

1 **Clinical and inflammatory characteristics of the European U-BIOPRED adult severe asthma**  
2 **cohort**

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106

107 **Message:** Patients with severe asthma have more airway inflammation, worse symptoms and lower  
108 lung function, despite higher doses of treatment.

109

110

111 **Abbreviations**

112 BMI; Body mass index

113 FeNO; Fraction of exhaled nitric oxide

114 FEV<sub>1</sub>; Forced Expiratory Volume in one second

115 FVC; Forced Vital Capacity

116 HC; Healthy non-smoking controls

117 ICS; Inhaled Corticosteroids

118 MMA; Mild/moderate non-smoking asthma

119 OCS; Oral corticosteroids

120 SAn; Severe non-smoking asthma

121 SAs/ex; Smokers and ex-smokers with severe asthma

122 U-BIOPRED: Unbiased Biomarkers for the Prediction of Respiratory Disease Outcome

123

124 **Abstract** (196)

125 U-BIOPRED is an EU consortium of 20 academic institutions, 11 pharmaceutical companies and 6  
126 patient organisations with the objective of improving the understanding of asthma disease  
127 mechanisms using a systems biology approach.

128

129 This cross-sectional assessment of adults with severe asthma, mild/moderate asthma and healthy  
130 controls from 11 European countries consisted of analyses of patient-reported outcomes, lung  
131 function, blood and airway inflammatory measurements.

132

133 Patients with severe asthma (non-smokers n=311 and smokers/ex-smokers n=110) had more  
134 symptoms and exacerbations compared to patients with mild-moderate disease (n=88) (2.5  
135 exacerbations versus 0.4 in the preceding 12 months,  $p<0.001$ ), with worse quality of life, and higher  
136 levels of anxiety and depression. They also had a higher incidence of nasal polyps and gastro-  
137 oesophageal reflux with lower lung function. Sputum eosinophil count was higher in severe asthma  
138 compared to mild-moderate asthma (median count 2.99% versus 1.05%,  $p=0.004$ ) despite treatment  
139 with higher doses of inhaled and/or oral corticosteroids.

140

141 Consistent with other severe asthma cohorts, U-BIOPRED is characterised by poor symptom  
142 control, increased co-morbidity and airway inflammation, despite high levels of treatment. It is well  
143 suited to identify asthma phenotypes using the array of 'omic' datasets that are at the core of this  
144 systems medicine approach.

145

146

## 147 **Introduction**

148 A substantial number of patients with asthma require systemic corticosteroids to control symptoms  
149 and/or suffer from poor control and frequent severe exacerbations despite currently available  
150 treatment (1, 2). Although recently-developed biologic compounds targeting cytokines of the Type 2  
151 pathways show promise (3, 4), identification of new treatment targets and the selection of patients  
152 best suited to respond to individual biologics is still hampered by a poor understanding of the  
153 physiological, pathological, and molecular heterogeneity of severe asthma (5, 6).

154

155 Severe asthma is a collection of disease entities with varying pathophysiological characteristics (7)  
156 that result in symptoms of cough, wheeze and breathlessness, with frequent exacerbations. To  
157 address the problem of phenotypic difference and heterogeneity, the Unbiased Biomarkers for the  
158 Prediction of Respiratory Disease Outcomes (U-BIOPRED) project was set up in 2009 as a public-  
159 private partnership within the framework of the Innovative Medicines Initiative (IMI), engaging  
160 academia, the pharmaceutical industry and patient groups. The aim of U-BIOPRED is to identify  
161 multi-dimensional phenotypes of severe asthma and new treatment targets using a combination of  
162 state of the art 'omics' (transcriptomic, proteomic, lipidomic and metabolomic) technologies  
163 applying a systems biology approach (8) thereby driving unbiased discovery in both adult and  
164 paediatric severe asthma (9). This novel approach is designed to make drug development more  
165 effective and efficient.

166

167 We present the baseline characteristics of the adult participants with severe asthma who form the  
168 majority of the U-BIOPRED cohort and compare these participants with those suffering from with  
169 mild-to-moderate disease, in terms of their clinical, symptomatic, functional and biomarker features.  
170 In a parallel paper the characteristics of the paediatric cohort are reported. These first publications  
171 of U-BIOPRED will serve as the reference documents for all subsequent publications using the 'omics  
172 technologies which are at the core of this programme.

173

## 174 **Methods**

### 175 ***Participants***

176 This was a multi-centre prospective cohort study recruiting from 16 clinical centres in 11 European  
177 countries. Details of the participating centres, assessments, and standard operating procedures are  
178 available in the online supplement. Prior to enrolment, participants with severe asthma were  
179 required to have been under follow-up by a respiratory physician for at least six months during  
180 which time assessments had been undertaken to optimise asthma control and assess medication

181 adherence (2). The study was approved by the ethics committee for each participating clinical  
182 institution, and adhered to the standards set by International Conference on Harmonisation and  
183 Good Clinical Practice. It is registered on *ClinicalTrials.gov*, (Identifier: NCT01982162). All participants  
184 gave written and signed informed consent.

185

### 186 **Adult Groups**

187 The definition of severe asthma used in this study was agreed at a U-BIOPRED consensus meeting  
188 (2). Participants with asthma had either airflow reversibility (increase in FEV<sub>1</sub> >12% predicted or  
189 200ml following inhalation of 400µg salbutamol), airway hyper-responsiveness (methacholine PC<sub>20</sub> <  
190 8mg/ml, or diurnal PEF amplitude % mean >8%), or a decrease in FEV<sub>1</sub> of 12% predicted or 200ml  
191 within 4 weeks after tapering maintenance treatment. Four groups were recruited:

192

193 A) Severe non-smoking asthma (SAn):

194 Participants in this group were non-smokers for at least the past 12 months, with a less than five  
195 pack-year smoking history, with asthma and uncontrolled symptoms defined according to GINA  
196 guidelines (10) and/or frequent exacerbations (more than two per year) despite high-dose inhaled  
197 corticosteroids (ICS) (ICS ≥ 1000µg fluticasone propionate/day or equivalent dose).

198

199 B) Smokers and ex-smokers with severe asthma (SAs/ex):

200 This group was defined as for the SAn group except that they were either current smokers or ex-  
201 smokers with a smoking history of at least 5 pack years.

202

203 C) Mild/Moderate non-smoking asthmatics (MMA):

204 Participants in this group were non-smokers for at least the past 12 months, with a less than five  
205 pack-year smoking history and had controlled or partially controlled asthma symptoms, as defined  
206 by the Global Initiative for Asthma (GINA), whilst receiving a dose of less than 500µg fluticasone  
207 propionate/day or equivalent.

208

209 D) Healthy non-smoking controls (HC):

210 These participants had no history of asthma or wheeze, had no other chronic respiratory disease,  
211 were non-smokers for at least the past 12 months with a smoking history of ≤ 5 pack years and their  
212 pre-bronchodilator FEV<sub>1</sub> was ≥ 80% predicted.

213

214 **Protocol and assessments**



215 Participants attended a screening visit to assess eligibility for the study (**Fig 1**). They underwent a  
216 baseline visit (up to 28 days later) and were invited to attend for an optional bronchoscopy, high  
217 resolution lung computed tomography and telemonitoring sessions. Spirometry, haematological  
218 profiles, and fraction of exhaled nitric oxide level (FeNO) at 50ml/sec were performed. Induced  
219 sputum was obtained (11) and differential sputum eosinophil and neutrophil counts measured  
220 following a standardised operating procedure. Sputum supernatants and cell pellets were collected.  
221 Allergic status was obtained by either skin prick testing or measurement of specific IgE to six  
222 common aeroallergens. Blood and urine samples were taken for lipidomic, proteomic and  
223 transcriptomic analyses for later assessment. An optional sample was taken for genetic analysis.  
224 Subsets of participants underwent plethysmographic measurements, high-resolution computed  
225 tomography (HRCT) and collection of exhaled breath for measurement of metabolites including  
226 volatile organic compounds, all for future analyses. All investigations were performed according to  
227 standardised operating procedures (online supplement).

228

229 Participants with severe asthma were reviewed at 12-18 months after enrolment and were also  
230 invited to attend if they experienced an exacerbation. At 12-24 months, they were contacted by  
231 phone or by post to obtain information on asthma control.

232

233 Data were entered on an electronic case report form. The study was run and monitored by  
234 Cromsource ([www.cromsource.com](http://www.cromsource.com)). Samples were sent to the Centre for Integrated Genomic  
235 Medical Research Biobank in Manchester, UK. Datasets were uploaded on to the tranSMART system,  
236 an open-source knowledge management platform for sharing research data (12) supported by the  
237 European Translational Information and Knowledge Management Services (eTRIKS) project.

238

239 The study aims are published on the U-BIOPRED home page ([www.europeanlung.org/en/projects-and-research/projects/u-biopred/home](http://www.europeanlung.org/en/projects-and-research/projects/u-biopred/home)).

241

#### 242 **Questionnaires**

243 The following were administered at baseline:

- 244 1. The Asthma Control Questionnaire (ACQ5) (13) to assess current asthma control.
- 245 2. The Asthma Quality of Life Questionnaire (AQLQ) (14) to assess quality of life and  
246 psychological morbidity.
- 247 3. The Hospital Anxiety and Depression Scale (HADS) (15).
- 248 4. Sino-Nasal Outcomes Test (SNOT20) (16) to measure upper airway symptoms.

249 5. The Epworth Sleepiness Scale (ESS) (17) to measure sleep and daytime drowsiness.

250 6. The Medicines Adherence Response Scale (MARS) (18) to measure adherence.

251

## 252 **Statistical Analysis**

253 Continuously distributed data were either summarised using mean (standard error; SE) if  
254 symmetrical, or median (inter-quartile range; IQR) values. Non-symmetrical variables all exhibited  
255 positive skew and were log-transformed prior to association testing. Missing data were not imputed.

256 P-values were calculated using a general linear model for continuous variables or a general logistic  
257 model for categorical variables. No adjustment for multiple testing was applied as the analyses were  
258 considered exploratory. Analyses were performed using R version 2.15.2 (R Core Team, 2012).

259

## 260 **Results**

261 A total of 610 adults were recruited over an 18-month period: 311, 110, 88 and 101 into the SAn,  
262 SAs/ex, MMA, HC groups respectively (**Table i**).

263

	N	severe non-smoking asthma n=311	smokers and ex-smokers with severe asthma n=110*	mild/moderate non-smoking asthma n=88	healthy non-smoking controls n=101	P-value
Age (yr)	Mean (SE) [N]	51.01 (0.8) [311]	54.51 (1.08) [110]	41.66 (1.65) [88]	38.85 (1.34) [101]	p<0.001
Age at Diagnosis(yr)	Median (IQR) [N]	20 (7_38) [302]	38 (20_48) [109]	14 (6_32) [83]	NA	p<0.001
Female	n/N (%)	205/311 (66)	56/110 (51)	44/88 (50)	39/101 (39)	p<0.001
BMI (kg/m <sup>2</sup> )	Mean (SE) [N]	29.11 (0.36) [311]	29.59 (0.6) [110]	25.73 (0.47) [88]	25.31 (0.36) [101]	p<0.001
BMI>30 (kg/m <sup>2</sup> )	n/N %	120/311 (38.6)	44/110 (40)	16/88 (18.18)	12/101 (11.88)	<0.001
Serum IgE (IU/ml)	Median (IQR) [N]	119.5 (45_342) [302]	126 (63_328) [104]	89.4 (49_244) [85]	23.45 (9_65) [98]	p<0.001
FEV <sub>1</sub> (%)	Mean (SE) [N]	67.5 (1.26) [308]	67.2 (1.84) [110]	89.5 (1.86) [87]	101.76 (1.29) [101]	p<0.001
FVC (%)	Mean (SE) [N]	87.2 (1.12) [308]	89.7 (1.74) [110]	104.5 (2.02) [87]	107.8 (1.3) [101]	p<0.001
FEV <sub>1</sub> /FVC ratio	Mean (SE) [N]	0.64 (0.01) [308]	0.61 (0.01) [110]	0.72 (0.01) [87]	0.79 (0.01) [101]	p<0.001
Exacerbations in Previous Year	Mean (SE) [N]	2.48 (0.13) [310]	2.55 (0.26) [110]	0.38 (0.08) [88]	NA	p<0.001
Pack Years	Median (IQR) [N]	2 (1_4) [47]	17.38 (10_26) [110]	4 (1_4) [13]	0.9 (0_3) [20]	<0.001
Intubation (Ever)	n/N (%)	35/307 (11)	6/109 (6)	0/87 (0)	NA	0.083
ICU Admission (Ever)	n/N (%)	80/307 (26)	18/109 (17)	1/86 (1)	NA	p<0.001
Atopy test positive	n/N (%)	213/272 (78.3)	62/87 (71.3)	72/78 (92.3)	36/78 (46.2)	p<0.001

264

265 **Table i**

266 Group Demographics

267 FEV<sub>1</sub>: Forced Expiratory Volume in one second

268 FVC: Forced Vital Capacity

- 269 IQR: Inter quartile range
- 270 ICU: Intensive care unit
- 271 NA: Not applicable
- 272 SE: Standard error
- 273 \* 42 current smokers and 68 ex-smokers
- 274

275 There were more females in the SAn group (66%) compared to the other asthma groups (50%), with  
276 the age of onset of asthma 18yrs later in SAs/ex compared with SAn. Participants with severe  
277 asthma had a higher BMI than those in MMA and HC groups and were older (**Table i**). Both severe  
278 asthma groups experienced 2.5 exacerbations in the preceding 12 months as compared with 0.4 in  
279 the MMA group ( $p < 0.001$ ). There was a higher rate of ICU admissions in the SAn participants  
280 compared to the SAs/ex group ( $p < 0.05$ ). Further split of the severe asthma groups based on current  
281 and ex-smoking is presented in the online supplement (Table S5).

282

### 283 Spirometry (**Table i**)

284 FEV<sub>1</sub> (% predicted or actual) was lower in the three asthma groups compared to the HC group  
285 ( $p < 0.001$ ), with the severe asthma groups having the lowest FEV<sub>1</sub>. FVC (% predicted or actual) was  
286 also lower in both the SAn and SAs/ex groups when compared to the MMA ( $p < 0.001$ ) and HC groups  
287 ( $p < 0.001$ ). The mean FEV<sub>1</sub>/FVC ratio was lower in those with severe asthma (0.64 and 0.61,  
288 respectively) compared to the MMA (ratio 0.72,  $p < 0.001$ ) and HC groups (ratio 0.79,  $p < 0.001$ )  
289 respectively).

290

### 291 Medications (**Table ii**)

292 Within the SAn and SAs/ex groups 46% and 45% respectively received daily OCS, and 17% and 16%  
293 respectively received anti-IgE therapy. Use of nebulised  $\beta$ -agonist was higher in the SAn and SAs/ex  
294 groups. Other classes of therapy were also used.

295

		severe non-smoking asthma N=311	smokers and ex- smokers with severe asthma N=110	mild/moderate non-smoking asthma N=88
Oral corticosteroid	n/N (%)	135/295 (45.8)	46/103 (44.7)	0/88 (0)
Prednisolone Equ. (mg)*	Mean (SE) [N]	13.2 (0.85) [122]	14.8 (1.81) [36]	NA (NA)
Inhaled corticosteroids	n/N (%)	310/311 (99.7)	110/110 (100)	87/88 (98.9)
Long acting beta agonist	n/N (%)	305/309(98.7)	109/110 (99.1)	2/88 (2.3)
Short acting beta agonist	n/N (%)	260/301 (86.3)	82/105 (78.1)	68/88 (77.3)
Injected corticosteroids	n/N (%)	19/284 (6.7)	1/97 (1.0)	0/88 (0)
Mucolytic	n/N (%)	31/286 (10.8)	18/100 (18.0)	0/88 (0)
Anti-histamine	n/N (%)	75/311 (24.1)	16/110 (14.6)	4/88 (4.5)
Antibiotic (excluding macrolide)	n/N (%)	11/288 (3.8)	4/98 (4.1)	0/88 (0)
Macrolide	n/N (%)	32/311 (10.3)	13/110 (11.8)	0/88 (0)
Long-acting muscarinic antagonist	n/N (%)	65/284 (22.9)	27/97 (27.9)	0/88 (0)
Short acting muscarinic antagonist	n/N (%)	127/292 (43.5)	48/104 (46.2)	0/88 (0)
Omalizumab	n/N (%)	50/287 (17.4)	16/98 (16.3)	0/88 (0)
Immunosuppressant	n/N (%)	9/311 (2.9)	4/110 (3.6)	0/88 (0)
Leukotriene modifier	n/N (%)	139/298 (46.6)	45/106 (42.5)	0/88 (0)
Cromones	n/N (%)	10/284 (3.5)	2/97 (2.1)	0/88 (0)
Anti-fungal agent	n/N (%)	5/311 (1.6)	1/110 (1.0)	0/88 (0)
Xanthine	n/N (%)	59/289 (20.4)	21/100 (21.0)	0/88 (0)
Nebulised beta-agonist	n/N (%)	82/284 (28.9)	24/97 (24.7)	2/88 (2.3)

296

297 **Table ii**

298 Medications

299 \* Hydrocortisone and Triamcinolone doses were converted to equivalent prednisolone dose

300 (4 healthy control participants took as required antihistamines)

301 Questionnaires (**Table iii**)

302 ACQ and AQLQ scores reflected worse asthma control and increased morbidity in both severe  
303 asthma groups with minimal impairment in the MMA group. A similar pattern was seen with anxiety  
304 and depression. There were more upper airway symptoms measured using the SNOT20 in both  
305 severe asthma groups compared with the MMA group. Similarly the ESS scores indicated that there  
306 was an increase in sleepiness in the severe asthma groups compared to only a very mild impairment  
307 in the MMA group.

308

		severe non- smoking asthma  N=311	smokers and ex- smokers with severe asthma N=110	mild/moderate non-smoking asthma N=88	Unadjusted P-value SA* vs. MMA
<b>Asthma control questionnaire (ACQ)</b>					
Mean ACQ5	Mean (SE) [N]	2.28 (0.07) [277]	2.23 (0.12) [96]	0.86 (0.07) [85]	p<0.001
Mean ACQ7	Mean (SE) [N]	2.67 (0.08) [277]	2.62 (0.12) [96]	1.01 (0.07) [85]	p<0.001
<b>Asthma quality of life questionnaire (AQLQ)</b>					
Total	Mean (SE) [N]	4.48 (0.07) [276]	4.44 (0.13) [92]	5.84 (0.1) [84]	p<0.001
Symptoms	Mean (SE) [N]	4.46 (0.08) [276]	4.36 (0.14) [92]	5.87 (0.1) [84]	p<0.001
Emotional	Mean (SE) [N]	4.63 (0.1) [276]	4.52 (0.16) [92]	5.98 (0.13) [84]	p<0.001
Environmental stimuli	Mean (SE) [N]	4.69 (0.09) [276]	4.57 (0.16) [92]	5.63 (0.14) [84]	p<0.001
Activity limitation	Mean (SE) [N]	4.35 (0.07) [276]	4.45 (0.13) [92]	5.81 (0.11) [84]	p<0.001
<b>Hospital and anxiety and depression score (HADS)</b>					
Total	Mean (SE) [N]	12.33 (0.54) [223]	13.64 (1.01) [72]	7.01 (0.7) [70]	p<0.001
Anxiety	Mean (SE) [N]	6.94 (0.3) [223]	7.71 (0.54) [72]	4.24 (0.41) [70]	p<0.001
Depression	Mean (SE) [N]	5.39 (0.28) [223]	5.93 (0.56) [72]	2.77 (0.39) [70]	p<0.001
<b>Sino-nasal outcome test 20 (SNOT 20)</b>					
Total	Mean (SE) [N]	31.76 (1.01) [283]	32.12 (1.92) [97]	15.42 (1.42) [83]	p<0.001
<b>Epworth sleepiness scale (ESS)</b>					
Total	Mean (SE) [N]	7.48 (0.26) [277]	7.95 (0.47) [95]	5.49 (0.41) [85]	p<0.001
<b>Medication adherence rating scale (MARS)</b>					
Total	Mean (SE) [N]	22.44 (0.14) [278]	22.17 (0.29) [94]	21.35 (0.4) [84]	0.002

310 **Table iii**

311 Questionnaires

312 \*SA represents SAn and SAs/ex groups combined



313 The MARS questionnaire scores for adherence to treatment recorded by the three asthma groups  
314 were in the range of 21 to 22, with the severe asthma groups recording higher scores ( $p < 0.005$ )  
315 indicating better adherence. The AQLQ score was correlated to several variables, including FEV<sub>1</sub> (95%  
316 CI 0.5\_0.7,  $p < 0.001$ ), FEV<sub>1</sub>/FVC (95% CI 1.14\_2.8,  $p < 0.001$ ), exacerbations in the previous year (95%  
317 CI -0.8\_-0.2,  $p < 0.001$ ), BMI (95% CI -0.006\_-0.002,  $p < 0.001$ ) and pack years smoked (95% CI -0.003\_-  
318 0.001,  $p < 0.001$ ) (**Figure 3**).

319

#### 320 Atopy and co-morbidities (**Table iv**)

321 There was a high incidence of atopy in the 4 groups, at 70% in the asthma groups and 46% in the HC  
322 group. The incidence of allergic rhinitis, hay fever and non-allergic rhinitis were highest in the  
323 asthma groups. The HC group were much less allergic with only a third reporting hay fever and only a  
324 sixth, rhinitis or eczema.

325

326

		severe non- smoking asthma N=311	smokers and ex- smokers with severe asthma N=110	Mild and moderate non- smoking asthma N=88	healthy non- smoking controls N=101	P-value	P-value SA* vs. MMA
Allergic rhinitis diagnosed	n/N (%)	164/277 (59.2)	44/101 (43.6)	42/70 (60)	5/30 (16.7)	p<0.001	0.442
Hayfever diagnosed	n/N (%)	135/284 (47.5)	51/100 (51)	46/73 (63.0)	10/33 (30.3)	0.019	0.024
Non-allergic rhinitis diagnosed	n/N (%)	42/284 (14.8)	17/101 (16.8)	8/72 (11.1)	1/34 (2.9)	0.090	0.356
Nasal polyps diagnosed	n/N (%)	103/291 (35.4)	34/101 (33.7)	7/76 (9.2)	3/34 (8.8)	p<0.001	p<0.001
Eczema diagnosed	n/N (%)	107/294 (36.4)	31/101 (30.7)	25/75 (33.3)	5/35 (14.3)	0.013	0.789
GORD diagnosed	n/N (%)	135/289 (46.7)	63/99 (63.6)	16/75 (21.3)	4/35 (11.4)	p<0.001	p<0.001

327

328 **Table iv.** Co-morbidities.

329 GORD: Gastro-oesophageal reflux disease

330 \*SA represents SAn and SAs/ex groups combined

331

332 The presence of nasal polyps was associated with severe asthma, regardless of smoking status (4-  
333 fold increased incidence in SAn and SAs/ex groups versus MMA group,  $p < 0.001$ ) (**Table iv**). No such  
334 association was seen with allergic or non-allergic rhinitis, hay fever or reported eczema. Gastro-  
335 oesophageal reflux disease was more common in severe asthma (46% SAn, 63% SAs/ex) than in  
336 MMA (21%) and HC (11%), with a greater incidence reported in the SAs/ex group versus the SAn  
337 group ( $p=0.004$ ).

338

339 Blood and sputum biomarkers (**Table v**)

340 Blood eosinophil counts were similar in all three asthma groups. Each group had a significantly  
341 higher blood eosinophil count than the HC group (SAn vs. HC  $p=0.002$ , SAs/ex vs. HC  $p=0.005$ , MMA  
342 vs. HC  $p<0.001$ ). Blood neutrophil counts were significantly higher in the severe asthma groups  
343 compared to the MMA group.

344

345

	N	severe non-smoking asthma N=311	smokers and ex-smokers with severe asthma N=110	Mild and moderate non-smoking asthma N=88	healthy non-smoking controls N=101	P-value	P-value SA* vs. MMA
Exhaled NO ppb	Median (IQR) [N]	26.5 (16_47) [290]	23.5 (12_42) [104]	25 (18_54) [87]	19.25 (13_29) [96]	<0.001	0.438
<b>Sputum</b>							
Sputum eosinophils (%)	Median (IQR) [N]	2.75 (0_19) [128]	4.13 (1_14) [53]	1.05 (0_3) [43]	0 (0_0) [41]	p<0.001	0.004
Sputum neutrophils (%)	Median (IQR) [N]	53.69 (34_75) [128]	55.15 (35_65) [53]	44.5 (26_62) [43]	39.56 (21_56) [41]	0.002	0.042
Sputum differential eosinophil count >1.9%	n (%) [N]	74 (57.81) [128]	32 (60.38) [53]	17 (39.53) [43]	1 (2.44) [41]	<0.001	0.026
<b>Blood</b>							
Blood eosinophils (%)	Median (IQR) [N]	2.94 (1_6) [302]	2.88 (1_5) [106]	3.00 (2_5) [88]	2.10 (1_3) [101]	0.001	0.295
Blood Eosinophils (absolute)	Median (IQR) [N]	0.2 (0.3) 302	0.22 (0.29) 106	0.23 (0.2) 88	0.1 (0.11) 101	0.001	0.295
Blood neutrophils (%)	Median (IQR) [N]	62 (55_70) [302]	61.75 (55_69) [106]	56.83 (52_63) [88]	57.34 (51_64) [101]	p<0.001	p<0.001
Blood neutrophils (absolute)	Median (IQR) [N]	4.73 (3.1) 302	4.97 (2.87) 106	3.64 (1.75) 88	3.03 (1.6) 101	p<0.001	p<0.001

346

347 **Table v**

348 Biomarkers in blood, sputum and exhaled air

349 \*SA represents SAn and SAs/ex groups combined

350

351 Sputum samples were provided and met criteria for analysis in 44.2% of the asthma participants and  
352 40.6% of the HC group. Median sputum eosinophil counts for the SAn, SAs/ex, MMA, and HC groups  
353 were 2.75%, 4.13%, 1.05% and 0% respectively (**Table v**). The sputum eosinophil count was higher in  
354 the two severe asthma groups combined compared to the mild/moderate asthma group (**Table v**,  
355 **Fig 4**).

356 There were no significant differences in differential sputum neutrophil counts between the two  
357 severe asthma groups, which when combined were significantly higher compared to the MMA group  
358 (**Table v**).

359 There was a significant negative association between log sputum eosinophils (Absolute or %) and  
360 FEV<sub>1</sub> (% predicted or actual value) when all cohorts were considered and an adjustment for age, sex  
361 and smoking was applied. There were significant negative associations between log blood  
362 eosinophils (%) and FEV<sub>1</sub>/FVC ratio (p=0.002) and between blood neutrophils (%) and actual FEV<sub>1</sub>  
363 (p=0.002) and FEV<sub>1</sub>/FVC ratio (p=0.026).

364 Exhaled Nitric Oxide (FeNO) (**Table v**)

365 FeNO levels in all asthma groups were higher than those in the HC group, but the FeNO levels in the  
366 severe asthma groups were not different from the levels in the MMA group. The presence of nasal  
367 polyps was associated with a higher FeNO (mean increase 2.1ppb, 95% CI 1.5\_2.9, p<0.001).

368

## 369 **Discussion**

370 In this large European cohort, patients with severe asthma experienced more symptoms, more  
371 exacerbations, higher levels of anxiety and depression, and a higher incidence of nasal polyps,  
372 gastro-oesophageal reflux symptoms and airflow obstruction than patients with milder disease. The  
373 clinical characteristics of asthma were present despite higher doses of treatment that included doses  
374 of inhaled corticosteroids equal or more than 1,000µg of fluticasone (or equivalent), with 45% of the  
375 combined severe asthma group receiving a daily dose of prednisolone. The characteristic features of  
376 the severe asthma U-BIOPRED cohort are similar to those reported in previous cohort studies (6, 19-  
377 21). While the entry criteria for severe asthma were comparable for most of these cohort studies,  
378 the ENFUMOSA study required a lower threshold with an ICS dose of ≥ 1,200µg of budesonide or  
379 beclomethasone with at least one exacerbation in the past year. Of these 5 cohorts, the current U-  
380 BIOPRED severe asthma cohort appears to be the most severe with a higher reported exacerbation  
381 rate of 2.5 per year, a reduced mean FEV<sub>1</sub> of 67.5% of predicted and a higher proportion of patients  
382 on oral corticosteroid therapy taking a mean dose of 14 mg/day.

383

384 One of the novel features of the U-BIOPRED cohort is the inclusion of a smoking and ex-smoking  
385 severe asthma group. Patients with asthma who smoke have been reported to have poorer disease  
386 control and a reduced therapeutic response to ICS (22), possibly through the induction of  
387 corticosteroid insensitivity (23, 24). However our analyses of the non-smoking and the smoking/ex  
388 smoking severe asthma groups identified few differences in demographics, airway physiology,  
389 inflammatory markers and asthma symptoms between these groups. In both groups, a similar  
390 percentage received oral corticosteroid therapy; they also had similar degrees of airflow  
391 obstruction. The slightly lower level of FeNO in the smoking/ex smoking group might be explained  
392 by an effect of current smoking (26). One notable difference is that asthma onset occurred on  
393 average 18 years later in the smokers and ex-smokers than in the non-smokers, and yet the degree  
394 of airflow obstruction measured was similar. One interpretation is that there may be a more rapid  
395 rate of loss of lung function in the patients with asthma who smoke. The significant correlation  
396 between AQLQ scores and the number of pack-years of smoking exposure would also support a  
397 contribution of cigarette smoke to impaired quality of life in this group. We also split the  
398 demographic data of the groups by smoking status rather than severity (see Table S5) in the online  
399 supplement. This revealed that current smokers had a lower BMI compared with ex and never  
400 smokers.

401

402 In agreement with the SARP study (20), patients with severe asthma (especially smokers) were less  
403 frequently atopic than those with mild/moderate disease. There was also a clear association of both  
404 nasal polyps and gastro-oesophageal reflux disease with disease severity, with approximately one-  
405 third and one half reporting polyps and reflux respectively, a finding that is in keeping with previous  
406 reports (5). Nasal polyps are commonly found in severe asthma, and are associated with a  
407 particularly severe phenotype. There is evidence that treating nasal polyps with anti-IgE therapy  
408 results in better asthma outcomes (25), however whether this is due to an effect on the underlying  
409 asthma or the polyps is unknown. The link with higher FeNO levels is in keeping with work showing  
410 that nasal polypectomy leads to a fall in FeNO (26).

411

412 Our findings are also similar to other studies published from severe asthma registries. In agreement  
413 with both the British Thoracic Society's (27) and Belgium's (28) severe asthma registries our patients  
414 are predominantly female, with a high BMI and evidence of fixed airflow obstruction. Moreover  
415 there are similarly high levels of reflux, nasal polyps and exacerbations despite greater levels of  
416 medication.

417

418 We found a greater degree of sputum eosinophilia in the two severe asthma groups compared to  
419 the mild-moderate asthma group. Up to 60% of patients in the two severe asthma groups had a  
420 differential sputum eosinophil of >1.9% (the established upper limit of normal for differential  
421 sputum eosinophil counts (29)). This percentage is similar to previous reports in severe asthma (21).  
422 The level of sputum eosinophilia observed in the mild/moderate asthma group are also similar to  
423 those reported previously (30).

424

425 The higher blood neutrophil count in participants with severe asthma may represent the effect of  
426 systemic corticosteroids which can increase blood neutrophil numbers. Sputum neutrophil counts  
427 were similar in the three asthma groups and were significantly higher than the in healthy control  
428 group. This similarly could represent the effect of corticosteroids although severe asthma has been  
429 linked to a higher level of sputum neutrophils (31, 32).

430

431 The impact and burden on our participants' health was noticeable with measures of symptoms and  
432 quality of life being far worse in severe asthma as compared to mild/moderate asthma, despite the  
433 use of higher doses and more classes of asthma treatment. Levels of anxiety and depression were  
434 also higher with severe asthma. There were significant relationships between quality of life  
435 measures and airflow obstruction, smoking history and BMI, supporting the contribution of these  
436 factors to an impairment of quality of life however the scatter of data reveals that these parameters  
437 are not closely related. The number of exacerbations experienced was greater than 2.5  
438 exacerbations per participant in both severe asthma groups in the preceding year. These findings  
439 highlight the need for an integrative assessment of clinical and physiological disease markers but  
440 additionally biological markers of disease in the assessment of severe asthma. For example, the  
441 finding that bariatric surgery has an effect on measures of airway hyper-responsiveness (33) and is  
442 associated with a lower all-cause mortality at 5 years particularly in younger, predominantly female  
443 populations (34) may point towards the need for specific and targeted intervention in people with  
444 severe asthma and obesity.

445

446 There are several limitations to our study. Firstly, there is no perfect way to assess treatment  
447 adherence; however, we only approached patients managed in a specialist respiratory clinic and only  
448 those who had been assessed to be adherent were eligible for the study. Furthermore MARS scores  
449 were high indicating good levels of self-reported adherence. Secondly, subjective or historical data  
450 were assessed by questionnaire which may be prone to recall bias. Thirdly, the success rate in  
451 obtaining adequate quality sputum for analysis was in the 42-50% range and the number of

452 bronchoscopies was relatively lower in the SA and SAs/ex groups. Thirdly due to the numerous  
453 formulations and inhaler devices used across Europe it was not possible to calculate the precise daily  
454 equivalent ICS dose for each participant and therefore these data are not shown, however high  
455 (>1000mcg FP) or low (<500mcg FP) dose was a study entry requirement for the severe and  
456 moderate groups respectively.

457

458 We have been successful in recruiting a substantial cohort of patients with the most severe asthma  
459 that has similar characteristics to previously-reported cohorts. This gives confidence that the U-  
460 BIOPRED consortium will define distinct phenotypes and endotypes of severe asthma. Matching  
461 these data to the 'omics' information with future unsupervised analyses will help identify new  
462 treatments for patients with severe asthma who currently have limited treatment options, and will  
463 improve our understanding of this important chronic disease.

464

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475

476



477

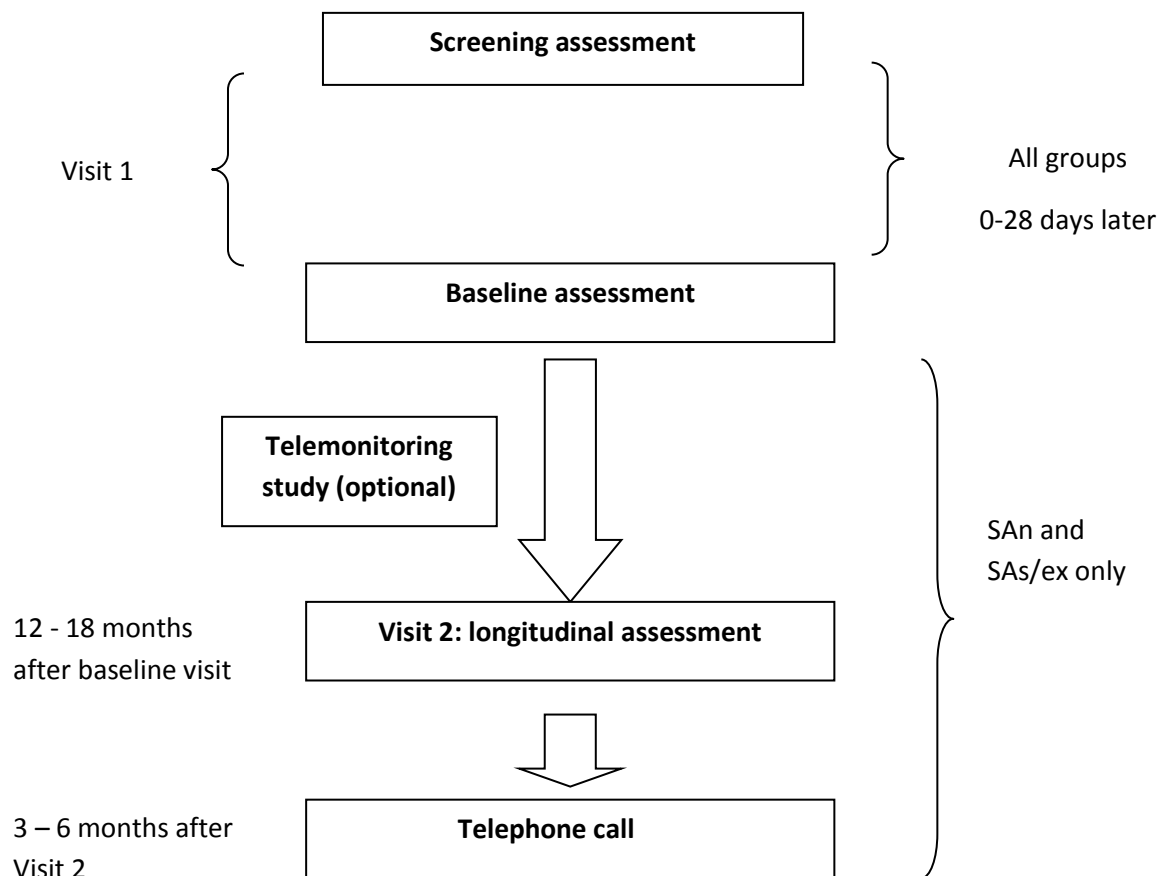
478 **Fig 1**

479 **Visit Schedule**

480

481

482



483

Figure 2

Consort diagram

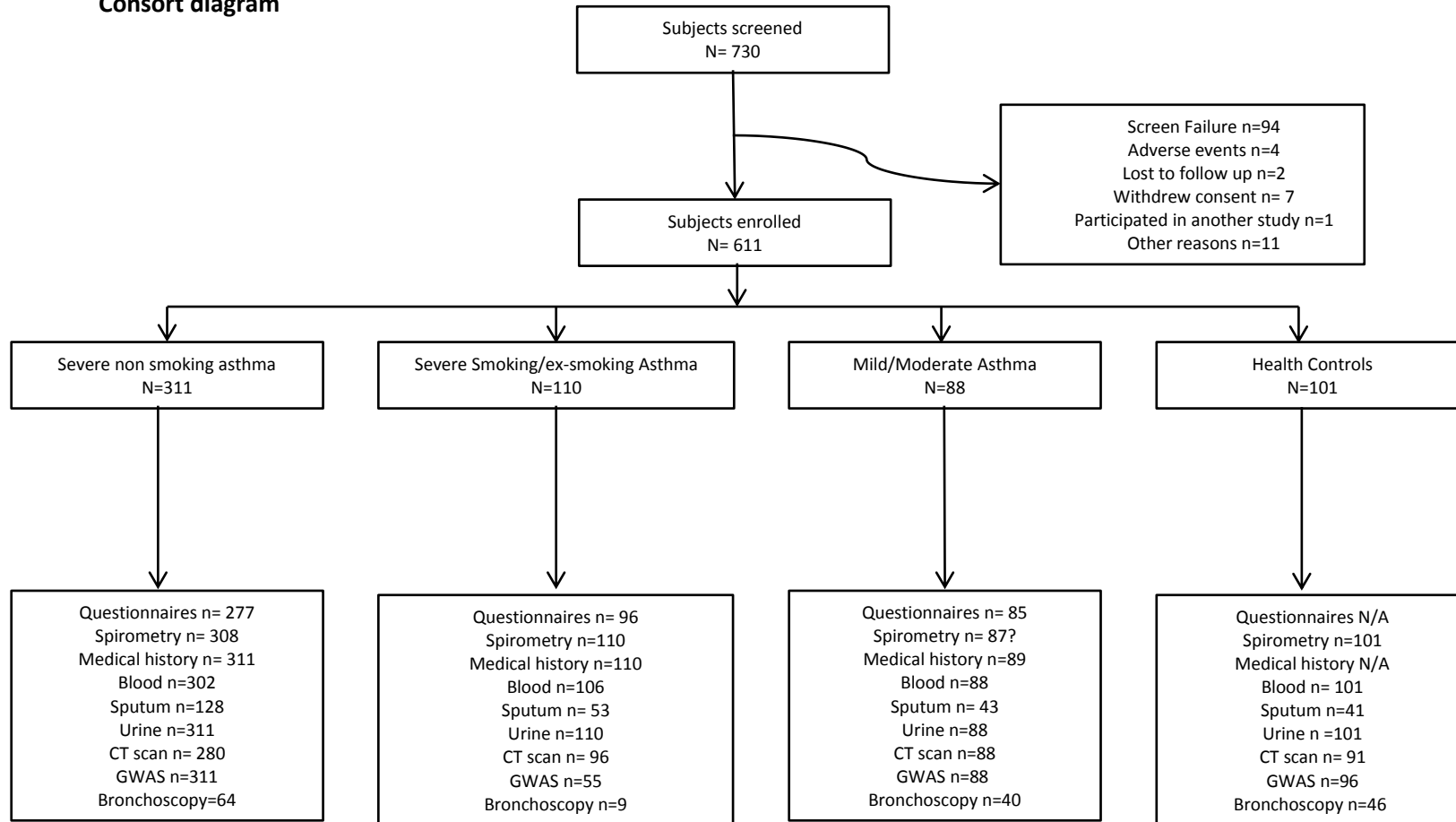
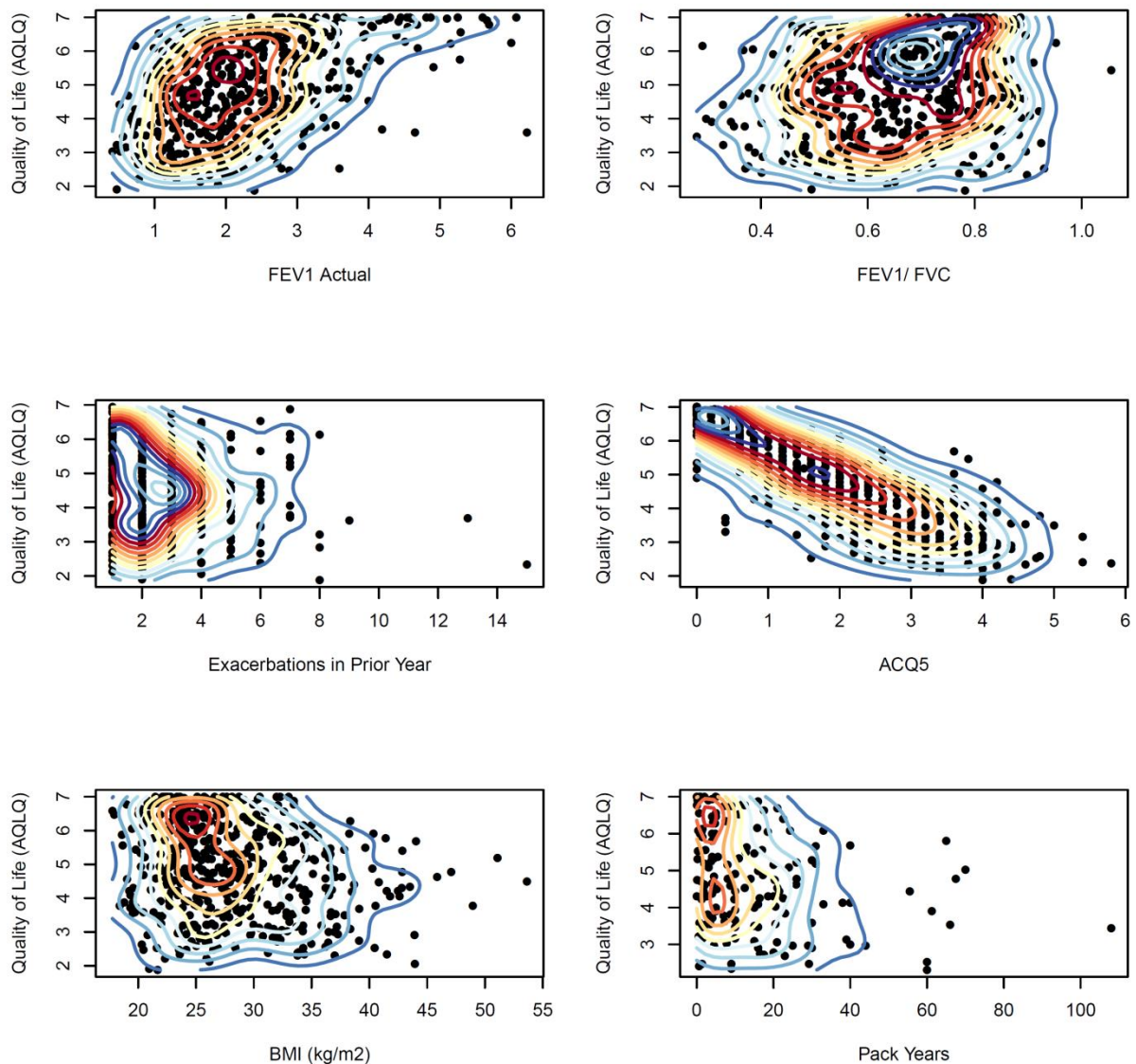


Figure 3

## Contour plots of AQLQ related to baseline demographics



Figures represent scatter plots describing the relationship between each factor and the asthma quality of life z-score. The contour lines are coloured blue to red, to indicate increasing density of points in the graph. The density was modelled using two-dimensional kernel density estimation. The contour plots show weak inverse relationships and particularly the scatter between quality of life and exacerbations, BMI and pack years, a strong inverse relationship between quality of life and asthma control and weak positive relationships between quality of life and measures of lung function.

Fig 4A

## Eosinophil counts by group

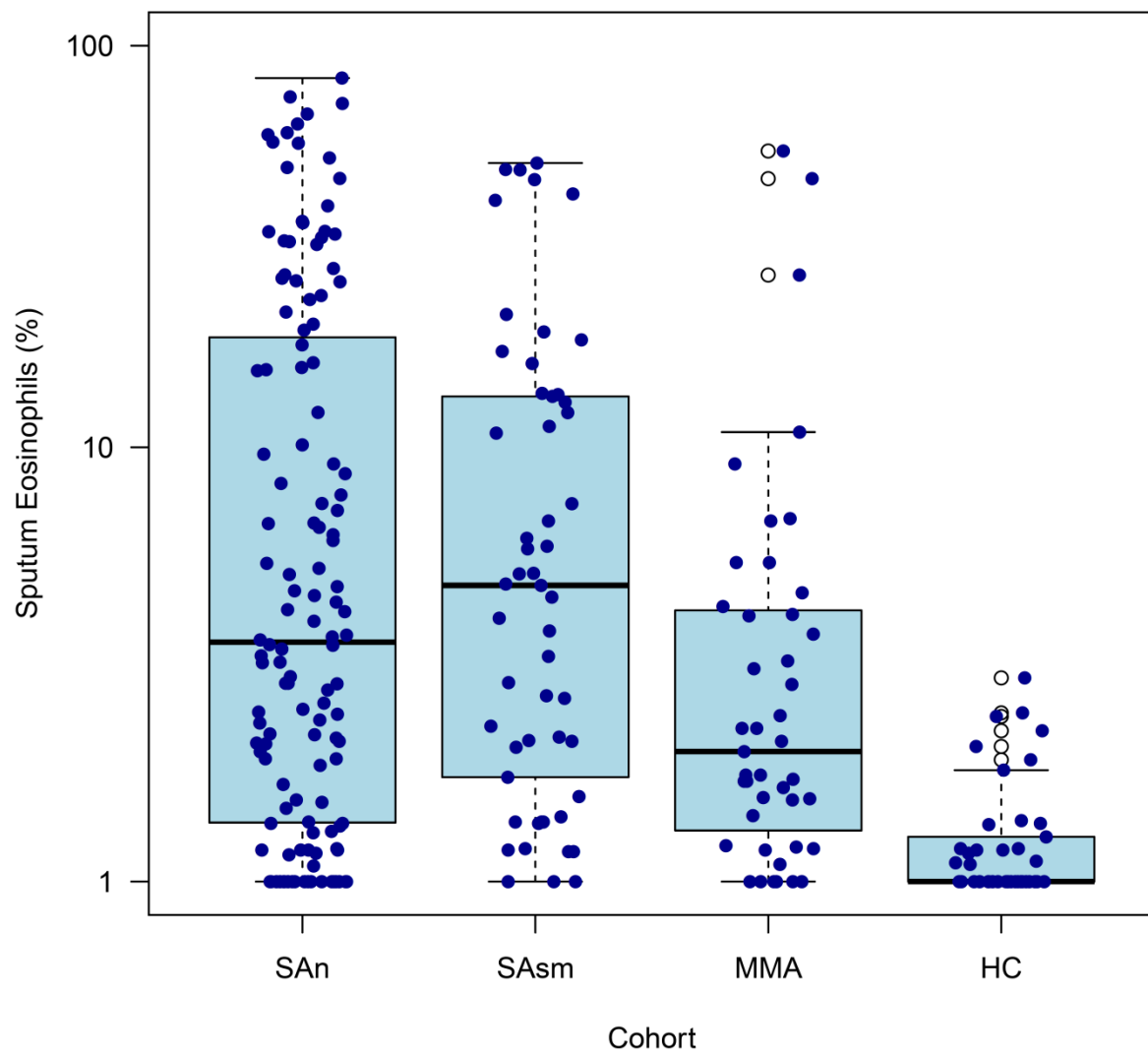


Fig 4B

## Neutrophil counts by group

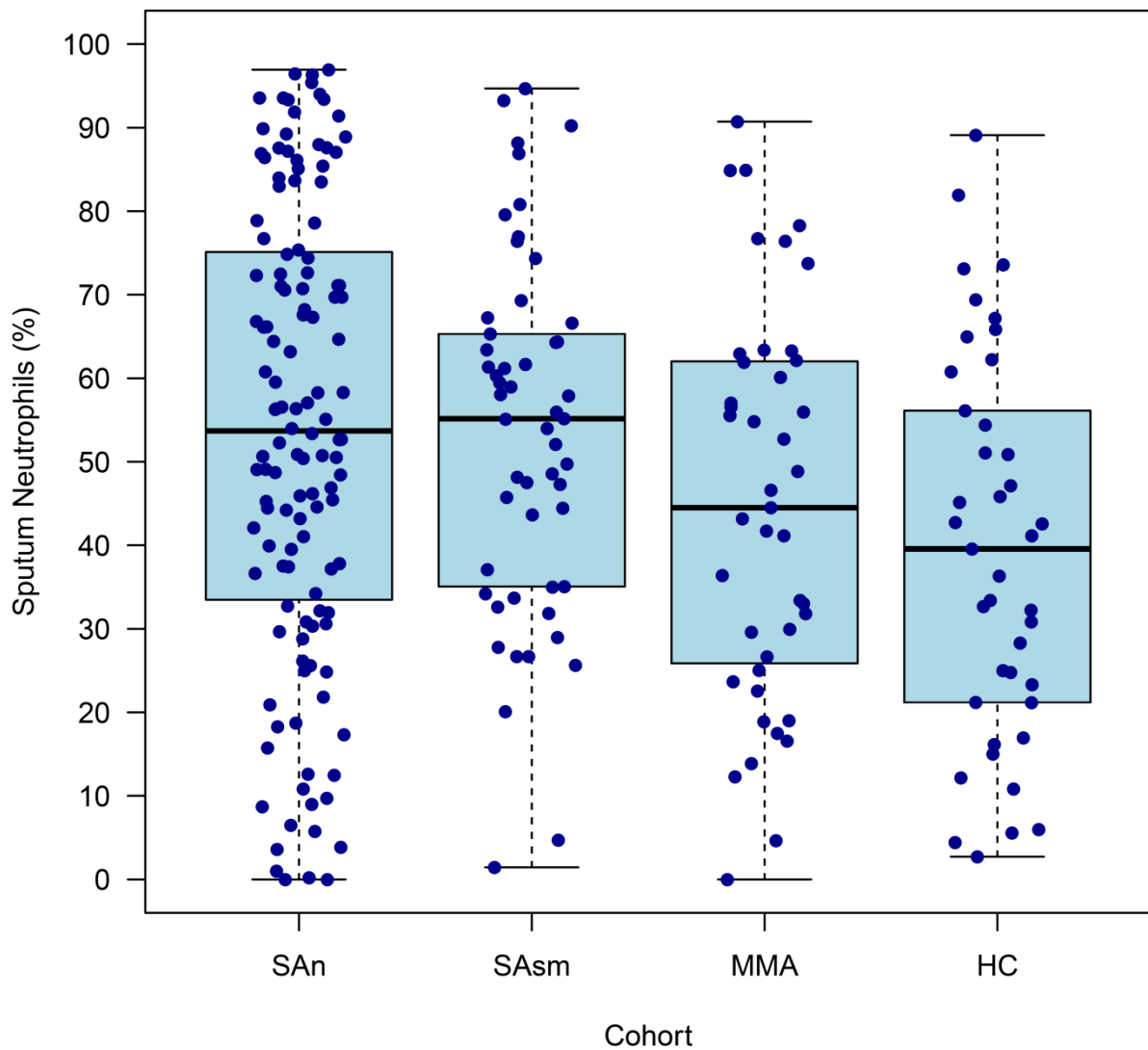


Fig 4

A) sputum eosinophil count and B) sputum neutrophil count, by cohort. The box and whisker plots are shaded in pale blue, with outliers denoted by open circles. The raw data are given by dark blue points overlaid.

SAn; Severe non-smoking asthma

SAs/ex; Smokers and ex-smokers with severe asthma

MMA; Mild/moderate non-smoking asthma

HC; Healthy non-smoking controls

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