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Clinical, morphological and molecular characterization of cancer phenotypes associated with chronic obstructive pulmonary disease (COPD): new prospective of target therapies

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ABSTRACT

BACKGROUND

Chronic obstructive pulmonary disease (COPD) and lung cancer are two catastrophic diseases, representing leading causes of morbidity and mortality worldwide.

Although the treatment has greatly improved both diseases continue to show increasing frequency and above all an unpredictable progression.

Several studies have firmly established a strict connection between COPD and lung cancer highlighting also the importance of the inflammatory response as a risk factor for both diseases.

The inflammatory paradigm is undoubtedly one of the most fascinating theories to connect COPD and lung cancer and it has acquired new impetus by the recent discoveries in the COPD pathogenesis. Emerging evidence in this context has emphasized the role of adaptive immune responses, possibly with an autoimmune component due to the recognition of pulmonary self-antigens modified by cigarette smoking and to the failure of mechanisms regulating immunological tolerance. In this context, COPD-associated cancers might have specific pathogenetic and morphological features, differently from tumours arising in non-COPD patients, due to the synergic effect of cigarette smoke and chronic inflammation.

AIM OF THE RESEARCH

This research project focuses on the study of lung cancer in patients with COPD compared to smokers without COPD and never smoker patients in order to identify eventual distinct clinical, morphological and molecular phenotypes.

MATERIALS AND METHODS

From 2010 to 2012, we prospectively enrolled patients with peripheral non small lung cancer submitted to anatomical lung resection (lobectomy, bilobectomy or pneumonectomy) associated with systematic lymphadenectomy. Patients with central airway cancer, secondary lung tumours or previously submitted to inductive treatment were excluded from the study.

According to respiratory functional tests and smoking history patients were then divided in 3 groups: COPD patients, smokers without COPD and never-smoker subjects with normal lung

function (FEV1/FVC ratio >70%). Each patient underwent a full clinical and instrumental assessment.

Morphological studies included detailed analysis of growth pattern (according to the latest revision of adenocarcinoma classification), cell proliferation (Ki67/MIB1 expression), parameters of intra-and peri-tumoral remodelling (inflammation, fibrosis and necrosis) and tumoural detection of interleukin-17 (IL-17) cytokine. Genetic analysis of EGFR and KRAS mutations was also performed in all cases.

RESULTS

In the study period, 66 patients who met the inclusion/exclusion criteria were initially enrolled: 16 COPD, 32 smokers without COPD and 18 never smokers.

As the selection criteria affected the predominant histologic profile with a clear predominance of the adenocarcinoma histotype (63% in COPD patients, 71% in smokers and 56% in never-smokers), we performed our investigations only in patients with this histology to obtain results not affected by different histotypes. Therefore the study group was composed of 43 patients (10 COPD, 23 smokers and 10 never-smokers), whose main demographic and functional parameters were comparable except for male/female ratio, reversed in never-smokers, and for lung function, reduced in COPD patients, as expected.

Given the specific aim the comparison of different clinical, morphological and molecular data was mainly performed within the category of smoking patients (COPD patients and smokers without COPD), while never smokers represented control group.

From a **clinical point of view** the most important differences concern the number of peripheral blood basophils and standard uptake value of positron emission tomography-computed tomography (SUV of PET-CT). COPD patients showed a significant higher number of basophils and lower SUV of PET-CT than smokers without COPD.

Concerning the **histological evaluation** adenocarcinoma of COPD patients showed a more frequent lepid pattern, less evident solid aspect and lower MIB1/Ki67 index than adenocarcinoma of smokers without COPD. A significant more extensive necrosis was found in adenocarcinoma of COPD and smokers without COPD compared to never-smokers. Finally although not statistically significant a stronger IL17 tissue expression was observed in COPD cases compared to smokers without COPD.

As regards **molecular data** the most interesting finding was a trend of less frequency of KRAS mutation in adenocarcinoma of COPD patients.

CONCLUSIONS

Adenocarcinoma in COPD patients presents clinical, molecular and morphological features of lower aggressiveness (higher number of basophils, low SUV of PET-CT, increased lepidic component, reduced solid pattern, lower cell proliferation and less frequent K-RAS mutation) compared to that of smokers without COPD.

Alternative mechanisms of carcinogenesis may be involved in the development/progression of lung cancer in COPD patients. Given the importance of inflammation in the pathogenesis of the disease other mechanisms, such as IL-17 pathway, mainly driving inflammatory mediated carcinogenesis might be crucial.

Additional knowledge of these mechanisms would be of considerable help in the fight against lung cancer especially concerning therapeutic perspectives, providing a rational basis for the development of targeted and more effective treatments.

RIASSUNTO

INTRODUZIONE

La BPCO e il tumore polmonare sono due malattie catastrofiche e rappresentano alcune tra le principali cause di morbilità e mortalità in tutto il mondo.

Sebbene il trattamento di queste patologie è notevolmente migliorato negli ultimi anni, esse continuano a presentare una crescente incidenza e soprattutto un andamento clinico non prevedibile a priori.

Lo stesso termine "tumore polmonare non a piccole cellule (NSCLC)" comprende un gruppo di malattie neoplastiche con caratteristiche cliniche e molecolari estremamente eterogenee.

Diversi studi hanno ormai fermamente stabilito la stretta connessione tra la BPCO e il cancro del polmone evidenziando anche l'importanza della risposta infiammatoria agli stimoli nocivi, in particolare il fumo di sigaretta, come fattore di rischio fondamentale per entrambe le malattie. La teoria infiammatoria è senza dubbio uno dei paradigmi più affascinanti per collegare la BPCO e il tumore polmonare e ha acquisito un nuovo impulso dalle più recenti scoperte nella patogenesi della BPCO.

Infatti, sono emerse in questo campo evidenze importanti che hanno sottolineato il ruolo fondamentale di risposte immunitarie adattative, anche con una componente autoimmune dovuta sia al riconoscimento di auto-antigeni polmonari modificati dal fumo di sigaretta sia al fallimento dei meccanismi che regolano la tolleranza immunologica. In questo contesto, le neoplasie a insorgenza in pazienti con BPCO, per effetto sinergico del fumo e di una specifica infiammazione cronica, potrebbero possedere specifiche caratteristiche patogenetiche e morfologiche, differenti da tumori di altre popolazioni non affette da BPCO.

SCOPO DELLA RICERCA

Questo progetto di ricerca si concentra sullo studio del cancro del polmone nei pazienti con BPCO comparandolo a due gruppi di controllo, composti da fumatori sani e pazienti non fumatori, al fine di individuare distinti fenotipi neoplastici dal punto di vista bioumorale, morfologico e molecolare.

MATERIALI E METODI

Dal 2010 al 2012, sono stati arruolati nello studio pazienti con NSCLC in sede periferica sottoposti a resezione polmonare anatomica (lobectomia, bilobectomia o pneumonectomia) associata a linfadenectomia sistematica. I pazienti con neoplasia a carico delle vie aeree centrali, con tumore polmonare secondario o precedentemente sottoposti a trattamento chemio-radioterapico sono stati esclusi dal progetto. Ogni paziente è stato sottoposto ad una completa valutazione clinica e strumentale, che ha compreso i test di funzionalità polmonare polmonari (i criteri GOLD sono stati utilizzati per identificare i pazienti con BPCO), radiografia del torace/TAC torace/¹⁸FDG PET-TC analisi del sangue.

I pazienti sono stati poi divisi in 3 gruppi in base alle prove funzionali respiratorie e alla storia di fumo: pazienti con BPCO, soggetti fumatori e pazienti non fumatori con funzione polmonare normale (rapporto FEV1/FVC > 70%).

Lo studio istologico della neoplasia è stato caratterizzato da: stadiazione pTNM, analisi morfometrica del pattern di crescita (secondo l'ultima revisione della classificazione del cancro del polmone), proliferazione cellulare (mediante valutazione dell'espressione di Ki67/MIB1), i parametri di rimodellamento intra- e peri-tumorale (infiammazione, fibrosi, necrosi) e la caratterizzazione del pattern citochinico di IL-17 a livello peri- e intra-tumorale.

Infine è stata eseguita l'analisi genetica delle mutazioni dei geni EGFR e KRAS.

RISULTATI

Nel periodo di studio sono stati inizialmente arruolati 66 pazienti che rispettavano i criteri di inclusione/esclusione, di cui 16 BPCO, 32 fumatori senza BPCO e 18 non fumatori. Poiché i criteri di selezione hanno profondamente condizionato il profilo istologico predominante, con una netta prevalenza dell'istotipo adenocarcinoma (63% nella BPCO, 71% nei fumatori e 56% nei non fumatori), abbiamo deciso di condurre la valutazione neoplastica funzionale, morfologica, molecolare solo nell'istologia prevalente.

Pertanto, il gruppo di studio è risultato composto da 43 pazienti (10 BPCO, 23 fumatori, 10 non fumatori), che presentavano comparabili dati demografici e funzionali ad eccezione del rapporto maschio/ femmina, invertito nei non fumatori, per la funzione polmonare, ridotta nei pazienti con BPCO. Dato l'obiettivo specifico, il confronto dei differenti dati clinici, morfologici e molecolari è stato svolto prevalentemente all'interno della categoria dei soggetti fumatori (pazienti affetti da BPCO e fumatori senza BPCO), mentre i pazienti con storia negativa di fumo hanno rappresentato il gruppo di controllo.

Da un punto di vista clinico, le più rimarcabili differenze sono emerse a livello del numero di basofili nel sangue periferico e del valore di standard uptake value (SUV) all'indagine PET-TC. Infatti i pazienti con BPCO hanno mostrato un numero significativamente superiore di basofili e un SUV inferiore rispetto ai soggetti fumatori senza BPCO.

Per quanto riguarda la valutazione istologica, gli adenocarcinomi nei pazienti con BPCO hanno presentato un aumento del pattern lepidico, con riduzione della componente solida e una più bassa espressione del Ki67/MIB1 rispetto ai tumori dello stesso istotipo insorti in soggetti fumatori senza BPCO. Si è evidenziata una maggiore rappresentazione della componente necrotica negli adenocarcinomi dei pazienti fumatori, con o senza BPCO, rispetto al gruppo dei non fumatori. Infine un forte ma non significativo aumento di IL-17 è stato osservata nei casi con BPCO rispetto ai fumatori.

L'analisi molecolare ha permesso di osservare, come dato più rilevante, un trend di ridotta frequenza di mutazione di KRAS negli adenocarcinomi dei pazienti con BPCO rispetto alle neoplasie del gruppo dei fumatori.

CONCLUSIONI

Gli adenocarcinomi correlati alla BPCO sono emersi presentare caratteristiche cliniche, morfologiche e molecolari di minore aggressività (aumento del numero di basofili, ridotto SUVmax alla PET-TC, aumento della componente lepidica, ridotti pattern solido e proliferazione cellulare e meno frequente mutazione di K-RAS) rispetto alle neoplasie insorte in pazienti fumatori senza BPCO. Vie alternative di carcinogenesi potrebbero essere coinvolte nello sviluppo/progressione del tumore polmonare dei pazienti con BPCO. Data l'importanza dell'infiammazione nella patogenesi di questa malattia polmonare, altri meccanismi, quale il pathway di IL-17, potrebbero essere cruciali per lo sviluppo cancerogenetico principalmente mediato dall'infiammazione. La conoscenza di questi meccanismi potrebbe essere di notevole aiuto nella lotta contro il tumore polmonare soprattutto per quanto riguarda nuove prospettive terapeutiche, fornendo le basi per sviluppare trattamenti mirati e con maggiore efficacia.

BACKGROUND

1.1. DEFINITION OF COPD

Chronic Obstructive Pulmonary Disease (COPD) is an inflammatory disorder of the lung, characterized by airflow limitation (bronchial obstruction), that is not fully reversible [1].

The airflow limitation is usually progressive and is associated with an abnormal inflammatory response of the lung to noxious particles or gases, primarily from cigarette smoking.

In the past COPD was known with different names. Bonet described a condition of “voluminous lungs” in 1679. In 1769, Giovanni Morgagni reported 19 cases where the lungs were “turgid” particularly from air. The first description and illustration of the enlarged airspaces in emphysema was provided by Ruysh in 1721. Matthew Baillie illustrated an emphysematous lung in 1789 and described the destructive character of the condition. In 1814 Badham used the word “catarrh” to describe the cough and mucus hypersecretion of chronic bronchitis that was first reported as a disabling disorder. He recognised that chronic bronchitis was a disabling disorder. René Laennec, the physician who invented the stethoscope, used the term “emphysema (1837) to describe lungs that did not collapse as usual because they were full of air and the airways were filled with mucus. In 1842, John Hutchinson invented the spirometer, which allowed the measurement of vital capacity of the lungs. However, his spirometer could only measure volume, not airflow. Tiffeneau in 1947, and Gaensler in 1950 and 1951, described the principles of measuring airflow [2].

The terms chronic bronchitis and emphysema were formally defined at the CIBA guest symposium of physicians in 1959. The term COPD was first used by William Briscoe in 1965 and has gradually overtaken other terms to become now the preferred name for this disease [3].

In 1977 Fletcher and Peto described COPD as an obstructive and hypersecretory chronic disorder of the airways, strictly related to cigarette smoke [4].

The 1980 and 90s saw a surge in the use of medication to manage the symptoms of COPD and restore pulmonary function. A major push in COPD education meant that smoking cessation and clean air awareness became prime focuses of self-care treatment.

Today it is known that a healthy lifestyle can help people with COPD to manage and improve their symptoms. Healthcare professionals stress the importance of diet, nutrition, and physical exercise as part of a COPD rehabilitation program. Over the years, physicians have done much to help understand the causes, diagnosis, and progression of COPD.

1.2 EPIDEMIOLOGY

COPD is a major cause of morbidity and mortality worldwide, resulting in a significant economic and social costs and growing [1].

It affects about 10% of the general population but the prevalence in heavy smokers may reach 50%.

The epidemiology varies considerably between countries and between population groups within the same nation and this difference may be related both to different exposure and individual susceptibility to risk factors and different methods and criteria for diagnostic characterization.

The BOLD Study (Burden of Obstructive Lung Disease) [5], a prevalence study conducted in 2007 on 9425 subjects in 12 different locations of the world, reported the following data:

- Higher prevalence in males.

- Prevalence of disease progressively increase with age in both males and females.

- Increased prevalence in parallel with the increase in the number of pack-years (number of cigarettes per day multiplied by years of smoking divided by 20).

- Substantial prevalence of COPD even in non-smokers: 11.3%.

Most of the epidemiological data collected in Western countries shows that the diagnosis of COPD is found in less than 6% of the population, but it is reasonable to assume that the disease still remains under-diagnosed.

On the one hand, the slow evolution of the natural history and lack of specificity of symptoms, especially in the early stages, are the cause of the delay with which patients come to medical attention. Secondly the population at risk is large, with absence of valid screening programs [6,7].

Since the disease was responsible for 4% of the total deaths, the WHO in 2001 placed this disease in fifth place among the leading causes of death in industrialized countries and in sixth place in the developing countries. It is expected that the prevalence and impact of the disease will increase in the coming decades, in parallel with the most exposure to recognized risk factors and aging population. By 2020, according to the BOLD Study, it is estimated that COPD will become the third leading cause of death in industrialized countries.

1.3 CLINICAL ASPECTS

1.3.1 LUNG SYMPTOMS

In the lungs, COPD is mainly characterized by the presence of progressive stress dyspnea that can evolve to respiratory failure associated with chronic bronchitis.

The obstruction of the small airways, responsible for the loss of lung function, has a slow and progressive development and typically occurs late, whereas the contribution of this part of the bronchial tree to expiratory flow is 10-15% of the total resistance [8]. Obstruction of the small airways and increased lung compliance determine an increase of exhalation duration, appearance of hyperinflation and increase in the residual volume. In addition to increase the functional dead space, hyperinflation causes a flattening of the diaphragm, thereby affecting its

contractile efficiency, and reduces the movements of the chest wall. This condition is clinically manifested with stress dyspnea and limitation of exercise tolerance [9].

The effort made by patients suffering from emphysema during exhalation, causes a pink colour in their faces, hence the term commonly used to refer to them, “Pink Puffers” [10]. With the progression of the disease, respiratory effort increases in association with expansion of the residual volume and an increase in the muscular work required for each breath. The result is a reduction in pulmonary and alveolar ventilation responsible for blood gas alterations such as chronic hypoxemia ($\text{PaO}_2 < 55 \text{ mmHg}$) and hypercapnia ($\text{PaCO}_2 > 45 \text{ mmHg}$) with a final result of pulmonary acidosis more or less offset by the retention of bicarbonate in the kidney [11].

Chronic bronchitis

Clinical condition characterized by the presence of cough and sputum production for at least three months a year for two consecutive years. These symptoms may occur in the natural history of COPD at different times and with varying severity. Hypersecretion of mucus in the proximal airways, responsible for chronic bronchitis, however, does not correlate with the decline in lung function (FEV) in patients with COPD [12].

Patients with advanced COPD that have primarily chronic bronchitis rather than emphysema were commonly referred to as “blue bloaters” due to the bluish colour of the skin and lips (cyanosis) [9].

CLINICAL FEATURES OF PREDOMINANT CHRONIC BRONCHITIS
AND PREDOMINANT EMPHYSEMA*

	Predominant Bronchitis	Predominant Emphysema
General appearance	Mesomorphic, overweight, dusky with suffused conjunctivae, warm extremities	Thin, often emaciated; pursed-lip breathing; anxious, prominent use of accessory muscles; normal or cool extremities
Age, years	40-55	50-75
Onset	Cough	Dyspnea
Cyanosis	Marked	Slight to none
Sputum	Copious	Scanty
Upper respiratory infections	Common	Occasional
Breath sounds	Moderately diminished	Markedly diminished
Cor pulmonale and right heart failure	Common	Only during bout of respiratory infection, and terminally
Radiograph	Normal diaphragm position; cardiomegaly, lungs normal or with increased bronchovascular markings	Small, pendulous heart; low, flat diaphragm; areas of increased radiolucency
Course	Ambulatory but constantly on verge of right-sided heart failure and coma	Incapacitating breathlessness punctuated by life-threatening bouts of upper respiratory infections; prolonged course, culminating in right heart failure and coma

TABLE 1.1 Clinical features of the principal COPD patterns, from Travis WD et al., atlas of nontumour pathology “Non-neoplastic disorders of the lower respiratory tract” (8).

Pulmonary Hypertension

The inefficient alveolar ventilation is associated with a reduction of vascular bed in the parenchyma which is on the base of hypoxic vasoconstriction. Even the destruction of alveolar septa in emphysema contributes to reduce the extension of the pulmonary capillary. An endothelial dysfunction is also present in pulmonary vessels, which correlates with the airways inflammation. The structural changes in the pulmonary arterioles occur progressively such as hypertrophy of the intima and in the second place of the tunica muscularis, and determine the persistent increase in resistance in the pulmonary microcirculation [13].

These alterations lead to the development of pulmonary hypertension, which may appear in the advanced stages of COPD (FEV <25% predicted), and it is responsible for right ventricular hypertrophy and, ultimately, for cor pulmonaris.

Exacerbations

Exacerbations have a significant impact on morbidity, disease progression, disability and socio-medical costs [14]. Bacterial or viral infections, often overlapping with other factors such as environmental pollutants are the principal cause of exacerbation. From a clinical point of view, exacerbations may lead to an increased cough, change in sputum characteristics, that becomes

more abundant and/or purulent, appearance of wheezing / whistling expiratory and worsening of dyspnea.

During the course of the disease, the frequency and severity of exacerbations tends to increase, resulting in accelerated decline in lung function.

1.3.2 COMORBIDITY AND SYSTEMIC INVOLVEMENT

Comorbidities contribute to the overall severity in individual patients (Global Initiative for Chronic Obstructive Lung Disease, Revision 2011) [1]. Moreover, COPD also produces significant systemic consequences mainly due to the development of the systemic inflammation.

Cardiovascular disease

These are mainly acute cardiovascular events due to atherosclerosis: endothelial dysfunction, that occurs in the pulmonary vessels, becomes progressively a systemic condition, along with the spread of the inflammatory reaction. CRP (C-reactive protein) is a key marker of systemic inflammatory response and the serum level correlates with the risk of acute cardiovascular events (CAD), such as myocardial infarction, unstable angina, and others. In COPD, CRP levels correlate with the severity of bronchial obstruction (ie, with the decline of FEV) and the risk of cardiac and pulmonary complications [15,16].

Metabolic syndrome

It is a complex disorder and an emerging clinical challenge, recognised clinically by the findings of abdominal obesity, elevated triglycerides, atherogenic dyslipidaemia, elevated blood pressure, high blood glucose and/or insulin resistance, for which the relative risk is increased (RR = 1.8) especially in women [17]. Patients with COPD often have one or more component of the metabolic syndrome and osteoporosis (70% of patients) which are, at least in part, independent from treatment with steroids and/or the decreased physical activity.

COPD patients have an increased risk of fractures, especially at vertebral level, and it has a direct effect on lung function with reduced vital capacity (estimated a loss of 7% for each vertebral fracture) [18].

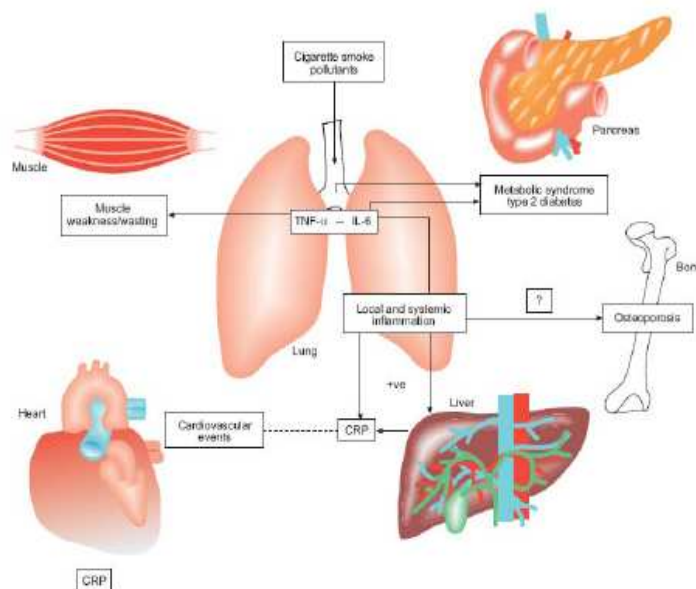


FIGURE 1.1 The central role of inflammation in comorbidity is associated with COPD. Inflammation appears to play a central role in the pathogenesis of COPD and other conditions that are increasingly being recognised as systemic inflammatory diseases. As part of the chronic inflammatory process, tumour necrosis factor (TNF)- α receptor polymorphisms are associated with increased severity of disease, possibly due to enhanced TNF- α effects. Also, C-reactive protein (CRP) levels can be increased directly by TNF- α and other cytokines. Elevated CRP and fibrinogen may be crucial in the pathogenesis of cardiovascular disease. Reactive oxygen species released as a result of COPD may enhance the likelihood of a patient developing cardiovascular disease, diabetes and osteoporosis. IL: interleukin;?: unknown; +ve: positive. From Fabbri LM et al, *Eur Respir J* 2008 (18).

1.4 DIAGNOSIS

1.4.1 HISTORY AND PHYSICAL EXAMINATION

The symptoms of chronic bronchitis (worsening dyspnea, cough and sputum production) may precede by many years the development of airflow obstruction.

This pattern offers a unique opportunity to identify smokers and other patients at risk for COPD, and to intervene when the disease is not yet a major health problem. In the presence of such symptoms and signs, a complete anamnestic evaluation should be obtained, including familiar history, presence of comorbidities and other respiratory diseases.

Particular interest must be reserved to risk factors exposure, especially cigarette smoking, other environmental factors and genetic predisposition or working.

1.4.2 SPIROMETRY

This is the most reproducible and objective measure available of airflow limitation. The spirometry should be performed after adequate dose of bronchodilator with short duration of action, that reduce the variability of the test. A value of $FEV_1 / FVC < 0.70$ after bronchodilator allows confirmation of persistent airway obstruction.

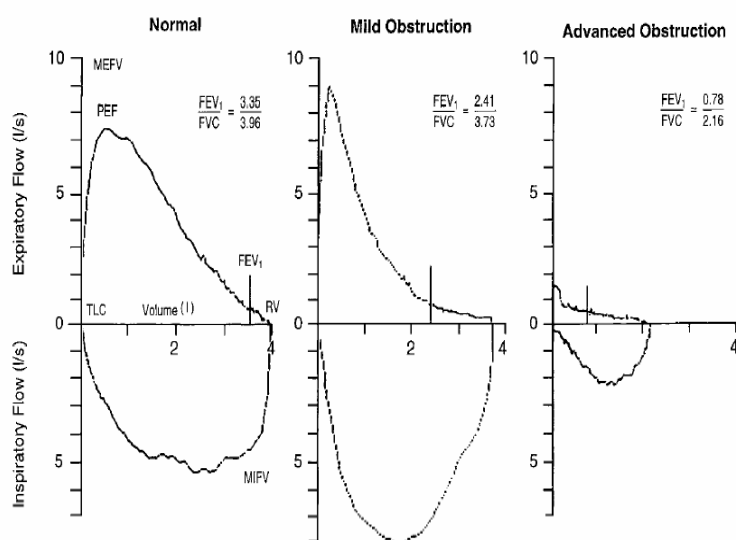


Figure 1.2 Graph of the curve spirometric flow / volume. In advanced stages of COPD, there is a marked reduction in FEV, FVC and maximal expiratory flow in general. The curve flow / volume in the expiratory phase flattens out. Even the inspiratory flow undergoes a reduction but less pronounced than that expiratory [19].

1.4.3 EVALUATION OF GRAVITY DISEASE

The new GOLD guidelines (Revision 2011) [1] underlines the diagnostic, therapeutic and prognostic importance of a more comprehensive assessment of disease severity, through the integration of anamnestic and clinical data with spirometry. In previous documents the severity of COPD was defined only on the basis of the degree of airway obstruction, assessed by spirometry in terms of value compared with the theoretical VEMS:

- GOLD 1: Mild, FEV₁ > 80% predicted
- GOLD 2: Moderate, 50% < FEV₁ < 80% predicted
- GOLD 3: Severe, 30% < FEV₁ < 50% predicted
- GOLD 4: Very Severe, FEV₁ < 30% predicted

One very important aspect is that only a minority of the smoking population (approximately 20% of the total) experience the rapid decline in lung function that leads to severe (GOLD-III) and very severe (GOLD-IV) COPD.

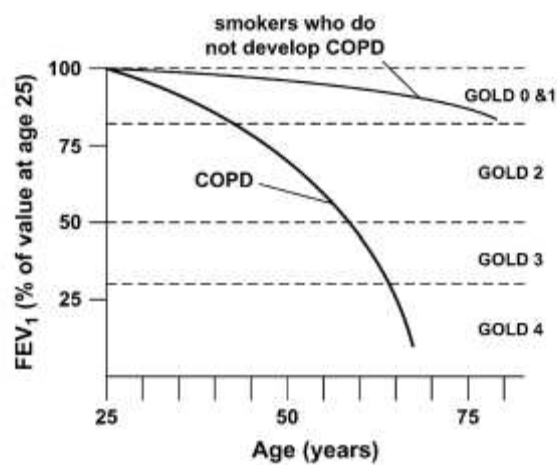


Fig. 1.3. The natural history of the FEV₁ decline in men followed by Fletcher and coll. (3) is shown with the GOLD severity stage superimposed as dotted horizontal lines. Modified from Curtis JL et al., Proc Am Thorac Soc 2007 (20).

According to the new guidelines, the two parameters are associated with other indices of disease severity, in order to quantify the risk of subsequent decline in lung function and mortality with greater predictive value. These indices are the severity of symptoms and the risk of exacerbations.

The severity of symptoms is determined on the basis of two questionnaires:

- the MMRC (british modified Medical Research Council) questionnaire, that assesses the degree of dyspnea and correlates with the risk of future mortality.
- The CAT (COPD Assessment Test), that provides a measure of the deterioration of health status in COPD through 8 questions.

The risk of subsequent exacerbations is based on the history of similar events recently treated.

The combination of the different parameters leads to define four groups of COPD patients:

- Group A: low risk, mild symptoms
- Group B: low risk, severe symptoms
- Group C: high risk, mild symptoms
- Group D: high risk, severe symptoms

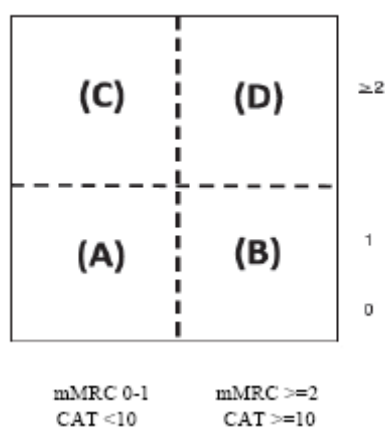


Figure 1.4 . Graph integrated for the assessment of disease severity according to the guidelines GOLD 2011 (1).

We have to note that patients in GOLD stages 3-4 are at increased risk of hospitalization or death, even in the absence of frequent exacerbations. For this reason it is reasonable to include these patients in the groups at "high risk".

1.4.4 ADDITIONAL INVESTIGATIONS

They are useful for diagnostic and especially for patient follow-up and they include:

1) Radiological investigations (Rx chest in two projections, CT scan), in particular for the diagnosis of emphysema and evaluation of its distribution in the lung.

The radiologic features of emphysema are hyperinflation and hyperlucency of the lung fields, destruction of lung parenchyma, flattening of hemidiaphragms, horizontal ribs and presence of bubbles.

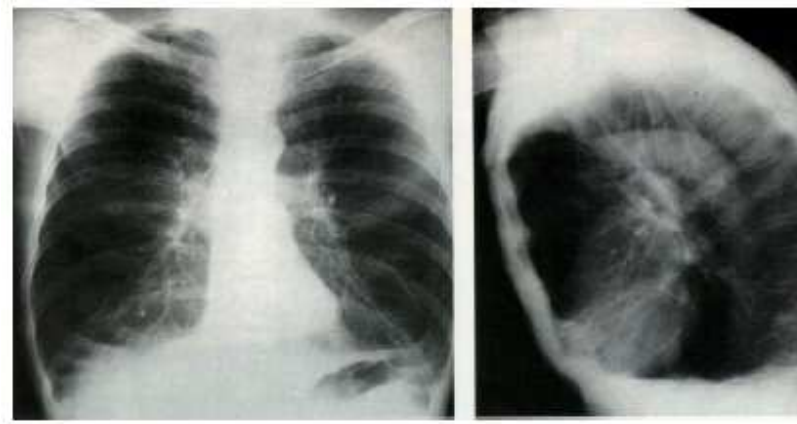


Figure 1.5 Chest x-ray in two projections that highlights the above described features of pulmonary emphysema.

- 2) Measurement of CO diffusion.
- 3) Pulse oximetry and blood gas analysis.
- 4) Screening for alpha1-AT deficiency
- 5) The exercise test (6-minute walking test)
- 6) Mixed indices, such as the BODE index, ranging from 0 to 10 [21], in which we evaluate the prognostic impact of global factors both pulmonary and extrapulmonary. The BODE index is an indicator of mortality risk in patients with COPD and for this reason it is particularly important for the decision of inclusion on the waiting list for lung transplantation [21]. In fact, subjects with a high BODE index (score of 7 to 10) show a mortality rate of 80% at 5 years, with a median survival of about 3 years, so they should be evaluated for lung transplantation. Patients with a BODE score of 5 to 6 would likely not derive a survival benefit from transplantation but may be candidates for early referral.

	0	1	2	3
FEV ₁ % pred	≥85	50-84	36-49	≤35
6MWD (m)	≥350	250-349	150-249	≤149
MMRC	0-1	2	3	4
BMI (kg.m ⁻²)	>21	≤21		

Total BODE Index score = 0 to 10 units

(FEV₁ % pred = predicted amount as a percentage of the forced expiratory lung volume in one second; 6MWD = six minute walking distance; MMRC = modified medical research council dyspnea scale; BMI = body mass index)

Table 1.2 Parameters for the calculation of the BODE index. BODE index is obtain through the sum of the scores of the 4 parameters evaluated.

1.5 MORPHOLOGICAL ASPECTS

The airflow limitation in COPD is a consequence of the functional framework pathological characteristic of the disease, which is mainly composed of two aspects: the small airway remodeling and pulmonary emphysema [1].

Chronic bronchitis is characterized by hyperemia, swelling, and oedema of the mucous membranes, frequently accompanied by excessive mucinous to mucopurulent secretions layering the epithelial surfaces and chronic inflammation at histology. Sometimes, heavy casts of secretions and pus fill the bronchi and bronchioles. Although the numbers of goblet cells slightly increase, the major pathological changes concern the bronchial glands, both in size and mucous production. The increased glandular volume can be assessed by the ratio of the thickness of the mucous gland layer to the wall between the epithelium and the cartilage (Reid index).

The Reid index (normally 0.4) is increased in chronic bronchitis, usually in proportion to the severity and duration of the disease. The bronchial epithelium may exhibit squamous metaplasia and dysplasia. There is marked narrowing of bronchioles caused by goblet cell metaplasia, mucus plugging, inflammation, and fibrosis. In the most severe cases, there may be obliteration of lumen due to fibrosis (bronchiolitis obliterans). As it was previously mentioned, these bronchiolar changes probably contribute to the obstructive features in bronchitis patients.

The American Thoracic Society defines emphysema as “a condition of the lung characterized by abnormal, permanent enlargement of the airspaces distal to the terminal bronchiole, accompanied by destruction of their walls” [22]. Fibrosis may be absent or subtle and mild. There are several types of emphysema according to its anatomic distribution within the lobule.

1. *Proximal acinar (centrilobular, centriacinar)* emphysema is characterized by distension and destruction mainly limited to the respiratory bronchioles, with relatively less change peripherally in the acinus. This pattern results from scarring and focal dilatation of the bronchioles and adjacent alveoli, which lead to an enlarged airspace or “microbulla” in the center of the secondary lung lobule. The central or proximal parts of the acini, formed by respiratory

bronchioles, are affected, whereas distal alveoli are spared. It is strongly associated with long-term cigarette smoking and predominantly involves the upper and posterior portions of the lung (Figure 1.6a-c).

2. *Panacinar (panlobular) emphysema* is characterised by diffuse, bilateral lung involvement in which the alveolar ducts as well as respiratory bronchioles are enlarged, causing uniform destruction of the alveolar tissue. This pattern is less common than proximal acinar emphysema and comprises several conditions including familiar emphysema associated with α 1-antitrypsin deficiency (AAT) (Figure 1.6b-d). Another key distinction between the two phenotypes of emphysema which has recently emerged is the inflammation characteristics [23]. Proximal acinar emphysema presents more numerous T-CD4 +, neutrophils, macrophages, but not T-CD8 + nor eosinophils. Furthermore we have an infiltration by mast cells, which is predominant in the smooth muscle of the airways and in the alveolar wall and correlates with the degree of bronchial reactivity [23].

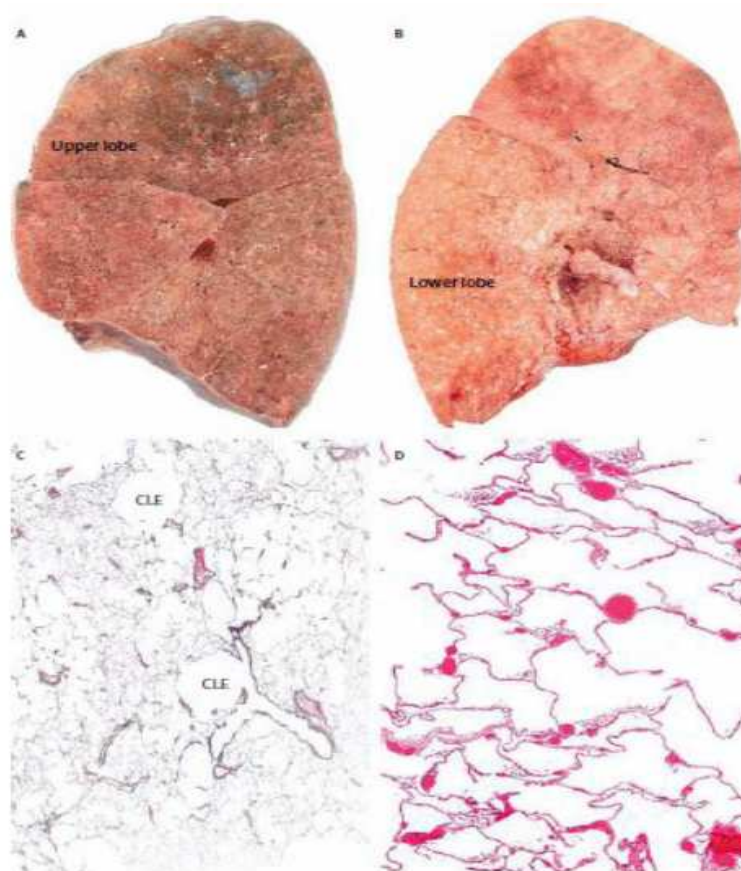


Fig. 1.6 On the left side macroscopic view (a) and histology (c) of centrilobular emphysema. On the right side macroscopic view (b) and histology (d) of panacinar emphysema.

3. Other forms include *distal acinar (paraseptal)* emphysema, which affects the periphery of the acinus, most often in the upper lobes beneath the pleura, and the *irregular* emphysema refers to airspace enlargement and lung destruction associated with a pulmonary scar.

1.6 PATHOGENESIS OF THE DISEASE

1.6.1 AETIOLOGY

Risk factors

Risk factors for COPD include both genetic factors and environmental exposures; the disease seems to arise from an interaction between these two types of factors.

- **GENETIC FACTORS:** the genetic factor that is best documented is a severe hereditary deficiency of alpha-1-antitrypsin, a prototype of protein inhibitors of serine proteases.

This serpin has a major role in inactivating neutrophil elastase and other proteases to maintain protease/anti-protease balance [24]. Reduction in anti-elastase defence (which might happen with severe deficiency of AAT) can unfavourably tip the elastase/anti-elastase balance unfavourably towards accelerated lung breakdown. In addition, AAT has important anti-inflammatory properties and this genetic defect could be responsible for the abnormal inflammatory response observed in COPD lungs.

The published COPD genetic association studies have focused on other candidate genes, identified by linkage analysis [25]. Variations in candidate genes, which are usually due to single-nucleotide polymorphisms (SNPs), have been investigated to detect association with COPD susceptibility.

Candidate genes involved in protease/anti-protease balance, oxidative stress, xenobiotic metabolism of toxins, and inflammatory or immune responses have been explored in this

field. The candidate genes listed in the following table (Table 1.4) resulted from targeted investigations based on linkage studies or pathophysiologic hypotheses [26].

Table 1—COPD Candidate Genes in Which Emphysema Has Been Investigated

Genes	Functional Class	Variants	Participants, No. (Emphysema Analysis)	Evidence for Emphysema Association
Matrix metalloproteinase 9 (MMP 9)	Metalloprotease: type IV and V collagenase	rs 3918242 (C-1562 T)	45 case patients/65 control subjects 84 case patients/55 control subjects	T allele associated with higher emphysema scores (Minematsu et al ¹¹ /2001) T allele associated with apical predominant emphysema among case patients with emphysema (Ito et al ¹⁵ /2005)
Microsomal epoxide hydrolase (EPHX1)	Xenobiotic metabolism: first-pass metabolism/detoxification of highly reactive species	rs 1051740 (T 113 C) rs 2234922 (A 139 C)	94 case patients/203 control subjects 282 case patients	C allele results in "slow" enzyme, associated with COPD and emphysema (Smith and Harrison ⁴² /1997) C allele results in "fast" enzyme, associated with reduced risk for apical predominant disease (DeMeo et al ¹⁴ /2007)
Heme oxygenase 1 (HMOX1)	Antioxidant: against heme- and non-heme-mediated oxidant damage	No. of (GT) ⁿ repeats at 5' end	101 case patients/100 control subjects	Long repeats (designated L-allele) putatively less inducible when exposed to oxidative stress; presence of L-allele associated with emphysema among case patients (Yamada et al ¹² /2000)
Glutathione S-transferase P1 (GST P1)	Xenobiotic metabolism: conjugates hydrophobic/electrophilic compounds	rs 1695 (A 105 C)	282 case patients	A allele (Ile) associated with COPD and with apical emphysema distribution (DeMeo et al ¹⁴ /2007)
Vitamin D binding protein (GC)	Immune response: in addition to binding vitamin D	1F allele	85 case patients	Presence of 1F allele associated with increased severity of emphysema (Ito et al ¹⁰ /2004)
β_2 -Adrenergic receptor (ADRB2)	β_2 -Adrenergic receptor	rs 1042713 (A 16 G)	111 case patients	C allele associated with airway diameter/percent wall area but not associated with emphysema in a Korean population (Kim ⁴⁹ /2009)
TNF- α (TNF)	Proinflammatory cytokine	rs 1800629 (G -308 A)	84 case patients 424 case patients	A allele putatively increases transcription; trend (p = 0.09) toward increased emphysema scores in a Japanese population (Sakao et al ¹⁷ /2002) Multiple SNPs tagging TNF showed no correlation to presence of emphysema in A1ATD patients from the United Kingdom (Wood et al ¹⁹ /2008)
Transforming growth factor- β_1 (TGFB1)	Autocrine peptide involved in proliferation/apoptosis and differentiation		70 case patients/99 control subjects	A haplotype block of eight SNPs was marginally associated with emphysema phenotype in Japanese subjects; however, no individual SNP showed a significant association (Ito et al ¹⁰ /2005)
Transforming growth factor- β receptor-3 (TCFBR3)	Glycoprotein receptor for transforming growth factor- β		527 probands/640 siblings (ICGN) 359 case patients/472 control subjects (NETT-NAS)	Genetic linkage analysis in families of emphysema predominant early-onset COPD probands led to increased evidence for linkage on chromosome 1 near TCFBR3; two intronic SNPs (rs2296621 and rs2291477) were associated with the presence and severity of COPD, with a trend toward increased emphysema in some of the studied populations (Hersh ⁴⁹ /2009)

NAS = Normative Aging Study; NETT = National Emphysema Treatment Trial.

TABLE 1.3 Candidate genes, besides AAT, with replicated associations to COPD, emphysema or related traits. From Wan ES et al., Chest 2009 (26).

- EXPOSURE TO ENVIRONMENTAL POLLUTION:

a) Active / passive / in utero cigarette smoking

This is the main risk factor for COPD. However, not all smokers develop clinically significant COPD, which suggests that other cofactors, host-related or not, may modify each individual risk. "Susceptible" smokers show a decline in lung function (FEV) of different degree, but generally progressive and stronger (60ml/year or more) than not susceptible smokers and non-smokers (30ml/year). In addition, smokers have a higher prevalence of respiratory symptoms and a higher mortality rate associated with COPD [4].

The smoke damage is cumulative and it depends on intensity and duration of exposure. The injury is due to an increase oxidative stress, resulting in airway epithelial cell damage and

activation of pro-inflammatory signals, responsible for recruitment of innate immunity cells and the perpetuation of tissue aggression [27,28].

b) Occupational dusts and chemicals.

Occupational exposures include organic and inorganic dusts and chemical agents and fumes. A statement published by the American Thoracic Society concluded that occupational exposures account for 10 to 20% of either symptoms or functional impairment consistent with COPD [29].

c) Indoor and outdoor air pollution.

There is evidence that indoor pollution from burning biomass fuels for cooking and heating is an important cause of COPD in many developing countries [30,31], but the exact pathogenetic mechanisms have not yet been elucidated.

Other recognized risk factors are:

- *Altered growth and development of respiratory system:* a low weight birth is associated with reduced values of lung function indices (FEV) in adulthood, implying a state of incomplete development of the respiratory system [32]. In addition, numerous studies have demonstrated the effect of pediatric recurrent respiratory infections on the risk of subsequent decline of FEV.
- *Sex:* historically COPD hits the males, as they are more exposed to cigarette smoke. The incidence and prevalence of the disease in women is growing, with the increase in smoking in recent decades [33].
- *Age and ageing:* epidemiological studies show that prevalence and impact of the disease increased with age [5]. Nevertheless, it is unclear whether aging is itself a risk factor for the disease or whether this association is an expression of cumulative and prolonged exposure to traditional risk factors.

- *Respiratory And Systemic Infections*: tuberculosis is a risk factor for the development of COPD. On the other hand it is seen that HIV infection accelerates the onset of emphysema smoke-related [34].
- *Socio-Economic Status*: There is a strong evidence of an inverse relationship between socioeconomic status and the risk of developing COPD, but the factors responsible for this association are not clear [35].
- *Asma*: A longitudinal study (Tucson Epidemiological Study of Airway Obstructive Disease) has shown that adults with asthma have a RR = 12 to develop COPD than non-asthmatics, adjusted for smoking status [36].
- *Chronic Bronchitis*: the study of Fletcher and colleagues showed that chronic bronchitis was not associated with decline in lung function [4]. Subsequent studies, however, have re-evaluated the role of mucus hypersecretion in the decline of FEV and the existence of an increased risk of developing COPD in young adults who smoke and who are suffering from chronic bronchitis [36].

1.6.2 PATHOGENETIC MECHANISM

Despite advances in care to date there are no therapies which halt or reverse the progressive and accelerated decline in lung function of this condition. Better means of preventing and treating COPD are urgently needed and a better understanding of the pathological bases is a necessary requirement.

The discovery of the AAT deficiency in the 60s led to the hypothesis that an imbalance between proteases and antiproteases in the lung could be the cause of lung damage in all COPD patients [37]. Smoking induces an increased number of neutrophils and macrophages in the lung and the release of proteolytic enzymes from these cells. The released proteases, not fully inhibited by antiproteases, lead to proteolysis of lung connective tissue (more specifically of elastin) and emphysema. However, the pathogenetic view of COPD has expanded significantly from what was understood in past decades. Moving beyond the original protease/antiprotease hypothesis, T-

lymphocytes have been identified as a key component of the inflammatory response, thus introducing the concept that adaptive immunity may be centrally involved in the pathogenesis of the disease [38]. Indeed, recent studies have convincingly shown that emphysema can be produced in experimental models, not only by manipulation of proteolytic pathways, but also targeting CD8 T-cell responses [39]. Distinctive immune features, such as lymphoid follicles rich in B and T cells are found in lung tissue of patients with COPD, at least in the severe stages of the disease [40]. Altered T and B cell responses are also seen in the peripheral blood of patients with COPD, indicating that immunological alteration extends beyond the lung.

Despite the well documented presence of inflammation in COPD, it is unclear what steps are involved in the genesis and progression of the disease. The new concept emerging from the most recent observations, to which substantial contribution was given by our group, is that persistence of the immune activation in COPD could be due to an autoimmune reaction secondary to the lung damage triggered by cigarette smoking (Fig. 1.7) [41]. Cigarette smoking can damage the epithelium and the products of epithelial cell injury can stimulate the innate immune response, acting as ligands for Toll like receptors (TLRs), especially TLR-4 and TLR-2.

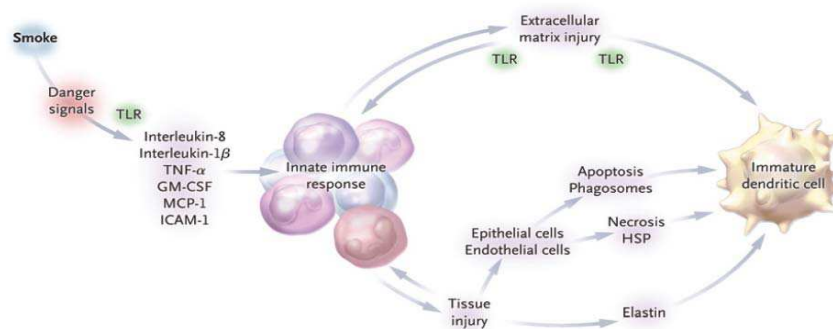


Fig. 1.7. Initial Response to Cigarette Smoke — Step 1

Cigarette smoke injures epithelial cells, which release “danger signals” that act as ligands for toll-like receptors (TLRs) in the epithelium. These actions trigger the production of chemokines and cytokines, which results in an innate inflammation. Products from the inflammatory cells may injure the extracellular matrix, leading to the release of TLR ligands and consequent TLR activation, which will promote further inflammation, tissue injury, and the production of antigenic substances. This chain of events may cause dendritic cells to mature and migrate to local lymph organs, where, if the conditions are favorable, T-cell activation may result, with progression of the disease. If the innate inflammation in step 1 is minimized or controlled, the inflammation will not progress to adaptive immunity, and the disease may be arrested. These processes are typical of smokers who have neither COPD nor Gold stage 1. GM-CSF denotes granulocyte–macrophage colony-stimulating factor, HSP heat-shock protein, ICAM-1 intercellular adhesion molecule 1, MCP-1 monocyte chemoattractant protein 1, and TNF tumor necrosis factor.

From Cosio MG et al., N Engl J Med 2009 (41).

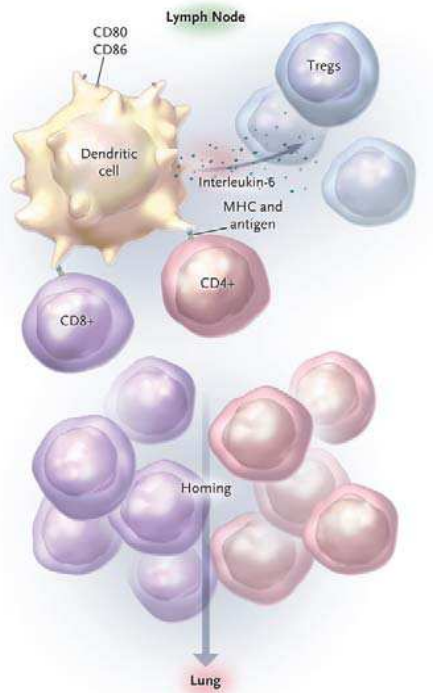


Fig. 1.7. Proliferation of T Cells — Step 2

When step 1 is successful, mature dendritic cells migrate to local lymphatic organs, whereupon stimulation by toll-like receptors (TLRs) leads to the expression of CD80–CD86 and cytokines, creating a propitious milieu for T cell antigen presentation and proliferation into effector CD4+ type 1 helper (Th1) T cells and cytolytic CD8+ T cells. Interleukin-6, secreted by the dendritic cells, favors the production of effector T cells by overcoming the signals from regulatory T (Treg) cells. Upon activation, effector T cells express tissue- specific chemokine receptors. Immune regulation or tolerance mechanisms will determine at this stage the degree of proliferation of T-cell effectors, homing, and eventually, disease severity. An absence of tolerance is associated with Gold stage 3 or stage 4, moderate tolerance with Gold stage 2, and full tolerance with Gold stage 1. MHC denotes major histocompatibility complex.

From Cosio MG et al., N Engl J Med 2009 (41).

The subsequent activation of alveolar macrophages and neutrophils, with their array of proteolytic enzymes, further damage the lung tissue disrupting the extracellular matrix and the resulting breakdown products, such as hyaluronate and bicyclan, can also ligate TLR- and TLR-4. In support of this proposed pathway is the finding in mice that lung inflammation induced by cigarette smoke depends on TLR-4 and MyD88, an adapter protein that stimulates NFκB [42]. These processes would result in release of normally sequestered autoantigens (proteins and DNA that may undergo modifications) which the adaptive immune system can recognize as foreign antigens triggering an immune reaction (Fig. 1.8).

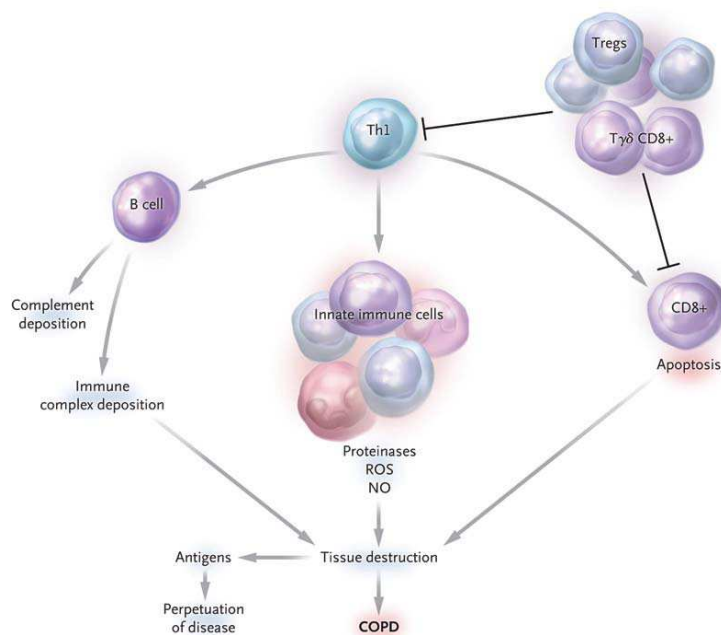


Fig. 1.8. The Adaptive Immune Reaction — Step 3

Inflammation (autoimmune) develops in the lung, consisting of CD4+ type 1 helper (Th1) T cells, cytolytic CD8+ T cells, and IgG-producing B cells. Regulatory T cells (Treg) and $\gamma\delta$ CD8+ T cells could modulate the severity of the adaptive immune inflammation. The resulting immune inflammation, induced by CD4+ Th1 T cells and consisting of activated innate immune cells producing oxidative stress and proteinases, along with cytolytic CD8+ T cells and B cells, leads to cellular necrosis and apoptosis, immune and complement deposition, tissue injury with airway remodelling, and emphysema, as well as the release of additional antigenic material, which perpetuates the process. In step 3, the full autoimmune process has developed, producing the most severe disease (Gold stage 3 and stage 4). NO denotes nitric oxide, and ROS reactive oxygen species.

From Cosio MG et al., *N Engl J Med* 2009 (41).

In this context some important cytokine may be involved to disease development and progression.

One of these is interleukin-32 (IL-32), which role may be of potential interest. IL-32 is a pro-inflammatory cytokine, which is upregulated in autoimmune diseases such as rheumatoid arthritis [43] and Chron's disease [44]. The cytokine has important functions in innate and adaptive immune responses. It synergizes with NOD1 and NOD2 ligands to stimulate IL-1 β and IL-6 release in a caspase-1-dependent manner. Proteinase 3 (PR3) is a specific IL-32 binding protein which cleaves the cytokine to enhance its activity. Interestingly, AAT is the main inhibitor of PR3 [45]. There is evidence that viral stimuli (influenza A or double-stranded RNA [poly(IC)] + IFN- γ) may trigger IL-32 expression and that cigarette smoke can potentiate the lung responses caused by these stimuli.

Of interest, we recently reported an increased expression of IL-32 in lung tissue of patients with COPD, where it was co-localized with TNF α and correlated with the degree of airflow obstruction, suggesting that IL-32 could be a potentially useful biomarker to evaluate the progression of the disease [46].

1.7 LUNG CANCER

Lung cancer is currently the leading cause of cancer death in the world with a 5-year survival of about 15%, despite the use of increasingly aggressive therapies [47]. Currently, lung cancer has a maximum incidence between 55 and 65 years and it is responsible for 32% of total deaths for cancer in men and 25% in women [47]. This high mortality closely linked to the characteristics of the tumor and to the delay in diagnosis of this disease which presents a rather low percentage of patients (approximately 20%) with early stage cancer.

The World Health Organization has included in the new classification of lung cancer 4 main histological types [48]; however, under therapeutic and prognostic purposes the most useful and widespread differentiation of lung cancer involves two major groups, defined as small cell lung carcinoma (SCLC) and non-small cell lung cancer (NSCLC).

Small cell lung cancer is an extremely malignant tumor, with a very high mortality rate and molecular pathogenic pattern completely different from NSCLC.

The NSCLC category can be divided in different histological types, among which the most important are the squamous cell carcinoma and the adenocarcinoma.

The adenocarcinoma classification was recently revised [49] with the abolition of the bronchioloalveolar carcinoma and mixed subtype adenocarcinoma. By contrast, invasive adenocarcinomas are classified by predominant pattern after using comprehensive histologic subtyping with lepidic, acinar, papillary, and solid patterns.

Adenocarcinoma incidence is increased in the last two decades, in parallel with the decrease in the incidence of squamous form [47]. In Europe, they account for 30% of bronchopulmonary

carcinomas while in North America they already are the most common histological type (about 50%). The increased incidence is probably related to the use of filter cigarettes and the consequent reduction in the size of the particulates and increase of nitrates. It has also been described the association with pre-existing lung scars caused by eg tuberculosis or pulmonary infarcts.

Worldwide, tobacco smoking is associated with more of 90% of cases of lung cancer [47] . In more developed countries, the incidence and mortality rates are generally declining, reflecting previous trends in smoking prevalence. However, in less developed countries lung cancer rates are predicted to continue to increase due to endemic tobacco use [50] . There is a clear role for genetics since only 15% of lifetime smokers develop lung cancer and 10% of lung cancers occur in never-smokers especially in women [51] and in Asiatic women in particular [52] .

One important hypothesis in these categories of patients is the presence of a genetic background that can predispose to lung cancer in the absence of external environmental pollutants [52].

In this field the two most important recognized oncogenes in tumour developing and growing are EGFR e KRAS.

Epidermal growth factor receptor (EGFR or ErbB-1) is a tyrosine kinase receptor belonging to the ErbB family; it is widely expressed in NSCLC (40-80% of NSCLC) and plays a significant role in carcinogenesis through improper activation of EGFR Tyrosin-Kinase (TK) domaine which results in increased malignant cell survival, proliferation, invasion and metastasis. The TK activity of EGFR may be dysregulated by various oncogenic mechanisms, including the presence of harbored mutations in the TK domain of EGFR [53-55]. Overall, the frequency of EGFR mutations is 5–20%, depending on the populations studied [56].

RAS is one of the important molecules in the downstream of EGFR signaling pathway. Wild-type forms of ras proteins have GTPase activity and their activation is strictly dependent on the bond with alternative forms of GDP (inactive) or GTP (active), allowing tight control of signal transduction [57]. These proteins acquire oncogenic function through point mutations at codons 12, 13, or 61 of exon 2 that abolish the GTPase function giving therefore a constitutionally active

conformation and deregulating thereby cell proliferation. For these reasons mutated *Ras* genes, especially *K-ras*, observed among 20~30% NSCLC patients, have been implicated in the pathogenesis and prognosis of lung cancer [58].

In 1991 Slebos and after ten years Ahrendt found that k-ras mutations are found almost exclusively in patients with lung adenocarcinoma and smoking history. In addition, Feng reported that exposure to this carcinogen induces alterations at the codon 12 of K-ras in the human bronchial epithelium, suggesting a direct molecular mechanism of smoking in human lung carcinogenesis. Moreover NSCLC patients with *K-ras* mutations are associated with unfavorable prognosis [59].

Concerning lung cancer treatment, surgery actually represents the cornerstone of therapy for NSCLC even if global resectability still ranges from 10 to 25% [60]. With strict pre-operative selection criteria, the healing rate in patients undergoing complete resection for NSCLC in early stage (I-II) ranges between 40 and 70% [61]. However, even in these cases, the possibility of loco-regional and/or distance recurrences is not an uncommon event, representing the main factor affecting long-term survival.

For this reason, the use of integrated approaches with adjuvant regimen of chemo- and / or radiotherapy has been recently introduced even in early stages in order to reduce this risk [62]. These integrated approaches have enabled a further increase in 5 year overall survival, but according to most authors, new therapeutic protocols based on traditional chemotherapy does not seem to offer further opportunities for improvement of results [63].

In addition to their carcinogenic activity, EGFR and KRAS have emerged as two of the most relevant targets for cancer treatment.

Gefitinib and erlotinib are small-molecule reversible tyrosine kinase inhibitors (TKIs) that selectively target EGFR. The presence of somatic mutations in the tyrosine kinase domain of *EGFR* (exon 18-21) involving the ATP-binding pocket of the receptor was shown to represent the most important predictive marker [53-55]. For these patients, the response rate to gefitinib

and erlotinib is approximately 75%, suggesting that these mutations, at least in part, drive malignant transformation [56,64].

In contrast, somatic mutations in exon 2 (codon 12-13) of *KRAS* have been associated with primary resistance to EGFR inhibitors. *EGFR* and *KRAS* mutations seem to be mutually exclusive, which is consistent with the idea that different alterations of the same pathway are involved in lung carcinogenesis [65].

Regarding lung cancer long-term results, TNM staging system [66], is currently the most important and accurate prognostic instrument. However, it still presents significant difficulties in predicting the effectively outcome of individual patients. Some authors has recently proposed the use of PET-CT scan, and in particular the maximum standard uptake value (SUVmax), as new tool to predict tumor biologic aggressiveness [67]. It is therefore understandable how a greater knowledge of molecular and histological cancer phenotypes associated to patient characteristics, able to predict individual outcomes and treatments, is therefore desirable.

1.8 COPD AND LUNG CANCER

It is becoming increasingly evident that COPD and lung cancer are strictly related.

Indeed COPD has recently established as important risk factor for developing lung cancer, even independently from the effects of smoking. Indeed, it has been shown that the risk of lung cancer is increased in smokers with COPD up to 6-fold compared to smokers with comparable cigarette exposure, but without COPD [68,69]. It has also been shown that the presence of COPD increased lung cancer mortality even among nonsmokers [70].

In addition, lung cancer is also a leading cause of morbidity and mortality in patients with COPD as 33% of patients died of lung cancer over a 14.5-year follow-up [71]. Furthermore 50–70% of patients diagnosed with lung cancer have spirometric evidence of COPD [72]]. As a practical consequence of the epidemiological associations between COPD and lung cancer, an important question is whether the relationship between lung cancer and COPD is subtype specific. A small

case – control study [73] showed that airflow obstruction is primarily a risk factor for squamous cell lung cancer (odds ratio = 3.49, 95% CI = 1.63 to 18.5; $P = .006$), whereas symptoms of chronic bronchitis without COPD is a risk factor (risk greater than fourfold) for adenocarcinoma of the lung. In a subset analysis, having concurrent bronchitic symptoms and COPD was associated with a more than threefold increased risk for squamous cell carcinoma. Despite these striking epidemiological associations, the mechanisms by which COPD can be linked to lung cancer are poorly understood.

1.8.1 Risk factors

Primarily, the main risk factor for the onset of both diseases is cigarette smoking, but they probably also share a common familial component and environmental risk factors other than smoking. This supposition is reinforced by the fact that only a fraction of smokers (around 15%) will develop lung cancer and/or COPD in their lifetime [1,74] which suggests a different individual susceptibility to the risk of lung cancer and/or COPD or time of disease onset.

Some common elements in the pathogenesis of COPD and lung cancer include oxidants, familiar and genetic predisposition (p53, Rb, K-ras), peptides and endopeptidases (bombesin-like), dysregulation of growth factor expression and among others inflammation.

While the mechanism of cigarette smoking damage in COPD has been previously treated, for lung carcinogenesis, the action is probably related to DNA mutations induction.

Carcinogenesis is a complex process characterised by the accumulation of multiple independent genetic alterations, often involving overexpression of oncogenes and loss of tumour suppressor genes. These genetic alterations disrupt the normal regulation of cell signalling pathways, essential for the control of cell growth, differentiation and apoptosis [75]. The observation of familial aggregation of emphysema dates back more than 200 years, and studies conducted 30 years ago reveal a familial aggregation of lung cancer associated with COPD that is not explained by α 1-antitrypsin genotype or smoking history [76,77]. Familial aggregation is well

documented for multiple cancers, although it has been difficult to separate the contributions of genetic differences and environmental exposures in the development of lung cancer.

A number of linkage studies [78] have investigated genes involved in both lung function and COPD and candidate susceptibility genes involved in COPD and lung cancer. These studies have led to the identification of markers on chromosome 6q for lung cancer and abnormal lung function and on chromosome 12 for lung cancer, COPD, and lung function, as well as numerous other candidate susceptibility genes involved in detoxification, immune regulation, matrix remodeling, DNA repair, cellular proliferation, and tumor suppression [78]. Among the genetic connections between COPD and lung cancer, it is particularly interesting to note that both the Z and the S alleles of the α 1-antitrypsin gene are more common in patients with lung cancer compared with the general population, as is a polymorphism in the neutrophil elastase gene [79,80], suggesting that an imbalance between neutrophil elastase and α 1-antitrypsin may contribute to the development of both COPD and lung cancer [81].

The role of epigenetic modifications in the common pathogenesis of COPD and lung cancer also deserves attention.

Although there is growing evidence implicating DNA methylation, histone deacetylation, and protein phosphorylation in lung cancer pathogenesis [82,83], this knowledge is only now being applied to COPD, alone or when associated with lung cancer [84].

It is plausible that several candidate risk genes and pathways identified by lung cancer studies may be shared by these two diseases and could constitute potential targets for the newly developed drugs (eg, demethylating agents and histone deacetylase – inhibiting agents) that modify epigenetic alterations.

The inflammatory paradigm is undoubtedly one of the most fascinating theories to connect COPD and lung cancer and it has acquired new impetus by the recent discoveries in the COPD pathogenesis.

1.8.2 The inflammatory theory

As described above, COPD is characterized by the accumulation of macrophages, CD4+ and CD8+ T cells, dendritic cells, B cells and neutrophils, particularly in smaller airways and lung parenchyma, and the severity of COPD is associated with the degree of infiltration by these inflammatory cells [1].

Moreover, chronic inflammation may play a significant role in the pathogenesis of lung cancer as a tumour promoter. A causal relation between inflammation and cancer was initially proposed by Galen and later by Virchow [85], who noticed the infiltration of leucocytes in malignant tissues and suggested that cancers arise at regions of chronic inflammation. The hypothesized carcinogenetic mechanism in COPD patients states that cigarette smoke initially creates a direct parenchymal damages that up-regulates the cytokines production, such as interleukin (IL)-1b, and the Th-1 cytokines, which can promote the inflammatory response through typical COPD T-lymphocytes and finally resulting in an overproduction of cytokines, such as IL-6, IL-8, and IL-10. Some of the latter mediators can inhibit apoptosis, interfere with cellular repair, and promote angiogenesis. All in all, chronic inflammation may play a pathogenic role in lung cancer by amplifying the initial mutagenic damage and enhancing both tumour growth and metastasis. IL-8, for example, has been shown to be pro-oncogenic, such as releasing BCL-2, but also suppresses oncogenes, such as p53, hence minimizing apoptosis while inducing cell transformation. In addition, among mechanisms that regulate the progression or suppression of tumor growth, the role of cellular senescence is becoming increasingly appreciated. Cellular senescence permanently arrests cell growth in tissues at risk for malignant transformation, particularly those that experience prolonged inflammation, such as the airways and the lung parenchyma in smokers with COPD. On the other hand, senescent cells undergo profound modifications which can have deleterious effects on the tissue microenvironment. The most significant of these alterations is the acquisition of a senescence-associated secretory phenotype (SASP) that promotes the secretion of several proinflammatory cytokines (particularly IL-6 and IL-8), growth factors and metalloproteinases, which could eventually promote tumor progression

and invasiveness. Of note, senescent cells have the ability to influence the populations of macrophages and lymphocytes infiltrating nascent tumors, shifting the balance from cell populations associated with tumor suppression (M1 macrophages, Th1 T-lymphocytes) to those that instead promote tumor progression (M2 macrophages, Th2 and T regulatory lymphocytes). The mechanism of perpetuation of damage typical of COPD could also explain in these patients the risk of lung cancer irrespective of active smoking [70]. In fact, after quitting smoking, lung cancer risk remains increased in patients with COPD, though this risk is superior in those who continue to smoke [70]. The close connection between COPD, inflammation, and lung cancer is even more apparent in light of recent findings that point to a possible relationship between inhaled corticosteroids and reduced lung cancer risk in COPD patients [86,87].

Complementary to this is the observation that the incidence of lung cancer is associated with specific stages of COPD severity. Lung cancer is assigned as the cause of death in 33% of patients with mild-to-moderate COPD with a decreased in patients with more severe disease [88,89].

According to this data, in our experience we did not observed cases of lung malignancy in patients with end-stage COPD candidates to lung transplantation, differently from other end-stage chronic inflammatory diseases such as scleroderma, sarcoidosis and especially pulmonary fibrosis.

For these reasons our research group has proposed a pathogenetic hypothesis that, in agreement with recent autoimmune theory, attributes to the peculiar inflammatory phenotype of COPD advanced stages a protective and surveillance role against neoplastic development.

As previously reported, the T regulatory cells and the associated cytokine pattern play a strategic role in the interaction between COPD and lung cancer, although a complete understanding of the process is still far away.

In this field, the analysis of TH17 cells and the IL17 cytokine could really represent the link between these two important diseases.

1.8.3 Pathway axys IL17- R23

Th17 cells are a critical component of the adaptive immune response and have also been implicated in chronic inflammatory diseases and autoimmune diseases [90], although their primary function appears to be the clearance of pathogens that are not adequately handled by Th1 and Th2 cells, especially extracellular bacteria and fungi [91]. The cytokine microenvironment influences the differentiation of naive T cells into Th17 cells. The Th17 lineage depends on the presence of TGF- β 1 and IL-6, found in elevated concentrations particularly in chronic obstructive pulmonary disease (COPD) [92].

In effect, an increase in Th17 cells was observed in patients with COPD compared with current smokers without COPD and healthy subjects. Human Th17 cells differ from other cell subsets in their potency to induce proinflammatory cytokines in bronchial epithelial cells, airway fibroblasts and smooth muscle cells and above all they secrete the cytokine IL-17 [93], which plays an important proinflammatory role in COPD.

The IL-17 family comprises of six members (IL-17A–17F) [94] and five receptors (IL-17RA–17RE) [95].

IL-17RA is the largest member of the IL-17R family and at least four ligands are mediated through this subunit [96].

The expression of IL-17RA is up-regulated by cytokines such as IL-15 (produced by a range of non-T-cells, including macrophages and IL-21 [97] on CD8⁺ T-cells, and so may play an important role in COPD.

IL-23 plays a key role in the maintaining and expanding the Th17 cell lineage over time to release IL-17 [98] and is secreted from APCs such as dendritic cells. IL23 is released in response to inflammatory signals and therefore is continually expressed in chronic inflammation.

IL-17 induces epithelial cells to produce antimicrobial peptides (such as β defensins) and numerous chemokines, such as TNF- α , IL-1 β , IL-6, GM-CSF, granulocyte colony-stimulating factor and IL-8 [99]. All these findings clearly indicate that IL-17 is able to generate an

intriguing crosstalk between the adaptive and innate immune systems, regulating an efficient immune response (Fig. 1.10) [100].

Many studies found an increased production of IL-17, especially IL-17A in COPD patients although its involvement in the development of COPD is poorly understood [101].

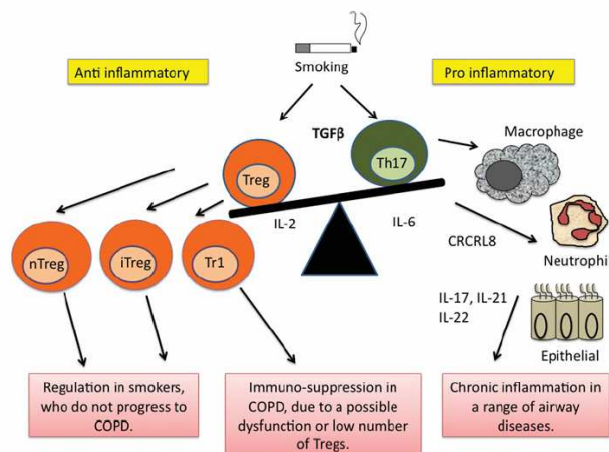


Fig. 1.10. Schematic representation of the balance between Treg-cells and Th17 cells in COPD. The type of immune response activated relies on the cytokine environment and the individual's immune system, leading to varied clinical outcomes. In some situations, there can be a switch of lineage from Treg-cells to Th17. TGF- β and IL-6 are crucial in the transition between lineages (From Lane N. (102).

In recent years, Th17 cells and IL-17A have been also associated with various human tumors, including NSCLC [103-105]. In particular the number of Th17 cells detected in tumors has been shown to positively correlate with microvessel density, as well as it was observed that high levels of IL-17A expression in tumor cell lines promoted angiogenesis, lymphangiogenesis and cell proliferation [106]. The action of IL-17 seems closely related to induction of VEGF. In fact, Li [107] has shown that the elevated expression of this growth factor in supernatant of adenocarcinoma cell lines were IL-17a concentration-dependent. In addition he found higher serum levels of IL-17a in patients with lung adenocarcinoma compared to healthy controls and higher positive expressions of IL-17a, IL-17Ra and VEGF in lung adenocarcinoma lesion tissues than pericancerous normal tissues.

Liu [108] has recently proposed a new potential mechanism of lung carcinogenesis for IL-17 through the formation of an M2-macrophage–dominant tumor microenvironment in non–small-cell lung cancer. Based on the finding that lung tumor tissues expressed significantly higher

levels of IL-17 and prostaglandin E2 (PGE2) than normal lung tissues, he speculated on the possibility that IL-17 served as a chemoattractant to recruit macrophages into the lung tumors while PGE2 induced differentiation of M2 macrophages.

Based on these fundamental premises, lung cancer associated with COPD seems to possess specific pathogenetic and morphological features, differently from cancers resulting from simple smoke damage.

Therefore, a better understanding of these phenotypes could lead to important consequences of great relevance in therapy and care of patients affected by COPD and lung cancer.

2. AIM OF THE RESEARCH

This research project focuses on the study of lung cancer in patients with COPD compared to control groups, composed by healthy smokers and non-smoking patients in order to identify eventual distinct biohumoral, morphological and molecular phenotypes.

3. MATERIALS AND METHODS

3.1 STUDY POPULATION

From January 2010 to December 2012 we prospectively enrolled in the study patients submitted to anatomical surgical resection for non small cell lung cancer (NSCLC) in the Thoracic Surgery Unit of Padua and Strasbourg.

Inclusion criteria were the follows:

- Patients undergoing major resections (lobectomy or pneumonectomy) associated with hilar-mediastinal lymphadenectomy; this decision was motivated from the need to have sufficient non neoplastic lung parenchyma to study the peri-neoplastic tissue remodeling.
- Patients with peripheral lung nodule.
- Patients with suspected or known primary NSCLC.

Exclusion criteria were the follows:

- Patients with inflammatory lesion or with metastatic pulmonary cancer.
- Patients with NSCLC involving central airways in order to avoid distortions of lung architecture or presence of local and/or systemic inflammatory reactions.
- Patients with chest wall involvement.
- Patients previously treated by chemo and / or radiation therapy.
- Past history of asthma or allergic rhinitis and acute upper respiratory tract infections.
- For COPD patients, presence of exacerbations during the month preceding surgery.

All patients included in the research project were subjected full clinical/biohumoral and functional analysis.

Clinical study

For each subject a detailed medical history with particular attention to respiratory symptoms, smoking history, concomitant medications and comorbidities was collected.

Venous blood samples, preserved in Ethylene Diamine Tetraacetic acid, were sent to the central laboratory to evaluate the levels of erythrocyte sedimentation rate and C-reactive protein as known systemic inflammation signs.

All subjects underwent pulmonary function tests (PFTs) that were performed and evaluated by an expert pneumologist following the most recent guidelines [109].

Briefly, the following parameters will be measured: vital capacity (CV), inspiratory capacity (IC), forced vital capacity (FVC), forced expiratory volume in the first second (FEV1), FEV1 to FVC ratio (FEV1/FVC), functional residual capacity (FRC), residual volume (RV) and total lung capacity (TLC). To assess the reversibility of the airway obstruction in subjects with a baseline FEV1/FVC <70 %, measurements will be repeated 15 minutes after the inhalation of 400 µg of salbutamol. The diffusing capacity for carbon-monoxide (DLCO) will be assessed using the single breath technique.

Arterial haemogasanalysis was performed to investigate gas exchange (oxygen partial pressure-PaO₂, carbon dioxide partial pressure-PaCO₂ and haemoglobin saturation-SHb).

A complete radiological assessment, including chest-x ray, Chest and Abdomen CT scan was performed in every patients both for oncological and research reasons.

In addition patients routinely underwent ¹⁸FDG PET-CT scan with measurement of primary lung tumor maximal standard uptake value (SUV_{max}), that is SUV on the highest image pixel in the parenchymal tumor regions.

In all patients bronchoscopic examination with bronchoalveolar lavage (BAL) was carried out. BAL, blood and fresh cancer tissue were preserved in frozen tissue bank for future immunohistochemical and molecular studies.

The collection and preservation of biological materials for frozen bank followed the guidelines of Regione Veneto and Azienda Ospedaliera di Padova.

Patients enrolled in the study were then divided **in 3 groups** according to respiratory functional tests and smoking history: **COPD patients, smoker patients without COPD and never-smoker subjects (with normal lung function)**.

COPD subjects were identified by the presence of fixed airflow limitation (forced expiratory volume in the first second (FEV1) to forced vital capacity (FVC) ratio < 70%). They will be subsequently staged on the basis of FEV1 values discriminating smokers with severe COPD (FEV1<50% predicted) from those with mild/moderate COPD (FEV1>50% predicted).

3.2 SURGICAL PROCEDURE AND LUNG SAMPLING

As previously reported, every patient underwent an anatomical resection associated with systematic lymphadenectomy.

Immediately after the surgical procedure, samples from the nodule and at least three from non-tumour subpleuric area (2 cm³) were preserved in RNAlater[®] and then stored in liquid nitrogen for molecular studies. The rest of lung parenchyma was sent to the Pathological Anatomy Laboratory and perfused using 4% formaldehyde via bronchial tree with constant pressure of 25 cm of water (Fig. 3.1).

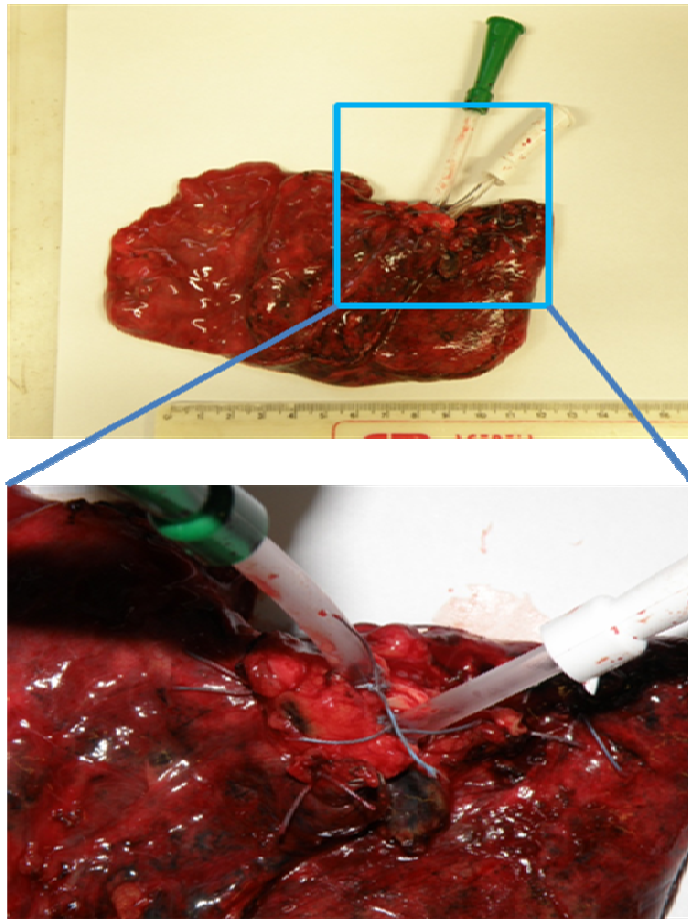


Fig. 3.1. Intra-operative view of a lobe specimen after resection with bronchial incannulation to perform selective perfusion using 4% formaldehyde with constant pressure of 25 cm of water.

The lung was then taken in a large container of formalin for at least 24 hours. This method of lobe perfusion allowed to obtain adequate lung samples for microscopic and morphometric diagnosis, avoiding on the one hand emphysema overestimation due to alveolar hyperdistensions and from the other areas of parenchymal atelectasis. Four samples were taken both from neoplastic areas and non neoplastic sites. After dehydration, samples will be embedded in paraffin and serial sections were cut for histological, immunohistochemical and molecular studies.

3.3 HISTOLOGICAL ANALYSIS

All histological and immunoistochemical parameters were blindly evaluated by the same pathologist (FC)

3.3.1 Pathological diagnosis

Tissue sections were stained with haematoxylin and eosin and Masson's stains. Histopathological evaluation of the tumor were done by an expert Pathologist according to the latest revision of lung cancer classification [48]: in particular, in case of adenocarcinoma, a precise quantification of the different histological pattern (lepidic, acinar, papillary and solid) percentage was done [49].

pTNM staging was conducted in accordance with the 7th edition of 2009 [66].

3.3.2 Morphological evaluations

In all neoplastic lesions were evaluated and quantified the following parameters:

- Inflammation, necrosis and fibrosis were quantified by an expert pathologist in peritumour and intratumour areas and the amount was expressed both semiquantitatively using a score system from 0 to 3 (no positivity, 0; mild, 1; moderate, 2; strong, 3) both as percentage of extension: 1: <30% of neoplastic and perineoplastic areas; 2: from 30 to 60% and 3>60%.

- Four μm -thick sections were cut and processed for immunohistochemical analysis of ki-67. Briefly after dewaxing and hydration, sections were incubated in citrate buffer 5 mM at pH 6.0 in a microwave oven for 30 min for antigen retrieval. Afterwards, sections were treated with blocking serum (Ultratech HRP Kit, Immunotech, Beckman Coulter, USA) and incubated for 60 min with the primary monoclonal antibody anti-ki-67 (MIB-1, Immunotech, Marseille, France) at a concentration of 1:50. Sections were subsequently incubated with a secondary biotinylated antibody for 10 min and then with streptavidin-biotin complex conjugated to horseradish peroxidase for 10 min (Ultratech HRP Kit, Immunotech, Beckman Coulter, USA). Immunoreactivity was visualized with diaminobenzidine (DAB, Dako, Glostrup, Denmark). Finally, the sections were counterstained with Mayer's hematoxylin. Negative controls for nonspecific binding were processed omitting the primary antibodies and

revealed no signal. Ki-67 positivity was evaluated by using an assisted computerized morphometric analysis counting at least 100 cells in the most representative areas. Data were expressed as number of positive cells/total cell count % and ki-67 quantification represented the ki-67 labeling index.

3.3.3 Characterization of the peri-and intra-tumor IL-17 cytokine pattern

For immunohistochemistry, deparaffinized sections were treated with 3% hydrogen peroxide in phosphate-buffered saline (PBS) for 10 minutes and incubated for 30 minutes at 90°C for antigen retrieval. Next, the specimens were individually incubated for 12 hours at 4°C with the polyclonal antibody diluted in 2% bovine serum albumin before use anti-IL-17 (1:100) (AbCAM). The sections were washed and incubated with the streptavidinperoxidase conjugate (Dako) for 30 minutes.

The slides were washed with HRP labeled polymer anti-rabbit (Dako) for 30' and immunoreactivity was visualized with 3-3'-diaminobenzidine (Dako). Negative controls for nonspecific binding were processed omitting the primary antibody and revealed no signal. The sections were counterstained with hematoxylin.

Data were expressed using a combined score from staining intensity (graded with score 1: mild, score 2: moderate, and score 3: strong) and percentage of positive cells.

3.4 GENETIC ANALYSES

All paraffin-embedded samples were evaluated by a pathologist in order to assess the tumour tissue quality and quantity. Molecular analysis was performed when the tumour tissue comprised more than 50% of the entire sample. DNA was extracted using a QIAmp DNA kit (Qiagen, Milan, Italy), according to the manufacturer's instructions. Agarose gel electrophoresis was used to control its quality and quantification was performed by spectrophotometry.

A total of 100 ng of genomic DNA was used for polymerase chain reaction (PCR) to amplify exons 18, 19, 20, 21 of *EGFR* gene and exon 2 of *KRAS* gene by using primers, sequences and amplification conditions according to previous studies (8, 17). PCR products were purified using a pre-sequencing kit (Amersham Biosciences, Little Chalfont Buckinghamshire, UK). Subsequently, they were sequenced with both forward and reverse primers using BigDye Terminator v1.1. Cycle Sequencing Kit (Applied Biosystems, Foster City, CA, USA). Automated sequencing was carried out using an ABI 3130xl DNA sequence (Applied Biosystems).

3.5 ELECTRONIC DATABASE

A multidisciplinary electronic database (Excel, Microsoft Office Professional 2007) was built following the privacy rules. The database contains all the clinical, radiological, functional, pathological and molecular data and was used for statistical analysis.

The database did not include any personal identification label and was held in secure password protected storage under the responsibility of the Unit Coordinators (Prof. Rea, Prof. Saetta and Prof. Calabrese) in accordance with the requirements of the Health Information.

The study, conforming to the Declaration of Helsinki, was approved by our local ethic committee and patients signed an informed consent

3.6 STATISTICAL ANALYSIS

All cases were coded and the measurements were made without knowledge of clinical data.

Normality of distribution for quantitative variables was assessed by means of Shapiro-Wilcox statistics. Non normal variables were log-transformed. Group data were expressed as mean and SEM, or as median and IQR when appropriate. To compare means from two different groups we

used the unpaired 2-tail Student's t test or one-way analysis of variance (ANOVA) for normal data; for non-normal data Mann-Whitney's U test or Kruskal-Wallis test were performed.

Categorical data were analyzed by the chi-square test.

Statistical analysis was performed using SAA statistical software version 9.1 (SAS institute, Carry, NC, USA). P values lower than 0.05 were considered statistically significant.

RESULTS

4.1 POPULATION CHARACTERISTICS

In the study period (January 2010 – December 2012) 74 patients were initially enrolled for the research, of which 66 have met the selection criteria, as in 8 cases a histological diagnosis of secondary malignancy was found.

According clinical/functional inclusion criteria the three study groups included: 16 COPD patients, 32 smokers without COPD and 18 never smokers.

The most important clinical data of the three groups are listed in table 4.1.

	COPD (16)	Smokers (32)	Never Smokers (18)	p
Age (years)	72 ± 7	68 ± 9.6	64 ± 12	NS
Sex (M/F)	13/3	25/7	6/12 #@	0.002
BMI (Kg/m ²)	27 ± 4	27 ± 4	27 ± 4	NS
Smoke Pack-year	46 (IQR 33-55)	35 (IQR 15-48)	0	<0.001
FEV1 (% of predict)	71 ± 14 #	93 ± 19	114 ± 19	<0.001
FEV1/VC (%)	58 ± 9 *#	75 ± 7	81 ± 6	<0.001
FEV1/FVC (%)	62 ± 10*#	81 ± 6	85 ± 8	<0.001
FVC (% of predict)	91 ± 18	90 ± 17	111 ± 17 #@	<0.001
RV (% of predict)	122 ± 18 *#	106 ± 18	106 ± 19	0.015
RV/TLC (%)	113 ± 17	109 ± 16	102 ± 15	NS
PaO2 (mmHg)	72 ± 7	75 ± 7	74 ± 7	NS
PaCO2 (mmHg)	39 ± 3	38 ± 3	37 ± 3	NS
WBC (x 10.9/L)	8.1 ± 3.2	7.7 ± 2.3	7.5 ± 2	NS
RBC (x 10.12/L)	4.8 ± 0.5	4.5 ± 0.6	4.6 ± 0.7	NS
Basophil cell (x 10.9/L)	0.04 (IQR 0.03-0.06)	0.02 (IQR 0.02-0.04)	0.02 (IQR 0.02-0.03)	0.002
ESR (mm/h)	28 ± 14	33 ± 20	38 ± 27	NS
CRP (mg/L)	5.9 (IQR 3-10.1)	3.5 (IQR 2.9-17.8)	3 (IQR 2.9-14.7)	NS
Adenocarcinoma histology (%)	73	61	56	NS

* COPD vs Smokers p < 0.05,

COPD vs Never smokers p < 0.05,

@ Smokers vs Never smokers p < 0.05

Table 4.1. Main demographic, clinical and functional patients characteristics. Data are expressed as mean and SEM, except Smoking pack-year, Basophil cell amount and CRP reported as median and IQR.

The three study groups presented homogeneous demographic characteristics (age, height, weight, BMI) except for sex, with a significant greater proportion of women in the never-smoker group. Regarding smoking history, COPD and smoker patients were comparable.

Concerning PFTs, as expected, COPD group evidenced a statistically reduction in most of the respiratory functional parameters (FEV1%, FEV1/VC, FEV1/FVC, PEF%, TLC%, RV%, FRC%) even if the Motley index (RV/TLC), sign of pulmonary hyperinflation from emphysema, did not differ in the three populations. Conversely, FVC% was higher ($111 \pm 17\%$) in never smoker patients compared to smokers without COPD ($90 \pm 17\%$, $p < 0.001$) and COPD patients ($91 \pm 18\%$, $p = 0.003$). According to GOLD staging for COPD patients, we reported 3 stage I and 7 stage II. Blood tests analyses did not show significant changes even for systemic inflammatory markers; the only altered value is an increase in percentage of basophil cells in COPD patients compared to smokers. The choice to include in the study only patients with peripheral lung cancer has undoubtedly led to a selection bias, affecting the predominant histologic profile. Indeed, in all the three populations there is a clear predominance of the glandular form, with percentages ranging from 63% in the COPD group, to 71 % in the smokers and 56% in never-smokers ($p = ns$). In order to obtain results not affected by different histotypes, we decided to analyze tumor functional, molecular and morphological assessment only in the prevalent histology, adenocarcinoma. Therefore, the study population was composed of 43 patients, of which 10 COPD patients, 23 smokers without COPD and 10 non-smokers. As shown in Table 4.2, the main demographic and functional parameters reflected those of the previous population, not showing significant differences among the 3 groups except for the male/female ratio, which result reversed in never smokers and for lung function, reduced in COPD patients. Even in this case, FVC% was higher ($115 \pm 11\%$) in never smoker patients compared to smokers ($93 \pm 17\%$, $p = 0.001$) and COPD ($91 \pm 19\%$, $p = 0.003$).

	COPD (10)	Smokers (23)	Never smokers (10)	P
Age (years)	71 ± 7	68 ± 7	68 ± 9	NS
Sex (M/F)	8/2	19/5	3/7 [@]	0.02
BMI (Kg/m ²)	27 ± 4	27 ± 4	28 ± 4	NS
Smoke Pack-year	47 ± 29	39 ± 32	0	0.001
FEV1 (% of predict)	71 ± 15 [*]	95 ± 19	119 ± 13	<0.001
FEV1/VC (%)	58 ± 8 [*]	74 ± 8	80 ± 3	<0.001
FEV1/FVC (%)	60 ± 10	80 ± 7	87 ± 10	<0.001
FVC (% of predict)	91 ± 19	93 ± 17	115 ± 11 [@]	0.003
RV (% of predict)	123 ± 17 [#]	107 ± 19	102 ± 20	0.059
RV/TLC (%)	114 ± 19	108 ± 18	98 ± 14	NS
PaO ₂ (mmHg)	75 ± 6	76 ± 7	74 ± 7	NS
PaCO ₂ (mmHg)	40 ± 3	39 ± 2	37 ± 3	NS
WBC (x 10.9/L)	7.3 ± 2.5	7.7 ± 2.7	7.8 ± 2.6	NS
RBC (x 10.12/L)	4.7 ± 0.5	4.5 ± 0.6	4.6 ± 0.6	NS
Basophil cell (x 10.9/L)	0.03 (IQR .03-0.06) [#]	0.02 (IQR 0.02-0.03)	0.02 (IQR 0.02-0.03)	0.048
ESR (mm/h)	27.6 ± 14	32.9 ± 21	33 ± 34	NS
CRP (mg/L)	6.4 (IQR 4.1-9.9)	3.4 (IQR 2.9-6.3)	3 (IQR 2.9-3.4)	NS
SUV max	5.83 ± 3.91	9.46 ± 5.77	7.31 ± 4.96	NS

* COPD vs Smokers p < 0.05,

COPD vs Never smokers p < 0.05,

@ Smokers vs Never smokers p < 0.05

Table 4.2. Main demographic, clinical and functional patients characteristics in patients with adenocarcinoma histology. Data are expressed as mean and SEM, except Basophil cell amount and CRP reported as median and IQR.

The analysis of tumor functional assessment through ¹⁸FDG PET-CT scan did not evidence significant differences in the three populations with values of mean SUVmax in COPD patients, smokers and never smokers group of 5.83 ± 3.91, 9.46 ± 5.77 and 7.31 ± 4.96, respectively.

Although not statistical significant, a strong trend of increase in the SUV max in smoker-related tumours compared to COPD-related cancers (9.46 ± 5.77 vs 5.83 ± 3.91, p=0.08) was observed.

4.2 HISTOLOGIC EVALUATION

According to WHO guidelines, a diagnosis of adenocarcinoma histology was found in all patients. In one case of never smokers group an additional well-differentiated stage I neuroendocrine tumor was observed in tissue specimen.

The maximal lung tumour diameter was measured and the mean values for each group was 3.11 ± 1.3 cm (COPD patients), 3.2 ± 0.1 cm (smokers) and 3.1 ± 1.1 cm (never smokers), without statistical difference.

Pathological staging of all patients was reported in table 4.3 and revealed a comparable stratification between the groups with a majority of early stages (p=n.s.).

Stage IIIA was due to mediastinal nodal involvement not evidenced at pre-operative investigation, except in one never smoker patient that was staged T4 for an ipsilateral lung nodule. The smoker patient without COPD with stage IV underwent to left upper lobectomy for two lung metastasis of a previous resected adenocarcinoma.

	COPD	Smokers	Never smokers
Stage IA	4 (40%)	5 (22%)	4 (40%)
Stage IB	3 (30%)	10 (43%)	1 (10%)
Stage IIA	1 (10%)	2 (9%)	3 (30%)
Stage IIB	1 (10%)	2 (9%)	0
Stage IIIA	1 (10%)	3 (13%)	2 (20%)
Stage IIIB	0	0	0
Stage IV	0	1 (4%)	0

Table 4.3. Stage distribution in study population without significant differences in distribution.

The evaluation of the histological analysis was reported in Table 4.4.

	COPD	Smokers	Never Smokers	P
Lepidic pattern (%)	25 (IQR 10-30)	0 (IQR 0-10)	25 (IQR 0-30)	0.01
Mucinous Lepidic Pattern (%)	0±0	4±14	0±0	NS
Acinary pattern (%)	44±22	44±32	49±26	NS
Papillary pattern (%)	0 (IQR 0-0)	0 (IQR 0-13)	15 (IQR 10-20)	NS
Solid pattern (%)	10 (IQR 0-20)	20 (IQR 6-70)	0 (IQR 0-0)	0.045
Ki67/MIB1 (%)	30±30	50±25	24±18	0.013
Necrotic remodelling	18±23	20±23	2±4	NS
Inflammatory remodeling	28±22	30±22	33±20	NS
Fibrotic remodelling	24±20	28±22	28±18	NS

Tab 4.4. Main histological results in three study populations. Data are expressed as mean and SEM, except lepidic, papillary and solid pattern reported as median and IQR.

We can observed a significant increase of the lepidic pattern (Fig. 4.1) percentage in COPD patients and never smokers tumours compared to smokers (25% (IQR 10-30%) vs 25% (IQR 0-30%) vs 0% (IQR 0-10%), respectively, p=0.01).

The mucinous lepidic variant was found only in smokers but no differences were observed with the other groups (0±0% vs 4±14 vs 0±0, respectively, p=NS).

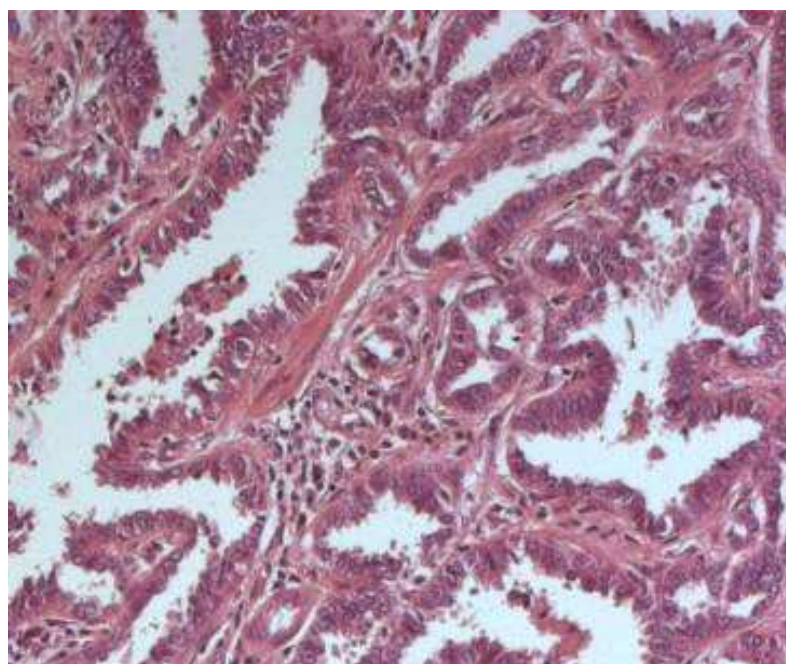


Fig. 4.1. Lepidic pattern of invasive adenocarcinoma: tumor cells, type II pneumocytes and Clara cells, grow along the surface alveolar walls.

At the same time, an increase of the solid component (Fig. 4.2) in smokers compared with other groups (20% (IQR 6-70%) vs 10% (IQR 0-20%) in COPD patients vs 0% (IQR 0-0%) in never smokers, $p=0.045$) was present.

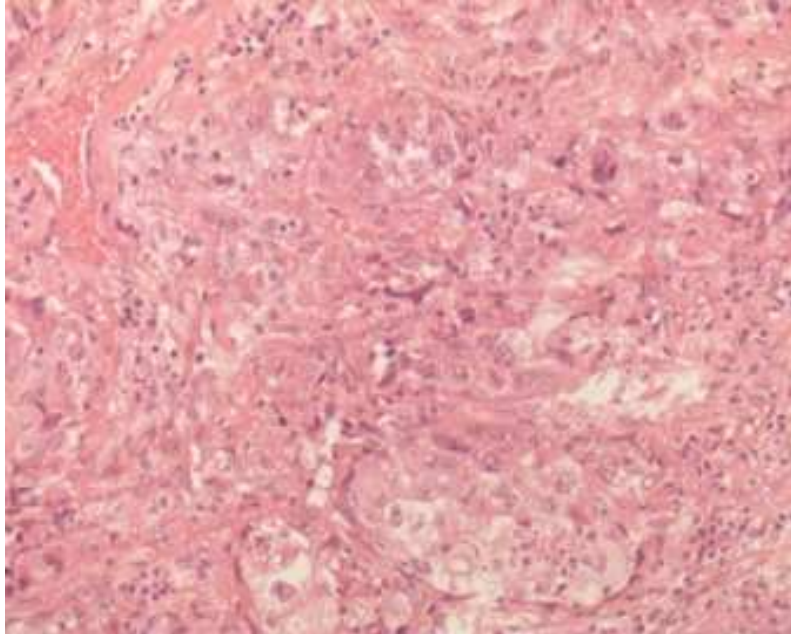


Fig. 4.2. Solid pattern of invasive adenocarcinoma: sheets of tumor cells with abundant cytoplasm and mostly vesicular nuclei with several conspicuous nucleoli.

No differences were observed for the intermediate histological components, the acinar ($44\pm 22\%$, $44\pm 32\%$, $49\pm 26\%$ for COPD patients, smokers and never smokers, respectively) and papillary pattern, including also micropapillary variant (0% (IQR 0-0%), 0% (IQR 0-13%) and 15% (IQR 10-20%) for COPD patients, smokers and never smokers, respectively) (Fig. 4.3).

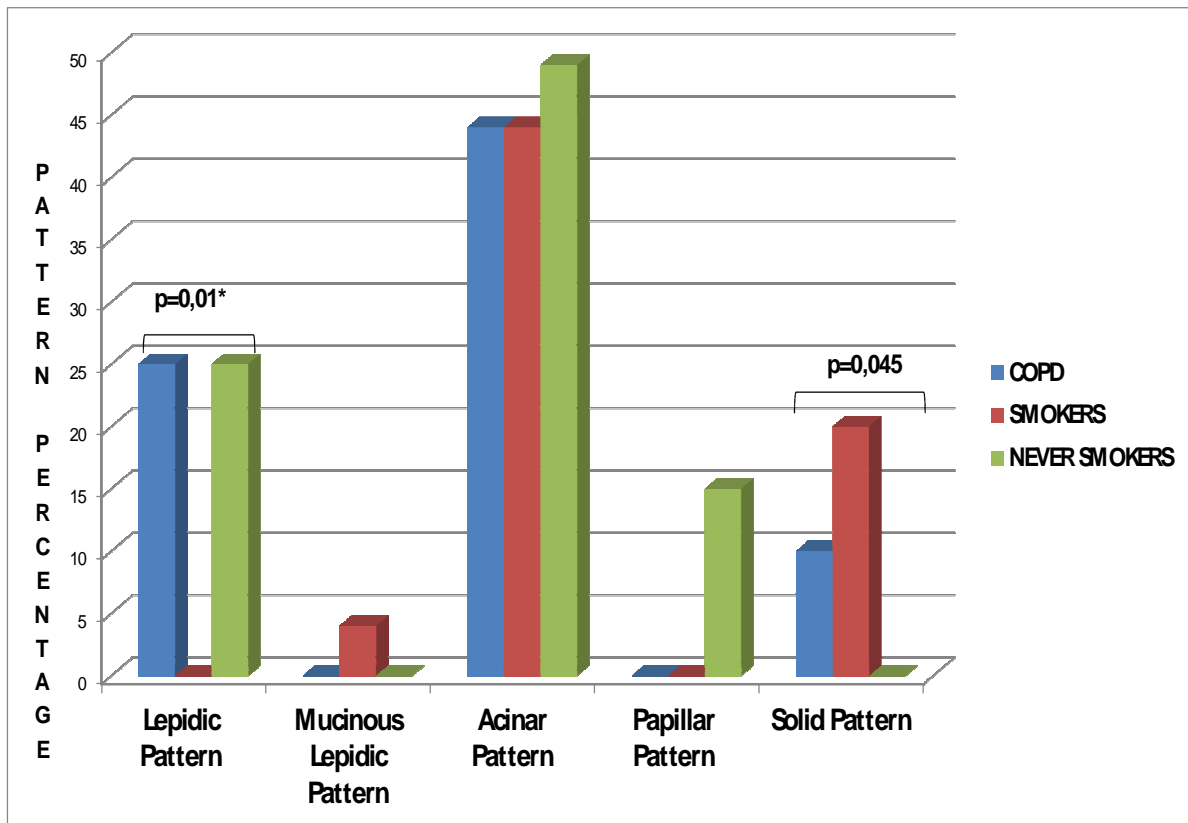


Fig. 4.3. Different lepidic, mucinous lepidic, acinar, papillary and solid pattern distribution in the three study groups.

Quantification of Ki67/MIB1 (Fig. 4.4) showed an higher replication tendency in smoker cases without COPD compared to COPD patients and never smokers ($50 \pm 25\%$ vs $30 \pm 30\%$ vs $24 \pm 18\%$, respectively, $p=0.013$) and was reported in Fig. 4.5.

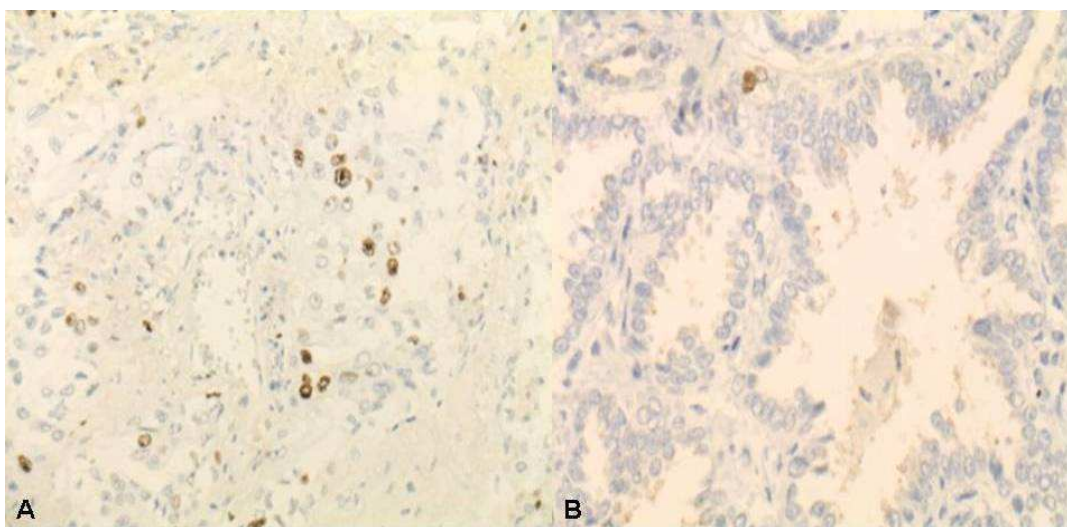


Fig. 4.4. Ki67/MIB1 expression in a smoker (A) and in COPD case (B).

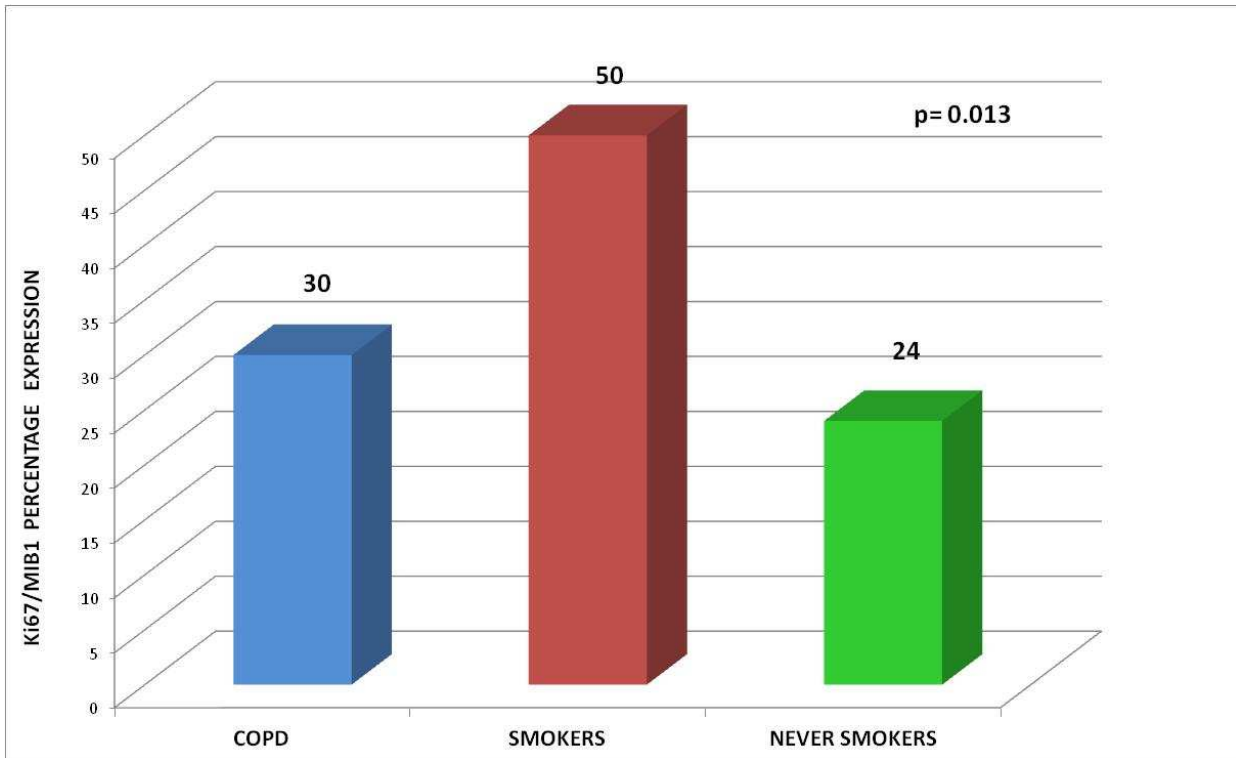


Fig. 4.5. Cell replication assessment by Ki67/MIB1 evaluation showed significant increase in smoker group compared to COPD patients and never smokers.

Finally, quantification of peri and intratumoral remodeling (Fig. 4.6) brought to an increase in the necrotic component in the group of smokers ($20 \pm 23\%$) and COPD ($18 \pm 23\%$), compared to never-smoker cases ($2 \pm 4\%$, $p = 0.03$ and $p = 0.05$, respectively). By contrary, there were no differences concerning the assessment of inflammation ($28 \pm 22\%$, $30 \pm 22\%$ and $33 \pm 20\%$) and fibrotic remodeling ($24 \pm 20\%$, $28 \pm 22\%$ and $28 \pm 18\%$).

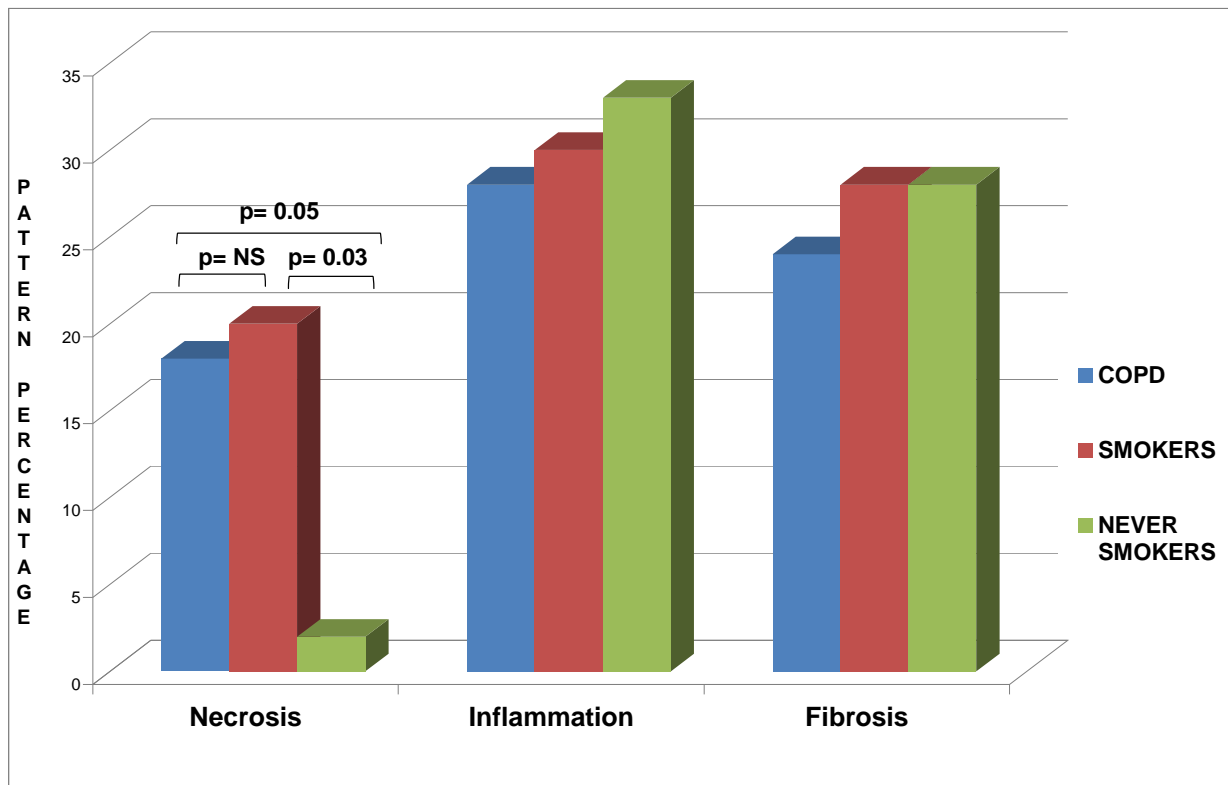


Fig. 4.6. Intra and peri-tumoral remodeling in the three groups.

4.3 GENETIC ANALYSES

The mutation analysis of EGFR and K-RAS genes was performed in all adenocarcinomas. Concerning EGFR, no differences were found between the three groups, although a trend of increased presence of this genetic alteration was observed in non-smokers patients (40%), compared to COPD patients (10%) and smokers (10%).

Analysing KRAS, as expected, we report a trend of increased mutation in smokers compared than never smokers (47% vs 10%, $p=0.09$). At the same time, a trend of difference was present even comparing the group of smokers to that of COPD, who had a smoking similar history (47% vs 30%, $p=NS$). Assuming that this trend could be evidenced, increasing the number of cases, we included in this analysis a retrospective series of 135 patients with a previous KRAS genetic evaluation, which PFTs and smoking history were reviewed in order to classify each case in the right group. We then looked at the new populations, reporting a significance higher percentage

of KRAS mutations in smoker patients without COPD compared to COPD patients and to never smokers (39% vs 15% vs 11%, $p < 0.001$, Fig. 4.4).

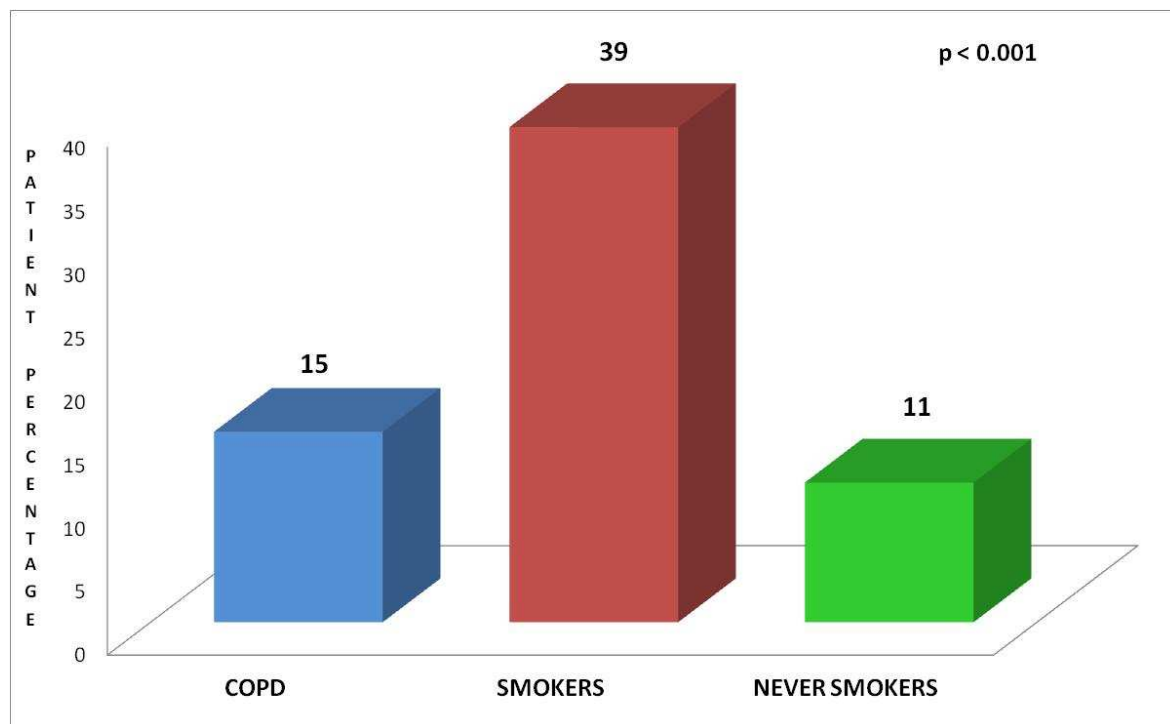


Fig. 4.7. K-RAS mutation prevalence in a combined retrospective and prospective population.

Analyzing the whole population of 46 cases, we found that the presence of KRAS mutations was associated to an higher value of Ki67/MIB1 (60% (IQR 30-70%) vs 25% (IQR 10-45%), $p=0.04$) and higher solid pattern percentage (20% (IQR 20-75%) vs 5% (IQR 0-20%), $p=0.02$) compared to wild-type form. In addition a negative trend was observed between K-RAS mutation and lepidic pattern percentage (0% (IQR 0-30%) vs 15% (IQR 0-30%), $p=0.125$ in KRAS mutated and wild-type form, respectively, Fig. 4.5).

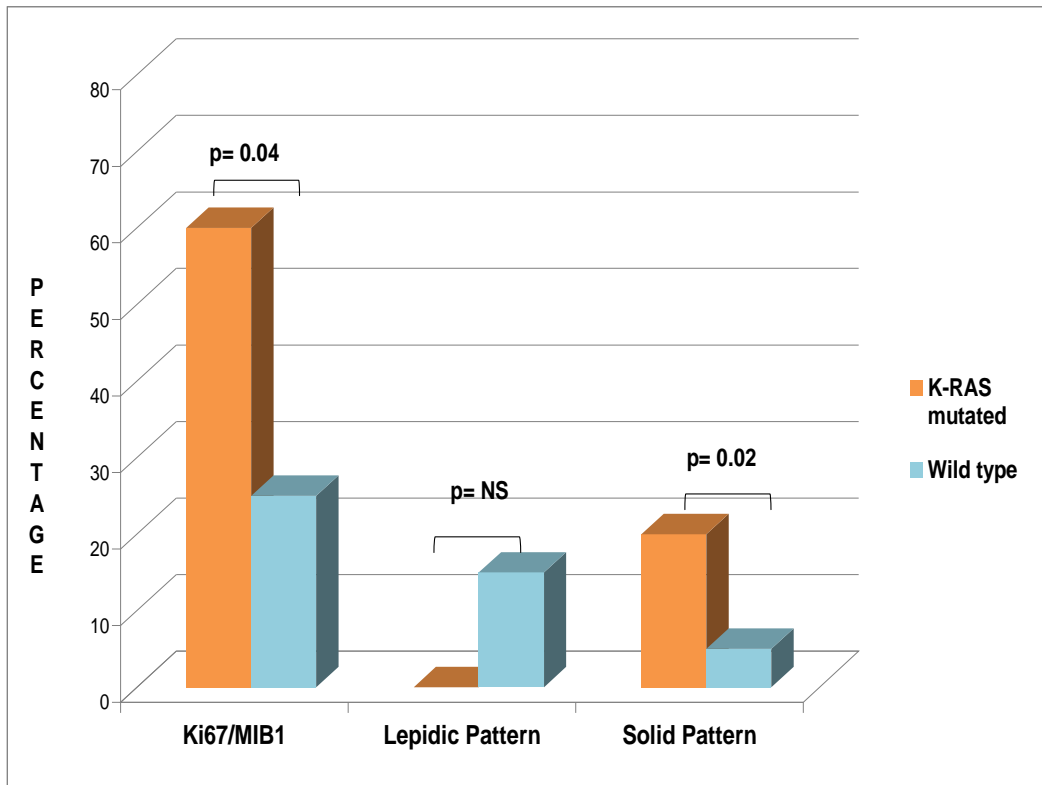


Fig. 4.8. Association of KRAs mutation with cell replication (Ki67/MIB1) and type of growth pattern (lepidic and solid).

4.4 CHARACTERIZATION OF THE PERI-AND INTRA-TUMOR IL-17 CYTOKINE PATTERN

Preliminary analysis of IL-17 expression performed in the groups of COPD patients and smokers without COPD showed a marked cytoplasmic infiltration both in lymphomonocyte and neoplastic cells. Although not significant a stronger IL-17 expression was observed in COPD cases compared to smokers without COPD (253 ± 50 vs 156 ± 109 , $p=NS$, Fig. 4.9 and Fig. 4.10).

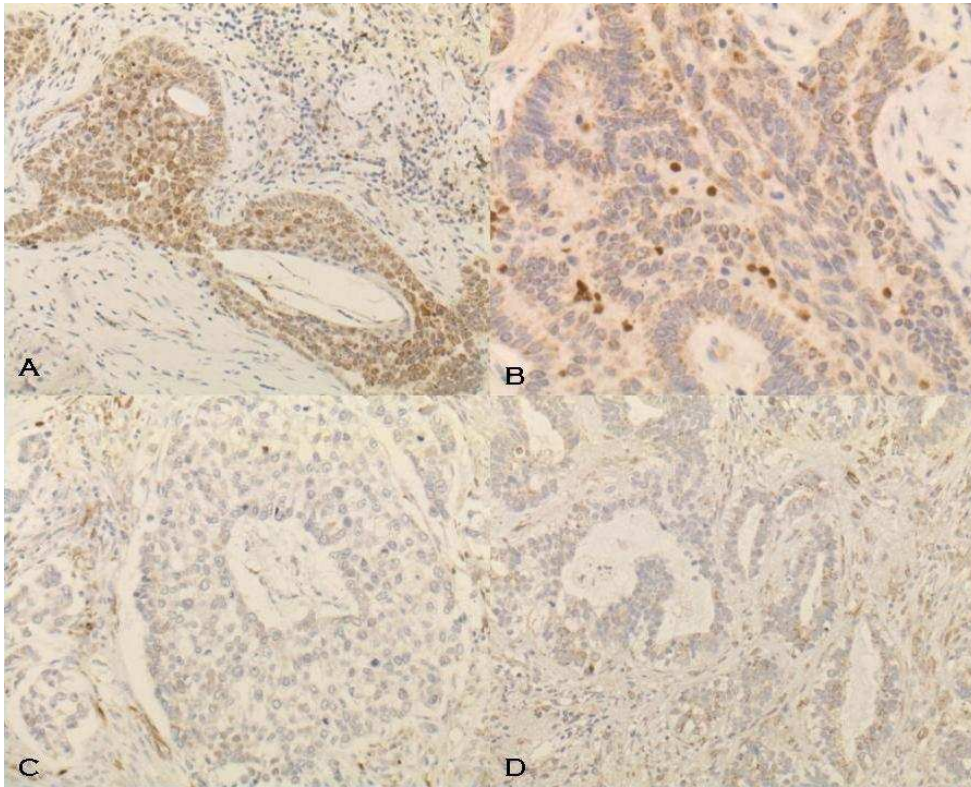


Fig. 4.9. *Histological view of IL-17 expression in COPD (A-B) and smoker without COPD(C-D) related tumours. A –B: both neoplastic cells and lymphocyte are strongly marked. C-D: only a few of neoplastic and lymphocyte show mild-moderate immunostaining.*

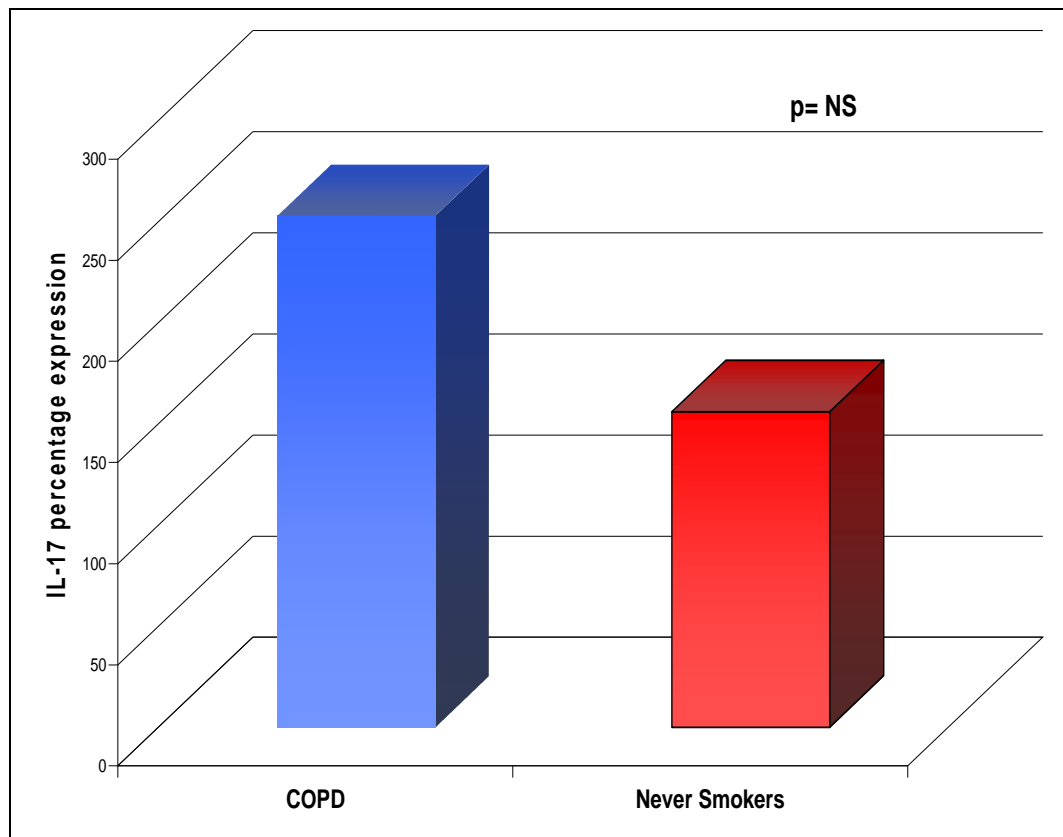


Fig. 4.10. *IL-17 expression assessment in COPD and smokers group.*

5. DISCUSSION

Lung cancer is the most prevalent tumour worldwide and a common cause of cancer-related death [47]. Evidence now demonstrates that pathological subtype classification of “non small-cell lung cancer” (NSCLC), a heterogeneous group of diseases, is central in selecting optimal treatment [110,111]. However, even if surgical and oncologic treatments have greatly improved in recent years overall based on better pathological and molecular diagnosis, lung cancer continues to present increasing frequency, poor long-term results and an unpredictable disease progression.

The co-presence of other diseases in individual patients may strongly influence the carcinogenetic process, making it not only dependent on the classical risk factors. In this context, the relationship that exists between COPD and lung cancer is undoubtedly important. Several epidemiological studies have now firmly established the connection between the two diseases highlighting how COPD is by far the greatest risk factor for lung cancer amongst smokers [68]. Skillrud et al first assessed the risk of lung cancer in patient with COPD compared to a matched case-control study and estimated that the cumulative probability of developing lung cancer within 10 years was 8.8% for those with COPD and 2% for patients with normal function [112]. This indicates that ~1% of patients with COPD develop lung cancer each year, while only 0.2% of patients with normal function develop lung cancer (five-fold increased risk of lung cancer). Several cross-sectional studies, together with recently published prospective studies show smokers with COPD (or reduced FEV1) have up to 4-to 6-fold increased risk of lung cancer when compared with smokers with normal function.

Furthermore 50-70% of patients diagnosed with lung cancer have spirometric evidence of COPD [113].

While the main risk factor for the onset of both diseases is represented by cigarette smoke exposure, they also share a common genetic predisposition which justifies the incidence of these diseases in only a fraction of smokers [74].

This increased risk of lung cancer among smokers with COPD is probably due to activation of common signaling pathways for both diseases by tobacco smoke as well as by smoking-induced chronic inflammation which is a risk factor for both diseases. A causal relation between inflammation and cancer has already been proposed by Galen and later by Virchow [85], who observed inflammatory cell infiltration in malignant tissues and suggested that cancers arise at regions of chronic inflammation. The longer the inflammation persists, the higher the risk of associated carcinogenesis [114].

The importance of this inflammatory response has recently been highlighted, inviting a broader look at the possible involvement of the immune system in the pathogenetic events of COPD. Emerging evidences in this context have emphasized the role of adaptive immune responses, possibly with an autoimmune component due to the recognition of pulmonary self-antigens modified by cigarette smoking and to the failure of mechanisms regulating immunological tolerance [41]. Thus it is not surprising that COPD is associated with increased risk of developing lung cancer due to the synergic effect of cigarette smoke and chronic inflammation. The histologic subtype more frequently associated with smoker COPD is squamous cell carcinoma however no other specific clinical, morphological and/or molecular details have been described in lung cancer associated with COPD.

In our research study we speculated that lung cancer developed in COPD patients has a different clinical/morphological/molecular phenotype compared to lung cancer of smokers without COPD and never smoker subjects. The comparison was mainly performed within the category of smoking patients: COPD versus smokers without COPD, with the main objective to assess the impact of COPD on the tumor phenotype. Never smokers were an important control group in our research study even if it should be noted that tumours arising in thi category of patients have peculiar genetic substrates and mechanisms. The findings of EGFR mutation balanced by the

low KRAS–mutation frequency in never-smoking cases enrolled in the project is proof of these characteristics.

The decision to perform the analysis only in adenocarcinomas was surely influenced by the high percentage of this histologic type determined by the selection bias of inclusion/exclusion criteria and by the need to obtain results not affected by different histotypes. However, this choice reflects the epidemiology trend of lung cancer with adenocarcinoma incidence that has now surpassed that of squamous cell carcinomas in both sexes, possibly due to changes in cigarette smoke composition and/or inhalation [115].

A fundamental point of our prospective study is represented by the fact that the study population included COPD and well matched smoker patients without COPD both for age, sex and smoking history, well known factors which may influence lung carcinogenesis. Among PFTs, excluding those specific for COPD (FEV1%, FEV1/VC, FEV1/FVC, PEF%, TLC%, RV%, FRC%), an approximately equal decrease of FVC percentage in patients with a smoking history (smokers or COPD) was observed. Functional impairment in these patients may be the spirometric expression of parenchymal damage caused by smoke exposure. With regard to biohumoral, and in particular to inflammatory indices, except for basophils, we did not observe significant difference between COPD and smoker patients. The raised number of blood basophils in COPD patients could reflect an anti-tumour immune response to circulating antigen. There are only few papers that focus on the significance of an increased number of basophils and cancer and even fewer for lung cancer. Basophils mediators are reported to attract eosinophils in the site of nematode infestation and arm them for more potent killing of the nematodes [116]. It is possible that the link between the numbers of circulating basophils and eosinophils in lung cancer patients lies in a similar interaction in relation to the tumour, since eosinophil infiltration of the tumours was associated with both blood eosinophilia and better prognosis [117].

Tumour functional assessment was evaluated through ¹⁸F-DG PET-scan which has been imposed as a strategic diagnostic and staging tool for lung cancer. ¹⁸F-DG PET-scan has recently been proposed as a prognostic examination in particular through the SUV index since it seems closely

related to tumour biological aggressiveness [118]. An important meta-analysis performed by Berghmans et al [67] in 2008 took into consideration 13 studies including 1474 patients with different stages of NSCLC. They reported that in 11 studies a high SUV was identified as a poor prognostic factor for survival although two studies found no significant correlation; in addition they estimated an overall combined hazard risk (HR) for the 13 reports of 2.27. Nair et al. recently published a clinical study where they analyzed the biologic basis for the use of FDG as a biomarker, founding that four of eight single genes associated with FDG uptake (LY6E, RNF149, MCM6, and FAP) were also associated with survival [119]. A lower SUV was detected in our COPD lung cancer. This is likely due to a less proliferative index of this neoplasia compared to lung cancer of smokers without COPD. Different works in the literature have reported a significant relation between tumour proliferation detected by MIB1/Ki67 and SUV, highlighting the significant value of FDG as an additional prognostic instrument [120]. The histological analysis of neoplastic patterns associated in our study groups revealed interesting data, emphasising significant differences.

From a morphological point of view adenocarcinoma of COPD patients showed less frequent solid and more frequent lepidic pattern and lower Ki67 proliferation. The more extensive necrosis present both in COPD patients and smokers with lung cancer seems to be the expression of direct smoking cellular damage. All these aspects would characterize this form as a less aggressive neoplastic phenotype. Regarding lepidic pattern, the mucinous variant was found only in the smokers and analysed separately for the other lepidic cases. Indeed the new classification of adenocarcinoma [49] has now divided the mucinous from the nonmucinous form, as multiple studies [121-123] indicated that tumours formerly classified as mucinous BAC have major clinical, radiologic, pathologic, and genetic differences from the tumours formerly classified as nonmucinous BAC. In particular, these tumours show a very strong correlation with KRAS mutation, whereas nonmucinous adenocarcinomas are more likely to show EGFR mutation and only occasionally KRAS mutation [124]. In addition, the neoplasms formerly termed mucinous BAC and now classified as invasive mucinous adenocarcinoma, has been recognized to have

invasive components in the majority of cases [125], leading to a different clinical prognosis compared to nonmucinous variants.

Concerning the molecular data the finding of less frequent K-RAS mutation in lung COPD patients was remarkable. This finding reinforces the concept that tumour related to COPD presented different biological behaviour. The impact of KRAS alterations on prognosis has already been studied in lung cancer and two different meta-analyses suggested an association of KRAS with poorer survival [126-127].

Our group in 2010 [128] also confirmed the impact of KRAS mutations on prognosis in a clinically selected group of advanced adenocarcinomas, highlighting the role of KRAS as an independent prognostic marker.

The analysis of the pathway axis of IL 17 seems to be an important link between COPD and lung cancer providing a possible pathogenetic substrate for both diseases. In our study we observed the increased cytokine expression in lung cancer of COPD cases.

A key role for this cytokine has already been emphasised in the context of autoimmune diseases [129], and more recently in COPD patients, where it seems to support the maintenance of chronic inflammation [99].

New evidences of this axis in the neoplastic field have been reported remarking its crucial role also in lung [130]. The relationship between IL-17 and cancer, whether beneficial or antagonistic, continues to be a controversial issue. The majority of studies reported an important influence of this cytokine on the induction of angiogenesis favoring vascular endothelial growth factor–VEGF-secretion and on neoplastic cell growth. [107] through the alteration of immunosurveillance component [108].

There are still no evidence in the literature that describe the presence and the role of this cytokine in adenocarcinoma associated to COPD. In our study, the small number of subjects certainly influenced the observed results and probably the lack of a significant difference. However, what seems to emerge is that the IL-17 pathway, recently demonstrated to be involved in the complex

immunological background of COPD, could play a key role in cancer development/progression of patients with COPD compared to smokers without COPD.

In conclusion the scenario that emerged from our research study is the presence in COPD patients of a different cancer to that of matched case smokers without COPD. Clinical, biohumoral, morphological and molecular findings of COPD lung cancer showed stigmata of lower aggressiveness and invasiveness lung cancer phenotype.

It might be speculated that alternative mechanisms of carcinogenesis are involved in the development/progression of COPD related lung cancer. Given the importance of inflammation in the pathogenesis of COPD the involvement of other pathways, such as IL-17, mainly driving inflammatory mediated carcinogenesis could be suspected.

The main limitation of our research is represented by the low number of patients especially in the COPD group. Moreover the absence of COPD advanced stages does not allow assessment of the presence of any differences in inflammatory and neoplastic phenotype within the GOLD staging. Finally, the short follow-up of these patients, considering also the early disease stages included in the study, does not allow us to perform any survival analysis.

Future research

The availability of large amount of materials properly stored in a dedicated tissue archive will be able to expand the knowledge of neoplastic phenotypes associated with study populations.

In this field, the characterization of immunophenotype inflammatory cell infiltration both in peritumoral inflammatory infiltrate and in distant areas with particular attention to lymphocytic, macrophagic and cytokine patterns will be of great importance.

The presence of liquid samples as blood and BAL will also lead to the research for possible peripheral biomarkers of disease that can provide predictive and prognostic information related to the neoplastic process. Finally, the analysis of medium-term results of survival and local or distant recurrence could support the preliminary data obtained in this study proposing a different lung cancer phenotype in COPD patients which show sign of less aggressiveness.

New knowledge of mechanisms at the basis of COPD lung cancer would be of considerable help in the fight against lung cancer both for individualized follow-up and for therapeutic perspectives, providing a rationale basis to develop targeted and more effective treatments to prevent or, at least, slow down cancer evolution (anti-inflammatory immunomodulatory or biological therapies).

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7. PRODUCTS OF THE RESEARCH

FULL PAPERS (Dr. Marco Schiavon)

1: Association of TMEM16A chloride channel overexpression with airway goblet cell metaplasia.

Scudieri P, Caci E, Bruno S, Ferrera L, Schiavon M, Sondo E, Tomati V, Gianotti A, Zegarra-Moran O, Pedemonte N, Rea F, Ravazzolo R, Galietta LJ.

J Physiol. 2012;590(Pt 23):6141-55.

2: Demonstration of persistent tumor cells 4 weeks after radiofrequency ablation of a pulmonary adenocarcinoma.

Renaud S, Schiavon M, Santelmo N, Massard G.

J Thorac Cardiovasc Surg. 2012;144(5):e111-3.

3: Increased tissue endothelial progenitor cells in end-stage lung diseases with pulmonary hypertension.

Schiavon M, Fadini GP, Lunardi F, Agostini C, Boscaro E, Calabrese F, Marulli G, Rea F.

J Heart Lung Transplant. 2012;31(9):1025-30.

4: Re-anastomosis of the anomalous segmental pulmonary vein during inferior bilobectomy.

Schiavon M, Antonacci F, Santelmo N, Massard G.

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5: Single-institution experience on robot-assisted thoracoscopic operations for mediastinal diseases.

Rea F, Schiavon M, Di Chiara F, Marulli G.

Innovations (Phila). 2011;6(5):316-22.

6: Does the use of extended criteria donors influence early and long-term results of lung transplantation?

Schiavon M, Falcoz PE, Santelmo N, Massard G.

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7: Synovial sarcoma: CT imaging of a rare primary malignant tumour of the thorax.

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Radiol Med. 2011;116(6):868-75.

8: Endobronchial valve for secondary pneumothorax in a severe emphysema patient.

Schiavon M, Marulli G, Zuin A, Nicotra S, Di Chiara F, De Filippis F, Fantoni U, Rea F.

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9: Multidisciplinary approach for advanced stage thymic tumors: long-term outcome.

Rea F, Marulli G, Di Chiara F, Schiavon M, Perissinotto E, Breda C, Favaretto AG, Calabrese F.

Lung Cancer. 2011;72(1):68-72.

10: Prognostic and predictive implications of EGFR mutations, EGFR copy number and KRAS mutations in advanced stage lung adenocarcinoma.

Bonanno L, Schiavon M, Nardo G, Bertorelle R, Bonaldi L, Galligioni A, Indraccolo S, Pasello G, Rea F, Favaretto A.

Anticancer Res. 2010;30(12):5121-8.

11: Everolimus as a new potential antiproliferative agent in aggressive human bronchial carcinoids.

Zatelli MC, Minoia M, Martini C, Tagliati F, Ambrosio MR, Schiavon M, Buratto M, Calabrese F, Gentilin E, Cavallesco G, Berdondini L, Rea F, degli Uberti EC.

Endocr Relat Cancer. 2010;17(3):719-29.

12: A new technique of diaphragmatic patch fixation in extrapleural pneumonectomy.

Schiavon M, De Filippis A, Marulli G, Rea F.

Eur J Cardiothorac Surg. 2010;38(6):798-800.

13: Pneumonectomy for lung cancer over the age of 75 years: is it worthwhile?

Zuin A, Marulli G, Breda C, Bulf R, Schiavon M, Rebusso A, Di Chiara F, Rea F.

Interact Cardiovasc Thorac Surg. 2010;10(6):931-5.

14: Geometric reconstruction of the right hemi-trunk after resection of giant chondrosarcoma.

Marulli G, Hamad AM, Schiavon M, Azzena B, Mazzoleni F, Rea F.

Ann Thorac Surg. 2010;89(1):306-8.

15: Innovative surgical technique of right upper bilobe transplantation.

Rea F, Hamad AM, Loy M, Breda C, Schiavon M, Marulli G.

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16: Chirurgia polmonare non oncologica

Rea F, Schiavon M, De Filippis AF.

Gasbarrini, Trattato di Medicina Interna, 1139-1150.

ABSTRACTS (Dr. Marco Schiavon)

1) Ruolo prognostico e predittivo dei geni EGFR e K-RAS nel tumore polmonare in stadio avanzato.

Marco Schiavon, Giuseppe Marulli, Andrea Zuin, Adolfo Favaretto, Cristiano Breda, Fabio De Filippis, Samuele Nicotra e Federico Rea

32° Congresso Nazionale SICT Catania 3-5 Giugno 2010

2) Sleeve lobectomy superiore destra: e' veramente una valida alternativa alla pneumonectomia destra nel tumore polmonare non a piccole cellule (NSCLC) localmente avanzato del lobo superiore?

Andrea Zuin, Marco Schiavon, Giuseppe Marulli, Cristiano Breda, Alessandro Rebusso, Francesco Di Chiara, Federico Rea, Francesco Sartori

32° Congresso Nazionale SICT Catania 3-5 Giugno 2010

3) Risultati a lungo termine nel trattamento multimodale del tumore di Pancoast: la nostra esperienza.

Breda Cristiano, Marulli Giuseppe, Zuin Andrea, Nicotra Samuele, De Filippis Fabio, Schiavon Marco, Rea Federico, Sartori Francesco.

32° Congresso Nazionale SICT Catania 3-5 Giugno 2010

4) Nuova tecnica di ancoraggio della protesi diaframmatica nell'intervento chirurgico di pleuropneumonectomia per mesotelioma pleurico maligno.

De Filippis A, Schiavon M, Marulli G, Breda C, Nicotra S, Rebusso A, Zuin A, Rea F

32° Congresso Nazionale SICT Catania 3-5 Giugno 2010

5) Risultati a medio termine della timectomia toracoscopica sx robotassistita nei pazienti affetti da miastenia gravis.

Marulli Giuseppe, Schiavon Marco, Breda Cristiano, Zuin Andrea, Di Chiara Francesco, Rebusso Alessandro, Sartori Francesco, Rea Federico.

32° Congresso Nazionale SICT Catania 3-5 Giugno 2010

6) Bronchovascular sleeve resection: an effective alternative option to pneumonectomy in NSCLC.

De Filippis A, Zuin A, Schiavon M, Marulli G, Rebusso A, Nicotra S, Loy M, Rea F.

23° Congresso Nazionale SPIGC, Forlì 2010.

7) Sleeve lobectomy versus pneumonectomy for NSCLC with N1 disease: analysis of short and long-term results and pattern of recurrence.

Schiavon M, Zuin A, Marulli G, De Filippis A, Battistella L, Rebusso A, Calabrese F, Rea F.

23° Congresso Nazionale SPIGC, Forlì 2010

8) Characterization of lymphoid follicles in patients with end-stage emphysema with or without AAT1 deficiency.

Calabrese F, Baraldo S, Bazzan E, Lunardi F, Ballarin A, Schiavon M, Turato G, Balestro E, Rea F, Cosio MG and Saetta M.

2010 American Thoracic Society Conference, May 14-19, New Orleans, Louisiana USA. Am J Respir Crit Care Med 181;2010:A3837.

9) Safety and long-term efficacy of left thoracoscopic robot-enhanced thymectomy in patients with myasthenia gravis

Marulli G, Schiavon M, Perissinotto E, Zuin A, Di Chiara F, Rebusso A, Battistella L, Rea F Valladolid *ITCVS Suppl. 1 to Vol. 11 (August 15, 2010), p 184*

10) il trapianto polmonare per fibrosi cistica: risultati dell'esperienza di un singolo centro.

Schiavon M, Marulli G, Zuin A, Breda C, Loy M, Nicotra S, Rebusso A, Calabrese F, Calabrese F, Rea F.

XXXIV° Congresso Nazionale SITO, Ancona 7-9 Novembre 2010

11) Laser application in thoracic surgery.

Schiavon M

XXIV° Congresso Nazionale SPIGC, Napoli 2011

12) Modified clinical pathway on flail chest: a preliminary experience.

Antonacci F, Schiavon M, Falcoz PE, Santelmo N, Massard G.

XXIV° Congresso Nazionale SPIGC, Napoli 2011

13) Stapler versus laser techniques for interlobar fissure completion during pulmonary lobectomy: a prospective randomized trial.

Rebusso A, Calabrese F, Battistella L, Schiavon M, Nicotra S, Loy M, Marulli G, Rea F.

XXIV° Congresso Nazionale SPIGC, Napoli 2011.

14) Anatomic segmentectomy for stage I NSCLC: safety and effectiveness.

Calabrese F, Rebusso A, Di Chiara F, Schiavon M, Nicotra S, Loy M, Marulli G, Rea F.

XXIV° Congresso Nazionale SPIGC, Napoli 2011.

15) Utilizzo dell'ECMO nella gestione della PGD post trapianto polmonare.

Di Chiara F, Marulli G, Breda C, Zuin A, Rebusso A, Battistella L, Calabrese F, Andriolo L.G., Bugana A., De Filippis AF, Nicotra S, Schiavon M, Loy M, Rea F.

XXXV° Congresso Nazionale SITO, Milano 2011

16) Donneur limite : comment la médecine factuelle peut nous aider!

Falcoz PE, Schiavon M.

IV° Journées de Transplantation Pulmonaire à Strasbourg - 14 - 15 octobre 2011

17) Experimental evaluation of a new system for laser tissue welding applied on damaged lung.

Schiavon M, Marulli G, Zuin A, Calabrese F, Breda C, Loy M, Nicotra S, Rea F.

XXVI° European Congress of EACTS, Barcellona 2012.

18) Is lobectomy really more effective than sublobar resection in surgical treatment of second primary lung cancer?

Zuin A, Andriolo L, Marulli G, Schiavon M, Nicotra S, Calabrese F, Romanello P, Rea F.

XXVI° European Congress of EACTS, Barcellona 2012.

19) Surgical And Neurological Outcomes After Left Thoracoscopic Robot-enhanced Thymectomy In 100 Consecutive Myasthenia Gravis Patients.

Marulli G, Schiavon M, Di Chiara F, Bugana F, Rebusso A, Perissinotto E, Calabrese F, Rea F. *Western Thoracic Surgical Association, Annual Meeting, Hawaii 2012.*

20) Studio sperimentale di un nuovo sistema laser-mediato (laser tissue welding) per l'aerostasi polmonare.

Schiavon M, Marulli G, Zuin A, Calabrese F, Breda C, Loy M, Nicotra S, Rea F. *Congresso Nazionale SICT, Pisa 2012.*

21) Allograft sternochondral replacement: a novel paradigm for the reconstruction of anterior chest wall?

Marulli G, Di Chiara F, Breda C, Loy M, Zuin A, Calabrese F, Schiavon M, Nicotra S, Rea F. *XX° European Conference on General Thoracic Surgery, ESTS, Essen 2012.*

CONGRESSES (Dr. Marco Schiavon)

1. International Conference on End-stage Lung Diseases: New Insights and Lung Transplantation, Padova 2010.

2. XXXII° Congresso Nazionale SICT (Società Italiana di Chirurgia Toracica), Catania 2010.

3. XXIII° Congresso Nazionale SPIGC (Società Polispecialistica Italiana Giovani Chirurghi), Forlì 2010.

4. XXXIV° Congresso Nazionale SITO (Società Italiana Trapianti d'Organo), Ancona 2010.

5. Corso di Aggiornamento: NSCLC. Target therapy: come cambia la gestione del paziente. Mogliano Veneto 2010.

6. XX° European Conference on General Thoracic Surgery, Marseille 2011.

7. XXIV° Congresso Nazionale SPIGC (Società Polispecialistica Italiana Giovani Chirurghi), Napoli 2011.

8. **IV° Journées de Transplantation Pulmonaire à Strasbourg - 14 - 15 ottobre 2011.**
9. **XX° European Conference on General Thoracic Surgery, ESTS, Essen 2012.**
10. **XXXII° Congresso Nazionale SICT (Società Italiana di Chirurgia Toracica), Pisa 2012.**
11. **XXVI° European Congress of EACTS, Barcellona 2012.**
12. **Biosurgery Symposium Hands-on Workshop, Vienna 2012.**
13. **Congresso Società Italiana per la Sicurezza e Qualità nei trapianti, Milano 2012.**
14. **I° congresso Nazionale Unità e Valore della Chirurgia Riunita, Roma 2012.**

AWARDS (Dr. Marco Schiavon)

1. **2010:** Best Communication in Thoracic Surgery at XXIII° SPIGC (Società Polispecialistica Italiana Giovani Chirurghi) National Congress. Forlì, 23 June 2010.
2. **2011:** Prize Cav. Albano e Cav. Antero Maran from Istituto Veneto di Scienze, Lettere ed Arti destinato to favour research on lung cancer surgical treatment for the project “ANALISI PROGNOSTICA E PREDITTIVA DEI GENI EGFR E K-RAS NELL’ADENOCARCINOMA POLMONARE IN STADIO PRECOCE”. Venezia, 29 May 2011.
3. **2012:** Prize “Mario Selli” at SICT National Congress for the project “STUDIO SPERIMENTALE DI UN NUOVO SISTEMA LASER-MEDIATO (LASER TISSUE WELDING) PER L’AEROSTASI POLMONARE”. Pisa, 17 June 2012.

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