

Supplementary material

Table S1. Patients' clinical data.

AP (n=141)	
<i>Routine blood biomarkers</i>	
Glucose (mg/dL)	136 (115-168)
Urea (mg/dL)	27 (19-33)
Creatinine (mg/dL)	0.79 (0.67-0.92)
<i>Arterial blood gas test</i>	
pH	7.41 (7.39-7.44)
pCO ₂	29.3 (26.3-33.0)
pO ₂	53.0 (48.2-62.2)
SatO ₂	86.0 (82.0-98.0)
HCO ₃ ⁻	18.7 (16.7-20.6)

Median and interquartile range (IR 25–75) are presented for all the variables. pH: Hydrogen potential. pCO₂: partial pressure CO₂. pO₂: Partial pressure Oxygen, SatO₂: Oxygen saturation. HCO₃⁻: Bicarbonate.

Table S2. Genetic association between SNPs and clinical data.

Gene/SNP	Rhinitis (n=52)	No rhinitis (n=80)	p	OR	CI (95%)
	GF (%)	GF (%)			
<i>IL13 rs20541</i>					
GG	22 (42.3)	33 (41.3)	0.33	1.0 (Ref)	
GA	23 (44.3)	42 (52.5)		0.81	0.39-1.72
AA	7 (13.4)	5 (6.2)		2.1	0.59-7.46
<i>IL13 rs1800925</i>					
CC	26 (50.0)	39 (48.8)	0.87	1.0 (Ref)	
CT	21 (40.4)	31 (38.7)		1.03	0.47-2.13
TT	5 (9.6)	10 (12.5)		0.74	0.23-2.43
<i>IL4 rs2070874</i>					
TT	16 (30.8)	24 (30.0)	0.82	1.0 (Ref)	
TC	26 (50.0)	37 (46.2)		1.05	0.47-2.36
CC	10 (19.2)	19 (23.8)		0.79	0.29-2.13
Gene/SNP	Dermatitis (n=6)	No dermatitis (n=126)	p	OR	CI (95%)
	GF (%)	GF (%)			
<i>IL13 rs20541</i>					
GG	3 (50.0)	52 (41.3)	0.72	1.0 (Ref)	
GA	3 (50.0)	62 (49.2)		0.84	0.16-4.33
AA	0	12 (9.5)		NA	NA
<i>IL13 rs1800925</i>					
CC	3 (50.0)	62 (49.2)	0.9	1.0 (Ref)	
CT	2 (48.3)	50 (39.7)		0.79	0.13-4.90
TT	1 (1.7)	14 (11.1)		1.31	0.14-4.52
<i>IL4 rs2070874</i>					
TT	1 (1.7)	34 (26.9)	0.18	1.0 (Ref)	
TC	5 (98.3)	58 (46.2)		2.93	0.33-26.15
CC	0	34 (26.9)		NA	NA
Gen/SNP	Allergies (n=34)	No allergies (n=98)	p	OR	CI (95%)
	GF (%)	GF (%)			
<i>IL13 rs20541</i>					
GG	11 (32.4)	44 (44.9)	0.41	1.0 (Ref)	
GA	20 (58.8)	45 (45.9)		1.78	0.76-4.14
AA	3 (8.8)	9 (9.2)		1.33	0.31-5.77
<i>IL13 rs1800925</i>					
CC	19 (55.9)	46 (46.9)	0.32	1.0 (Ref)	
CT	10 (29.4)	42 (42.9)		0.54	0.22-1.29
TT	5 (14.7)	10 (10.2)		1.13	0.34-3.76
<i>IL4 rs2070874</i>					
TT	13 (38.2)	22 (22.4)	0.16	1.0 (Ref)	
TC	15 (44.1)	48 (48.9)		0.53	0.22-1.30
CC	6 (17.7)	28 (28.6)		0.36	0.12-1.11

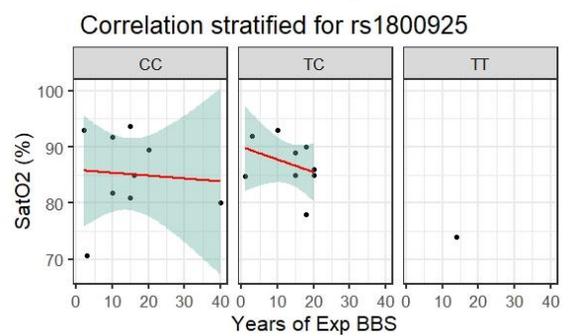
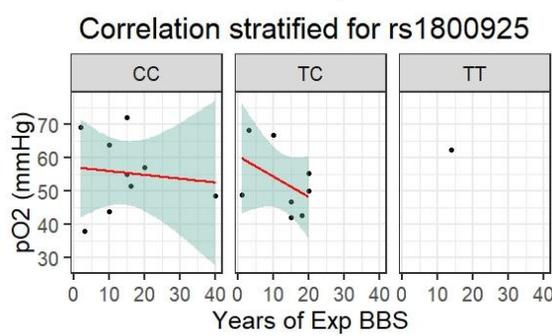
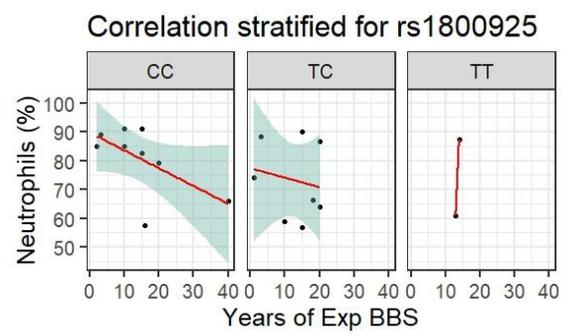
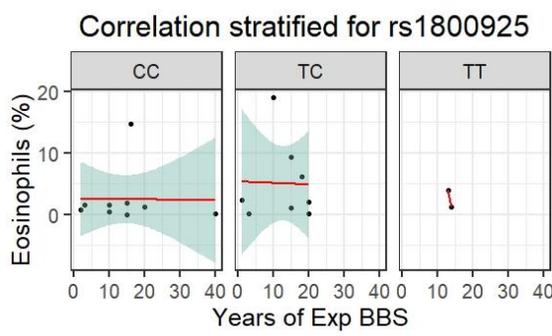
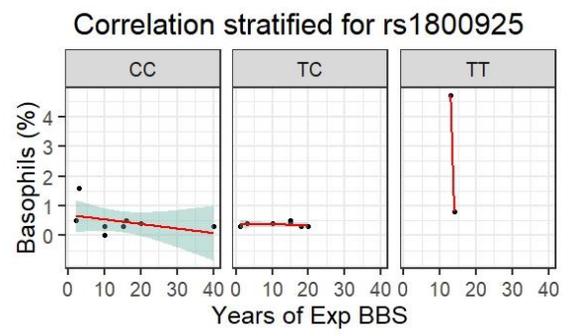
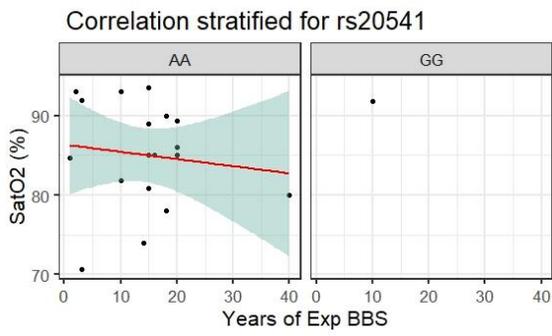
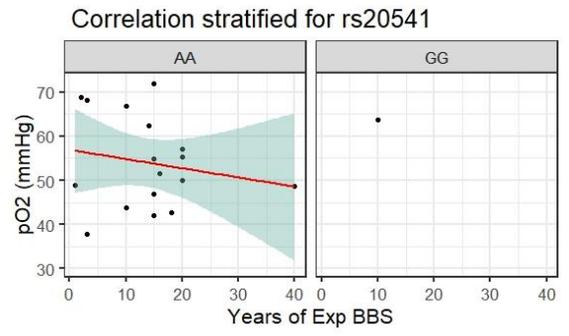
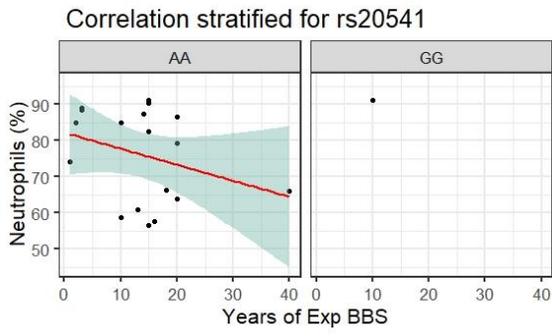
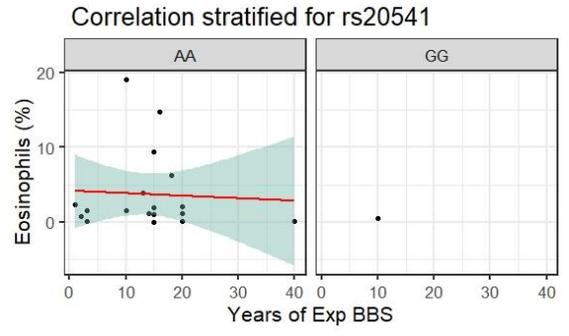
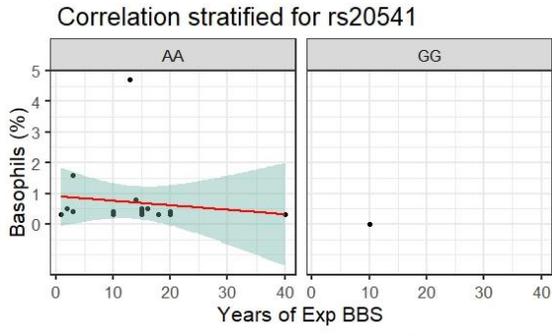
Genotype frequencies were compared by Fisher's exact test. For statistical significance was considered a $p < 0.05$. Ref: Genotype employed as a reference for statistical analysis. CI: Confidence intervals. OR: Odds ratio. NA: Not apply.

Table S3. Genetic association between SNPs and clinical data.

Total leukocytes					
Gene/SNP	Normal (n=73)	High (n=52)	p	OR	CI (95%)
	GF (%)	GF (%)			
<i>IL13</i> rs20541					
GG	29 (39.7)	24 (46.2)	0.41	1.0 (Ref)	
GA	39 (53.4)	22 (42.3)		1.47	0.69-3.11
AA	5 (6.9)	6 (11.5)		0.69	0.19-2.54
<i>IL13</i> rs1800925					
CC	35 (47.9)	26 (50.0)	0.08	1.0 (Ref)	
CT	33 (45.2)	17 (32.7)		1.57	0.71-3.49
TT	5 (6.9)	9 (17.3)		0.4	0.12-1.33
<i>IL4</i> rs2070874					
TT	20 (27.4)	13 (25.0)	0.87	1.0 (Ref)	
TC	35 (47.9)	24 (46.2)		0.95	0.40-2.26
CC	18 (27.7)	15 (28.8)		0.78	0.29-2.08
Neutrophils					
Gene/SNP	High (n=90)	Normal (n=35)	p	OR	CI (95%)
	GF (%)	GF (%)			
<i>IL13</i> rs20541					
GG	40 (44.4)	13 (37.1)	0.68	1.0 (Ref)	
GA	41 (45.6)	19 (54.3)		0.7	0.31-1.61
AA	9 (10.0)	3 (8.6)		0.98	0.23-4.15
<i>IL13</i> rs1800925					
CC	42 (46.7)	18 (51.4)	0.7	1.0 (Ref)	
CT	36 (40.0)	12 (34.3)		1.29	0.55-3.02
TT	9 (10.0)	5 (14.3)		0.77	0.23-2.63
<i>IL4</i> rs2070874					
TT	23 (25.6)	10 (28.6)	0.6	1.0 (Ref)	
TC	41 (45.6)	18 (51.4)		0.99	0.39-2.50
CC	26 (28.9)	7 (20.0)		1.62	0.53-4.94
Lymphocytes					
Gene/SNP	High (n=109)	Normal (n=16)	p	OR	CI (95%)
	GF (%)	GF (%)			
<i>IL13</i> rs20541					
GG	47 (43.1)	6 (37.5)	0.82	1.0 (Ref)	
GA	53 (48.6)	8 (50.0)		0.85	0.27-2.61
AA	9 (8.3)	2 (12.5)		0.57	0.09-3.31
<i>IL13</i> rs1800925					
CC	51 (46.8)	9 (56.2)	0.39	1.0 (Ref)	
CT	43 (39.5)	4 (25.0)		1.89	0.54-6.59
TT	11 (10.1)	3 (18.8)		0.64	0.15-2.78
<i>IL4</i> rs2070874					
TT	30 (27.5)	3 (18.8)	0.73	1.0 (Ref)	
TC	50 (45.9)	8 (50.0)		0.62	0.15-2.53
CC	28 (25.7)	5 (31.2)		0.56	0.12-2.56
Eosinophils					
Gene/SNP	High (n=62)	Normal (n=36)	p	OR	CI (95%)

	GF (%)	GF (%)			
IL13 rs20541					
GG	24 (38.7)	17 (47.2)		1.0 (Ref)	
GA	32 (51.6)	16 (44.4)	0.71	1.41	0.59-3.36
AA	6 (9.7)	3 (8.3)		1.41	0.31-6.49
IL13 rs1800925					
CC	32 (51.6)	18 (50.0)		1.0 (Ref)	
CT	22 (35.5)	12 (33.3)	0.82	1.03	0.41-2.56
TT	6 (9.7)	5 (17.7)		0.67	0.18-2.52
IL4 rs2070874					
TT	16 (25.8)	9 (25.0)		1.0 (Ref)	
TC	28 (45.2)	17 (47.2)	0.98	0.92	0.33-2.55
CC	18 (29.0)	10 (27.8)		1.01	0.32-3.11
Gene/SNP	Low (n=25)	Normal (n=36)	p	OR	CI (95%)
	GF (%)	GF (%)			
IL13 rs20541					
GG	11 (44.0)	17 (47.2)		1.0 (Ref)	
GA	13 (52.0)	16 (44.4)	0.89	1.25	0.43-2.60
AA	1 (4.0)	3 (8.3)		1.54	0.08-27.36
IL13 rs1800925					
CC	9 (36.0)	18 (50.0)		1.0 (Ref)	
CT	13 (52.0)	12 (33.3)	0.38	2.16	0.71-6.65
TT	3 (12.0)	5 (17.7)		1.2	0.23-6.19
IL4 rs2070874					
TT	8 (32.0)	9 (25.0)		1.0 (Ref)	
TC	13 (52.0)	17 (47.2)	0.55	0.86	0.26-2.84
CC	4 (16.0)	10 (27.8)		0.45	0.10-2.01
Basophils					
Gene/SNP	High (n=113)	Normal (n=9)	p	OR	CI (95%)
	GF (%)	GF (%)			
IL13 rs20541					
GG	49 (43.4)	2 (22.2)		1.0 (Ref)	
GA	55 (48.7)	6 (66.7)	0.46	0.37	0.01-1.94
AA	9 (7.9)	1 (11.1)		0.36	0.03-4.49
IL13 rs1800925					
CC	56 (49.6)	3 (33.3)		1.0 (Ref)	
CT	41 (36.3)	5 (55.6)	0.54	0.43	0.09-1.94
TT	13 (11.5)	1 (11.1)		0.69	0.06-7.25
IL4 rs2070874					
TT	29 (25.7)	4 (44.4)		1.0 (Ref)	
TC	54 (47.8)	4 (44.4)	0.39	1.86	0.43-7.99
CC	30 (26.5)	1 (11.1)		4.13	0.44-39.26

Genotype frequencies were compared by Fisher's exact test. For statistical significance was considered a $p < 0.05$. Ref: Genotype employed as a reference for statistical analysis. CI: Confidence intervals. OR: Odds ratio.



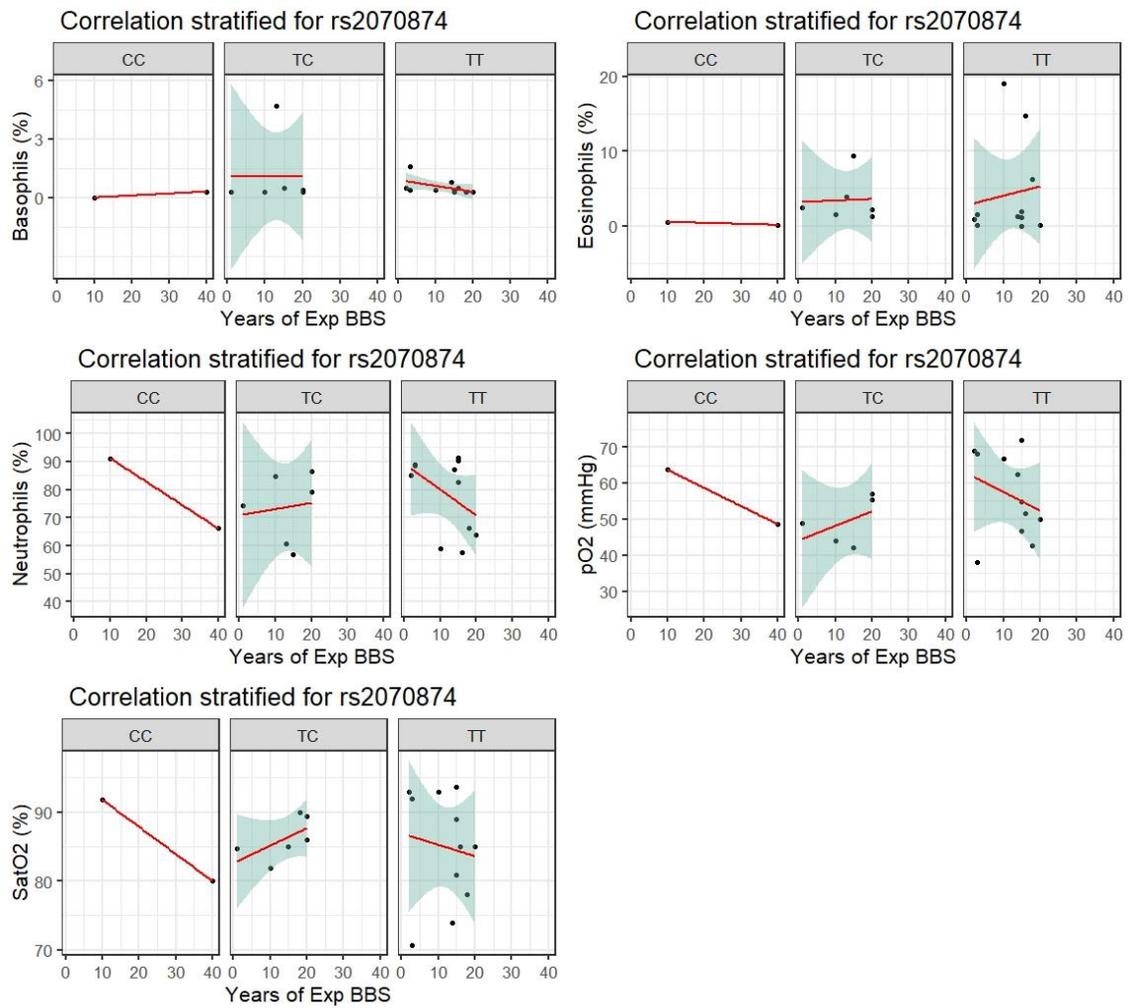


Figure S1. Correlation between cellularity and SNPs in the context of years of exposition to BBS. Blue areas show standard error (95%).

Table S4. Correlation between SNPs and cellularity/arterial blood gas values in the context of BBS.

Gene/SNP	Variable	ρ	p	CI 95%	
Neutrophils					
<i>IL13</i> rs1800925	TT	NA	NA	NA	
	TC	-0.12	0.800	(-0.73 - 0.59)	
	CC	-0.66	0.055	(-0.92 - 0.02)	
	Eosinophils				
	TT	NA	NA	NA	
	TC	-0.19	0.656	(-0.76 - 0.54)	
	CC	-0.07	0.855	(-0.70 - 0.62)	
	Basophils				
	TT	NA	NA	NA	
TC	-0.37	0.368	(-0.85 - 0.45)		
CC	-0.23	0.545	(-0.78 - 0.51)		
pO₂					
TT	NA	NA	NA	NA	
TC	-0.20	0.627	(-0.76 - 0.53)		
CC	-0.06	0.881	(-0.66 - 0.59)		
SatO₂					
TT	NA	NA	NA	NA	
TC	-0.22	0.569	(-0.75 - 0.48)		
CC	-0.16	0.682	(-0.72 - 0.52)		
Neutrophils					
<i>IL13</i> rs20541	GG	NA	NA	NA	
	GA	NA	NA	NA	
	AA	-0.25	0.318	(-0.63 - 0.23)	
	Eosinophils				
	GG	NA	NA	NA	
	GA	NA	NA	NA	
	AA	-0.10	0.702	(-0.53 - 0.37)	
	Basophils				
	GG	NA	NA	NA	
	GA	NA	NA	NA	
	AA	-0.37	0.128	(-0.72 - 0.13)	
	pO₂				
	GG	NA	NA	NA	NA
	GA	NA	NA	NA	NA
	AA	-0.14	0.602	(-0.57 - 0.35)	
SatO₂					
GG	NA	NA	NA	NA	
GA	NA	NA	NA	NA	
AA	-0.07	0.772	(-0.51 - 0.39)		
Neutrophils					

<i>IL4</i> <i>rs2070874</i>	TT	-0.37	0.259	(-0.77 - 0.22)
	TC	0.26	0.618	(-0.61 - 0.85)
	CC	NA	NA	NA
Eosinophils				
	TT	0.10	0.762	(-0.50 - 0.64)
	TC	-0.26	0.618	(-0.85 - 0.61)
	CC	NA	NA	NA
Basophils				
	TT	-0.62	0.042	(-0.90 - 0.02)
	TC	0.22	0.682	(-0.64 - 0.83)
	CC	NA	NA	NA
pO₂				
	TT	-0.41	0.206	(-0.80 - 0.21)
	TC	0.56	0.322	(-0.46 - 0.94)
	CC	NA	NA	NA
SatO₂				
	TT	-0.20	0.549	(-0.70 - 0.42)
	TC	0.75	0.084	(0.00 - 0.96)
	CC	NA	NA	NA

Data obtained from Spearman's Rho correlation. pO₂: Partial pressure Oxygen, SatO₂: Oxygen saturation, NA: Not applicable. CI: Confidence intervals. ρ = Ro value.