

The Relationship Between Causative Microorganisms and Airway Obstruction in Patients with COPD Exacerbation

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Abstract

Objective: Chronic obstructive pulmonary disease (COPD) is a common, preventable and treatable disease. Exacerbations and comorbidities contribute to the severity of the disease. In our study, we aimed to evaluate the relationship between causative microorganisms and airway obstruction in patients hospitalized for COPD exacerbation.

Methods: Sputum and blood cultures of 75 patients, who were admitted to Nevşehir State Hospital Chest Diseases Unit between January 2015 and January 2016 and who were diagnosed as COPD exacerbation, were analyzed retrospectively. The relationship between microorganisms in sputum culture and airway obstruction was evaluated. Forced expiratory volume in one second (FEV₁) value was used to evaluate airway obstruction.

Results: Of the 75 patients evaluated in our study, 45 (60%) were male and 30 (40%) were female. The mean age was 61.2 years. Isolated microorganisms were as follows: *H. influenzae* in 12 (16%) patients, *S. pneumoniae* in nine (%12) patients, *M. catarrhalis* in six (8%) patients, *P. aeruginosa* in four (5.3%) patients, *E. coli* in three (4%) patients, *K. pneumoniae* in three (4%) patients, and methicillin-sensitive *S. aureus* in one (1.3%) patient. No statistically significant difference was found between the microorganisms in terms of airway obstruction (p>0.05).

Conclusion: We did not find a correlation between the causative microorganism and airway obstruction in patients hospitalized for COPD exacerbation. **Keywords:** COPD exacerbation, sputum culture, airway obstruction, FEV₁

INTRODUCTION

Chronic obstructive pulmonary disease (COPD) is a preventable and treatable disease characterized by the chronic inflammatory response of the airways and lungs against harmful gases and particles, leading to a progressive limitation of airflow. Exacerbations and comorbidities that occur during the course of the disease cause an increase in the severity of the disease (1).

Exacerbation of COPD has been described as a condition characterized by acute deterioration of the patient's respiratory tract symptoms leading to drug change (1). There is a loss of pulmonary functions due to exacerbations, which leads to impaired quality of life and an increase in morbidity and mortality. The etiology of COPD exacerbations consists of tracheobronchial infections (50-70%) and air pollution (10%), and the etiology cannot be detected in about 30% of exacerbations. In patients with COPD, exacerbations should be prevented and treated in order to prevent disease progression and to maintain quality of life.

In our study, we aimed to evaluate tracheobronchial infectious agents in sputum culture, and the relationship between cautious microorganisms and airway obstruction in patients hospitalized for COPD exacerbation.

METHODS

Seventy-five patients, who were admitted to Nevşehir State Hospital Chest Diseases Unit with COPD exacerbation between January 2015 and January 2016, were evaluated retrospectively.



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Patients who were not treated with antibiotics one month before admission were evaluated. Patients with cystic fibrosis, asthma and bronchiectasis were excluded from the study.

Sputum and blood cultures that were obtained for the isolation of microbiological agent causing acute exacerbation of COPD were examined. Sputum samples with polymorphonuclear leukocyte count >25 and mouth epithelial cells <10 in all areas were evaluated. Pulmonary function tests performed in the stable period of three months before the acute exacerbation were evaluated. Forced expiratory volume in 1 second (FEV₁) value was used to evaluate airway obstruction. Patients with positive sputum culture were divided into two groups according to FEV₁ (FEV₁ >50% and FEV₁ <50%).

Statistical Analysis

Statistical analyzes were performed using IBM SPSS Statistics 22 software. Mean and standard deviation was used for numerical data when evaluating the study data. Frequency distributions were calculated for comorbidities and isolated microorganisms from sputum cultures. The chi-square test was used to evaluate the relationship between FEV_1 value and the causative microorganism.

RESULTS

Of the 105 patients evaluated in our study, 30 patients were not included in the study because of antibiotic use up to one month before hospitalization and having comorbidities such as asthma and bronchiectasis. Of the 75 patients evaluated in our study, 45 (60%) were male and 30 (40%) were female. In our study, the mean age of patients was 61.2 ± 16.3 (range, 40-85).

The complaints of the patients were as follows: increased dyspnea in 65 (86%) patients, cough in 60 (80%) patients, increased sputum in 55 (73%) patients, and change in sputum color in 36 (48%) patients. While 50 (66%) patients continued smoking, 25 (33%) had quitted smoking.

The mean FEV_1 was 1.35 ± 0.75 L (47.7% ±20.68), FEV_1 /forced vital capacity (FVC) was $50.4\%\pm15.2$, and FVC was 1.79 ± 0.79 L (56.81 ±17.45).

Twenty-eight (37%) of the 75 patients had hypertension, 24 (32%) had diabetes mellitus, 18 (24%) had congestive heart failure, 12 (16%) had atherosclerotic heart disease, four (5%) had lung cancer, four (5%) had obstructive sleep apnea and three (4%) had depression (Table 1).

Isolated microorganisms were as follows: *H. influenzae* in 12 (16%) patients, *S. pneumoniae* in nine (%12) patients, *M. catarrhalis* in

six (8%) patients, *P. aeruginosae* in four (5.3%) patients, *E. coli* in three (4%) patients, *K. pneumoniae* in three (4%) patients, and methicillin-sensitive *S. aureus* in one (1.3%) patient. Normal throat flora elements were observed in 37 (49.3%) patients (Table 2).

Positivity was detected in only four of the blood cultures obtained during hospitalization of the patients. Isolated microorganisms were *S. pneumoniae* in one (1.3%) patient, *E. coli* in one (1.3%) patient, *H. influenzae* in one (1.3%) patient and *K. pneumonia* in one (1.3%) patient.

 FEV_1 was >50% in 18 (47%) patients and <50% in 20 (53%) patients with positive sputum culture (Table 3).

As they are the most common isolated microorganisms in acute bacterial COPD exacerbations, *S. pneumoniae*, *H. influenzae* and *M. catarrhalis* growth in patients with $FEV_1 > 50\%$ were compared with the group with $FEV_1 < 50\%$ (group with severe airway obstruction). There was no statistically significant difference between the two groups in terms of sample distribution (p>0.05).

Patients with $FEV_1 > 50\%$ and <50% were compared in terms of gram (+) and gram (-) bacteria growth. There was no statistically significant difference between the two groups in terms of gram (+) and gram (-) bacteria growth (p>0.05).

Table 1. Comorbidities of the study cohort						
Comorbidity	N	umber	%			
Hypertension	28	}	37			
Diabetes mellitus	24	-	32			
Congestive heart failure	18	}	24			
Atherosclerotic heart disease	12		16			
Lung cancer	4		5			
Obstructive sleep apnea	4		5			
Depression	3		4			
Table 2. Distribution of bacteria isolated in sputum culture						
Bacteria		Number	%			
H. influenzae		12	16			
S. pneumoniae	9		12			
M. catarrhalis		6	8			
P. aeruginosae		4	5.3			
E. coli		3	4			
K. pneumoniae		3	4			
S. aureus (methicillin-sensitive)		1	1.3			
Normal throat flora elements		37	49.3			
			1			

Total

100

75

DISCUSSION

Exacerbation of COPD is characterized by acute deterioration of the patient's respiratory tract symptoms leading to drug change (1). Exacerbations have negative effects on morbidity and mortality. As the disease progresses, the frequency and severity of exacerbations increase and the quality of life of COPD patients is further reduced.

The etiology of COPD exacerbations consists of tracheobronchial infections (bacterial agents 40-50%, viral agents 30-40%, atypical bacterial agents 5-10%) and air pollution (10%), and the etiology cannot be detected in about 30% of exacerbations (2). Respiratory pathogens can be detected in the respiratory tract in the stable period of the disease, so the role of infectious agents in exacerbations is controversial. In bronchoscopic studies, it was observed that approximately half of the patients had higher concentrations of bacteria in the lower respiratory tract during exacerbation than the stable period of the disease (3). In our study, we detected 50.7% bacterial infection in the etiology of exacerbations.

In a study to determine the indication for antibiotics in COPD exacerbations, the basic criterion for antibiotic indication was found to be sputum purulence. In this study, 86 patients with positive sputum culture were evaluated and 73 of them had purulent sputum (sensitivity 84-94%, specificity 77-84%) (4). In our study, sputum purulence was increased in 36 (48%) patients. According to the Committee on Publication Ethics study, it is recommended that antibiotics should be given in case of visualization of bacteria with gram staining, decrease in respiratory function, and two or more exacerbation history within last year, and antibiotic treatment is not recommended in the absence of any of these criteria (absence of all these criteria has 100% negative predictive value) (5).

Table 3. Causative microorganisms and their distributions according to \ensuremath{FEV}_1						
Causative microorganism	FEV ₁ >50%	FEV ₁ <50%	Total	р		
H. influenzae	5 (42%)	7 (58%)	12	0.148		
S. pneumoniae	5 (55.5%)	4 (44.5%)	9	0.164		
M. catarrhalis	3 (50%)	3 (50%)	6	0.213		
E. coli	1 (33.3%)	2 (66.6%)	3	0.394		
K. pneumoniae	1 (33.3%)	2 (66.6%)	3	0.394		
P. aeruginosae	2 (50%)	2 (50%)	4	0.369		
<i>S. aureus</i> (methicillin-sensitive)	1 (100%)	-	1	0.512		
Total	18 (47%)	20 (53%)	38	0.102		
p>0.05, FEV ₁ : Forced expiratory	volume in 1	•				

The most frequently causative bacterial agents responsible for exacerbations are H. influenzae, S. pneumoniae and M. catarrhalis. Enteric gram (-) bacteria and P. aeruginosa are increasingly isolated in patients with severe airway obstruction, hypoxemia, malnutrition, comorbidity, frequent hospitalization and antibiotic use (6, 7, 8). In our study, similar to literature, the most frequently isolated infectious agents were H. influenzae in 12 (16%) patients, S. pneumoniae in nine (12%) patients and *M. catarrhalis* in six (8%) patients. In recent years, studies that are supported by serological diagnostic methods reported that Chlamydia pneumoniae and Mycoplasma pneumoniae might be responsible for a significant proportion of COPD exacerbations (4-34%), and atypical bacteria were not detected in studies using polymerase chain reaction (9).

Viruses are responsible for 15-40% of COPD exacerbations caused by infectious causes and a significant proportion of these infections are found together with bacterial infections (10, 11). In a study by Seemungal et al., (12) they stated that 64% of patients with COPD exacerbations had common cold 18 days before exacerbation and that rhinoviruses were the most common respiratory tract virus with a rate of 58%. In our study, no growth was detected in sputum culture of 37 (49.3%) patients and we think that atypical bacteria, viruses or non-infectious causes may cause exacerbation in these patients.

Eller et al. (13) reported that the microorganisms isolated in the exacerbations of infective COPD might change with the degree of impairment of pulmonary function. They reported that most of the microorganisms isolated in patients with FEV, values >50% were S. pneumoniae, H. influenzae and M. catarrhalis, and that gram-negative microorganisms such as Enterobacteriaceae and Pseudomonas spp. were more common in cases with severe airway obstruction with a $FEV_1 < 35\%$. Miravitles et al., (14) found higher rates of *Pseudomonas* spp. and H. influenza in patients with COPD exacerbation and FEV, <50%. In our study, we found no significant difference in the distribution of causative microorganisms in cases with severe airway obstruction (FEV₁ <50%) and FEV₁ >50%. We believe that the retrospective nature of our study and the low number of causative microorganisms in the sputum culture may be effective on this result.

CONCLUSION

We aimed to evaluate the relationship between tracheobronchial infectious agents and airway obstruction in patients with COPD exacerbation. We did not find a relationship between the distribution of cautious microorganisms and airway obstruction in COPD exacerbations. We believe that this is due to the low number of patients with isolated causative agents.

Ethics

Ethics Committee Approval: Retrospective study.

Informed Consent: Retrospective study.

Peer-review: Externally peer-reviewed.

Conflict of Interest: No conflict of interest was declared by the authors.

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