

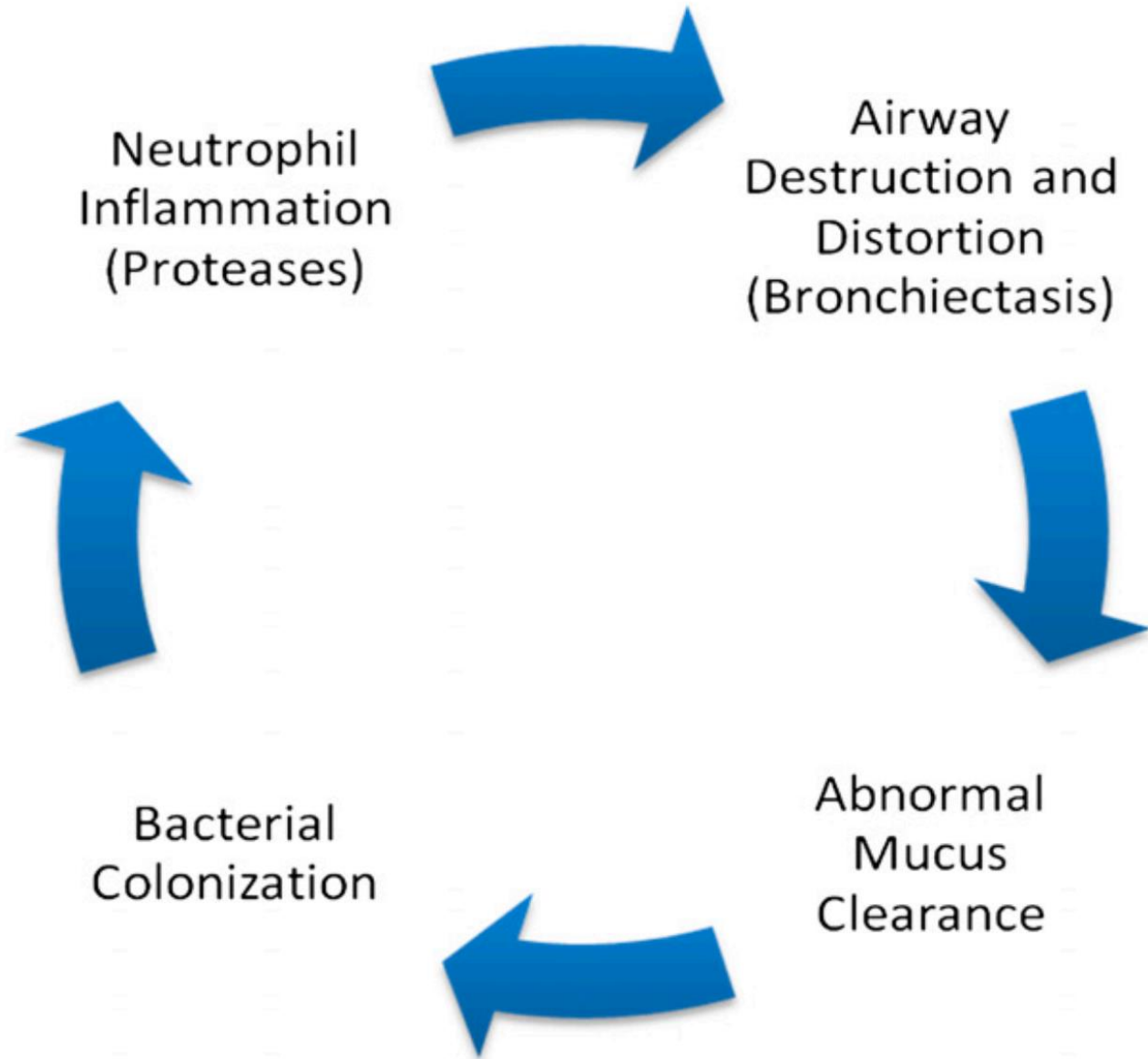
# Advances in Diagnosis and Evaluation of Bronchiectasis

이정규

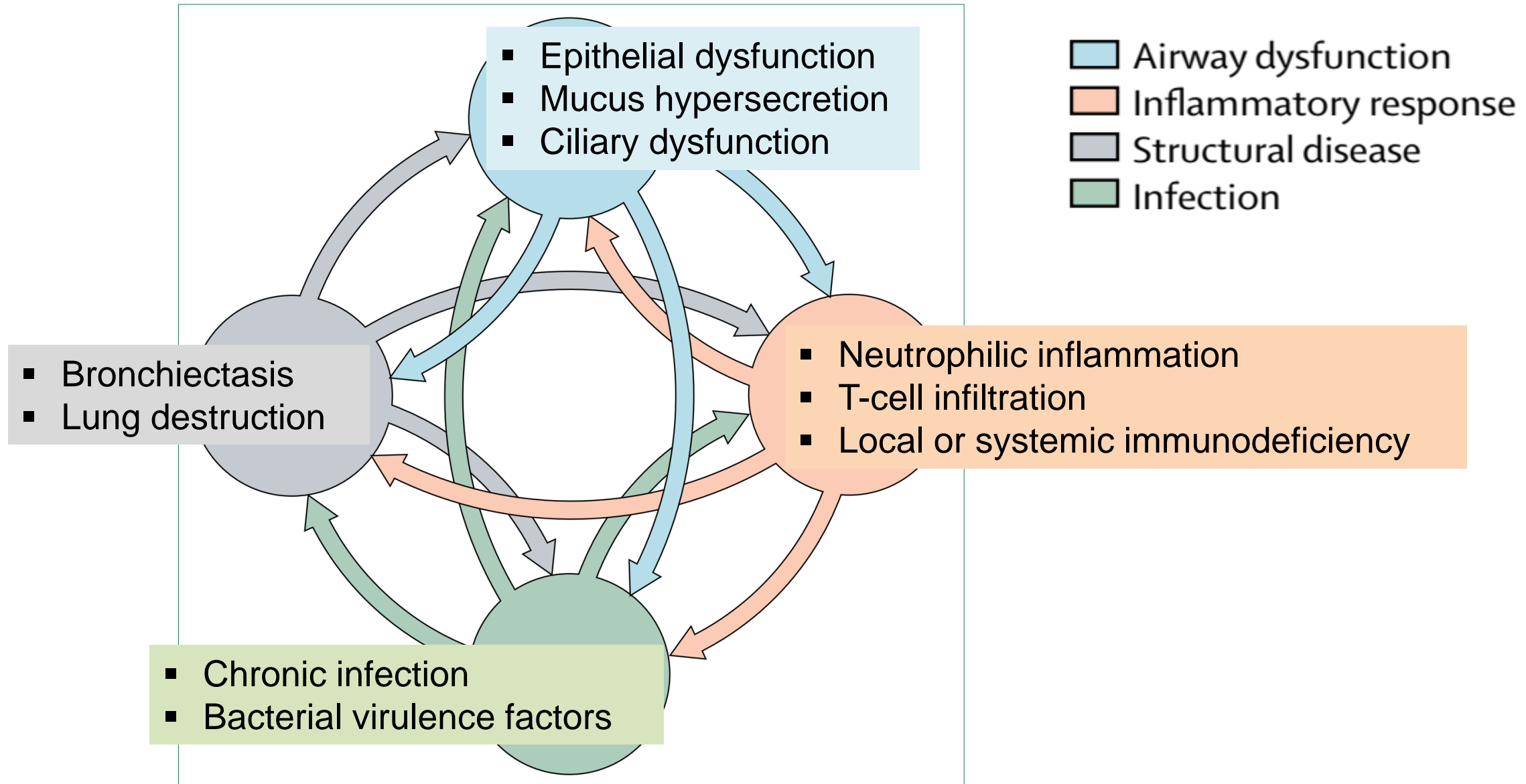
서울대학교 의과대학

서울특별시보라매병원 호흡기내과

# Vicious cycle of bronchiectasis pathogenesis

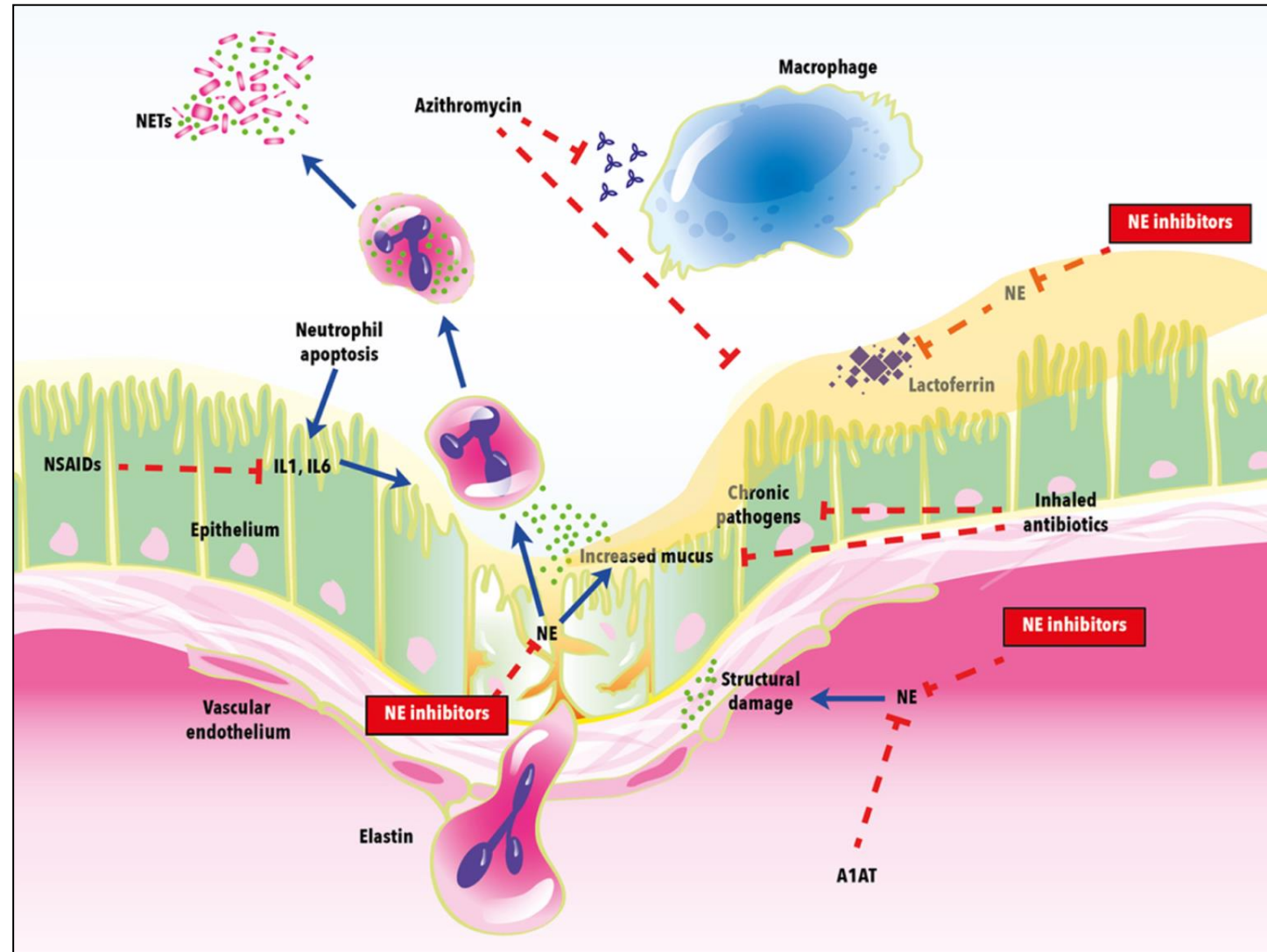


# Recent model for pathogenesis of bronchiectasis



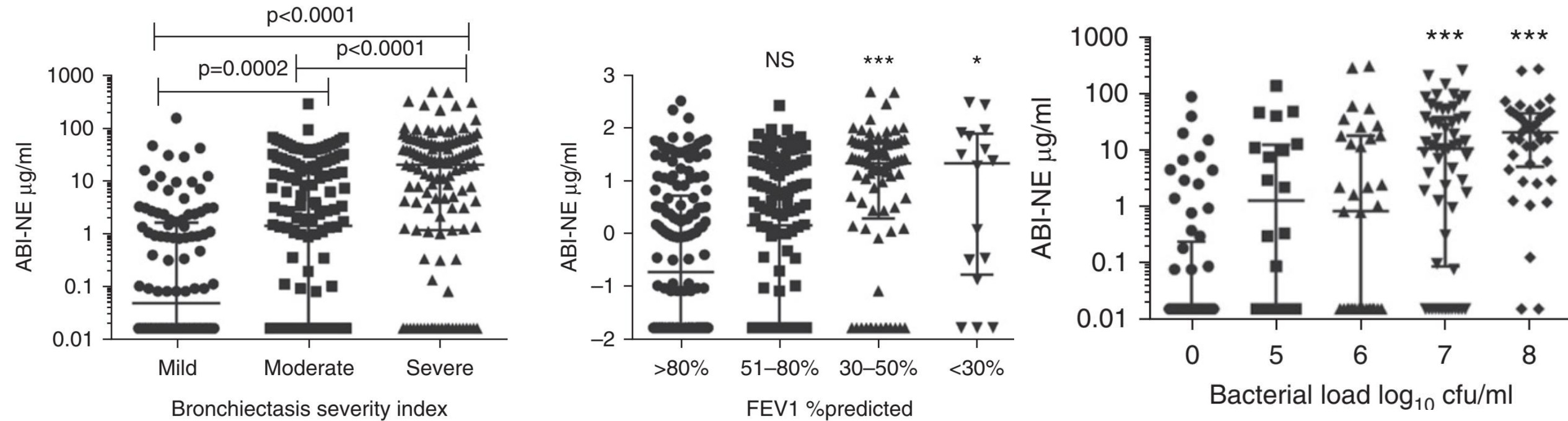
# Neutrophil elastase in bronchiectasis

- Proteolytic enzyme belonging to the chymotrypsin-like family of the serine-proteinases
- Roles in both impaired mucus clearance and chronic bacterial infections leading to a serious neutrophilic activation
- Key determinant of tissue damage

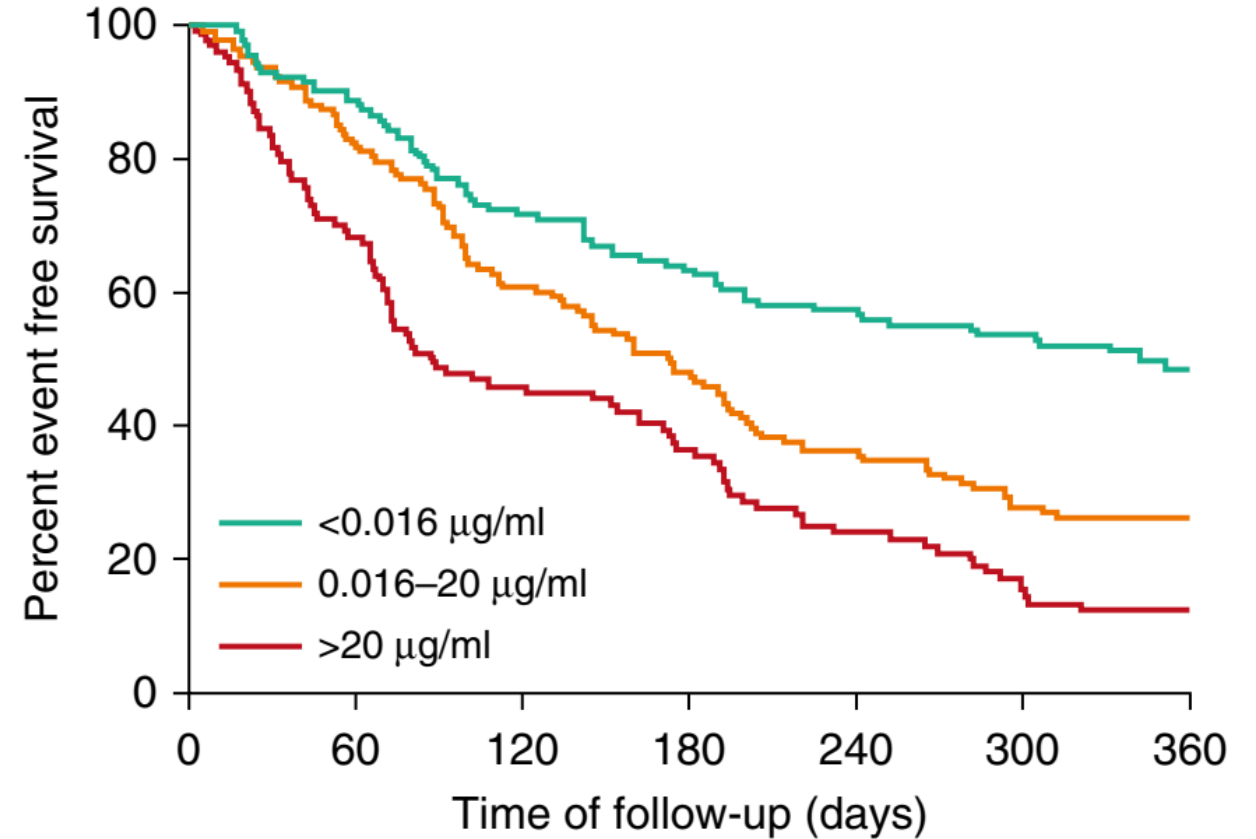
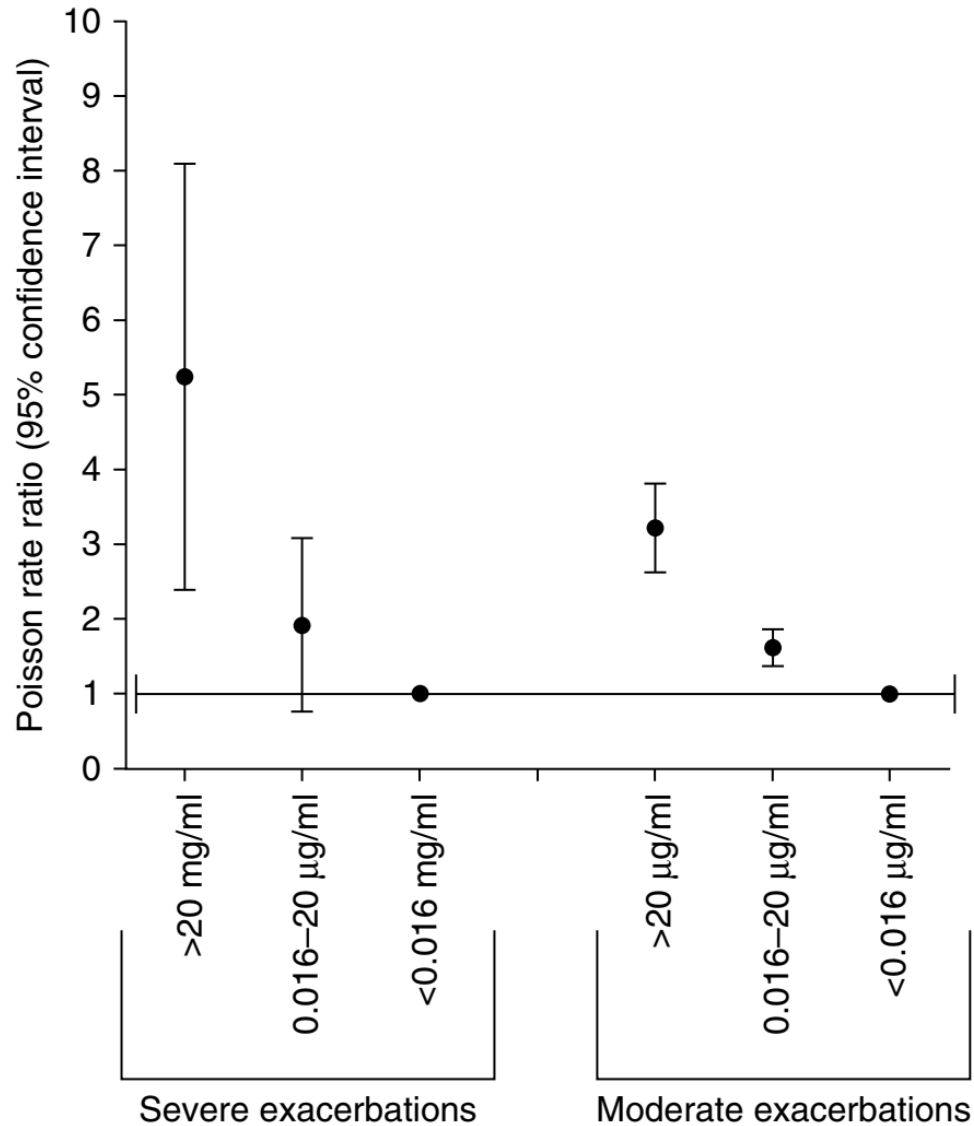


# NE activity and disease severity

- 433 bronchiectasis patients from UK prospective cohort
- sputum NE activity using ProteaseTag active NE immunoassay



# NE activity and clinical course

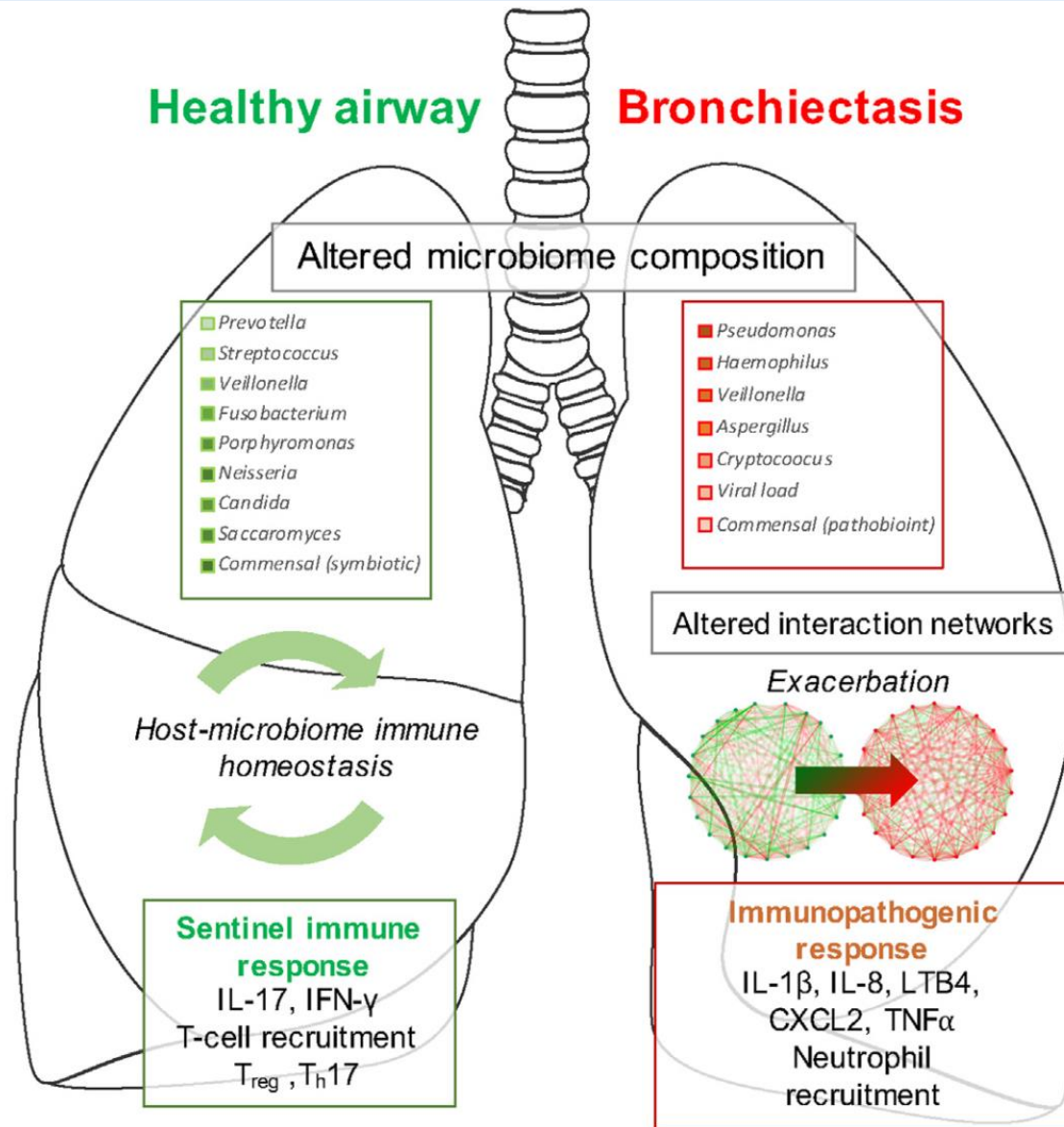


# Clinical characteristics according to sputum NE level

➤ 266 bronchiectasis patients from Southern European cohort

	Sputum NE level			P-value
	Low (0-6 $\mu$ g/ml)	Medium (7-20 $\mu$ g/ml)	High (>30 $\mu$ g/ml)	
Age	65 (51-75)	63 (53-73)	64 (54-74)	0.45
Bronchiectasis severity index	6 (4-9)	7 (4-10)	8 (5-12)	0.009
Sputum volume	6 (5-20)	15 (5-50)	25 (7-75)	<0.001
QoL respiration	74.1 (66.7-81.5)	70.4 (59.3-77.8)	66.7 (51.9-74.1)	0.04
Chronic infection with <i>P.aeruginosa</i>	12 (14.0)	21 (27.6)	44 (47.3)	<0.001
FEV1 <50% predicted	8 (8.9)	9 (12.0)	22 (23.2)	0.02

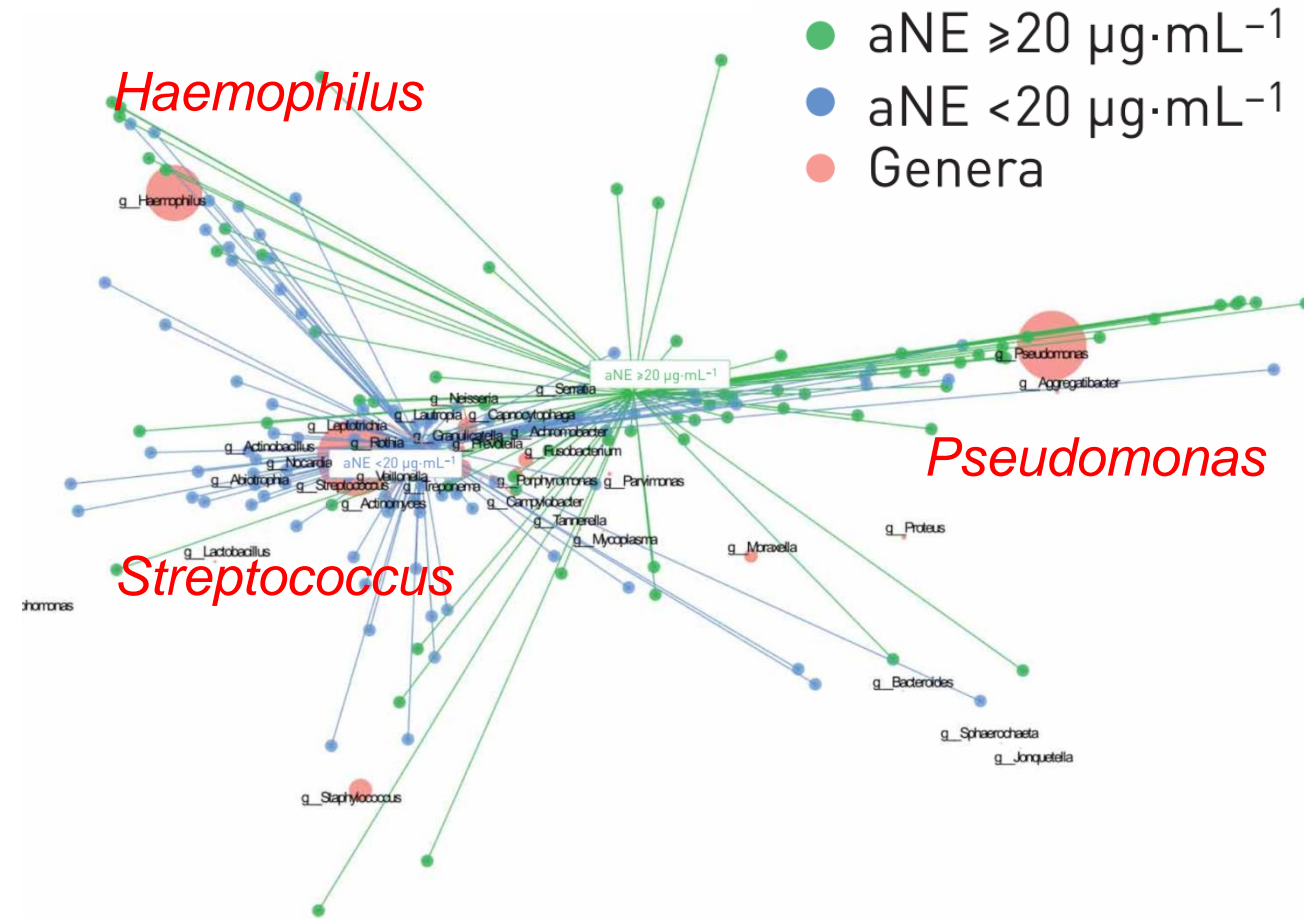
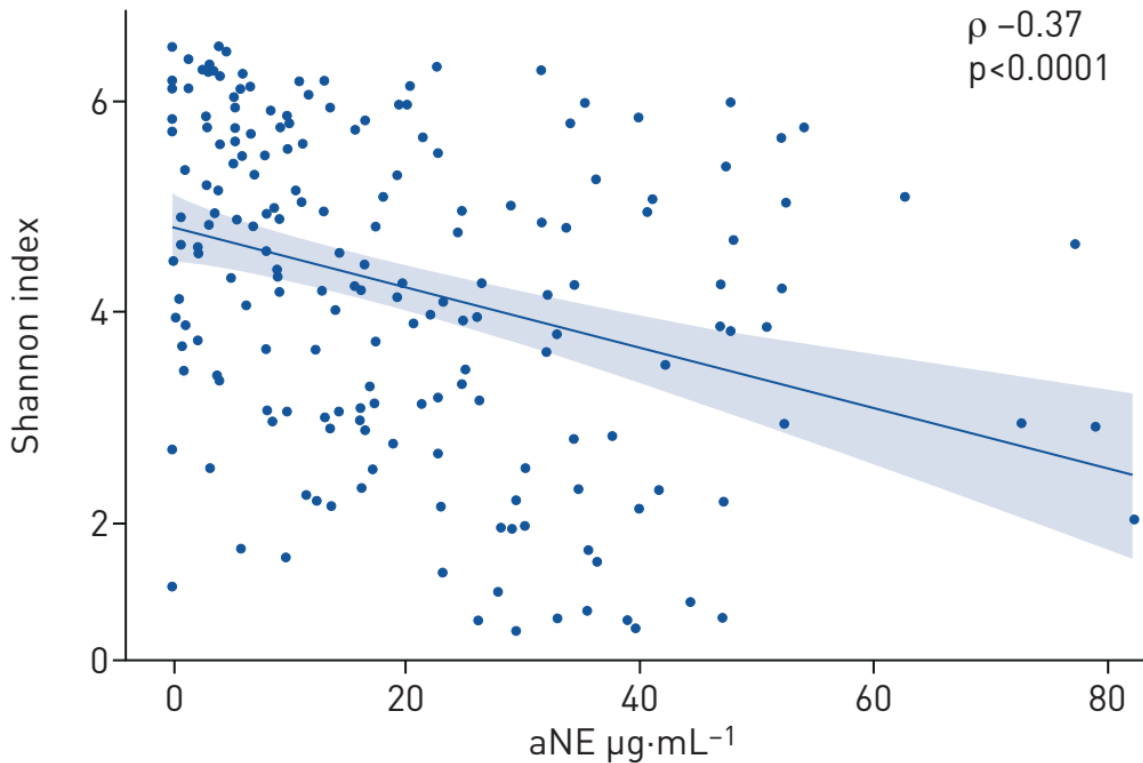
# Microbiome in bronchiectasis



# Sputum NE activity and microbiome

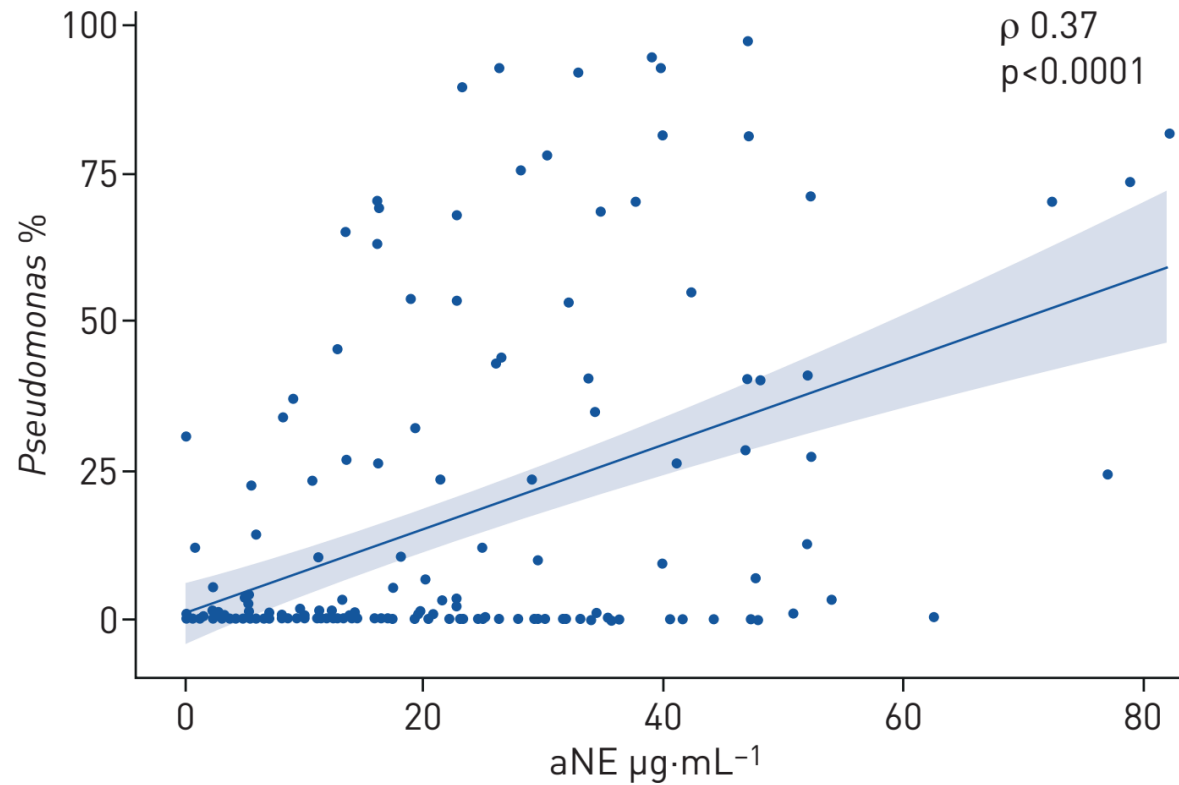
- 185 adult bronchiectasis patients with clinical stable status, Italy
- Sputum NE level measurement and microbiome analysis

## Microbiome $\alpha$ -diversity

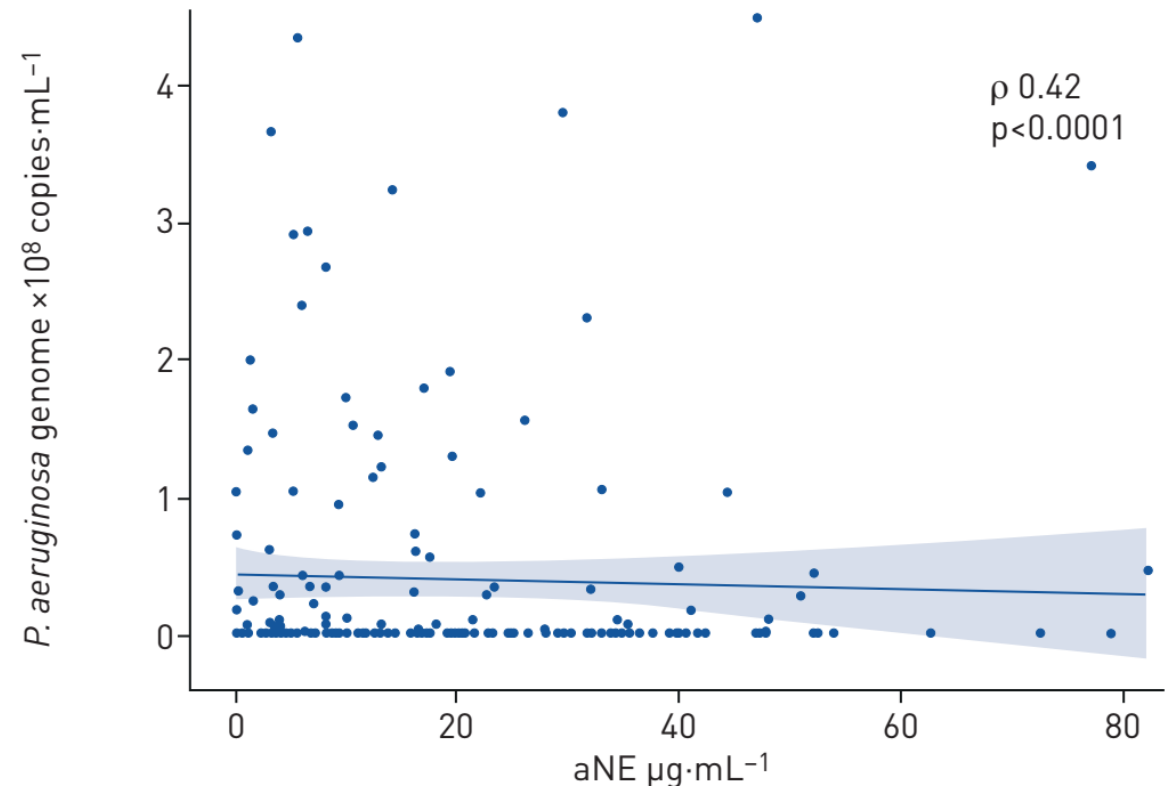


# Sputum NE activity and microbiome

Relative abundance of *P. aeruginosa*

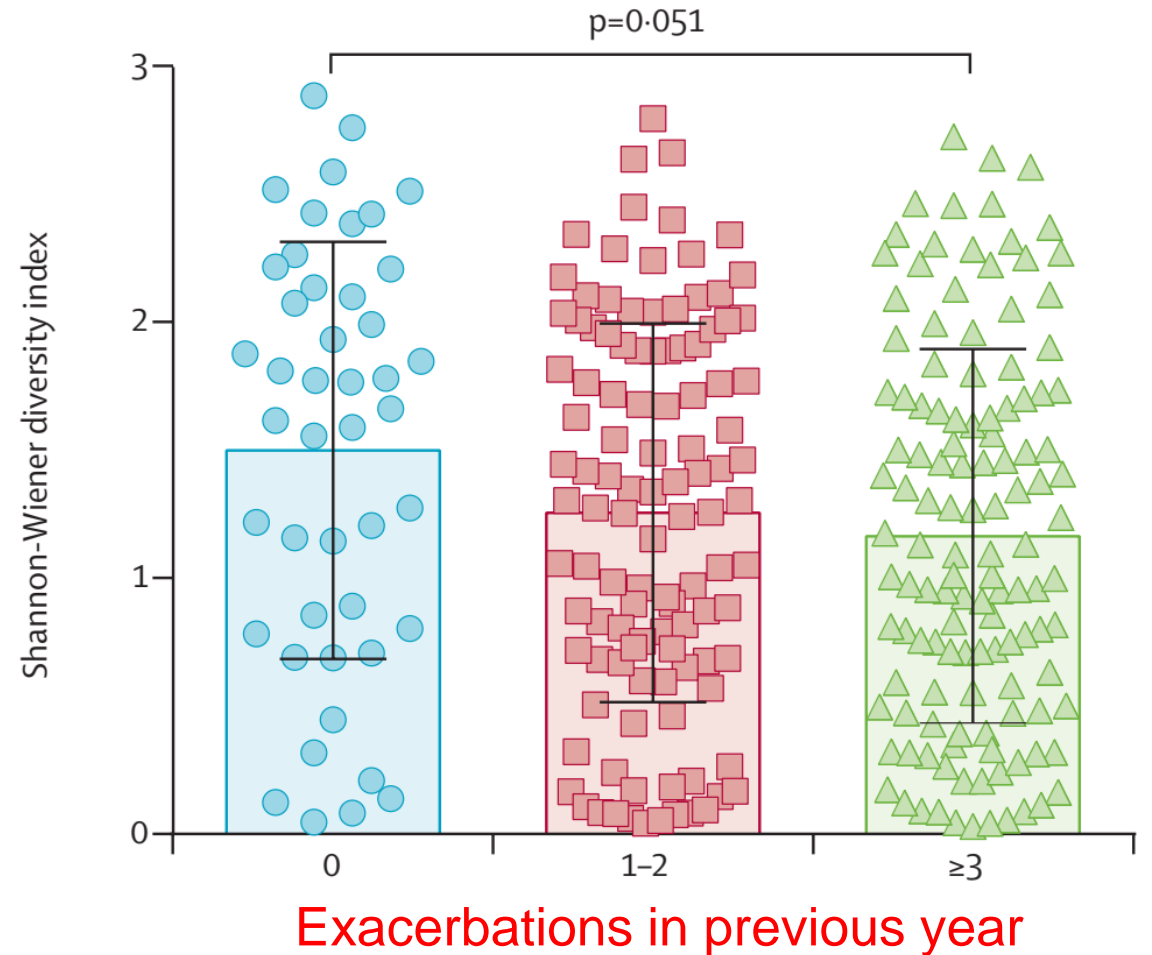
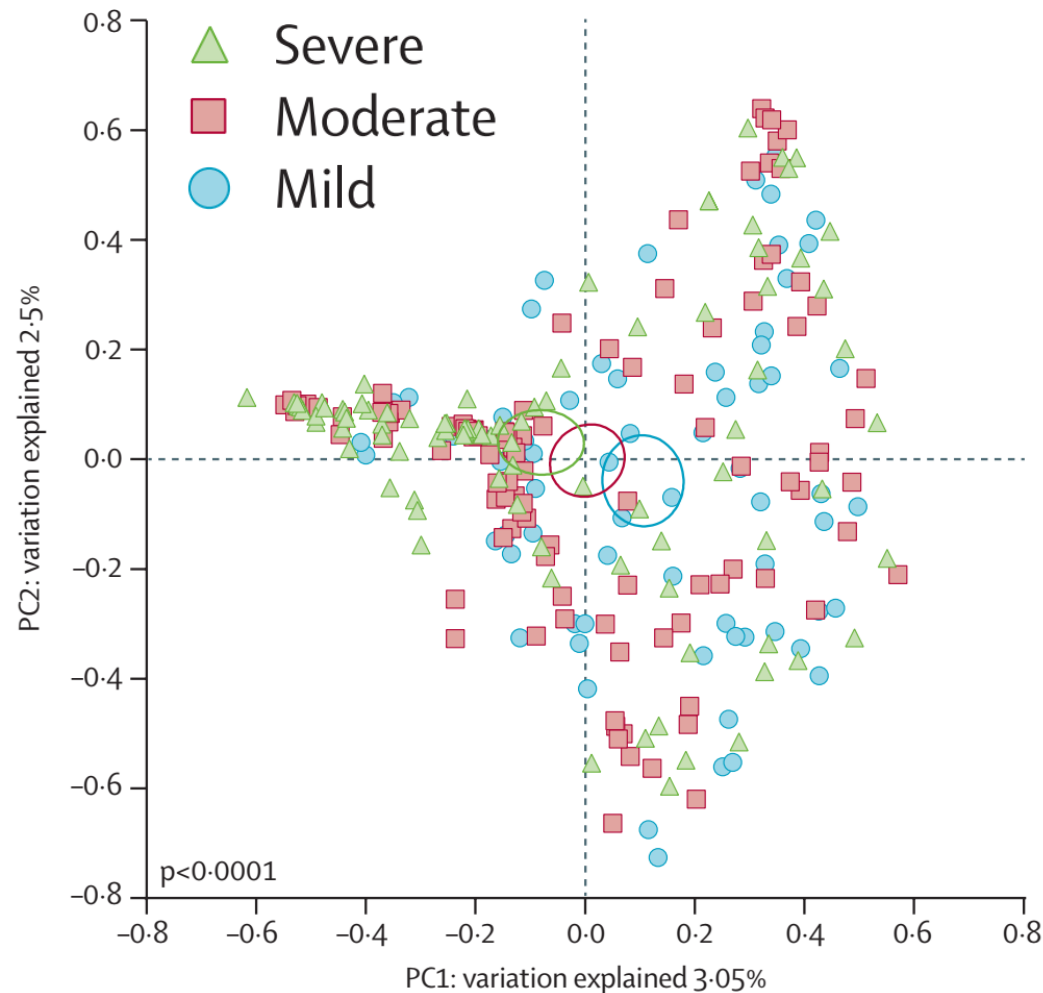


Bacterial burden of *P. aeruginosa*



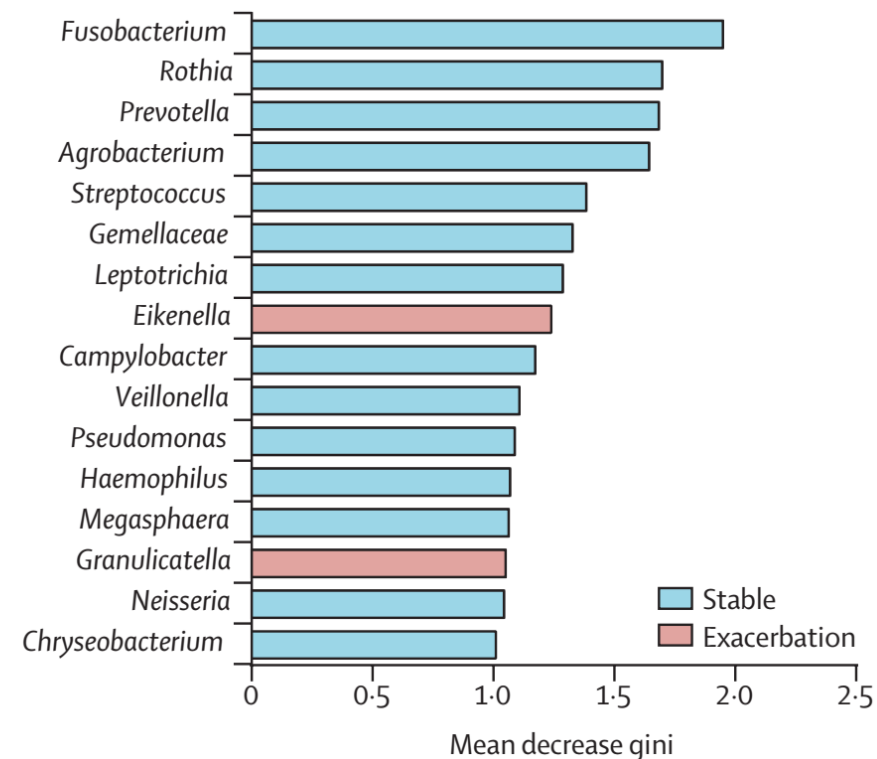
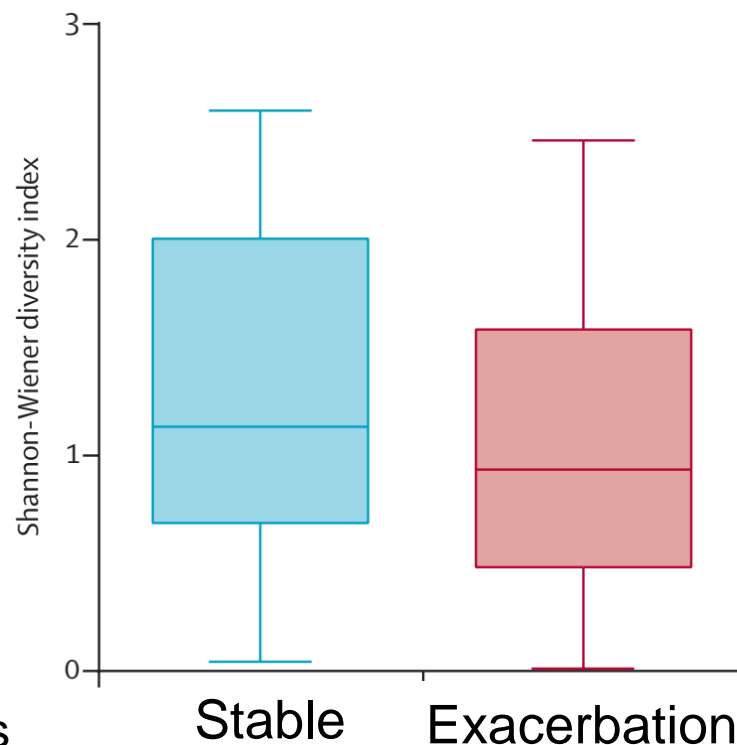
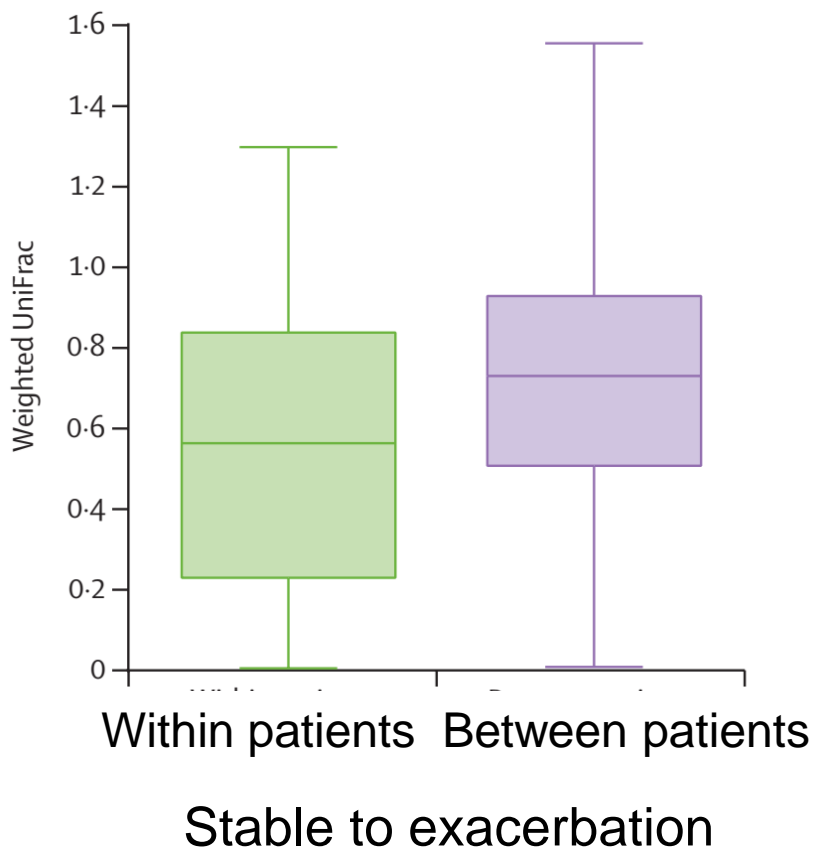
# Sputum microbiome and bronchiectasis severity

- 281 bronchiectasis patients with clinical symptom
- Repeated sputum samples at stable and exacerbation visits



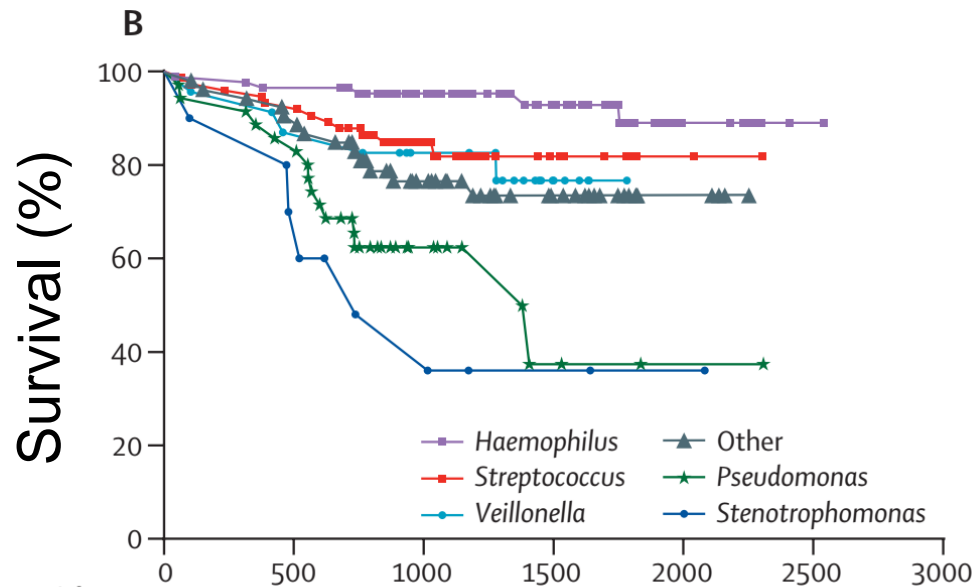
# Sputum microbiome according to exacerbation

Differences in the microbiome between stable state and exacerbation

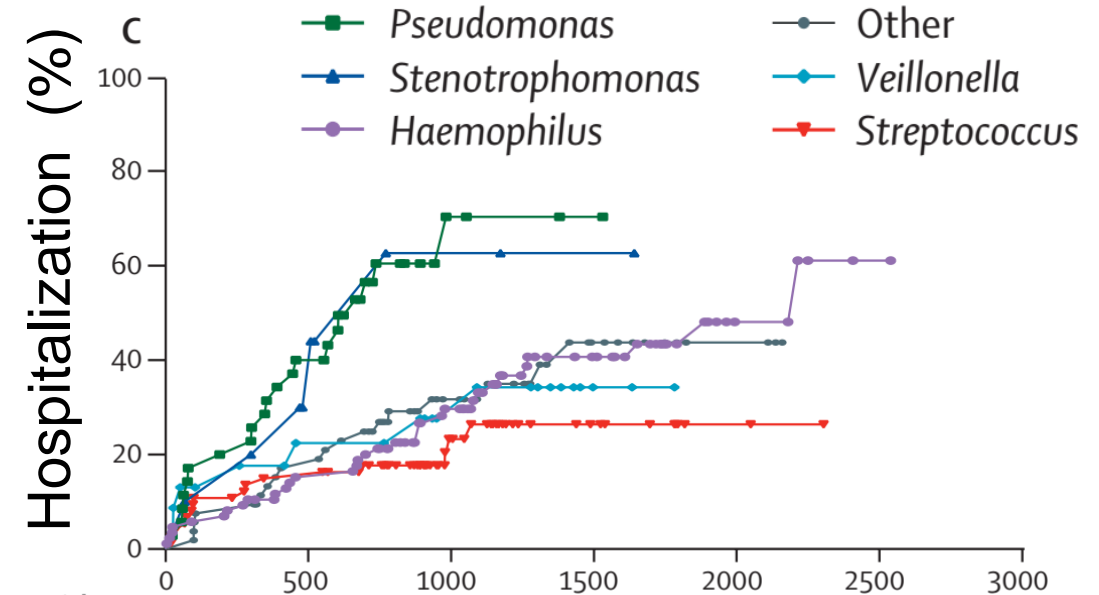


# Microbiome profile and longitudinal outcome

Based on the dominant organism identified at baseline sampling



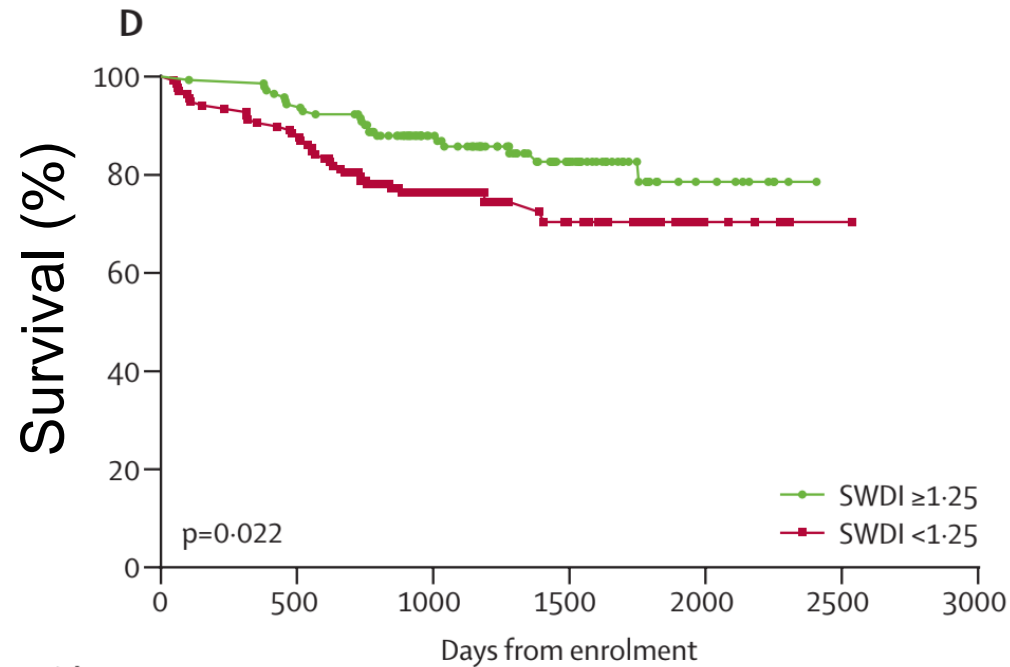
	0	500	1000	1500	2000	2500	3000
<b>Number at risk (number censored)</b>							
Haemophilus	86 (0)	83 (0)	58 (24)	35 (22)	7 (27)	1 (6)	0 (1)
Streptococcus	74 (0)	69 (0)	30 (33)	10 (19)	2 (8)	0 (2)	0 (0)
Veillonella	23 (0)	20 (0)	16 (3)	6 (9)	0 (6)	0 (0)	0 (0)
Other	53 (0)	48 (0)	31 (10)	16 (14)	4 (12)	0 (4)	0 (0)
Pseudomonas	35 (0)	30 (0)	9 (13)	3 (4)	1 (2)	0 (1)	0 (0)
Stenotrophomonas	10 (0)	7 (0)	4 (1)	2 (1)	1 (1)	0 (1)	0 (0)



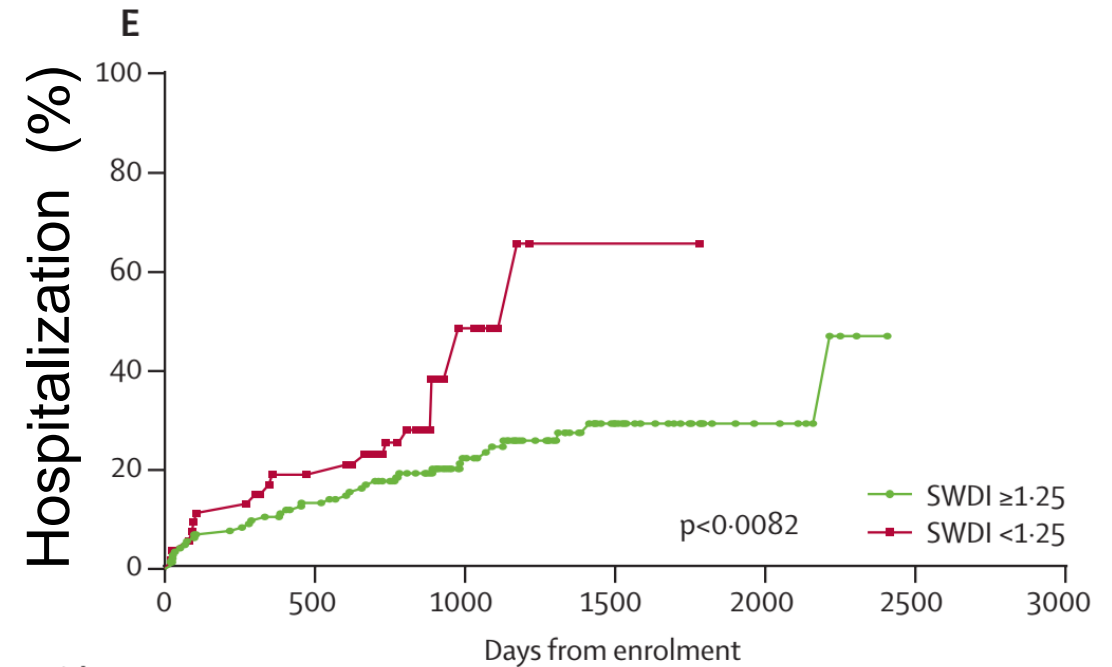
	0	500	1000	1500	2000	2500	3000
<b>Number at risk (number censored)</b>							
Pseudomonas	35 (0)	21 (0)	3 (11)	1 (2)	0 (1)	0 (0)	0 (0)
Stenotrophomonas	10 (0)	5 (2)	2 (1)	1 (1)	0 (1)	0 (0)	0 (0)
Haemophilus	86 (0)	71 (2)	47 (13)	26 (15)	5 (19)	1 (3)	0 (1)
Other	53 (0)	43 (1)	25 (11)	10 (12)	3 (7)	0 (3)	0 (0)
Veillonella	23 (0)	16 (2)	11 (4)	3 (7)	0 (3)	0 (0)	0 (0)
Streptococcus	74 (0)	62 (1)	27 (31)	9 (17)	2 (7)	0 (2)	0 (0)

# Microbiome profile and longitudinal outcome

Based on the microbiome  $\alpha$ -diversity (Shannon-Wiener diversity index)



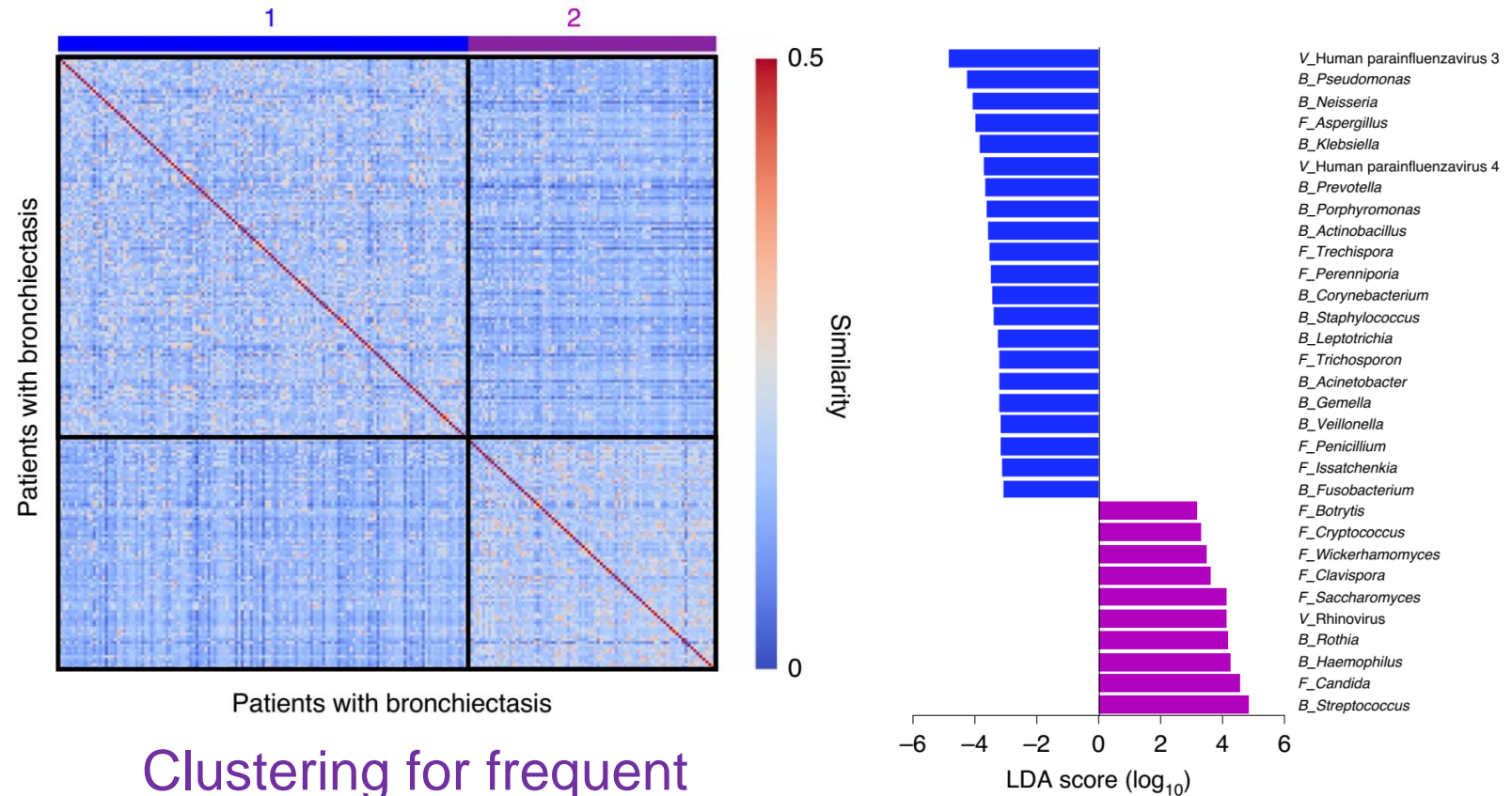
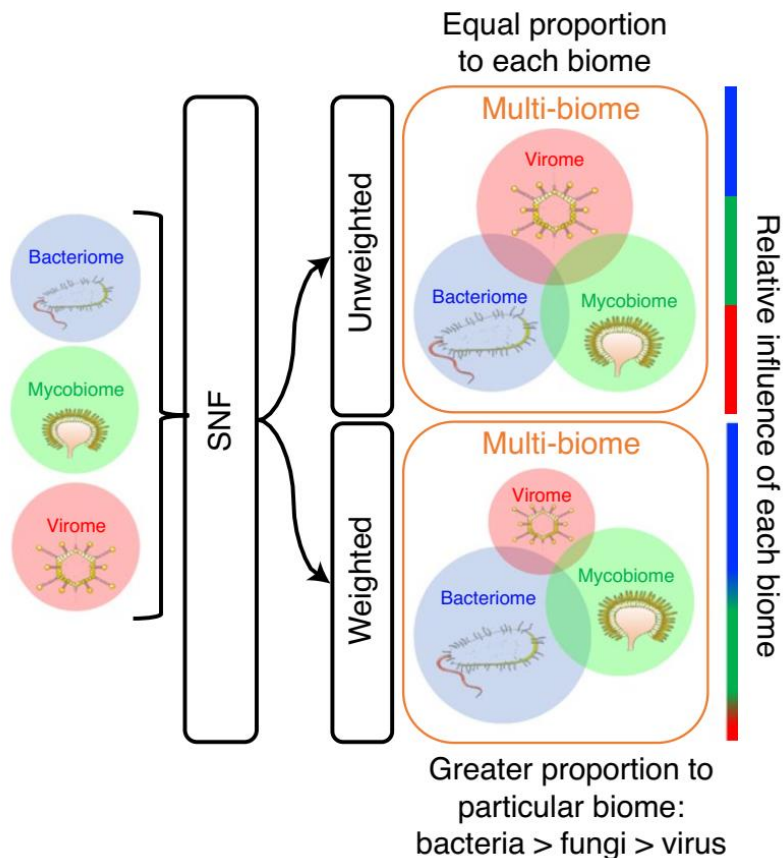
Number at risk	0	500	1000	1500	2000	2500	3000
SWDI $\geq 1.25$	143 (0)	135 (0)	84 (42)	40 (40)	9 (30)	0 (9)	0 (0)
SWDI $< 1.25$	138 (0)	122 (0)	64 (42)	32 (29)	6 (26)	1 (5)	0 (1)



Number at risk	0	500	1000	1500	2000	2500	3000
SWDI $< 1.25$	138 (0)	97 (5)	44 (32)	19 (19)	2 (15)	0 (1)	0 (1)
SWDI $\geq 1.25$	143 (0)	121 (3)	71 (39)	31 (35)	8 (23)	0 (7)	0 (0)

# Multi-biome intergration in bronchiectasis exacerbation

- 217 bronchiectasis patients from Asian and European cohort (CAMEB)
- Respiratory samples with integrated multi-biome analysis (bacteriome/mycobiome/virome)



Clustering for frequent exacerbator phenotype

# Multi-biome and clinical outcome

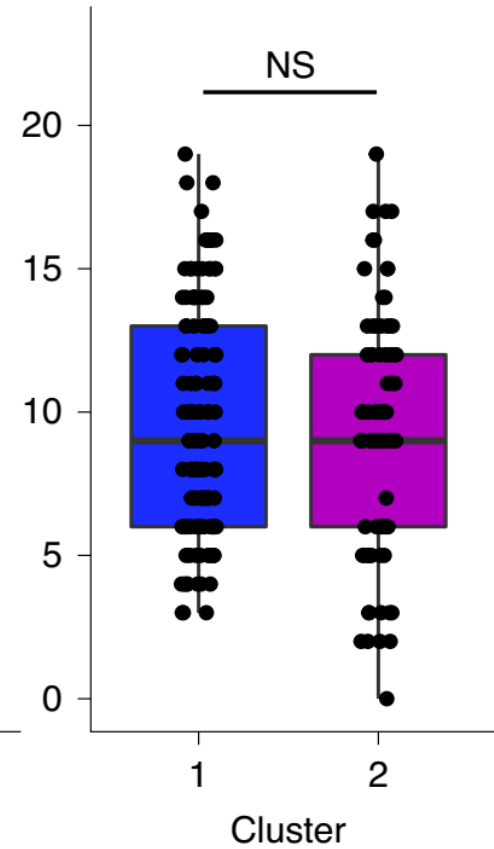
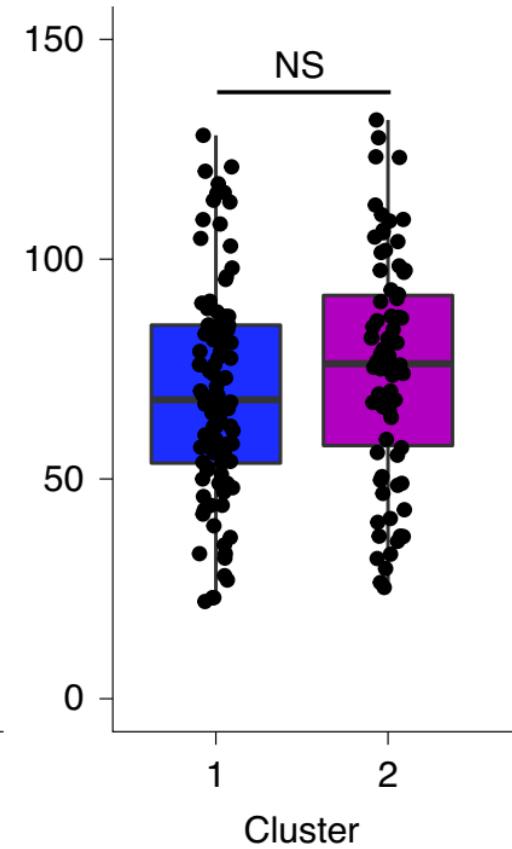
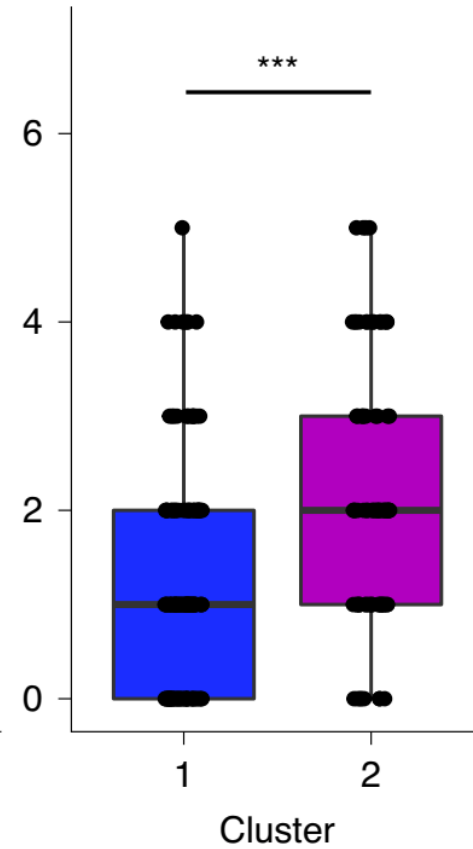
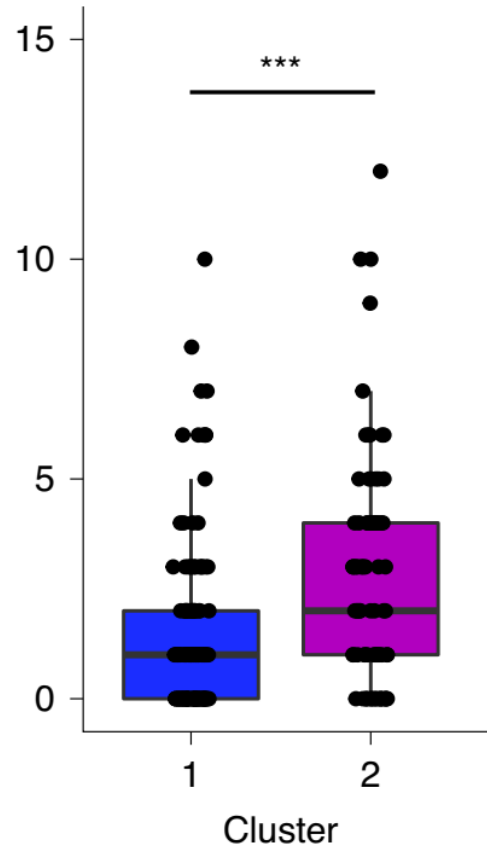
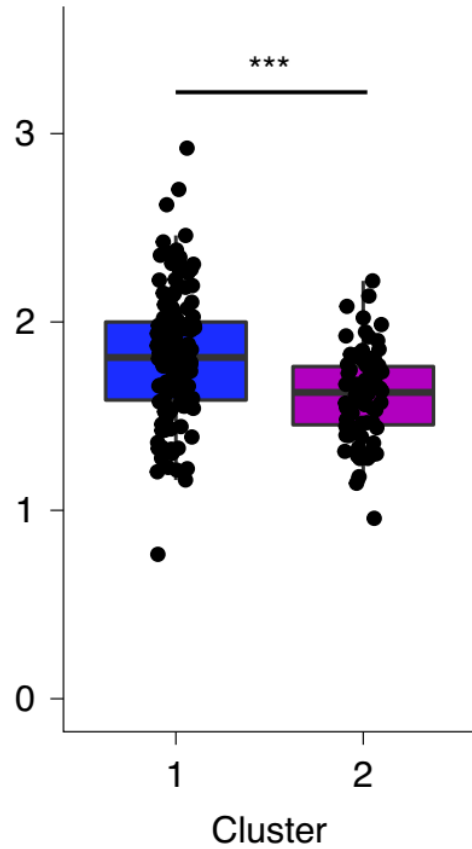
Shannon diversity index

Exacerbations

mMRC score

FEV1(% predicted)

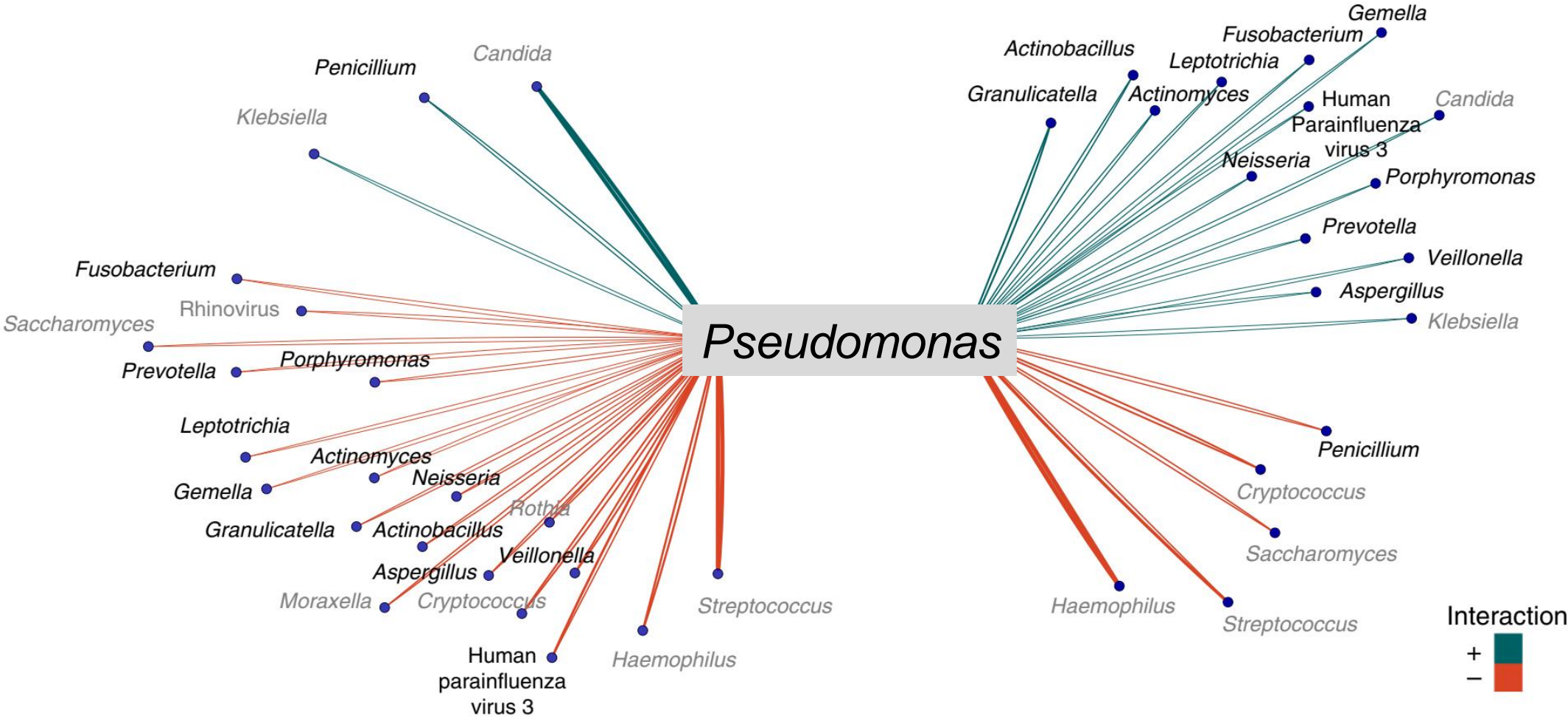
BSI



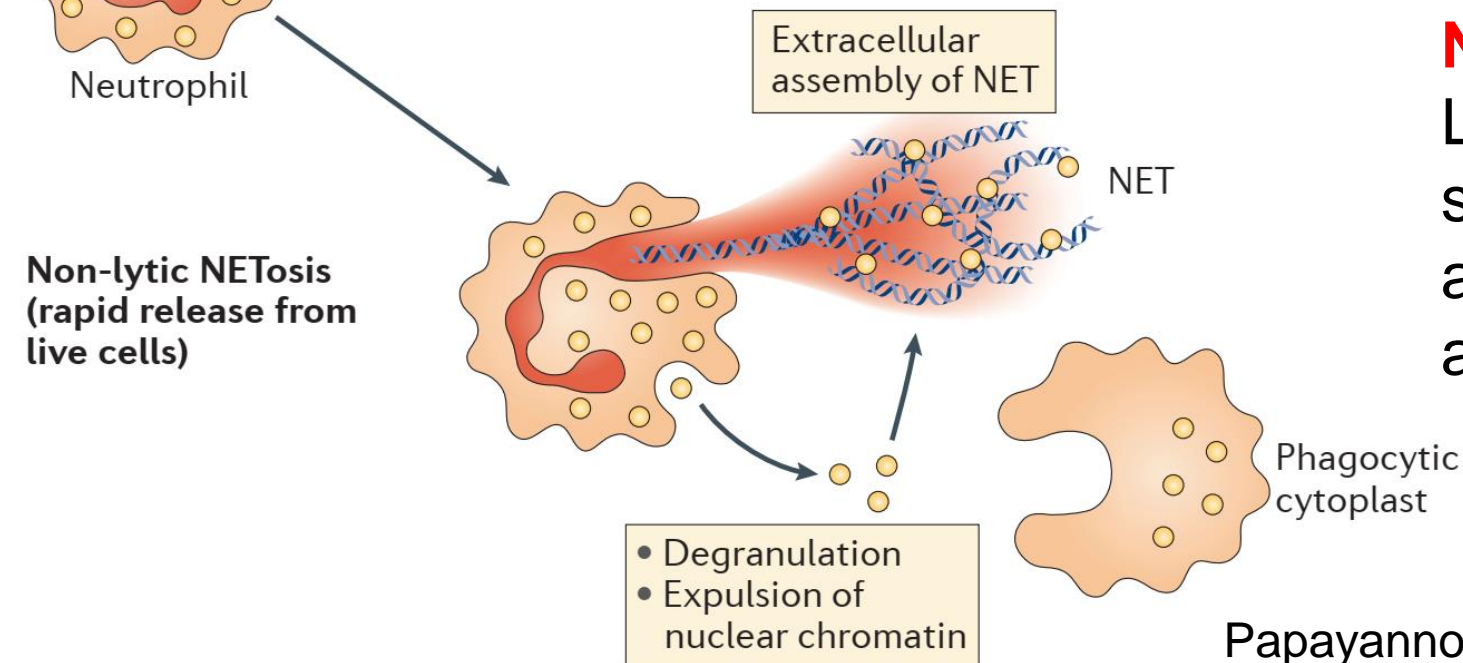
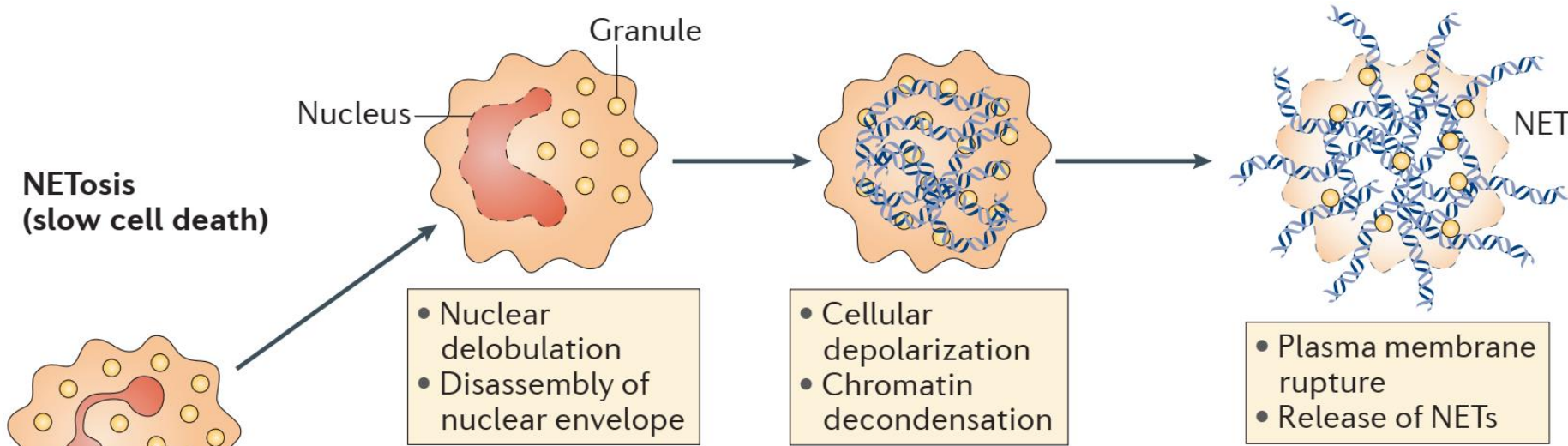
# Pseudomonas-interaction network in exacerbation clustering

Low exacerbation frequency

High exacerbation frequency



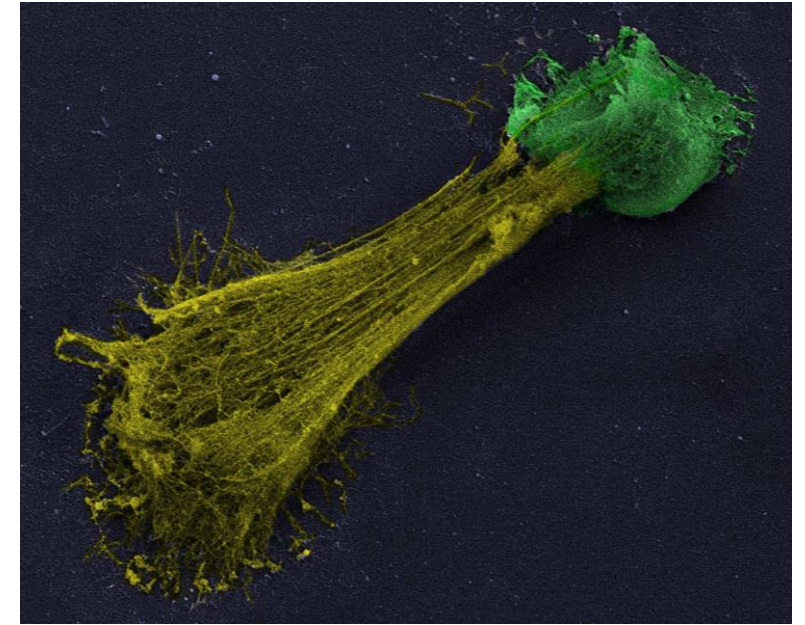
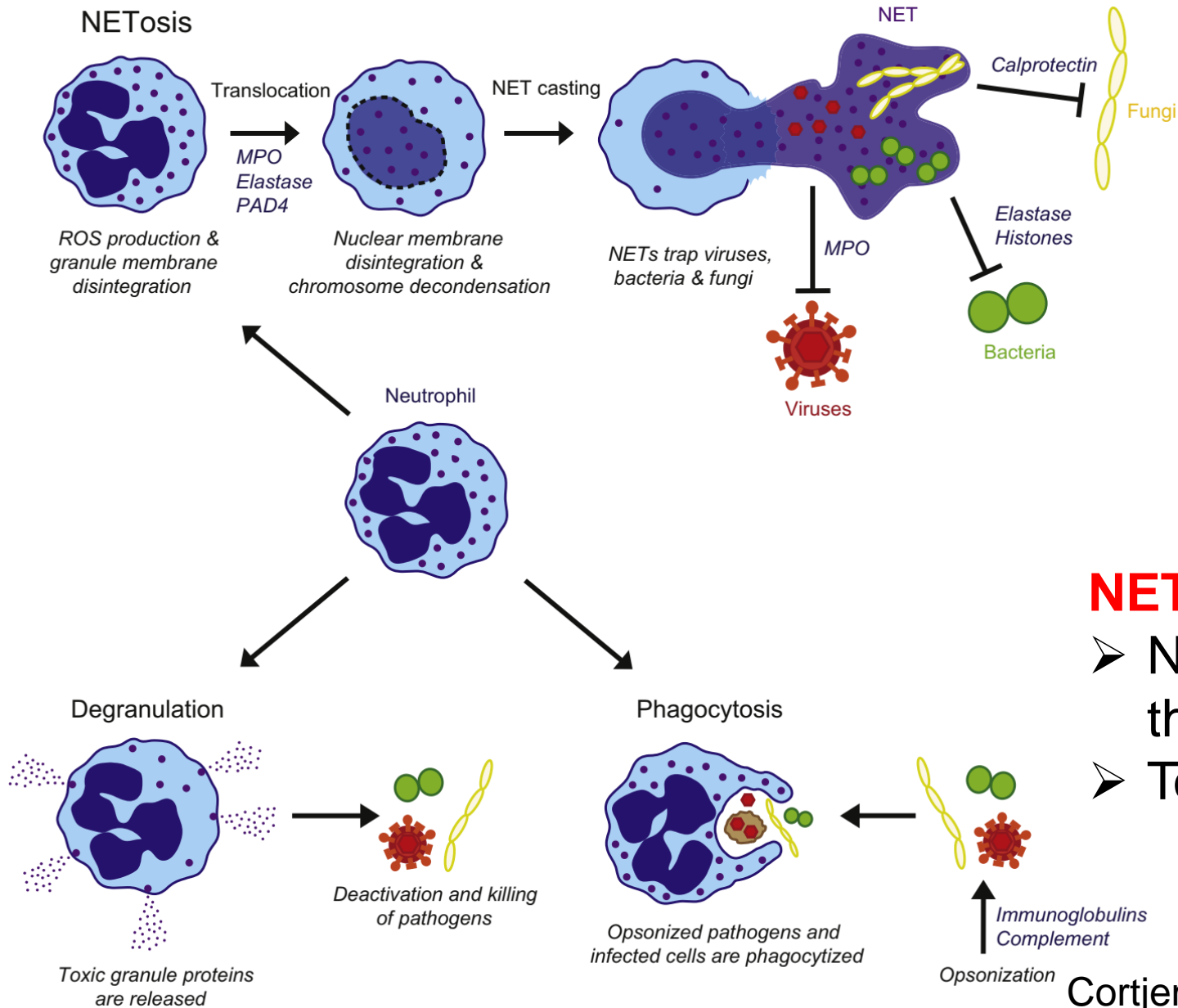
# Neutrophil extracellular traps in immunity



## NET formation pathways

Large, extracellular, web-like structures composed of cytosolic and granule proteins, assembled on a scaffold of decondensed chromatin

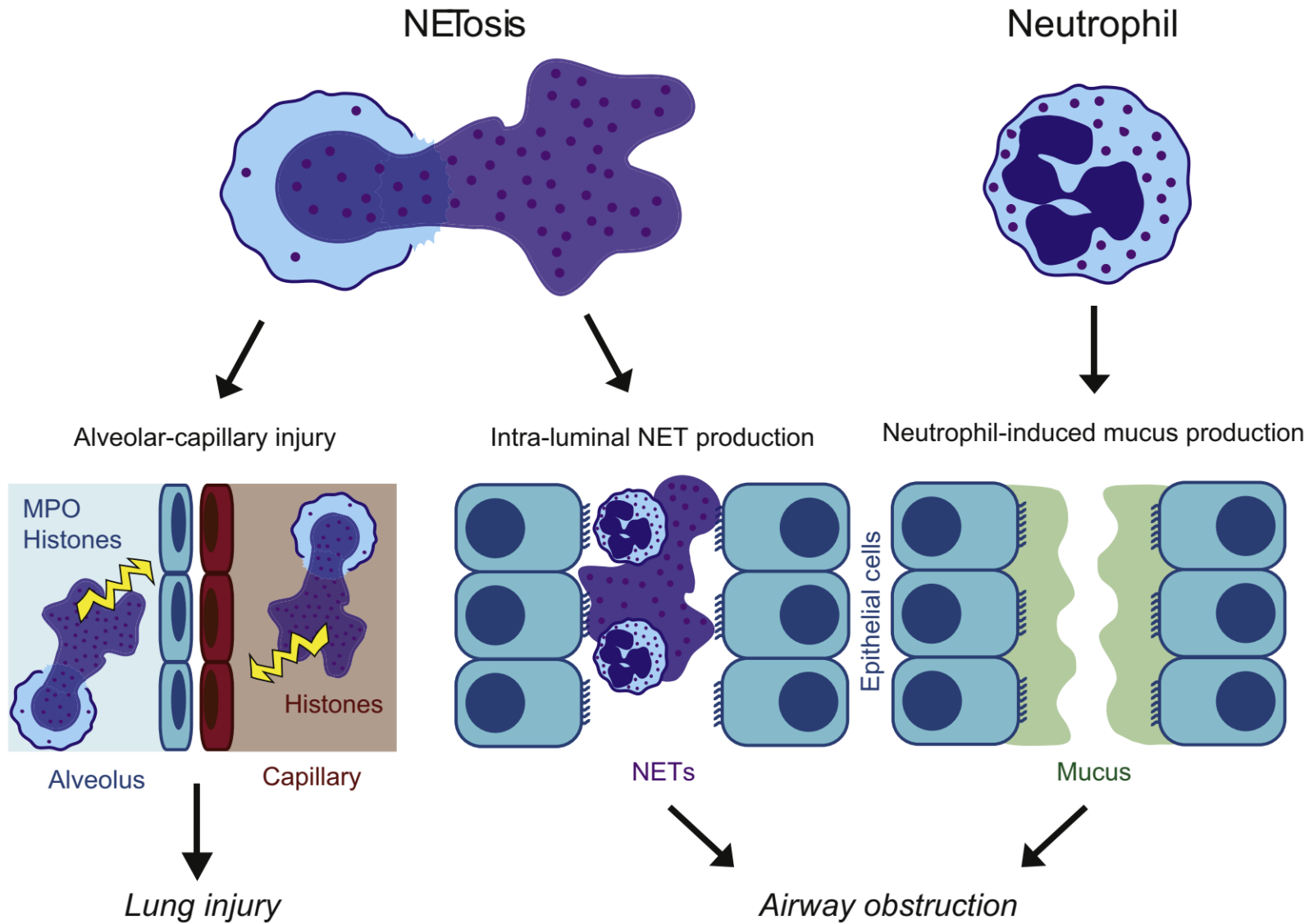
# NET in respiratory disease



## NETosis

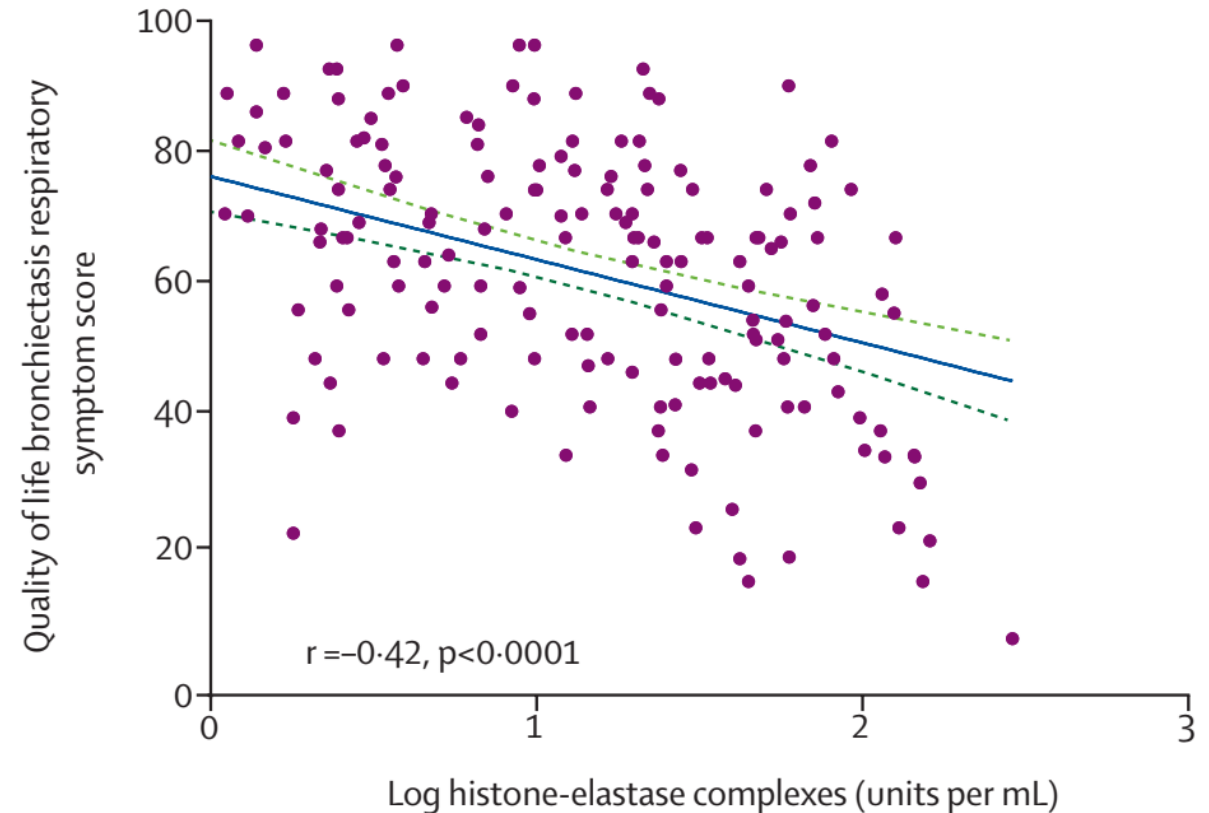
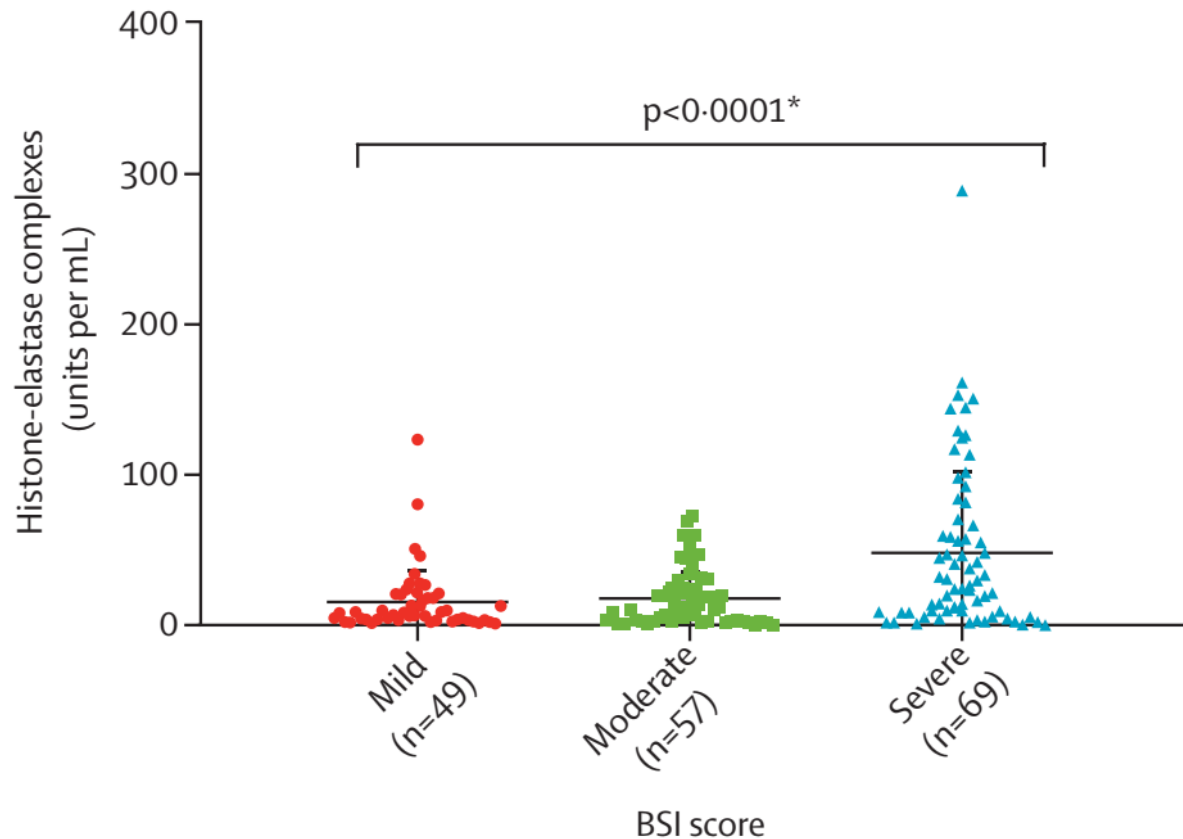
- NET release occurring primarily through a cell death process
- To trap and/or neutralize pathogens

# Potential immunopathological roles of NETs



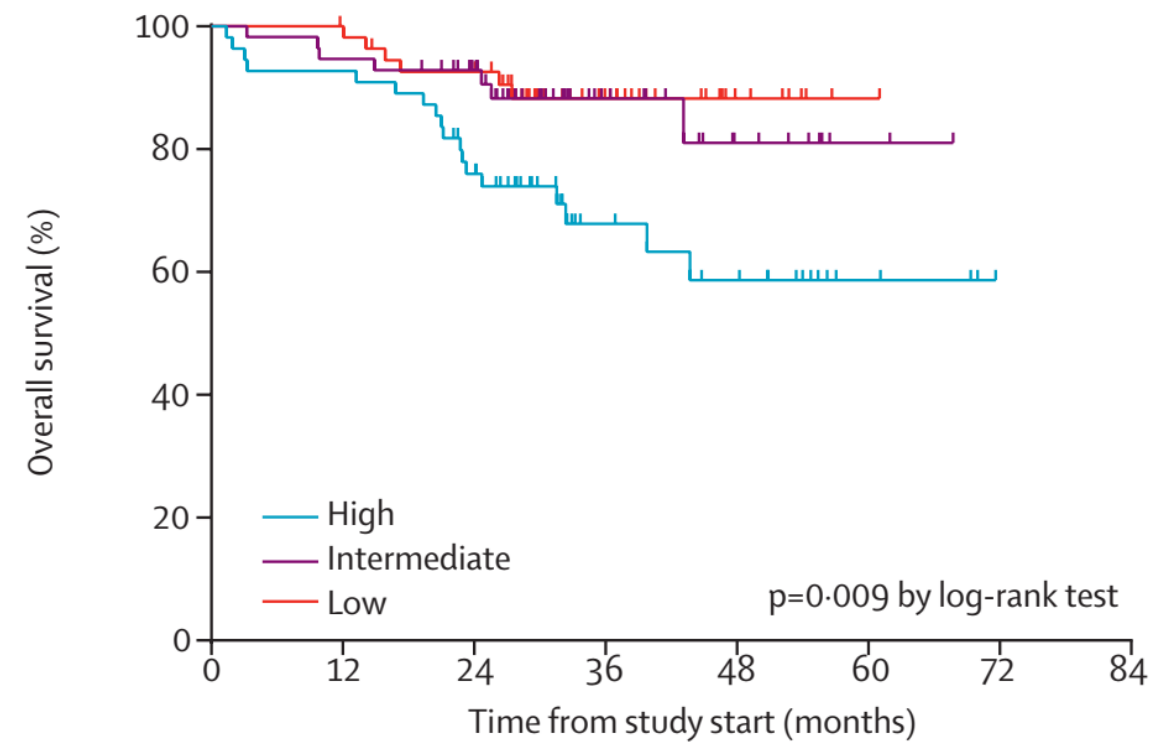
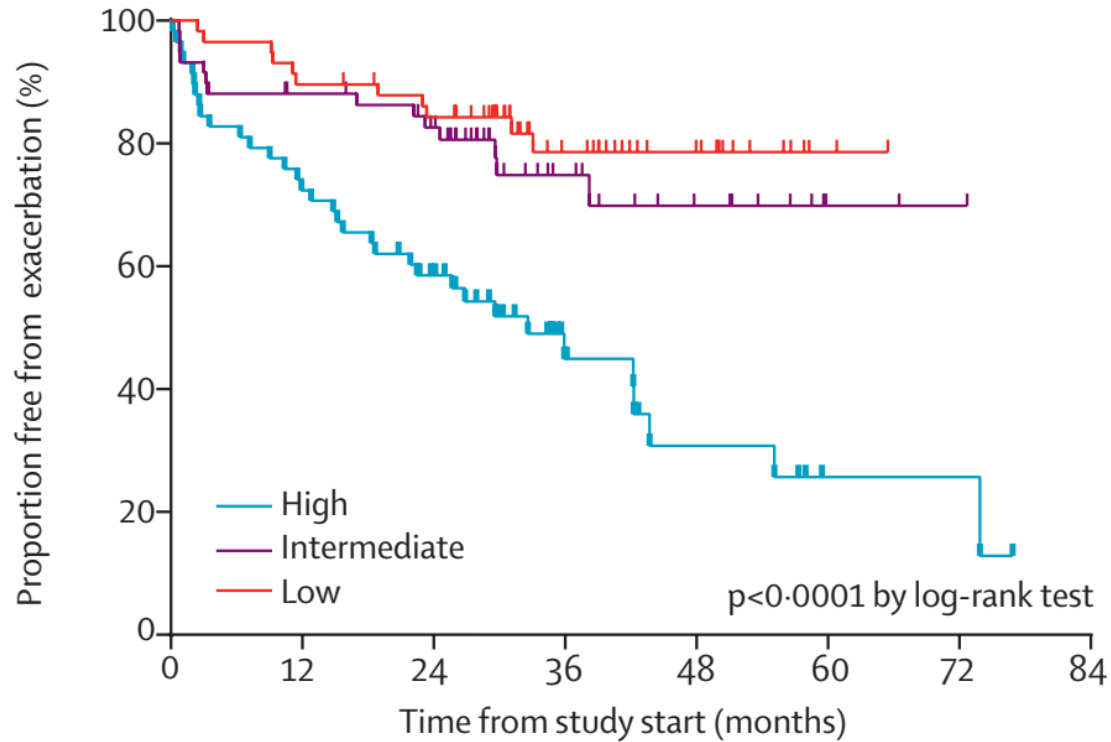
# NET and bronchiectasis severity

- Patients with clinically stable, CT-confirmed idiopathic or post-infective bronchiectasis, without antibiotic treatment for 4 weeks before enrollment
- Sputum samples for proteomic analysis
  - ✓ NET quantification with immunoassay for histone-elastase complexes



# NET and longitudinal outcome

- Association with severe bronchiectasis, symptoms, frequent exacerbations, and future outcomes



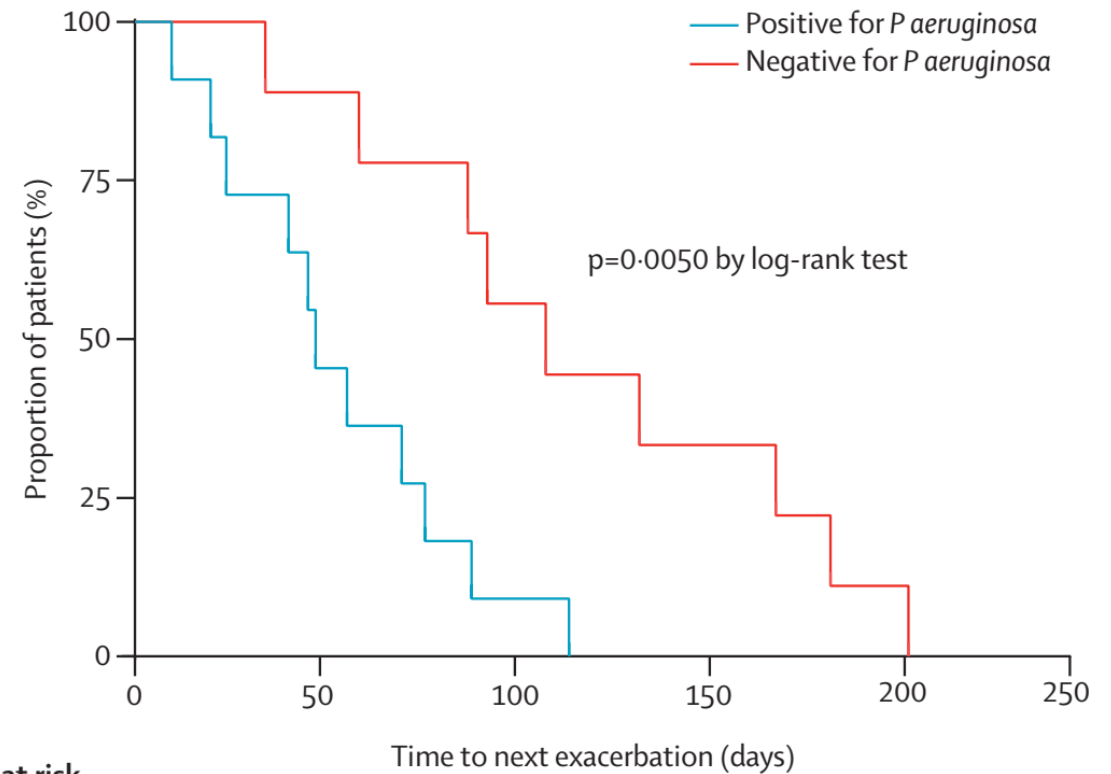
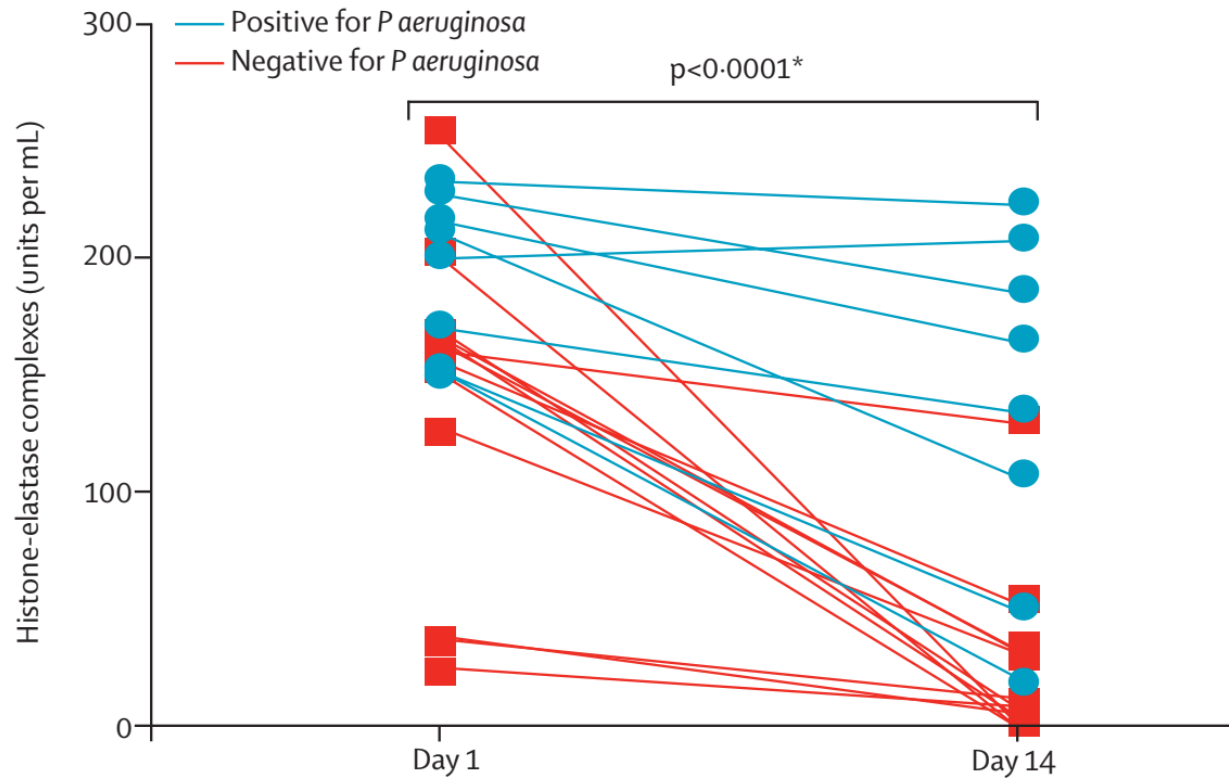
**Number at risk  
(number censored)**

High	58 (0)	43 (0)	31 (4)	12 (18)	7 (20)	3 (23)	3 (23)	0 (24)
Intermediate	59 (0)	50 (3)	44 (6)	18 (29)	10 (36)	3 (43)	2 (44)	0 (45)
Low	58 (0)	53 (0)	48 (2)	25 (23)	14 (34)	3 (45)	1 (47)	0 (47)

High	58 (0)	55 (0)	48 (24)	19 (26)	14 (34)	7 (37)	4 (40)	0 (40)
Intermediate	59 (0)	57 (0)	53 (4)	21 (33)	12 (41)	4 (49)	2 (51)	0 (52)
Low	58 (0)	58 (0)	53 (2)	27 (26)	14 (39)	3 (50)	1 (52)	0 (52)

# NET change and antibiotic response

- Antibiotic treatment reduces NETs
- Patients with *P aeruginosa* infection: higher baseline NETs and less of a reduction in NETs, which correlates with a worse clinical outcome

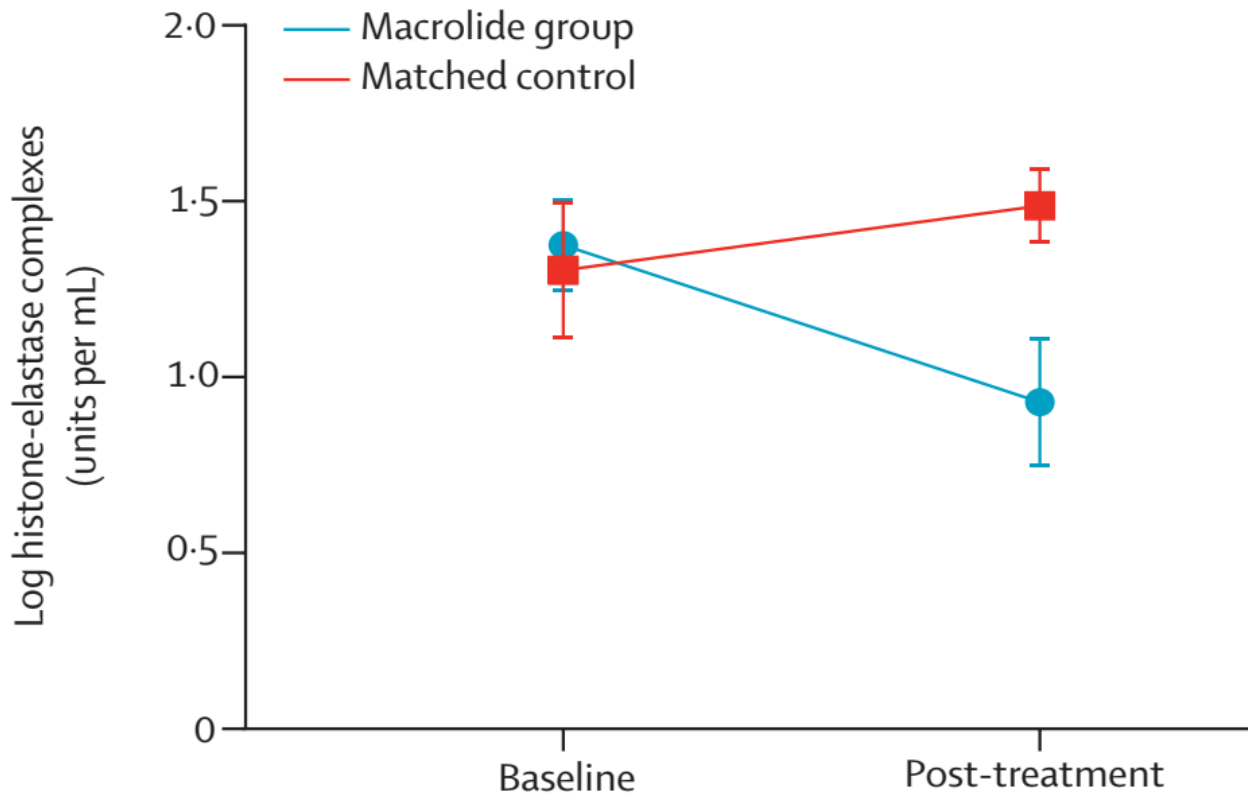


	0	50	100	150	200	250
Positive for <i>P aeruginosa</i>	11	6	2	0	0	0
Negative for <i>P aeruginosa</i>	9	9	6	4	0	0

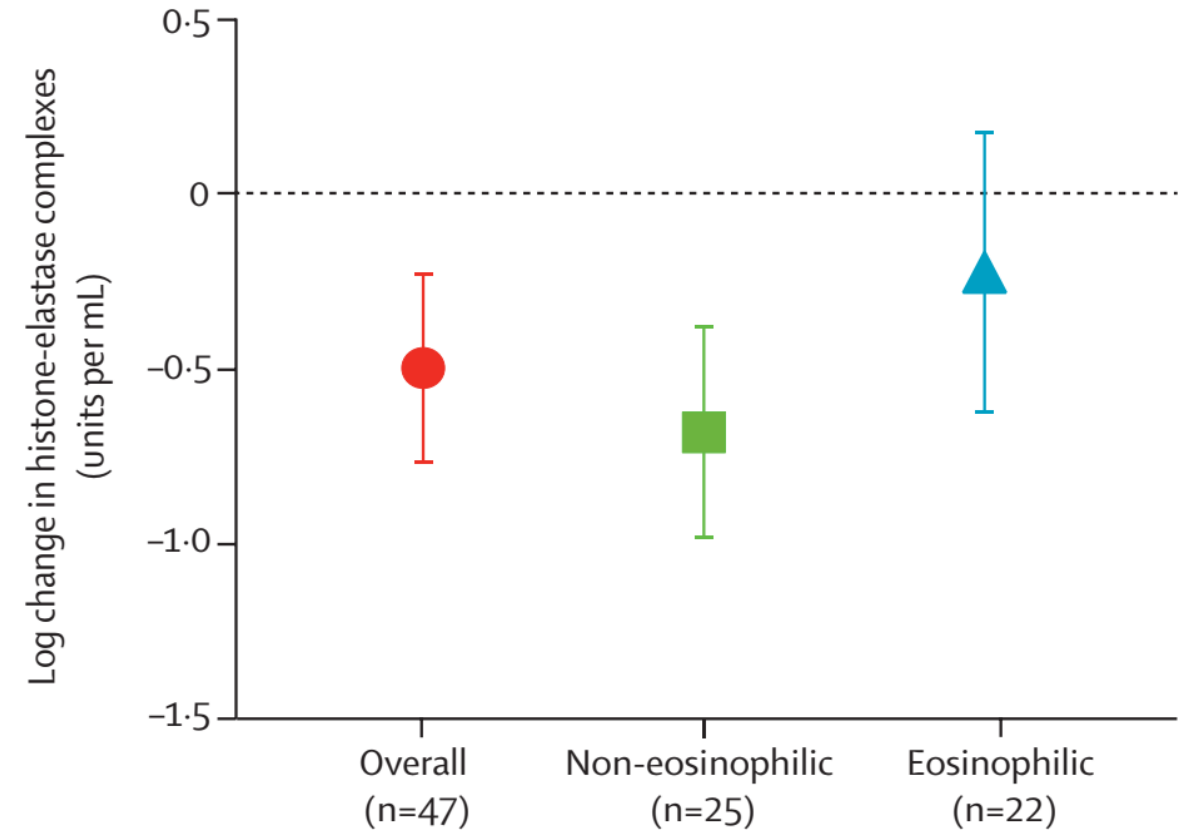
# NET change according to macrolide therapy

- Changes in NET concentrations in response to long-term macrolide therapy in bronchiectasis and asthma

Bronchiectasis group



Asthma group



# Exacerbation as a prognostic factor

- Key events in the natural history of bronchiectasis
- **Exacerbation**
  - ✓ Requirement for antibiotics in the presence of one or more symptoms of increasing **cough**, increasing **sputum volume**, worsening **sputum purulence**, worsening **dyspnea**, increased **fatigue/malaise**, **fever**, and **hemoptysis**
- **Severe exacerbation**
  - ✓ Unscheduled hospitalizations or emergency department visits for exacerbations or complications

# Exacerbation as a prognostic factor

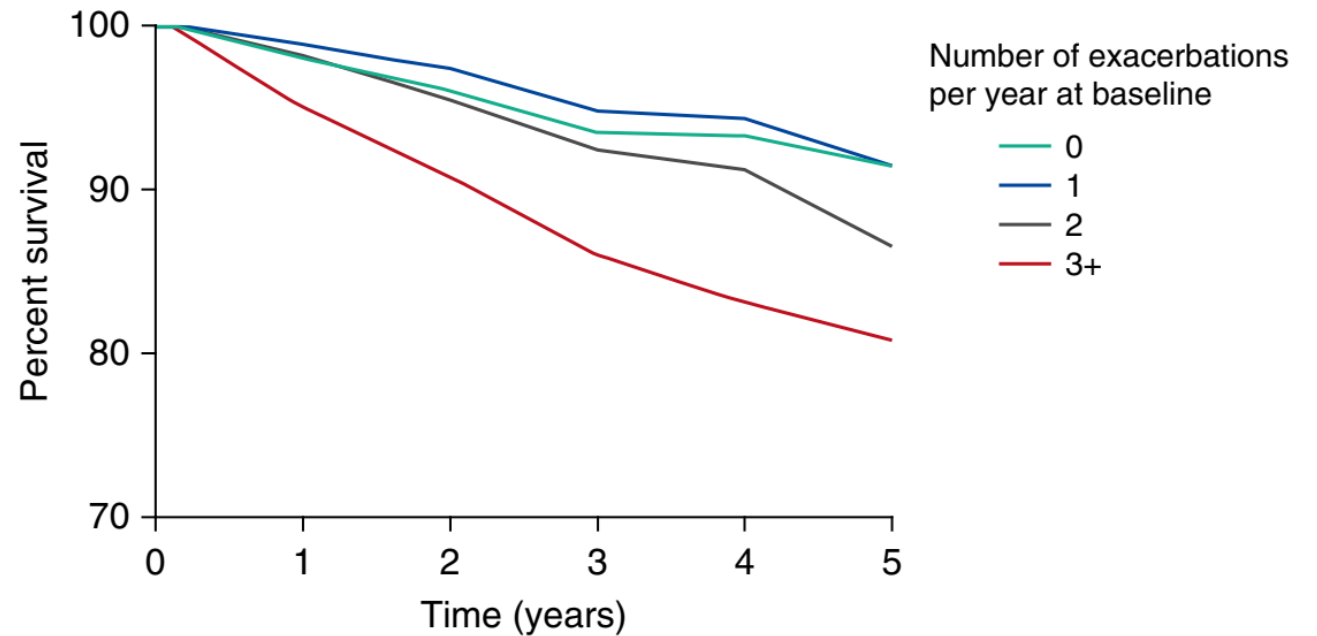
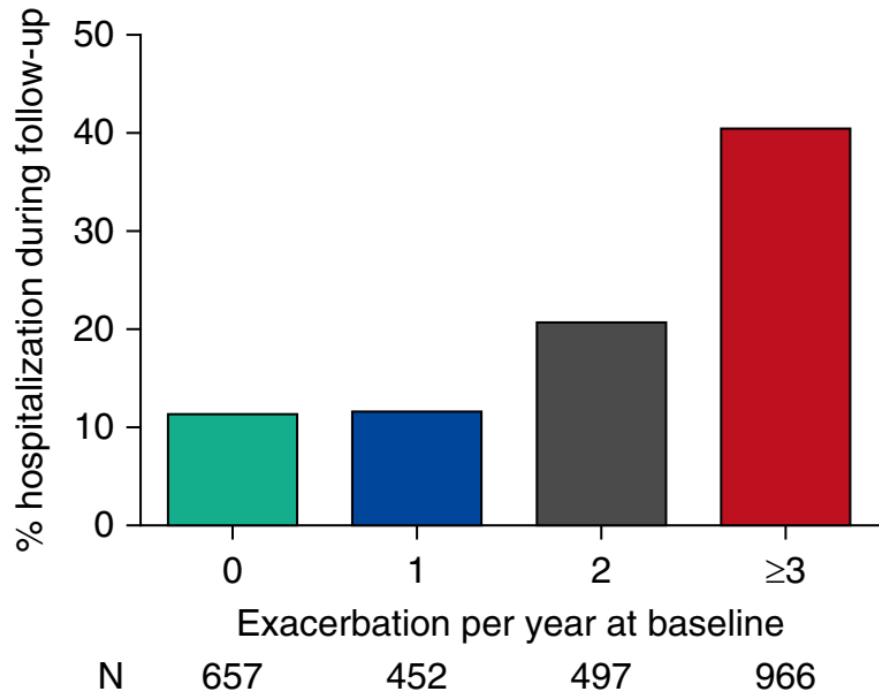
➤ 2,572 bronchiectasis patients in Europe and Israel, up to 5 years of follow-up

## Incident rate ratios for exacerbation frequency during follow-up

	aIRR (95% CI)	P-value
<b>0 Past history of exacerbations</b>	<b>1.0 (ref)</b>	
<b>1 Past history of exacerbations</b>	<b>1.81 (1.54-2.12)</b>	<b>&lt;0.001</b>
<b>2 Past history of exacerbations</b>	<b>3.07 (2.62-3.60)</b>	<b>&lt;0.001</b>
<b>3 Past history of exacerbations</b>	<b>5.18 (4.51-5.95)</b>	<b>&lt;0.001</b>
FEV1 % predicted (per 10%)	0.96 (0.94-0.98)	0.001
<i>Haemophilus influenzae</i>	1.13 (1.01-1.28)	0.040
<i>Pseudomonas aeruginosa</i>	1.20 (1.04-1.40)	0.010
COPD	1.43 (1.22-1.67)	<0.001

# Exacerbation as a prognostic factor

Follow-up hospitalization rates and annual survival in groups on the basis of baseline exacerbations



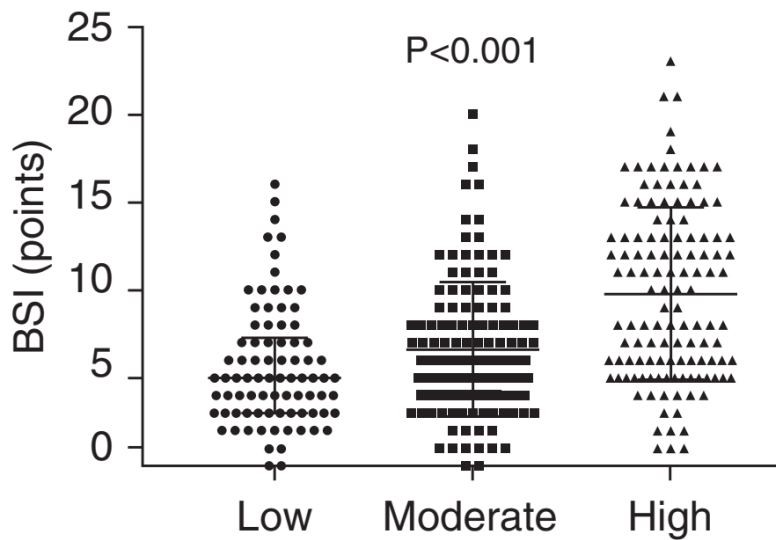
## Numbers at risk

0	657	654	600	554	522	153
1	452	444	402	381	351	134
2	497	490	437	407	376	196
3 or more	966	958	836	771	694	365

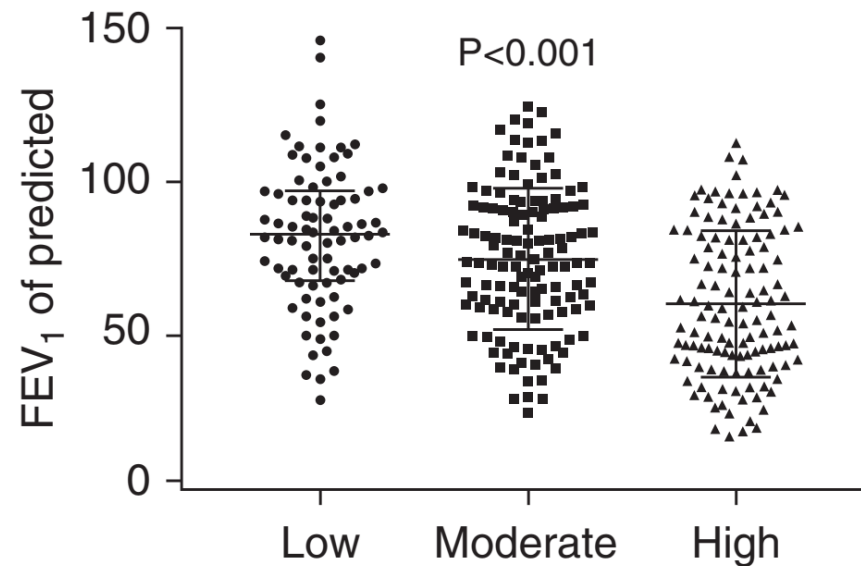
# Symptom burden and disease severity

- 333 bronchiectasis, observational cohort in Scotland
- Measurement of daily symptom burden using SGRQ symptom score
  - ✓ Low, <40 points; Moderate, 40-70 points; High, >70 points

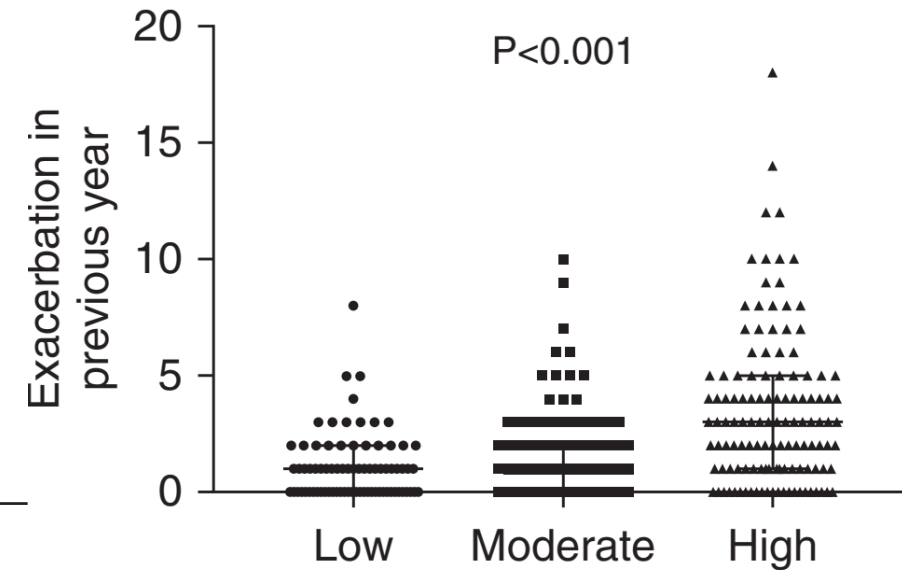
Bronchiectasis severity index



FEV<sub>1</sub> % predicted



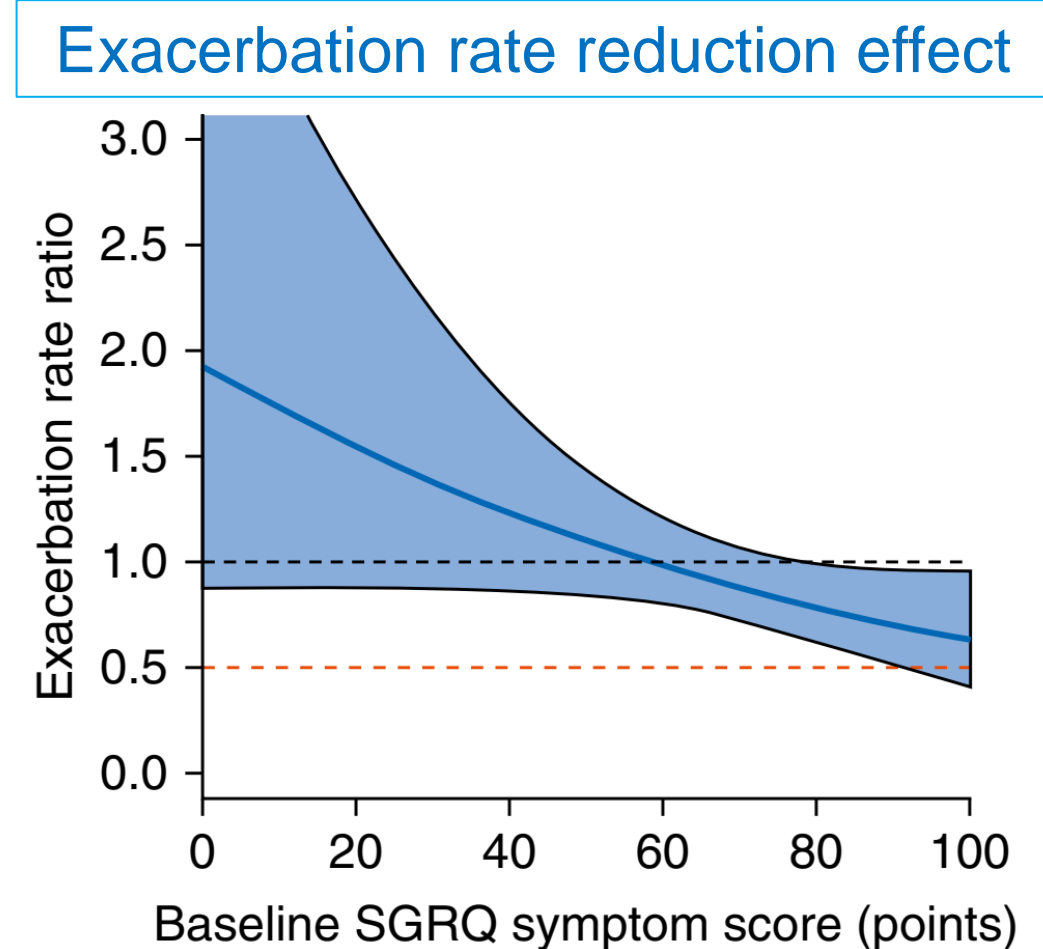
Exacerbation in previous year



# Symptom burden and exacerbation risk

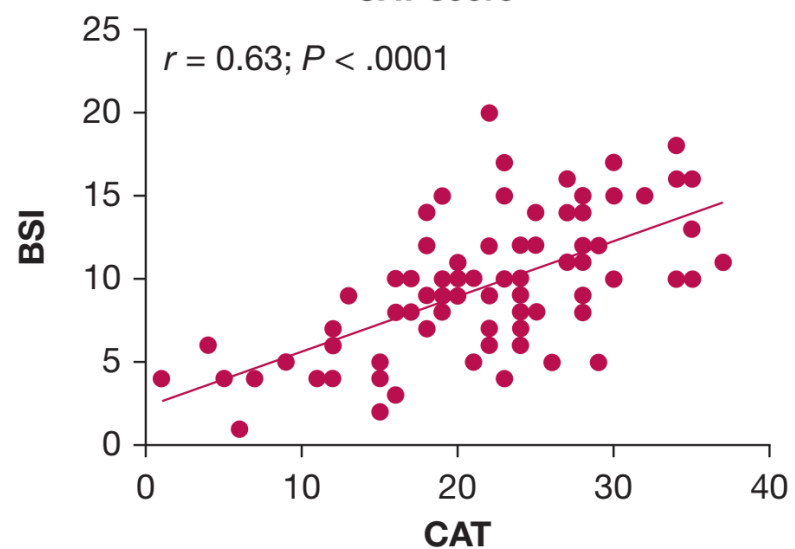
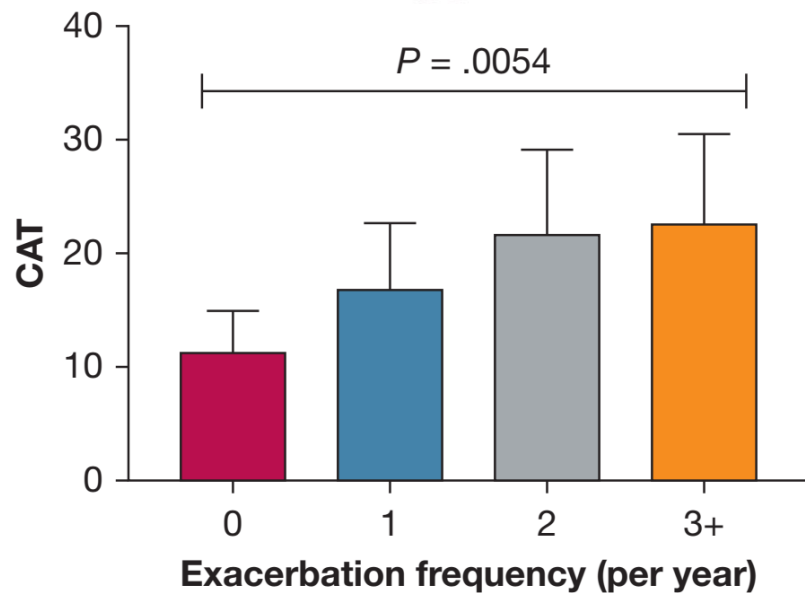
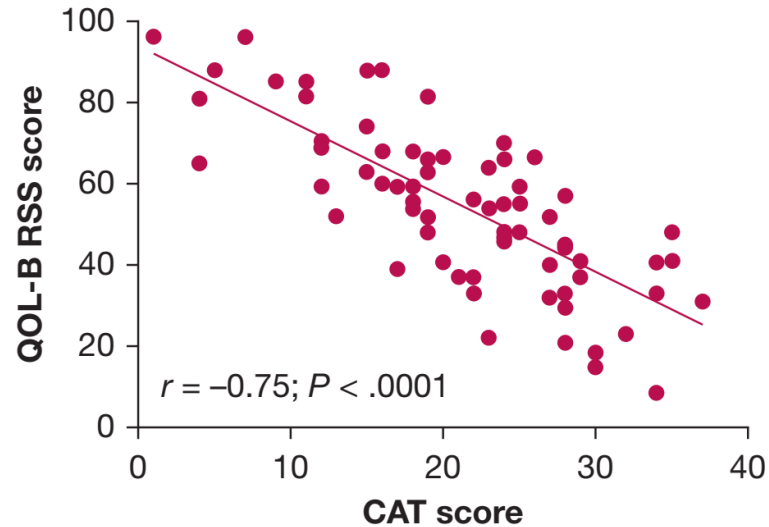
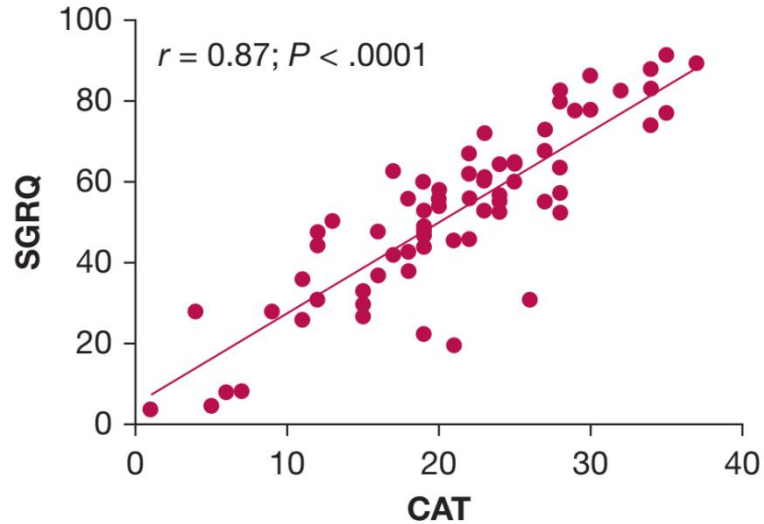
- *Post hoc* analysis of RCT of inhaled dry power mannitol treatment
  - ✓ Inhaled mannitol 400mg bid vs. low-dose control, 1:1, 52wk

SGRQ Symptom Score	Exacerbation		Time to first exacerbation	
	aRR (95% CI)	P-value	aRR (95% CI)	P-value
Uncategorized	1.09 (1.02-1.17)	0.017	1.09 (1.02-1.15)	0.009
Categorized				
Low	1 (ref)		1 (ref)	
Moderate	1.50 (0.92-2.45)	0.10	1.73 (1.07-2.80)	0.03
High	1.75 (1.07-2.86)	0.03	1.78 (1.09-2.90)	0.02

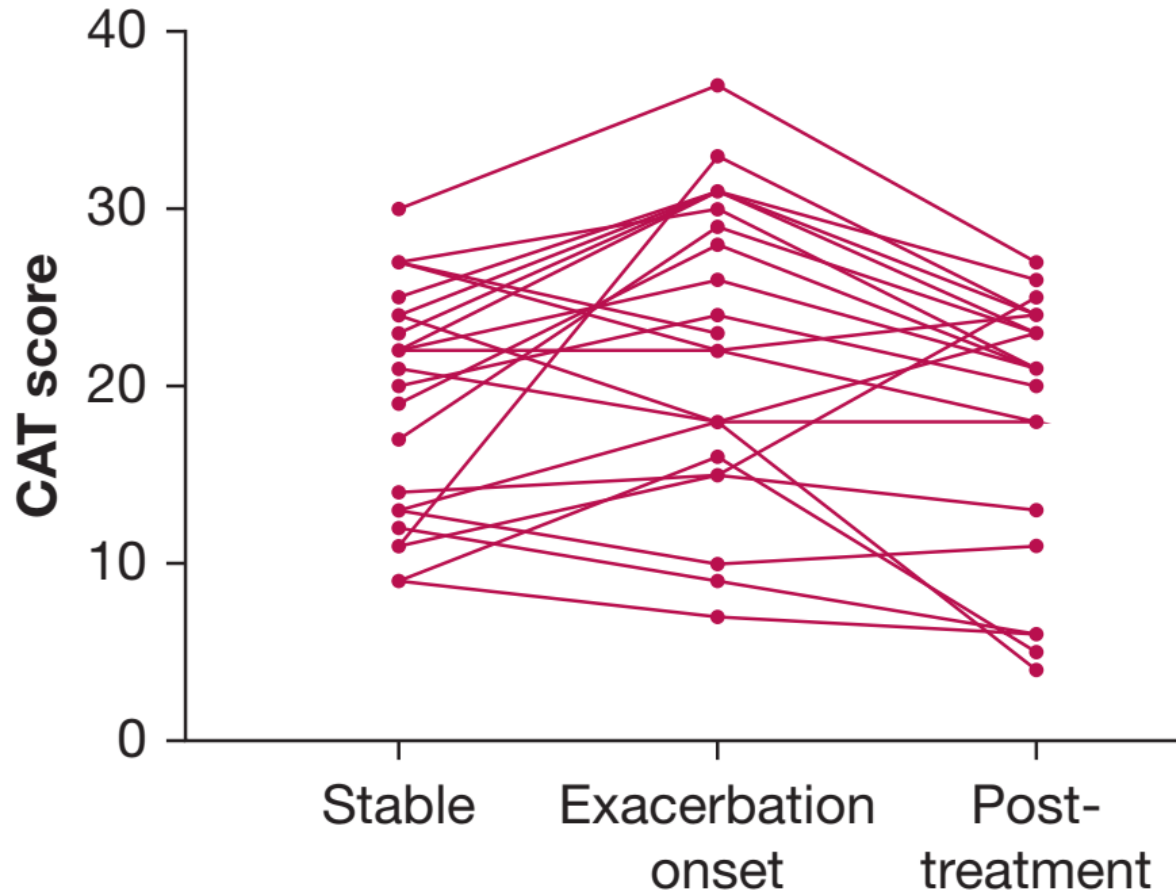


# COPD assessment test as an outcome measure

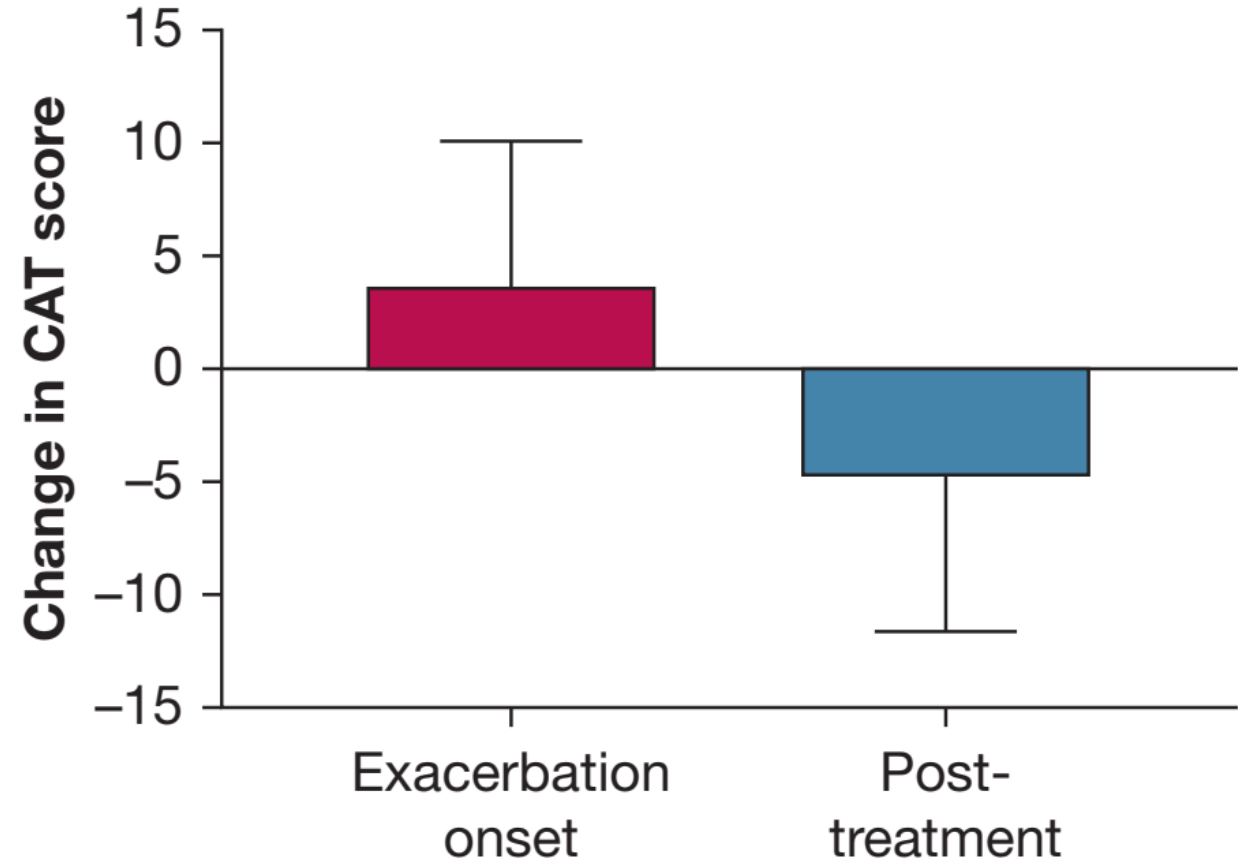
➤ Bronchiectasis patient without any history of COPD in TRIBE study



# CAT according to exacerbation



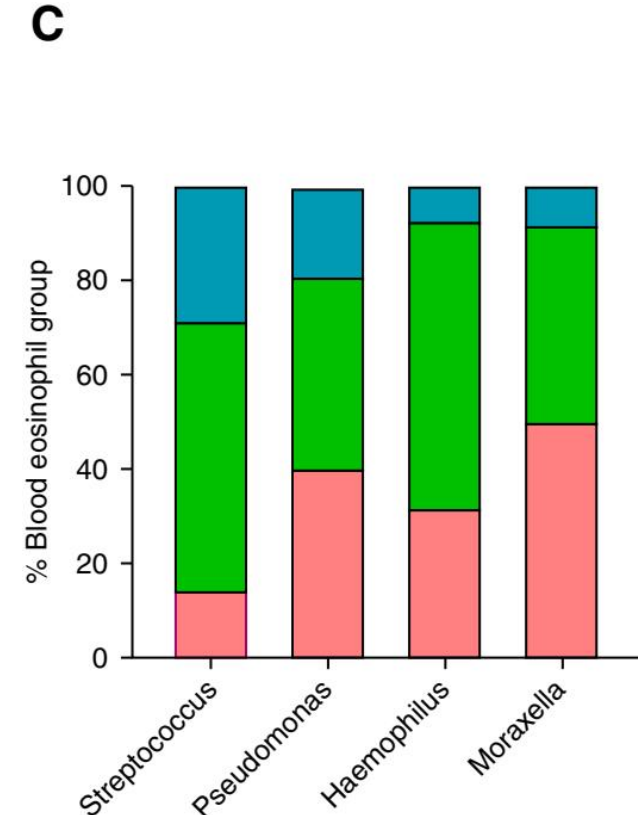
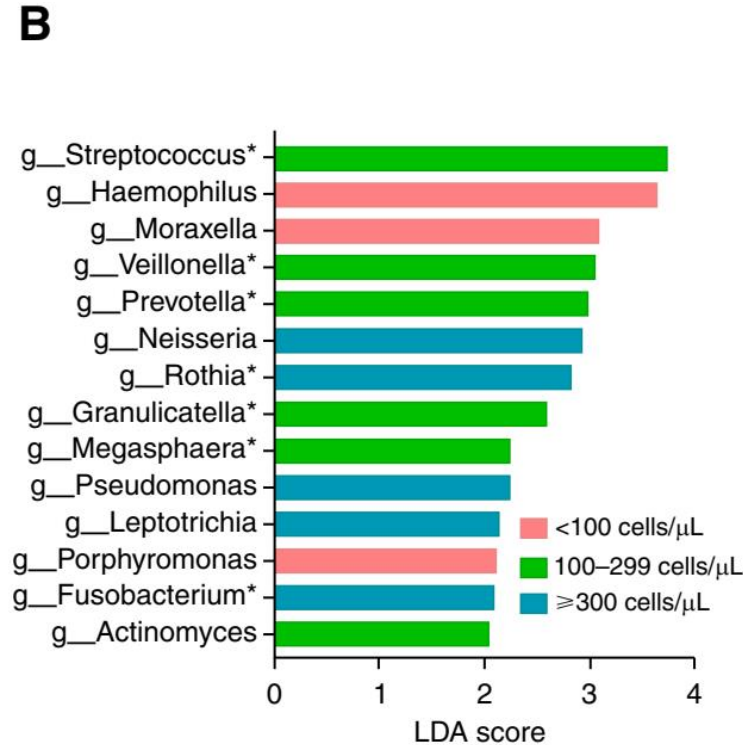
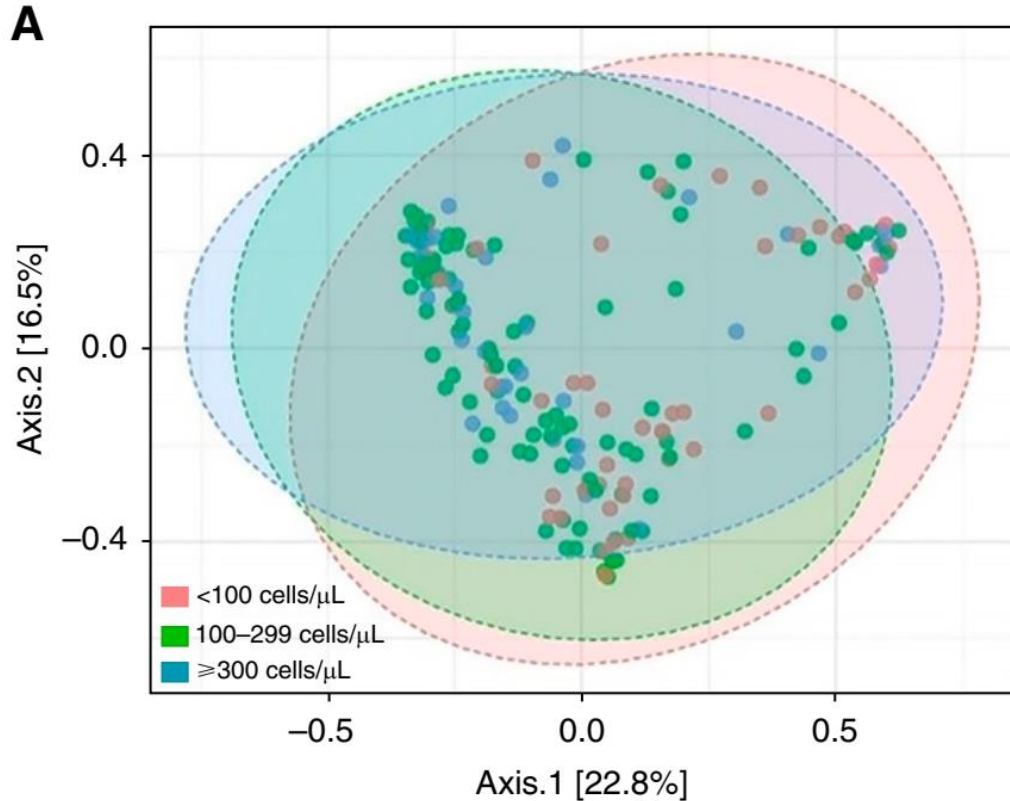
Proposed minimum clinically important difference (MCID): 4 points



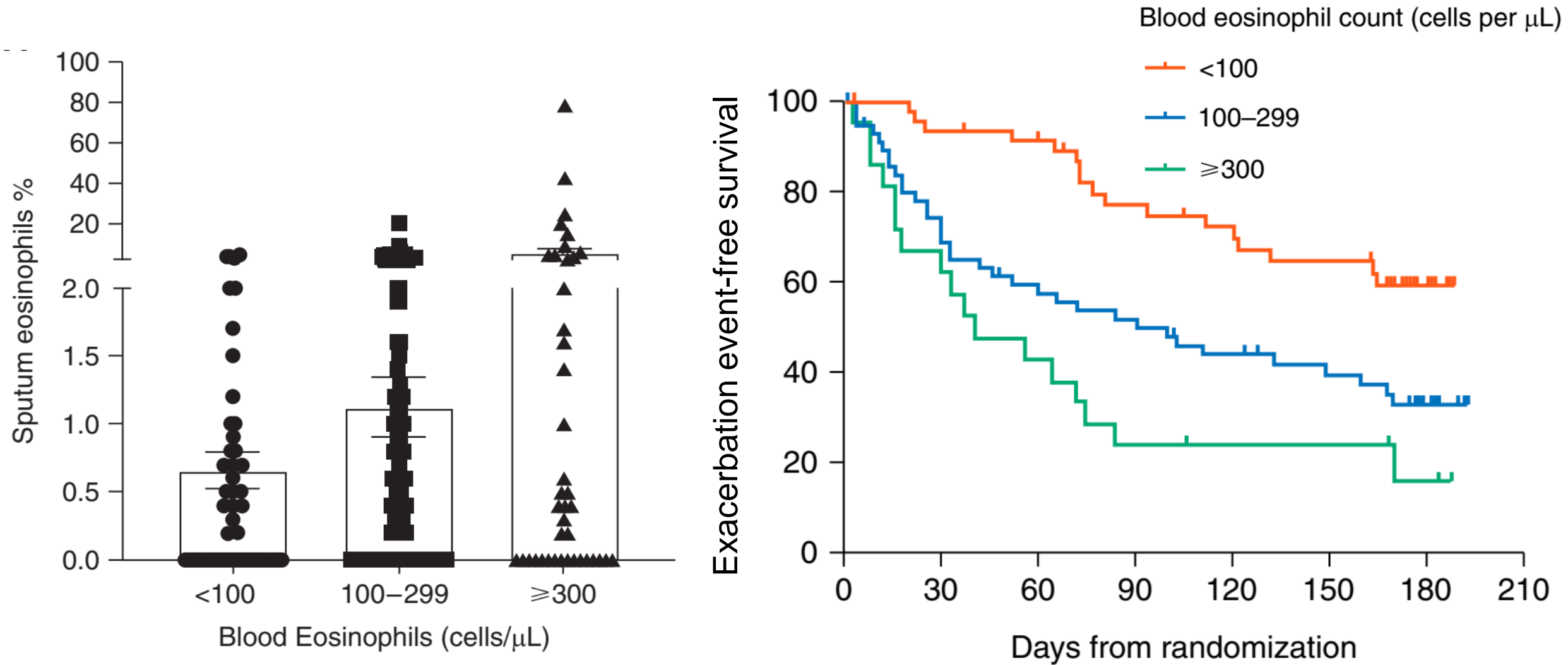
# Eosinophilic phenotype

- 1,007 bronchiectasis patients without asthma, European multicohort
- Patients with blood eosinophil counts  $\geq 300$  cells/ $\mu\text{L}$ : 22.6%

## Relationship between the sputum microbiome and blood eosinophil counts



# Eosinophilic phenotype and exacerbation



# Blood eosinophils and ICS response

- 86 clinically stable bronchiectasis patients without asthma, CF, and ABPA
- Baseline blood eosinophils: low (<3%) vs. high (≥3%)
- ICS (fluticasone propionate) 250µg/500µg 2 times/day vs. no treatment, 6 months

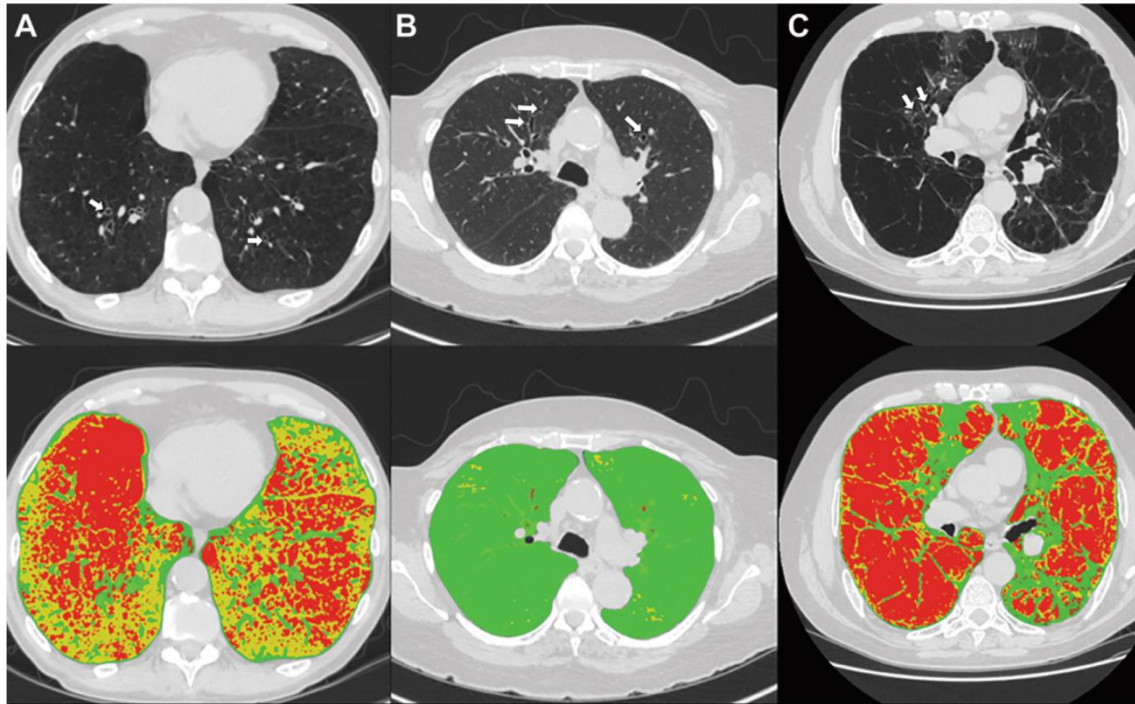
	Low eosinophil group			High eosinophil group		
	FP+ (n=13)	FP- (n=10)	P-value	FP+ (n=44)	FP- (n=19)	P-value
Change SGRQ ≥4 points	4 (33.3)	1 (11.1)	0.34	<b>21 (47.7)</b>	0 (0.0)	<0.001
SGRQ total change	-2.5 (-5.3 to +2.7)	-0.7 (-2.0 to +1.6)	0.54	<b>-3.1 (-8.9 to +2.8)</b>	+1.6 (0.4 to +4.2)	0.003
FEV1 at 6 months %	68.4±19.6	58.5±15.3	0.33	61.4±20.3	72.6±18.5	0.21
MRC (3-4) at 3 months	3 (23.1)	1 (10.0)	0.60	2 (4.6)	4 (21.1)	0.06
Exacerbation at 6 months	7 (70.0)	3 (37.5)	0.34	12 (35.3)	6 (40.0)	0.75

# Criteria and definitions for the radiological and clinical diagnosis of bronchiectasis in adults for use in clinical trials: international consensus recommendations

	Mean scores
Inner airway–artery diameter ratio $\geq 1.5$	3.50/4.00
Outer airway–artery diameter ratio $\geq 1.5$	3.21/4.00
Lack of tapering	3.00/4.00
Visibility of airways in the periphery	2.89/4.00
Inner airway–artery diameter ratio $\geq 1.1$	2.75/4.00
Inner airway–artery diameter ratio $\geq 1.0$	2.71/4.00
Outer airway–artery diameter ratio $\geq 1.1$	2.13/4.00
Outer airway–artery diameter ratio $\geq 1.0$	1.88/4.00

# Small airway disease and emphysema in bronchiectasis

- 737 former and current smokers (387 bronchiectasis) from COPDGene study
- **Parametric response mapping** on paired expiratory and inspiratory CT scans to identify and quantify the extent of SAD and emphysema



Emphysema: red; SAD: yellow  
Normal areas: green

	COPD only	BE without COPD	BE with COPD
PRM <sup>NORMAL</sup> (%)	66.7 (45.5-79.8)	87.5 (82.0-92.9)	58.6 (42.2-73.8)
PRM <sup>SAD</sup> (%)	24.8 (16.0-33.6)	9.7 (6.4-13.8)	27.6 (19.2-34.6)
PRM <sup>EMPHY</sup> (%)	4.1 (1.2-12.6)	0.4 (0.1-1.2)	7.6 (2.6-18.4)
Bronchiectasis severity score		3.0 (2.0-4.0)	3.0 (2.0-4.0)

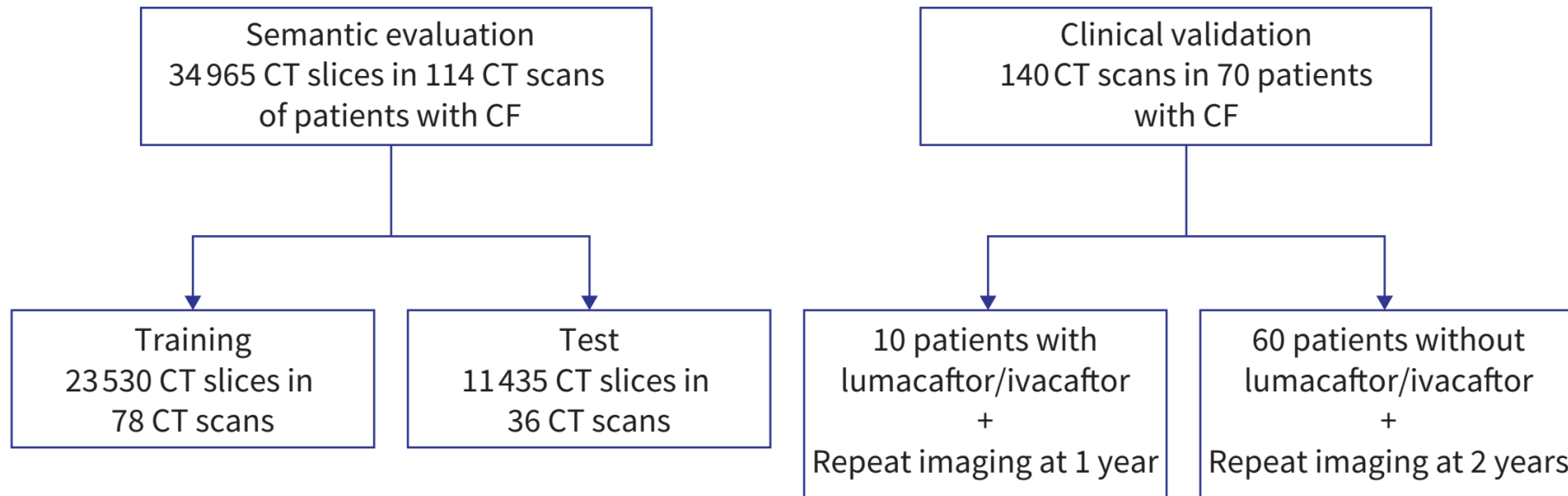
# SAD/emphysema and exacerbation

## SAD and emphysema at PRM with exacerbation in heavy smokers with bronchiectasis and/or COPD

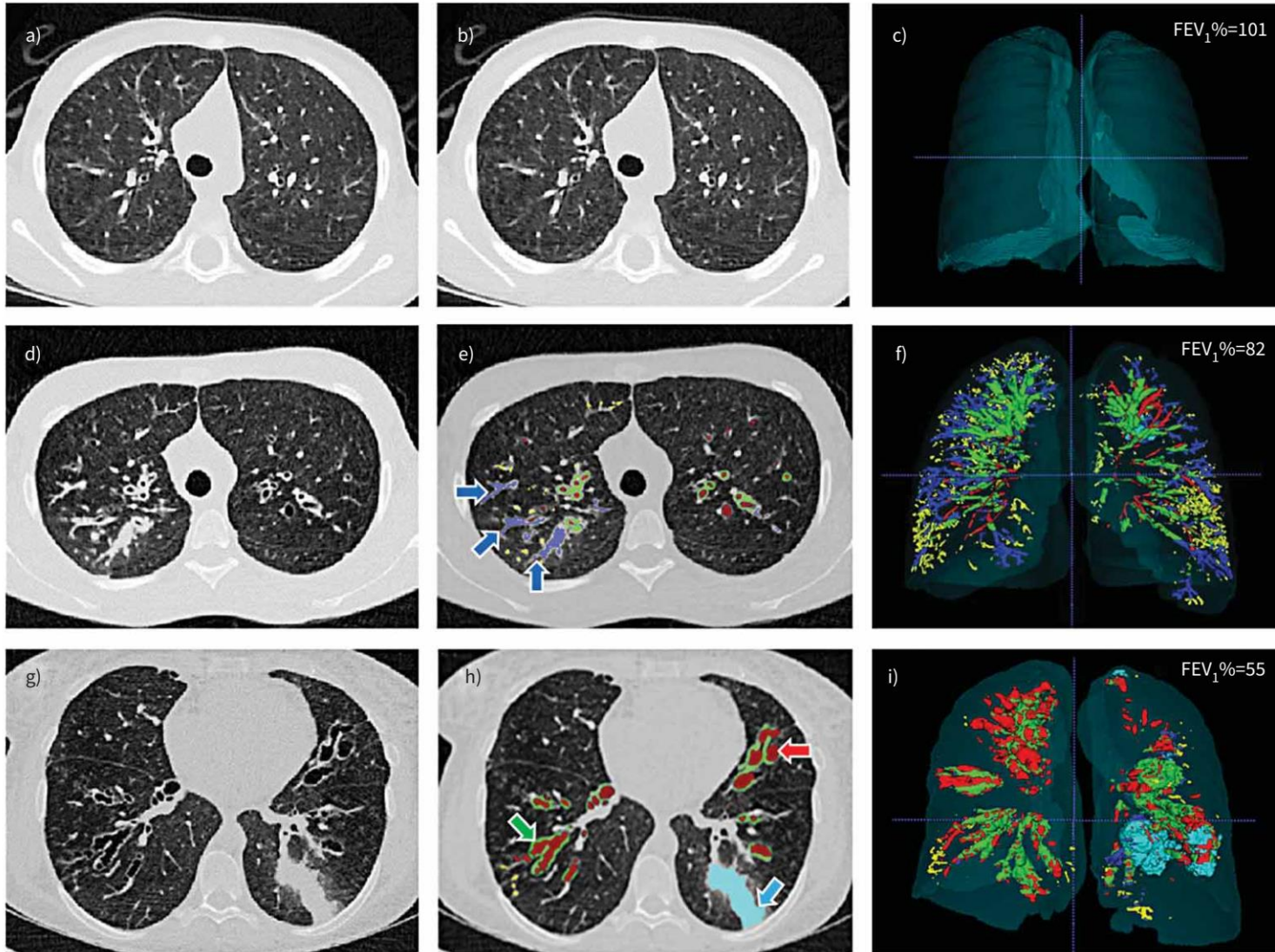
Group and Parameter	Exacerbation		Severe Exacerbation	
	Relative Risk	<i>P</i> Value	Relative Risk	<i>P</i> Value
CT-derived bronchiectasis with and without COPD ( <i>n</i> = 387)				
PRM <sup>SAD</sup> ≥15.6% vs <15.6%	1.89 (1.54, 2.33)	<.001	2.02 (1.40, 2.90)	<.001
PRM <sup>EMPH</sup> ≥5% vs <5%	1.37 (1.13, 1.66)	.001	0.98 (0.69, 1.38)	.89
COPD only ( <i>n</i> = 350)				
PRM <sup>SAD</sup> ≥15.6% vs <15.6%	0.97 (0.82, 1.15)	.72	1.14 (0.82, 1.58)	.45
PRM <sup>EMPH</sup> ≥5% vs <5%	1.61 (1.38, 1.87)	<.001	1.68 (1.27, 2.22)	<.001
CT-derived bronchiectasis with COPD ( <i>n</i> = 197)				
PRM <sup>SAD</sup> ≥15.6% vs <15.6%	1.67 (1.23, 2.27)	.001	1.83 (1.08, 3.11)	.03
PRM <sup>EMPH</sup> ≥5% vs <5%	1.51 (1.20, 1.91)	<.001	0.96 (0.64, 1.44)	.84
CT-derived bronchiectasis without COPD ( <i>n</i> = 190)				
PRM <sup>SAD</sup> ≥15.6% vs <15.6%	1.67 (1.16, 2.41)	.007	2.40 (1.20, 4.83)	.01
PRM <sup>EMPH</sup> ≥5% vs <5%	0.91 (0.27, 3.02)	.87	1.42 (0.17, 11.90)	.75

# Artificial intelligence for quantifying lung changes

- Cystic fibrosis patients aged 4-54 years with/without lumacaftor/ivacaftor, cystic fibrosis transmembrane conductance regulator (CFTR) modulator
- **AI-driven measurements:** algorithm using three 2D convolutional neural networks trained for the semantic labelling of bronchiectasis, peribronchial thickening, bronchial mucus, bronchiolar mucus and collapse/consolidation



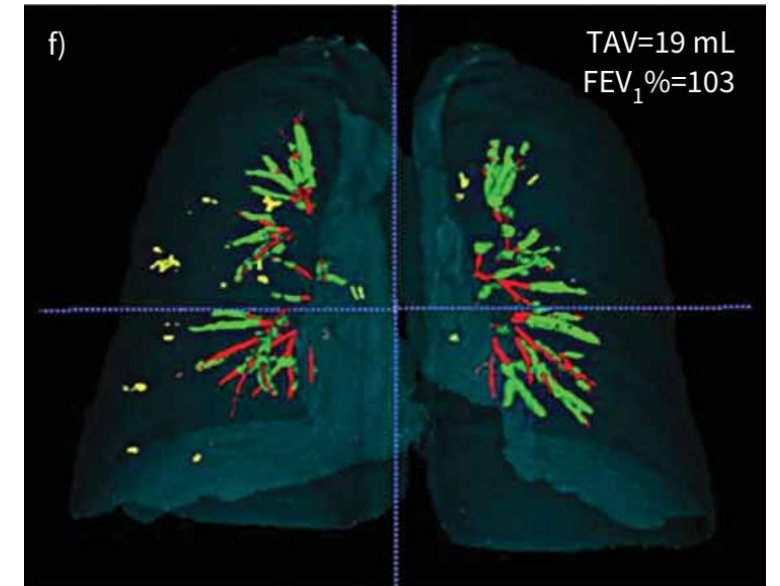
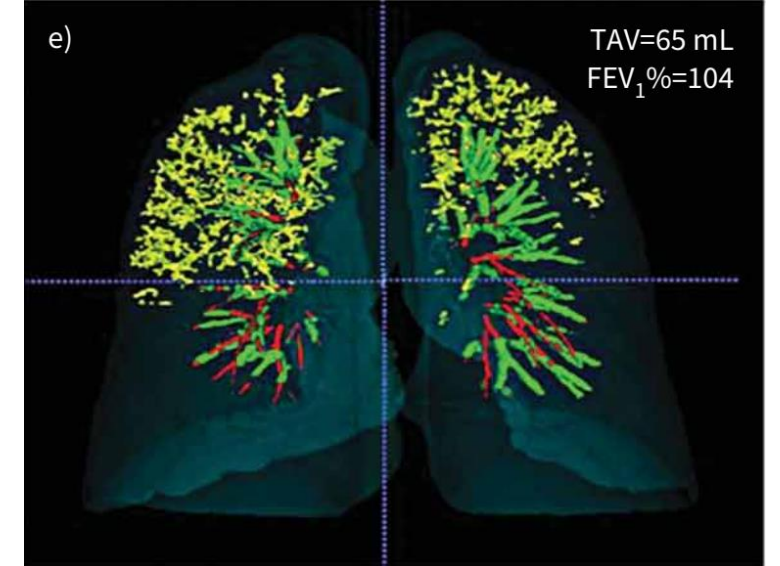
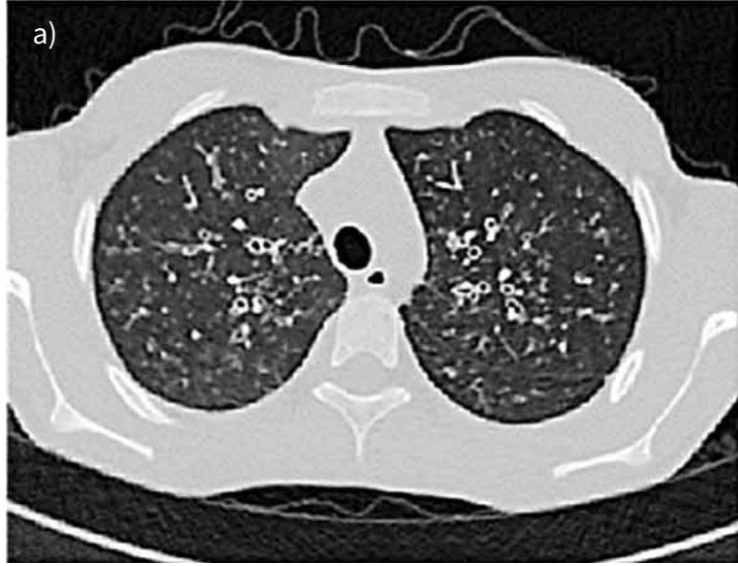
# AI-driven semantic labelling



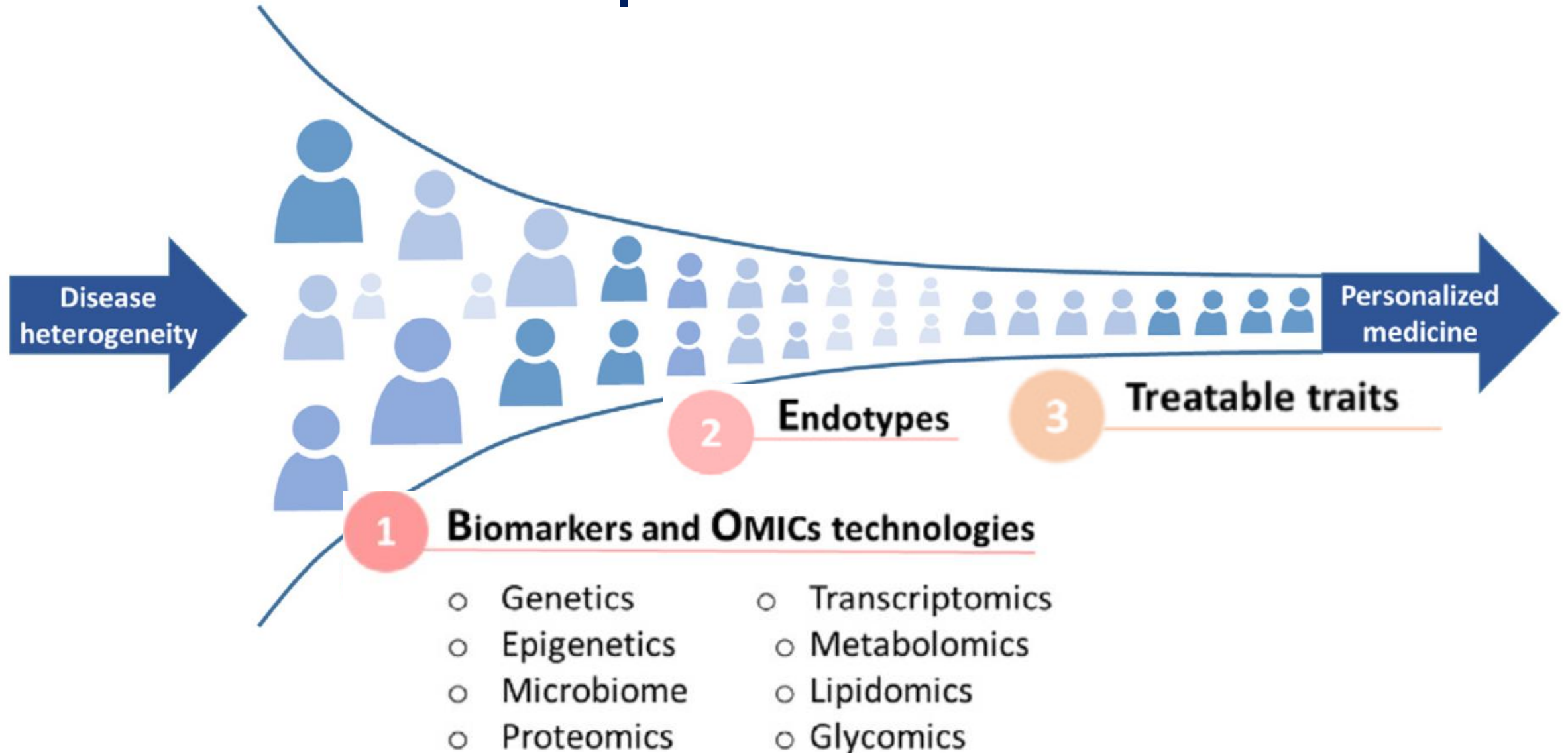
Axial CT slices  
→ AI-driven semantic labelling  
→ Integration all individual 2D labellings over the entire CT

**Blue:** central mucus plugs  
**Red:** mucus-free lumen dilatations  
**Green:** peribronchial thickening  
**Cyan:** consolidation

# AI-imaging before and after treatment



# Future directions of bronchiectasis research toward a personalized medicine



# Summary

## ➤ Biomarker

- ✓ Neutrophil elastase activity
- ✓ Microbiome
- ✓ Neutrophil extracellular trap

## ➤ Clinical evaluation

- ✓ Exacerbation
- ✓ Symptom burden
- ✓ COPD assessment test
- ✓ Eosinophilic phenotype and ICS response

## ➤ Radiology

- ✓ Small airway disease/emphysema
- ✓ Artificial intelligence for quantification

*Thank You for Your Attention*

