# **compmed:** A new command for estimating causal mediation effects with non-adherence to treatment allocation

Anca Chis Ster, Sabine Landau, Richard Emsley
Department of Biostatistics and Health Informatics, King's College London, United Kingdom

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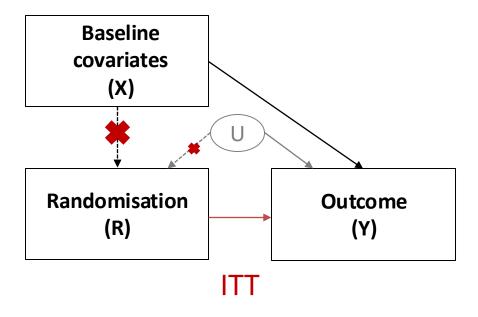




# Motivation for handling nonadherence in mediation

- Trials in mental health often evaluate complex interventions, such as therapy or psychotherapy
- One guideline for evaluating complex interventions is to understand the treatment mechanism, i.e., understand how the treatment works in practice
- Trials in mental health are faced with the challenge of nonadherence to treatment allocation
- It's currently unclear how to account for nonadherence in a mechanism evaluation, or how to practically implement this within a statistical package, e.g., **STATA**

### Background: The challenge of non-adherence in trials



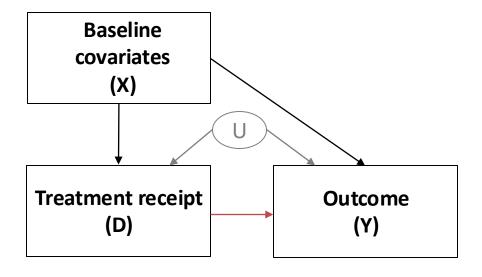
X, U = measured or unmeasured variables

ITT = Intention-to-treat

CACE = Complier Average Causal Effect

NDE = Natural Direct Effect

### Background: The challenge of non-adherence in trials



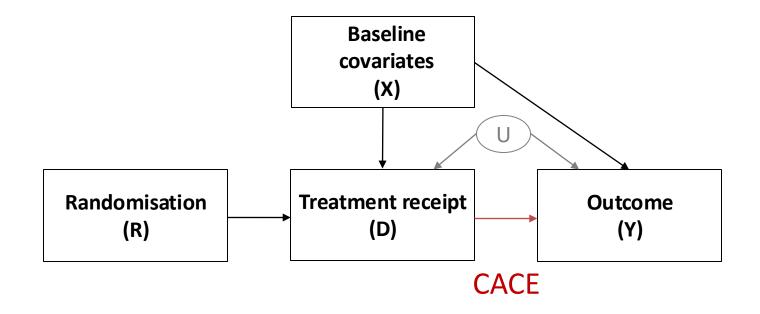
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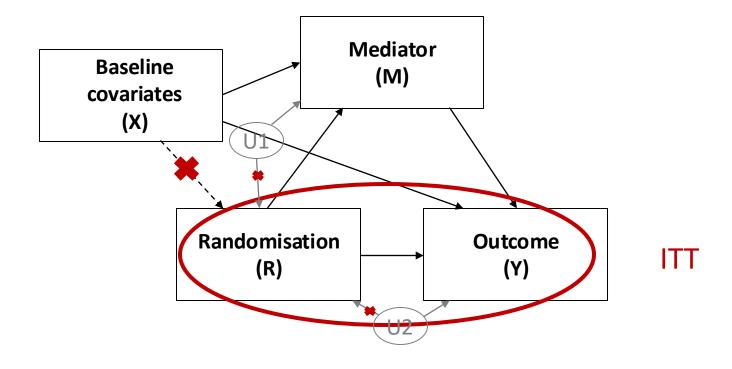
### Background: The challenge of non-adherence in trials



X, U = measured or unmeasured variables ITT = Intention-to-treat

CACE = Complier Average Causal Effect

NDE = Natural Direct Effect
NIE = Natural Indirect Effect

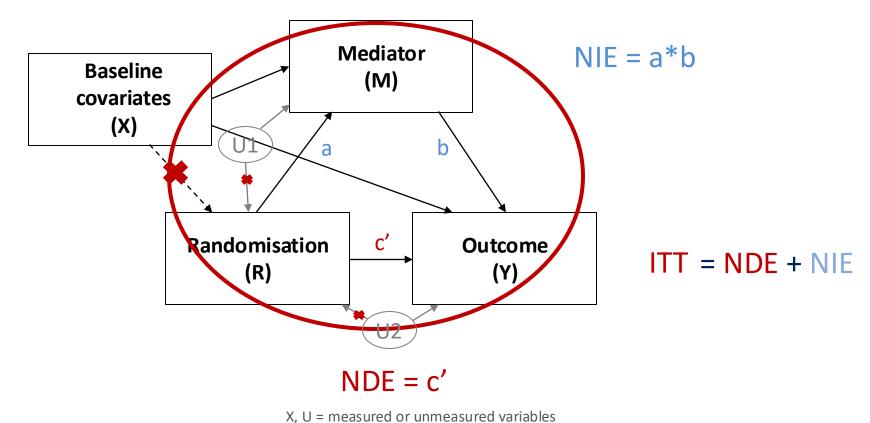


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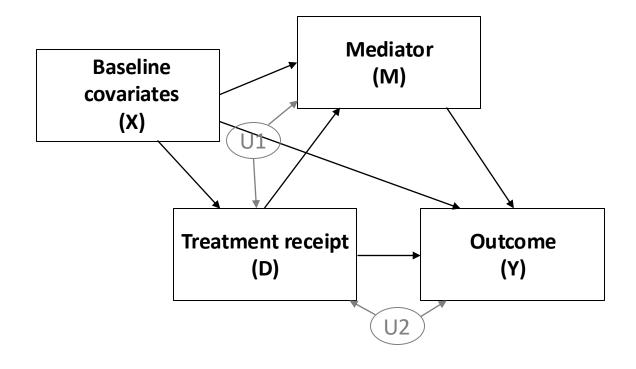
NDE = Natural Direct Effect



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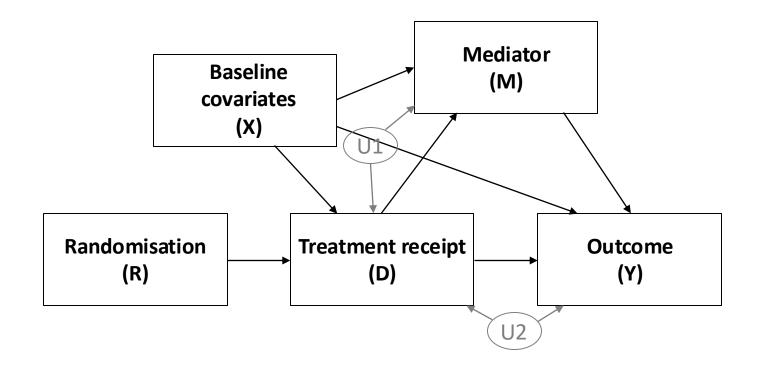


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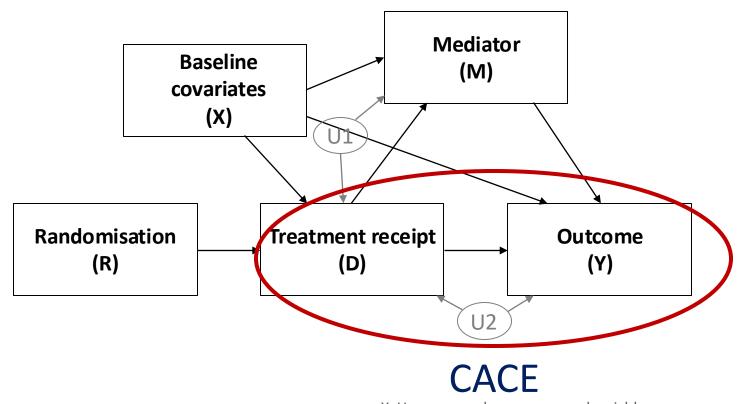
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### Combining mediation and non-adherence: Identification



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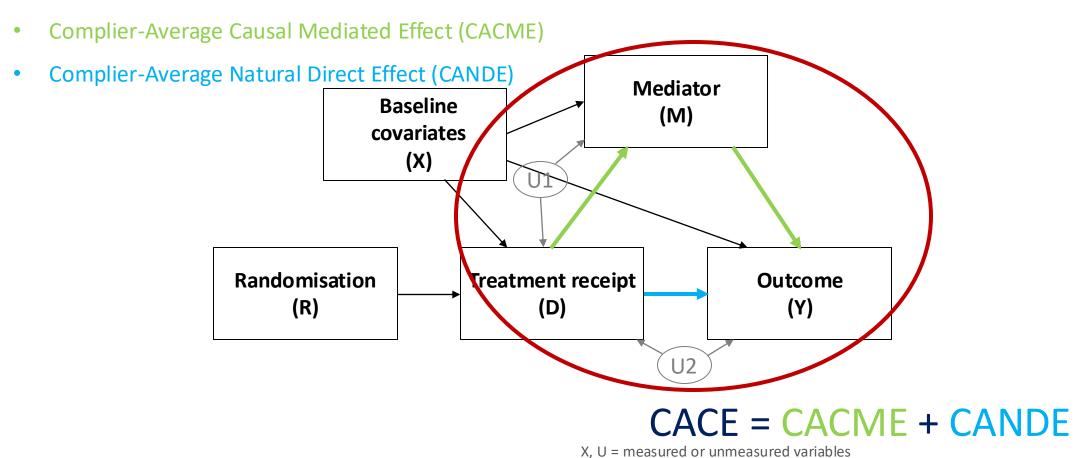
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### Combining mediation and non-adherence: Identification

#### The CACE can be partitioned into a:



Anca Chis Ster
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anca.m.chis ster@kcl.ac.uk

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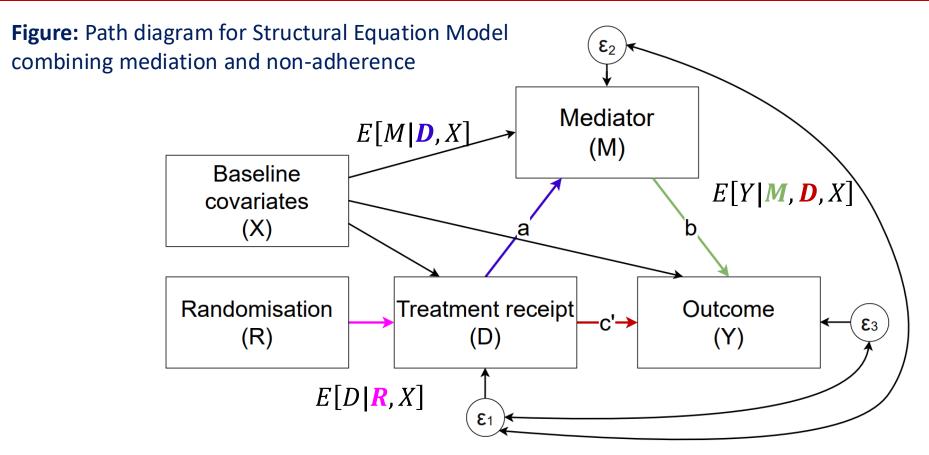
NDE = Natural Direct Effect

### Combining mediation and non-adherence: Assumptions

#### The CACME and CANDE can be identified under:

- (1) Conditionally ignorable treatment assignment
  - No variables that influence the randomisation variable
- (2) Monotonicity
  - No individuals who would receive the opposite intervention to the one offered
- (3) Exclusion restriction for non-compliers
  - Randomisation cannot directly influence the mediator or outcome variables
- (4) Conditionally ignorable observed mediator among compliers
  - No unmeasured confounding between the mediator and outcome

### Combining mediation and non-adherence: Estimation



CACME = a \* bCANDE = c'

## Illustrating example: The AVATAR study

- 150 participants randomised 1:1 to receive AVATAR
   therapy or supportive counselling for psychosis related
   symptoms
- The primary outcome was the total score on the
   Psychotic Symptom Rating Scales at 12 weeks and was analysed with the ITT principle
- 84% compliance (attended ≥3 of 6 sessions)
- Mediator of interest is the participants acceptance-based attitudes in relation to their auditory hallucinations



### Estimation via -sem-

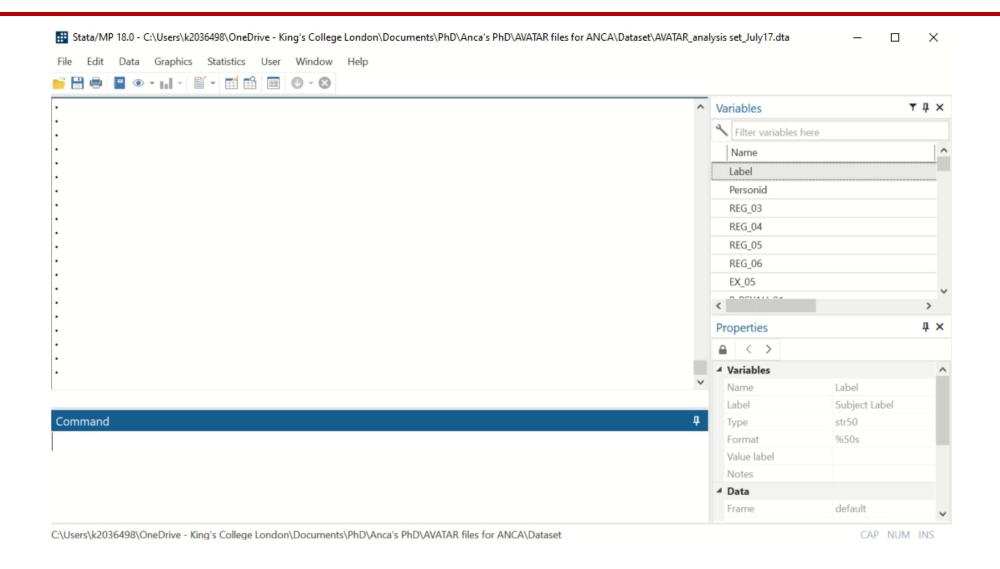
```
sem (R \rightarrow D) (D \rightarrow M) (D M \rightarrow Y), cov (e.D*e.M) cov (e.D*e.Y)
```

- The path (R -> D) implements IV theory and accounts for the endogeneity in D
- The covariances cov (e.D\*e.M) and cov (e.D\*e.Y) are essential as they allow for unmeasured confounding between D-M and D-Y

### estat teffects

- Produces many results and paths many are not relevant
- Can be difficult to identify which paths correspond to the CACME and CANDE

### Estimation via the -sem- command



### The -compmed- command: Motivation

- Estimating the CACME and CANDE requires
  - (1) Knowledge of SEMs and the SEM Stata package
  - (2) Fitting the correct Structural Equation Model
  - (3) Identifying the correct pathways that correspond to the CACME and CANDE
- compmed offers a standardised approach for estimating the CACME and CANDE in
   Stata using a single, more intuitive, and user-friendly programme.

# The -compmed- command: Syntax

```
compmed Y, mvar(M) dvar(D) rvar(R) cvars(varlist)
[, vce(vcetype) FULLoutput]
```

cvars (varlist) determines the list of covariates that are included in the outcome and mediator models.

vce (vcetype) calculates the standard error of the estimator. vcetype can be oim (observed information matrix), robust (Huber/White/sandwich estimator), bootstrap, or cluster (generalized Huber/White/sandwich estimator). If the option is not specified, the default is oim.

**FULLoutput** reports the full decomposition of effects into total, direct, and indirect effects, along with standard errors obtained by the delta method (Sobel, 1987). This option is equivalent to the 'estat teffects' command that is for use after running an the sem command in Stata.

## The -compmed- command: Demonstration



Anca Chis Ster King's College London anca.m.chis ster@kcl.ac.uk

### Missing data consideration

- Both -sem- and -compmed- undertake a complete-case analysis, i.e.,
   observations with missing values are dropped from the analysis
- The analysis is therefore valid provided the missing data are MCAR, or MAR (provided all variables that drive missingness are in the analysis model)
  - ✓ Information on non-adherence, a common predictor of missingness, is already included in the analysis model
- A Monte Carlo simulation study demonstrates that unbiased estimates of CACME,
   CANDE, and CACE can be obtained under the MCAR and MAR scenarios explored
   (full results and details not described here)

### **Final remarks**

- The CACE can be partitioned into a CACME and CANDE under a given set of assumptions and can be estimated with linear SEMs
- A new Stata program, compmed, provides a practical tool for undertaking causal mediation analysis with non-adherence
  - 1. Fits the correct SEM model
  - 2. Automatically identifies the paths that correspond to the CACME and CANDE
  - 3. Outputs a nice and simple table with these estimates

### Questions



anca.m.chis\_ster@kcl.ac.uk @anca\_chisster