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# **Supporting Information**

- 2 Poor Regulation Implications in a Low and Middle
- 3 Income Country Based on PAHs Source
- <sup>4</sup> Apportionment and Cancer Risk Assessment

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#### 20 Source Apportionment using Positive Matrix Factorization

21 This model works by solving the chemical mass balance (CMB) equation:

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$$X_{ij} = \sum_{k=1}^{p} g_{ik} f_{kj} + e_{ij}$$
(Eq.1)

where,  $X_{ij}$  refers to the concentration for the  $j_{th}$  specie in the  $i_{th}$  sample; p represents the number of factors;  $g_{ik}$  is the relative contribution of factor k to  $i_{th}$  sample;  $f_{kj}$  refers to the profile factor of each source for the  $j_{th}$  specie; and  $e_{ij}$  is the PMF residual error for the  $i_{th}$  sample and the  $j_{th}$  species not fitted by the model.<sup>25</sup>

An optimal source apportionment analysis is accomplished when the minimal value of  $Q_{robust}$ (Eq.2) for a given p is found.  $Q_{robust}$  is the goodness of fit parameter calculated excluding outliers. However,  $Q_{true}$  is the goodness of fit parameter calculated including all collected data. For a stable and reliable solution,  $Q_{robust}$  and  $Q_{true}$  should be comparable with a ratio of  $Q_{true}/Q_{robust}$  less than 1.5, meaning that the outliers are not affecting the modeling of the system.<sup>23</sup>

32 
$$Q = \sum_{i=1}^{n} \sum_{j=0}^{m} {\binom{e_{ij}}{u_{ij}}}^2$$
(Eq.2)

33 Here, n refers to the number of samples and m to the number of species; and  $u_{ij}$  refers to the 34 uncertainty of the measured concentration for the  $i_{th}$  sample and the  $j_{th}$  species.<sup>27</sup>

To evaluate the robustness solution for the system, additional steps are done such as signal-tonoise ratio, base model, bootstrap (BS), displacement (DISP), and bootstrap-displacement (BS-DISP) that are offered by the model, further explanation about these steps are found in the supporting information. 39 In this study, 55 samples of 16 PAHs were placed in the EPA PMF 5.0 model with their40 uncertainties to estimate the different source contributions to the measured PAHs.

41 Measurement uncertainty  $(u_{ij})$  is very critical to determine the optimal solution to the analytical 42 equation by minimizing the function Q. In the absence of samples replicates, the 43 analytical uncertainties cannot be calculated, however uncertainties can be estimated based on 44 uncertainty equations.

45 Four of the most commonly used equations (Table 1) were tested to identify the best uncertainties46 to our study.

PMF Uncertainty (u <sub>ij</sub> )	References
	20
$\sigma ij = 0.05 \times Xij + Dij$	20
$U_{ij} = 0.1N_{ij} + \frac{D_{ij}}{3}$	29
$\sqrt{(Error Fraction \times Conc.)^2 + MDL^2}$	30
$\sigma i j = C1(Nij + Nj)$ then, $si j = \sigma i j + C3Ni j$	31, 32

47 Table S1: Methods for calculating uncertainties for PMF analyses

48

49 Where:  $X_{ij}$  and  $N_{ij}$  are the concentrations of the species;  $D_{ij}$  and MDL are the detection limits;  $C_1$ 50 and  $C_3$  are constants between 0.01-0.05 and 0.1-0.5 respectively;  $\bar{N}j$  is the average concentration 51 of species.

52 Each equation was tested based on the five essential steps, explained in the supporting information,

53 to determine which uncertainty calculation compliments the concentrations.  $\sigma i j = C_1(N_{ij} + \bar{N}j)$  then,

54  $s_{ij}=\sigma_{ij}+C_3N_{ij}$  showed the best fit for our data, with  $C_1$  and  $C_3$  being 0.02 and 0.2 respectively, which 55 were determined by trial-and-error tests.

To evaluate the solution for the system, five essential steps are required: Signal-to-noise ratio, base model, bootstrap, displacement, and bootstrap-displacement. The signal-to-noise ratio indicates if the measurements are above or within the noise (detection limit) of the data. The model categorizes the signals as "strong" if S/N > 2, "weak" if  $0.2 \le S/N \le 2$ , and "bad" if S/N < 0.2.<sup>1</sup> If a species is classified as weak, then it is down weighted by the PMF model, and if a species is classified as bad, it is excluded from the PMF analysis.

62 The base model run determines the  $Q_{true}$  and  $Q_{robust}$ . The ratio of  $Q_{true}/Q_{robust}$  should be smaller than 63 1.5 to indicate that the outliers are not affecting the modeling of the system.<sup>2</sup> Another output of 64 the base model run is the residual analysis, which is the degree of adjustment of each species by 65 the model. An ideal result is when the residual analysis is normally distributed between ±3 for 66 each species.<sup>3</sup>

67 The bootstrap (BS) calculation validates the base model by predicting the results using the 68 developed model. A model represents a good fit if the bootstrap analysis shows values of 70-75 69 and higher for each factor.<sup>2</sup>

The displacement (DISP) runs the solution infinite times to give the maximum variation in  $Q_{robust}$ (dQmax). A model is considered a good fit if the displacement analysis returns a dQmax less than 1% of the initial  $Q_{robust}$ .<sup>3</sup>

73 The last step to be discussed is the bootstrap-displacement (BS-DISP) run. A combination of the 74 bootstrap and displacement analysis evaluates the effect of random errors and returns with a 75 decrease in  $Q_{\text{robust}}$ . If the decrease in  $Q_{\text{robust}}$  is less than 0.5% then the model is a good fit and the 76 analyses can be continued.<sup>3</sup>

#### 77 Number of Factors from PMF model

78 Using Eq. 4 from Table 1 with  $C_1$  and  $C_3$  determined as 0.02 and 0.2, respectively, 2 to 6 factors

79 were tested to identify the number of sources that contributed to PAHs using PMF version 5.0.

80 The optimal number of three factors was chosen based on the signal-to-noise ratio, Qtrue/Qrobust

81 ratio of 1, scaled residuals normally distributed between  $\pm 3$ , BS of 96%, 96%, and 92%, DISP and

82 BS-DISP decrease in dQmax by less than 1% and 0.5%, respectively.

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### 84 Cancer Risk Calculation

#### 85 Table S2: Toxicity Equivalence Factor (TEF) for each PAH

PAHs	TEF <sup>4</sup>	Carcinogenic Group (IARC
		2018)
Naphthalene (Nap)	0.001	2B
Acenaphthylene (Acy)	0.001	-
Acenaphtene (Ace)	0.001	3
Fluorene (Flu)	0.001	3
Phenanthrene (Phe)	0.001	3
Anthracene (Ant)	0.01	3
Fluroanthene (Flt)	0.001	3
Pyrene (Pyr)	0.001	3
Benzo[a]anthracene (BaA)	0.1	2B
Chrysene (Chr)	0.01	2B
Benzo[k]fluoranthrene (BkF)	0.1	2B
Benzo[b]fluoranthrene (BbF)	0.1	2B
Benzo[a]pyrene (BaP)	1	1
Benzo[g,h,i]perylene (BghiP)	0.01	3
Dibenz[a,h]anthracene (DBahA)	1	2A
Indeno[1,2,3-c,d]pyrene (IP)	0.1	2B

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## 87 PAHs concentrations and Weather Data

88 Table S3: Total PAHs with the recorded temperature, wind speed and direction for each sampling

89 day.

Sampling Date	Total PPAH	Temperature	Wind Speed	Wind Direction
	$(ng/m^3)$	(°C)	(m/s)	
11/16/2016	15.4	23.0	6.8	S/SW
11/18/2016	11.4	22.0	2.8	N/NE
11/24/2016	12.3	23.0	0.5	E/SE
11/30/2016	11.0	22.0	7.2	S/SE
12/5/2016	14.4	21.0	2.0	E/NE
12/11/2016	14.6	19.0	2.6	S
12/17/2016	16.0	16.0	1.3	E/NE
12/28/2016	14.3	16.0	3.9	S
1/2/2017	15.2	14.0	7.9	S/SE
1/8/2017	12.7	16.0	10.6	S
1/14/2017	11.1	16.0	0.7	SE
1/24/2017	14.9	17.0	0.9	E/SE
1/30/2017	15.8	13.0	3.1	Е
2/6/2017	16.3	17.0	1.9	E/SE
2/12/2017	9.6	16.0	1.9	E/SE
2/20/2017	17.8	16.0	2.4	E/NE
2/26/2017	15.6	13.0	2.7	E/NE
3/5/2017	9.5	18.0	4.4	N/NE
3/11/2017	11.3	19.0	10.0	S/SW
3/17/2017	10.2	17.0	7.3	S
3/23/2017	10.1	21.0	3.6	W
3/29/2017	8.5	21.0	3.6	N/NE
4/4/2017	10.6	21.0	4.9	N/NE

4/10/2017	7.8	21.0	5.4	N
4/16/2017	8.3	20.0	4.8	SW
4/22/2017	8.1	26.0	1.2	Е
4/26/2017	9.7	25.0	5.0	Ν
5/2/2017	8.4	22.0	4.7	SW
5/8/2017	10.0	25.0	5.4	SW
5/14/2017	9.6	25.0	5.2	W/SW
5/20/2017	8.1	24.0	7.6	S/SW
6/11/2017	6.8	27.0	5.2	SW
6/17/2017	6.5	28.0	5.5	W/SW
6/23/2017	6.6	28.0	4.2	W/SW
6/29/2017	7.7	29.0	4.8	W/SW
7/6/2017	5.1	29.0	4.7	W/SW
7/12/2017	6.9	31.0	4.6	W/SW
7/18/2017	5.3	31.0	4.3	W
7/24/2017	5.5	31.0	4.2	W/SW
8/8/2017	6.3	30.0	4.2	N/NW
8/14/2017	6.2	30.0	4.5	W/SW
8/21/2017	6.1	30.0	3.9	W
8/27/2017	6.6	32.0	5.2	S/SW
9/2/2017	8.3	28.0	3.8	SW
9/8/2017	9.1	28.0	4.2	Ν
9/14/2017	9.3	30.0	4.0	Ν
9/20/2017	9.3	29.0	4.8	SW
10/10/2017	10.5	23.0	4.1	SW

10/16/2017	9.2	23.0	2.8	N
10/19/2017	10.2	28.0	1.5	E/SE
10/25/2017	7.5	25.0	2.6	SW
11/10/2017	13.3	22.0	1.7	E/NE
11/14/2017	11.4	22.0	3.3	S/SW
11/18/2017	15.6	23.0	1.8	E/SE
11/22/2017	13.3	18.0	2.6	W/NW

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91 Figure S1 shows the negative correlation between total PAHs and wind speed, where during high 92 wind speed, the total sum of PAHs captured is low and vice versa. The wind speed and direction 93 were obtained from Earth NullSchool (<u>https://earth.nullschool.net/</u>). Samples were collected on 94 non-rainy days. This is in good accordance with data recorded by other studies.<sup>5-7</sup>



96 Figure S1: Variation of concentration of total PAHs with respect to wind speed over the period 97 extended between Nevember 2016 and Nevember 2017

97 extended between November 2016 and November 2017

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99 References

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- 102 1. P. Paatero and P. K. Hopke, Analytica Chimica Acta, 2003, 490, 277-289.
- 103 2. C. Belis, O. Favez, M. Mircea, E. Diapouli, M.-I. Manousakas, S. Vratolis, S. Gilardoni,
- 104 M. Paglione, G. Močnik, D. Mooibroek, S. Takahama, R. Vecchi, P. Paatero, P. Salvador
- 105and S. Decesari, European guide on air pollution source apportionment with receptor106models Revised version 2019, 2019.
- 107 3. G. Norris and R. Duval, EPA Positive Matrix Factorization (PMF) 5.0 Fundamentals and
  108 User Guide,

109https://cfpub.epa.gov/si/si\_public\_file\_download.cfm?p\_download\_id=523318&Lab=NE110RL).

- 111 4. I. C. Nisbet and P. K. Lagoy, *Regulatory toxicology and pharmacology*, 1992, 16, 290300.
- 113 5. A. C. Amarillo and H. Carreras, *Atmospheric Pollution Research*, 2016, 7, 597-602.
- M. Amodio, M. Caselli, G. de Gennaro and M. Tutino, *Environmental Research*, 2009, **109**, 812-820.
- A. Nadali, M. Leili, A. Bahrami, M. Karami and A. Afkhami, *Ecotoxicology and Environmental Safety*, 2021, 209, 111807.

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