# SU/A/AIT RESEARCH

# Microbial, inflammatory and clinical determinants of Acute Exacerbations of Chronic obstructive pulmonary disease



Sara Dias



**Alda Marques** 



Ana Sousa



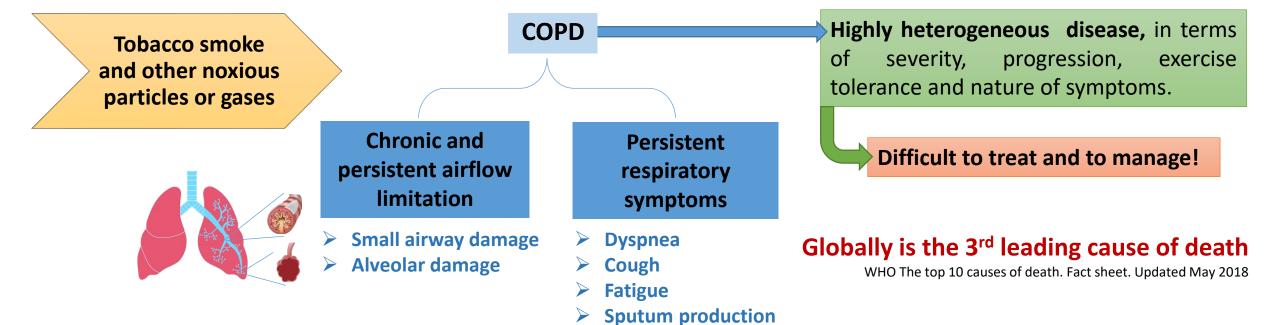
Laboratório de Investigação e Reabilitação Respiratória



## SU/A /A iT RESEARCH



# **Background:**



#### **ACUTE EXACERBATIONS (AECOPD)**

Transient and apparently stochastic periods of increased symptoms.

Are typically **treated** with a combination of antibiotics and steroids **in a non-specific manner, without understanding its aethiology.** 

# **Research challenges:**

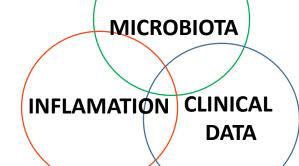
What is the role of lung microbiota in the onset of AECOPD?

Is the pathoadaptation of the resident microbiota an essential trigger of AECOPD?

What is the advantage of combining multi-type data (microbial, inflammatory and clinical) to predict AECOPD?

# Hypothesis: DYSBIOSIS of the commensal airway/lung microbiota PRECEDES THE ONSET OF AECOPD



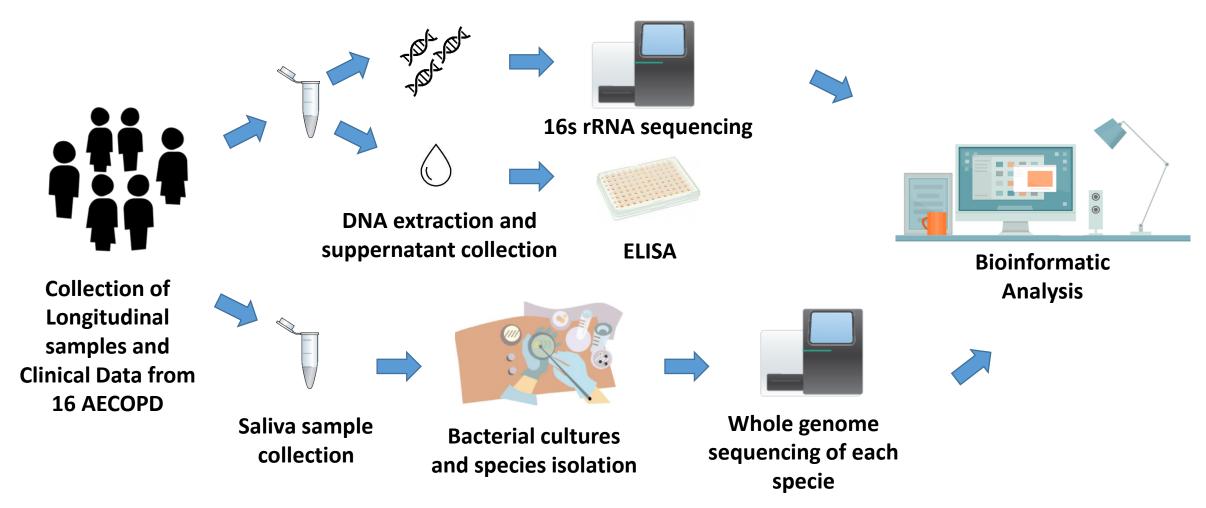




# SU/A A IT



**Research method and techniques:** 



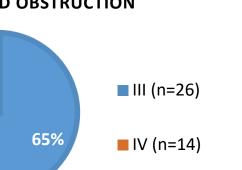
## SUM MIT RESEARCH



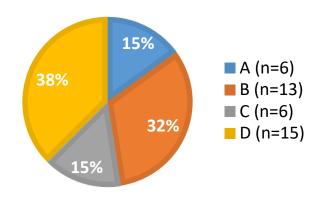
<sup>5</sup> GOLD 2018



#### OUR SAMPLE (n=40)



#### **GOLD CLASSIFICATION**



#### **GOLD Classification**

#### (Exacerbation history/symptoms)

≥2 or ≥1 leading to hospital admission	С	D
0-1 not leading to hospital admission	Α	В
	mMRC 0-1	mMRC ≥2
	CAT<10	CAT ≥10

**GOLD Obstruction** 

(Airflow limitation)









I-Mild FEV<sub>1</sub>≥80%

**II-Moderate III-Severe** 50%≤FEV<sub>1</sub><79% 30%≤FEV<sub>1</sub><49%

**IV-Very severe**  $FEV_1 < 30\%$ 

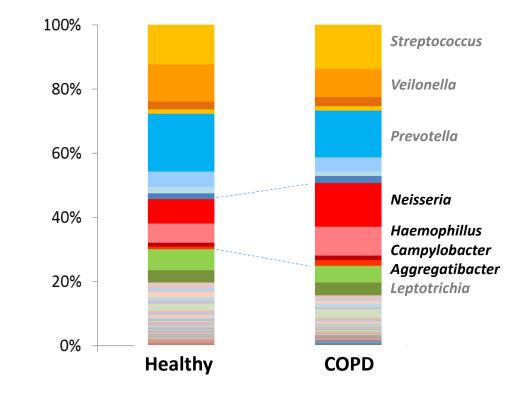




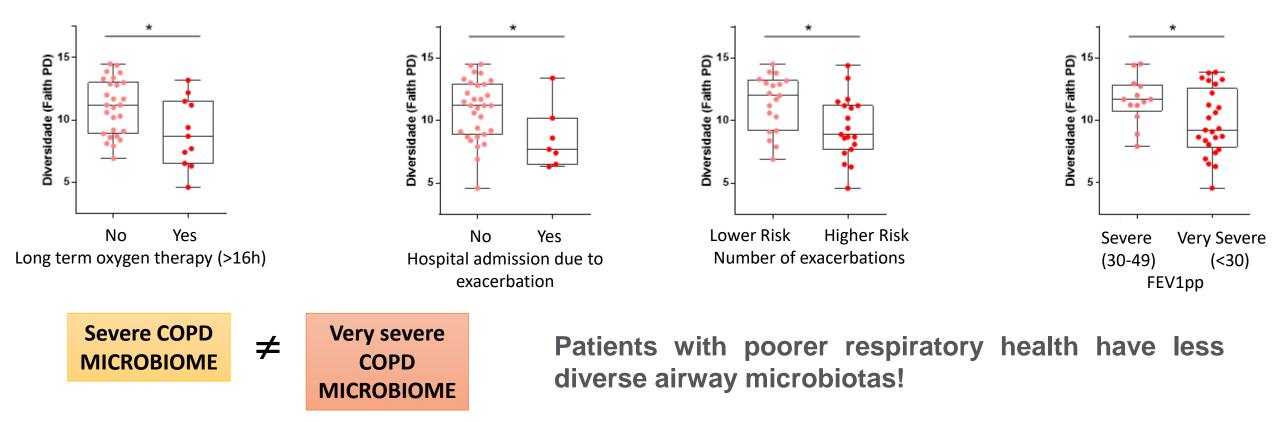
**Results: Microbiota** diversity and composition is different GN02 BD1\_5 among patients and healthy controls

Gammaproteobacteria

EALTHY



Patients with COPD have an enrichment of *Proteobacteria* phylum, specifically of *Neisseria*, *Haemophilus*, *Campylobacter* and *Aggregatibacter* genera.



# **Results:**

It was also possible to stratify patients according to several clinical parameters:





## SU/A /A iT RESEARCH



## **Current impact:**



Sara Dias, Filipa Machado, Carla Valente, Lilia Andrade, Alda Marques, Ana Sousa 2018, "Airway microbiota diversity and composition correlates with severity of Chronic Obstructive Pulmonary Disease (COPD)" (submitted Abstract for Mechanisms of Acute Exacerbation of Respiratory Disease, 17th Lung Science Conference Science Lung Conference 2018)



Ana Sousa, Sara Dias, Filipa Machado, Carla Valente, Lilia Andrade and Alda Marques 2018, "Airway Microbiota diversity and composition correlates with the severity of Chronic Obstructive Pulmonary Disease" (Oral Communication in XXXIV Congresso Nacional de Pneumologia 2018)

# SU/A AAIT RESEARCH



# **Acknowledgements:**

**Bioinformatic facility** 

Miguel Pinheiro



**ESSUA** escola superior de saúde



#### universidade de aveiro



### • Funding:

This work was funded by Programa Operacional de Competitividade e Internacionalização – POCI, through Fundo Europeu de Desenvolvimento Regional - FEDER (POCI-01-0145-FEDER-007628), Fundação para a Ciência e Tecnologia (PTDC/DTP-PIC/2284/2014 and PTDC/SAU-SER/28806/2017) and under the project UID/BIM/04501/2013.



