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*Exploiting causality methods for
knowledge discovery
from observational data*

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Why study CAUSALITY?

Why study CAUSALITY?

Because

CORRELATION IS NOT CAUSATION

CORRELATION IS NOT CAUSATION



CORRELATION IS NOT CAUSATION

..but sometimes it gives strong hints



Because

CORRELATION IS NOT CAUSATION

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... and understanding how things work (causation) is the ultimate goal of science

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1. Which variables are “important” in my scenario?
2. How does changing one variable affect the system?

... and understanding how things work (causation) is the ultimate goal of science

1. Which variables are “important” in my scenario?
≈ Feature selection

2. How does changing one variable affect the system?

Correlational approaches fail at discovering “important” variables

Linear regression

$$\hat{y}_i = w_0 + w_1 x_1 + \dots + w_N x_N$$

Trained by minimizing the mean squared error

$$\mathbf{w} = \underset{\mathbf{w}}{\operatorname{argmin}} \frac{1}{N} \sum_i (y_i - \hat{y}_i)^2$$

Correlational approaches fail at discovering “important” variables

Linear regression

$$\hat{y}_i = w_0 + w_1 x_1 + \cdots + w_N x_N$$

LASSO focuses on \mathbf{w} such that

$$\mathbf{w} = \underset{\mathbf{w}}{\operatorname{argmin}} \frac{1}{N} \sum_i (y_i - \hat{y}_i)^2 - \lambda \sum_i \|w_i\|$$

Correlational approaches fail at discovering “important” variables

Solution obtained minimizing MSE

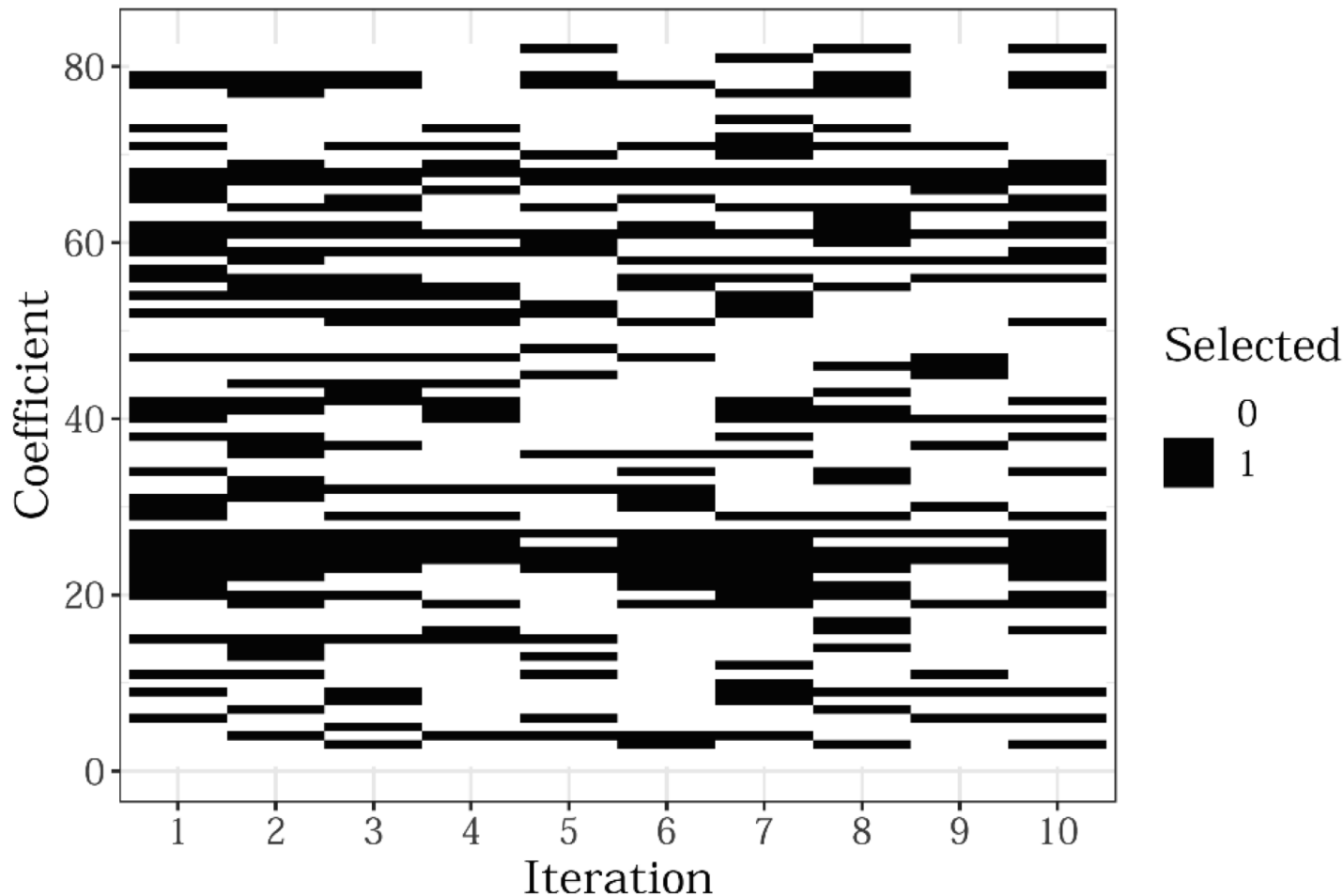
$$\hat{y}_i = w_0 + w_1 x_1 + w_2 x_2 + \cdots + w_{N-1} x_{N-1} + w_N x_N$$

Solution obtained minimizing LASSO

$$\hat{y}_i = w_0 + w_1 x_1 + w_2 x_2 + \cdots + w_{N-1} x_{N-1} + w_N x_N$$

Correlational approaches fail at discovering “important” variables

FIGURE 2: INCLUDED PREDICTORS IN A LASSO REGRESSION ACROSS TEN SAMPLES FROM THE SAME POPULATION



Gillis, T.B. and Spiess, J.L., 2019. Big data and discrimination. *The University of Chicago Law Review*, 86(2), pp.459-488

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1. Which variables are “important” in my scenario?

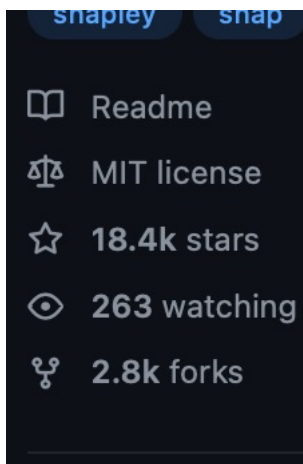
2. How does changing one variable affect the system?

≈ Interpretation

Correlational approaches fail at discovering effects



SHAP



A unified approach to interpreting model predictions

[SM Lundberg](#), [SI Lee](#) - *Advances in neural information ...*, 2017 - [proceedings.neurips.cc](#)

Understanding why a model makes a certain prediction can be as crucial as the prediction's accuracy in many applications. However, the highest accuracy for large modern datasets is often achieved by complex models that even experts struggle to interpret, such as ensemble or deep learning models, creating a tension between accuracy and interpretability. In response, various methods have recently been proposed to help users interpret the predictions of complex models, but it is often unclear how these methods are related and ...

☆ Save [Cite](#) Cited by 10824 [Related articles](#) [All 17 versions](#) [↔](#)

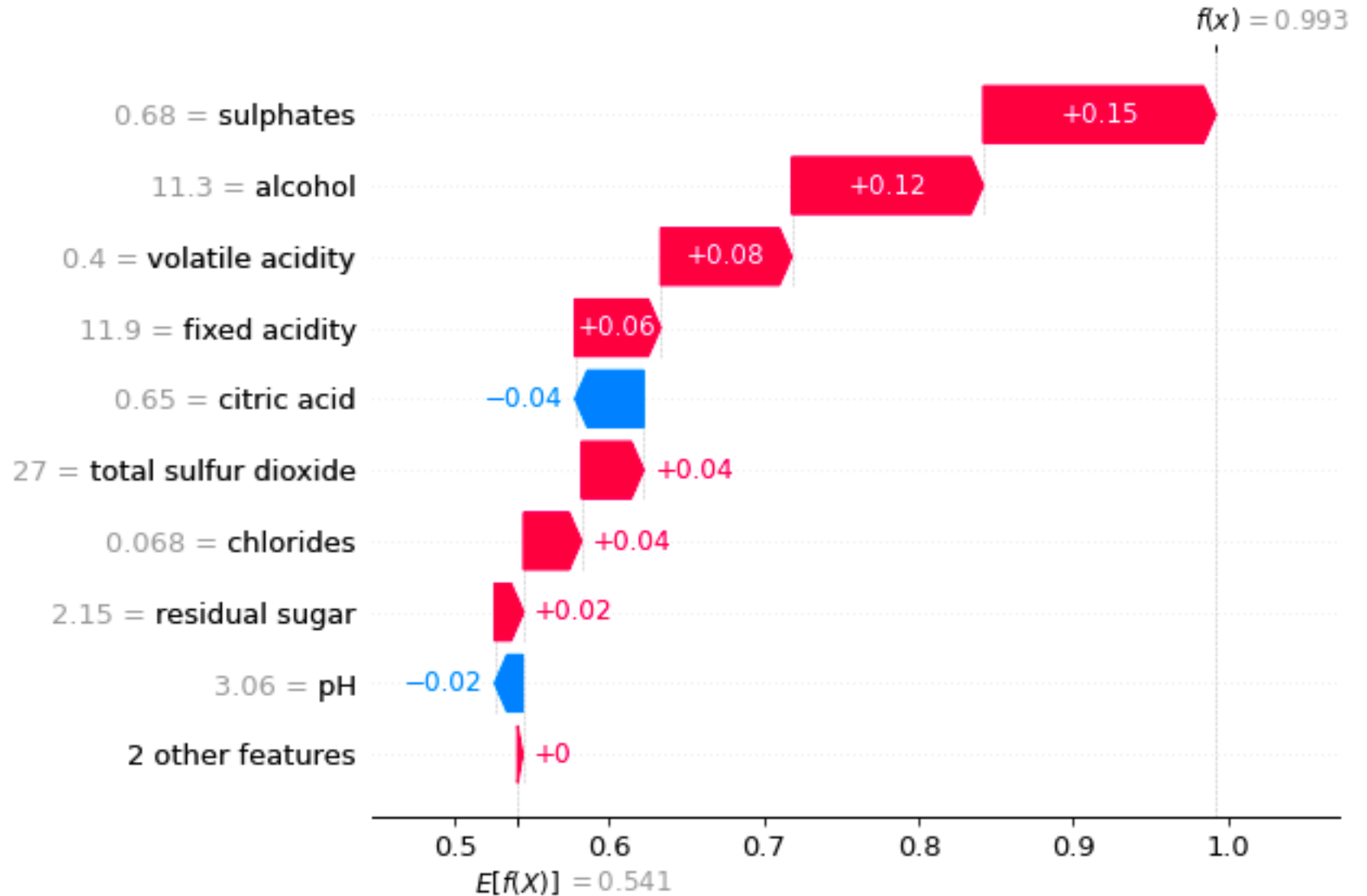
[\[PDF\] neurips.cc](#)

Correlational approaches fail at discovering effects



SHAP

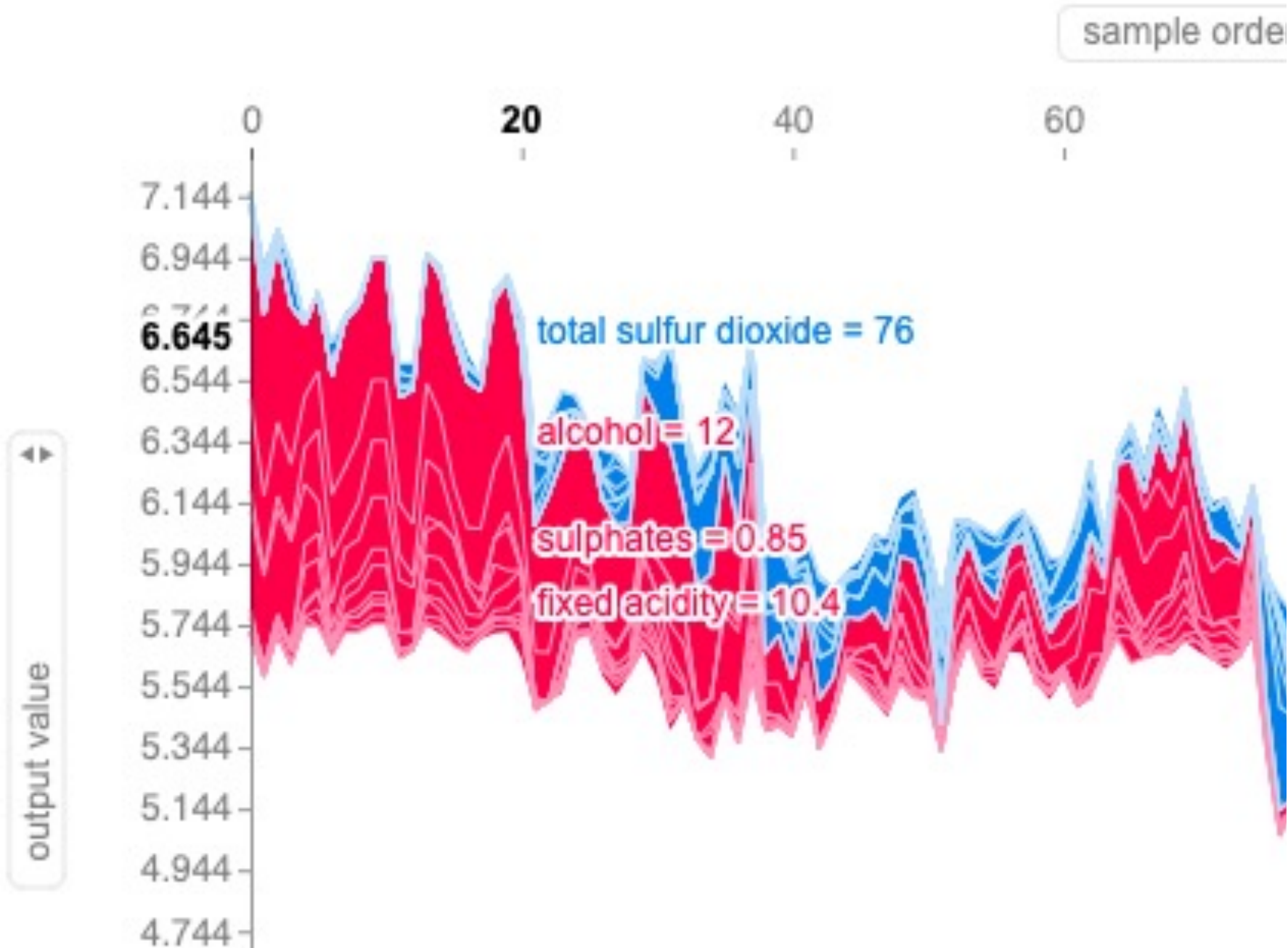
The prediction is 0.9925615



Correlational approaches fail at discovering effects



SHAP



Correlational approaches fail at discovering effects

```
[96] #!pip install shap
import shap
import numpy as np
import sklearn

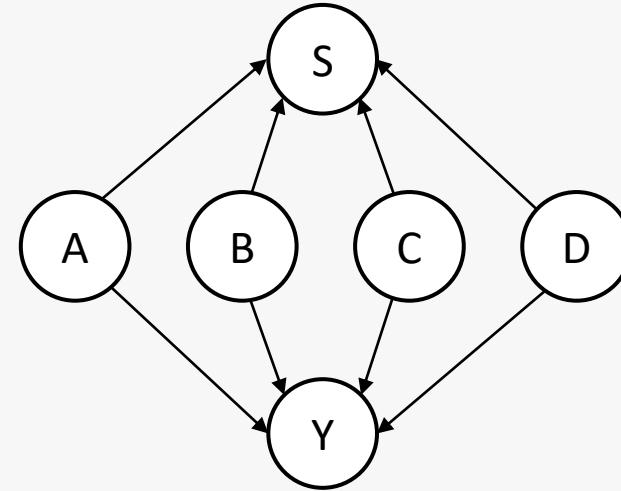
shap.initjs()

n_el = 250
p = 0.8
a = np.random.binomial(1,p,n_el)
b = np.random.binomial(1,p,n_el)
c = np.random.binomial(1,p,n_el)
d = np.random.binomial(1,p,n_el)
s = (A  $\wedge$  B  $\wedge$  C  $\wedge$  D)  $\oplus$  B(0.1)

y = (A  $\wedge$  B  $\wedge$  C  $\wedge$  D)  $\oplus$  B(0.1)

X = np.reshape(np.array((a,b,c,d,s)),(5,-1)).T

lr = sklearn.linear_model.LogisticRegression().fit(X,y)
```



Correlational approaches fail at discovering effects

✓
0s



`n_el = 1000`

`p = 0.8`

`a2 = np.random.binomial(1,p,n_el)`

`b2 = np.random.binomial(1,p,n_el)`

`c2 = np.random.binomial(1,p,n_el)`

`d2 = np.random.binomial(1,p,n_el)`

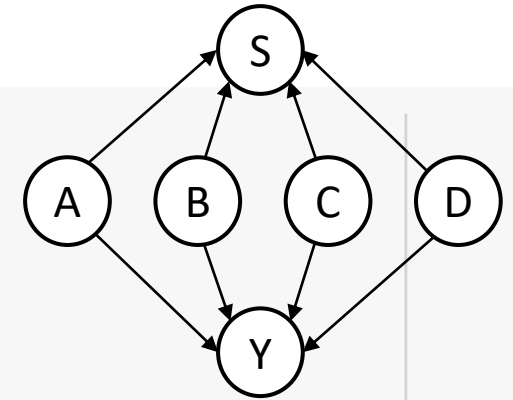
`s2 = (A2 ∧ B2 ∧ C2 ∧ D2) ⊕ B(0.1)`

`y2 = (A2 ∧ B2 ∧ C2 ∧ D2) ⊕ B(0.1)`

`X2 = np.reshape(np.array((a2,b2,c2,d2,s2)),(5,-1)).T`

`lr.score(X2,y2)`

↪ 0.842



`lr.coef_`

Coefficients of

A

B

C



D

S

`array([[1.11347288, 1.28590981, 0.98832866, 1.25790585, 2.09644493]])`

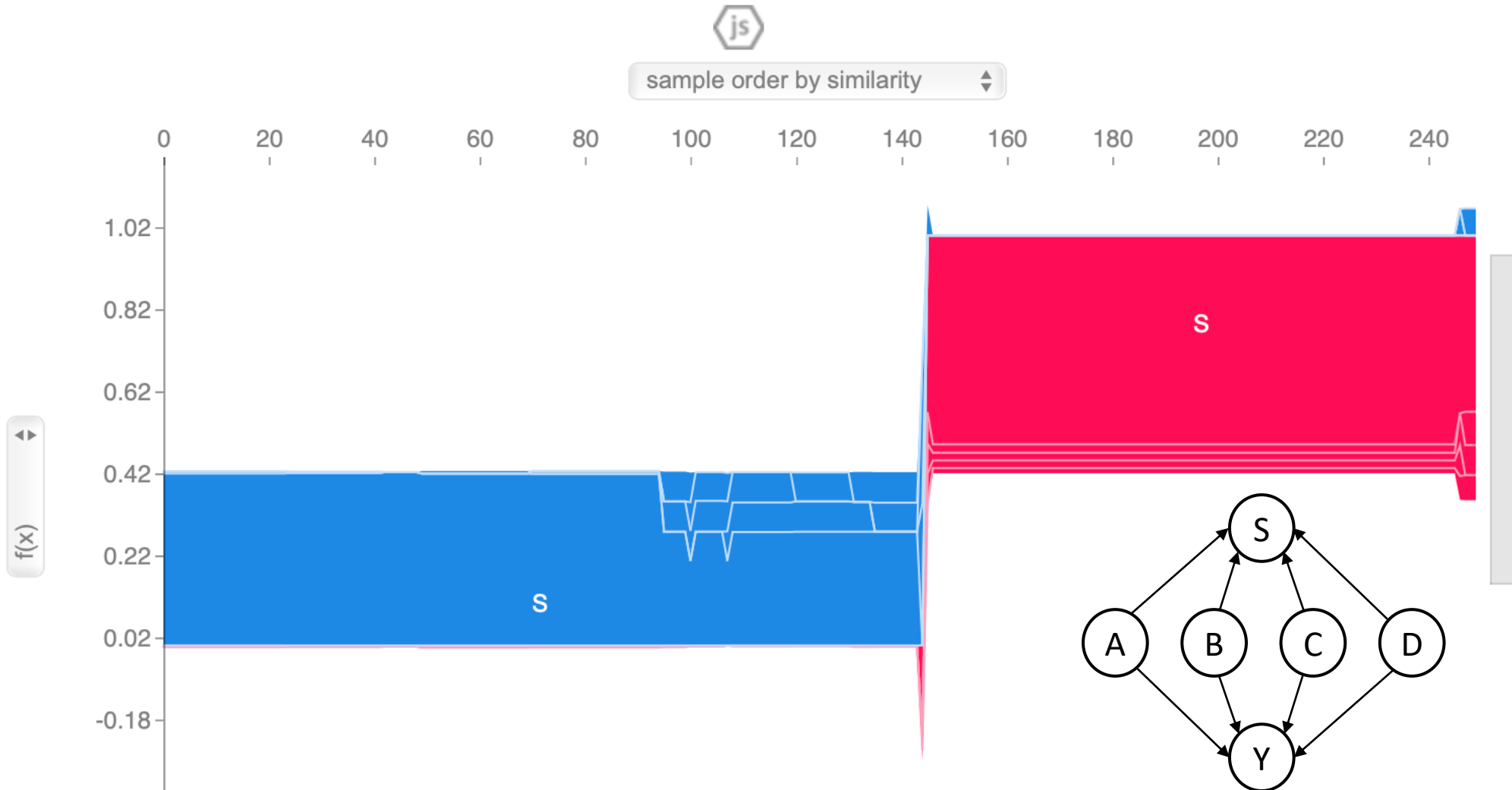
Correlational approaches fail at discovering effects

✓
0 s

[100]

```
shap.initjs()
```

```
shap.force_plot(explainer.expected_value, shap_values, pd.DataFrame(X, columns=["A", "B"
```



...and understanding how things work (causation) is the ultimate goal of science

1. Which variables are “important” in my scenario?
2. How does changing one variable affect the system?

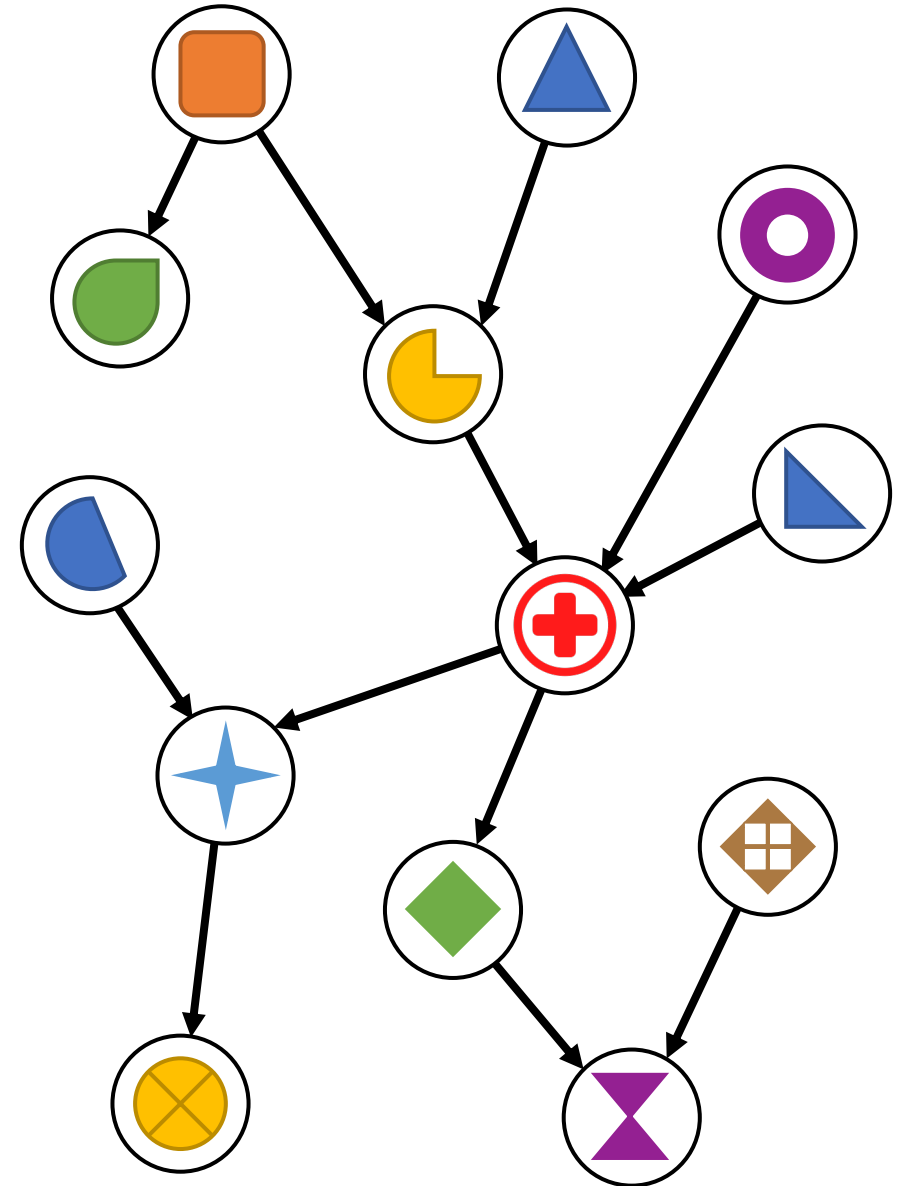
Causal Bayesian Networks
are convenient ways to model
causal relationships among
variables

$$BN = \langle G, p \rangle$$

where

$$p(X_i) = f(pa(X_i), \epsilon)$$

Interventional probability

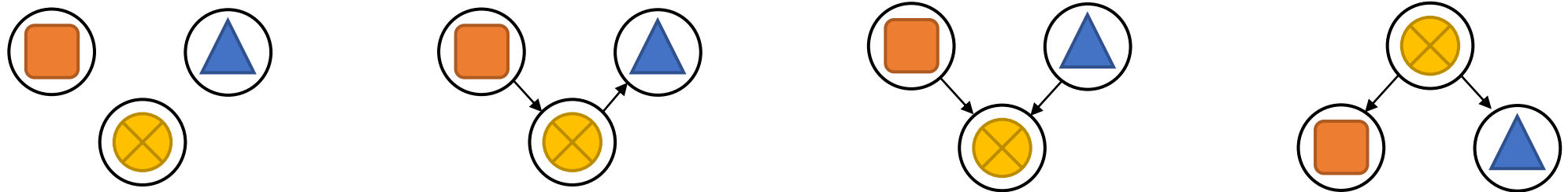


(Some) main objectives of causal techniques:

- Which variables are “important” in my scenario?

Structure discovery:

How are variables linked?



- How does changing one variable affect the system?

Effect estimation: How changing  affects  ?

Develop algorithms to:

- (Structure discovery) Discover causally related variables to a target
- (Effect estimation) Evaluate effect of causal rules

From observational data and providing guarantees on the results

Causal BNs are built using **interventional data** (e.g. setting variable $X_i = x_i$)

Observational data is much more common

Th. [informal] If spurious dependencies are removed, observational and interventional probability distributions are equivalent

Develop algorithms to:

- (Structure discovery) Discover causally related variables to a target
- (Effect estimation) Evaluate effect of causal rules

From observational data and providing guarantees on the results

Statistical guarantees are NOT just evaluation of performances

Guarantees are fundamental to gain users trust

Typically, algorithms with guarantees focus on False Discoveries (or False Positives)

Develop algorithms to:

- (Structure discovery) Discover causally related variables to a target
- (Effect estimation) Evaluate effect of causal rules

From observational data and providing guarantees on the results

Develop algorithms to:

- **(Structure discovery) Discover causally related variables to a target**
- (Effect estimation) Evaluate effect of causal rules

From observational data and providing guarantees on the results

Develop algorithms to:

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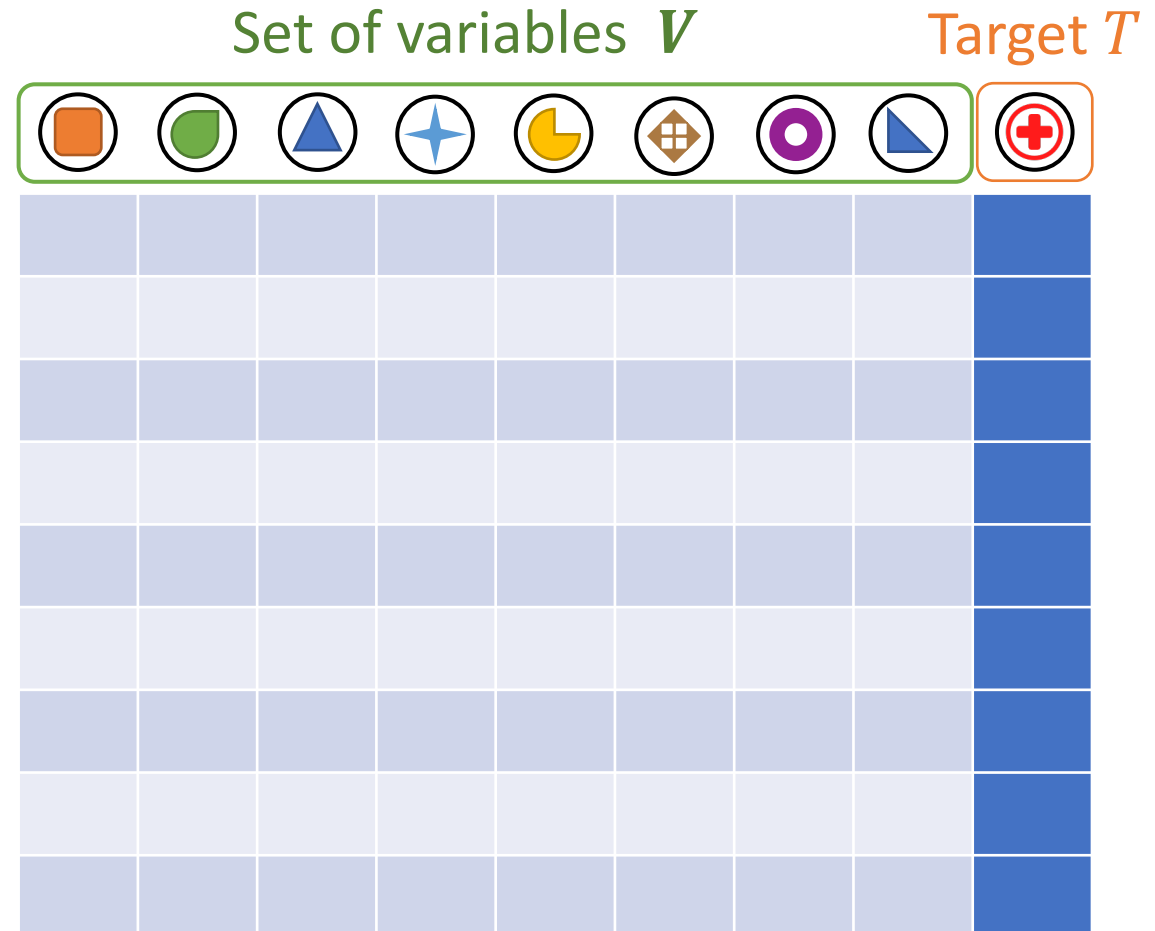
Simionato Dario, and Fabio Vandin. "Bounding the Family-Wise Error Rate in Local Causal Discovery using Rademacher Averages."

Accepted at ECML PKDD 2022 – **Best paper award**

Problem Definition





Task: Given a dataset of observations of variables V , find those **causally** related to T with **guarantees** on the result (e.g. no false positives)

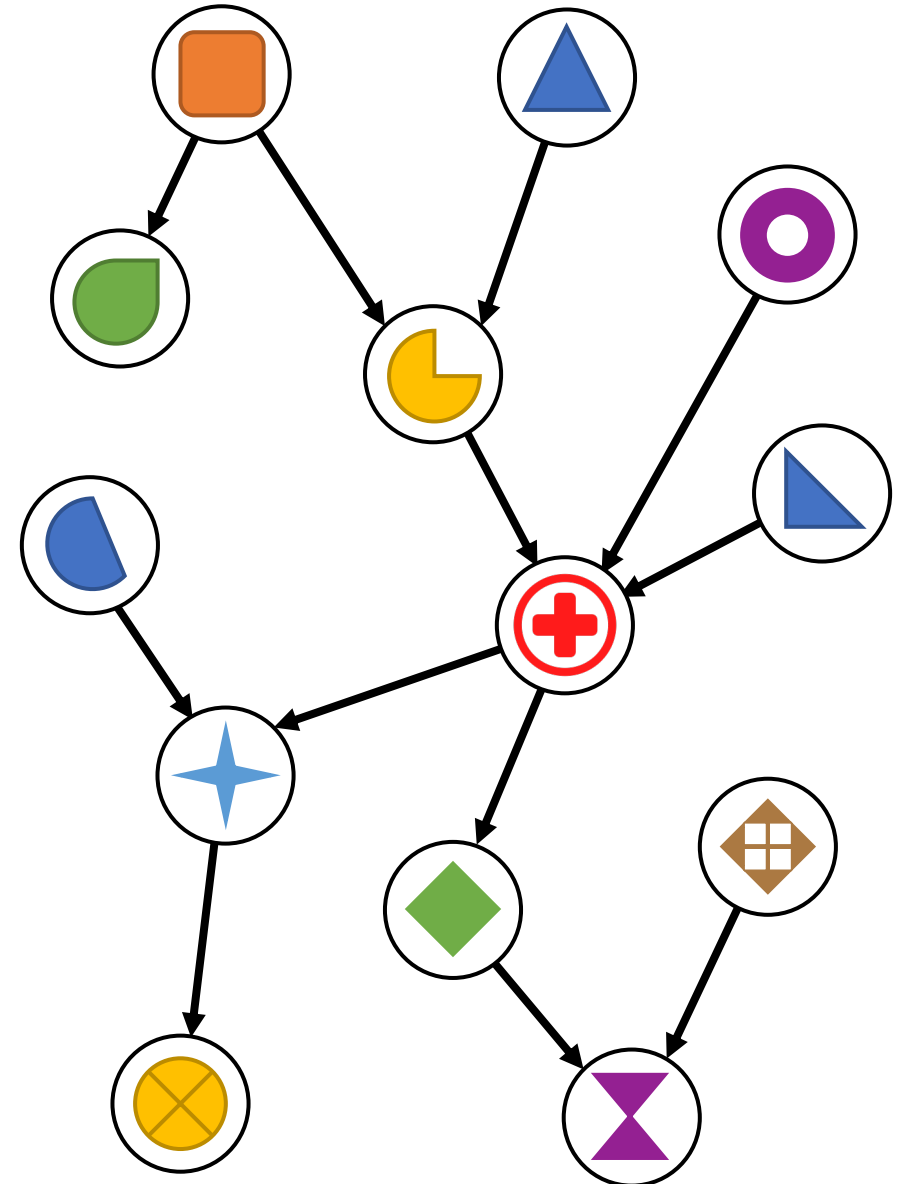
Useful in: Biology, medicine, neuroscience



Problem Definition





Causal Bayesian Networks represent cause-consequence relations between variables

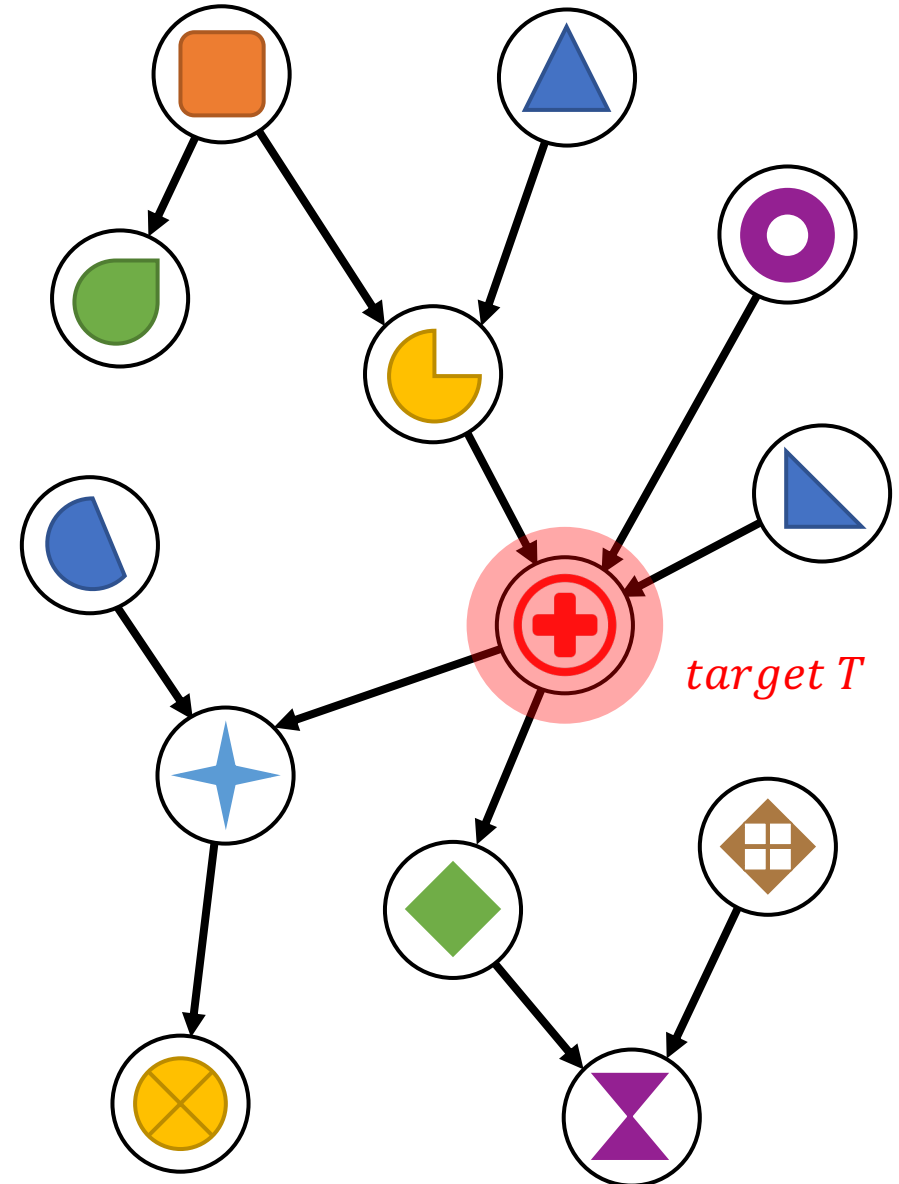
Informally, if  is a **cause** of , then fixing all variables values and changing the value of  leads to a variation on the values of 



Problem Definition

Causal Bayesian Networks represent cause-consequence relations between variables

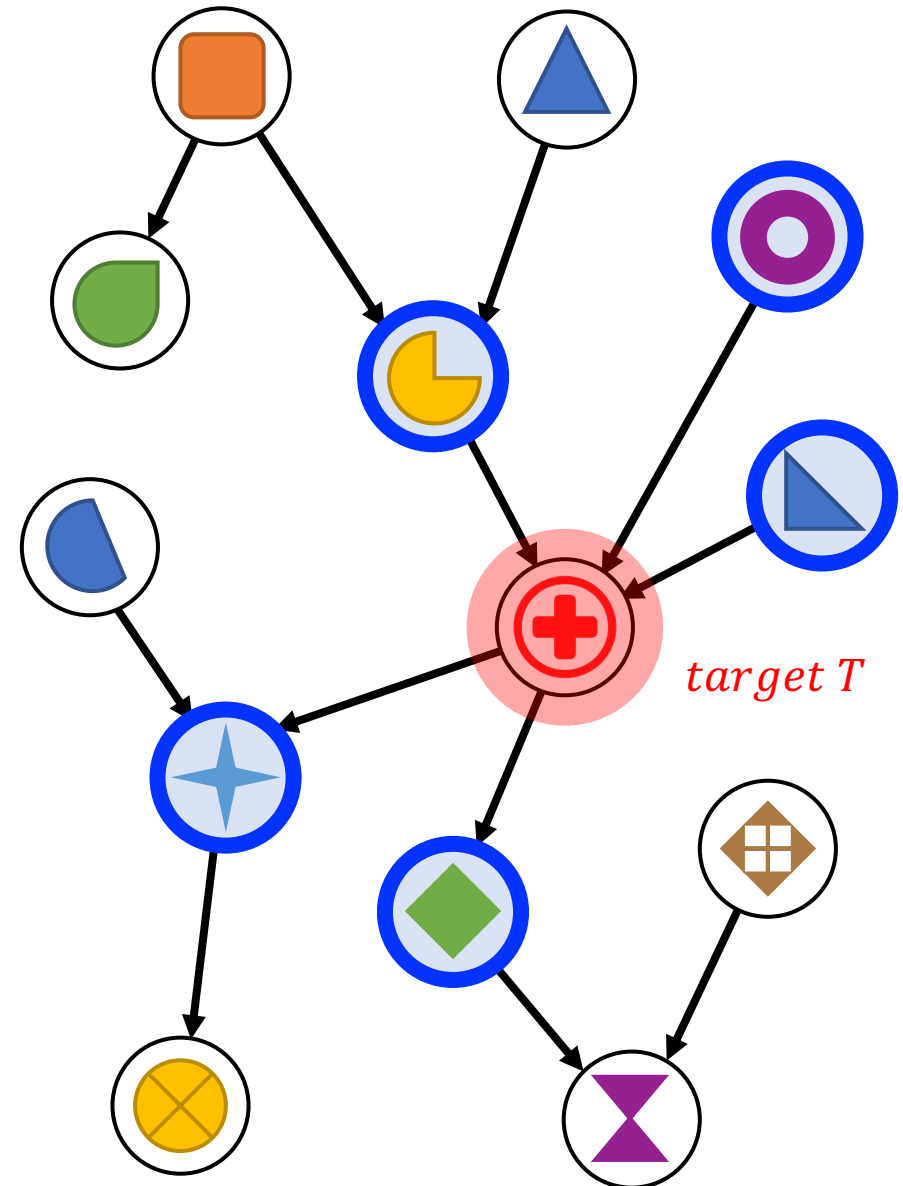
Informally, if  is a **cause** of , then fixing all variables values and changing the value of  leads to a variation on the values of 



Local causal discovery focuses on:

- **Parent – Children set of T** $PC(T)$
Parents(T) + Children(T)

Contains direct causes and consequences of T



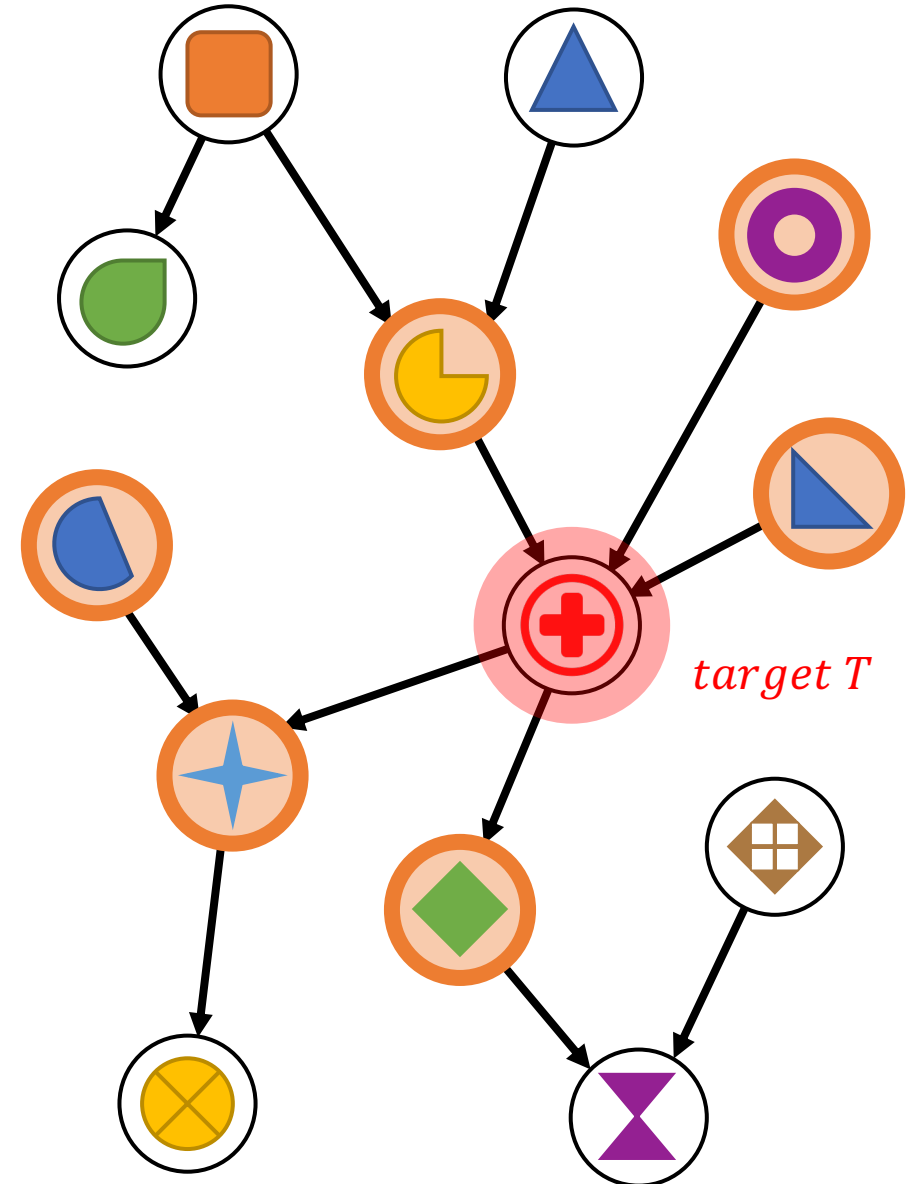
Local causal discovery focuses on:

- **Parent – Children set of T** $PC(T)$
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Contains direct causes and consequences of T

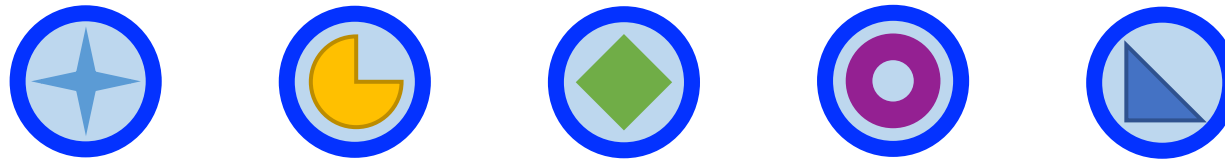
- **Markov Boundary of T** $MB(T)$
 $PC(T)$ + Spouses(T)

Optimal set for the prediction of T



Given a dataset and a target T , our task is to discover:

- **Parent – Children set of T** $PC(T)$

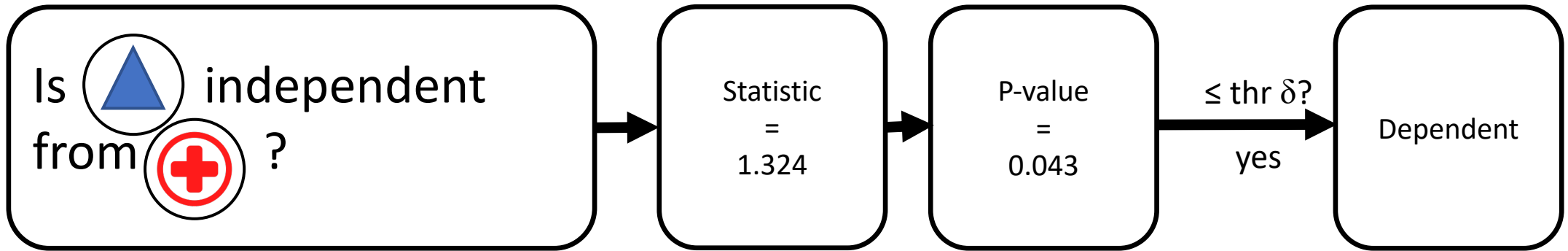


- (or) **Markov Boundary of T** $MB(T)$

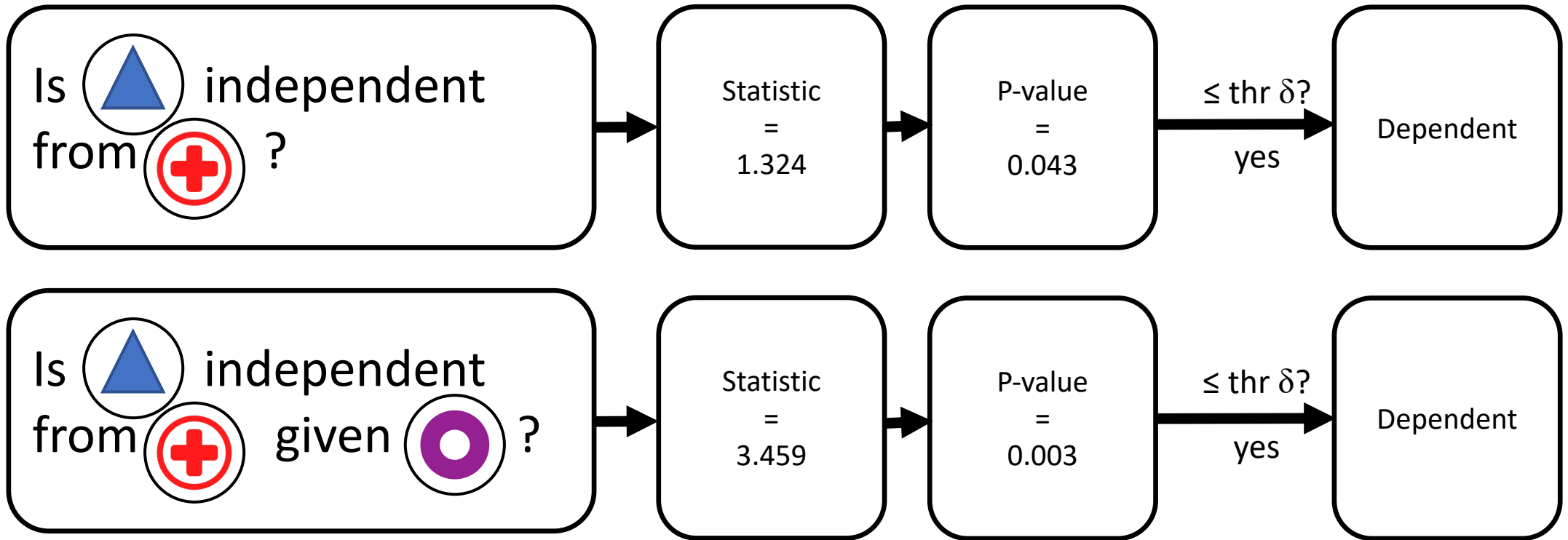


Several approaches are proposed in the literature
[Pena et al. '07], [Aliferis et al. '10], [Aliferis et al. '03]

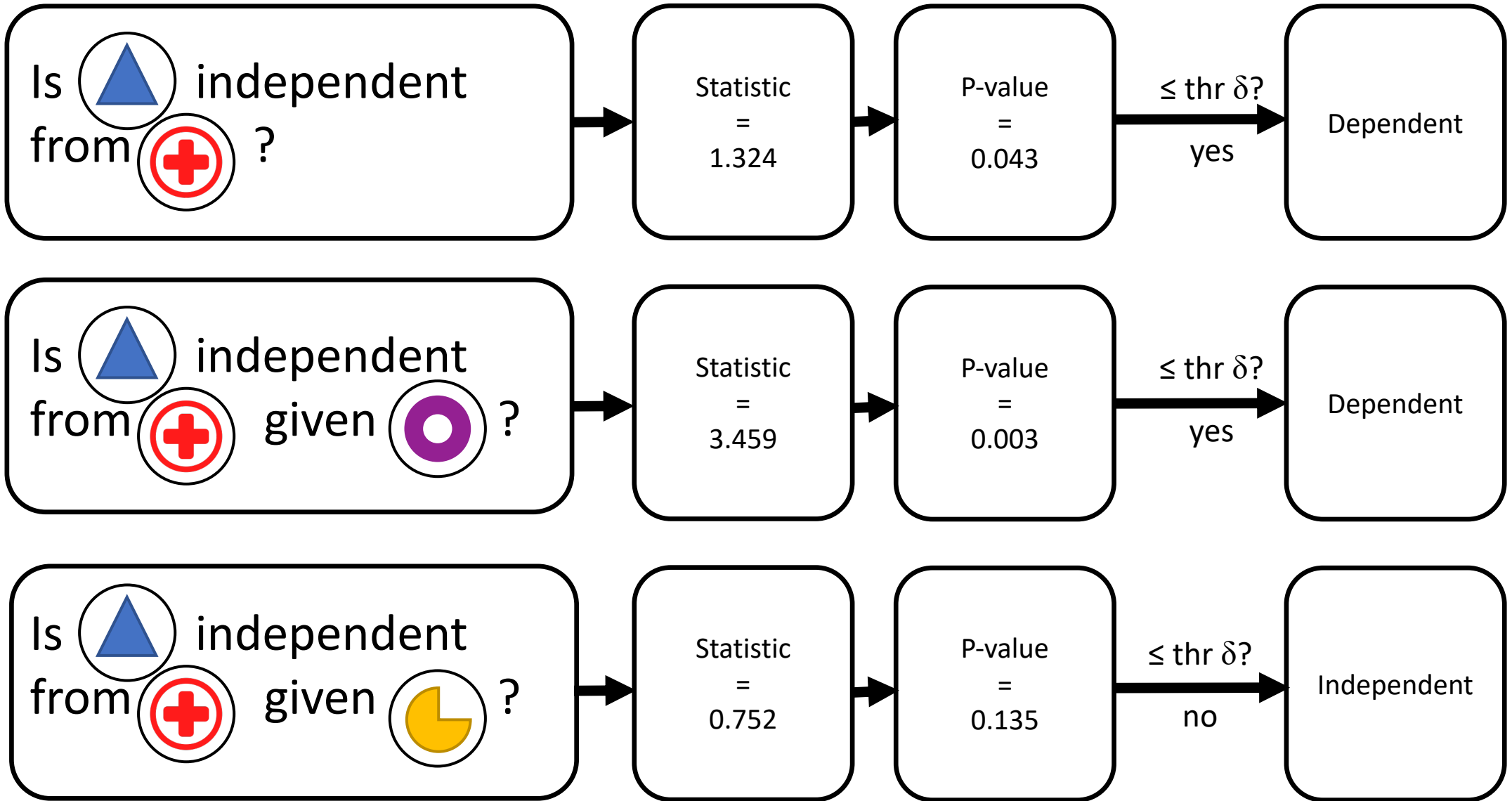
Statistical Testing and Guarantees



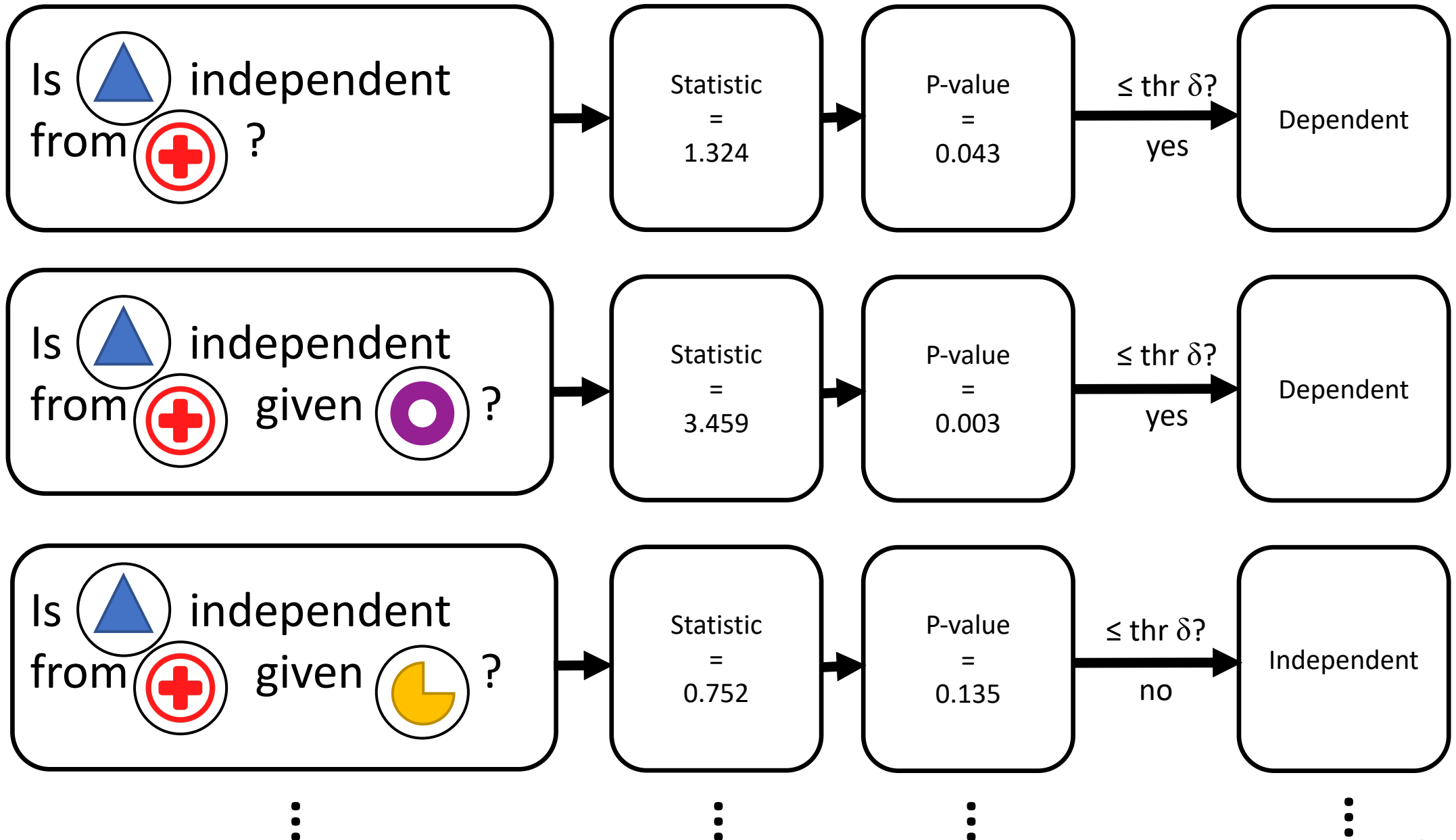
Statistical Testing and Guarantees



Statistical Testing and Guarantees



Statistical Testing and Guarantees



When testing N hypotheses

Number of false positives $\sim \text{Bin}(N, \delta)$

$$P(\text{At least one false positive}) = 1 - (1 - \delta)^N \geq \delta$$

In our problem $N = |\mathbf{V}|(|\mathbf{V}| + 1)2^{|\mathbf{V}|-2}$ is the number of possible conditional independence tests.

Approaches with guarantees typically focus on bounding

False Discovery Rate

$$FDR = E \left[\frac{\text{Number false discoveries}}{\text{Total discoveries}} \right]$$

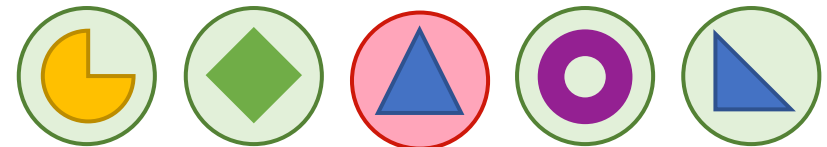
Family-Wise Error Rate

$$FWER = P(\text{Return at least one false positive})$$

Correct solution



Typical solution (controlling the *FDR*)



Typical solution (controlling the *FWER*)



Approaches with guarantees typically focus on bounding

False Discovery Rate

$$FDR = E \left[\frac{\text{Number false discoveries}}{\text{Total discoveries}} \right]$$

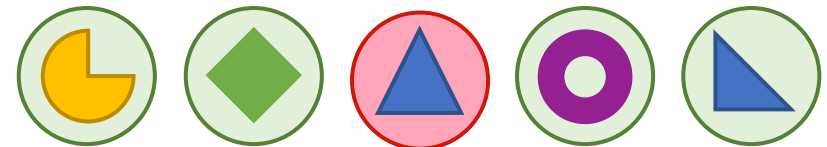
Family-Wise Error Rate

$$FWER = P(\text{Return at least one false positive})$$

Correct solution



Typical solution (controlling the *FDR*)



Typical solution (controlling the *FWER*)



Our focus

Assume perfect detection of dependencies and independencies

[Pena et al. '07], [Aliferis et al. '10], [Aliferis et al. '03]

- Unfeasible and untestable assumptions

Bound the **false discovery rate**

[Tsamardinos and Brown '08]

- May still return false positives

- Proved that **SoA algorithms cannot control the *FWER*** by correcting for multiple hypothesis testing
- Developed **RAveL-PC and RAveL-MB**: the first algorithms for local causal discovery with **guarantees on the *FWER***
- Implemented bounds on *FWER* exploiting **classical corrections** and **data-dependent** bounds based on **Rademacher averages**
- **Tested** RAveL-PC and RAveL-MB both on **synthetic** and **real-world** datasets

Th. [informal] State of the art algorithms (GetPC, PCMB and IAMB) control the *FWER* if they correct for multiple hypothesis testing and some strong assumptions (infinite power) are met.

We showed some examples in which **removing such assumptions** may lead to returning **false positives in output**

Our algorithms:

- Formulate the local discovery task using **only independence tests** (and not dependence tests)
- **Apply suitable corrections** (Rademacher, Bonferroni) to control for the *FWER*

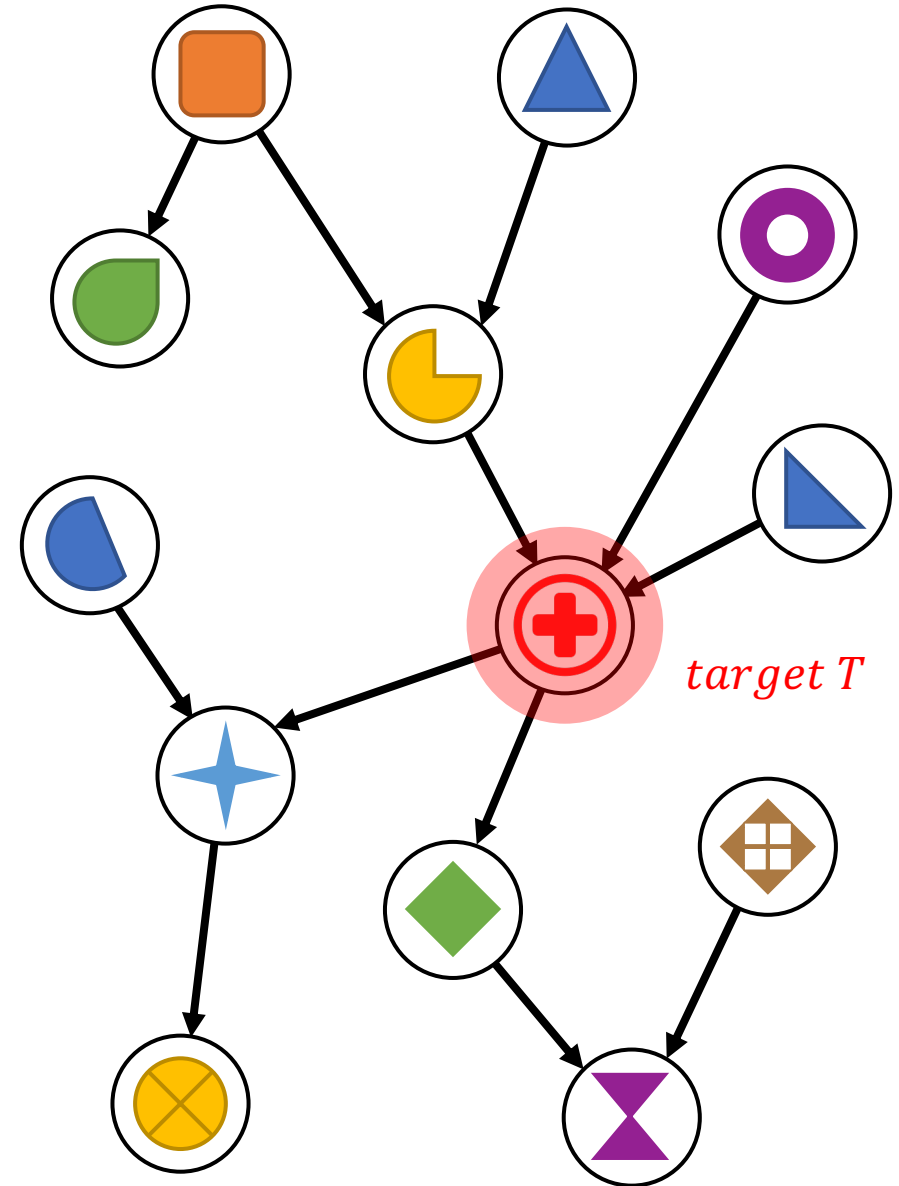
Our algorithms:

- Formulate the local discovery task using **only independence tests** (and not dependence tests)
- **Apply suitable corrections** (Rademacher, Bonferroni) to control for the *FWER*

Th. [informal] RAveL-PC and RAveL-MB effectively control the *FWER* below a given threshold δ .

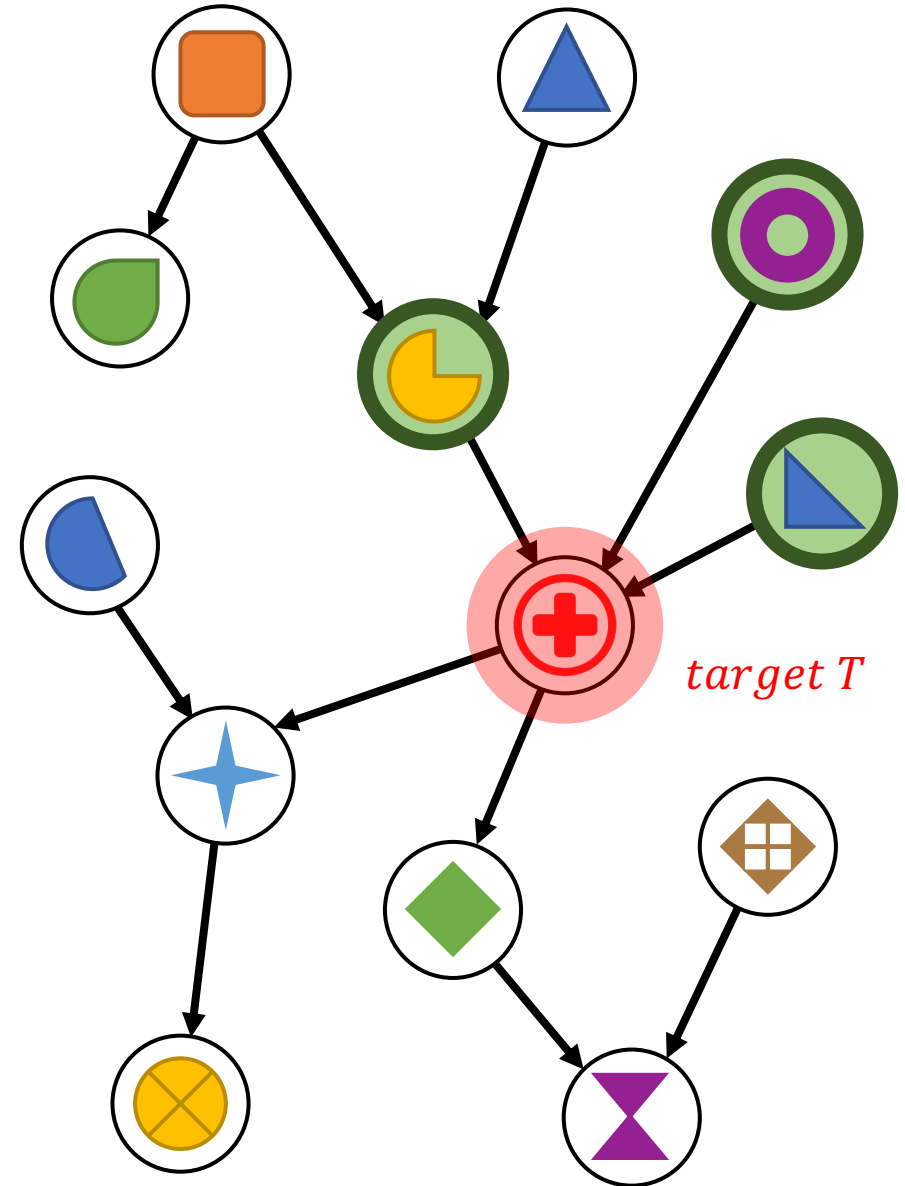
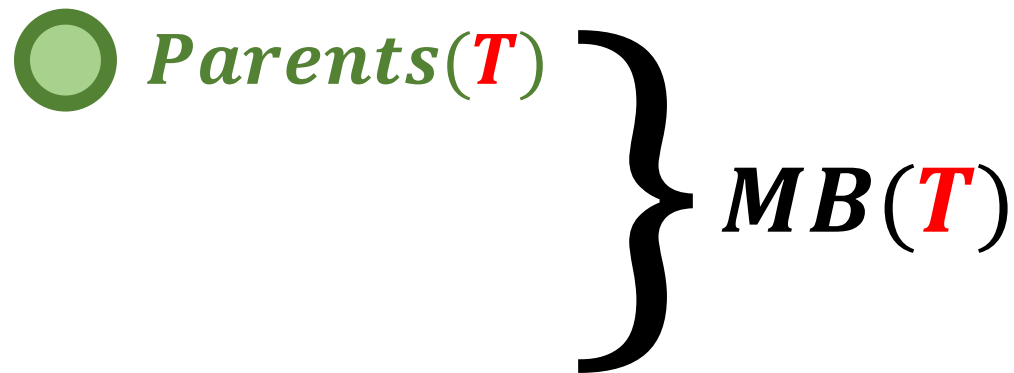
RAveL-MB at a Glance

Task: Discover a subset of $MB(T)$ without returning any false positive



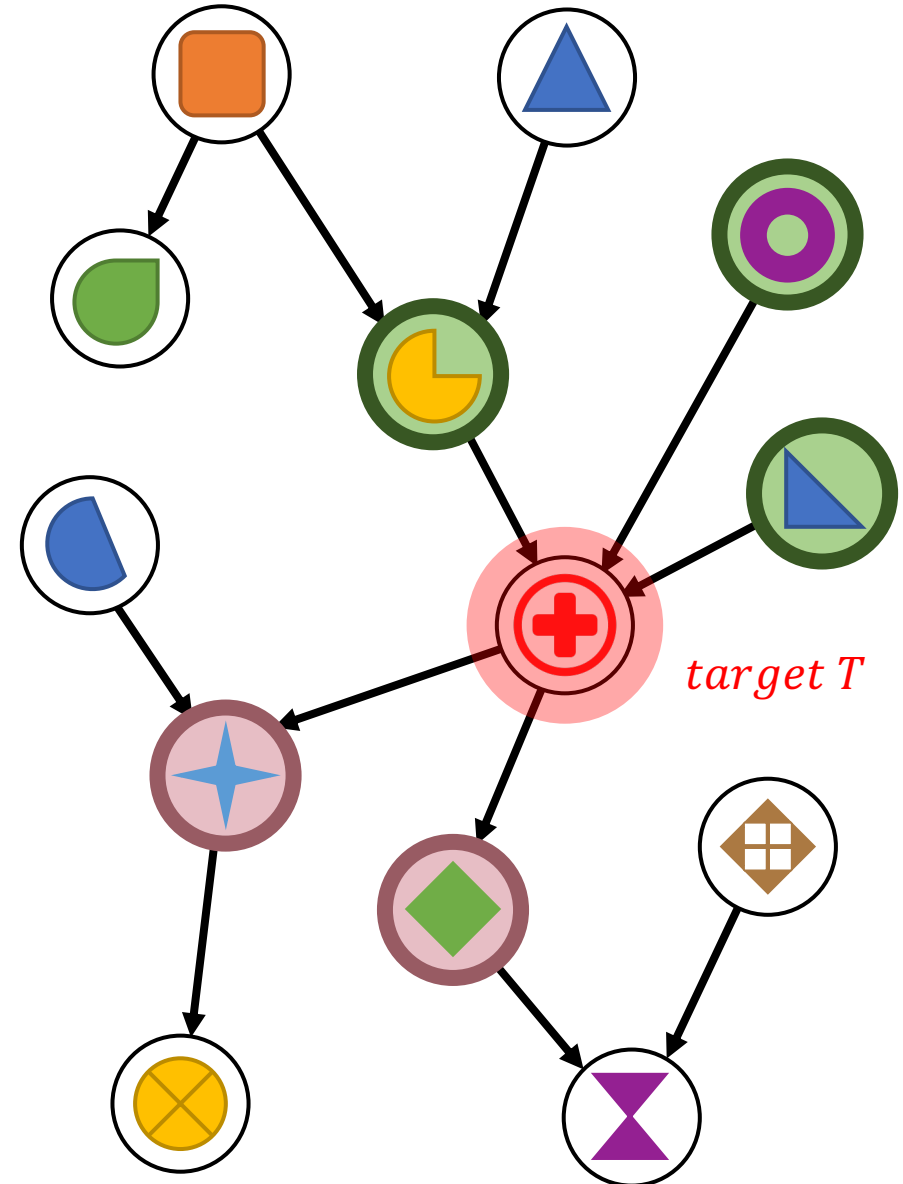
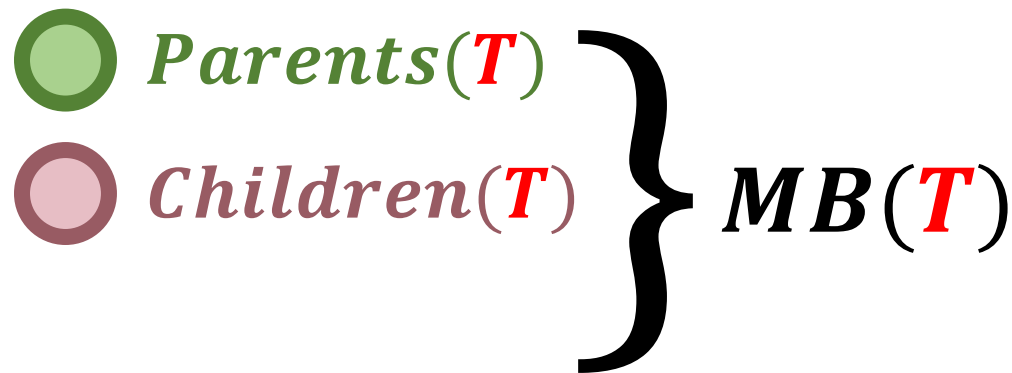
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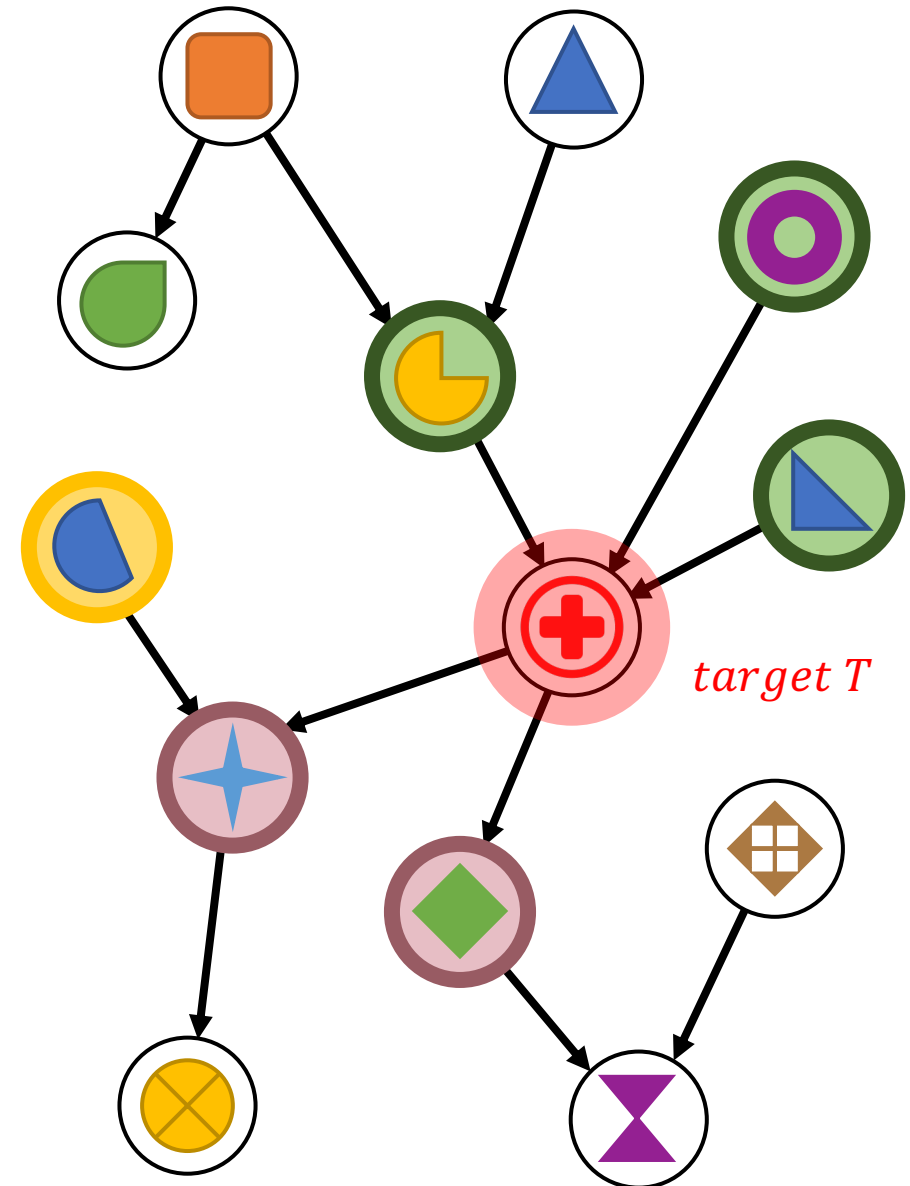
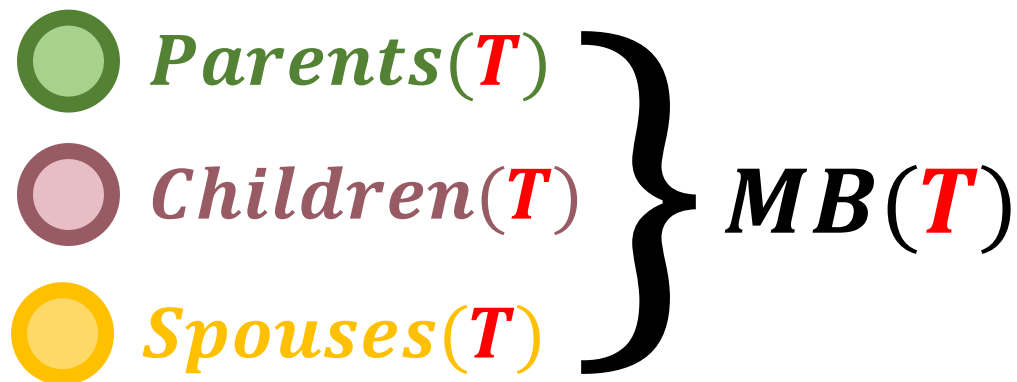
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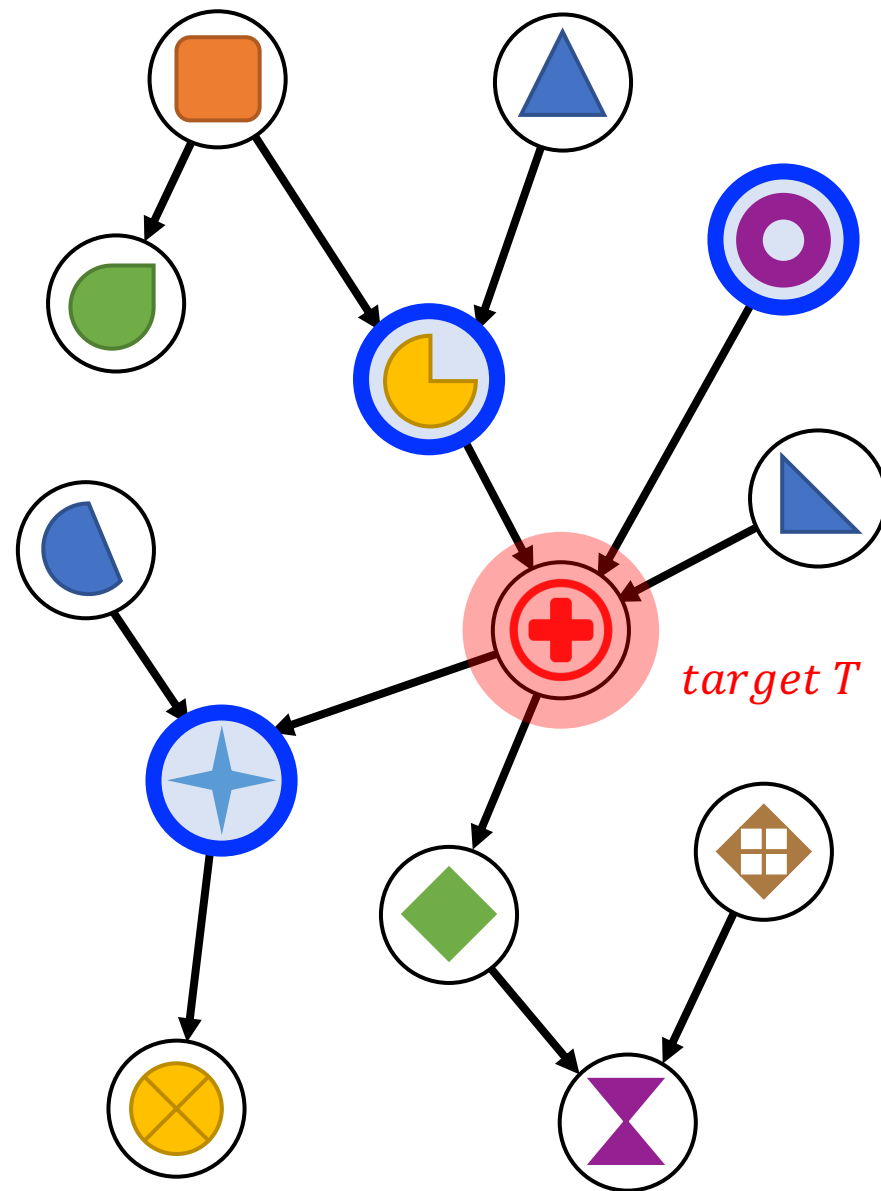
RAveL-MB at a Glance

Task: Discover a subset of $MB(T)$ without returning any false positive



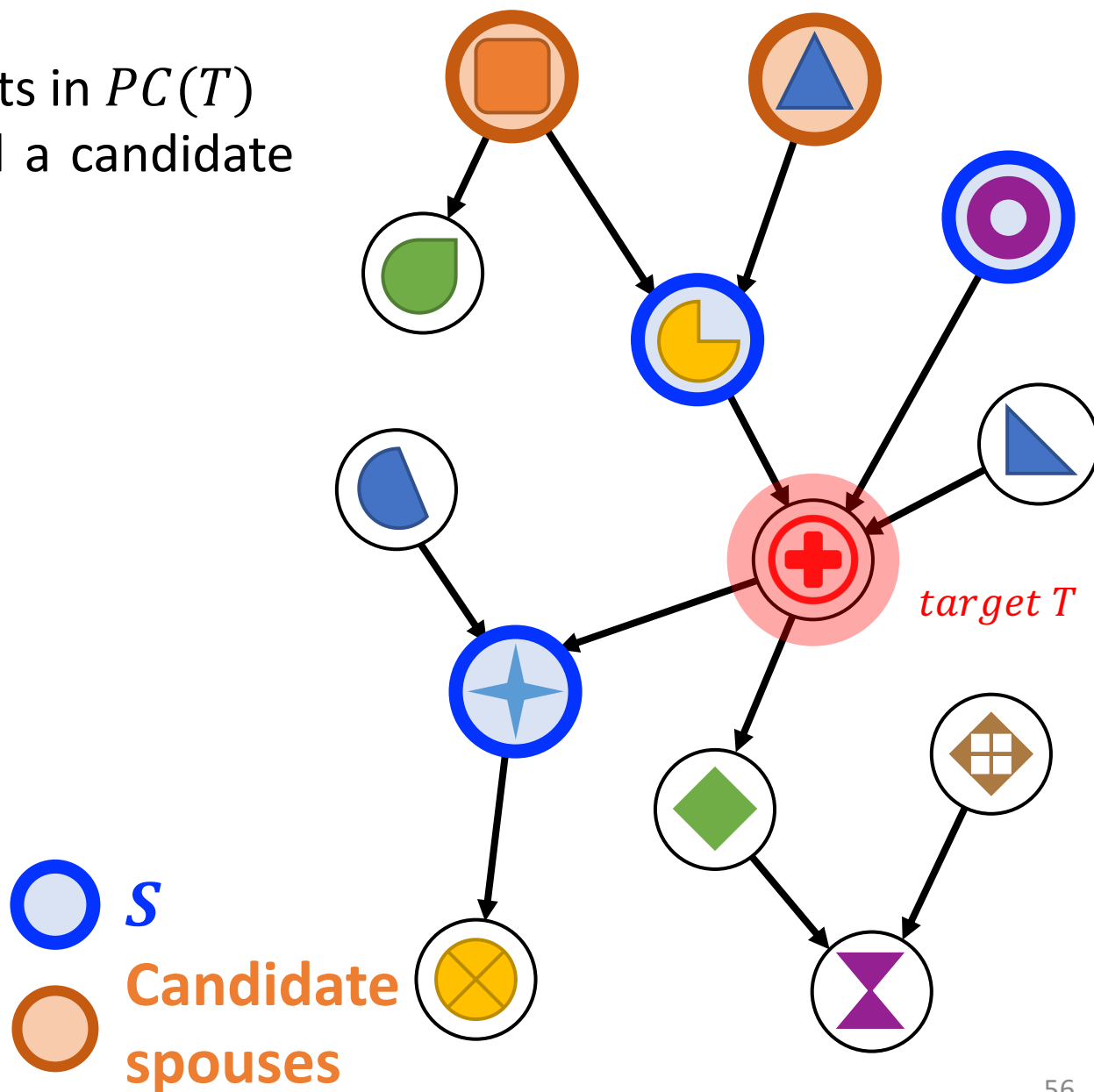
RAveL-MB at a Glance

1- Discover a subset S of elements in $PC(T)$



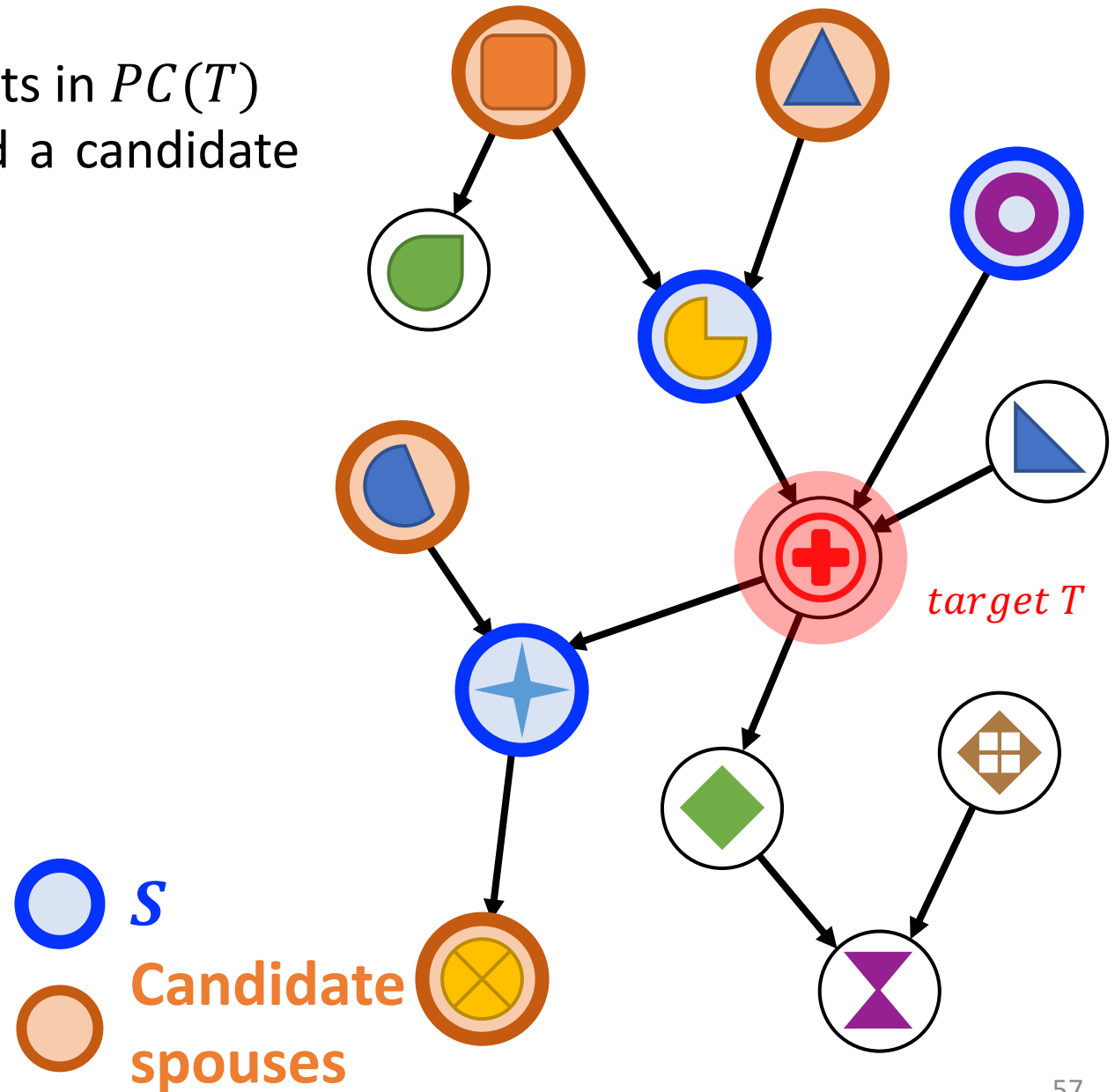
RAveL-MB at a Glance

- 1- Discover a subset \mathcal{S} of elements in $PC(T)$
- 2- For each element X in \mathcal{S} find a candidate set of spouses in $PC(X)$



RAveL-MB at a Glance

- 1- Discover a subset \mathcal{S} of elements in $PC(T)$
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RAveL-MB at a Glance

- 1- Discover a subset \mathcal{S} of elements in $PC(T)$
- 2- For each element X in \mathcal{S} find a candidate set of spouses in $PC(X)$
- 3- For each candidate spouse Y with children X , test the spouse condition

Previous approaches:

Dependence test

$$Y \perp\!\!\!\perp T \mid \mathbf{Z} \text{ and } Y \not\perp\!\!\!\perp T \mid \mathbf{Z} \cup \{X\}$$

for all $\mathbf{Z} \subseteq \mathbf{V} \setminus \{X, Y, T\}$

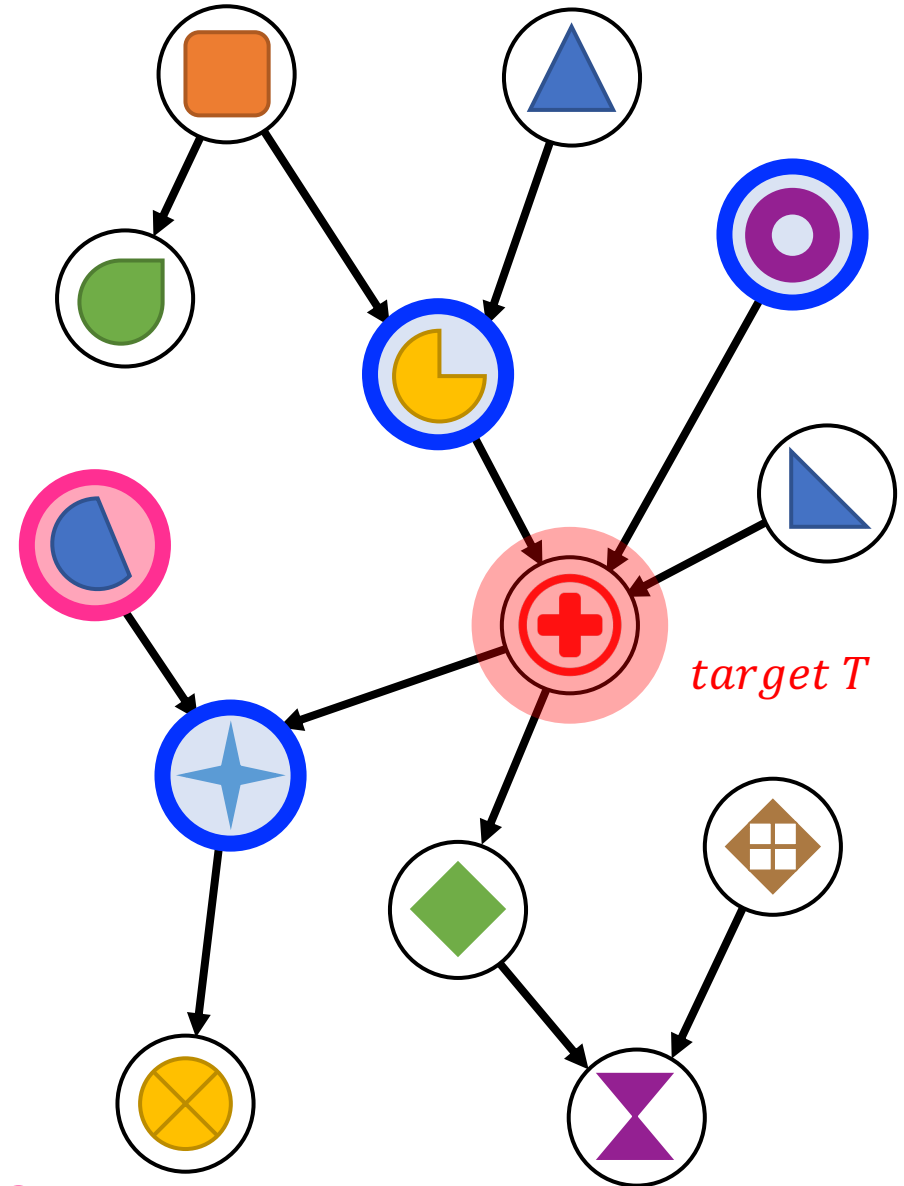
Independence test

$$Y \not\perp\!\!\!\perp T \mid \mathbf{Z} \cup \{X\}$$

Our formulation (equivalent):

$$Y \not\perp\!\!\!\perp T \mid \mathbf{V} \setminus \{Y\}$$

● \mathcal{S}
● Actual spouses



RAveL-MB at a Glance

- 1- Discover a subset \mathcal{S} of elements in $PC(T)$
- 2- For each element X in \mathcal{S} find a candidate set of spouses in $PC(X)$
- 3- For each candidate spouse Y with children X , test the spouse condition

Previous approaches:

Dependence test

$$Y \perp\!\!\!\perp T \mid \mathbf{Z} \text{ and } Y \not\perp\!\!\!\perp T \mid \mathbf{Z} \cup \{X\}$$

for all $\mathbf{Z} \subseteq \mathbf{V} \setminus \{X, Y, T\}$

Independence test

$$Y \not\perp\!\!\!\perp T \mid \mathbf{Z} \cup \{X\}$$

Our formulation (equivalent):

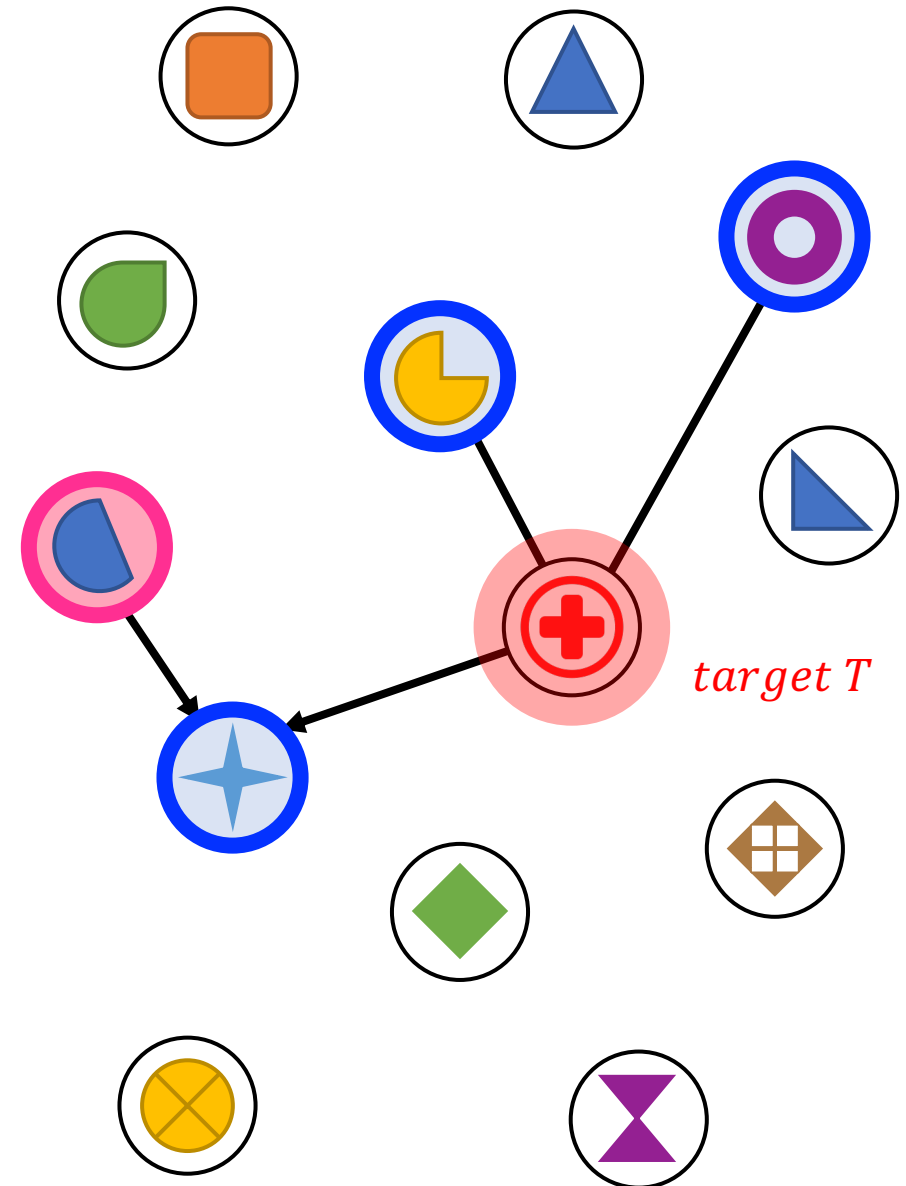
$$Y \not\perp\!\!\!\perp T \mid \mathbf{V} \setminus \{Y\}$$



\mathcal{S}



Actual spouses



- **Bonferroni correction** on the threshold δ

Classical correction for multiple hypotheses testing uses a modified threshold δ/N on each test

May be too strict if N is big (and in our case N is exponential on $|V|$)

- **Rademacher averages** to bound each test statistic

Provide data-dependent bounds

Rademacher Averages - Idea

Given a family of functions \mathcal{F} and a dataset S ,
Rademacher averages upper bound with high
probability the Supremum Deviation $D(\mathcal{F}, S)$

Empirical sample mean

**Rademacher
estimate**

$$D(\mathcal{F}, S) = \sup_{f \in \mathcal{F}} \left| \hat{E}_S[f] - E[f] \right| \leq 2\tilde{R} + O\left(\frac{1}{\sqrt{m}}\right)$$

Expectation

**Dataset
size**

Given a family of functions \mathcal{F} and a dataset S ,
Rademacher averages upper bound with high
probability the Supremum Deviation $D(\mathcal{F}, S)$

$$D(\mathcal{F}, S) = \sup_{f \in \mathcal{F}} \left| \hat{E}_S[f] - E[f] \right| \leq 2\tilde{R} + O\left(\frac{1}{\sqrt{m}}\right)$$

Empirical sample mean Rademacher estimate

Expectation Dataset size

They can lower bound **simultaneously** each
independence test statistic for providing guarantees
on the *FWER*

Given two normalized vectors of observations \mathbf{x}, \mathbf{y}

Pearson's r coefficient $r_{\mathbf{x}, \mathbf{y}} = \frac{\sum_{i=0}^{m-1} x_i y_i}{(m - 1)}$

Given two normalized vectors of observations \mathbf{x}, \mathbf{y}

Pearson's r coefficient $r_{\mathbf{x}, \mathbf{y}} = \frac{\sum_{i=0}^{m-1} x_i y_i}{(m-1)}$

By defining

$$r_{\mathbf{x}, \mathbf{y}}(s_i) = \frac{m}{(m-1)} x_i y_i$$

We get

$$r_{\mathbf{x}, \mathbf{y}} = \widehat{\mathbb{E}}_S[r_{\mathbf{x}, \mathbf{y}}(s_i)]$$

Given two normalized vectors of observations \mathbf{x}, \mathbf{y}

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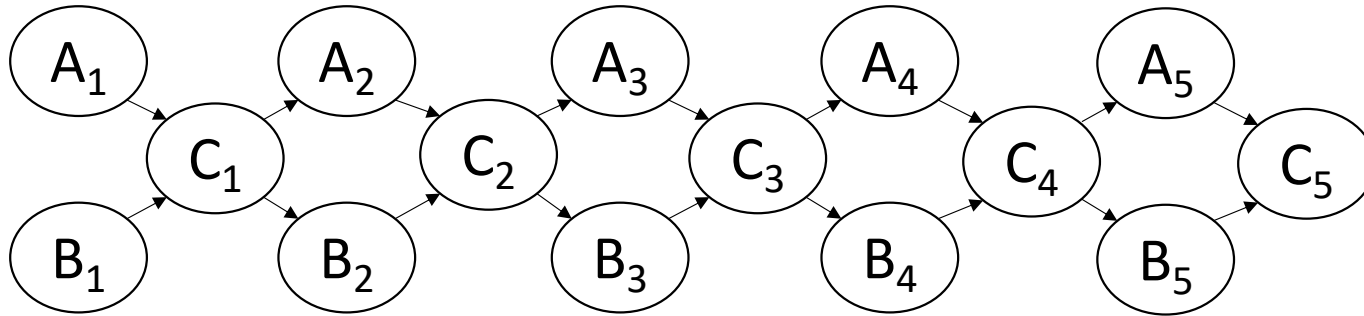
By defining

$$\mathcal{F} \ni r_{x,y}(s_i) = \frac{m}{(m-1)} x_i y_i$$

We get

$$r_{x,y} = \widehat{\mathbb{E}}_S[r_{x,y}(s_i)]$$

The same approach works with other test statistics



$$A_1 \sim N(0, 1)$$

$$B_1 \sim N(0, 1)$$

$$C_1 \sim -3 + 2A_1 + 3B_1 + N(0, 1)$$

$$\forall i \in [2, 3, 4, 5]$$

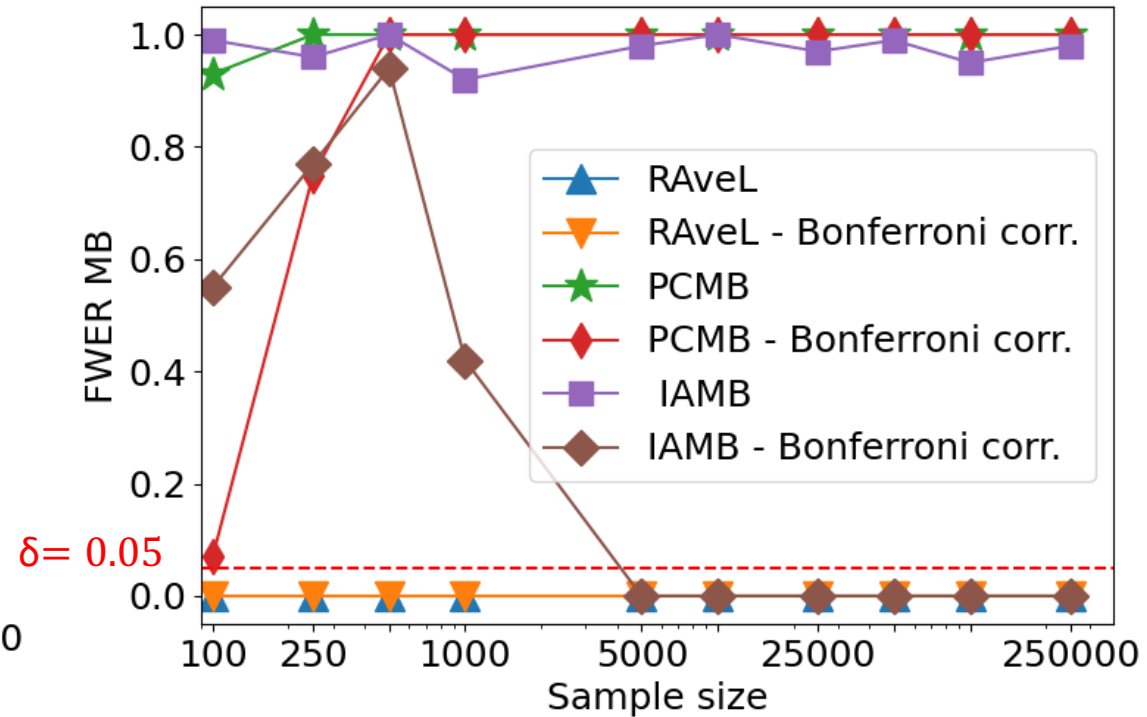
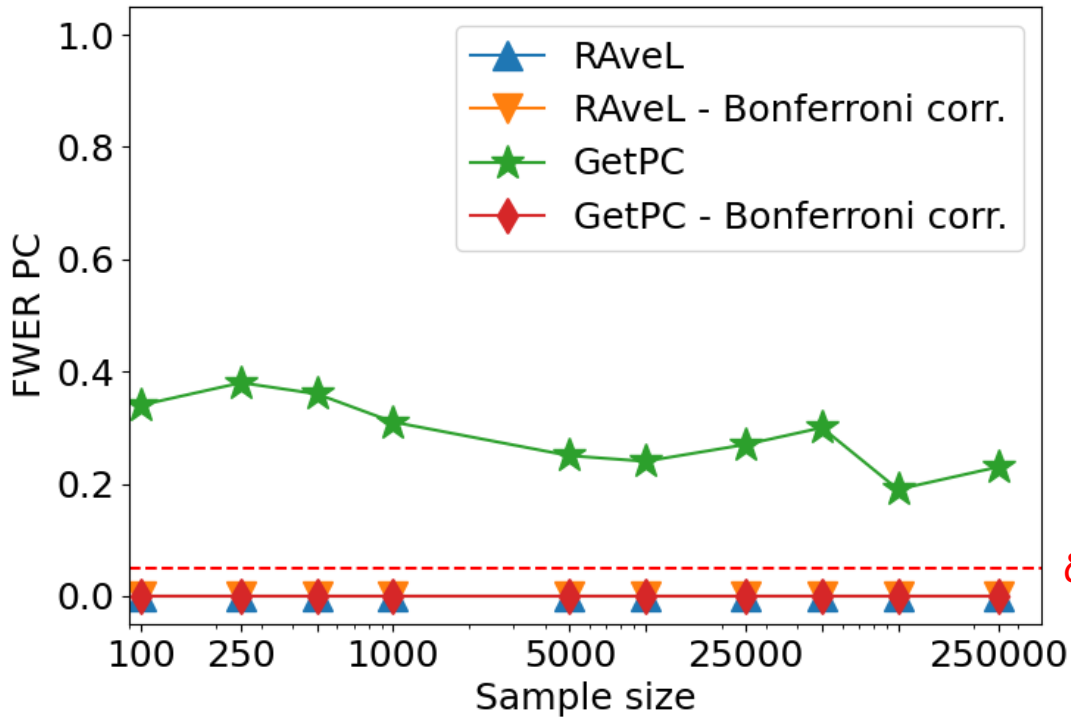
$$A_i \sim +3 + 3C_{i-1} + N(0, 1)$$

$$B_i \sim -1 - 2C_{i-1} + N(0, 1)$$

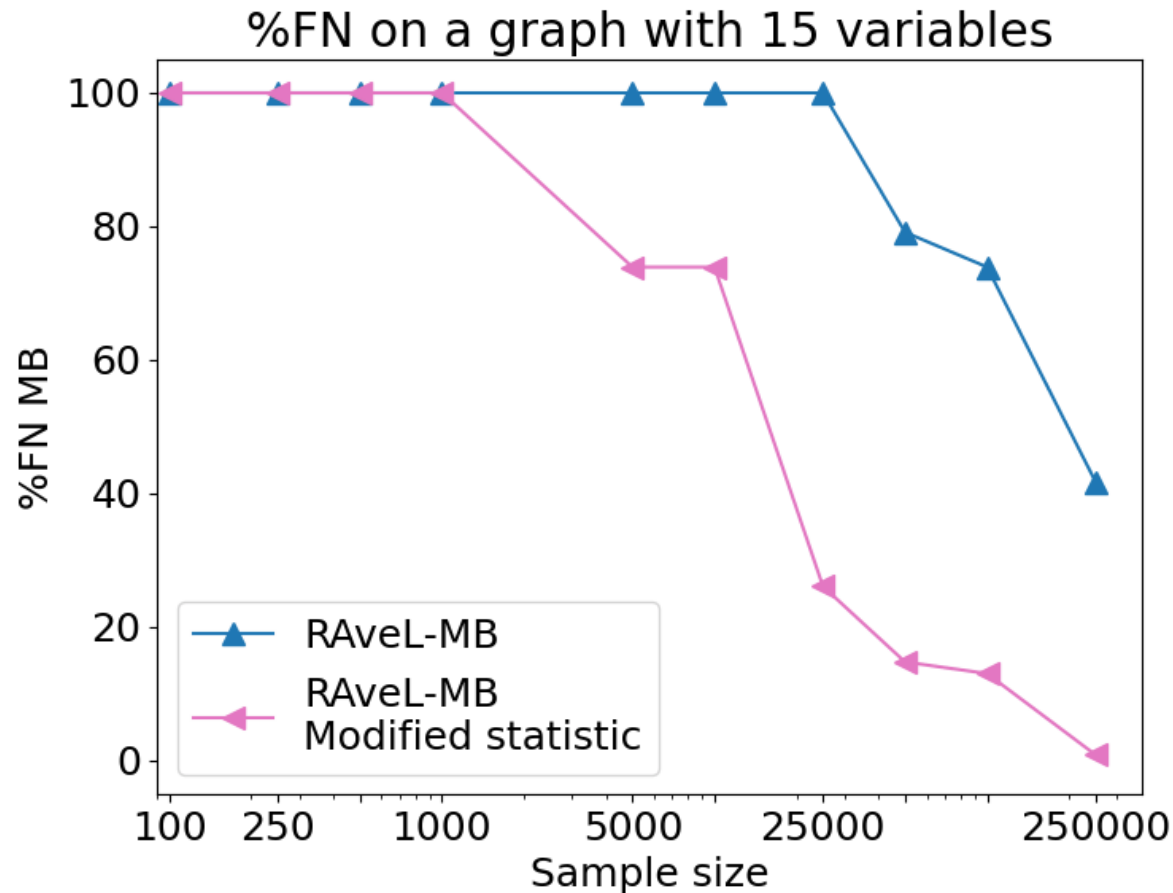
$$C_i \sim -3 + 2A_i + 3B_i + N(0, 1)$$

- On each run, test the SoA and RAveL algorithms on every variable
- Analyse results of each iteration:
 - No False Positives only if **all the outputs** (one per variable) **of the run** do not contain false positives
 - Count mean percentage of false negatives

RAveL-PC and RAveL-MB effectively control the *FWER*

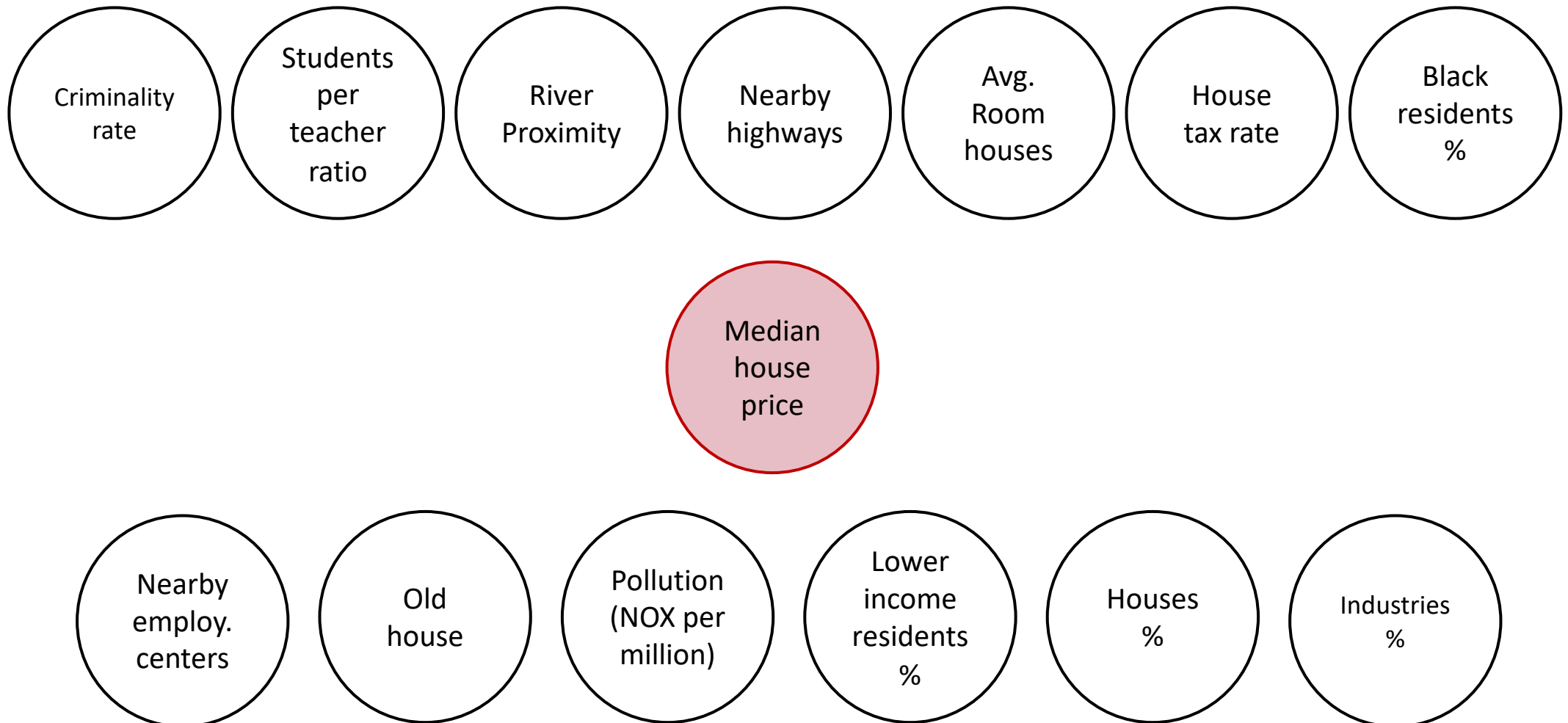


Tests with increasing number of variables

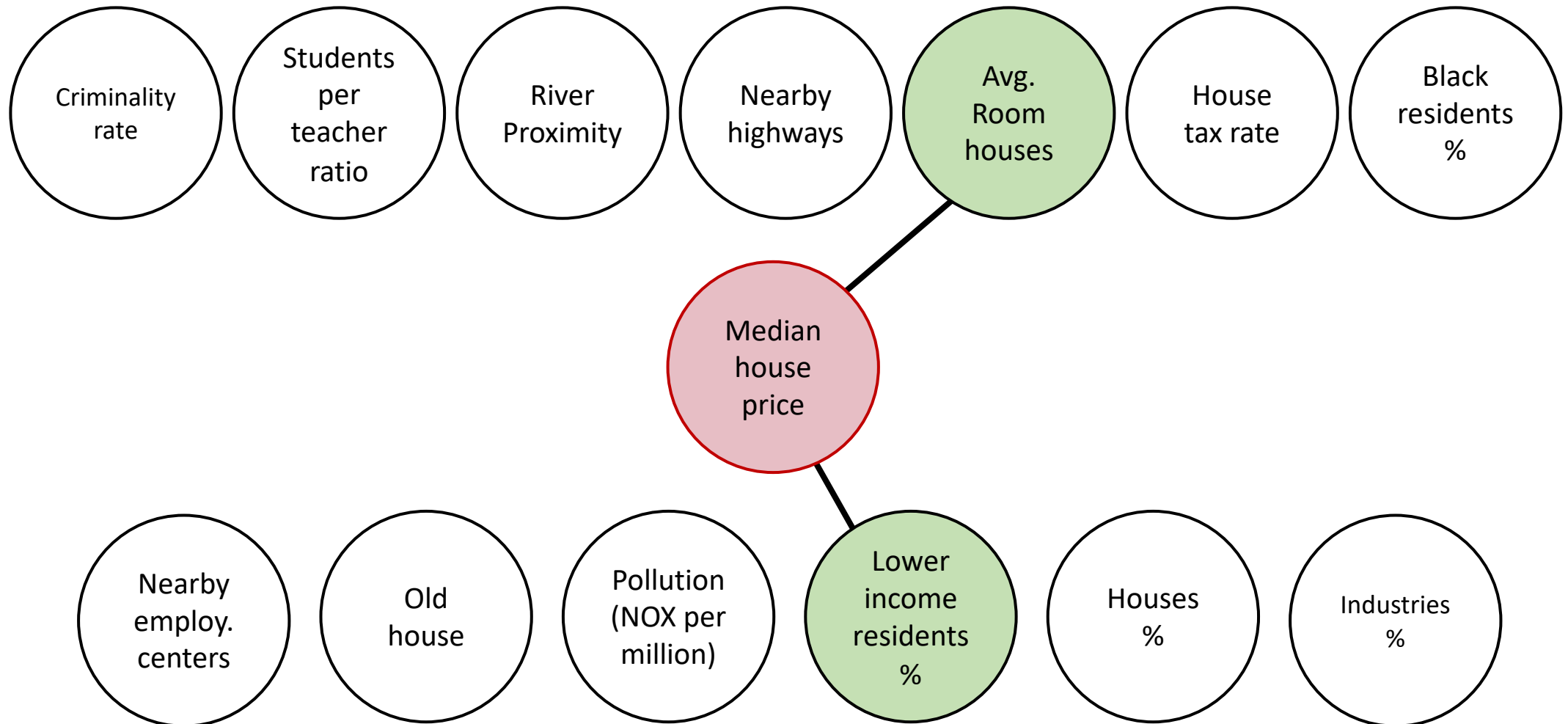


Test statistic choice heavily affects Rademacher based corrections

Tested RAveL-PC and RAveL-MB on **Boston housing** dataset with $\delta = 0,01$.



Tested RAveL-PC and RAveL-MB on **Boston housing** dataset with $\delta = 0,01$.

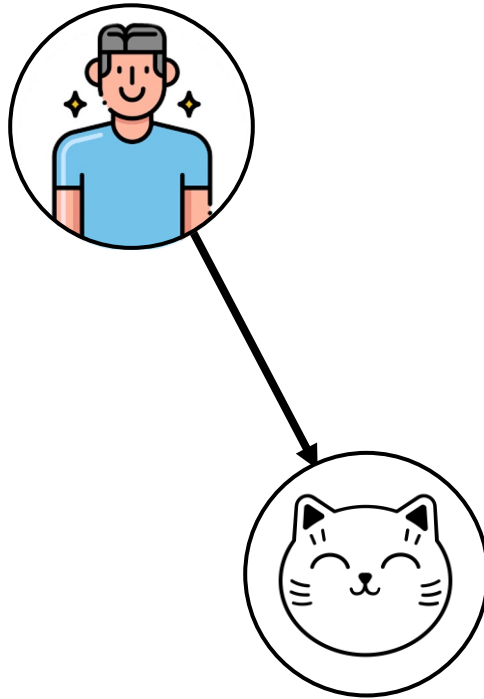


Develop algorithms to:

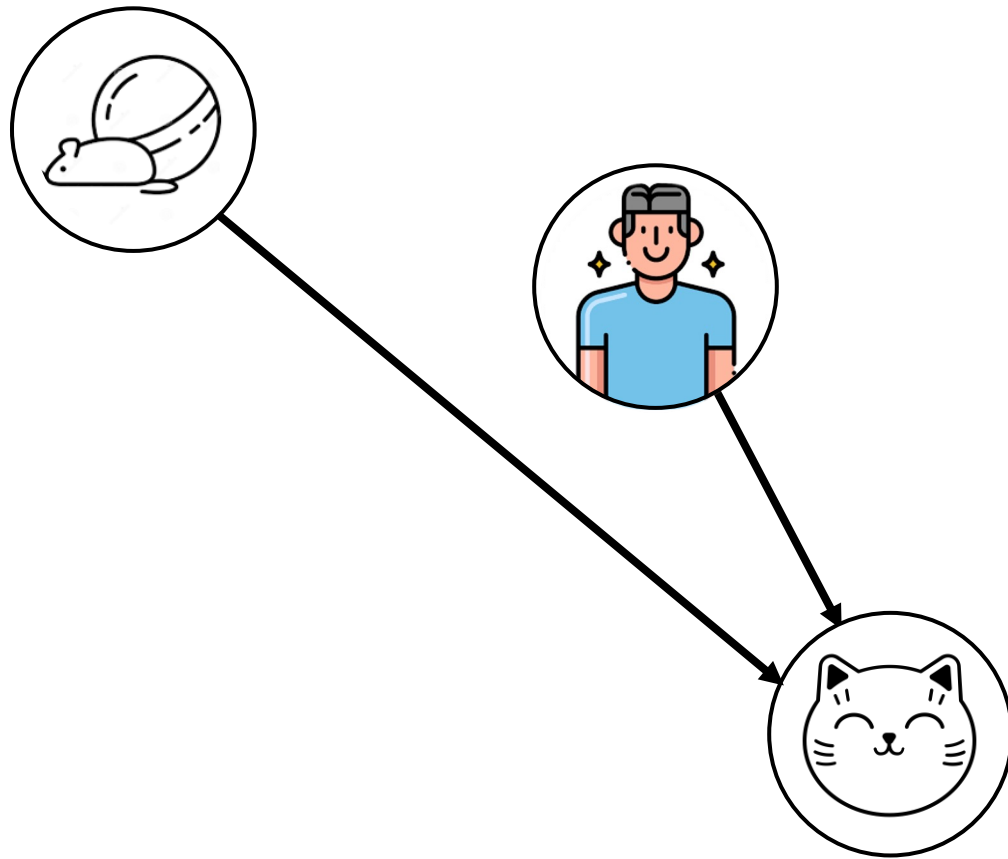
- **(Structure discovery) Discover causally related variables to a target**
- (Effect estimation) Evaluate effect of causal rules

From observational data and providing guarantees on the results

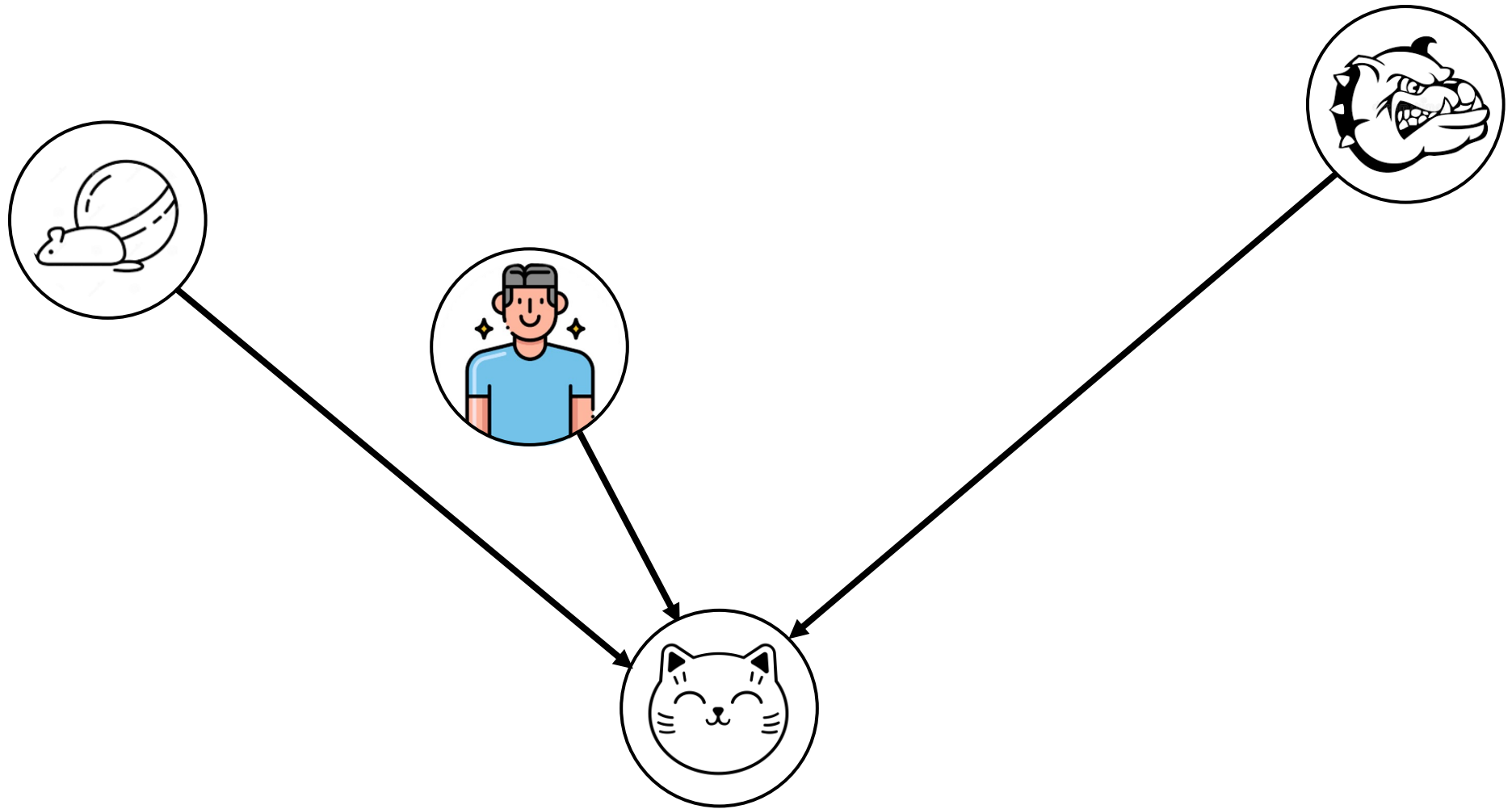
Very serious research question



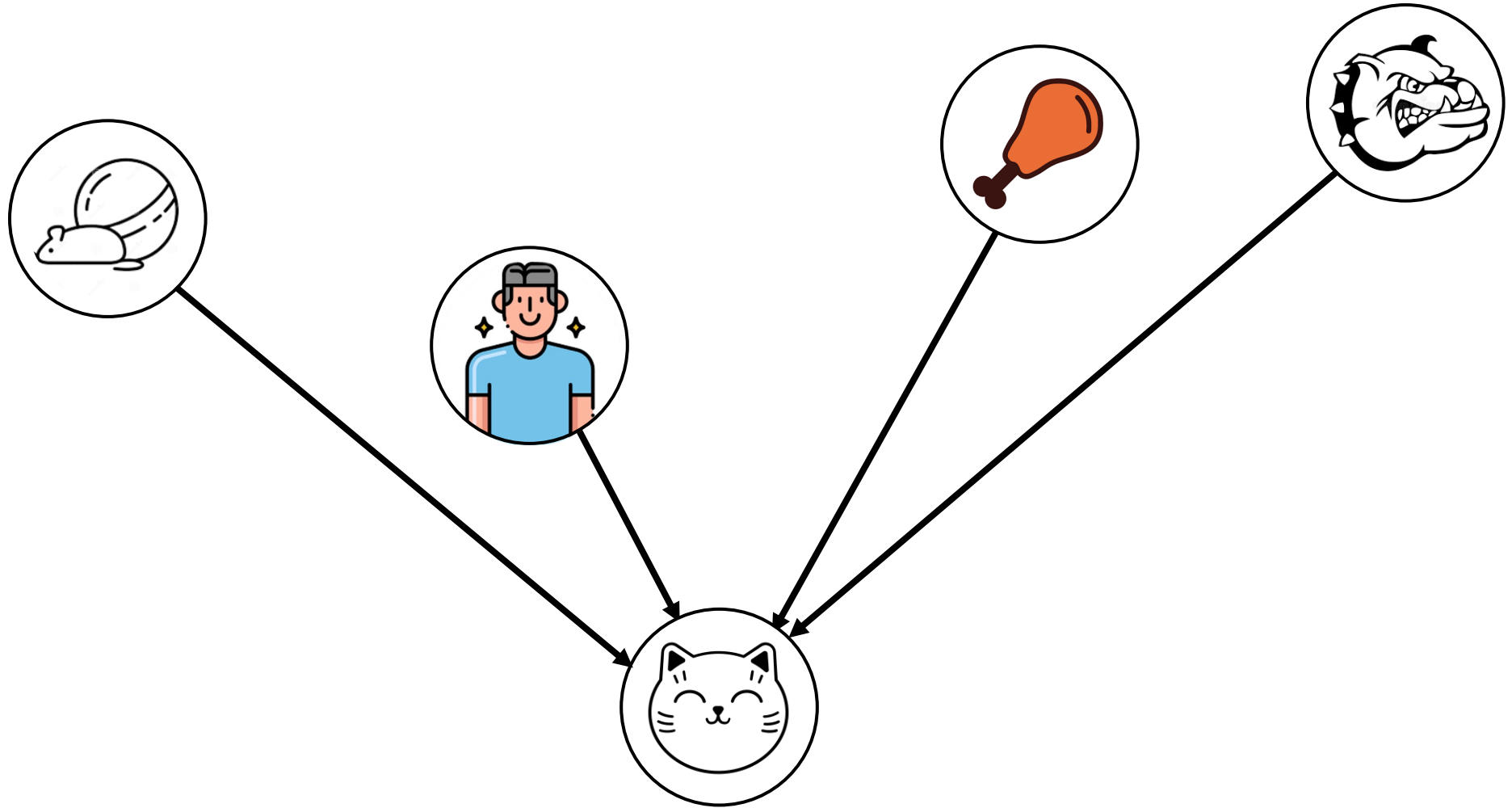
Very serious research question



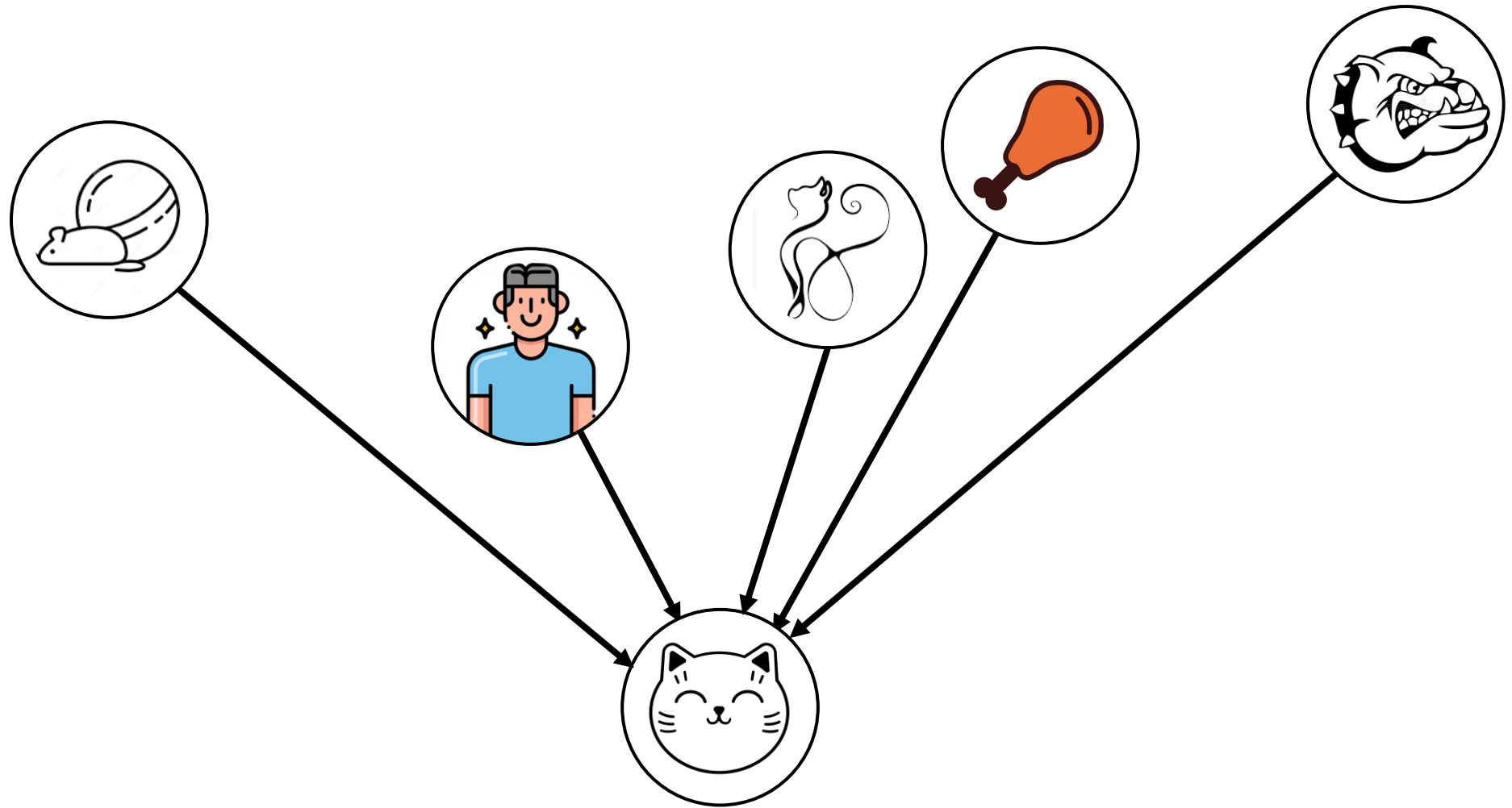
Very serious research question



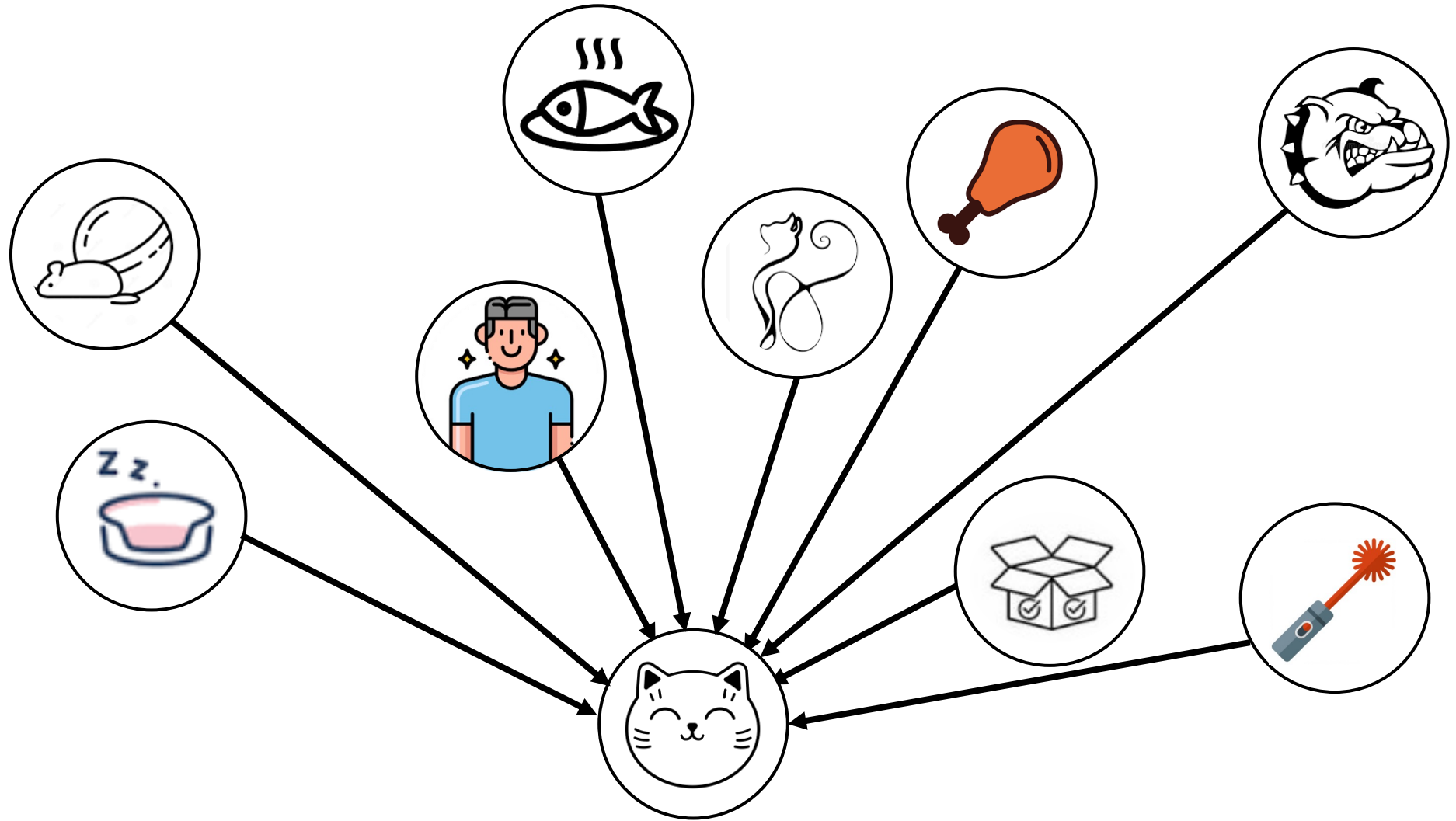
Very serious research question



Very serious research question



Very serious research question



Develop algorithms to:

- (Structure discovery) Discover causally related variables to a target
- **(Effect estimation) Evaluate effect of causal rules**

Submitted paper at ECCB 2023 and work in progress in collaboration with:

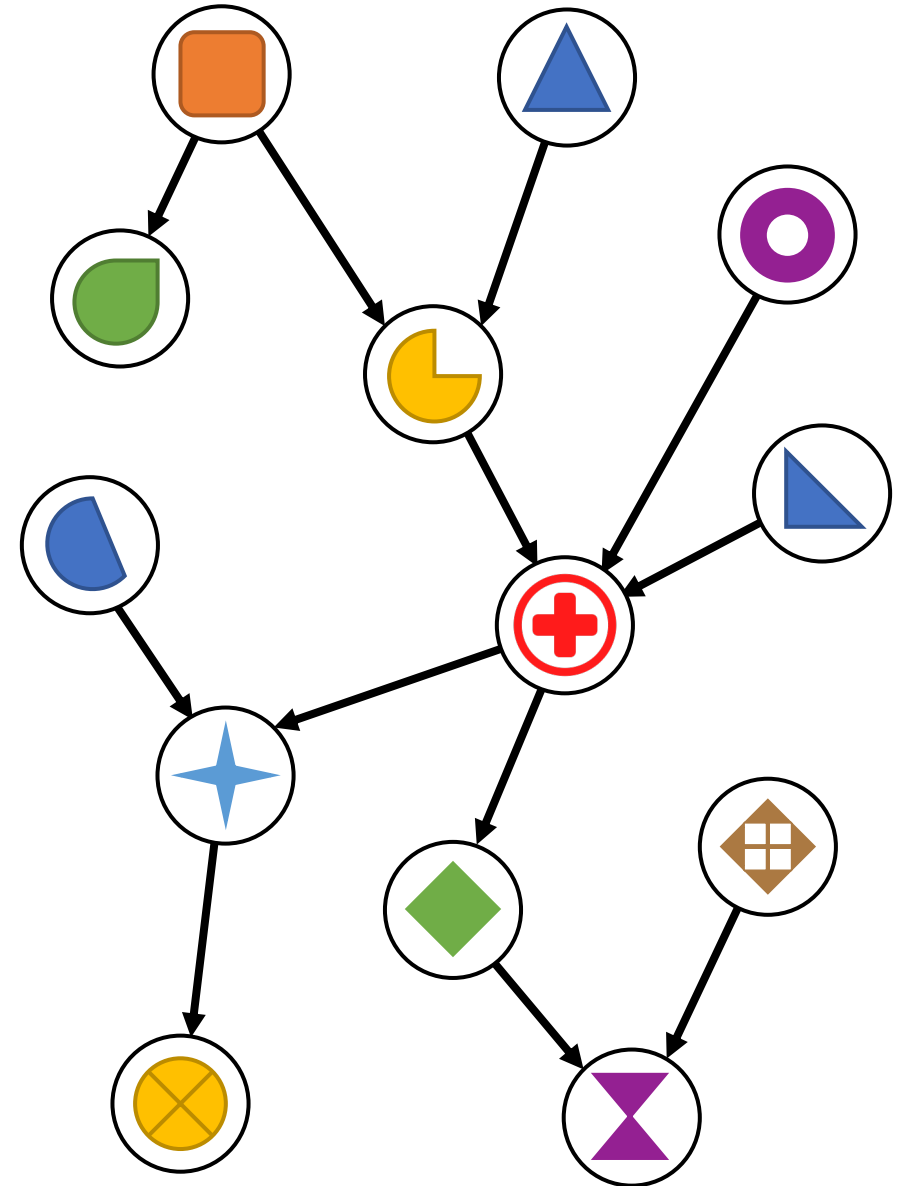
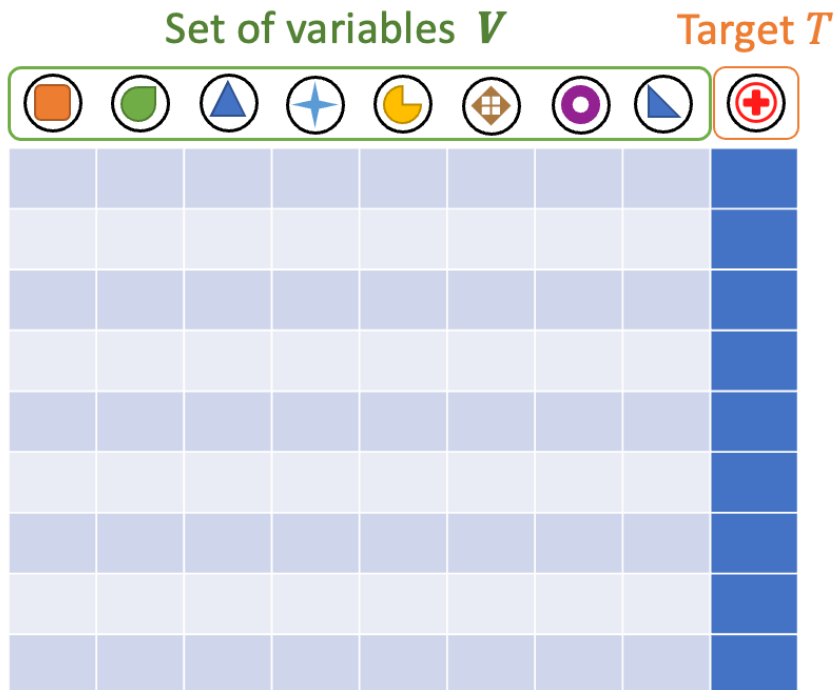
Antonio Collesei, PhD s. (IOV)

Paola Donolato, MS

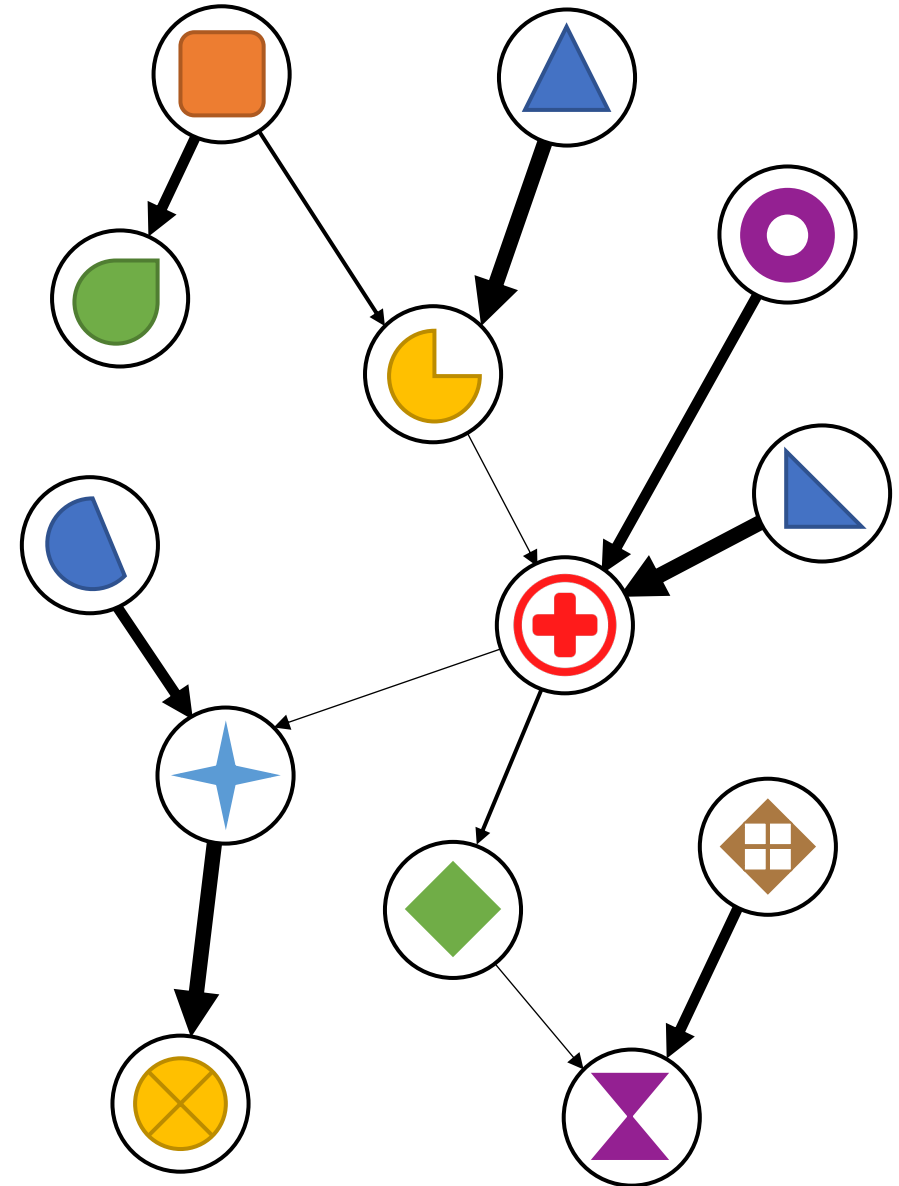
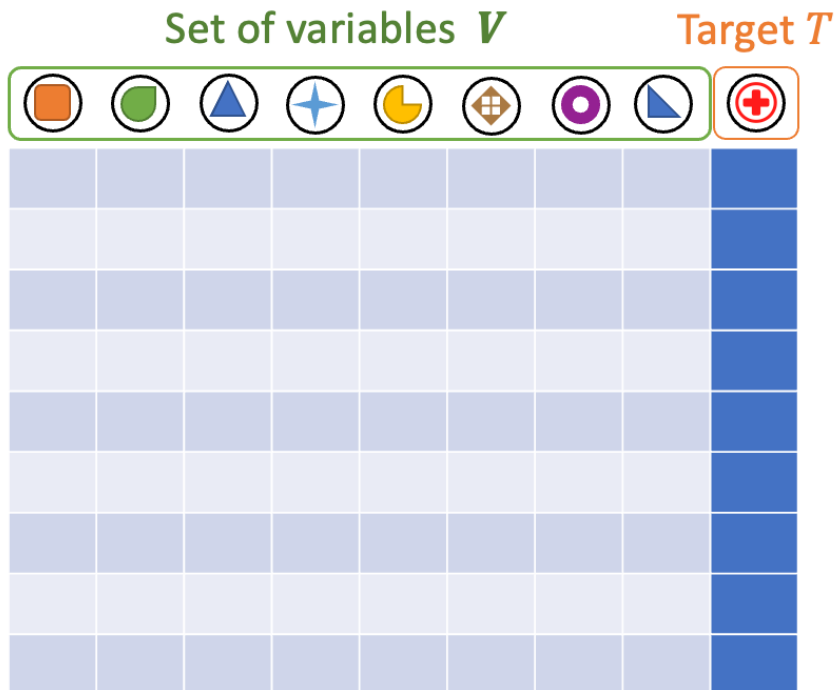
Federica Miglietta, MD and PhD s. (IOV)

Fabio Vandin, Full Professor

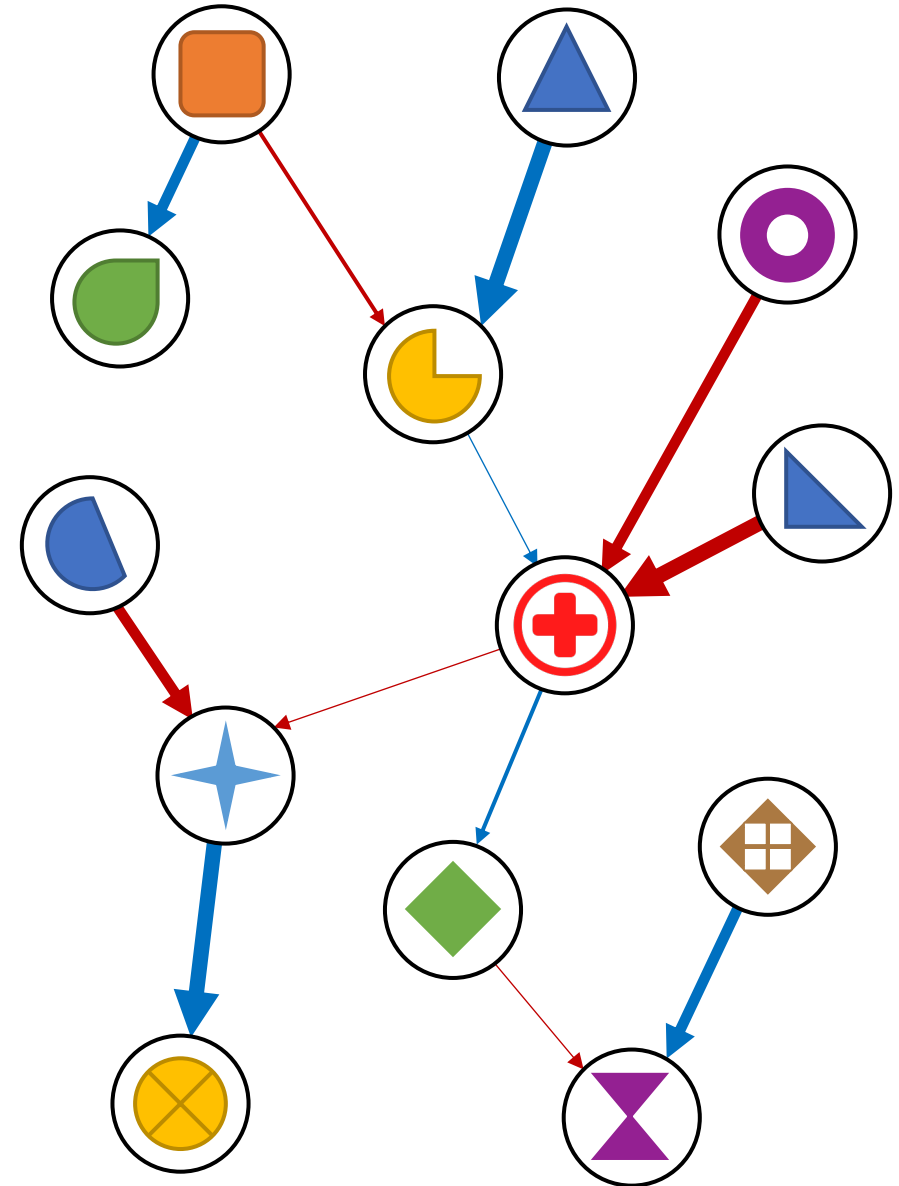
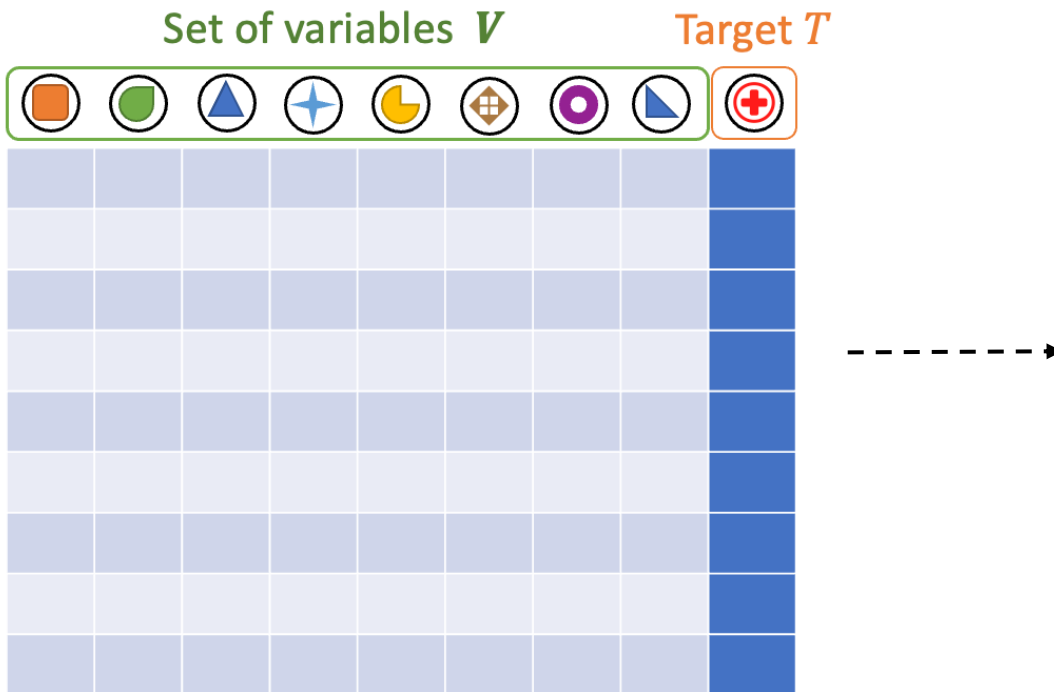
Causal Bayesian Networks represent cause-consequence relations between variables



Causal Bayesian Networks represent cause-consequence relations between variables

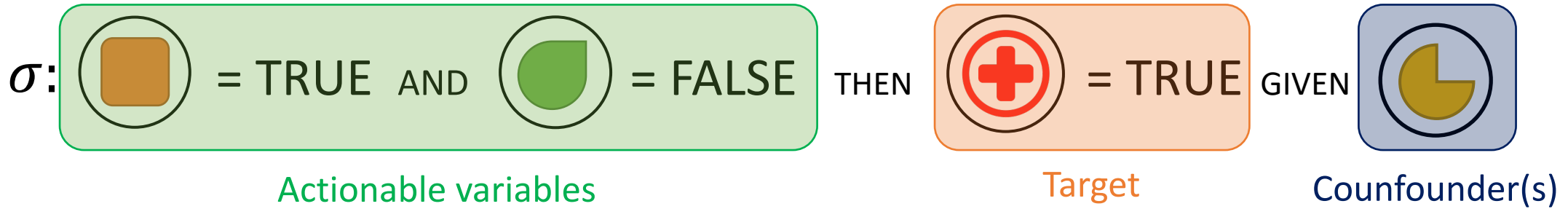


Causal Bayesian Networks represent cause-consequence relations between variables

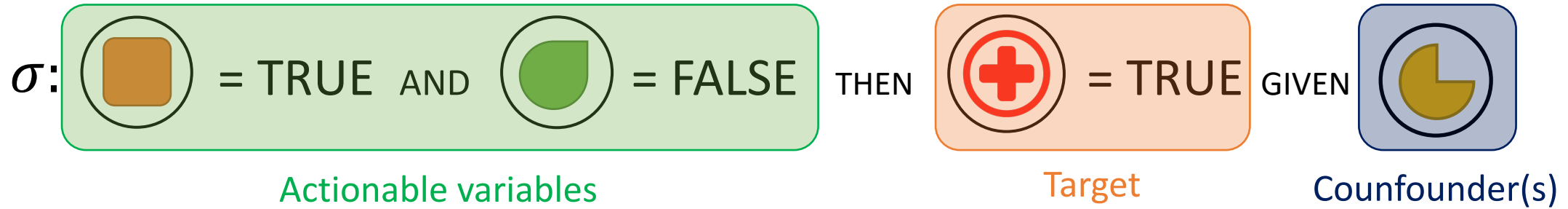


Positive impact
 Negative impact

Problem Definition

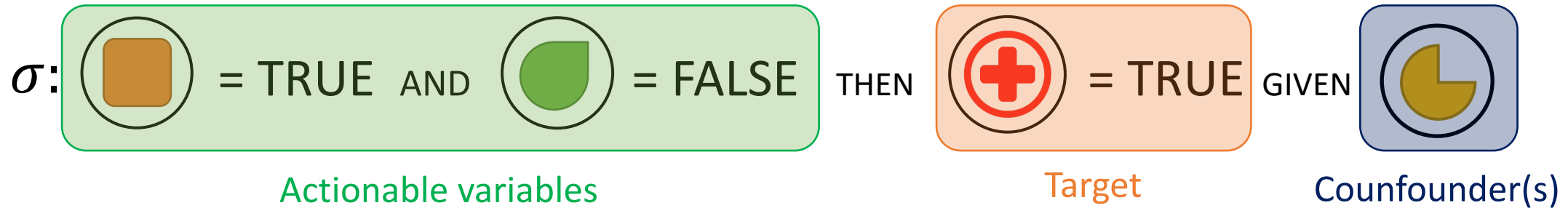


Problem Definition



$$e(\sigma) = p\left(\text{⊕} = TRUE \mid \sigma, \text{Ⓚ}\right) - p\left(\text{⊕} = TRUE \mid \bar{\sigma}, \text{Ⓚ}\right)$$

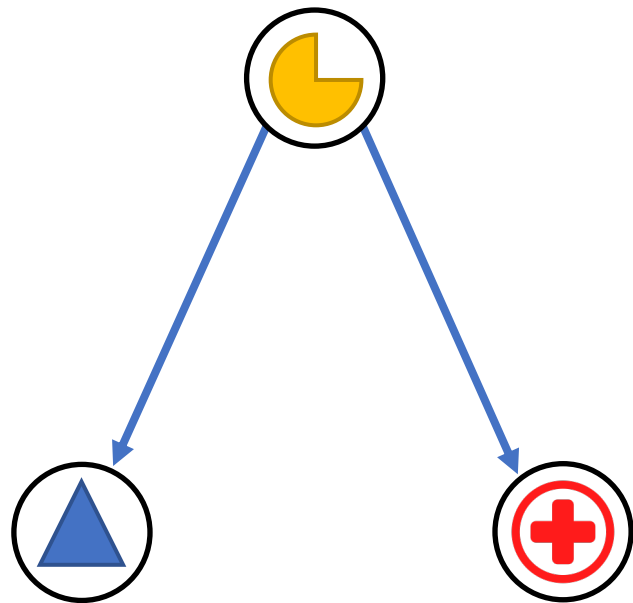
Problem Definition



$$e(\sigma) = p\left(\textcircled{+} = TRUE \mid \sigma, \textcircled{\text{L}}\right) - p\left(\textcircled{+} = TRUE \mid \bar{\sigma}, \textcircled{\text{L}}\right)$$

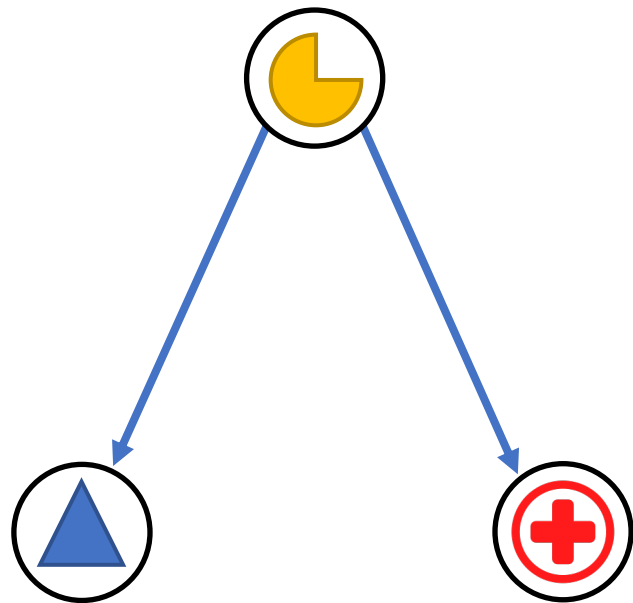
Task: Given a dataset of observations of variables V , find the top- k rules $\sigma_1^*, \dots, \sigma_k^*$ with **the highest causal effect** on T with **guarantees** on the result (e.g. no false positives)

We have to deal with confounder variables



- Positive impact
- Negative impact

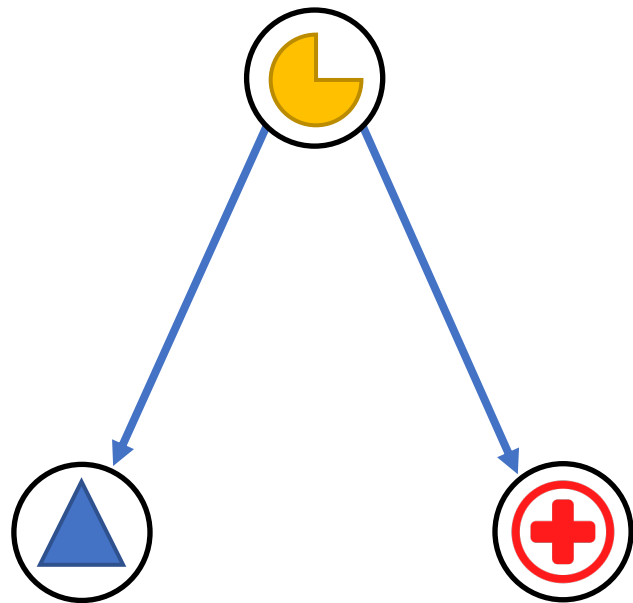
We have to deal with confounder variables



Positive impact
 Negative impact

0.5	0.7	1.5
1.4	1.8	2.0
2.7	3.0	3.7
2.3	2.9	3.2
1.8	1.9	2.3
1.6	1.9	2.2
0.6	0.9	1.4
0.3	0.5	1.0
1.1	1.3	1.4

We have to deal with confounder variables



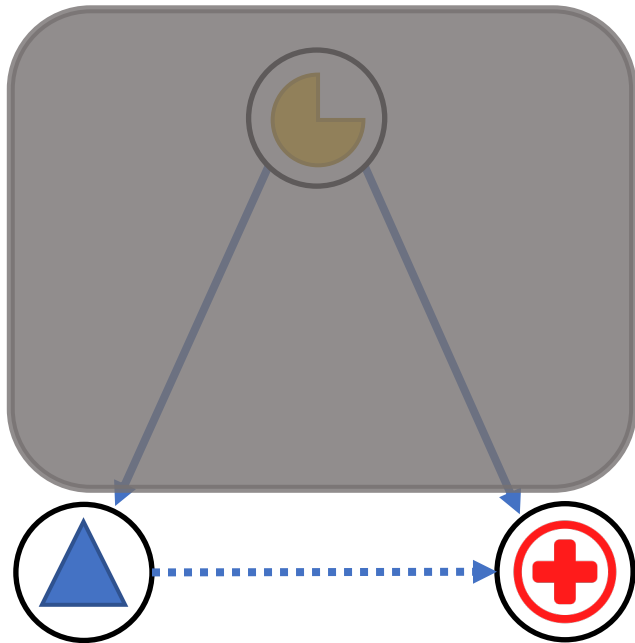
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Our focus

and
 are correlated

We have to deal with confounder variables



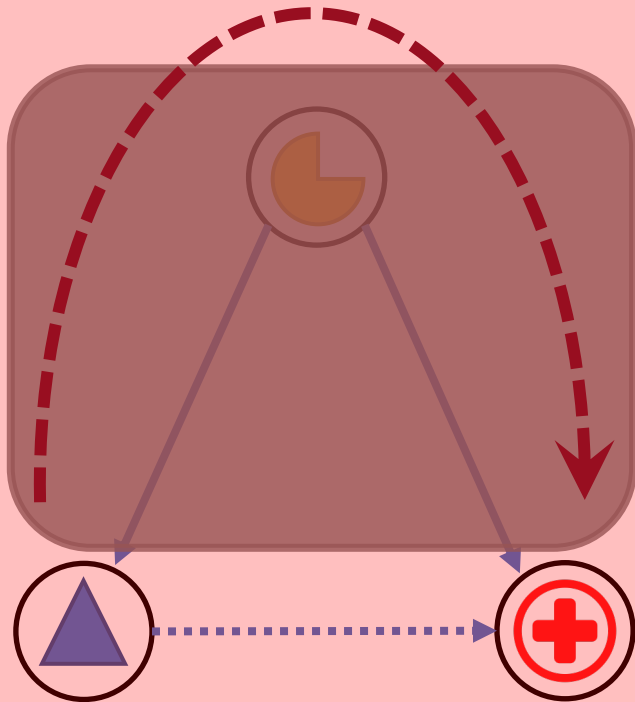
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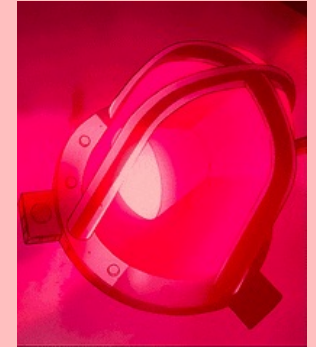
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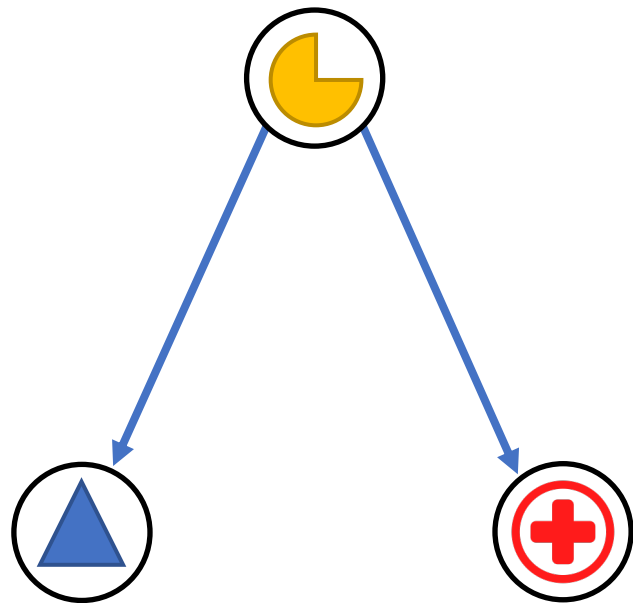
Our focus



and
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**CORRELATION
IS NOT
CAUSATION**

We have to deal with confounder variables



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Our focus

and are correlated

and are **NOT** correlated by conditioning on

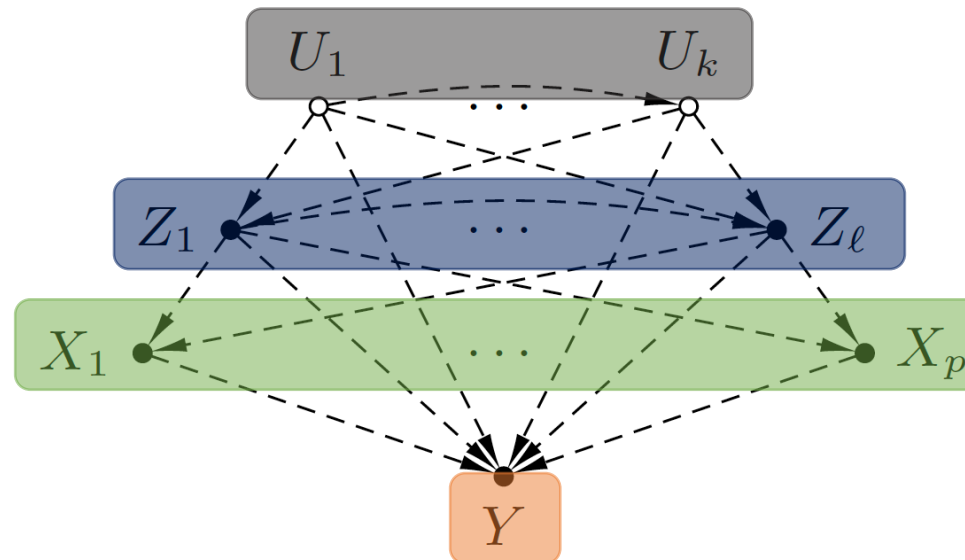


We are currently working with Breast Cancer Data

Set of confounders Z

Set of actionables X

Target T



We are currently working with Breast Cancer Data

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Germline genomic alterations

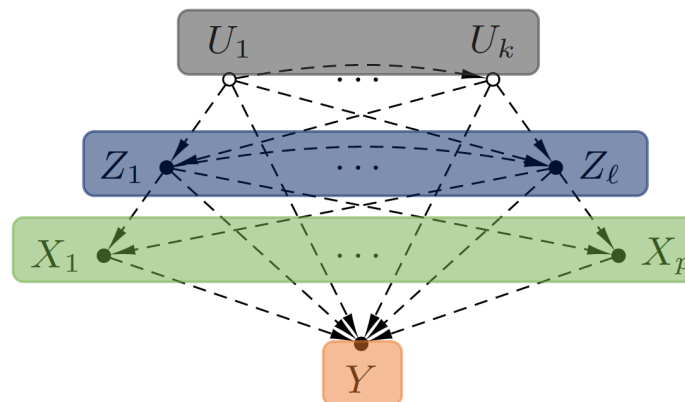
Somatic genomic alterations

Cancer type

Sex

Age

...



We are currently working with Breast Cancer Data

Set of confounders Z

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Target T



Germline genomic alterations

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Cancer type

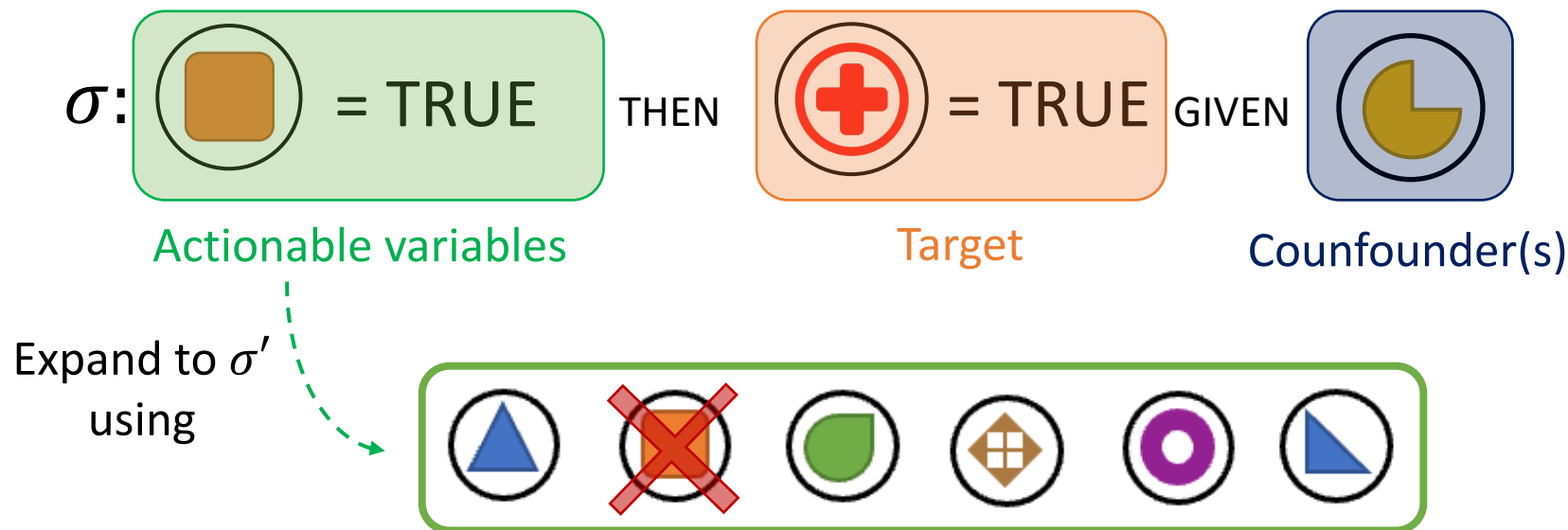
Sex

Age

...

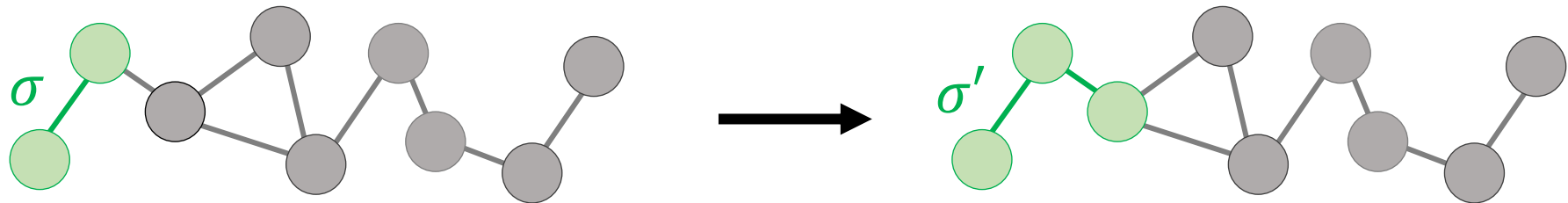
$TP53_{som} = 1 \wedge ERBB2_{loh} = 1 \rightarrow Basal \mid history\ other\ malignancy$

[Budhathoki et al.] proposed the **reliable** rule effect estimation framework and developed a branch and bound algorithm for the discovery task.



Budhathoki, K., Boley, M. and Vreeken, J., 2021. Discovering reliable causal rules. In *Proceedings of the 2021 SIAM International Conference on Data Mining (SDM)* (pp. 1-9). Society for Industrial and Applied Mathematics.

- Guarantees for multiple hypothesis testing
- Prove that the general problem is NP-hard
- Exploit dependency graph G for rule expansion



- Extensive experiments on breast cancer data
- Correction for multiple hypothesis testing issues – Currently working on data dependent corrections with Rademacher Averages

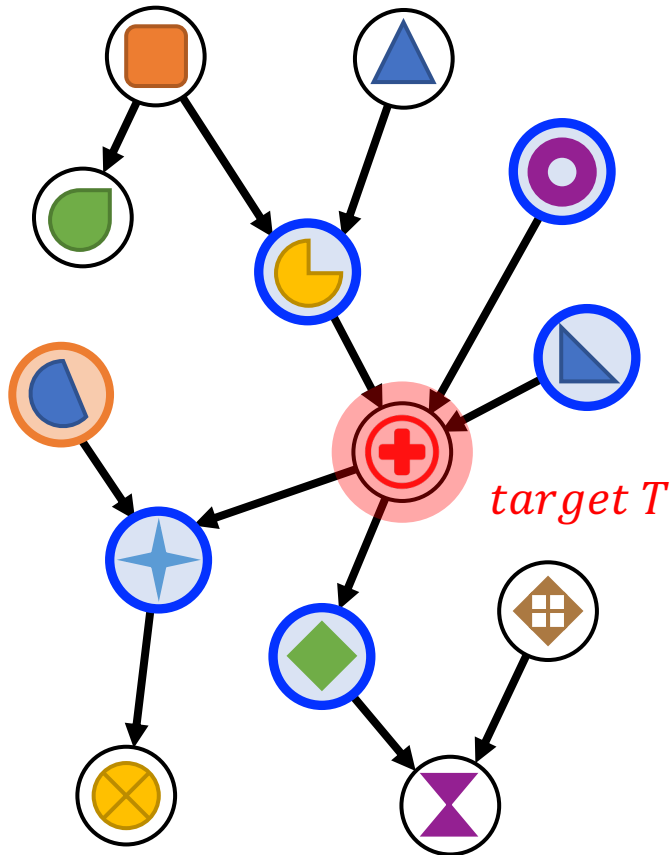
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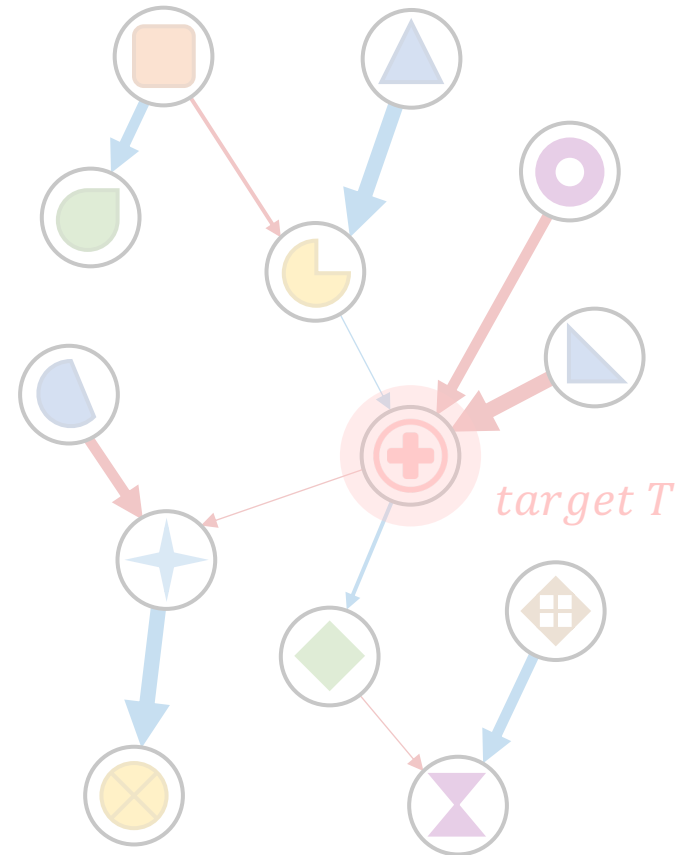
1 - Structure discovery

Discover causally related variables to a target



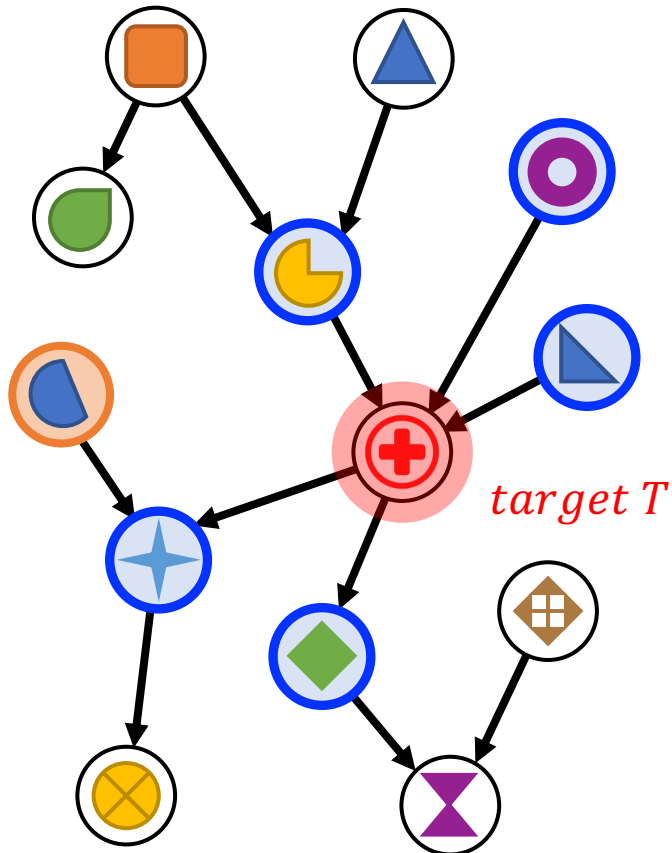
2 - Effect estimation

Evaluate effect of causal rules



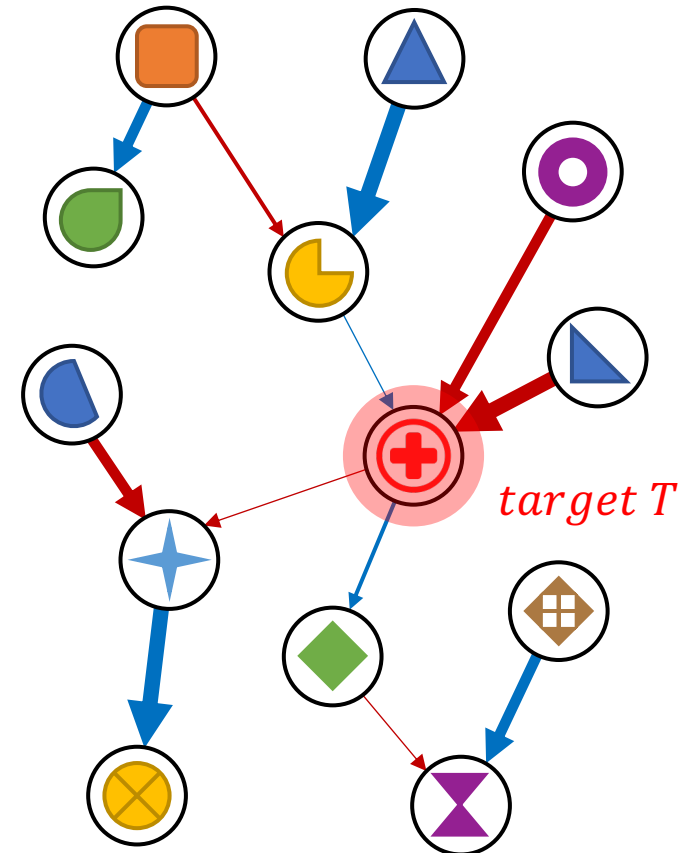
1 - Structure discovery

Discover causally related variables to a target



2 - Effect estimation

Evaluate effect of causal rules



Hopefully true correlation example



Hopefully true correlation example



Exploiting causality methods for knowledge discovery from observational data

Q & A TIME

 **Dario Simionato**

dario.simionato@phd.unipd.it

Padua, Italy

Code available at: <https://github.com/VandinLab/RAveL>