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Profile of exposures and lung function in adults with asthma: an exposome approach in the EGEA study

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Abstract

Background: Environmental research on multifactorial health outcomes calls for exposome approaches able to assess the joint effect of multiple exposures.

Objective: Our aim was to identify profiles of exposure to lifestyle/environmental factors associated with lung function in adults with asthma using a cluster-based approach.

Methods: We used data from 599 adults of the Epidemiological study on the Genetics and Environment of Asthma, bronchial hyperresponsiveness and atopy (EGEA) (mean age 39.0 years, 52% men) who ever had asthma. Exposures to 53 lifestyle/environmental factors were assessed by questionnaires or geographic information systems-based models. A two-step approach was developed: 1) exposome dimension reduction by selecting factors showing association with forced expiratory volume in 1 second (FEV₁) (p <0.20) in an exposome-wide association study (ExWAS), 2) clustering analysis using the supervised Bayesian Profile Regression (sBPR) to group individuals according to $FEV₁$ level and to their profile of exposure to a reduced set of uncorrelated exposures (each paired correlation<0.70) identified in step 1.

Results: The ExWAS identified 21 factors showing suggestive association with FEV₁ (none) significant when controlling for multiple tests). The sBPR conducted on 15 uncorrelated exposures identified in step 1, revealed 3 clusters composed of 30, 115 and 454 individuals with a mean \pm SD FEV₁(%pred) of 79% \pm 21, 90% \pm 19 and 93% \pm 16, respectively. Cluster 1 was composed of individuals with heavy smoking, poor diet, higher outdoor humidity and proximity to traffic, while cluster 2 and 3 included individuals with moderate/low levels of exposure to these factors.

Discussion: This exposome study identified a specific profile of joint lifestyle and environmental factors, associated with a low $FEV₁$ in adults with asthma. None of the exposures revealed significant association when considered independently.

Words count: 272

Key words: Lung function**,** asthma, exposome, Bayesian Profile Regression

1 **Introduction**

2 Over decades, several studies reported associations between exposure to various environmental 3 factors and asthma related outcomes. The most consistent associations are observed for tobacco 4 smoke (Strachan and Cook, 1998; Vignoud et al., 2011), allergens and sensitizers (house-dust 5 mites, moulds, domestic pets, …), (Wright and Phipatanakul, 2014) traffic-related air pollution 6 (Guarnieri and Balmes, 2014) and occupational exposures to asthmagens (Tarlo and Lemiere, 7 2014). Other exposures are suspected to be associated with asthma, such as use of cleaning 8 products at home (Zock et al., 2007) and climate factors (Correia Junior et al., 2017). Finally, 9 there are emerging concerns for other exposures including greenness or lifestyle factors such 10 as diet and physical activity (Fuertes et al., 2016; Sbihi et al., 2017; Varraso and Camargo, 11 2016). Although asthma is a multifactorial disease in which environmental and lifestyle factors 12 are suspected to interact, most of previous environmental studies are limited by the fact that 13 they focused on a single exposure or family of exposures (Vineis, 2018).

14 The concept of exposome was first proposed by C. Wild in 2005 to encompass life-course 15 environmental exposures (including lifestyle factors), from the prenatal period onwards (Wild, 16 2005). It was developed to highlight the need for more comprehensive approaches to address 17 the role of the environment in complex diseases (Schwartz and Collins, 2007) carrying the 18 expectation that the use of holistic and data-driven approaches can expand our understanding 19 of the aetiology and long-term trajectories of complex diseases.

20 The exposome includes a large number of covariates that are inter-related and aggregated, in 21 particular within exposure families (Tamayo-Uria et al., 2019). Among the different statistical 22 approaches to deal with such exposome data (Santos et al., 2020), a comprehensive 23 methodology based on clusters representing joint covariate profiles as opposed to individual 24 risk factors offers new avenues in environmental research on multifactorial diseases (Molitor 25 et al., 2010). This type of clustering approach has already been applied in the field of asthma to 26 disentangle the asthma phenotypic heterogeneity by simultaneously considering several clinical 27 and individual characteristics (Boudier et al., 2013; Haldar et al., 2008; Siroux et al., 2011). In 28 the context of the exposome-respiratory health association, although a recent study from Agier 29 *et al.* considered a large number of exposures using an exposome approach to evaluate their 30 association with lung function in European children, their work did not consider cluster-based 31 analyses to account for combined exposures (Agier et al., 2019). A recent study applied a 32 cluster-based analysis to account for joint prenatal exposures to chemicals in asthma and lung 33 function in 319 children, but the study was limited to a single family of exposures (phenols and 34 phthalates) (Berger et al., 2020). Because some of these risk factors for respiratory health might 35 have a stronger effect among individuals with asthma than among individuals without asthma, 36 as clearly demonstrated for smoking (Polosa and Thomson, 2013), investigating the exposome-37 lung function association in individuals with asthma might be more powerful than in the general 38 population.

39 In the frame of the Epidemiological Study on the Genetics and Environment of Asthma (EGEA) 40 and based on a detailed exposome in adulthood, we aimed to identify the joint effect of lifestyle 41 and environmental exposures on lung function in adults with asthma using a supervised 42 clustering approach.

43

44 **Methods**

45 *Study population*

46 EGEA is a French cohort based on three surveys over 20 years. At EGEA1 in early 1990s, 388 47 cases with asthma were recruited in chest clinics, so as their first-degree relatives $(n=1244)$ and 48 415 population-based controls. All participants were born in France as well as their two parents. 49 The population was invited to participate to the 12-year and 20-year follow-ups (EGEA2 and 50 EGEA3). At each survey, individuals responded to a detailed questionnaire on respiratory 51 health and environment, based on international standardised tools (Kauffmann and Dizier, 52 1995). The present study focuses on adults with ever asthma at EGEA2 (2003-2007) who 53 performed a spirometry. Ever-asthma was defined at EGEA1 by being recruited as an asthma 54 case in chest clinics or, for first degree relatives, by a positive answer to either "Have you ever 55 had attacks of breathlessness at rest with wheezing?" or "Have you ever had asthma attacks?".

56 *Spirometry testing*

57 Spirometry was performed using a flow-volume spirometer (SpiroDyn'R; SAS Dyn'R, Aix-en-58 Provence, France) according to the American Thoracic Society and European Respiratory 59 Society guidelines (Miller et al., 2005). Pre-bronchodilator forced expiratory volume in 1 s 60 (FEV1) was expressed in percentage of the predicted value using the Global Lung Function 61 Initiative reference equations, thereafter referred as $FEV₁%$ (Quanjer et al., 2012) to account 62 for the major effects of age, height and gender in $FEV₁$.

63 *Characterisation of the exposome*

64 Data regarding lifestyle factors were assessed mostly through self-completed questionnaires, 65 which provided information about 9 exposure families: indoor air (3 variables), tobacco 66 exposure (4 variables), season of examination (season at which individual performed the 67 spirometry, 1 variable), pets (2 variables), occupational exposures (2 variables, assessed 68 through a job-exposure matrix), household cleaning products (2 variables), rurality (2 69 variables), socio-economic status (SES) (assessed at the contextual level by the French 70 Deprivation Index (FDep) from current geocoded residential address, built to capture health 71 inequalities at ecological level (Temam et al., 2017), 1 variable), and physical activity and diet 72 (2 variables, assessed through the Metabolic Equivalent Tasks (METs) and the Alternative 73 Healthy Eating Index-2010 (AHEI-2010) diet score, respectively). Moreover, a broad range of 74 environmental exposures was assessed through geographic information system (GIS) models 75 that allow allocating a spatio-temporal estimate of exposure from geographical residential 76 coordinates. These GIS-based exposures covered 8 families: altitude (1 variable), building and 77 population densities (3 variables), atmospheric pollutants (4 variables), natural spaces (8 78 exposures), weather (6 variables), roads/cycle paths (3 variables), ambient ultra violet radiation 79 (UV) (3 variables) and health services (6 variables). The GIS-based models are resting on 80 National Geographic Institute (IGN) maps, the French National Institute for Statistics and 81 Economic Research (INSEE) data, Météo-France data, geospatial models, land cover databases 82 and satellite data. All these exposures were calculated from the current geocoded residential 83 addresses. Most of these exposures was assessed during the last 12 months before the 84 spirometry examination, except for some exposures assessed by GIS-based models for which 85 models of exposure were not available for the relevant time window (see Exposome assessment 86 method in the Supplementals).

87 In total, 53 exposure variables (19 lifestyle and 34 environmental), a priori candidate risks 88 factors in respiratory health, were assessed at EGEA2.

89 *Statistical analysis*

90 Figure 1 shows the statistical workflow. Exposure variables were transformed (to approach 91 normality) or dichotomized and missing values were imputed using the method of chained 92 equations (White et al., 2011) and 5 complete datasets were generated. All continuous 93 exposures variables were standardized by the inter-quartile range (IQR) to allow between-94 covariate comparison of regression estimates. Correlations between exposure variables were 95 presented through heatmaps. Correlations were assessed through Pearson coefficient for two 96 quantitative variables, polychoric coefficient for two qualitative variables and polyserial 97 coefficient for a quantitative and a qualitative variable. A second heat map assessed absolute 98 within (diagonal) and between (off-diagonal) families of exposure median correlations.

99 We proposed a two-step approach to evaluate the impact of the exposome on lung function. 100 The first step of the association study consisted in the exposome dimension reduction to allow 101 a better stability of the clustering model applied in step 2. Exposome-Wide Association 102 Analysis (ExWAS) was performed for FEV1% by fitting independent mixed linear regression 103 models for each exposure variable, with adjustment for the study centre, gender, age, height 104 and socio-professional category as fixed effects and family as random effect (Patel et al., 2010). 105 Factors showing association with $FEV₁%$ (p-value < 0.2) were selected in the reduced set of 106 exposures.

107 The second step consisted in identifying profiles of co-exposures associated with low lung 108 function. A supervised cluster analysis, *i.e.* simultaneously considering in the model the 109 outcome and exposures, was conducted using the supervised Bayesian Profile Regression 110 (sBPR) clustering method that relies on Markov Chain Monte Carlo sampling methods 111 (200,000 iterations) (Molitor et al., 2010). The sBPR, that requires to *a priori* select a unique 112 set of variables, was applied in the reduced set of uncorrelated (<0.7) exposures identified in 113 step 1 by the ExWAS, FEV1% and included confounders used in the ExWAS analysis (age, 114 gender, height, centre of inclusion, number of smoked pack-years and socio-professional 115 category) as covariates. When the absolute correlation between two of the exposures identified 116 in step 1 was above 0.70, we retained the variable with the lowest number of missing data, and 117 when these two numbers were similar we retained the variable with the lowest p-value in the 118 ExWAS analysis. Socio-demographic characteristics and exposure factors were compared 119 between clusters to identify the factors mainly driving the clustering. This analysis was also 120 performed after adjustment for age, gender and centre to identify exposures associated with 121 clusters independently of these 3 variables. Moreover, phenotypic characteristics at EGEA2, 122 but also at EGEA3 about 7 years later, were compared between clusters.

123 Because this clustering method relies on a stochastic process, we tested the stability of our final 124 model by running the model with 10 different seeds (all with 200,000 iterations). Each model 125 identified 3 clusters and the median [interquartile range] number of subjects belonging to the 126 same cluster whatever the seed used was 0.86 [0.73-0.94] (data not shown).

127 All analyses were performed using R software (version 3.5). We used the software package 128 *mice* for multiple imputation, *polycor* for correlation coefficients, *lme4* for mixed models and 129 *PReMiuM* to fit BPR models.

130 **Results**

131 *Description of the population*

132 Among 1,903 individuals included in the EGEA2 study, 698 individuals ever had asthma. 133 Among them, 99 were not analysed further because they had not performed spirometry or had 134 missing data for their residential address or did not complete the environmental questionnaire 135 (Figure 2). Excluded individuals had similar characteristics than the 599 included individuals, 136 except that they were younger (mean \pm SD age 39 \pm 6 *vs.* 34 \pm 18 years, p=0.01, for included *vs.* 137 excluded individuals, respectively, see supplemental Table E1). Main characteristics of the 138 studied population are presented in Table 1. The 599 included individuals belonged to 356 139 families, and half of them were men and had never smoked. Two thirds of the population were 140 executives or technicians. The studied population had a mean \pm SD FEV₁% of 92% \pm 17% and 141 88% had either at least one asthma attack or used asthma treatment in the past 12 months.

142 *Description of the exposome*

143 Descriptive statistics, transformation used and number of missing values for each exposure 144 variable are presented in Table 2. Most of the environmental exposures showed high between-145 centre variability (see supplemental Table E2). Overall, correlations between exposures were 146 higher within-families of exposure (median [IQR] absolute correlation = 0.46 [0.28-0.76]) than

147 between-families of exposures (median [IQR] absolute correlation = 0.08 [0.04-0.13]) (Figures 148 3 and 4). As an example, the three variables belonging to the building and population densities 149 family had a median absolute 2-way correlations of 0.78 while the median of the absolute 2- 150 way correlations between any of these three variables and the two variables belonging to the 151 rurality family was 0.26.

152 *Exposome dimension reduction from ExWAS results*

153 Among the 53 exposure variables, the ExWAS analysis identified 21 exposures from 10 154 families of exposures associated with FEV₁% with a p-value < 0.20 (Supplemental Table E3 and 155 Figure E1). After the examination of the correlation matrix of these 21 variables, 15 156 uncorrelated exposure variables from 9 exposure families were retained for the inclusion in the 157 sBPR analysis. Among them, 7 had a positive association with FEV1% (distance to nearest 158 green space and major road, humidity rate, population density, AHEI-2010 score, the second 159 quintile of the French Deprivation Index (FDep) score (*vs.* the first quintile), and having at least 160 one pharmacy in their surrounding space), while 8 had negative association with $FEV₁%$ (\geq 1 161 major road and/or one cycle path in a 100m buffer, be born in a rural municipality, mean annual 162 wind ≥2 m/s, distance to nearest blue space, mean annual temperature, season of examination 163 (reference=spring) and intermediate exposure to passive smoking (*vs.* no exposure)). None of 164 the association tested in the ExWAS was significantly associated with $FEV₁$ % when correcting 165 for multiple comparison (Supplemental Figure E1).

166 *Identification of profiles of joint exposures associated with FEV1% using the sBPR*

167 The sBPR, conducted on FEV1 and on the 15 uncorrelated exposures identified in the previous 168 step, identified 3 clusters composed of 30, 115 and 454 individuals, respectively. Lifestyle and 169 environmental characteristics of the three clusters are described in Table 3. Individuals 170 belonging to cluster 1 (n=30), that shows the lowest mean $FEV₁%$ (mean \pm SD 79% \pm 21%), were 171 more often heavy smokers and had more often an unhealthy diet, ≥ 1 major road in a 100m

172 buffer surrounding their home, a higher outdoor humidity rate and less pharmacy surrounding 173 their residential address. Regarding individuals assigned in cluster 2 (n=115, mean \pm SD FEV₁%) $174 = 90\% \pm 19\%$, they had the same mean age and mean pack-years than individuals from cluster 175 1, but had a healthier diet and were exposed to lower rate of humidity than individuals belonging 176 to other clusters. Finally, individuals belonging to cluster 3 (n=454, mean \pm SD FEV₁% = 177 93% ± 16 %) were younger, more often never-smokers but more often exposed to passive 178 smoking, and lived closer to blue spaces. After adjustment on age, gender and centre, 179 comparison of exposures between the 3 clusters remained overall similar, except for the 180 distance to nearest green space that became significantly lower in cluster 1 as compared to the 181 other clusters.

182 *Phenotypic characterization of the clusters at EGEA2 and about 7 years later at EGEA3*

183 Individuals from cluster 1 had a statistically significant lower prevalence of allergic 184 sensitization and higher rate of neutrophils at EGEA2 than individuals from other clusters 185 (Table 4). Although not statistically significant, individuals belonging to cluster 1 more often 186 reported use of inhaled steroids in the past 12 months as compared to those from cluster 2 and 187 3 (57%, 36% and 40% respectively). No difference was observed for the number of asthma 188 symptoms and the total Asthma Quality of Life Questionnaire score. Comparison of phenotypic 189 characteristics observed about 7 years later at EGEA3 (n=502, Supplemental Table E4) between 190 the three clusters did not show statistically significant differences, although individuals 191 belonging to cluster 1 tended to have more often asthma exacerbation over the past 12 months 192 than those belonging to cluster 2 and 3 (35% vs 18% and 21%, respectively, Table 4).

193

194 **Discussion**

195 To our knowledge, this is the first study that addresses the impact of the exposome on lung 196 function in adults with asthma by considering simultaneously a broad range of lifestyle factors 197 and social, built and natural environment. While none of the factors revealed significant 198 association with FEV_1 when considered independently in an $ExWAS$ approach, the profile 199 regression method revealed 3 profiles of joint exposures that show strong differences in terms 200 of FEV1% level and exposure profiles including lifestyle (smoking, diet), social (at the area and 201 individual levels), built (proximity to major road, pharmacy) and natural (blue space, humidity) 202 environments.

203 *Methodological considerations*

204 A major novelty in our study relies on the comprehensive analytical approach used, allowing 205 considering the outcome and a large set of exposures simultaneously to cluster individuals. 206 Indeed, until now most exposome studies considered exposure variables individually while, in 207 their daily life, individuals are simultaneously exposed to a large number of various factors, 208 which may jointly affect health. Thus, joint exposures were considered to identify clusters of 209 adults with asthma sharing the same exposure profiles. This method has the advantage to deal 210 with inter-correlated variables, as it is the case in exposome data. Nevertheless, this method did 211 not allow us to consider the family structure of our data. Thus, we cannot rule out that the family 212 design, although considered in the first step based on ExWAS, led to confounding issues 213 through shared environment and genetic factors within families. In addition, the BPR method 214 relies on Markov chain Monte Carol sampling methods which might lead to instability issue, in 215 particular with increased number of variables. Although we attempted to limit the model 216 instability by reducing the exposome dimension, avoiding very strong correlation between 217 exposures and running the algorithm for 200,000 iterations, this remains a limitation. (Molitor 218 et al., 2010). Finally, clustering methods being based on the entire set of exposure variables that 219 are considered simultaneously to identify clusters, the model is highly dependent of the set of 220 variables including in the model and the addition or the deletion of one of these variables could 221 lead to substantial changes in results. Strong *a priori* hypotheses on the input variables are 222 therefore needed and sensitivity analyses on different set of variables should be avoided.

223 *Results interpretation*

224 Interestingly, the sBPR method allowed identifying joint exposures that formed subpopulations 225 at risk of higher or lower FEV1%, while the ExWAS resting on traditional regression models 226 revealed no statistically significant association after considering multiple comparisons. Mixture 227 of exposures is expected to have higher impaired health effects than each individual exposure 228 because the effects of each factor are superimposed (Kortenkamp and Faust, 2018). 229 Accordingly, cluster 1, that was associated with the lowest $FEV₁$, was characterized by 230 smoking, unhealthy diet and exposure to traffic that are susceptible to be involved in similar 231 biological pathways, such as for example inflammation and oxidative stress. A synergistic 232 effect cannot be dismissed, as suggested by a study indicating a modifier effect of antioxidants 233 on lung function in Mexican children exposed to high levels of air pollutants (Romieu et al., 234 2002), although environment-by-environment interactions have been poorly investigated.

235 As previously mentioned, our study is the first to address the impact of joint exposures on 236 respiratory health, and therefore, results are difficult to compare with previous literature. 237 Among the wide range of lifestyle, social, built and natural environment factors that we took 238 into account in our exposome approach, only some of them play a major role in the 239 identification of the clusters. Most of these factors, taken individually, have previously been 240 identified, or were suggested to be involved, in asthma characteristics. For example, previous 241 studies showed that smoking reduces lung function, in both individuals with and without asthma 242 (Lange et al., 1998). A greater AHEI-2010, an overall measure of diet quality, has been linked 243 with improved or fewer asthma symptoms and greater asthma control (Andrianasolo et al., 244 2018; Chiuve et al., 2012; Li et al., 2017). The presence of at least 1 major road in a 100m 245 buffer surrounding the house, a proxy of traffic-related air pollution, was associated with lower 246 FEV1 and exacerbations in asthma patients (Balmes et al., 2009; Guarnieri and Balmes, 2014). 247 For other factors (distance from pharmacies, annual wind and outdoor humidity) results are 248 more novel and warrant further investigation. Besides these factors identified in our clusters, 249 some factors suggested to be associated with asthma characteristics or lung function decline in 250 the literature, such as molds exposure (Pekkanen et al., 2007; *WHO Guidelines for Indoor Air* 251 *Quality*, 2009) or domestic exposure to cleaning products (Dumas et al., 2017; Le Moual et al., 252 2012), were not identified as major determinants by our study. Overall, our findings underline 253 the interest of the exposome, as a complementary approach in environmental epidemiology to 254 better understand how co-exposures may impact respiratory health.

255 Regarding the phenotypic differences between the three clusters we found that cluster 1, 256 showing the lowest $FEV₁$ % value, included more subjects without allergic sensitisation and 257 with higher level of neutrophils as well as a higher subsequent risk for asthma exacerbations, 258 indicating that it might be a group characterized by a more severe form of the disease.

259 *Strengths and weaknesses*

260 Our study has a number of strengths. The main one, as previously discussed, lies in the fact that 261 the comprehensive method applied allows for the first time to investigate the effect of multiple 262 exposures together on lung function and does not simply adjust for co-exposures. By allowing 263 to consider all possible interactions between the set of exposures, the BPR approach used allows 264 to make a step forward in addressing interactions, usually restricted to approach considering 265 interactions between pairs of exposures. In addition, the analysis was conducted in a well-266 characterized cohort of adults with asthma with strong attention paid to the quality of the 267 objective lung function measures (Siroux et al., 2016). All exposure data were collected through 268 validated questionnaires or state of the art GIS-based models.

269 We acknowledge that our study also suffers from several weaknesses. First, although our study 270 is one of the first attempts to integrate multiple families of exposures, our exposome does not 271 include all exposures relevant for lung function. For example, some chemical exposures 272 (pesticides, phenols, phtalates, perfluorinated compounds, …) for which there is emerging 273 concern regarding a potential impact on respiratory health (Agier et al., 2019; Hoppin et al., 274 2009), concentration of aeroallergens, such as pollens, and early life exposures were not 275 available. Secondly, all considered exposures were either self-reported in questionnaires or 276 derived from GIS-based models, which could imply a lack of accuracy in the exposure 277 assessment. Nevertheless, GIS used in this study were based on high spatial resolution models, 278 below 1km x 1km for most of them. Third, although we collected through questionnaire 279 phenotypic characteristics of individuals 7 years after their participation to the EGEA2, the 280 current analysis is mainly a cross-sectional study and longitudinal studies are warranted to 281 assess the impact of exposome on lung function trajectories. Fourth, the sBPR method, that 282 identifies the best partition of a population without controlling for the size of the clusters, 283 resulted in a cluster with few individuals ($n = 30, 5\%$ of the population), consistent with 284 previous studies that applied this method (Coker et al., 2017). Interestingly, this specific cluster 285 identified a subpopulation with a specific co-exposure profile associated with a mean $FEV₁%$ 286 10 points lower in average than individuals from the other clusters, therefore a restricted 287 population that could benefit from a targeted reinforced surveillance. Fifth, our population was 288 essentially composed of Caucasian individuals living in France which is a developed country. 289 Therefore, our results cannot be generalized to individuals from developing countries or 290 individuals of other ethnic groups. Finally, we acknowledge that a replication study in an 291 independent cohort would be particularly relevant, but we were unable to identify a cohort of 292 adults with asthma, in which both similar information on a large set of exposures and the 293 measure of lung function were available.

294 *Clinical and public health perspectives*

295 While previous studies identified factors individually associated with lung function in adults 296 with asthma, here we demonstrated that identifying comprehensive profiles of social, lifestyle 297 and environmental exposure, whose effects on lung function are likely to interact, can help to 298 get better insight into the etiology of multifactorial health outcomes. In terms of public health, 299 our results support that complementary approaches to simple public health messages focusing 300 on a unique exposure to prevent health issues at the general population level (no or quit 301 smoking, daily consumption of 5 fruits or vegetables, …) are warranted. Approaches aimed at 302 identifying restricted subpopulations at very high risk of disease development (or worsening) 303 given their exposure profile could provide further policy relevant information. From clinical 304 perspectives, methods for more systematic assessment of exposome, including questionnaires 305 and digital health tools (apps) need to be developed and validated to be deployed in routine 306 asthma care (Bédard et al., 2019). Then, practitioner could identify patients belonging to co-307 exposure profile associated with higher risk of low lung function and adapt accordingly 308 healthcare management. Secondary and tertiary preventions in this particular subpopulation 309 could help to limit costs related to their disease.

310

311 **Conclusion**

312 This exposome study identified a subpopulation of adults with asthma with a specific profile of 313 joint environmental and lifestyle factors associated with a strong decrease of FEV1. By 314 characterizing the joint effect of multiple lifestyle and environmental risk factors in lung 315 function, the present study goes a step further in the understanding of the complex role of 316 environment in asthma and led to the identification of a specific asthma subgroup that could 317 benefit from a targeted reinforced surveillance. A such comprehensive exposome approach, 318 easily transposable to any multifactorial disease, is expected to help unravelling the impact of 319 environment on health.

320

321

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Figure captions

Fig. 1 Workflow of the study

Fig. 2 Flowchart of the population

Fig. 3 Heatmap of the between-exposure correlations

Fig. 4 Heatmap of the absolute within- and between-families of exposure correlations *The diagonal of the heat map assesses the median of the 2-way absolute correlations between exposures variables belonging to the same family (within-family correlations) while the offdiagonal assesses the median of the absolute correlation between each exposure variable of two different family of exposures (between-family correlations)*.

Variable	Number of individuals with missing values	Range $(min - max)$	Mean \pm SD or n (%)
Age (years)	θ	$16 - 79$	$39 + 16$
Height (cm)	$\mathbf{0}$	145 - 192	172 ± 9
Gender	$\boldsymbol{0}$		
Female, n $(\%)$			288 (48)
Male, n $%$			311(52)
Socio-professional category	3		
No profession, n $(\%)$			84 (14)
Executive, n $%$			189 (32)
Technician, n (%)			254 (42)
Farmer/Labourer, n (%)			72(12)
Centre	$\boldsymbol{0}$		
Paris, n $(\%)$			184 (31)
Lyon, n $(\%)$			94 (16)
Marseille, n $(\%)$			89 (15)
Montpellier, n $(\%)$			47(8)
Grenoble, n $(\%)$			185(31)
SD: standard deviation			

Table 1. Main characteristics of the studied population (n=599)

SD: standard deviation ; AHEI: Alternative Healthy Eating Index; METs: Metabolic Equivalent of Tasks ; NO₂: nitrogen dioxide ; PM_{2.5}: particles matters with aerodynamical diameter less than 2.5 µm; PM₁₀: particles matters with aerodynamical diameter less than 10 µm; O₃: ozone; NDVI: normalized difference vegetation index; UV: ultraviolet *: Daily values averaged on the 365 days before performing the spirometry; \dagger : index of area composition and richness; \ddagger : number of health services in the IRIS ("Ilots") Regroupés pour l'Information Statistique") where an IRIS represents an area covering between 1,800 and 5,000 inhabitants.

γ .	Cluster 1	Cluster 2	Cluster 3		
	$n = 30$	$n = 115$	$n = 454$	\mathbf{p}	p adjusted ^{\ddagger}
Age (years)	$51 \pm 14^{\dagger}$	$48 \pm 13^{+}$	36 ± 16	< 0.001	$\overline{}$
Height (cm)	170 ± 10	169 ± 8	170 ± 9	0.52	0.68
Gender				0.12	
Female, $n\frac{6}{6}$	9(30)	55 (48)	224 (49)		
Male, $n\frac{6}{6}$	21(70)	60(52)	230(52)		
Socio-professional category				< 0.001	0.22
No profession, $n\frac{0}{0}$	0(0)	$2(2)^{+}$	82(18)		
Executive, $n\frac{0}{0}$	10(33)	43 (37) [†]	136(30)		
Technician, n(%)	17(57)	49 (43) [†]	188(41)		
Farmer/Labourer, n(%)	3(10)	$21(18)$ [†]	48(11)		
Centre				< 0.001	
Paris, $n\frac{6}{6}$	$16(53)$ ^{*†}	$20(17)^{+}$	148 (33)		
Lyon, $n\frac{6}{6}$	$8(27)$ ^{*†}	$25(22)^{+}$	61(13)		
Marseille, n(%)	$4(13)^{*^+}$	22 (19) [†]	63(14)		
Montpellier, $n\frac{6}{6}$	$1(3)^{*}$	$12(10)^{+}$	34(7)		
Grenoble, n(%)	$1(3)^{*}$	$36(31)$ [†]	148 (33)		
Season of examination				0.23	0.17
Spring, $n\frac{6}{6}$	13(43)	33(29)	108(24)		
Summer, $n\frac{6}{6}$	6(20)	25(22)	107(24)		
Automn, $n\frac{6}{6}$	8(27)	28 (24)	132(29)		
Winter, $n\frac{6}{6}$	3(10)	29(25)	107(24)		
Number of pack-years	$23 \pm 28^{\dagger}$	17 ± 10^{4}	1 ± 2	< 0.001	< 0.001
Passive smoking				0.005	0.17
None, $n\frac{6}{6}$	$18(60)^{\dagger}$	45 (39) [†]	194(43)		

Table 3. Sociodemographic characteristics and exposures of the 3 clusters obtained by Bayesian Profile Regression supervised on FEV1 (%predicted).

FEV1: Forced expiratory volume in 1s; FVC: forced vital capacity; AHEI: Alternative Healthy Eating Index; FDep: French Deprivation Index. *: $p < 0.05$ vs cluster 2;[†]: $p < 0.05$ vs cluster 3;[‡]p-value adjusted on age, gender and centre

Table 4. Phenotypic characteristics at EGEA2 and EGEA3 according to the 3 environmental clusters obtained by the supervised Bayesian Profile Regression

*: $p < 0.05$ vs cluster 2; $\dot{p} < 0.05$ vs cluster 3; \dot{p} : exacerbation was defined by use of oral steroids or hospitalization or emergency visit for asthma in the last 12 months; §: current asthma defined by the presence of >1 asthma attacks or use of asthma treatment in the past 12 months; ¹¹: Asthma symptoms score as defined by Sunyer et al [38] by the sum of five symptoms over the past 12 months (wheezing or whistling in the chest, woken up with a feeling of chest tightness, attack of shortness of breath at rest, attack of shortness of breath that came on following strenuous activity, woken by an attack of shortness of breath; **: allergic sensitization was defined by >1 positive skin prick test response to any of the 12 allergens (cat, *Dermatophagoides pteronyssinus, Blattela germanica*, olive, birch, *Parietaria judaica*, timothy grass, ragweed pollen, *Aspergillus species*, *Cladosporium herbarum, Alternaria tenuis*, and cypress)

Figure 1.

Figure 2.

Figure 3.

Figure 4.

Graphical abstract

FE' ($predictf$)

Online Data Supplement

Profile of exposures and lung function in adults with asthma: an exposome approach in the EGEA study

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Online methods

Part I. Study population and spirometry

Study population

EGEA is a French cohort based on three surveys over 20 years. Initially (EGEA1 study), 388 cases with asthma were recruited in the early 1990s in five cities (Paris, Lyon, Marseille, Montpellier and Grenoble), so as their first-degree relatives (n=1244) and 415 population-based controls. Two follow-ups were conducted in 2003-2007 (EGEA2) and 2011-2013 (EGEA3). At each survey, individuals responded to a detailed questionnaire on respiratory health and environment, based on international standardised tools, as previously described (Kauffmann et al., 1997; Kauffmann and Dizier, 1995). The EGEA collection and program management was certified ISO 9001 from 2006 to 2018.

Because part of geographic information systems (GIS)-based models did not fit for the EGEA1 study period and spirometry was not part of EGEA3 protocol, the present study focuses on adults with ever asthma at EGEA2. Ever-asthma was defined by a positive answer to either "Have you ever had attacks of breathlessness at rest with wheezing?" or "Have you ever had asthma attacks?" or being recruited as an asthma case in chest clinics.

Spirometry testing

Spirometry was performed using a flow-volume spirometer (SpiroDyn'R; SAS Dyn'R, Aix-en-Provence, France) according to the American Thoracic Society and European Respiratory Society guidelines (Miller et al., 2005). Standardized operating procedures were implemented and controlled, including calibration of all equipment before each measurement, and quality control visits by study coordinators to ensure correct protocols by field staff. Participants were asked to perform three to nine attempts to provide at least three technically acceptable spirometry manoeuvres. Pre-bronchodilator forced expiratory volume in 1 s (FEV_1) was expressed in percentage of the predicted value using the Global Lung Function Initiative (GLI-2012) reference equations, thereafter referred as FEV1% (Quanjer et al., 2012).

Part II. Exposome assessment method

In total, 53 exposure variables (19 lifestyle or social factors and 34 environmental factors) were assessed.

Lifestyle and social factors

Data regarding behavioural factors were assessed mostly through self-completed questionnaires and provided information about 9 families of exposure: indoor air (3 variables), tobacco exposure (4 variables), season of examination (season at which individual performed the spirometry, 1 variable), pets (2 variables), occupational exposure (2 variables, assessed through a job-exposure matrix), household cleaning products (2 variables), rurality (2 variables), socioeconomic status (SES) (assessed at the contextual level by the French Deprivation Index (FDep) from geocoded residential address, built to capture health inequalities at ecological level (Temam et al., 2017), 1 variable), and physical activity and diet (2 variables, assessed through the Metabolic Equivalent Tasks (METs) and the 2010 Alternative Healthy Eating Index (AHEI-2010) score, respectively).

Indoor air

Three variables were collected to characterise indoor air. **Water damage in the last 12 months** was defined by a positive answer to the question: "Has there been water damage in your building in the last 12 months". Presence of **moulds in the bedroom/living-room/bathroom** was defined by positive answers to the questions: "Has there ever been any mould on any surface inside the house (food excluded)?" and "has this happened in the last 12 months?" and if the room was bedroom, living-room and/or bathroom. **Use of gaz for cooking** was defined by the answer "gas" to the question: "Currently, what is the fuel used for cooking?".

Tobacco

Detailed information on smoking history was collected, included the number of cigarettes/pipe/cigars/cigarillos smoked per day for different time periods if the consumption of tobacco smoking has changed over time. From these data the number of **pack-years** (as a quantitative and divided in 3 classes (0; >0-10; >10-20; >20) was calculated and the **smoking status** was defined: never smokers were defined as those never smoking or having smoked on average less than one cigarette, cigar, cigarillos or one pipe a day for a year. Former smokers had stopped smoking since at least a month. No exposure to **passive smoking** was defined as exposure to tobacco smoke for less than one hour a day (all sources of exposure combined), intermediate exposure was defined as reporting exposure to tobacco smoke at home or work less than one hour a day and high exposure was defined as being exposed >1h a day (all sources of exposure combined).

Season

Season of examination was assessed from the date at which individuals performed spirometry and defined in 4 categories: spring (March $20th$ to June $20th$), summer (June $21th$ to September $22th$), autumn (September $23th$ to December $21th$) and winter (December $22th$ and March 19th).

Pets

Having cat/dog currently was defined by at least one positive answer to the following questions: "Do you have a cat?" and "Do you have a dog?". **Having cat/dog in childhood** was defined by a positive answer to the question: "In childhood, did you have a pet?" and have chosen the answer "cat" or "dog" to the question: "Was it a dog? A cat? A rodent? A bird? Other?".

Occupational exposure to asthmagens

Occupational exposures to 18 asthmagens (high or low molecular weight agents, mixed environment) and to low risk agents (possible low level exposure to irritants or antigens) were evaluated at EGEA2 through an asthma-specific Job-Exposure Matrix (Kennedy, 2000) for both the last job and all occupational history. In addition, an expert re-evaluation step was performed to improve occupational exposure evaluation (Dumas et al., 2011).

Cleaning products

The variable **use of bleach** was defined according to the answer at the question: "how many days a week did you use bleach?" where the possible answers were: "never; <1 day/week; 1-3 days/week; 4-7 days/week". For the present analysis, this variable was recorded in the three following categories: "<1 day/week; 1-3 days/week; 4-7 days/week" (Matulonga et al., 2016). **Use of sprays** was defined by the exposure to any of the eight types of sprays (furniture, glasscleaning, carpet, mopping the floor, oven, ironing, air-refreshing, other use) and was composed of three categories: "none; 1/week; ≥2/week" (Le Moual et al., 2012).

Rurality

Be **born in a rural municipality** was defined as those reporting having lived in a municipality of less than 5000 inhabitants at birth.

Using data from the General Agricultural Census (GAC), the commune at birth was categorised as rural when the number of inhabitants was lower than 5,000 (to have a sensitive definition of rural).

Ever-living in a rural municipality was defined as those reporting having lived in a municipality of less than 5000 inhabitants during their life. These variables were calculated from the zip code of each municipality the participants reported to have lived since birth. Social

The **French Deprivation Index** (FDep), defined at the geographical IRIS level (division of the French territory of homogeneous size defined by 2000 inhabitants per unit), integrates four dimensions: the median income per unit of consumption in the household, the percentage of graduates in the population over 15, the percentage of workers in the labor force and the unemployment rate. Finally, this index was divided into 5 quintiles from a large french cohort (E3N: Etude épidémiologique auprès des femmes de la Mutuelle Générale de l'Education Nationale) (Temam et al., 2017).

Physical activity and diet

Dietary data were collected at EGEA2 using a 118-item self-administered food frequency questionnaire (FFQ), which is based on a validated French questionnaire (Bonifacj et al., 1997). Participants were asked to indicate their frequency of consumption of all food items by checking one of eight given categories, including "never or ≤ 1 time/month", "1-3

times/month", "1 time/week", "2–4 times/week", "5–6 times/week", "1 time/day" 2–3 times/day", and "≥ 4 times/day", for a standard portion size of each item. Standard portion sizes were listed for each food. The selected frequency category for each food item was converted to a daily intake. For example, a response of "1 serving/week" was converted to 0.14 servings/d. Through the use of the French food-composition data from the Vitamin and Antioxidant Element Supplementation (SU.VI.MAX) survey (Hercberg et al., 1998), the average daily intake of nutrients was calculated by multiplying the frequency of consumption of each item by its nutrient content per serving and totaling the nutrient intake for all food items. Out of the 1,601 participants from EGEA2, 35 adolescents (less than 16 years-old) were firstly excluded (no FFO in adolescents). Among adults ($n = 1,566$), we excluded 109 participants who did not fill the FFQ, 60 who left more than 4 items blank on the diet questionnaire, and 161 participants with an implausibly high ($> 4,200$ kcal/d for men; $> 3,500$ kcal/d for women) or low total ($<$ 800 energy intake kcal/d for men; < 500 kcal/d for women) (Varraso et al., 2012). In total, dietary data were available for 1,236 adults at EGEA2.

The scoring criteria of the **Alternate Healthy Eating Index 2010 (AHEI-2010)** was described in detail elsewhere (Chiuve et al., 2012). Briefly, the AHEI-2010 score includes 11 components. A high AHEI-2010 score denotes high intake of vegetables, fruits, whole grains, nuts and legumes, long-chain Ω-3 fatty acids and polyunsaturated fatty acids; moderate intake of alcohol; and low intake of sugar-sweetened drinks and fruit juice, red/processed meat, *trans* fat, and sodium. For each component, a minimal score of 0 and a maximal score of 10 was allocated based on its intake, and intermediate values scored proportionally. The total AHEI-2010 score sums scores of all the 11 components, ranging from 0 to 110; a higher score represents a healthier diet. In the EGEA study, we calculated a modified AHEI-2010 including all the items except for *trans* fat, which is not available in the French food composition table from the SU.VI.MAX study. Besides, cereal fiber intake was calculated instead of whole grain intake, because only intakes of whole wheat bread and of brow rice have been recorded as whole grains in the EGEA2 study. An intake of 15 g cereal fiber/day was considered as ideal based on the scoring criteria of the Alternate Healthy Eating Index (AHEI). Therefore, the highest value for the AHEI-2010 diet score in the EGEA study was 100 (rather than 110) (Li et al., 2017).

Assessment of habitual physical activity at EGEA2 was based on five questions on the usual distance walked daily (<500, [500-2,000], and \geq 2,000 m), the average number of flights of stairs climbed daily (0, [1-4], and \geq 5), the average amount of time spent weekly doing vigorous recreational activity (such as jogging, aerobic, swimming, cycling, tennis, soccer, digging in the garden) (0, [1-2], [3-4] and \geq 5 hours), moderate recreational activity (such as cleaning, walking, golf, slight gardening) (0, [1-2], [3-4] and ≥5 hours) and light recreational activity (such as reading, watching TV, office work, driving, eating) (0, [1-2], [3-4] and \geq 5 hours), based on the questionnaire from Paffenbarger *et al.* (Paffenbarger et al., 1993). Physical habits at work (at work, you are seated most of the day; at work, you are standing, but your activity does not require any particular physical effort; at work, your activity requires physical efforts (lifting heavy loads, etc.) was also recorded based on the questionnaire from Baecke *et al.* (Baecke et al., 1982). Total physical activity was estimated by multiplying the metabolic equivalent cost (MET) of each activities according to the Compendium of Physical Activities (Ainsworth et al., 2000) by their frequency and duration, and assessed in **metabolic equivalents per week (METs/w**).

Environmental exposures

A broad range of environmental exposures was assessed through geographic information systems (GIS) that allow allocating a spatio-temporal estimate of exposure from geographical residential coordinates. Environmental exposures covered 8 families: altitude (1 variable), building and population densities (3 variables), atmospheric pollutants (4 variables), natural spaces (8 exposures), weather (6 variables), roads/cycle paths (3 variables), ambient ultra violet radiation (UV) (3 variables) and health services (6 variables). The GIS-based models are resting on National Geographic Institute (IGN) maps, the French National Institute for Statistics and Economic Research (INSEE) data, Météo-France data, geospatial models, land cover databases and satellite data. All these exposures were calculated from the geocoded residential address. Altitude

Altitude (m) at the residential address of each participant was assessed from the ALTI database from IGN ("Institut de l'Information Géographique et Forestière"). Data were available for the year 2016 and had a spatial resolution of 25m.

Building and population densities

Building density was calculated within **100- and 300-meters buffers** by dividing the area of building cover (km^2) by the area of each buffer (km^2). The area of building cover was estimated from the 2016 data of the BDTOPO database (IGN).

Population density around residential address was calculated as the number of inhabitants in an area of **200*200m** surrounding the home address with the INSEE ("Institut Nationale de la Statistique et des Etudes Economiques") database.

Atmospheric pollutants

Chronic exposure to four atmospheric pollutants (nitrogen dioxide $(NO_2, g/m^3)$, particulate matter with an aerodynamic diameter of less than 2.5 μ m (PM_{2.5}, g/m³) and of less than 10 μ m $(PM_{10}, g/m^3)$, and ozone $(O_3, g/m^3)$) was assessed using the "GAZELAIR" model (Bentayeb et al., 2014). This model was defined through a geostatistical (Kriging) analysis, including data from the CHIMERE chemistry-transport model, a validated model in Europe (Alonso et al., 2011; Valari et al., 2011) and provides annual data from 1989 to 2008. The 12-month average levels of air pollutants at the participant's residential address before the lung function test was estimated.

Natural spaces

Number of blue spaces in a 300m buffer, number of green spaces in a 300m buffer, distance to nearest blue space, and the distance to nearest green space were assessed using the 2006 Corine Land Cover (CLC) database (https://www.geoportail.gouv.fr/donnees/corineland-cover-2006). This database provides information about land use in French territory, with a spatial resolution of 100m. Green spaces were defined as "green urban areas" (class 1.4.1) and blue spaces as "water bodies" (class 5). Distances from the nearest blue/green spaces were defined as the distance between residential address and nearest blue/green space identified in

the CLC database. Numbers of blue/green spaces in a 300m buffer were defined as the number of object of classes 5/1.4.1 in a 300m buffer surrounding residential address.

The **Shannon Evenness Index (SEI) in a buffer of 300m** was also calculated from the CLC database. The SEI is the degree of mixing of different types of land uses (such as residential, commercial, entertainment, and office development). A higher value indicates a more even distribution of land between the different types of land uses. This index was calculated as follow:

$$
\frac{-\sum_{i=1}^{m}(p_i * ln\ p_i)}{\ln m}
$$

with:

 p_i : = proportion of the area occupied by land use type (class) i

m = number of land use types (classes) present in the study area

The SEI was calculated in a 300m buffer around residential address.

Normalized Difference Vegetation Index (NDVI) quantifies vegetation by measuring the difference between near-infrared (which vegetation strongly reflects) and red light (which vegetation absorbs). NDVI values range from +1.0 to -1.0. Areas of barren rock, sand, or snow usually show very low NDVI values (for example, 0.1 or less). Sparse vegetation such as shrubs and grasslands or senescing crops may result in moderate NDVI values (approximately 0.2 to 0.5). High NDVI values (approximately 0.6 to 0.9) correspond to dense vegetation such as that found in temperate and tropical forests or crops at their peak growth stage. Negative values of NDVI (values approaching -1) correspond to water.

NDVI in a **100m, 300m and 500m buffers** were derived from the LANDSAT data. The original data set had a spatial resolution of 30m and provided daily values of NDVI from 1986 to 2017. We calculated the NDVI as the average of daily values from June to August in the year at which individuals performed spirometry.

Weather

Wind (m/s) and **relative humidity** (%) were calculated from the SAFRAN model (Météo-France). This model had a spatial resolution of 8km and provided daily values. The daily values of the 365 days preceding the day at which individuals performed spirometry were averaged to obtain the mean annual value.

Absolute humidity (g/m^3) was calculated as follow:

wvp_mb = $6.11* (10**(((7.5 * dpt24h)/(237.7 + dpt24h))))$ ah gm3 = $(1000*(100 * wvp mb))/((273.15 + tempc24h)* 461.5)$

where dpt24h is the dewpoint 24h and tempc24h was the average temperature in the last 24h $(in °C)$

Minimal, maximal and mean daily **temperature** were estimated by an updated version (Hough *et al.*, submitted) of a hybrid spatio-temporal model based on Moderate Resolution Imaging Spectroradiometer (MODIS) satellite surface temperature data (Kloog et al., 2017). Briefly, daily satellite surface temperature data (in 1 km2 grid cells) were obtained, and calibrated with air temperature data from Météo-France monitors, with adjustment for spatiotemporal predictors (NDVI, altitude, etc). These daily model predictions were used to generate temperature exposure estimates for every study participant, based on his home address.

Roads/cycle paths

Distance to nearest major road, number of major roads in a 100m buffer and **number of cycle paths in a 100m buffer** were derived from the BDTOPO database of IGN (http://professionnels.ign.fr/bdtopo). Major roads comprised "highways", "nearly-highways", "highway ramp" and "dual carriageway". BDTopo data were derived in 2016.

Ambient ultra‐violet (UV) radiation

Daily, weekly and monthly measurements of UV radiation (as **erythemal UV, Vitamin-D and DNA damaging UV**) at home at 0.5 x 0.5 degree resolution were derived from total ozone column measurements by the Global Ozone Monitoring Experiment-2 (GOME-2) instrument onboard the Meteorological Operational satellite program (MetOp) and SCIAMACHY instrument onboard ENVISAT satellite, assimilated into global ozone fields (Temis project, http://www.temis.nl/uvradiation/UVindex.html), and averaged over the year before the spirometry examination.

Health services

Health services information were derived from INSEE data: (https://www.insee.fr/fr/statistiques/3568611?sommaire=3568656). Data were collected in 2017 at the IRIS ("Ilots Regroupés pour l'Information Statistique") scale, where an IRIS is composed of nearly 5,000 inhabitants (https://www.insee.fr/fr/metadonnees/ definition/c1523). This database provided information regarding 42 types of health services. We individually considered presence of "**Medical practitioner**", "**pulmonogist**" and "**pharmacy**" in the IRIS while we grouped other health services in 3 classes. "**Other specialists"** was defined as the sum of cardiologists, dermatologists/venerelogists, gastroenterologists hepatologists, psychiatrists, ophthalmologists, otolaryngologists, paediatricians, radiologists, stomatologists, gynaecologists and dental surgeons in the IRIS. "**Paramedics"** were defined as the sum of midwives, nurses, physiotherapists, speech therapists, orthoptists, chiropodists, orthosist, occupational therapist, psychomotor therapist, dietician and psychologists in the IRIS. "**Other health services"** was defined as the sum of health care organization for short, medium and long stays, psychiatric hospitals, cancer centres, emergency rooms, maternities, health centres, psychiatric outpatient clinics, centre for preventive medicine, dialysis clinic, hospitalization at home, multidisciplinary health centre, medical analysis laboratories, ambulances, blood transfusion centre and thermal buildings.

Part III. Statistical analysis

Transformation of exposure variables

For each exposure variable X, eight histograms were plotted: X, $log_{10}(X)$, $log_2(X)$, $1/X^2$, $1/X$, $1/\sqrt{X}$, \sqrt{X} and X^2 . The best transformation to achieve normality was chosen on the basis of these graphs and, if no transformation worked, the exposure variable was dichotomized (where the cut-off was the median value). Transformation was chosen to obtain reasonably symmetric distributions and, if possible, the same transformation was used for all exposure variables from the same exposure family.

Imputation of missing data

Missing values of exposures and cofounders (median (IQR) missing values = 0 (0-3)) were imputed using the method of chained equations (van Buuren and Groothuis-Oudshoorn, 2011; White et al., 2011) and 5 complete datasets were generated (Stuart et al., 2009). The imputation model included outcomes variables, centre of recruitment and all exposure variables with an absolute correlation greater than 0.2 with the exposure variable to be imputed. Predictive mean matching was used to impute continuous variables and logistic and multinomial regressions were used to impute binary and categorical exposures, respectively. M=5 complete datasets were created. Imputed missing values several times allows to account for uncertainty in the final standard errors, confidence intervals and p-value (Stuart et al., 2009).

Standardisation of exposure variables

Each quantitative transformed exposure variable was standardised by the interquartile range so as to allow between-covariate comparison of regression estimates. In that order, each transformed variable was divided by its $75th$ percentile (3rd quartile) minus 25th percentile (1st) quartile).

Exposome description

All exposure variables were described as mean (standard deviation) or n (%) for quantitative and qualitative variables, respectively. A first heat map presented correlations between exposures. They were assessed through Pearson coefficient for two quantitative variables, polychoric coefficient for two qualitative variables and polyserial coefficient for a quantitative and a qualitative variable. A second heat map assessed absolute within (diagonal) and between (off-diagonal) families of exposure median correlations.

Exposome dimension reduction from ExWAS results

This step aimed to restrict the number of exposure variables to be included in the supervised Bayesian Profile Regression (sBPR) method to allow a good model stability. In that order, an Exposome-Wide Association Analysis (ExWAS) was performed to identify exposures individually associated with $FEV₁%$. This method consists in fitting independent mixed linear regressions models for each exposure variable, with adjustment for the centre of inclusion, gender, age, height and socio-professional category as fixed effects and family as random effect. The Rubin's rule was used to combine the estimates from each of the 5 imputed datasets into one overall estimate together with an associated variance that incorporates both the within

and between imputation variability (Marshall et al., 2009; Rubin, 2004)[25,26]. Factors showing suggestive association with $FEV1\%$ (p-value ≤ 0.2) were selected in the reduced set of exposures. In addition, to address the significance of each factor independently, the number of independently effective tests was calculated with the Li formula (Li et al., 2012) applied to the correlation matrix of exposures variables and p-values of the association tests were compared to the significance threshold of 0.05 divided by this number (i.e. $1.71*10^{-3}$) to correct for multiple hypothesis testing.

Identification of profiles of joint exposures associated with FEV1% using sBPR

The second step consisted in identifying profiles of co-exposures associated with low lung function. In that order, a supervised cluster analysis, *i.e.* simultaneously considering in the model the outcome and exposures, was conducted using the sBPR clustering method (Molitor et al., 2010). This model is fitted using Markov Chain Monte Carlo sampling methods and outputs a different clustering of the data at each iteration of the sampler, thus coherently propagating uncertainty (Gilks et al., 1998). Here, the algorithm was run for 200,000 iterations. The sBPR was applied in the reduced set of exposures identified in step 1 by the ExWAS, FEV1%, and included variables used as confounders in the ExWAS analysis (age, gender, height, centre of inclusion, number of smoked pack-years and socio-professional category) as covariates. When two exposures variables with a p-value < 0.20 in the ExWAS analysis had an absolute correlation coefficient > 0.70 (Ratner, 2009), only the variable with the lowest p-value in the ExWAS analysis was retained. This step allowed us to obtain well-characterized clusters with specific profiles of exposure. Socio-demographic characteristics and exposure factors were compared between clusters to identify the factors mainly driving the clustering. This analysis was also performed after adjustment for age, gender and centre to identify exposures associated with clusters independently of these 3 variables. Moreover, phenotypic characteristics at EGEA2, but also at EGEA3 about 7 years later, were compared between clusters.

As this method relies on a stochastic process, results may not be stable (final results depend from the initialization step). To test the stability of our final model, we ran it with 10 different seeds (all with 200,000 iterations). All these 10 seeds identified 3 clusters and the median [interquartile range] number of subjects remaining in the same cluster, independently from the seed, was 0.86 [0.73-0.94] (data not shown).

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Supplemental Tables

Supplemental Table E1. Characteristics of included and excluded individuals with asthma.

Supplemental Table E2. Comparisons of exposures by inclusion centre

AHEI: Alternative Healthy Eating Index; METs: Metabolic Equivalent of Tasks; NO₂: nitrogen dioxide; PM_{2.5}: particles matters with aerodynamical diameter less than 2.5 μ m; PM₁₀: particles matters with aerodynamical diameter less than 10 μ m; O₃: ozone; NDVI: normalized difference vegetation index; UV: ultraviolet

*: Daily values averaged on the 365 days before before performing the spirometry; \dagger : index of area composition and richness; \dagger : number of health services in the IRIS ("Ilots") Regroupés pour l'Information Statistique") where an IRIS represents an area covering between 1,800 and 5,000 inhabitants.

Supplemental Table E3. Results of the ExWAS analyses assessing associations between each exposure variable and $FEV₁(\%$ predicted)

All linear regressions were adjusted for age, gender, height, socioprofessional category, inclusion centre as fixed effects and familly as random effect.

ExWAS: Exposome-Wide Association Study ; FEV₁: forced expiratory volume in 1s ; sd: standard deviation ; AHEI: Alternative Healthy Eating Index; METs: Metabolic Equivalent of Tasks ; NO2: nitrogen dioxide ; $PM_{2.5}$: particles matters with aerodynamical diameter less than 2.5 μ m; PM₁₀: particles matters with aerodynamical diameter less than 10 µm; O3: ozone; NDVI: normalized difference vegetation index; UV: ultraviolet

*: Daily values averaged on the 365 days before performing the spirometry; \dagger : index of area composition and richness ; [‡]: number of health services in the IRIS ("Ilots Regroupés pour l'Information Statistique") where an IRIS represents an area covering between 1,800 and 5,000 inhabitants

 $\ddot{\cdot}$: exacerbation was defined by use of oral steroids or hospitalization for asthma or emergency visit for asthma in the last 12 months; FEV₁: forced expiratory volume in 1 second

Online Figures

Supplemental Figure E1. Volcano plot of the coefficient estimates for exposure variables versus p value in the ExWAS analysis of the exposures-FEV₁ (%predicted) association.

The solid line corresponds to a p-value of 0.20, the dashed line corresponds to a p-value of 0.05 (alpha) and the dotted line corresponds to a p-value of $1.71*10⁻³$ (alpha corrected for multiple comparisons)