

EXPLORATORY MEDIATION ANALYSIS WITH MANY POTENTIAL MEDIATORS

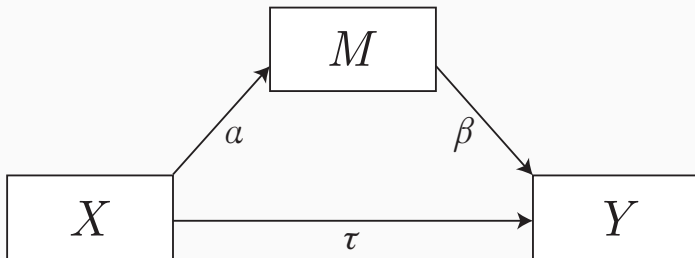
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MEDIATION

Q: When is M a mediator?



MacKinnon et al. (2002):

1. Causal steps: α & β
2. Difference in coefficients: $\tau - \tau|M$
3. Product of coefficients: $\alpha \times \beta$

VanderWeele (2015, p. 46): *“Also take into account $X \cdot M$ interaction!”*

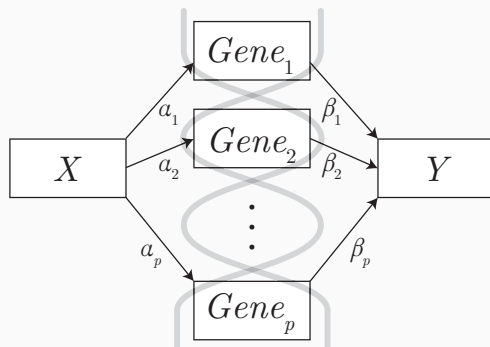
Theory-based **decision functions** using data from X, M, Y :

$$\mathcal{D}: \{\mathbf{x}, \mathbf{m}, \mathbf{y}\} \mapsto \{0, 1\}$$

(0 = not mediator, 1 = mediator)

MANY MEDIATORS

Q: When is *Gene_i* a mediator?

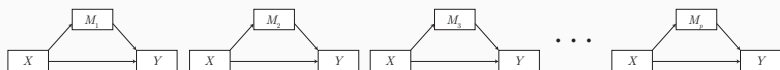


Preacher and Hayes (2008):

1. Fit the full SEM so your parameter estimates take all mediators into account
2. Select mediators using the estimated parameters

$\mathcal{D}(\mathbf{x}, \mathbf{m}_i, \mathbf{y})$ conditional on M_{-i}

p single mediator models



for (i in 1:p) $\mathcal{D}(\mathbf{x}, \mathbf{m}_i, \mathbf{y})$

The “filter” method (Guyon and Elisseeff, 2003)

Good

- Simple
- Quick
- Flexible

Bad

- Assumes uncorrelated mediators: won't work if mediation only visible conditionally

Jacobucci et al. (2016): We can now penalise SEM parameters

$$F_{\text{regsem}} = F_{\text{ML}} + \lambda P(\cdot)$$

Serang et al. (2017): We can use this to select mediators! Put a lasso penalty on α and β

The “XMed” method

Good

- "Full" SEM
- Does not assume uncorrelated mediators
- Regularisation is hip

Bad

- What are we actually optimising for?
- Find M for which α OR β but we want α AND β .

Our contribution:

$\mathcal{D}(\mathbf{x}, \mathbf{m}_i, \mathbf{y})$ conditional on M_{-i}

Insight from regularisation literature (Hastie et al., 2015):

conditional parameter == parameter estimated on **residual**

Idea:

```
1 sel ← rep(0, p)
2
3 while (!convergence) {
4   for (i in 1:p) {
5     x_res ← x - M[, sel] %*% beta_x_sel
6     y_res ← y - M[, sel] %*% beta_y_sel
7     sel[i] ← decisionFunction(x_res, M[, i], y_res)
8   }
9 }
```

COORDINATE-WISE MEDIATION FILTER

for each mediator Coordinate-wise
perform the decision function Mediation
throw it out if 0 Filter

conditional on the other selected mediators

repeat until convergence

Good

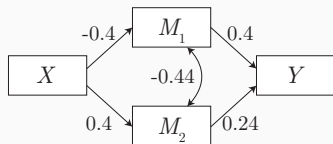
- Uses theoretically relevant \mathcal{D}
- Does not assume uncorrelated mediators

Bad

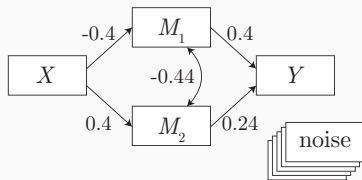
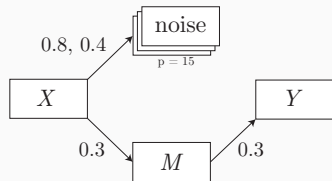
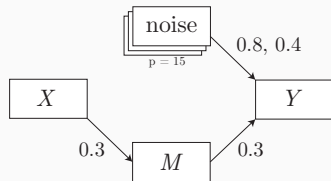
- Nonconvergence

SIMULATION

Conditional-only



High-dimensional

Noise (α paths)Noise (β paths)

IMPLEMENTATION

```
> devtools::install_github("vankesteren/cmfilter")
> library(cmfilter)
> res <- cmf(x, M, y, verbose = TRUE)

# CMF Algorithm
#
# -----
#
# 1 0 0 1 1 1 1 0 1 0 0 0 0 1 1 0 0
# 1 1 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
# 0 1 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
# 0 1 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
#
# Algorithm converged
#
# -----
```

CONCLUSION

- Novel method for selecting among many mediators
- Flexible choice of \mathcal{D}
- Conditional on M_{-i}
- Stable in boundary cases
- Traditional power/type-I tradeoff

- Group lasso
- Bayesian regularisation prior on $\alpha \times \beta$

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- Jacobucci, R., Grimm, K. J., and McArdle, J. J. (2016). Regularized Structural Equation Modeling. *Structural Equation Modeling*, 23(4):555–566.
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QUESTIONS

Suppression

Method	Power (M_1)	Power (M_2)
Full SEM	1.00	0.99
Filter	0.99	0.13
XMed	1.00	0.99
CMF	1.00	0.91

Noise (α paths)

Method	Power	FPR	PPV
Full SEM	0.20	0.11	0.11
Filter	0.27	0.09	0.17
XMed	0.67	0.34	0.12
CMF	0.17	0.06	0.17

Noise (β paths)

Method	Power	FPR	PPV
Full SEM	0.08	0.01	0.32
Filter	0.44	0.02	0.58
XMed	0.49	0.12	0.22
CMF	0.41	0.02	0.58

High-dimensional data

Method	Power (M_1)	Power (M_2)	FPR	PPV
Full SEM	NA	NA	NA	NA
Filter	0.91	0.07	2.4e-3	0.30
XMed	NA	NA	NA	NA
CMF	0.82	0.06	1.8e-3	0.32