

Suppose P is faithful to DAG G^0 . At each stage, the skeleton of G^0 must be a subgraph of \hat{G} . Edges (j, k) remaining at termination will have

$$Z_j \perp\!\!\!\perp Z_k \mid Z_S \quad \forall S \subseteq (\text{adj}(\hat{G}, k) \setminus \{j\}) \cap (\text{adj}(\hat{G}, j) \setminus \{k\})$$

In particular, $Z_j \perp\!\!\!\perp Z_k \mid Z_S \quad \forall S \subseteq (\text{adj}(G^0, k) \setminus \{j\}) \cap (\text{adj}(G^0, j) \setminus \{k\})$

So such j, k must be adjacent in G^0 , so \hat{G} will be the skeleton of G^0 .

Prop 36 Suppose we have $j - l - k$ in the skeleton of a DAG (j, k not adjacent).

i) If $j \rightarrow l \leftarrow k$, then any S that d-separates j and k cannot contain l .

ii) If $\exists S$ that d-separates j and k but $l \notin S$, then $j \rightarrow l \leftarrow k$.

To find α -structures, we perform

For all $j - l - k$ in \hat{G} do

If $l \notin S(j, k)$, then orient $j \rightarrow l \leftarrow k$

Other edges can be oriented using e.g. the acyclicity of DAGs.

A simple version of the PC algorithm replaces querying conditional independencies with conditional independence tests. If the data are multi-variate normal, these tests are based on partial correlation.

$$\text{Corr}(Z_j, Z_k \mid Z_S) = 0 \iff Z_j \perp\!\!\!\perp Z_k \mid Z_S$$

Given $x_1, \dots, x_n \stackrel{iid}{\sim} Z$, can estimate the partial correlation by regressing X_j and X_k on X_S and computing the correlation between the residuals.

4 Multiple testing

In many modern applications, often interested in testing many hypotheses H_1, \dots, H_m simultaneously when m_0 of them are true nulls.

| | Claim non-sig | Claim sig | Total |
|-------------|---------------|-----------|-----------|
| True nulls | N_{00} | N_{01} | m_0 |
| False nulls | N_{10} | N_{11} | $m - m_0$ |
| Total | $m - R$ | R | m |

Suppose we have p-values p_1, \dots, p_m and $H_i : i \in I_0$ are the true nulls, so

$$P(p_i \leq \alpha) \leq \alpha \quad \forall \alpha \in [0, 1], i \in I_0 \quad |I_0| = m_0$$

Traditionally, approaches have aimed to satisfy

$$\text{Familywise error rate (FWER)} = P(N_{01} \geq 1) \leq \alpha$$

Bonferroni correction rejects H_i if $p_i \leq \frac{\alpha}{m}$.

Thm 37 Using Bonferroni correction

$$P(N_{01} \geq 1) \leq E(N_{01}) \leq \frac{m \cdot \alpha}{m} \leq \alpha$$

Markov

Proof

$$E(N_{01}) = E\left(\sum_{i \in I_0} \mathbb{1}_{\{p_i \leq \frac{\alpha}{m}\}}\right)$$

$$= \sum_{i \in I_0} \underbrace{P(p_i \leq \frac{\alpha}{m})}_{\leq \frac{\alpha}{m}}$$

$$\leq \frac{m \cdot \alpha}{m} \leq \alpha$$

□

4.1 Closed testing procedure

Given family of hypotheses $\{H_i\}_{i=1}^m$ define its closure to be

$$\{H_I : I \subseteq \{1, \dots, m\}, I \neq \emptyset\}$$

where $H_I = \bigcap_{i \in I} H_i$ is the intersection hypothesis i.e. H_I is the hypothesis that all

$H_i, i \in I$ are true.