

Real-world evaluation of molecular testing and treatment patterns for EGFR mutations in non-small cell lung cancer in Latin America

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Abstract. Lung cancer is a leading cause of cancer-related deaths in Latin America, with non-small cell lung cancer (NSCLC) being the most prevalent. The current study aimed to report real-world data on epidermal growth factor receptor (EGFR) mutational testing and treatment regimens at diagnosis and progression in patients with metastatic NSCLC across four Latin American countries (Argentina, Chile, Colombia and Uruguay). A retrospective, multicenter,

observational study was conducted in patients with NSCLC using medical records from participating countries. The study population was categorized into two cohorts: Cohort 1 comprised of newly diagnosed, treatment-naïve patients with stage IV NSCLC; and cohort 2 comprised of stage IV NSCLC EGFR mutation (EGFRm)-positive patients who had progressed after first- or second-generation EGFR-tyrosine kinase inhibitor (TKI) treatment. Measures included demographic variables, health characteristics, treatment regimen, molecular testing rate and turnaround time at diagnosis and at progression for cohorts 1 and 2, respectively. Descriptive statistics were used to summarize all study measures. Of the 462 patients enrolled, 431 were newly diagnosed or treatment naïve with metastatic NSCLC. In cohort 1, the majority of patients with private health insurance (57.31%) underwent molecular diagnosis while only 41.3% of patients within the public sector had access to testing. The average molecular testing rate in cohort 1 varied across countries, with Argentina having the highest testing rate (79%) and Uruguay the lowest (27.63%). EGFRm was observed in 22% of patients. Cohort 2 comprised 31 patients who had progressed after first- or second-generation EGFR-TKI treatment and of these, only 22 (70.97%) underwent testing after progression. Access to molecular testing is still a challenge impacting the choice of first-line treatment in Latin American patients with NSCLC.

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Abbreviations: NSCLC, non-small cell lung cancer; EGFR, epidermal growth factor receptor; TKI, tyrosine kinase inhibitor; PDL-1, programmed death-ligand 1; ALK, anaplastic lymphoma kinase; CNS, central nervous system

Key words: lung cancer, epidermal growth factor receptor, non-small cell lung cancer, Latin America, tyrosine kinase inhibitor

These findings underline the unmet needs of ensuring early diagnosis, molecular profiling and use of correct treatment to alleviate NSCLC burden in the region.

Introduction

Over the last century, lung cancer has advanced as a major global burden and the predominant cause of cancer-related deaths worldwide (1). According to the GLOBOCAN 2018 analysis, lung cancer contributed to 11.6% of the total diagnosed cancer cases, with a mortality rate of 18.4% (2). As expected, Latin America shows a similar pattern, with lung cancer being the leading cancer type in the region imposing a major public health issue. Among all cancers, lung cancer mortality ranks number one in Argentina and Uruguay and number two in Chile and Colombia (3).

The most prevalent form of lung cancer accounting for almost 85% of all lung cancers is non-small cell lung cancer (NSCLC) (4,5). In most cases, lung cancer is diagnosed at advanced or metastatic stage where surgical resection is not amenable and survival is low (6,7). According to a recent Economist Intelligence Unit report, apart from tobacco usage, late detection, and insufficient access to screening and treatment are the leading causes of lung cancer mortality in Latin America (3).

Screening techniques such as low-dose helical computed tomography and broad molecular profiling are key to reducing cancer-based mortality (8-10). The implementation of these techniques require well-structured diagnostic approaches and good public health systems. Platinum-based therapy has been the standard of care in advanced NSCLC. However, the discovery of new molecular alterations and development of respective targeted treatments represent a major improvement over conventional chemotherapy when applied to appropriately selected patient populations. Testing of mutation specific to epidermal growth factor receptor (*EGFR*) and *BRAF V600E*, rearrangement of *ROS1* genes (*c-ros* oncogene 1) and *ALK* (anaplastic lymphoma kinase) as well as alteration in programmed death-ligand 1 (*PDL-1*) expression are critical factors for advanced or metastatic NSCLC treatment strategy. Currently, targeted therapies and checkpoint inhibitors are being widely approved for these specific genetic aberrations. Furthermore, mutations in genes like *MET* exon 14 and *HER2* (human epidermal growth factor receptor 2) and fusion genes involving *RET* (rearranged during transfection) and *NTRK1* (neurotropic tropomyosin receptor kinase 1) are also evolving as important targets or biomarkers (11).

Apart from high mortality rates, lung cancer in Latin America is also associated with high direct and indirect costs (3,12). Although the Latin American countries have national treatment guidelines, real-world data on molecular testing and treatment pattern are limited (3). A lack of data on existing practices hinders identification of the actual requirements and effective economic management.

The current study presents real-world data on the management of patients with stage IV NSCLC in four Latin American countries including Argentina, Colombia, Chile, and Uruguay. This retrospective study aimed to evaluate the molecular testing and treatment patterns in a cohort of newly diagnosed, treatment-naïve patients with advanced or metastatic NSCLC

and in another cohort of EGFR mutation-positive patients who had progressed after EGFR-tyrosine kinase inhibitor (TKI) treatment.

Materials and methods

Study design. This retrospective, multicenter study was conducted in patients with stage IV NSCLC from the four Latin American countries including Argentina, Colombia, Chile, and Uruguay diagnosed from January 1, 2017 to December 31, 2017. All data were collected using patient medical records and transferred from patients' medical record book to a secure web-based electronic case report form by the physician or trained study personnel in accordance with outlined data entry procedures.

The study was performed in accordance with ethical principles that are consistent with the Declaration of Helsinki, International Conference on Harmonization-Good Clinical Practice (ICH GCPs), and the applicable legislation. The final study protocol was approved in writing by the Ethics Committee (EC) for each site in all the countries that participated.

Study population. The study population was categorized into two cohorts: cohort 1 comprised newly diagnosed and treatment-naïve patients aged ≥ 18 years with stage IV NSCLC; cohort 2 comprised stage IV NSCLC EGFR mutation-positive patients who had progressed after first- or second-generation EGFR-TKI treatment.

Furthermore, all included patients were required to provide written informed consent as per local regulations. There were no minimum reporting requirements for the medical record to be eligible other than those specified for each cohort. Data were collected post diagnosis of advanced or metastatic NSCLC in cohort 1 and post EGFR-TKI progression in cohort 2 up to maximum follow-up registered in the medical record book.

Measures. The demographic variables examined in the study included age, gender, smoking status, and health insurance status for both cohorts. The health characteristics reported include ECOG and histology type. Molecular testing rate, biopsy site, and turnaround time at diagnosis for cohort 1 and at progression for cohort 2 were reported. Molecular profiling including EGFR mutations and subsequent treatment strategies were followed for both cohorts.

Statistical analysis. No formal hypothesis testing was incorporated in the study. Descriptive statistics (mean, standard deviations for continuous variables and frequencies and percentages for categorical variables) was used to summarize all study measures.

Results

Demographic and clinical characteristics in the treatment-naïve cohort. Of the 462 patients enrolled in the present study, 431 (93.2%) were newly diagnosed or treatment naïve with advanced or metastatic NSCLC from Argentina ($n=195$), Colombia ($n=96$), Chile ($n=64$), and Uruguay ($n=76$). The mean age of patients in cohort 1 was 66 ± 11 years;

60% of the patients were men and 57.3% had private insurance coverage. Among all the study nations, Argentina had the highest private insurance coverage (87%) and Uruguay reported the lowest private insurance coverage (8%).

Most of the patients with NSCLC were found to either have a smoking history (41%) or were active smokers (37%), whereas a small proportion of patients (16.4%) had no smoking history. The highest percentage of current smokers among the four nations was observed in Uruguay (62%), followed by Argentina (39%) and Chile (28%) (Table I). The majority of patients in cohort 1 (71.0%) had adenocarcinoma histology. The metastatic sites reported were highest in bone (41.8%), followed by lymph nodes (36.7%), brain (20.4%), adrenal (13.7%), and liver (14.3%). Tumor characteristics were similar in almost all the countries, including high rates of bone metastasis and adenocarcinoma histology, except in Uruguay which had lower rates of adenocarcinoma.

Among all patients diagnosed in stage IV, Performance Status (PS) of almost half (48%) showed PS 0-1; 14% of the patients were PS 2, whereas only 7.7% of the patients were severely or completely disabled (PS 3 or 4). Most patients were diagnosed in advanced or metastatic stages (stage III: 9%; stage IV: 76%) and only 5% were diagnosed at early stages (I-II).

Molecular testing. On an average, the frequency of molecular testing was 66%, with Argentina having the highest testing rate (79%) with a population belonging primarily to the private sector (87%) and Uruguay having the lowest testing rate (28%) but with a population belonging primarily to the public sector (91%). The mean turnaround time for testing was 12.4 days, with Colombia reporting the longest turnaround time (Argentina, 10; Colombia, 20.4; Chile, 10; Uruguay, 13.4). Formalin fixed paraffin embedded tissue was commonly used for sample processing and further testing (55.68%; Argentina, 68.72%; Colombia, 46%; Chile, 64%; Uruguay, 27.6%). The percentage of samples collected as blood (2.3%), fresh frozen tissue (1.1%), or cytology (0.46%) was relatively less. Unknown sample collection procedures accounted for 40.1% of the patients. PDL-1 testing was performed in the majority of patients (89.1%) (Table II).

Molecular profile and treatment patterns. Among all the patients, the frequency of EGFR mutations in NSCLC was 22.2% (Argentina, 26%; Colombia, 26%; Chile, 8.3%; Uruguay, 19.0%). The most frequent EGFR mutation was exon 19 deletion (47%), followed by exon 21 L858R (36%). The frequency of ALK mutation in the tested population was 2.4%, followed by KRAS (1.4%), MET (1.4%), and ROS1 (1.04%) (Table III).

Chemotherapy was the most common first-line treatment (51%), specifically platinum-based chemotherapy (Table SI) followed by targeted therapy (12.5%) and immunotherapy (3.7%). Immunotherapy was mainly based on immune check point inhibitors (Table SII). Unknown first-line treatment procedures or untreated patients accounted for 32.9%.

Demographic and clinical characteristics of the progressed NSCLC cohort. A total of 31 patients from Argentina (n=25, Colombia (n=5), and Chile (n=1) with stage IV NSCLC who had progressed after first- or second-generation EGFR-TKI

treatment were included in cohort 2. The majority of these patients were female (77.4%), with a mean age of 58±12 years; all of them had adenocarcinoma histology with private health insurance (77.4%) and no smoking history (64.5%) (Table IV).

Gefitinib was the most common previous TKI (41.9%) used, followed by afatinib (35.48%) and erlotinib (22.58%). The most frequent tumor metastasis was observed in the bone (32.3%), followed by brain (16.13%), lymph nodes (16.13%), liver (16.13%), and adrenal gland(s) (9.68%). Approximately 19.35% of patients with targeted therapy showed no restricted disability (PS=0), whereas 22.58% showed limited physical disability (PS=1).

In cohort 2, 71% of patients underwent a post-progression diagnostic test. The reasons for not testing included physicians' decision and severity of disease. Thorax was the most common site of progression observed in majority of the patients (61.3%), followed by lymph nodes (19.4%) and sinonasal cancer (SNC) (19.4%). The most frequent biopsy method used at progression was Circulating Tumor DNA (ctDNA) (63.6%) (Table V).

Molecular profile and treatment at progression. Half of the tested patients were found to be T790M positive. Ten (32%) patients received osimertinib in the second line, whereas another 5 (16%) received it as third-line treatment. Patients who did not progress with T790M mainly received chemotherapy and less frequently immunotherapy in the second or third line (Table VI).

Discussion

Lung cancer is the main cause of cancer-related deaths in Latin America (13). Data scarcity on lung cancer related to clinical practices impedes progress in eliminating the disease burden. Thus, real-world data are becoming increasingly important in the clinical care of cancer patients. This retrospective real-world study evaluated the current situation of lung cancer diagnosis including molecular testing patterns and treatment regimens followed for the management of NSCLC in Argentina, Colombia, Chile, and Uruguay. The study focused on two cohorts: newly diagnosed stage IV NSCLC patients and patients with EGFR mutations who had progressed after EGFR-TKI therapy.

Of the 431 patients included in the treatment-naïve cohort, almost 90% were diagnosed at advanced stages and only 8% were diagnosed at early stages. Although the study data do not allow conclusions to be drawn because of the characteristics of the inclusion criteria, early diagnosis is still an unmet need in Latin America (13). The reasons for this include low threshold for respiratory symptoms, especially among heavy smokers; delayed consultation; and lack of programs for lung cancer screening in the region.

Despite international guidelines emphasizing the importance of molecular testing, poor access to testing has been widely reported in Latin America (14). Although two-third of the treatment-naïve cohort underwent molecular testing, this study observed disparity in testing rates among countries, from as low as 28% in Uruguay to almost 79% in Argentina. The differences between the testing of genetic mutations and the search for tumor biomarkers (66% for genetic alterations vs. 89% for PDL-1 testing) are also evident, even though the

Table I. Demographics and clinical characteristics of the newly diagnosed treatment-naïve cohort.

Characteristics	Argentina (n=195)	Colombia (n=96)	Chile (n=64)	Uruguay (n=76)	Total (n=431)
Age, mean \pm SD	66 \pm 11.00	67 \pm 10.50	67 \pm 8.96	66 \pm 9.92	66 \pm 11.00
Sex, n (%)					
Male	105 (53.8)	62 (64.6)	34 (63.5)	56 (73.7)	257 (59.6)
Female	90 (46.1)	34 (35.4)	30 (46.9)	20 (26.3)	174 (40.4)
Health insurance status, n (%)					
Private	169 (86.7)	49 (51.0)	23 (35.9)	6 (7.9)	247 (57.3)
Public	25 (12.8)	44 (45.8)	41 (64.1)	69 (90.8)	179 (41.5)
Unknown	0 (0.0)	3(3.1)	0 (0.0)	1 (1.3)	5 (1.7)
Smoking status, n (%)					
Current	76 (39.0)	19 (19.8)	18 (28.1)	47 (61.8)	160 (37.1)
Former	73 (37.4)	55 (57.3)	27 (42.2)	21 (27.6)	176 (40.8)
Never	35 (17.9)	17 (17.7)	13 (20.3)	6 (7.9)	71 (16.5)
Unknown	11 (5.6)	5 (5.2)	6 (9.4)	2 (2.6)	24 (5.8)
Family history of lung cancer, n (%)					
Yes	19 (9.7)	3 (3.1)	4 (6.2)	3 (3.9)	29 (6.7)
No	113 (57.9)	66 (68.7)	23 (35.9)	36 (47.4)	238 (55.2)
Unknown	63 (32.3)	25 (26.0)	37 (57.8)	37 (48.7)	164 (38.0)
Histology type, n (%)					
Adenocarcinoma	155 (79.5)	71 (74.0)	51 (79.7)	30 (39.5)	307 (71.1)
Squamous cell	22 (11.3)	15 (15.6)	8 (12.5)	14 (18.4)	59 (13.7)
Others or unknown	18 (9.2)	10 (10.4)	5 (7.8)	32 (42.1)	65 (15.1)
Stage at original diagnosis, n (%)					
IA	5 (2.6)	0 (0.0)	0 (0.0)	0 (0.0)	5 (1.7)
IB	7 (3.6)	0 (0.0)	0 (0.0)	0 (0.0)	7 (1.6)
IIA	1 (0.5)	2 (2.1)	0 (0.0)	2 (2.63)	6 (1.4)
IIB	4 (2.0)	0 (0.0)	0 (0.0)	0 (0.0)	4 (0.9)
IIIA	10 (5.1)	2 (2.1)	0 (0.0)	1 (1.3)	13 (3.0)
IIIB	14 (7.2)	4 (4.2)	1 (1.6)	0 (0.0)	18 (4.2)
IIIC	1 (0.5)	0 (0.0)	1 (1.6)	3 (3.95)	5 (1.2)
IVA	45 (23.1)	33 (34.4)	22 (34.4)	12 (15.8)	112 (26.0)
IVB	108 (55.4)	55 (57.3)	40 (62.5)	58 (76.3)	261 (60.6)
Site of metastasis at diagnosis, n (%)					
Brain/CNS	36 (18.5)	17 (17.7)	17 (26.6)	18 (23.8)	88 (20.4)
Bone	84 (43.1)	37 (38.5)	28 (43.7)	31 (40.8)	180 (41.8)
Lymph node	71 (36.1)	27 (28.1)	27 (42.2)	33 (43.4)	158 (36.7)
Adrenal	26 (13.3)	11 (11.5)	8 (12.5)	14 (18.4)	59 (13.7)
Liver	20 (10.3)	9 (9.4)	13 (20.3)	20 (26.3)	62 (14.4)
ECOG PS, n (%)					
0	41 (21.0)	16 (16.7)	13 (20.3)	12 (15.8)	82 (19.0)
1	52 (26.7)	33 (34.4)	24 (37.5)	17 (22.4)	126 (29.2)
2	25 (12.8)	18 (18.7)	6 (9.4)	13 (17.1)	62 (14.4)
3	10 (5.1)	4 (4.2)	0 (0.0)	12 (15.8)	26 (6.0)
4	3 (1.5)	1 (1.0)	0 0 (0.0)	3 (3.9)	7 (1.6)
Unknown	64 (32.8)	24 (25.0)	21 (32.8)	19 (25.0)	128 (29.7)

ECOG scores are divided as follows: 0, fully active, able to carry out all pre-disease performance without restriction; 1, restricted in physically strenuous activity, but ambulatory and able to carry out work of a light or sedentary nature (for example, light housework and office work); 2, ambulatory and capable of all self-care but unable to carry out any work activities (up and active for >50% of waking hours); 3, capable of only limited self-care, confined to bed or chair for >50% of waking hours. CNS, central nervous system; ECOG, Eastern Cooperative Oncology Group; PS, performance status.

Table II. Molecular testing frequency for the treatment-naïve cohort.

Parameter	Argentina (n=195)	Columbia (n=96)	Chile (n=64)	Uruguay (n=76)	Total (n=431)
Testing rate, %	78.8	64.6	75.0	27.6	65.7
PDL-1 testing rate, %	91.2	87.5	75.0	98.7	89.1
Turnaround time in days, mean	9.9	20.4	10.0	13.5	12.4
Blood (plasma/serum)	4 (2.0)	6 (6.2)	0 (0.0)	0 (0.0)	10 (2.3)
Cytology	0 (0.0)	2 (2.1)	0 (0.0)	0 (0.0)	2 (0.5)
Formalin fixed paraffin embedded tissue	134 (68.7)	44 (45.8)	41 (64.1)	21 (27.6)	240 (55.7)
Fresh frozen tissue	4 (2.0)	0 (0.0)	1 (1.6)	0 (0.0)	5 (1.2)
Unknown/no treatment, n (%)	52 (26.8)	44 (45.8)	22 (34.4)	55 (72.4)	173 (40.1)

PDL-1, programmed death-ligand 1.

search for oncogenic drivers and the consequent targeted therapy have been shown to increase patient survival and should therefore be the first diagnostic approach to treatment choice (15).

The frequency of *EGFR* mutations (22%) observed in our study is comparable with other Latin American studies that reported 26% in patients with NSCLC (16). Despite these limitations, the mean turnaround time for testing was consistent with that recommended by international guidelines (17). While treatment for advanced stages of the disease is moving toward immuno-oncology therapy (monotherapy or combined with chemotherapy) and targeted therapies in the majority of countries, first-line treatment in the Latin American region is still based on chemotherapy and target agents. In this study, the most commonly used first-line treatment for newly diagnosed patients with stage IV NSCLC was platinum-based chemotherapy, followed by target agents and immunotherapy. However, as a second-line of treatment strategy, immunotherapy was the preferred choice, followed by chemotherapy and target agents.

Since 2013, clinical guidelines have strongly recommended repeated molecular testing upon progression on first- and second-generation *EGFR*-TKIs to detect mutations such as T790M (17). Third-generation TKI osimertinib has led to a paradigm shift in the management of patients with metastatic NSCLC who develop T790M resistance mutation. The need for tumor genotyping is thus highly supported at the time of diagnosis and at disease progression (18). In previous studies among patients with NSCLC who had progressed after first-line *EGFR*-TKI therapy, approximately 60% were T790M positive (19,20). Our study shows a comparable proportion of T790M-positive patients (59%). However, the small sample size for patients who showed disease progression in this cohort might cause bias for this result and should be interpreted cautiously.

For molecular genotyping at progression, either tissue (36%) or blood (63.6%) sample was used for *EGFR* testing. The possible reason for blood samples being commonly used could be the ease of sample collection and the difficulty in obtaining tissue in some patients with disease progression. Both techniques have shown correlation in mutation

assessment, irrespective of the biopsy method used (21,22). Nevertheless, not all patients in this study who received first- or second-generation *EGFR*-TKIs and progressed were tested for T790M prior to initiating subsequent therapy. Of the 31 patients with NSCLC who showed progression after treatment, only 22 (71%) were tested, and among these, 13 had a positive resistance mutation T790M, which was found either alone or in combination with baseline sensitizing mutation, and received osimertinib. In addition, two patients received osimertinib without confirmation of T790M at progression because of central nervous system (CNS) compromise based on preclinical and clinical activity shown in patients with CNS metastases (4). Of note, at the time of this study, osimertinib was approved in Argentina, Colombia, and Chile but not in Uruguay for second-line treatment with T790M resistance mutation. The most common second- or third-line treatment for patients without T790M mutation was chemotherapy, followed by a very limited use of immunotherapy because of the poor activity of these agents in oncogene-driven NSCLC (23).

Approximately 30% of the patients who had progressed after *EGFR*-TKI treatment could not be re-biopsied and, of those tested, 42% progressed with T790M. This study highlights the importance of choosing the most optimal treatment option for patients with *EGFR*-mutated NSCLC in the first-line setting.

Strengths and limitations. The study sheds light on real-world practices with respect to lung cancer diagnosis and management in Latin America. The data are collected from medical records and closely resemble present clinical practices in the four study countries. It is important, however, to consider the implications of these results in line with the study limitations.

The sample size varies among the study countries, which could introduce bias. The results of this study among patients who showed progression should be interpreted cautiously considering the small sample size. The retrospective nature of the study could also reflect recall biases and other forms of measurement. Furthermore, given the cross-sectional design of the study, statements of causality cannot be made from the

Table III. EGFR mutation frequency, molecular profile and treatment patterns of the treatment-naïve cohort.

Parameter	Argentina (n=195)	Columbia (n=96)	Chile (n=64)	Uruguay (n=76)	Total (n=431)
Tested population, n	154	62	48	21	288
EGFR mutations, n (%)					
Exon 18	3 (1.9)	1 (1.6)	1 (2.1)	0 (0.0)	5 (1.2)
Exon 19 deletion	17 (11.0)	10 (16.1)	2 (4.2)	1 (4.8)	30 (10.4)
Exon 20 insertion	1 (0.5)	1 (1.6)	0 (0.0)	0 (0.0)	2 (0.7)
T790M <i>de novo</i>	1 (0.6)	0 (0.0)	0 (0.0)	0 (0.0)	1 (0.3)
Other	1 (0.6)	0 (0.0)	0 (0.0)	0 (0.0)	1 (0.3)
Exon 21 L858R	15 (9.7)	4 (6.4)	1 (2.1)	3 (14.3)	23 (8.0)
Other	2 (1.3)	0 (0.0)	0 (0.0)	0 (0.0)	2 (0.7)
Total EGFR mutations, n (%)	40 (26.0)	16 (25.8)	4 (8.3)	4 (19.0)	64 (22.2)
ALK, n (%)	4 (2.0)	1 (1.0)	1 (1.6)	1 (4.8)	7 (2.4)
ROS1, n (%)	3 (1.5)	0 (0.0)	0 (0.0)	0 (0.0)	3 (1.0)
KRAS, n (%)	4 (2.0)	0 (0.0)	0 (0.0)	0 (0.0)	4 (1.4)
MET, n (%)	4 (2.0)	0 (0.0)	0 (0.0)	0 (0.0)	4 (1.4)
Treatment pattern, n (%)					
Chemotherapy					
First line	107 (54.9)	41 (42.7)	41 (64.1)	31 (40.8)	219 (51.0)
Second line	14 (7.2)	6 (6.3)	6 (9.4)	3 (3.9)	24 (5.6)
Third line	0 (0.0)	0 (0.0)	0 (0.0)	3 (3.9)	3 (0.7)
Systemic therapy, n (%)	152 (77.9)	60 (62.5)	53 (82.8)	36 (47.4)	301 (69.8)
Targeted EGFR-TKI therapy, n (%)					
First line	24 (12.3)	13 (13.5)	4 (6.3)	3 (3.9)	44 (10.2)
Second line	11 (5.6)	3 (3.1)	1 (1.6)	0 (0.0)	15 (3.5)
Third line	1 (0.5)	0 (0.0)	0 (0.0)	0 (0.0)	1 (0.2)
Targeted therapy, ALK/ROS1 inhibitors					
First line	6 (3.1)	2 (2.1)	2 (3.4)	0 (0.0)	10 (2.3)
Second line	1 (0.5)	0 (0.0)	0 (0.0)	0 (0.0)	1 (0.2)
Third line	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)
Immunotherapy, n (%)					
First line	12 (6.2)	1 (1.0)	3 (4.7)	0 (0.00)	16 (3.7)
Second line	16 (8.2)	3 (3.1)	7 (10.9)	5 (6.6)	29 (6.7)
Third line	3 (1.5)	0 (0.0)	0 (0.0)	0 (0.0)	3 (0.7)
Surgeries, n (%)	23 (11.8)	12 (12.5)	6 (9.4)	3 (3.9)	44 (10.2)
Radiotherapy, n (%)	60 (30.7)	29 (30.2)	22 (34.4)	35 (46.0)	146 (33.9)
Treatment for CNS metastasis, n (%)					
Whole brain radiotherapy	19 (9.7)	12 (12.5)	8 (12.5)	11 (14.5)	50 (11.6)
Brain surgery	6 (3.1)	3 (3.1)	1 (1.6)	4 (5.3)	14 (3.2)
Stereotactic radiosurgery	6 (3.1)	1 (1.0)	2 (3.13)	0 (0)	9 (2.1)
Unknown/no treatment, n (%)					
First line	46 (23.6)	39 (40.6)	14 (21.9)	42 (55.3)	142 (32.9)
Second line	153 (78.5)	84 (87.5)	50 (78.1)	68 (89.5)	360 (83.5)
Third line	191 (97.9)	96 (100.0)	64 (100.0)	73 (96.1)	425 (98.6)

First-line chemotherapy regimens include platinum duplets (96%), platinum duplets plus bevacizumab (3%) or monotherapies (2%). Second-line chemotherapy includes platinum duplets for patients receiving immune oncology or target agents in the first line; docetaxel and maintenance regimens (pemetrexed). Immunotherapy refers to either pembrolizumab, nivolumab, nivolumab + ipilimumab or atezolizumab ALK/inhibitors, anaplastic lymphoma kinase inhibitors. CNS, central nervous system; EGFR, epidermal growth factor receptor; KRAS, Kirsten rat sarcoma viral oncogene homolog; MET, mesenchymal-epithelial transition; ROS1, reactive oxygen species inhibitors; TKI, tyrosine kinase inhibitor.

Table IV. Demographics and clinical characteristics of the cohort with non-small cell colorectal cancer progression.

Characteristic	Argentina (n=25)	Colombia (n=5)	Chile (n=1)	Total (n=31)
Age, mean \pm SD	56.84 \pm 12.49	63.8 \pm 8.40	68.00 (NA)	58.32 \pm 12.01
Sex, n (%)				
Female	18 (72.0)	5 (100.0)	1 (100.0)	24 (77.4)
Male	7 (28.0)	0 (0.0)	0 (0.0)	7 (22.6)
Health insurance status, n (%)				
Private	22 (88.0)	2 (40.0)	0 (0.0)	24 (77.4)
Public	1 (4.0)	3 (60.0)	1 (100.0)	5 (16.1)
Unknown	2 (8.0)	0 (0.0)	0 (0.0)	2 (6.4)
Smoking status, n (%)				
Current	2 (8.0)	0 (0.0)	1 (100.0)	3 (9.7)
Former	3 (12.0)	1 (20.0)	0 (0.0)	4 (12.9)
Never	17 (68.0)	3 (60.0)	0 (0.0)	20 (64.5)
Unknown	3 (12.0)	1 (20.0)	0 (0.0)	4 (12.9)
Family history of cancer, n (%)				
Yes	0 (0.0)	1 (20.0)	0 (0.0)	1 (3.2)
No	18 (72.0)	3 (60.0)	0 (0.0)	21 (67.7)
Unknown	7 (28.0)	1 (20.0)	1 (100.0)	9 (29.0)
Previous TKI, n (%)				
Afinib	11 (44.0)	0 (0.0)	0 (0.0)	11 (35.5)
Erlotinib	5 (20.0)	2 (40.0)	0 (0.0)	7 (22.6)
Gefitinib	9 (36.0)	3 (60.0)	1 (100.0)	13 (41.9)
Histology type, n (%)				
Adenocarcinoma	25 (100.0)	5 (100.0)	1 (100.0)	31 (100.0)
Squamous cell	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)
Other or unknown	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)
Site of metastasis at progression n (%)				
Brain/CNS	4 (16.0)	1 (20.0)	0 (0.0)	5 (16.1)
Bone	7 (28.0)	3 (60.0)	0 (0.0)	10 (32.3)
Lymph node	5 (20.0)	0 (0.0)	0 (0.0)	5 (16.1)
Adrenal	2 (8.0)	1 (20.0)	0 (0.0)	3 (9.7)
Liver	4 (16.0)	1 (20.0)	0 (0.0)	5 (16.1)
ECOG, n (%)				
0	6 (24.0)	0 (0.0)	0 (0.0)	6 (19.3)
1	4 (16.0)	3 (60.0)	0 (0.0)	7 (22.6)
2	3 (12.0)	1 (20.0)	0 (0.0)	4 (12.9)
3	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)
4	1 (4.0)	0 (0.0)	0 (0.0)	1 (3.2)
Unknown	11 (44.0)	1 (20.0)	1 (100.0)	12 (38.7)

ECOG scores were as follows: 0, fully active, able to carry out all pre-disease performance without restriction; 1, restricted in physically strenuous activity, but ambulatory and able to carry out work of a light or sedentary nature (for example light housework and office work); 2, ambulatory and capable of self-care but unable to carry out any work activities (up and active for >50% of waking hours); 3, capable of only limited self-care, confined to bed or chair for >50% of waking hours. CNS, central nervous system; ECOG, Eastern Cooperative Oncology Group; TKI, tyrosine kinase inhibitor.

results, and temporal trends in the relationships between study variables cannot be ascertained. Limitations in data analysis arise as a result of the finite set of measured variables and there is the possibility that unmeasured variables may have an impact on these outcomes.

This real-world study provides insight into diagnosis, mutation testing, and treatment patterns in patients with metastatic NSCLC at diagnosis as well as in patients who had progressed after EGFR-TKI treatment in four Latin American countries. The molecular profile testing rate at stage IV NSCLC is low.

Table V. Post-progression test frequency and characteristics of the cohort with non-small cell lung cancer progression.

Parameter	Argentina (n=25)	Colombia (n=5)	Chile (n=1)	Total (n=31)
Post-progression diagnostic test, n (%)				
Yes	19 (76.0)	3 (60.0)	0 (0.0)	22 (71.0)
No	5 (20.0)	2 (40.0)	1 (100.0)	8 (25.8)
Unknown	1 (4.0)	0 (0.0)	0 (0.0)	1 (3.2)
Biopsy method at progression, n (%)				
Primary tumor	3 (12.0)	1 (20.0)	0 (0.0)	4 (18.2)
Metastasis site	3 (12.0)	1 (20.0)	0 (0.0)	4 (18.2)
ctDNA (liquid biopsy)	13 (52.0)	1 (20.0)	0 (0.0)	14 (63.6)
Site of progression, n (%)				
Within the thorax	15 (60.0)	3 (60.0)	1 (100.0)	19 (61.3)
Lymph nodes	6 (24.0)	0 (0.0)	0 (0.0)	6 (19.3)
SNC	4 (16.0)	2 (40.0)	0 (0.0)	6 (19.3)
Bone and liver	8 (32.0)	3 (60.0)	0 (0.0)	11 (35.5)

ctDNA, circulating tumor DNA; SNC, sinonasal cancer.

Table VI. Epidermal growth factor receptor mutation frequency, molecular profile and treatment patterns in the cohort with non-small cell colorectal cancer progression.

Parameter	Total (n=31)
Tested population, n (%)	22 (71.0)
T790M mutations, n	2
T790M + exon 19 deletion, n	7
T790M+ L858R, n	4
Total T790M mutations, n	13
Treatment at progression, n	
Chemotherapy	
Second line	5
Third line	4
Immunotherapy	
Second line	1
Third line	1
Osimertinib, n (%)	
Second line	10 (32.3)
Second line with T790M mutation	8 (80.0)
Third line	5 (16.1)
Third line with T790M mutation	5 (100.0)

Similarly, the high percentage of patients who continue to receive chemotherapy as the only treatment in first line, along with the low use of immunotherapy or latest generation target therapies, implies significant lacunae not only in the awareness of testing but also in working to unlock access barriers that prevent achieving high-quality treatments. Differences in access to healthcare systems and economic barriers generate wide disparities in diagnosis and access to therapy, which are easily visible with real-world data.

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Availability of data and materials

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

Authors' contributions

CM acquired, analyzed and interpreted the data, and wrote, reviewed and edited the manuscript. MC, BO, RG, AS, PH, SS, BC, OF, BS, GCV, GA, KE, ZM, CJ and OE were involved in data acquisition and revised the manuscript. GS contributed to the writing, review and editing of manuscript, as well as study conceptualization. RF wrote the original draft of the manuscript, provided project administration and conceptualized the current study. RF, GCV and CM confirm the authenticity of all the raw data. All authors read and approved the final manuscript.

Ethics approval and consent to participate

The current study was approved by the Ethics Committee for each site in all the countries that participated, including The Research Ethics Committee of the Alexander Fleming Specialized Medical Institute (CEIAF); IRB Bioethics

Initiative and Reflection; CIEIS Italian Hospital; Research Ethics Committee Hospital Italian La Plata; CEMIC Research Ethics Committee; Committee of Independent Ethics of the German Hospital; CAICI-CIAP Ethics Committee; Scientific Ethics Committee of the Eastern Metropolitan Health Service; Clinical Scientific Ethics Committee; Santa Maria Clinical Research Ethics Committee of the Clinica del Country Research Ethics Committee of the San Vicente de Paul Hospital Foundation; Ethics and Research Committee of IMAT Oncomedica S.A; Ethics Committee in Biomedical Research Valle del Lili Foundation; Research Ethics Committee CAIMED S.A.S; Ethics and Research Committee of the National Institute of Cancerology E.S.E; and the Ethics Committee of the Hospital de Clinicas. Written informed consent was acquired from patients when required by local legislation.

Patient consent for publication

Not applicable.

Competing interests

Claudio Martin works as Speaker in Roche, MSD, BMS, Boehringer Ingelheim, Astra Zeneca and AB; MSD, BMS, Roche, Pfizer. Mauricio Cuello declared no conflict of interest and no disclosures to inform. Olga Barajas holds consulting or Advisory role in AstraZeneca; Bristol-Myers Squibb; MSD Oncology; Roche/Genentech and reports research Funding from Bristol-Myers Squibb; MSD Oncology; Sanofi Aventis Travel, Accommodations, Expenses-Roche; Lilly. Gonzalo Recondo, Sandra Aruachan, Herman Perroud and Susana Sena declared no conflict of interest and no disclosures to inform. Carlos Bonilla is in the Advisory board of MSD, Bristol, Eli Lilly, Novartis, AstraZeneca and Speaker in Merck-Serono, Astra-Zeneca, Eli Lilly. Patrocinio de viajes para congresos médicos: Amgen, Merck-erono, Pfizer, MSD, Roche, Tecnofarma, Participación ensayos clínicos: MSD, Roche, BMS y Astra-Zeneca. Francisco Orlandi holds Consulting or Advisory role in AstraZeneca; Bristol-Myers Squibb; Bristol-Myers Squibb; Lilly; MSD Oncology; Pfizer; Roche/Genentech and in Speakers' Bureau of AstraZeneca/MedImmune and reports research funding from Amgen; Astellas Medivation; AstraZeneca/MedImmune; Boehringer Ingelheim; Bristol-Myers Squibb; Celltrion; Genentech/Roche; MSD Oncology. Susana Berutti, Garcia Cocco Virginia, and Alvaro Gomez declared no relationship to disclosure. Ernesto Korbenfeld is doing research in AstraZeneca. Maycos Zapata reports personal fees from Astra Zeneca, Bristol-Myers Squibb, MSD, Roche and non-financial support from Bristol-Myers Squibb, MSD, Roche. Juan Cundom is in the Member Board of MSD Latin America and consulting for Boehringer Ingelheim, MSD and works also as Investigator in clinical trials pharma sponsored; MSD, Eli Lilly. Eric Orellana is employed as Investigator: AstraZeneca, Nektar Therapeutics, Merck Sharp Dohme MSD, Sanofi and Advisory Board and Consultant: AstraZeneca, Roche Genentech and declared no financial conflicts. Susana Goncalves and Florencia Reinhold works for AstraZeneca, Medical Department.

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