

## MINIMAL SUFFICIENT ADJUSTMENT SET

We explored whether the potential confounders included in *model 1* and *model 2* represent the “*minimal sufficient adjustment set*” to estimate the total effect of each exposure on the outcomes (i.e. the smallest group of measured covariates that needs to be included in order to eliminate confounding). We used directed acyclic graphs (DAGs) [1] in DAGitty (*dagitty.net*) (**figure S1** and **figure S2**). DAGs help to minimize the magnitude of the bias in the estimates, to avoid the risk of over-adjustment and to establish whether the statistical models used are the most parsimonious.

The DAG analysis supported the assumption that the minimal sufficient adjustment set contains grandparents’ education level, fathers’ age, education level and occupational class, mother’s smoking before or after offspring’s birth, and offspring’s age, education level, sex and smoking (“*education\_GP*”, “*age\_F*”, “*education\_F*”, “*occupation\_F*”, “*smoke\_M*”, “*age\_O*”, “*education\_O*”, “*sex\_O*” and “*smoke\_O*” in **figure S1** and **figure S2**).

## UNMEASURED CONFOUNDING

We evaluated the impact of unmeasured confounding [2] on the estimate of the natural direct and indirect effects of fathers’ and grandmothers’ smoking on offspring’s lung function, using the Umediation package ([github.com/SharonLutz/Umediation](https://github.com/SharonLutz/Umediation)) in R3.6.1. Umediation makes it possible to simulate unmeasured confounding of the exposure-outcome, exposure-mediator and mediator-outcome relationships in order to investigate how the results would change if up to two unmeasured confounders were included in the mediation models.

We carried out the simulation analyses as follows:

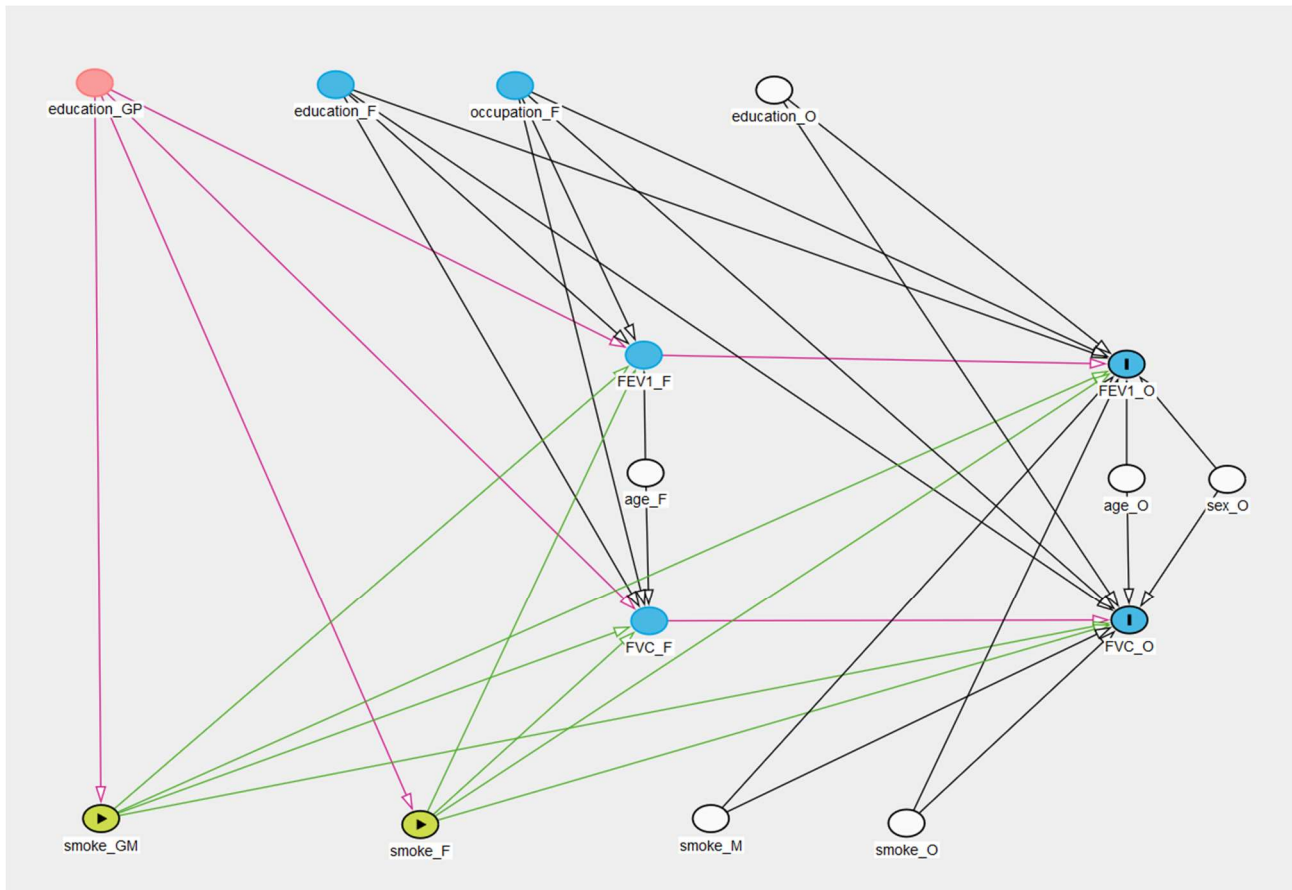
- In *model 1* and *model 2*, we subdivided the paths in order to simulate unmeasured confounding within a single-exposure, single-mediator, single-outcome framework. Then, we added up to two unmeasured normally distributed confounders with mean 0 and variance 0.001 to the models (“ $U_1$ ” and “ $U_2$ ” variables). **Figure S3** shows how the data were simulated for “*smoke\_GM*” (exposure), “*FEV1\_F*” (mediator) and “*FEV1\_O*” (outcome).
- As inputs for the simulations, we used the beta regression coefficients obtained from the cluster-robust (cluster variable: centre) linear models, defined as follows:
  - “*FEV1\_O*” as the outcome and “*smoke\_GM*”, “*education\_F*”, “*FEV1\_F*”, “*occupation\_F*”, “*smoke\_F*”, “*smoke\_M*”, “*age\_O*”, “*education\_O*”, “*sex\_O*”, “*smoke\_O*” as covariates;
  - “*FVC\_O*” as the outcome and “*smoke\_GM*”, “*education\_F*”, “*FVC\_F*”, “*occupation\_F*”, “*smoke\_F*”, “*smoke\_M*”, “*age\_O*”, “*education\_O*”, “*sex\_O*”, “*smoke\_O*” as covariates;
  - “*FEV1/FVC\_O*” as the outcome and “*smoke\_GM*”, “*education\_F*”, “*FEV1/FVC\_F*”, “*occupation\_F*”, “*smoke\_F*”, “*smoke\_M*”, “*age\_O*”, “*education\_O*”, “*sex\_O*”, “*smoke\_O*” as covariates;
  - “*FEV1\_F*” as the outcome and “*education\_GP*”, “*smoke\_GM*”, “*age\_F*”, “*education\_F*”, “*occupation\_F*”, “*smoke\_F*” as covariates;
  - “*FVC\_F*” as the outcome and “*education\_GP*”, “*smoke\_GM*”, “*age\_F*”, “*education\_F*”, “*occupation\_F*”, “*smoke\_F*” as covariates;
  - “*FEV1/FVC\_F*” as the outcome and “*education\_GP*”, “*smoke\_GM*”, “*age\_F*”, “*education\_F*”, “*occupation\_F*”, “*smoke\_F*” as covariates;
  - “*smoke\_GM*” as the outcome and no covariates (null model);
  - “*smoke\_F*” as the outcome and no covariates (null model).

The beta regression coefficients were estimated from 800 bootstrap samplings of one offspring per parent ( $n = n^\circ$  offspring =  $n^\circ$  fathers = 274). This was done to avoid the “ $2 \rightarrow 2 \rightarrow 1$ ” mediation pattern.

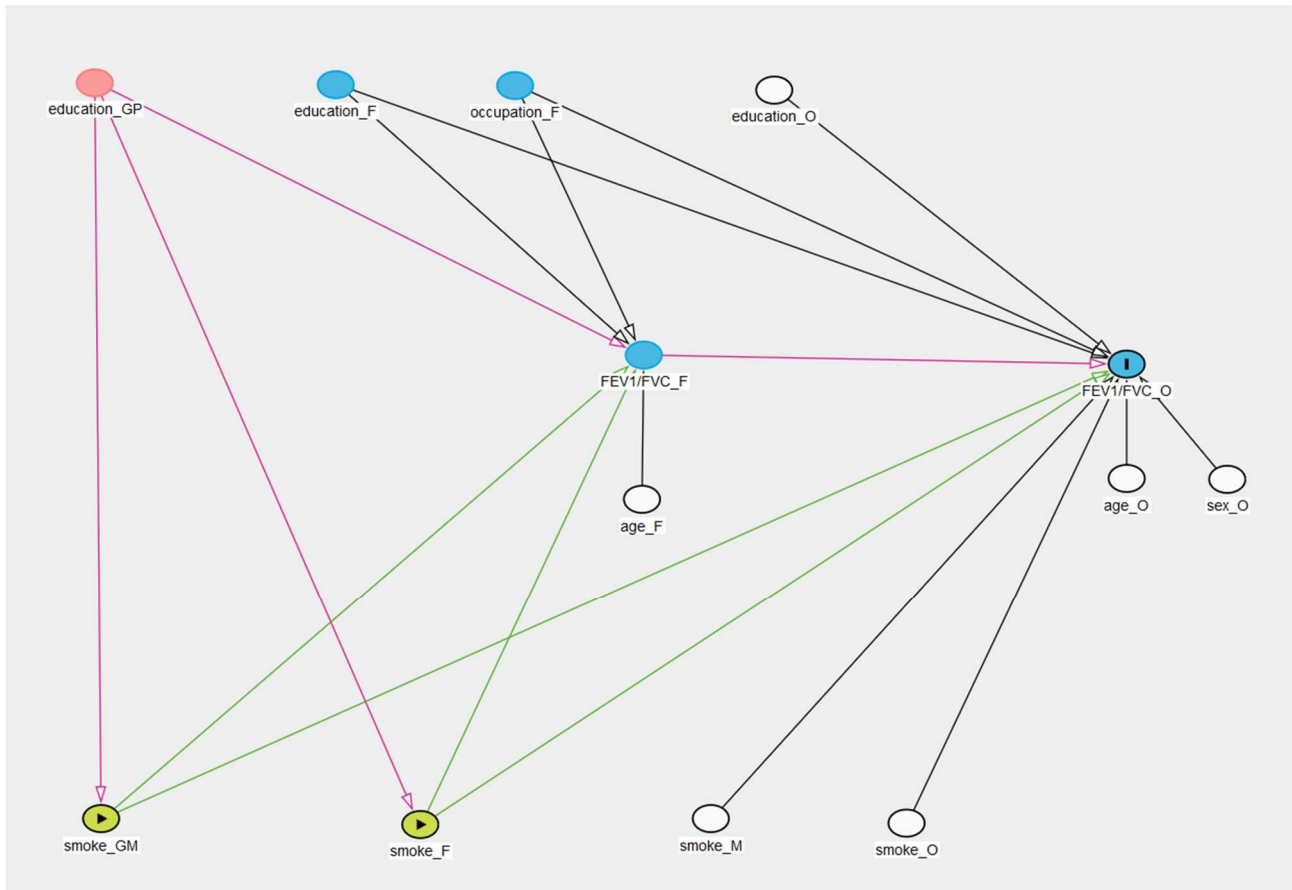
- We carried out the simulations under multiple scenarios for the effects (beta regression coefficients) of the unmeasured confounder “ $U_1$ ” on “*smoke\_GM*” and “*smoke\_F*” (exposure;  $\beta_{U \rightarrow E}$ ), on “*FEV1\_F*”, “*FVC\_F*” and “*FEV1/FVC\_F*” (mediator;  $\beta_{U \rightarrow M}$ ), and on “*FEV1\_O*”, “*FVC\_O*” and “*FEV1/FVC\_O*” (outcome;  $\beta_{U \rightarrow O}$ ), by fixing  $\beta_{U \rightarrow E} = \beta_{U \rightarrow M} = \beta_{U \rightarrow O} = 0, 1, 3, 5, 7$  and  $9$ . We repeated the simulations by adding the second unmeasured confounder “ $U_2$ ” to the models under the same assumptions.
- We selected 1,000 simulation runs and 1,000 Monte Carlo draws for the nonparametric bootstrap in each of the simulation analyses.

## REFERENCES

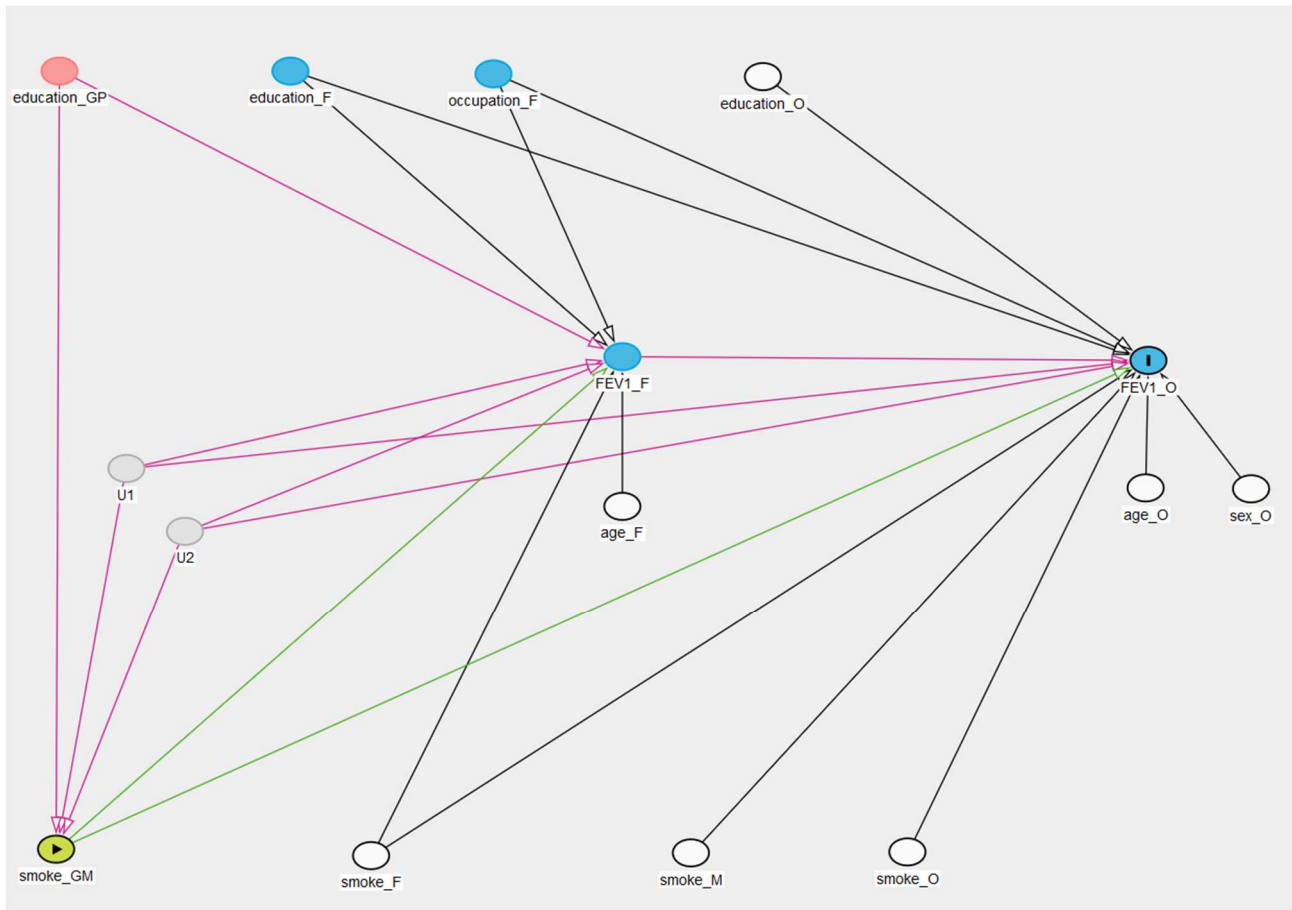
- 1 Greenland S, Pearl J, Robins JM. Causal diagrams for epidemiologic research. *Epidemiology* 1999; 10: 37-48.
- 2 Lutz SM, Thwing A, Schmiede S, *et al.* Examining the role of unmeasured confounding in mediation analysis with genetic and genomic applications. *BMC Bioinformatics* 2017; 18: 344.



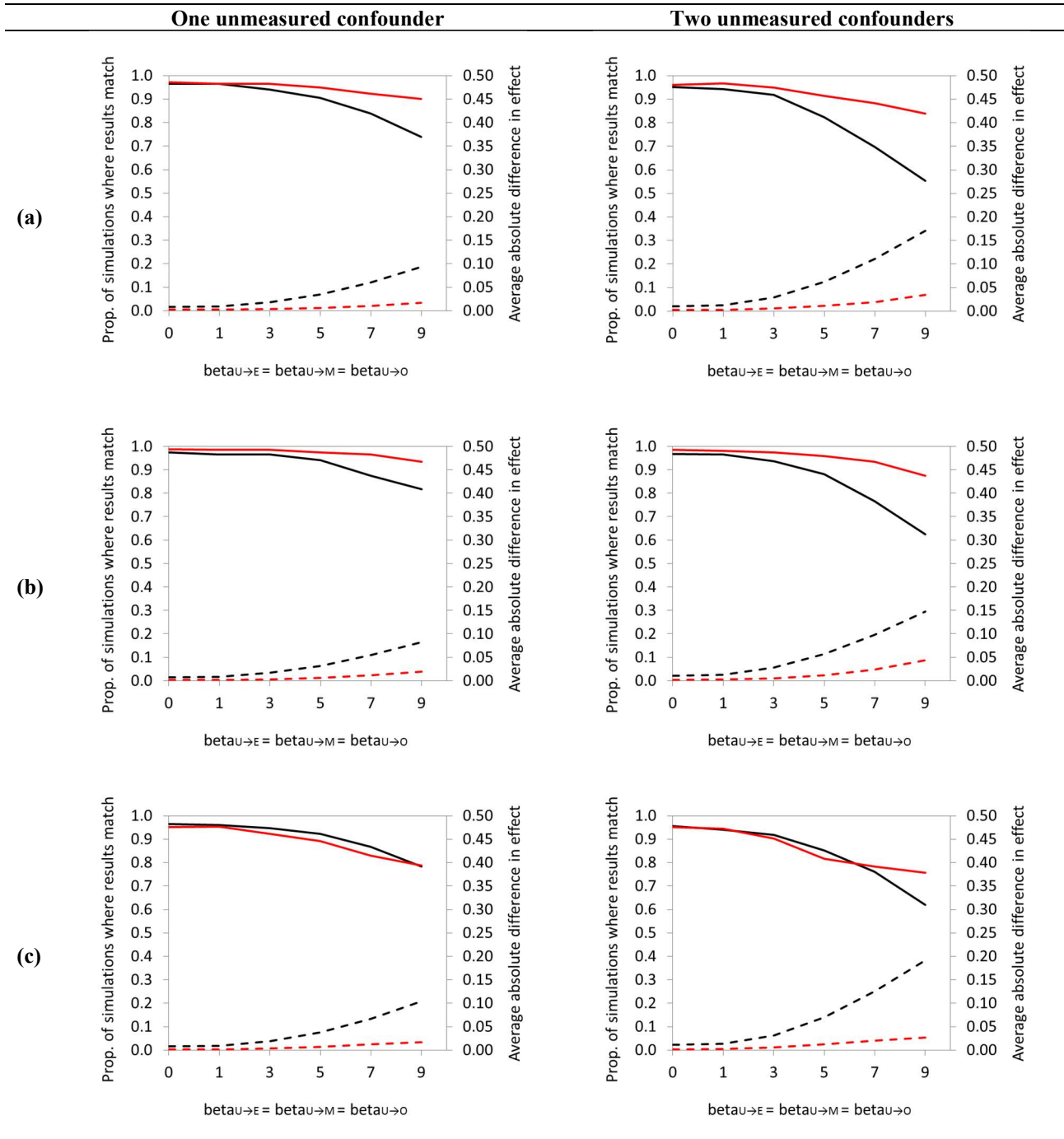
**Figure S1.** Directed acyclic graph used to check if the potential confounders included in *model 1* represent the “*minimal sufficient adjustment set*”. Grandparents (generation G0): “education\_GP”, grandparents’ education level; “smoke\_GM”, grandmother’s smoke. Father/mother (generation G1): “age\_F”, father’s age; “education\_F”, father’s education level; “FEV1\_F”, father’s FEV<sub>1</sub> z-score; “FVC\_F”, father’s FVC z-score; “occupation\_F”, father’s occupational class; “smoke\_F”, father’s smoke; “smoke\_M”, mother’s smoke. Offspring (generation G2): “age\_O”, offspring’s age; “education\_O”, offspring’s education level; “FEV1\_O”, offspring’s FEV<sub>1</sub> z-score; “FVC\_O”, offspring’s FVC z-score; “sex\_O”, offspring’s sex; “smoke\_O”, offspring’s smoke.



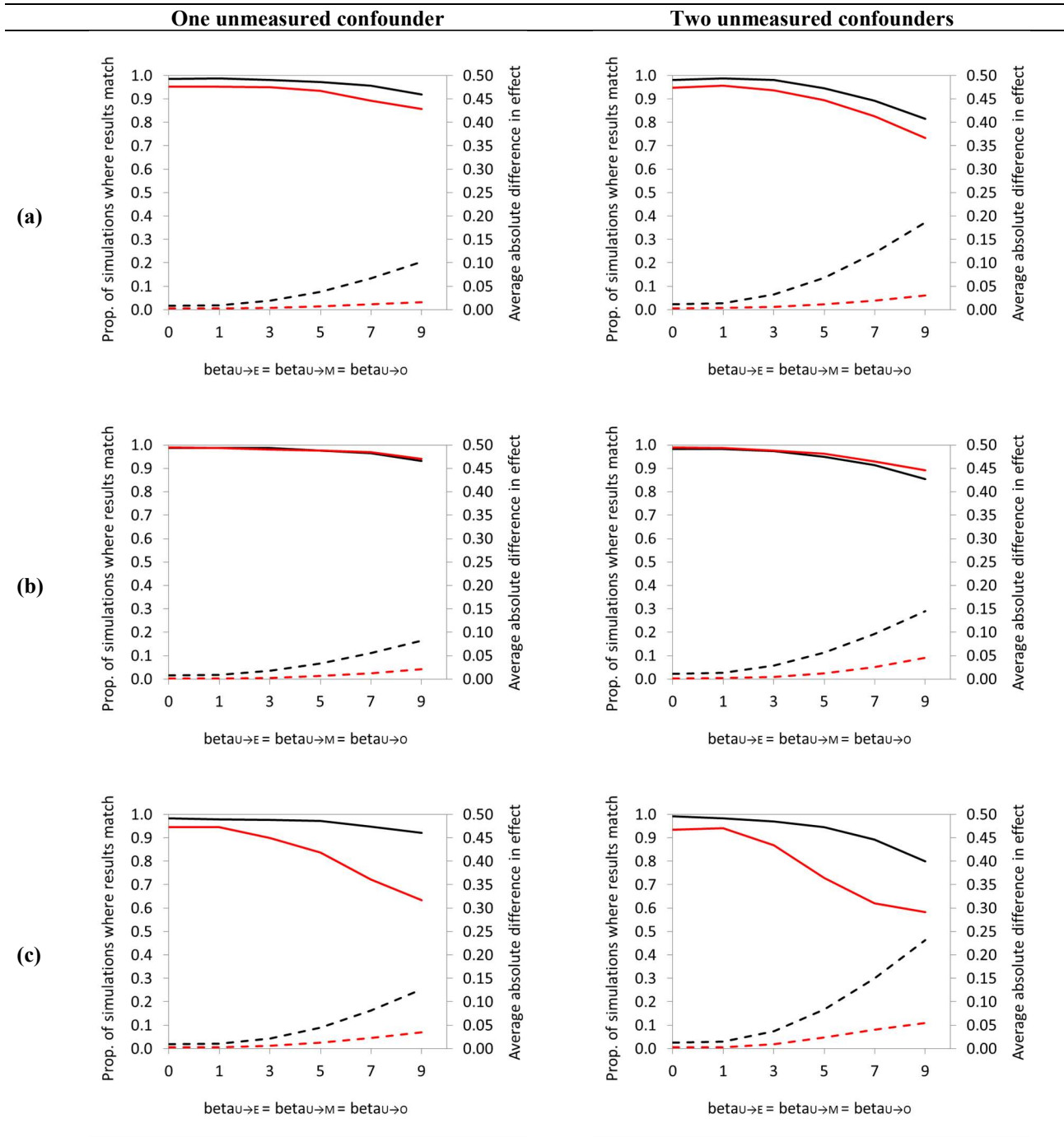
**Figure S2.** Directed acyclic graph used to check if the potential confounders included in *model 2* represent the “*minimal sufficient adjustment set*”. Grandparents (generation G0): “education\_GP”, grandparents’ education level; “smoke\_GM”, grandmother’s smoke. Father/mother (generation G1): “age\_F”, father’s age; “education\_F”, father’s education level; “FEV1/FVC\_F”, father’s FEV<sub>1</sub>/FVC z-score; “occupation\_F”, father’s occupational class; “smoke\_F”, father’s smoke; “smoke\_M”, mother’s smoke. Offspring (generation G2): “age\_O”, offspring’s age; “education\_O”, offspring’s education level; “FEV1/FVC\_O”, offspring’s FEV<sub>1</sub>/FVC z-score; “sex\_O”, offspring’s sex; “smoke\_O”, offspring’s smoke.



**Figure S3.** Directed acyclic graph used to simulate the impact of two unmeasured confounders (“U<sub>1</sub>” and “U<sub>2</sub>”) on the direct and indirect effects of grandmothers’ smoking in pregnancy on offspring’s FEV<sub>1</sub> z-score. Grandparents (generation G0): “education\_GP”, grandparents’ education level; “smoke\_GM”, grandmother’s smoke. Father/mother (generation G1): “age\_F”, father’s age; “education\_F”, father’s education level; “FEV1\_F”, father’s FEV<sub>1</sub> z-score; “occupation\_F”, father’s occupational class; “smoke\_F”, father’s smoke; “smoke\_M”, mother’s smoke. Offspring (generation G2): “age\_O”, offspring’s age; “education\_O”, offspring’s education level; “FEV1\_O”, offspring’s FEV<sub>1</sub> z-score; “sex\_O”, offspring’s sex; “smoke\_O”, offspring’s smoke.



**Figure S4.** Proportion of Monte Carlo simulations where results match (solid line) and average absolute difference (dashed line) in the average direct (black line) and indirect (red line) effects of fathers' smoking initiation in prepuberty on offspring's lung function (whether one or two unmeasured confounders are included or excluded from the models). Outcomes: (a) FEV<sub>1</sub>, (b) FVC and (c) FEV<sub>1</sub>/FVC z-scores.



**Figure S5.** Proportion of Monte Carlo simulations where results match (solid line) and average absolute difference (dashed line) in the average direct (black line) and indirect (red line) effects of grandmothers' smoking in pregnancy on offspring's lung function (whether one or two unmeasured confounders are included or excluded from the models). Outcomes: (a) FEV<sub>1</sub>, (b) FVC and (c) FEV<sub>1</sub>/FVC z-scores.

## SUPPLEMENTARY INFORMATION ON THE EUROPEAN COMMUNITY RESPIRATORY HEALTH SURVEY (ECRHS)

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**Ethics approval:** Ethics approval was obtained by all centres from the appropriate ethics committees: **Antwerp City and Antwerp South:** Adviescommissie Medische Ethiek UZA-UA (CME); **Tartu:** Research Ethics Committee of the University of Tartu, Estland (N° 209T-17); **French centres:** Comité de protection des personnes, Sud V Est (N° 2011-A00013-38); **German centres:** Ethik-Kommission der Bayerischen Landesärztekammer (N° 10015); **Reykjavik:** National Biotech Committee of Iceland (NBCI) (N° VSNb2011090016/03.11); **Pavia:** Fondazione IRCCS Policlinico 'San Matteo' (N° P-20110024215); **Turin:** Comitato Etico dell'Azienda Sanitaria Locale TO/2 di Torino (N° 569/09/08); **Verona:** Comitato Etico per la Sperimentazione dell'Azienda Ospedaliera Istituti Ospitalieri di Verona (N° 1393); **Bergen:** Universitetet i Bergen, Regional komité for medisinsk og helsefaglig forskningsetikk, Vest-Norge (REK Vest) (N° 2010/759); **Albacete:** Comité de Ética e Investigación de Complejo Hospitalario de Albacete (N° 04/09); **Barcelona:** Comité Ético de Investigación Clínica del Instituto Municipal de Asistencia Sanitaria, Barcelona, Spain (N° PS09/00716); **Galdakao:** Comité Ético de Investigación del Hospital de Galdakao, Spain (N° 20101104); **Huelva:** Comisión de Investigación del Hospital Juan Ramón Jiménez de Huelva (N° 20090417); **Oviedo:** Comité Ético de Investigación Clínica Regional, Hospital Universitario Central de Asturias (N° 20110415); **Swedish centres:** Ethics Committee at the Medical Faculty, Uppsala University (N° 1999/313 and 2010/068); **Basel:** Swiss Academy of Medical Sciences and the ethics committee of Basel (N° PV123/00,157/00); **UK centres:** NRES Committee London - Stanmore (REC Reference 11/LO/0965 IRAS number 70769).