

Reproducibility of paper:

Lin, L., Yi, X., Liu, H. et al. The airway microbiome mediates the interaction between environmental exposure and respiratory health in humans. Nat Med 29, 1750–1759 (2023).

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Packages used

```
if (!require("pacman", quietly = TRUE)) {  
  install.packages("pacman")  
}  
  
pacman::p_load(  
  tidyverse, # Used for basic data handling and visualization.  
  dagitty, #Used in conjunction with https://www.dagitty.net/ to create  
  #directed acyclic graph to inform statistical modelling.  
  lavaan, #Used to create correlation matrix to assess conditional independencies.  
  CBPS, #Used to calculate non-parametric propensity scores for IPW.  
  WeightIt, #Used to calculate inverse probability weights.  
  boot # Calculate bootstrap confidence intervals.  
)
```

Session and package dependencies

R version 4.3.3 (2024-02-29 ucrt)
Platform: x86_64-w64-mingw32/x64 (64-bit)
Running under: Windows 11 x64 (build 22631)

Matrix products: default

locale:

[1] LC_COLLATE=Spanish_Mexico.utf8 LC_CTYPE=Spanish_Mexico.utf8
[3] LC_MONETARY=Spanish_Mexico.utf8 LC_NUMERIC=C
[5] LC_TIME=Spanish_Mexico.utf8

time zone: Europe/Berlin

tzcode source: internal

attached base packages:

[1] stats graphics grDevices utils datasets methods base

other attached packages:

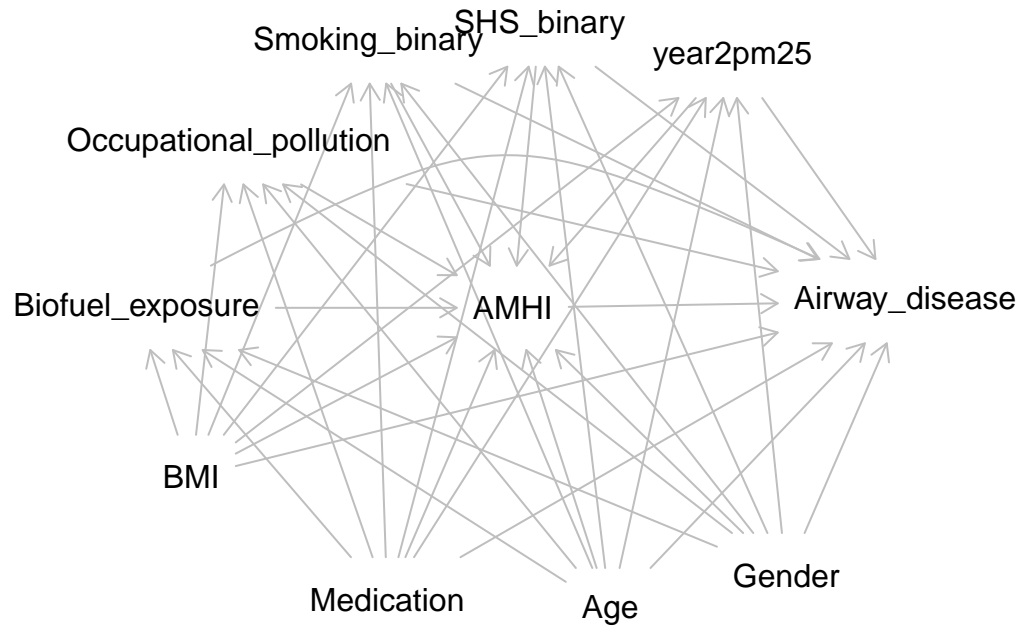
[1] boot_1.3-30	WeightIt_1.0.0	CBPS_0.23
[4] glmnet_4.1-8	Matrix_1.6-5	numDeriv_2016.8-1.1
[7] nnet_7.3-19	MatchIt_4.5.5	MASS_7.3-60.0.1
[10] lavaan_0.6-17	dagitty_0.3-4	lubridate_1.9.3

[13]	forcats_1.0.0	stringr_1.5.1	dplyr_1.1.4
[16]	purrr_1.0.2	readr_2.1.5	tidyr_1.3.1
[19]	tibble_3.2.1	ggplot2_3.5.0	tidyverse_2.0.0
[22]	pacman_0.5.1		

The summary of criticisms to the paper and rationale for the following analyses are summarized in the [powerpoint presentation](#) contained in the *mediation* directory.

DAG

The following directed acyclic graph (DAG) was reconstructed based on assumptions available in the paper for the construction of regression models in mediation analyses, by using the [DAGitty website](#). The DAG is saved and sourced from the accompanying script **DAG.r**



Testing of conditional independencies in DAG:

This procedure was performed as suggested in [this article](#).

Implied conditional independencies:

```

Age _||_ BMI
Age _||_ Gndr
Age _||_ Mdct
BMI _||_ Gndr
BMI _||_ Mdct
Bfl_ _||_ Occ_ | Age, BMI, Gndr, Mdct
Bfl_ _||_ SHS_ | Age, BMI, Gndr, Mdct

```

Bfl_ _||_ Smk_ | Age, BMI, Gndr, Mdct
 Bfl_ _||_ y225 | Age, BMI, Gndr, Mdct
 Gndr _||_ Mdct
 Occ_ _||_ SHS_ | Age, BMI, Gndr, Mdct
 Occ_ _||_ Smk_ | Age, BMI, Gndr, Mdct
 Occ_ _||_ y225 | Age, BMI, Gndr, Mdct
 SHS_ _||_ Smk_ | Age, BMI, Gndr, Mdct
 SHS_ _||_ y225 | Age, BMI, Gndr, Mdct
 Smk_ _||_ y225 | Age, BMI, Gndr, Mdct

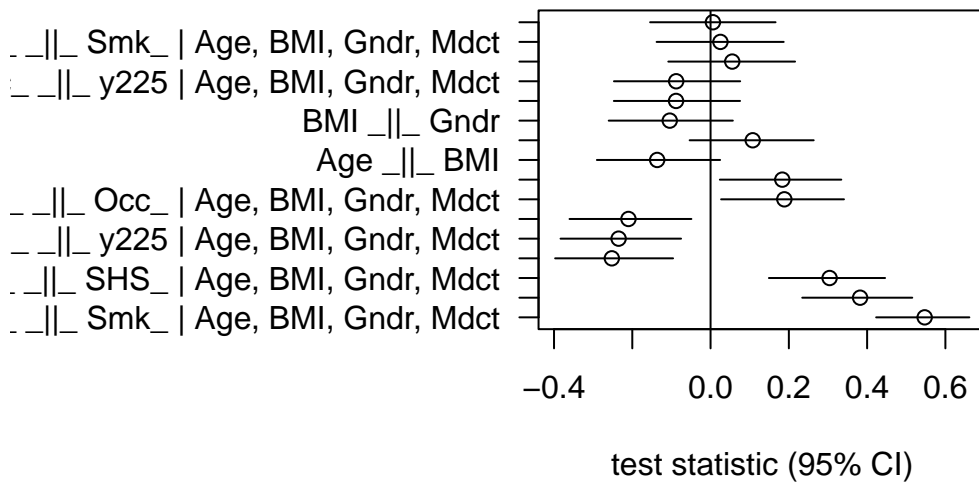
	Bfl_xp	Occpt_	Smkng_	SHS_bn	yr2p25	Age	BMI	Medctn
Biofuel_exposure	1.000							
Occupational_pollution	0.304	1.000						
Smoking_binary	0.171	0.342	1.000					
SHS_binary	0.154	0.296	0.052	1.000				
year2pm25	-0.196	-0.122	-0.267	-0.040	1.000			
Age	0.159	0.112	0.141	-0.032	-0.005	1.000		
BMI	-0.220	-0.035	-0.125	-0.125	0.026	-0.136	1.000	
Medication	0.372	0.458	0.231	0.245	0.002	0.006	-0.252	1.000
Gender	0.026	0.328	0.939	-0.148	-0.214	0.108	-0.104	0.183
Airway_disease	0.216	0.104	0.003	0.012	0.287	0.059	-0.155	0.519
AMHI	-0.110	-0.134	-0.025	-0.041	-0.125	-0.089	-0.078	-0.392
	Gender	Arwy_d	AMHI					

Biofuel_exposure
 Occupational_pollution
 Smoking_binary
 SHS_binary
 year2pm25
 Age
 BMI
 Medication
 Gender 1.000
 Airway_disease 0.048 1.000
 AMHI 0.007 -0.548 1.000

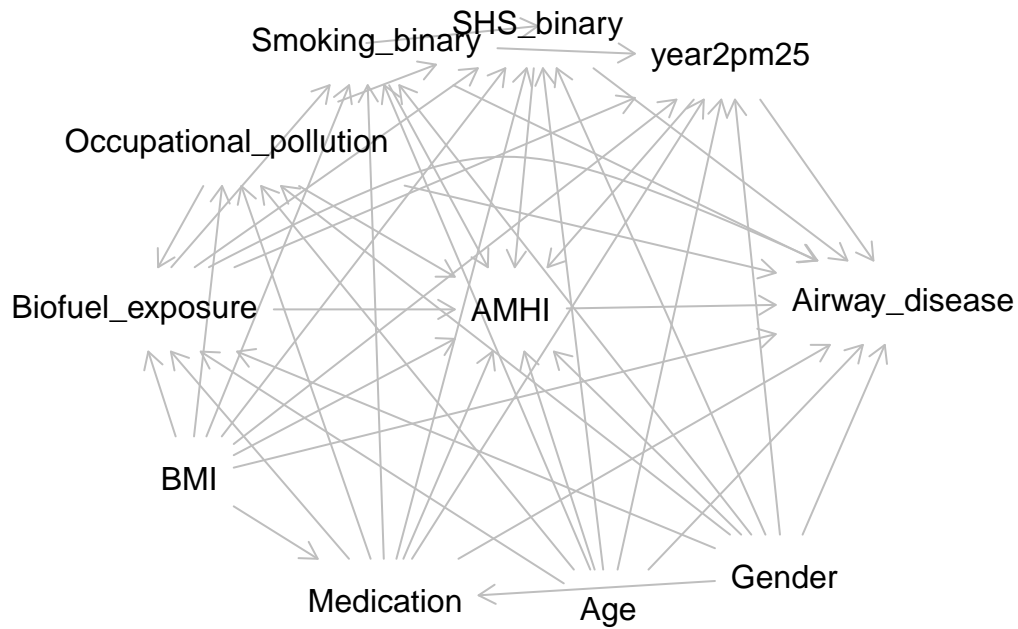
	estimate	p.value	2.5%
Age _ _ BMI	-0.136414439	9.604998e-02	-0.29047053
Age _ _ Gndr	0.107697031	1.899115e-01	-0.05348771
Age _ _ Mdct	0.005961319	9.423807e-01	-0.15445367
BMI _ _ Gndr	-0.104254367	2.045743e-01	-0.26025039
BMI _ _ Mdct	-0.252231108	1.774505e-03	-0.39715968
Bfl_ _ _ Occ_ Age, BMI, Gndr, Mdct	0.188498630	2.252495e-02	0.02687314

Bfl_ _ _ SHS_	Age, BMI, Gndr, Mdct	0.055390463	5.072971e-01	-0.10803097
Bfl_ _ _ Smk_	Age, BMI, Gndr, Mdct	0.382088405	1.485078e-06	0.23422305
Bfl_ _ _ y225	Age, BMI, Gndr, Mdct	-0.234927411	4.199311e-03	-0.38334693
Gndr_ _ _ Mdct		0.183276828	2.460759e-02	0.02371177
Occ_ _ _ SHS_	Age, BMI, Gndr, Mdct	0.304120705	1.729680e-04	0.14904018
Occ_ _ _ Smk_	Age, BMI, Gndr, Mdct	0.025186032	7.632278e-01	-0.13782960
Occ_ _ _ y225	Age, BMI, Gndr, Mdct	-0.087881979	2.920506e-01	-0.24686925
SHS_ _ _ Smk_	Age, BMI, Gndr, Mdct	0.547029400	2.073739e-13	0.42304675
SHS_ _ _ y225	Age, BMI, Gndr, Mdct	-0.088189061	2.903589e-01	-0.24716023
Smk_ _ _ y225	Age, BMI, Gndr, Mdct	-0.209778365	1.088447e-02	-0.36038277
		97.5%		
Age_ _ _ BMI		0.02438008		
Age_ _ _ Gndr		0.26349879		
Age_ _ _ Mdct		0.16607269		
BMI_ _ _ Gndr		0.05695887		
BMI_ _ _ Mdct		-0.09584441		
Bfl_ _ _ Occ_	Age, BMI, Gndr, Mdct	0.34083839		
Bfl_ _ _ SHS_	Age, BMI, Gndr, Mdct	0.21592797		
Bfl_ _ _ Smk_	Age, BMI, Gndr, Mdct	0.51534956		
Bfl_ _ _ y225	Age, BMI, Gndr, Mdct	-0.07535500		
Gndr_ _ _ Mdct		0.33402330		
Occ_ _ _ SHS_	Age, BMI, Gndr, Mdct	0.44585615		
Occ_ _ _ Smk_	Age, BMI, Gndr, Mdct	0.18688523		
Occ_ _ _ y225	Age, BMI, Gndr, Mdct	0.07564652		
SHS_ _ _ Smk_	Age, BMI, Gndr, Mdct	0.66120201		
SHS_ _ _ y225	Age, BMI, Gndr, Mdct	0.07533880		
Smk_ _ _ y225	Age, BMI, Gndr, Mdct	-0.04899982		

Local tests results plot:



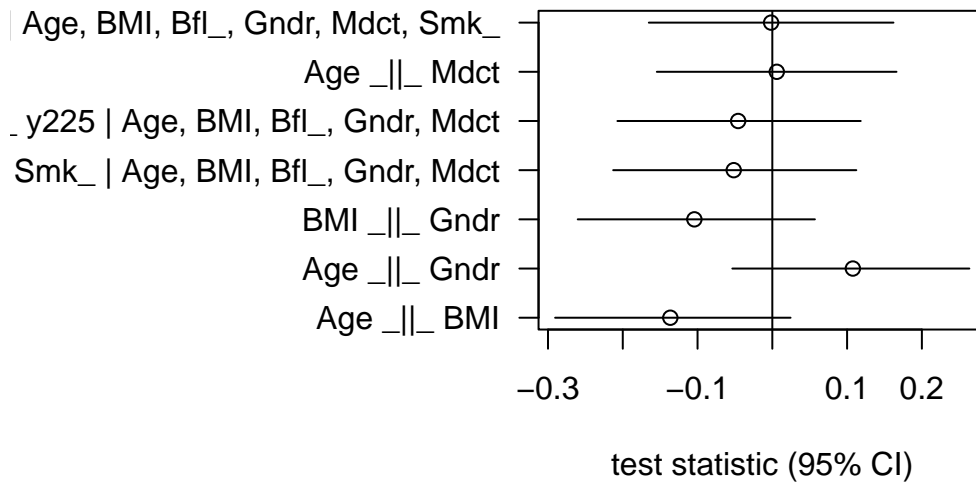
I updated the DAG according to the procedure described by [Ankan, 2021](#).



	estimate	p.value
Age_ BMI	-0.136414439	0.09604998

Age _ _ Gndr	0.107697031	0.18991153
Age _ _ Mdct	0.005961319	0.94238067
BMI _ _ Gndr	-0.104254367	0.20457426
Occ_ _ _ Smk_ Age, BMI, Bfl_, Gndr, Mdct	-0.051607761	0.53820853
Occ_ _ _ y225 Age, BMI, Bfl_, Gndr, Mdct	-0.045672563	0.58600780
SHS_ _ _ y225 Age, BMI, Bfl_, Gndr, Mdct, Smk_	-0.001613923	0.98471005
	2.5%	97.5%
Age _ _ BMI	-0.29047053	0.02438008
Age _ _ Gndr	-0.05348771	0.26349879
Age _ _ Mdct	-0.15445367	0.16607269
BMI _ _ Gndr	-0.26025039	0.05695887
Occ_ _ _ Smk_ Age, BMI, Bfl_, Gndr, Mdct	-0.21285615	0.11234779
Occ_ _ _ y225 Age, BMI, Bfl_, Gndr, Mdct	-0.20716543	0.11821834
SHS_ _ _ y225 Age, BMI, Bfl_, Gndr, Mdct, Smk_	-0.16515452	0.16201228

Local tests results plot:



Biofuel

Inverse probability weights

Weights for exposure:

Weights for mediator:

Overall weight:

Confidence interval

BOOTSTRAP CONFIDENCE INTERVAL CALCULATIONS

Based on 10000 bootstrap replicates

CALL :

```
boot.ci(boot.out = boot_results, type = "all")
```

Intervals :

Level	Normal	Basic
95%	(-28.0075, 29.2961)	(-4.0017, 4.8129)

Level	Percentile	BCa
95%	(-4.1996, 4.6150)	(-1.5276, 13.7922)

Calculations and Intervals on Original Scale

Occupational pollution

Inverse probability weights

Weights for exposure:

Weights for mediator:

Overall weight:

Confidence interval

BOOTSTRAP CONFIDENCE INTERVAL CALCULATIONS

Based on 10000 bootstrap replicates

CALL :

```
boot.ci(boot.out = boot_results, type = "all")
```

```

Intervals :
Level      Normal      Basic
95%  ( -962.870,  957.094 )  ( -22.483,   8.908 )

Level      Percentile      BCa
95%  (  -16.420,   14.971 )  ( -47254.558,   -1.360 )
Calculations and Intervals on Original Scale
Warning : BCa Intervals used Extreme Quantiles
Some BCa intervals may be unstable

```

PM 2.5

Inverse probability weights

Weights for exposure:

Weights for mediator:

Overall weight:

Confidence interval

```

BOOTSTRAP CONFIDENCE INTERVAL CALCULATIONS
Based on 10000 bootstrap replicates

```

```

CALL :
boot.ci(boot.out = boot_results, type = "all")

```

```

Intervals :
Level      Normal      Basic
95%  ( -48.9454,  48.3801 )  ( -0.8053,   0.6633 )

Level      Percentile      BCa
95%  ( -0.7096,  0.7590 )  ( -0.6328,   0.9072 )
Calculations and Intervals on Original Scale

```

Smoking

Inverse probability weights

Weights for exposure:

Weights for mediator:

Overall weight:

Confidence interval

BOOTSTRAP CONFIDENCE INTERVAL CALCULATIONS

Based on 10000 bootstrap replicates

CALL :

```
boot.ci(boot.out = boot_results, type = "all")
```

Intervals :

Level	Normal	Basic
95%	(-52.7994, 53.0275)	(-5.9490, 6.3779)

Level	Percentile	BCa
95%	(-5.7300, 6.5969)	(-2.1082, 16.5905)

Calculations and Intervals on Original Scale

Second hand smoking

Inverse probability weights

Weights for exposure:

Weights for mediator:

Overall weight:

Confidence interval

BOOTSTRAP CONFIDENCE INTERVAL CALCULATIONS

Based on 10000 bootstrap replicates

CALL :

```
boot.ci(boot.out = boot_results, type = "all")
```

Intervals :

Level	Normal	Basic
95%	(-61.014, 72.596)	(-2.389, 13.715)

Level	Percentile	BCa
95%	(-7.930, 8.173)	(1.141, 1095.965)

Calculations and Intervals on Original Scale

Warning : BCa Intervals used Extreme Quantiles

Some BCa intervals may be unstable

Table results

	Exposure	ci_lower	ci_upper	confidence_interval
1	Biofuel_exposure	0	4.6149729	0, 4.61
2	Occupational_pollution	0	14.9713783	0, 14.97
3	PM25	0	0.7589639	0, 0.76
4	Smoking	0	6.5969129	0, 6.6
5	Secondhand_smoking	0	8.1734864	0, 8.17

