



**TLR4 up-regulation and reduced Foxp3 expression in mechanically ventilated smokers with obstructive chronic bronchitis.**

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Manuscripts

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3 1 **TITLE: TLR4 up-regulation and reduced Foxp3 expression in mechanically ventilated**  
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5 2 **smokers with obstructive chronic bronchitis.**  
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49 22 **RUNNING HEAD: Innate immunity and immune regulation in acute respiratory failure.**  
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51 23  
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53 24 **KEY WORDS:** toll like receptors, Foxp3, chemokines, smokers, respiratory failure.  
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3 26**LIST OF ABBREVIATIONS**4  
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6 27 COPD=Chronic obstructive pulmonary disease; C= Controls; CB= chronic bronchitis; S= smokers;7  
8 28 Mini-BAL= mini-bronchoalveolar lavage; TLR= Toll like receptor; Foxp3=forkhead box P3; IP-9  
10 29 10= interferon gamma induced protein 10; IL-8= interleukin 8.11  
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**CONFLICTS OF INTEREST**

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Elisabetta Pace- Competing interests: None declared.

Maria Ferraro - Competing interests: None declared.

Antonino Giarratano- Competing interests: None declared.

Chiara Cipollina-Competing interests: None declared.

Mark Gjomarkaj- Competing interests: None declared.

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## SUMMARY

**Background.** Chronic bronchitis (CB) is a risk factor in chronic obstructive pulmonary disease (COPD) for accelerated lung function decline and increased mortality. The lung and systemic inflammatory and immunological profile of COPD patients with CB which acutely experience respiratory failure upon a disease exacerbation is unknown.

**Methods.** In this study, we explored the expression of Foxp3 by western blot analysis, TLR4 by immunocytochemistry and the concentrations of IP-10 and IL-8 by ELISA in the mini-bronchoalveolar lavages (mini-BAL) and in the peripheral blood of patients with respiratory failure requiring intubation and mechanical ventilation. The recruited subjects were separated into three different groups: smokers with CB and COPD (COPD, n=18), smokers with CB but without COPD (S, n=8) and patients without CB and without COPD (C, n=10).

**Results.** In mini-BAL of COPD group, Foxp3 and IP-10 were significantly reduced while TLR4 was significantly increased in comparison to C. TLR4 was also increased in mini-BAL of S. In COPD peripheral blood, Foxp3 was reduced in comparison to C but no significant differences were observed for TLR4 and for IP-10. No significant differences were observed for IL-8 concentrations in the mini-BAL and in the blood of the recruited patients. The mini-BAL TLR4 expression correlated with the Clinical Infective Pulmonary Score.

**Conclusions.** In exacerbated COPD patients with respiratory failure, lung and systemic reduced immune regulatory events (low Foxp3 expression) and lung increased innate immunity responses (high TLR4 expression) occur. These events may contribute to the increased inflammatory events leading to respiratory failure.

## INTRODUCTION

Chronic obstructive pulmonary disease (COPD) is a heterogeneous disease, is associated with pulmonary and extra-pulmonary clinical manifestations and includes different clinically relevant subtypes (1). One of the COPD subtypes is characterized by chronic bronchitis (CB). CB is a risk factor for accelerated lung function decline in COPD, increased hospitalization, and increased mortality (2).

Cigarette smoke represents the most important risk factor for CB (3). Chronic oxidative stress of cigarette smoking induces mucus secretion and the increased mucus viscosity renders the airways susceptible to bacterial infections, a hallmark of CB (3).

A key component of the innate immunity and of the innate defence mechanisms against infections is represented by the toll like receptor (TLR) family (4) .

A recent hypothesis regarding COPD pathogenesis suggests as “step 1 of the disease” the activation of innate responses by injured tissue components (5). Products derived from epithelial cell injury can act as ligands for TLR4 and TLR2, thus amplifying inflammatory responses within the airways.

Cigarette smoke is able to increase the expression of TLR4 and to orientate the activation of TLR4 toward an increased release of IL-8 and a reduced release of interferon gamma-induced protein 10 (IP-10) in bronchial epithelial cells (6).

In the airways of COPD patients mechanically ventilated due to acute respiratory failure, there is an increased expression of TLR4 and an increased chemotactic activity toward neutrophils but a reduced concentration of IP-10 with a reduced chemotactic activity toward lymphocytes (7). The increased airway inflammation in the airways of COPD patients may also be sustained by the impairment of immune regulatory events and in particular may be linked to the reduced expression of the forkhead box P3 (Foxp3), a transcription factor crucially involved in T regulatory activities. COPD patients have, in small airways, decreased numbers of Foxp3 positive cells that negatively correlate with airflow obstruction (8) (9). Although COPD is associated with lung and systemic

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3 87 inflammation, it is unknown whether the alteration in the innate and immune regulatory events  
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5 88 observed in the airways of COPD patients are also present in the systemic compartment.  
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9 89 The objectives of this study were to investigate the immune regulatory events and the inflammatory  
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11 90 and the host defence responses in the airways and in the systemic compartment of COPD patients  
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13 91 with CB. Exacerbated COPD patients who experienced an acute respiratory failure requiring  
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15 92 endotracheal intubation and mechanical ventilation were compared to patients requiring also  
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17 93 intubation and mechanical ventilation but without COPD and without CB. In the recruited patients  
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19 94 the expression of TLR4, the expression of Foxp3 and the release of specific chemokines (IP-10 and  
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21 95 IL-8) were investigated.  
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## MATERIALS AND METHODS

### 98 Patient population

99 This study was conducted at the ICU of the Department of Anesthesiology, Reanimation and  
100 Emergency of the University of Palermo, Italy. Local ethic committee permission and informed  
101 written consent from either the patient or closest relatives were obtained. The patients were  
102 classified into the following groups: 1) subjects without smoking history, without history of  
103 previous CB or chronic pulmonary diseases including asthma and COPD and with acute respiratory  
104 failure upon surgery for abdominal or thoracic aneurysm, (Controls =C; n=10); 2) smoking subjects  
105 (>15 pack-year) with CB, without COPD and with acute respiratory failure upon surgery for  
106 abdominal or thoracic aneurysm (S; n=8); 3) smoking subjects (>15 pack-year) with CB with  
107 COPD (GOLD-1-2) and with acute respiratory failure upon treatment failure of an acute  
108 exacerbation (COPD; n=18). The COPD patients fulfilled the diagnostic criteria of COPD (10) with  
109 a post-bronchodilator obstruction ( $FEV_1 < 80\%$  predicted, and  $FEV_1/FVC$  ratio  $< 70\%$ ). Patients  
110 with x-ray or clinical evidence of sepsis or pneumonia at the time of mini-bronchoalveolar lavage  
111 (mini-BAL) collection were not included. CB were defined on the basis of symptoms and in  
112 particular productive cough lasting more than three months in more than two years. All recruited  
113 subjects required mechanical ventilation and underwent therapy with antibiotics and systemic  
114 corticosteroids (no significantly different doses among the patients included in the three groups).  
115 The antibiotics were adjusted to cover any identified pathogens on the basis of antibiograms. COPD  
116 patients, before COPD exacerbation, were undergoing therapy with bronchodilators but not with  
117 corticosteroids. Exacerbations were defined as previously reported (10) and were treated with  
118 antibiotics and systemic corticosteroids. At the ICU admission, data for Clinical Pulmonary  
119 Infective Score (CPIS), the simplified Acute Physiology Score (SAPS II) and sepsis-related organ  
120 failure assessment (SOFA) were collected from each recruited patient. Paired mini-BAL and blood  
121 samples were collected from all participants. Microbiology of mini-BAL was also assessed.



**122 mini-BAL collection and processing**

123 Distal lung fluid samples (mini-BAL) were obtained using BAL Cath system (by Kimberly Clark)  
124 within 1 h from the intubation. The protected catheter was blindly advanced through the  
125 endotracheal tube until it was wedged into a distal airway and aliquots of 10 ml of sterile 0.9%  
126 NaCl were instilled and gently suctioned (recovered volume about 70% of the instilled volume).  
127 Mini-BAL samples were filtered through a sterile gauze and then centrifuged at 1300 rpm for 10  
128 min to separate cells from supernatants. Total and differential (diff-quick staining) cell counts were  
129 assessed. The cell fraction was used for immunocytochemistry and western blot experiments. The  
130 supernatants were assessed for cytokine levels.

**131 Blood samples**

132 Blood samples (10 ml) were collected from C, S, COPD subjects and then processed for obtaining  
133 plasma and peripheral blood mononuclear cells (PBMC). PBMC were isolated from blood s by  
134 Ficoll-Hypaque (Pharmacia) gradient centrifugation. The cells were suspended in RPMI 1640 tissue  
135 culture medium (Invitrogen Life Technologies) supplemented with 1% heat-inactivated FCS  
136 (Invitrogen Life Technologies), 2 mM L-glutamine, 20 mM HEPES, 100 U/ml penicillin, 100 µg/ml  
137 streptomycin,  $5 \times 10^{-5}$  M 2-ME and 85 µg/ml gentamicin. Purity and viability were tested using  
138 trypan blue exclusion.

**139 Immunocytochemistry**

140 The expression of TLR4 was evaluated using a rabbit polyclonal antibody (Santa Cruz  
141 Biotechnology). Immunocytochemistry was performed using AP-LSAB2 (DAKO, Glostrup,  
142 Denmark) kit following the manufacturer's instructions and new Fuchsin as chromogenic substrate  
143 (DAKO) (cytoplasmic red staining). Negative controls were performed using rabbit or mouse  
144 negative control immunoglobulins (DAKO). Data are expressed as percentage of positive cells.

**145 Western blot analysis**

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3 146 The expression of Foxp3 was evaluated by western blot analysis as previously described (11) with  
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5 147 minor modifications. 40 µg of total protein were loaded in the gel. All blots were probed using a  
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7 148 goat polyclonal antibody anti-Foxp3 (1:100) (Santa Cruz Biotechnology, Santa Cruz, CA).  
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9 149 Membranes were then stripped and incubated with goat polyclonal anti-β-actin (Sigma) as  
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11 150 housekeeping protein to normalize differences in protein loading. Revelation was performed with an  
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13 151 enhanced chemiluminescence system (GE Healthcare, Chalfont St. Giles, UK) followed by  
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15 152 autoradiography. Negative controls were performed in the absence of primary antibody or including  
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17 153 an isotype control antibody. Data are expressed as densitometric arbitrary units by correction with  
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19 154 the density of the bands obtained for beta-actin.  
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#### 23 155 **Measurement of IL-8 and IP-10**

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26 156 The concentrations of IL-8 and IP-10 in plasma from C, S, CB-COPD subjects were determined by  
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28 157 an enzyme-linked immunosorbent assays (ELISA) following the manufacturer's instructions  
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30 158 (Quantikine; R&D Systems, Minneapolis, MN).  
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#### 32 159 **Statistics**

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35 160 Data are expressed as median (25-75 percentiles). Kruskal Wallis test was performed for  
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37 161 comparisons among patient groups. A non-parametric Mann Whitney test was then applied as the  
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39 162 initial Kruskal Wallis test was significant. The Wilcoxon test was used for comparisons between  
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41 163 mini-BAL and autologous peripheral blood in each recruited patient. The Spearman test was used  
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43 164 for correlations. P< 0.05 was accepted as statistically significant.  
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## RESULTS

### 167 **Demographic characteristics of the subjects.**

168 The demographic characteristics of the three study groups are shown in Table 1. SAPS II and SOFA  
169 scores revealed no significant differences in the recruited patients. The CPIS score was significantly  
170 higher in COPD than in C (Table 1). No significant differences for CPIS score were shown in S in  
171 comparison to C and to COPD. The total and the differential cell counts of mini-BAL are shown in  
172 Table 2. Significantly higher numbers of total cells and of neutrophils were present in COPD  
173 patients. **Microbiology of mini-BAL was shown in table 3.**

### 175 **Expression of TLR4 in cells from mini BAL and from peripheral blood cells.**

176 The percentage of TLR4 positive cells was significantly higher in mini-BAL cells from COPD and  
177 from S in comparison to mini-BAL from C. The percentage of TLR4 positive cells was significantly  
178 higher in mini-BAL cells from COPD in comparison to mini-BAL from S and to autologous  
179 peripheral blood (figure 1) **(table 4)**. No significant differences in TLR4 expression were observed  
180 in peripheral blood cells among C, S and COPD (figure 1) **(table 4)**.

### 181 **Concentrations of IL-8 and of IP-10 in mini BAL and in peripheral blood cells.**

182 The concentrations of IL-8 were significantly higher in mini-BAL of all the recruited patients (C, S  
183 and COPD patients) in comparison to autologous peripheral blood (figure 2). No significantly  
184 different concentrations of IL-8 were observed in mini-BAL and in peripheral blood among C, S  
185 and COPD (figure 2). The IP-10 concentrations were significantly reduced in mini-BAL from  
186 COPD in comparison to C and S and in comparison to autologous peripheral blood (figure 3). The  
187 IP-10 concentrations were significantly increased in peripheral blood from COPD in comparison to  
188 mini-BAL from C. No significantly different concentrations of IP-10 were observed in peripheral  
189 blood between S and C (figure 3).

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3 190 **Expression of Foxp3 in mini BAL and in peripheral blood cells.**

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5 191 In COPD and in S the expression of Foxp3 in mini-BAL and in peripheral blood was significantly  
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7 192 lower than in C (figure 4 A-B) (table 4). In COPD the expression of Foxp3 in mini-BAL and in  
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9 193 peripheral blood was significantly lower than in S (figure 4 A-B) (table 4). No significant  
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11 194 differences were observed in the expression of Foxp3 in mini-BAL and in autologous peripheral  
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13 195 blood in all the recruited patients.

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16 196 **Correlations**

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18 197 Finally, we tested whether the observed alterations in TLR4, Foxp3 and IP-10 in both lung and  
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20 198 systemic compartments correlate with clinical scores of severity. CPIS correlates with mini-BAL  
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22 199 TLR4 expression (figure 5) but not with the other markers (Foxp3 and IP-10) (data not shown). No  
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24 200 significant correlations were observed between SOFA or SAPS II score with any of the tested  
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26 201 markers (data not shown).

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3 203 **DISCUSSION**

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5 204 COPD is a heterogeneous disease and includes different clinically relevant subtypes. A divergent  
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7 205 distribution of parenchymal (emphysema) and bronchial airway (chronic bronchitis) (CB) disease  
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9 206 contributes to the phenotypic heterogeneity in COPD (12). COPD is associated not only with an  
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11 207 abnormal inflammatory response in the lung but also with systemic inflammation, including  
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13 208 systemic oxidative stress, activation of circulating inflammatory cells and increased circulating  
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15 209 levels of inflammatory cytokines (12). The low-grade systemic inflammation present in COPD  
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17 210 patients may be responsible for the systemic clinical manifestations of the disease including  
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19 211 malnutrition, muscle wasting, osteoporosis, cardiovascular diseases, type II diabetes, anaemia and  
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21 212 depression (13).

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23 213 This study explored whether lung and systemic inflammation occur concurrently and similarly in  
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25 214 patients with acute respiratory failure with and without CB or COPD. We demonstrated for the first  
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27 215 time that patients with both CB and COPD who underwent an acute exacerbation have a different  
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29 216 inflammatory profile (IP-10 levels) and a different alteration in the innate immune responses (TLR4  
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31 217 expression) in the airways and in the systemic compartment. Differently, in these patients a similar  
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33 218 alteration in the immune regulatory events (low Foxp3 expression) was observed in the airways and  
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35 219 in the systemic compartment. The novel aspects underlined by the present study are related to the  
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37 220 accurate selection of patients with a similar phenotype (CB) and a similar exposure to risk factor  
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39 221 (cigarette smoke) in order to limit the variability of the obtained results and to better identify the  
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41 222 role of the observed alterations in airway obstruction. Most of the studies report data from smoker  
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43 223 or COPD patients with different phenotypes.

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45 224 CB is defined on the basis of chronic cough and sputum due to mucus hypersecretion and  
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47 225 histologically, it is characterised by airway inflammation, hypertrophy of submucosal mucus  
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49 226 secreting glands, and goblet cell hyperplasia (14). CB is a risk factor in COPD for accelerated lung  
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51 227 function decline, increased hospitalization, and increased mortality (2). In this regard it has been  
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53 228 demonstrated that the presence of CB may compromise the sterility of distal airway supporting the  
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3 229 hypothesis of natural progression from the simple mucus hypersecretion to purulent hypersecretion  
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5 230 and obstructive bronchitis ( Hogg, Lancet 2004). Here, it is showed that the expression of TLR4 is  
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7 231 increased in COPD patients with CB and acute exacerbation in the airways but not in the systemic  
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9 232 compartment and this alteration is observed at lower extent also in smokers with CB but without  
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11 233 COPD and without acute exacerbation. TLR4 expressed is associated in these patients with the  
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13 234 presence of neutrophils which represent the predominant cell type in COPD patients. The data  
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15 235 provided extend and integrate previous results from our group showing that in the airways of COPD  
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17 236 patients with acute respiratory failure, there is an increased expression of TLR4 (7). Our data  
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19 237 demonstrate that TLR4 expression is higher in mini-BAL cells in comparison to blood compartment  
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21 238 suggesting an up-regulation of TLR4 at the transit from blood into the airway compartment. This  
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23 239 phenomenon seems to be specific for TLR4 since it has been previously demonstrated that in COPD  
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25 240 patients the expression of TLR2 is lower on sputum neutrophils in comparison to blood  
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27 241 compartment indicating a down-regulation of TLR2 at the transit from blood into the airway  
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29 242 compartment (16).  
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35 243 TLR4 signaling, through MyD88 and IRAK1, plays a predominant role as a regulator of smoke-  
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37 244 induced protease production (17). Furthermore, CSE increase the expression of TLR4 but not of  
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39 245 TLR2 and modify the functional activation of TLR4 generating an imbalance between cytokines  
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41 246 with opposite functions such as IL-8 and IP-10 (6). IP-10 concentrations in mini-BAL of smoker  
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43 247 COPD who are mechanically ventilated for acute exacerbation are reduced in comparison to another  
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45 248 group of patients mechanically ventilated but not smokers and without COPD (7). We confirm here  
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47 249 the presence of reduced concentrations of IP-10 within the airways of mechanically ventilated and  
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49 250 exacerbated COPD patients and demonstrate for the first time that within the systemic compartment  
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51 251 in the same patients an increased concentration of IP-10 is observed in comparison to patients with  
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53 252 acute respiratory failure but without CB and COPD. Nasal epithelial cells obtained from smokers  
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55 253 create an overall cytokine microenvironment that after infection with influenza suppresses the  
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3 254 concentrations of IP-10 (18). When the bronchial epithelial cells were exposed to CSE, the release  
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5 255 of IP-10 decreases while the release of IL-8 increases (6). Although no significantly different  
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7 256 concentrations of IL-8 were observed in mini-BAL and in peripheral blood between COPD, S and C  
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9 257 within the airways the elevated concentrations of IL-8 are not balanced by elevated IP-10  
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11 258 concentrations. The prevalence of IL-8 may in turn sustain the influx of neutrophils into the airways  
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13 259 thus triggering innate immunity responses, while IP-10 attracts monocytes and lymphocytes (19)  
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15 260 thus promoting activation of adaptative responses to efficiently and specifically limit microbial  
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17 261 invasion and to restrain the harmful effects of prolonged neutrophil activation. The findings that  
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19 262 mini-BAL from COPD had reduced IP-10 concentrations and reduced chemotactic activities  
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21 263 toward lymphocytes (7) might contribute to explain why the differential cell counts of mini-BAL  
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23 264 from COPD failed to have lymphocytes and prompted us to explore whether lymphocyte regulatory  
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25 265 activities were altered in these patients. CD4<sup>+</sup> Foxp3<sup>+</sup> regulatory T lymphocytes (Treg) are a  
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27 266 subclass of CD4<sup>+</sup> T cell receptor (TCR)  $\alpha\beta$ <sup>+</sup> T cells that are essential to preserve immune  
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29 267 homeostasis (20), (21). Stable expression of the transcription factor Foxp3 is a prerequisite for the  
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31 268 maintenance of suppressive properties in CD4<sup>+</sup> regulatory T cells. Foxp3 mRNA expression is not  
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33 269 itself sufficient for stable Foxp3 protein expression (22). The epigenetic modifications, such as  
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35 270 histone modification or DNA methylation, control regulatory T cells by controlling Foxp3 gene  
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37 271 expression through altering the accessibility of the Foxp3 locus and by acetylating or deacetylating  
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39 272 Foxp3 protein, thereby enabling the epigenetic regulation of Foxp3 target genes. Absence of the  
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41 273 Foxp3 transcription factor at systemic level leads to the rapid development of fulminant multiorgan  
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43 274 autoimmunity. A decreased Foxp3 expression in COPD patients and smokers parallels the  
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45 275 aggravation of the disease (23). Patients with moderate or severe COPD upon fluticasone and  
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47 276 salmeterol combination therapy show an increased proportion of Foxp3+Tregs in the total  
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49 277 peripheral blood CD4+T cell population (24). Smokers with normal lung function and COPD  
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51 278 patients have increased numbers of Foxp3-positive cells in large airways but they have decreased  
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53 279 numbers of Foxp3-positive cells in small airways (25). In the patients with acute respiratory failure  
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3 280 recruited in the present study, the decreased expression of Foxp3 is present in smokers and in  
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5 281 COPD in the distal airways and in the systemic compartment suggesting that the alterations in the  
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7 282 immune regulatory activities are early events in the disease and may contribute to the systemic  
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9 283 effects of the disease. It is conceivable that cigarette smoke may contribute to induce epigenetic  
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11 284 modifications leading to the reduced expression of Foxp3 in smokers and in COPD smokers at both  
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13 285 the airways and systemic compartment. No study has already specifically addressed this point yet.  
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17 286 Finally, the relevance of the observed alterations in the severity of the recruited patients was  
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19 287 assessed. Several commonly used scoring systems exist assessing the severity of the disease in  
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21 288 critically ill patients by predicting mortality. In the present study the recruited patients were  
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23 289 classified on the basis of their SOFA, SAPS II and CPIS score. The mortality in elderly patients  
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25 290 was higher than that of the younger patients and SAPS II (26) was an independent predictor of  
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27 291 mortality in elderly patients with sepsis (27). The SOFA score, widely used in many cardiac  
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29 292 surgical intensive units, is used for grading organ dysfunction or failing organ system (28) .  
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31 293 Prognostic relevance of the SOFA score in combination with inflammatory parameters was also  
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33 294 found in a recent study conducted by Zügel *et al.* (29). The CPIS score is calculated on the basis of  
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35 295 points assigned for various signs and symptoms of pneumonia (eg, fever and extent of oxygenation  
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37 296 impairment) and a CPIS >6 may serve as a surrogate tool to facilitate the diagnosis of ventilator-  
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39 297 associated pneumonia (30). In the present study, recruited patients have no significant different  
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41 298 SOFA or SAPS II scores and these scores did not correlate with any of the tested markers. The only  
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43 299 differences among the recruited patients were related to the CPIS score and CPIS score correlates  
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45 300 with TLR4 mini-BAL expression. Future studies on a larger cohort of patients are needed to clarify  
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47 301 whether the assessment of TLR4 expression in miniBAL may improve the predictive value of CPIS  
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49 302 to early identify patients with ventilator-associated pneumonia.  
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56 303 In conclusion, patients with both CB and COPD who underwent an acute exacerbation have a  
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58 304 different alteration in the levels of IP-10 and in TLR4 expression in the airways and in the systemic  
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3 305 compartment while they have a similar alteration in Foxp3 expression in the airways and in the  
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5 306 systemic compartment. The inflammatory profile of the COPD patients in the present study may be  
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7 307 the result of the failure to properly clear the inflammatory cells after the exacerbation of the disease.  
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312 Elisabetta Pace and Maria Ferraro designed the study, performed the statistical analysis of the data e

313 wrote the manuscript and declares that they have had access to and takes responsibility for the

314 integrity of the data and the accuracy of the data analysis.

315 Chiara Cipollina performed all the experiments of the study and participated to the interpretation of

316 the data.

317 Antonino Giarratano contributed to the patient selection, collected and managed biological

318 samples.

319 Mark Gjomarkaj contributed to the interpretation of the data and to the writing out of the

320 manuscript.

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413 **LEGENDS TO THE FIGURES**

414 **Figure 1. Increased expression of TLR4 in mini-BAL but not in the peripheral blood of S and**  
415 **of COPD.** Mini-BAL cells and paired blood samples were recovered from C (n=10), from S (n=8)  
416 and from COPD (n=18) patients. The expression of TLR4 was assessed by immunocytochemistry  
417 using an anti-TLR4 polyclonal antibody; \* p<0.05 vs C; \*\* p<0.05 vs S; # p<0.05 vs autologous  
418 peripheral blood. Data are expressed as median (25-75 percentiles).

419 **Figure 2. Absence of differences in the concentrations of IL-8 in mini-BAL and in the**  
420 **peripheral blood of S and of COPD.** Mini-BAL supernatants and paired blood samples were  
421 recovered from C (n=10), from S (n=8) and from COPD (n=18) patients. IL-8 concentrations were  
422 measured by ELISA as described in "materials and methods" and are expressed as pg/ml. Data are  
423 expressed as median (25-75 percentiles).

424 **Figure 3. Reduced concentrations of IP-10 in mini-BAL but not in the peripheral blood of**  
425 **COPD.** Mini-BAL supernatants and paired blood samples were recovered from C (n=10), from S  
426 (n=8) and from COPD (n=18) patients. IP-10 concentrations were measured by ELISA as  
427 described in "materials and methods" and are expressed as pg/ml. \* p<0.05 vs C; \*\* p<0.05 vs S; #  
428 p<0.05 vs autologous peripheral blood. Data are expressed as median (25-75 percentiles).

429 **Figure 4. Reduced expression of Foxp3 in mini-BAL and in the peripheral blood of S and of**  
430 **COPD.** Mini-BAL cells and paired blood samples were recovered from C (n=10), from S (n=8) and  
431 from COPD (n=18) patients. Total proteins were extracted and analysed for Foxp3 expression by  
432 western blot analysis. Membranes were then stripped and incubated with goat polyclonal anti-β-  
433 actin.. **A.** Densitometric analysis of Foxp3 expression. Signals corresponding to Foxp3 on the  
434 various western blots were semiquantified by densitometric scanning, normalized and expressed  
435 after correction with the density of the band obtained for beta-actin (mean±SD). \* p < 0.05. **B.**  
436 Representative western blot analysis for Foxp3 expression from C, S and COPD subjects. Data are  
437 expressed as median (25-75 percentiles).

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3 438 **Figure 5. Correlations between the expression of TLR4 in miniBAL and CPIS score.** The  
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5 439 expression of TLR4 in mini-BAL of C (n=10), S (n=8) and COPD (n=18) was correlated with  
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7 440 CPIS score by Spearman Correlation test.  
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**Table 1: Demographic and clinical characteristics of the subjects:**

<b>Characteristic</b>	<b>C</b>	<b>S</b>	<b>COPD</b>	<b>P value</b>
<b>No. of subjects</b>	<b>10</b>	<b>8</b>	<b>18</b>	
Age (yr)	73 (72-78)	79.5 (76-82)	76 (71-81)	n.s.
Male/Female	7/3	5/3	11/7	n.s.
Cigarette smoke Packs/years	-	45±20	50±21	n.s.
FEV1( % of predicted normal)	85±2.5	87±3.9	67±5*	*p<0.05 COPD vs C and S
FEV1/FVC ( % of predicted normal)	77±3	76±2	65±2*	*p<0.05 COPD vs C and S
CPIS	2.3±1.1	3.3±1.4	3.7±0.9*	*P<0.05 COPD vs C
SOFA	9.1±1.2	8.7±1.3	8.9±1	n.s.
SAPS II	53.6±7.8	59.5±11	56±7	n.s.

**Data are expressed as percentiles (age) or as mean ± SD.**

1 **Table 2: Total and differential cell counts of mini-BAL**

	<b>C =10</b>	<b>S=8</b>	<b>COPD=18</b>	<b>P value</b>
<b>Mini-BAL cells</b>				
Total number (X1000)/ml	536±269	919±167*	3,2±2,400*	* p<0.05 COPD and S vs C
% Neutrophils	46±30	48±17	83±13*	*p<0.05 COPD vs C and S
% Macrophages	51±29	52±16	17±13*	*p<0.05 COPD vs C and S
% Lymphocytes	2.7±1.3	0*	0*	*p<0.05 COPD and S vs C

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5 Results are expressed as mean ± SD.

Table 2: Microbiology of miniBAL

	C with MiniBAL cultures Positive (n=4)	C with Mini BAL cultures Negative (n=6)	CFU
Patient#1		X	
Patient#2		X	
Patient#3	Candida		$\geq 10^3$ /ml
Patient#4	Candida		$\geq 10^3$ /ml
Patient#5		X	
Patient#6		X	
Patient#7	Candida		$\geq 10^3$ /ml
Patient#8		X	
Patient#9		X	
Patient#10	Candida		$\geq 10^3$ /ml
	S with MiniBAL cultures Positive (n=3)	S with Mini BAL cultures Negative (n=5)	CFU
Patient#1		X	
Patient#2		X	
Patient#3	Candida		$\geq 10^3$ /ml
Patient#4	Candida		$\geq 10^3$ /ml
Patient#5		X	
Patient#6		X	
Patient#7		X	
Patient#8	Acinetobacter Baumannii		$\geq 10^3$ /ml
	COPD with MiniBAL cultures positive (n=7)	COPD with Mini BAL cultures negative (n=11)	CFU
Patient#1	Candida		$\geq 10^3$ /ml
Patient#2		X	
Patient#3		X	
Patient#4	Acinetobacter Baumannii		$\geq 10^3$ /ml
Patient#5		X	
Patient#6	Candida		$\geq 10^3$ /ml.
Patient#7		X	
Patient#8	Pseudomonas Aeruginosa		$\geq 10^3$ /ml
Patient#9		X	
Patient#10	Acinetobacter Baumannii		$\geq 10^3$ /ml
Patient#11		X	
Patient#12		X	
Patient#13	Candida		$\geq 10^3$ /ml
Patient#14		X	
Patient#15		X	
Patient#16	Pseudomonas Aeruginosa		$\geq 10^3$ /ml
Patient#17		X	
Patient#18		X	

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3 **Table 2: Total and differential cell counts of mini-BAL**  
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	<b>C =10</b>	<b>S=8</b>	<b>COPD=18</b>	<b>P value</b>
<b>TLR4 expression in Mini-BAL cells</b> (% of positive cells)	21.9±6.4	39.8±3.8	64.88±4.8	*p<0.05 COPD and S vs C
<b>TLR4 expression in peripheral blood cells</b> (% of positive cells)	25.4±6.5	17.5±1.76	26.8±6.37	n.s.
<b>Foxp3 expression in Mini-BAL cells</b> (arbitrary units)	2.5±0.24	1.83±0.12*	1.39±0.41*	*p<0.05 COPD and S vs C
<b>Foxp3 expression in peripheral blood cells</b> (arbitrary units)	3.79±0.3	2.36±0.36	1.75±0.53	*p<0.05 COPD and S vs C

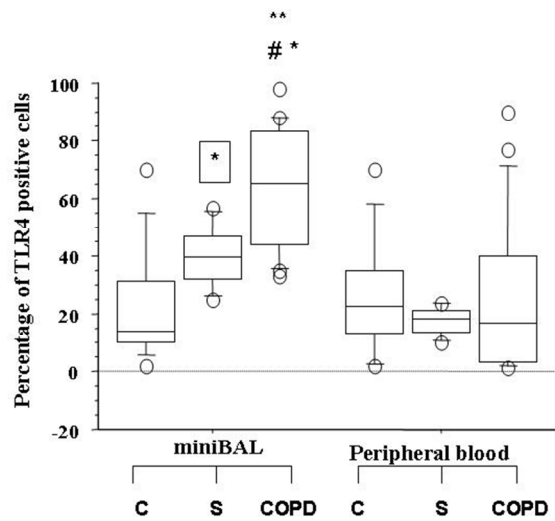
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5 Results are expressed as mean ± SE.

6 \* Mann Whitney test

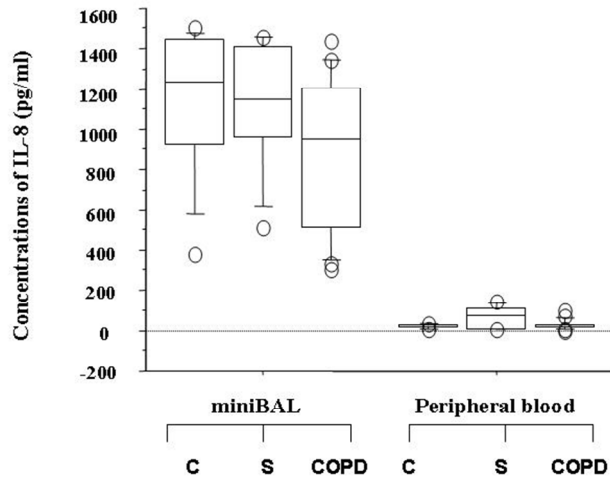
Figure 1

TLR4 expression in miniBAL and in Peripheral blood



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Figure 2  
IL-8 Concentrations in miniBAL and in  
Peripheral blood

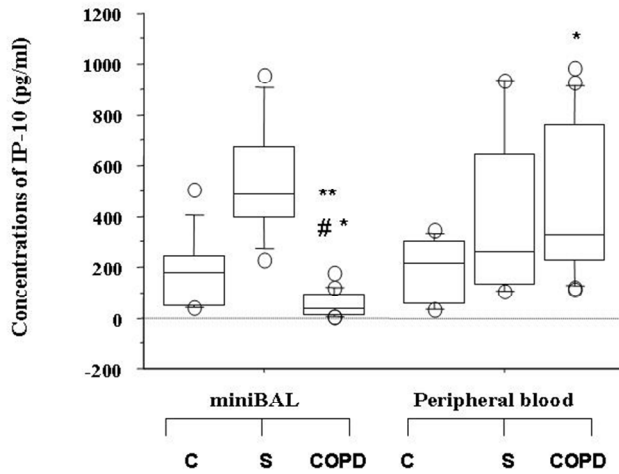


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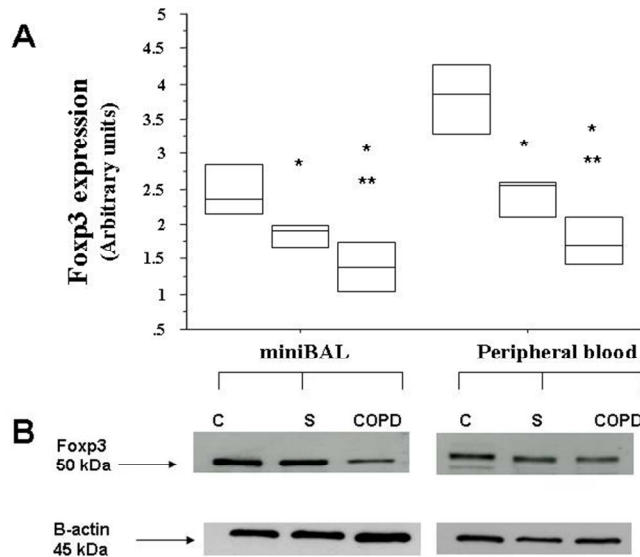
**Figure 3**  
**IP-10 Concentrations in miniBAL and in**  
**Peripheral blood**



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**Figure 4**  
**Foxp3 expression in miniBAL and in Peripheral blood**



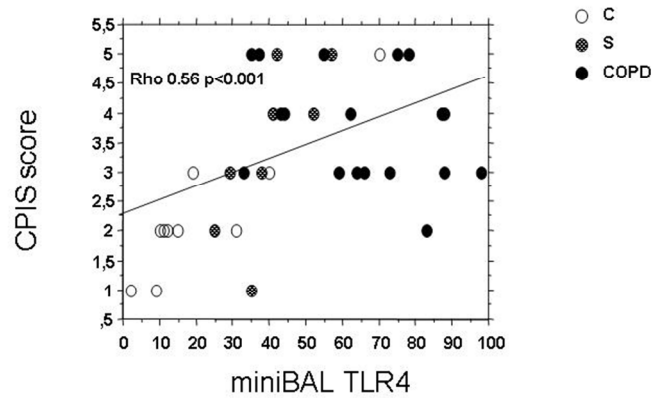
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**Figure 5**  
**TLR4 expression in miniBAL correlates with**  
**CPIS score**



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