clinical trial and recruitment into open trials is encouraged (1).

An example of a rare oncogenic driver, for which no TKIs have been approved yet, is the *RET* gene rearrangement, although several TKIs are under investigation. Choosing the right treatment may be challenging in these patients, especially when no RET-TKI is available within a clinical trial.

We present a case of a patient with a stage IV NSCLC with a low PD-L1 expression, high TMB and a *RET* gene rearrangement, as identified by fluorescence *in situ* hybridisation (FISH) analysis, who progressed after first line chemotherapy. We will describe the challenges in choosing the best second line treatment for this patient. After description of the case, we will summarize the current evidence for ICI in *RET* gene rearranged positive NSCLC, as well as the literature for RET-TKIs.

We present the following article in accordance with the CARE reporting checklist (available at <http://dx.doi.org/10.21037/pcm-20-26>).

Case presentation

A 65-year old man, diagnosed with a cT4N3M1b stage IVB (TNM7) poorly differentiated NSCLC (differentiation between adenocarcinoma and squamous cell carcinoma initially not possible), was referred in March 2017 to our clinic for a second opinion regarding treatment options because of his recently diagnosed NSCLC. He had no relevant medical history, except for a history of smoking (100 pack years). At initial diagnosis, he had several bone and liver metastases. At that time, despite two attempts to obtain enough tissue, molecular analysis was not possible due to poor DNA quality in the obtained tissue samples. Liquid biopsy for molecular analysis was not available in the clinic at that time. After discussion with the patient, no third attempt to obtain tissue for molecular analysis was performed. First line clinical trials were not enrolling, and first line treatment with gemcitabin/cisplatin was started, with stable disease after four cycles (August 2017). In September 2017 he presented with disease progression, and gave consent to obtain a new biopsy for molecular analysis, as well as a biopsy for the CPCT-02 trial (NCT01855477). This is a trial in which fresh biopsies and blood samples are collected from patients with metastatic cancer, which are then analysed by whole genome sequencing (WGS) to obtain a mutational profile. The histological biopsy revealed a NSCLC, subtype adenocarcinoma, TTF-1 positive. The in-hospital molecular analysis showed no *EGFR*, *KRAS*, *BRAF* and *MET*-exon 14 skipping mutations (tested with PCR and pyrosequencing). PD-L1 [immunohistochemistry (IHC), 22C3 clone] expression in tumour cells was less than 1%. FISH showed no *ROS1* nor *ALK* rearrangement and no *MET* amplification was detected. FISH analysis for the detection of RET gene rearrangements was performed using the Vysis 10q11 RET Break-Apart FISH Probe (Abbott Laboratories, Chicago, Illinois, USA), which revealed loss of the green probe signal, located at the 5' centromere side of the common RET translocation breakpoint, in ~58% of the tumour cells, while the red probe signal at the 3' telomere side of the gene (near the tyrosine kinase domain) was maintained. The latter indicates the presence of a *RET* gene rearrangement. At that time, no clinical trials involving RET-TKI's were available in which the patient could be included. In October 2017, it was therefore decided to start ICI (nivolumab) as a second line therapy. During treatment, WGS results from the CPCT-02 trial became available, and it turned out that the patient's tumour material had a high TMB, as indicated

Timeline of diagnosis and treatment

Diagnosis stage IVb NSCLC	Start palliative chemotherapy (gemcitabin/cisplatin)	Stable disease after 4 cycles of chemotherapy	Progressive disease	Start Nivolumab	Stop nivolumab	Follow-up
March 22, 2017	May 2, 2017	August 3, 2017	September 1, 2017	October 11, 2017	August 8, 2019	Ongoing (June 24, 2020)

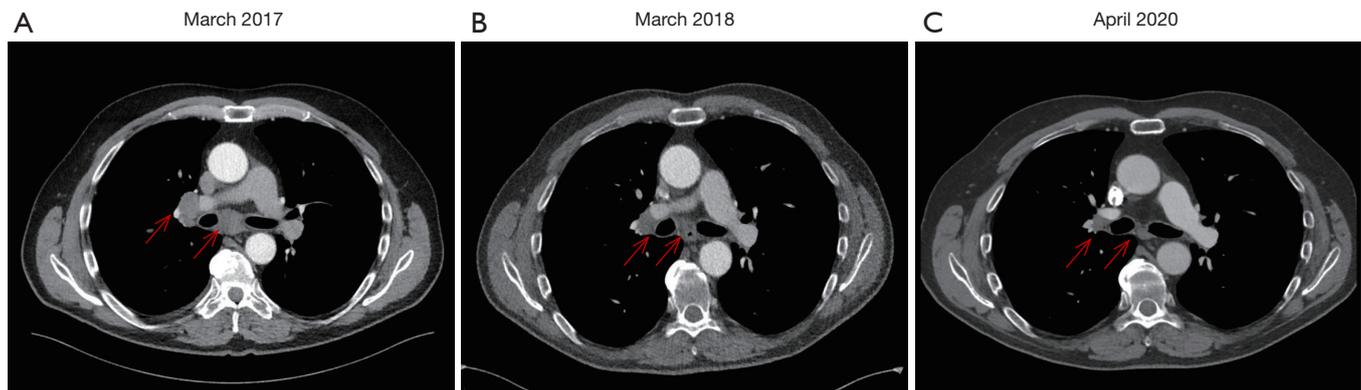


Figure 1 Timeline picture. Red arrows: (A) lymphadenopathy at diagnosis; (B) 5 months after start nivolumab treatment; (C) 20 months after stop immunotherapy.

by the presence of 527 non-synonymous somatic missense mutations per whole genome DNA (cut-off for high TMB: >140) (10). Among the detected genomic variants, a single frame shift mutation located in the RB1 gene [c.1768delT, p.(Cys590fs)] was identified, and considered pathogenic based on the mutation type, as well as the detected variant allele frequency in relation to the tumour cell percentage (~40%). The mutation had a variant allele frequency of 36%, most likely suggestive for loss of heterozygosity (LOH) of the other RB1 allele. After 7 cycles of nivolumab the CT-scan showed a nearly complete remission. After 2 years of treatment, the nivolumab was discontinued because of a persistent grade 2 arthritis, despite treatment with methotrexate and low dose corticosteroids. Nine months after discontinuation of nivolumab, the patient still has an on-going response (disease course is depicted in *Figure 1*).

All procedures performed in studies involving human participants were in accordance with the ethical standards of the institutional and national research committees and with the Helsinki Declaration (as revised in 2013). Written informed consent was obtained from the patient for publication of this manuscript and any accompanying images.

Discussion

Here, we discuss a case of a metastatic NSCLC patient,

progressing on first line chemotherapy, and presentation with a RET translocation, low PD-L1 and high TMB in 2017. The patient obtained durable benefit with ICI.

How can we put the data of this patient into perspective, with the currently available knowledge?

RET detection

The *RET* gene was identified in 1985, and encodes a receptor tyrosine kinase consisting of three domains: extracellular, transmembrane, and intracellular. RET activation can occur through mutations and fusions. Activation of RET leads to auto phosphorylation of RET and downstream cell signalling through intracellular pathways such as mitogen-activated protein kinase (MAPK), phosphoinositide 3-kinase (PI3K)/AKT and Janus kinase (JAK)/signal transducer and activator of transcription (STAT) pathways which leads to uncontrolled cellular proliferation (11,12).

RET rearrangements were initially found in thyroid cancers, but also occur in 1–2% of NSCLC patients. In NSCLC, RET fusions are more commonly involved than single nucleotide variations (SNVs). In NSCLC at least 12 fusion partner genes have been identified so far, most common are the KIF5B-RET and CCDC6-RET (11). In contrast with our case, RET rearrangements are mainly observed in younger patients and never or light smokers

(13,14). Consistent with our case, RET rearrangements are usually found in adenocarcinoma or in poorly differentiated tumours. Furthermore, RET point mutations and fusions are found in medullary and papillary thyroid carcinoma, spitzoid neoplasms, pancreatic cancer, ovarian cancer and colorectal cancers (15-17).

In general, different methods exist for detecting genomic rearrangements, each method having its advantages and drawbacks with respect to sensitivity, specificity, hands-on time and costs. Whereas ALK and ROS1 rearrangements are commonly detected using IHC and/or FISH, RET translocations cannot be adequately detected by IHC, as currently no validated antibody is available for diagnostic application (11). Based on the break-apart probe signal ('classic pattern'), FISH allows detection of gene rearrangements, however, no fusion genes can be detected or specified. As illustrated in our study, in some cases a loss of the 5' probe signal (centromere side) might be observed, while the 3'probe signal (telomere side) is maintained. Such probe pattern suggests the presence of a potential RET gene rearrangement or deletion, but does not clarify whether a RET gene fusion is present, resulting in an active fusion transcript (18). Moreover, with respect to the ROS1 translocation FISH, it was previously shown that the presence of isolated extra 3' probe signal can lead to false positive FISH interpretation. By comparing ROS1 FISH with IHC and RNA sequencing, it was shown that in 10 cases FISH analyses indicated a translocation based on the presence of isolated 3'probe signal, whereas the IHC as well as RNA sequencing outcomes were negative (19). RNA sequencing allows detection of gene rearrangements that result in a fusion-transcript, and therefore is a reliable approach to characterize genomic rearrangements that are likely to result in oncogenic events. At the time of WGS sequencing in our case, the WGS bioinformatics pipeline was not suitable for detecting large genomic rearrangements, and therefore no such genomic alterations were reported. Currently, different commercial NGS RNA fusion gene panels exist (e.g., Archer FusionPlex, Illumina Trusight RNA panels, Oncomine Comprehensive assays, QuantideX NGS RNA assay), which in contrast to the sequencing of a complete transcriptome, allow detection of a limited number of target gene-transcripts in a relatively cost effective manner. RNA sequencing libraries can be generated based on hybrid capture enrichment, multiplex amplicons or anchored multiplex PCR, where sequencing methodologies differ in their ability to detect RNA fusion transcripts and characterize known and novel gene fusion

partners (20). Yet, characterization of the specific gene fusion partner could be of significant value in near future, when highly selective RET inhibitors might become the new standard in treatment. As an example, with respect to crizotinib treatment in NSCLC patients with an EML4-ALK fusion, it was shown that even the specific variant of the EML4-ALK fusion transcript (variant 3/5 versus other EML4-ALK fusion transcripts) influenced progression free survival and overall survival (21).

The frame shift mutation in the RB1 gene [c.1768delT, p.(Cys590fs)], as detected in the DNA of our patient, could have therapeutic consequences when considering inclusion in ongoing RET-TKI trials, as RB1 functions downstream of RET, suppressing cell-cycle progression.

Treatment of RET rearranged NSCLC

Developing a TKI for treatment of RET rearranged patients has been challenging, as is described below.

Ret inhibitors

Initially, multi-target TKIs, such as vandetanib, cabozantinib, lenvatinib, alectinib, sunitinib, ponatinib, nintedanib, regorafenib and sorafenib were evaluated for the treatment of RET-rearranged NSCLC patients (22,23).

The first documented results of RET-TKI therapy were with cabozantinib in 2013 (n=3) (24). Two patients experienced a partial response (PR) and the third had prolonged stable disease (SD). The phase II trials in NSCLC, with cabozantinib (25), lenvatinib (26) and vandetanib (27,28) (*Table 1*), showed ORR of 16–53%, a median PFS of 4.5–7.3 months and a median OS of 9.9–11.6 months. For the other multi-kinase inhibitors used in the treatment of RET-rearranged NSCLC even more limited data is available. Unfortunately, multi-kinase inhibitors are associated with high rates of adverse effects (AEs) due to their anti-VEGFR (hypertension, proteinuria) and anti-EGFR activity (rash, diarrhoea). In 70% of the cases treatment discontinuation or dose reductions because of toxicity were necessary.

The new selective RET inhibitors, selpercatinib (LOXO-292) and pralsetinib (BLU-667) are much more promising and showed impressive activity with a low percentage of severe AE (SAEs) in phase I/II trials. Selpercatinib is being investigated in the LIBRETTO-001 trial for RET rearranged solid tumours. Preliminary data of 105 NSCLC patients show an ORR of 68% (95% CI: 58–76) in the primary analysis and a median PFS of 18.4 months

Table 1 Multi-kinase inhibitors (MKI) selective tyrosine kinase inhibitors (TKI)

Author, year	Histology	Phase trial	RET inhibitor	Number of NSCLC patients	ORR % (95% CI)	mPFS, months (95% CI)	mOS, months (95% CI)
MKI's							
Drilon <i>et al.</i> 2016	NSCLC	II	Cabozantinib	26	28 (12 to 49)	5.5 (3.8 to 8.4)	9.9 (8.1 to not reached)
Yoh <i>et al.</i> 2017 (LURET)	NSCLC	II	Vandetanib	19	53 (28 to 77)	4.7 (2.8 to 8.5)	11.1 (9.4 to not reached)
Hida <i>et al.</i> 2017	NSCLC	II	Lenvatinib	25	16 (N/A)	7.3 (3.6 to 10.2)	Not evaluable (5.5 to not reached)
Lee 2017	NSCLC	II	Vandetanib	17	18 (N/A)	4.5 (N/A)	11.6 (N/A)
Selective TKI's							
Drilon <i>et al.</i> 2019 (LIBRETTO-001)	NSCLC, TC	I/II	Selpercatinib	105	68% (58 to 76)	18.4 (12.9–24.9)	N/A
Gainor <i>et al.</i> 2019 (ARROW)	NSCLC, TC, other	II	Pralsetinib	120	58% (N/A)	N/A	N/A

RET, rearranged during transfection; ORR, objective response rate; CI, confidence interval; mPFS, median progression free survival; mOS, median overall survival; NSCLC, non-small cell lung cancer; N/A, not available; TC, thyroid cancer.

(95% CI: 12.9–24.9). Anti-RET activity was seen regardless of RET fusion partners, previous chemotherapy or cerebral metastasis. The ORR for cerebral metastasis was 91% (95% CI: 59–100). The tolerability was good with mostly grade 1–2 AEs (diarrhoea, dry mouth) with a discontinuation rate of 1.7% (17). Pralsetinib is being investigated in the ARROW study. Preliminary findings report an ORR of 58% (n=120), regardless of RET fusion partners and previous treatment of multi-kinase inhibitors. 40% of the patient had baseline brain metastases and pralsetinib led to a response of the brain metastasis in 7 out of 9 patients. Most of the AEs were grade 1–2 (increased AST/ALT, hypertension, constipation and fatigue) (29). RXDX-105 is a VEGFR-sparing, multikinase inhibitor with activity against RET and to a lesser degree against wild-type BRAF and BRAF V600E. Because RXDX-105 is VEGFR-sparing it has lower toxicity compared to other multikinase inhibitors, resulting in less dose modifications and/or discontinuation. In a phase I/Ib study ORR was 67% in those with non-KIF5B RET partners, while KIF5B RET fusions showed no response. Most common toxicities were fatigue, diarrhoea, hypophosphatemia and rash (30). Recently the FDA approved selpercatinib for the treatment of patients with metastatic RET fusion-positive NSCLC, metastatic RET-mutant medullary thyroid cancer (MTC) and metastatic RET fusion-positive thyroid cancer (31).

Although these selective *RET* inhibitors show promising results, acquired resistance will eventually occur and can limit long-term efficacy. *RET* solvent front mutations (G810R, G810S, G810C), evaluated in circulating tumour DNA (ctDNA), have recently been found to cause acquired resistance to selpercatinib and pralsetinib (23). Comparable with acquired EGFR and ALK mutations in TKI treated EGFR-mutated/ALK rearranged patients, these mutations were detected before clinical or radiographic progression (32). To overcome resistance to selpercatinib and pralsetinib, new inhibitors that have both activity against gatekeeper and solvent front mutations are being developed and investigated (33).

Ongoing or planned first line studies in RET rearranged NSCLC are for example the randomized phase III LIBRETTO-431 (NCT04194944, selpercatinib versus SoC platinum-doublet chemotherapy +/- pembrolizumab) and the randomized, open-label, phase III first AcceleRET Lung Study (NCT04222972, pralsetinib versus SoC platinum-doublet chemotherapy +/- pembrolizumab). The single arm phase II ALERT study (NCT03445000) is evaluating the activity of alectinib in pre-treated NSCLC patients. The single arm phase 2 LIBRETTO-321 study (NCT04280081) investigates the effectiveness of selpercatinib in patients with advanced solid tumours, including *RET* Fusion. The phase 1/2 study of TPX-0046 (NCT04161391),

Table 2 RET and immunotherapy

Author, type of study	RET+ patients (N)	PD-L1+	TMB (median) and type of test used	ORR in RET	mPFS (months, 95% CI)	mOS (months, 95% CI)	mTTP (months, 95% CI)
Mazieres <i>et al.</i> (retrospective)	16	6/16, mean 26% (0–80%)	N/A	6%	2.1 (1.3–4.7)	21.3 (3.8–28)	N/A
Offin <i>et al.</i> (retrospective)	74	0% in 58% (15/26); 1–49% in 23% (6/26); ≥50% in 19% (5/26)	44 patients with known TMB (MSK-IMPACT); median TMB: 1.75 mut/Mb (range, 0–9.65)	13 patients, 0%	3.4 months (2.1–5.6)	OS compared between ICI naïve (n=46) and ICI (n=16): no difference (HR 1.4, 95% CI: 0.7–2.9)	N/A
Hegde <i>et al.</i> (retrospective, abstract only)	72 (19 ICI, 53 non-ICI); 29 NSCLC	Available for 7 patients: 5/7 PD-L1 ≥1%	N/A	N/A	N/A	N/A	10.5 (6.7–30.1); HR: ICI vs. non-ICI 1.73 RETf and 8.0 RETm
Guisier <i>et al.</i> (retrospective)	9	Positive in 33%; >50% in 22%; negative 56%; unknown in 11%	N/A	37.5%	7.6 (2.3 to NR)	NR (26.8 to NR)	N/A

N, number; PD-L1, programmed death ligand 1; RET, rearranged during transfection; patients; patients; TMB, tumor mutational burden; ORR, objective response rate; mPFS, median progression free survival; CI, confidence interval; mOS, median overall survival; mTTP, median time to progression; N/A, not available; NR, not reached; ICI, immune checkpoint inhibitor; HR, hazard ratio; NSCLC, non-small cell lung cancer; RETf, RET fusions; RETm, RET mutations.

a RET/SRC inhibitor in solid tumours with a RET rearrangement is enrolling. A phase 1 study with RXDX-105 (NCT03784378) for the treatment of patients with NSCLC harbouring a *RET* Fusion and patients with ovarian cancer harbouring a *BRAF* mutation is planned but not recruiting yet. BOS172738 (NCT03780517) is being investigated in a phase I study including patients with NSCLC and Medullary Thyroid Cancer (MTC) with RET rearrangements. The LUNG-MAP Sub-Study, NCT04268550, is a phase II study evaluating the activity of selpercatinib in metastatic or recurrent NSCLC patients with *RET* Fusion. Cabozantinib (NCT01639508) is still being investigated in a phase II trial including NSCLC patients with *RET* fusion, ROS1 or NTRK Fusions or increased MET or AXL Activity. Last, cabozantinib is also being investigated in the phase II CRETA trial (NCT04131543) enrolling pre-treated, advanced *RET*-rearranged NSCLC patients.

RET-immunotherapy

Limited data is available regarding the immune environment and ICI efficacy in RET positive NSCLC (Table 2) (8,23, 34–36). In a retrospective study including 74 patients with RET rearranged lung cancer, the expression of PD-L1 (22C3 clone) was lower than 50% in 21 of 26 patients (81%), with a PD-L1 expression of 0% in 58% (n=15/26) cases and PD-L1 expression of 1–49% in 23% (n=6/26). In 44 patients, TMB was determined with MSK-IMPACT assay: median TMB was 1.75 mutations/Mb, which was significantly lower than the median TMB of RET wild-type NSCLCs, 5.27 mutations/Mb (n=3,631). The RET rearranged patients showed no response to ICI, and the median PFS was 3.4 months (34). In the six RET rearranged NSCLC patients included in the Immunotarget cohort, the median percentage of PD-L1 expression was 26 (range, 0–80%) (8). The ORR was only 6%, median PFS was 2.1 months. Another study included 785 patients with NSCLC who underwent surgery, all were tested for oncogenic drivers and PD-L1 expression. None of the six RET rearranged patients had a PD-L1 expression of more than 1% (37). A retrospective study of Guisier *et al.* showed 9 patients with RET-arrangements, with a PD-L1 expression of >50% in 2 patients (22%) and a negative PD-L1 status in 5 patients (56%). This study showed higher response rate in patients with a RET rearrangement (36). These limited data suggest that generally, patients with RET rearranged NSCLC have both low PD-L1 expression and low TMB, and do not respond well to ICI monotherapy,

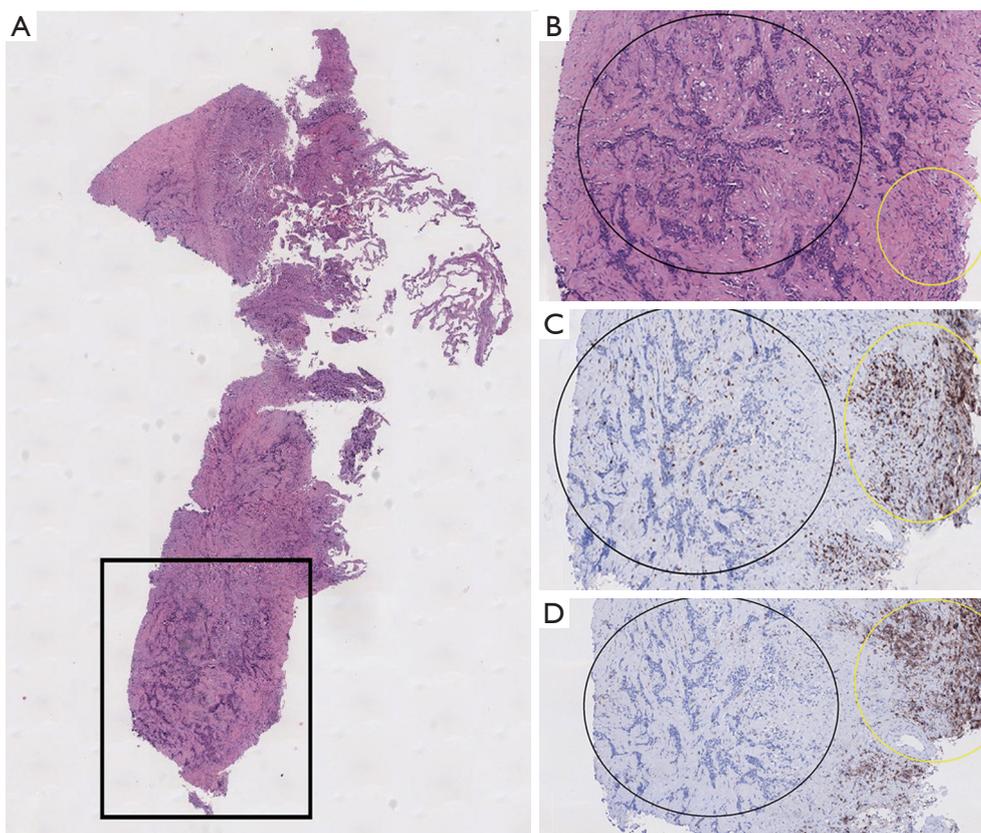


Figure 2 Lung biopsy. (A) Overview of lung biopsy (HE, $\times 20$). (B) Details of the inset of A (HE, $\times 100$). Black circle shows area of tumor and yellow circle shows inflammatory infiltration. (C,D) Immunohistochemical stains of CD45 (leucocyte common antigen) and CD3, respectively ($\times 100$). Inflammatory cells are stained brown. In the tumor area (black circle) there are only few discrete inflammatory cells, mainly composed of T lymphocytes. Yellow circle shows reactive inflammation which is not related to the tumor.

except in the small study of Guisier *et al.* Our patient indeed had a low PD-L1 level, but high TMB and obtained durable benefit with nivolumab. The high TMB could be related to his smoking status (100 pack years). Furthermore, high TMB is associated with ICI benefit (38), and this could be the explanation in our case too. To further evaluate the cancer immune environment in our patients, we evaluated the percentage of tumour infiltrating lymphocytes (TILs) in the sample obtained before start of nivolumab. TILs are associated with improved survival of NSCLC patients upon ICI treatment (39). However, the tumour tissue of our patient showed only a very low number of TILs (*Figure 2*). Detailed immune environment data are lacking for RET rearranged NSCLC. In general, it has been described that activated RET can activate the Wnt pathway which acts inhibitory on the recruitment of dendritic cells (17,40). Based on our case and the Guisier study, some patients with

RET rearranged NSCLC respond well to ICI, and more data are needed to identify this special subset.

The role of ICI combined with chemotherapy in RET rearranged NSCLC is not clear, and hopefully ongoing clinical trials will shed some light. For example, the POSEIDON trial (NCT04322591) is evaluating chemotherapy with or without a PD-1 antibody in RET fusion positive NSCLC. Furthermore, the LIBRETTO-431 and AcceleRET Lung have in their SoC arm the option to combine platinum-doublet chemotherapy with pembrolizumab.

Conclusions

RET rearranged NSCLC has been difficult to treat for years with disappointing results of MKIs. However, the new selective RET TKIs are very promising in early

phase clinical trials with high ORR and durable responses, combined with low toxicity. Phase III trials are on-going. Tumours with RET fusions and mutations often express low PD-L1 and have low TMB and ICI monotherapy data are disappointing in this patient population. It is not clear yet whether combination therapies with chemotherapy and ICI are effective in RET rearranged NSCLC.

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