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Latent PCMCI



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machine _{in} climate learning ⁱⁿ science

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Content

Recap: PCMCI



Fundamental Idea Examples

LPCMCI

- Setting 2 Idea
 - Algorithm

Conclusion



Weaknesses

I. Recap PCMCI: Fundamental Idea and Example

Recap: PCMCI

Goal: learn causal graph of given variables with conditional independence tests

- FullCI has low detection power and estimates low effect size (too many conditionals)
- PC (Peter-Clarke) has high false positive rate for high autocorrelation (broken iid assumption)
- → PCMCI:
 - First PC: Remove irrelevant conditions \rightarrow learn superset of parents $\hat{\mathscr{P}}(X_j)$ for each variable
 - Then MCI (Momentary Conditional Independence):

MCI:
$$X_{t-\tau}^i \perp X_t^j \mid \widehat{\mathscr{P}}(X_t^j) \setminus \{X_{t-\tau}^i\}, \widehat{\mathscr{P}}(X_{t-\tau}^i).$$

Establish cond. independence

Reduce false positive rate

Start with complete undirected graph Three steps:

- 1. Identify the skeleton
- 2. Identify immoralities and orient them
- Orient qualifying edges that are incident on colliders



PC Example: Identify the skeleton



PC Example: Orient Edges



We have found all colliders



https://www.researchgate.net/figure/Anexample-for-PC-algorithm-Figure-from-S pirtes-et-al-2000-Chapter-5_fig7_32182 8295

Source:

Estimated Graph

I. Latent PCMCI: Setting, Idea and Algorithm

Assumptions: Faithfulness, Causal Markov Condition, Causal Stationarity

 $V_t^j = f_j(pa(V_t^j), \eta_t^j) \quad ext{with } j = 1, \dots, ilde{N} \ .$

Only observe $\{X_1,\ldots,\,X_N\}\subseteq ig\{V_1,\ldots,\,V_{ ilde N}ig\}$

 $\mathscr{M}(\mathscr{G})$ Maximal Ancestral Graph (MAG) after marginalizing over unobserved variables and all t'<t - τ_{max}

 $\mathscr{P}(\mathscr{G})$ Maximally informative Partial Ancestral Graph, which can be obtained by running FCI orientation rules (enforce time order and repeating adj.)

May include contemp. links and unobs. confounders

Detection power of true link depends on

- Sample size
- CI tests' sign. level
- CI tests' estim. dimension
- Effect size $\min_S I(A, B \mid S)$ over used cond. sets S

Problem often: low signal-to-noise ratio (e.g. autocorrelation) due to unfortunate choice of conditioning sets leading to low effect size,

Leads to missing links, which lead to false positives and wrong orientations

- → Carefully choose conditioning sets:
 - Discard known non-ancestors
 - Include known ancestors ALWAYS

by entangling edge removal and orientation phase

multiple iterations k, collecting knowledge

Algorithm 1 LPCMCI

Require: Time series dataset X = {X¹,..., X^N}, maximal considered time lag τ_{max}, significance level α, CI test CI(X, Y, S), non-negative integer k
1: Initialize C(G) as complete graph with Xⁱ_{t-τ} ◦^L→X^j_t (0 < τ ≤ τ_{max}) and Xⁱ_{t-τ} ◦² ◦ X^j_t (τ = 0)
2: for 0 ≤ l ≤ k − 1 do
3: Remove edges and apply orientations using Algorithm S2
4: Repeat line 1, orient edges as Xⁱ_{t-τ} ²→X^j_t if Xⁱ_{t-τ} ^{*}→X^j_t was in C(G) after line 3
5: Remove edges and apply orientations using Algorithm S2
6: Remove edges and apply orientations using Algorithm S3

7: return PAG $\mathcal{C}(\mathcal{G}) = \mathcal{P}(\mathcal{G}) = \mathcal{P}(\mathcal{G})_{statAO}^{\tau_{\max}}$

After line 5, all links are correctly removed for pairs of variables where one is an ancestor of the other

- Introduce node ordering and 'middle marks' to save what is known about the link
- There exist several orientation rules for orienting and deleting edges
- Iterate until all middle marks have been removed or reached hyperparam. k
 - In practice for finite samples, larger k does not imply better results :((see river example)

Algorithm S2 Ancestral removal phase

```
Require: LPCMCI-PAG \mathcal{C}(\mathcal{G}), memory of minimal test statistic values I^{\min}(\cdot, \cdot), memory of sepa-
     rating sets SepSet(\cdot, \cdot), time series dataset \mathbf{X} = {\mathbf{X}^1, \dots, \mathbf{X}^N}, maximal considered time lag
     \tau_{\max}, significance level \alpha, CI test CI(X, Y, S)
 1: repeat starting with p = 0
 2:
          for -1 < m < \tau_{\max} do
               for all ordered pairs of variables (X_{t-\tau}^i, X_t^j) adjacent in \mathcal{C}(\mathcal{G}) with X_{t-\tau}^i < X_t^j do
 3:
                    if (m = -1 \text{ and } i \neq j) or (m \ge 0 \text{ and } \tau \neq m \text{ or } i = j) then continue with next pair
 4:
 5:
                    \mathcal{S}_{def} = pa(\{X_{t-\tau}^i, X_t^j\}, \mathcal{C}(\mathcal{G}))
                    if the middle mark is '?' or 'L' then
 6:
                         S_{search} = apds_t(X_t^j, X_{t-\tau}^i, \mathcal{C}(\mathcal{G})) \setminus S_{def}, ordered according to I^{\min}(X_t^j, \cdot)
if |S_{search}| < p then update middle mark with 'R' according to Lemma [S8]
 7:
 8:
 9:
                         for all subsets S \subseteq S_{search} with |S| = p do
                              (p-value, I) \leftarrow CI(X_{t-\tau}^i, X_t^j, S \cup S_{def})
10:
                              I^{\min}(X_{t-\tau}^{i}, X_{t}^{j}) = I^{\min}(X_{t}^{j}, X_{t-\tau}^{i}) = \min(|I|, I^{\min}(X_{t-\tau}^{i}, X_{t}^{j}))
11:
                              if p-value > \alpha then
12:
13:
                                   mark edge for removal, add S \cup S_{def} to SepSet(X_{t-\tau}^i, X_t^j)
14:
                                   break innermost for-loop
15:
                    repeat lines 6 - 14 with X_{t-\tau}^i and X_t^j as well as 'R' and 'L' swapped
16:
               remove all edges that are marked for removal from \mathcal{C}(\mathcal{G})
17:
           if any edge has been removed in line 16 then
18:
               run Alg. S4 using [APR, MMR, R8', R2', R1', R9', R10'], orient lagged links only
19:
               let p = 0
20:
           else increase p to p+1
21: until there are no other middle marks than '!' or '' (empty)
22: run Alg. S4 using [APR, MMR, R8', R2', R1', R0'd, R0'c, R3', R4, R9', R10', R0'b, R0'a]
23: return \mathcal{C}(\mathcal{G}), I^{\min}(\cdot, \cdot), SepSet(\cdot, \cdot)
```



A) Improves with larger k if assumptions hold

B-D) LPCMCI better for contemp. links for k large enough

Controls FPR

Figure 2: Results of numerical experiments for (A) LPCMCI(k) for different k, LPCMCI compared to SVAR-FCI and SVAR-RFCI for (B) varying autocorrelation, for (C) number of variables N, and for (D) maximum time lag τ_{max} (other parameters indicated in upper right of each panel).

Conclusion: LPCMCI

- Like PCMCI for setting with contemporaneous links and confounders
 - → More general and realistic
- Idea: Always condition on known parents
- Uses middle marks and complicated link orientation and removal rules
- Higher recall for highly autocorr. time series
- order-independent, returns optimal PAG for infinite sample size
- additional hyperparam. k (number of iterations)
- no finite sample guarantees (conv. rate arbitrarily slow for almost unfaithful distr.)

LPCMCI: https://arxiv.org/abs/2007.01884

PCMCI: https://advances.sciencemag.org/content/5/11/eaau4996

Thank you!



Important Realisations

- Weak autocorrelation, better recall
- Correct false positive rates in MCI step
- Assumptions: Causal Markov Condition, Faithfulness, causal stationarity, no contemporaneous causal links, no hidden variables Then: consistent
- LPCMCI: Causal Markov Condition, Faithfulness, causal stationarity •
- Findings of non-causality hold for weaker assumptions (faithfulness, test flexible enough), thus more reliable
- Fast PC: only choose conditions with highest association
- Choose tau_max large, only longer runtime, not higher dim. Small alpha means few false positives, but less true positives
- Convergence rate to consistency can be made arbitrarily slow by almost unfaithful distributions
- Autocorrelation breaks iid assumption in tests, thus in ParCorr t distr. Has fewer degrees of freedom hence get more false positives: adjust the degrees of freedom in some way, using pre-whitening, or by block-shuffling. While these approaches help to some extent for the simple bivariate case, they fail in the multivariate case that is relevant for causal discovery https://arxiv.org/abs/1407.0742 (MCI good option)