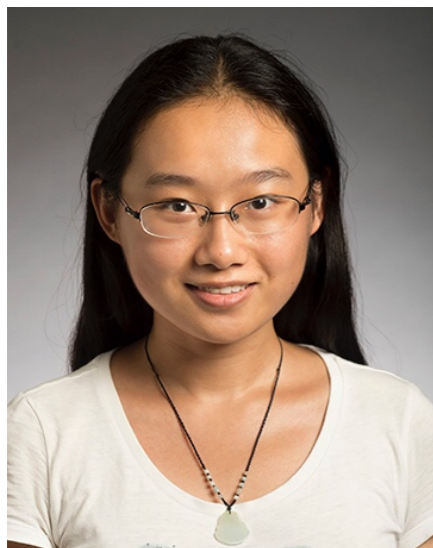


Familywise error rate control by interactive unmasking

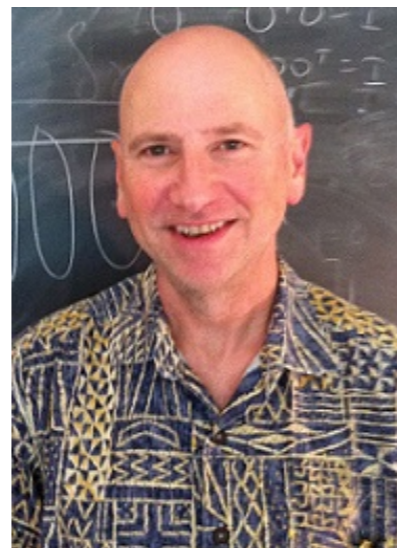
Boyan
Duan



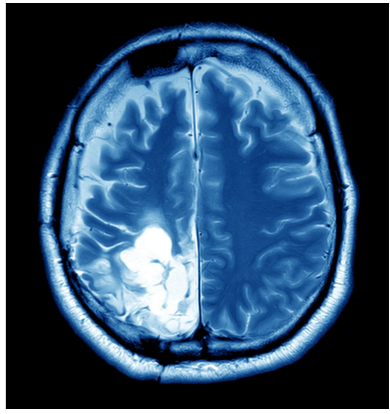
Aaditya
Ramdas



Larry
Wasserman

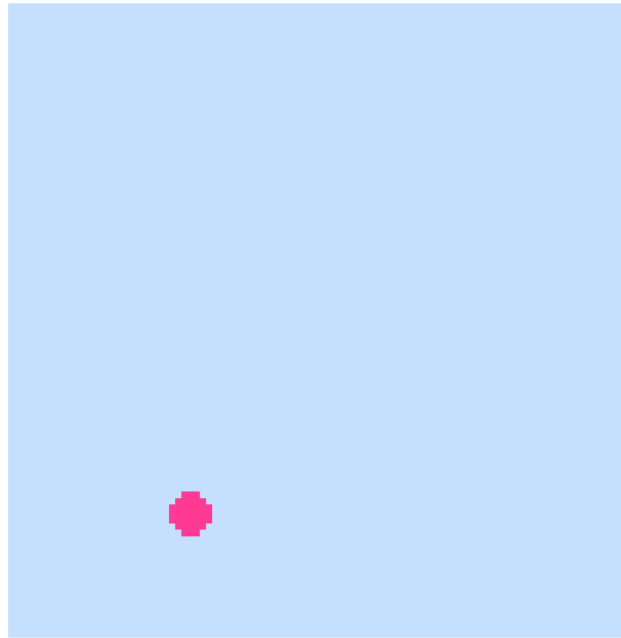
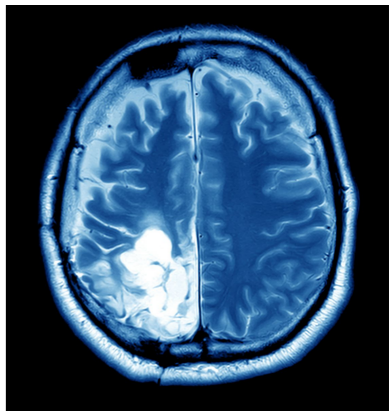


Motivating example: tumor detection in brain image



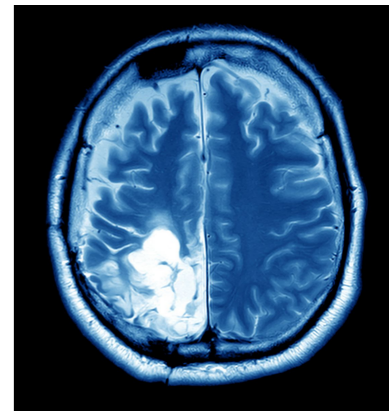
Motivating example: tumor detection in brain image

Eg. $Z_i \sim N(\mu_i, 1)$ for each pixel and test $H_i : \mu_i \leq 0$.

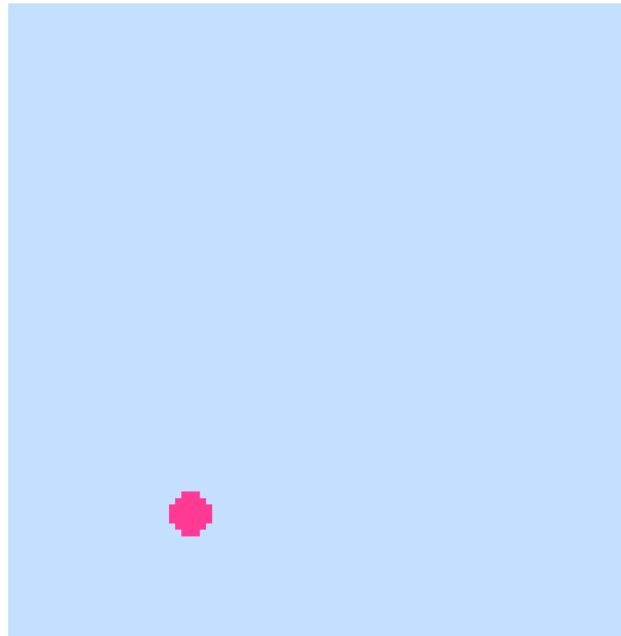


Underlying true μ_i

Motivating example: tumor detection in brain image



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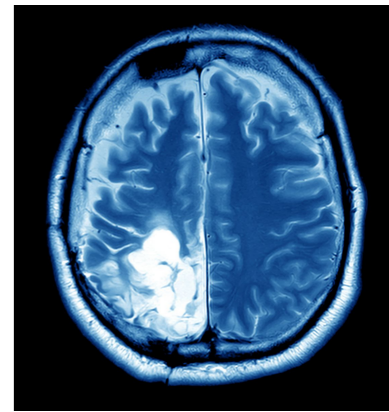


Underlying true μ_i

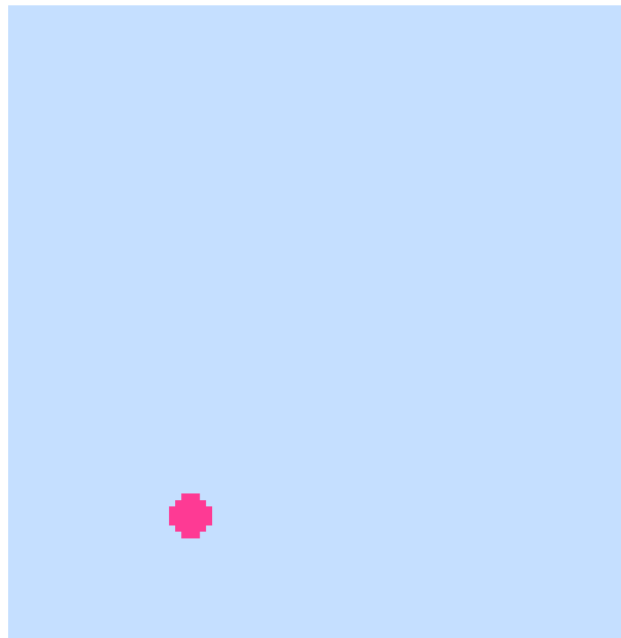
Observed p -values

$$P_i = 1 - \Phi(Z_i)$$

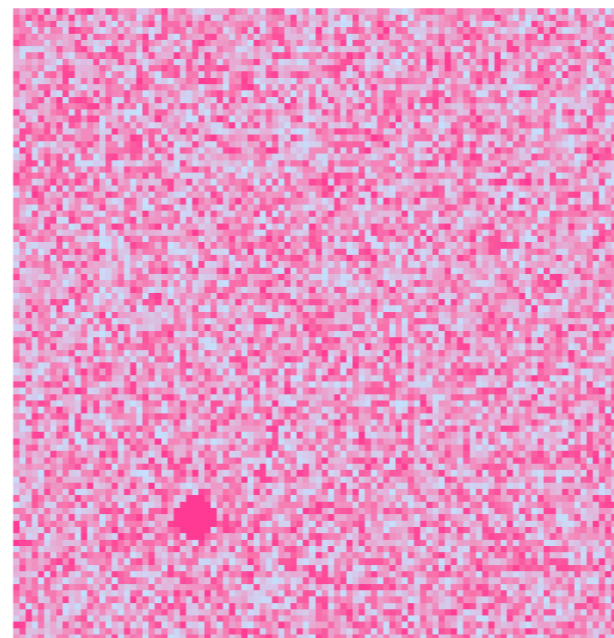
Motivating example: tumor detection in brain image



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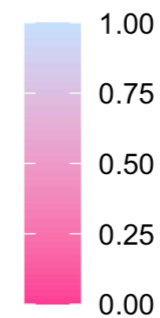


Underlying true μ_i

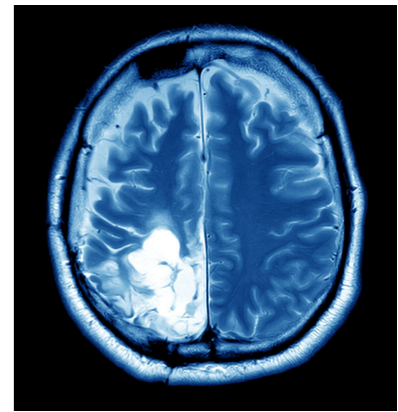


Observed p -values

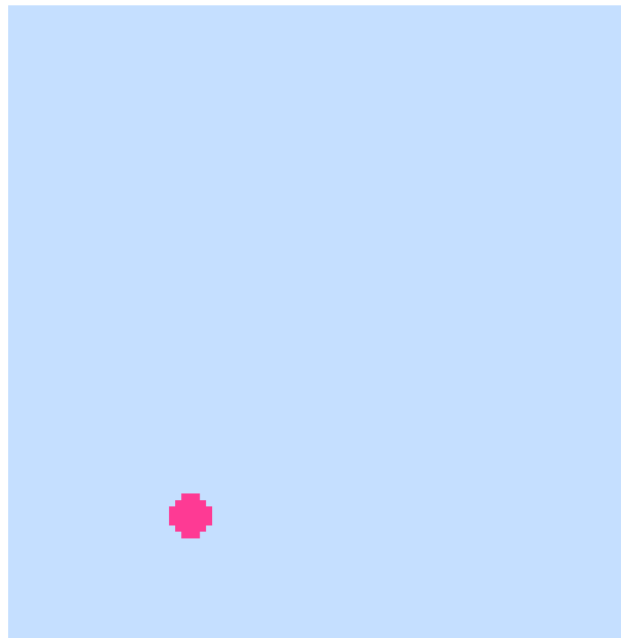
$$P_i = 1 - \Phi(Z_i)$$



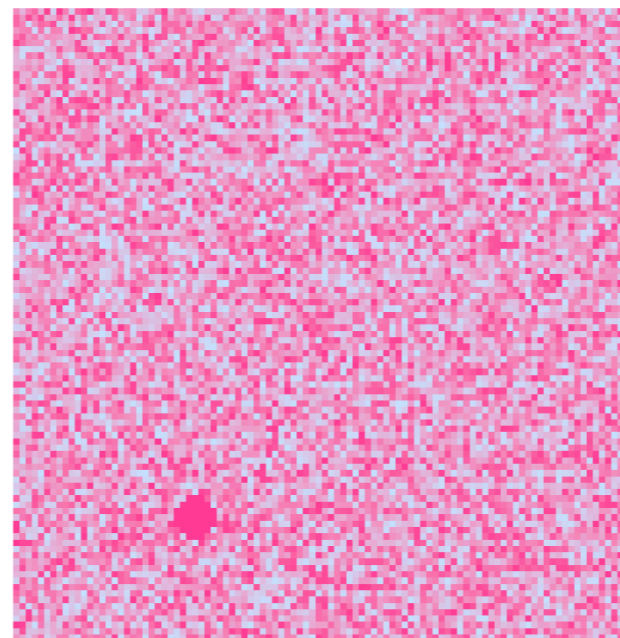
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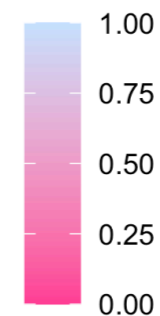


Underlying true μ_i



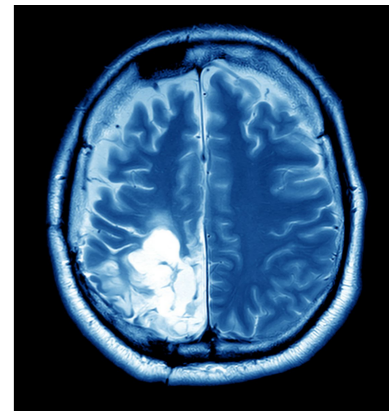
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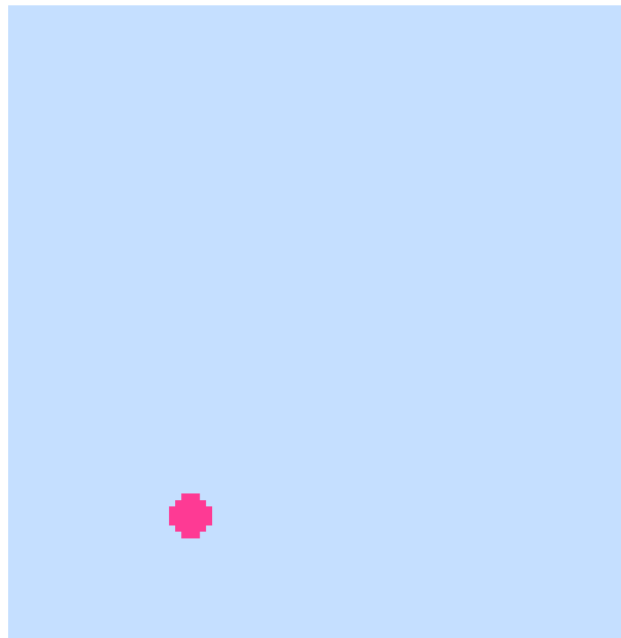


Task: identify non-nulls (decide whether to reject each H_i),

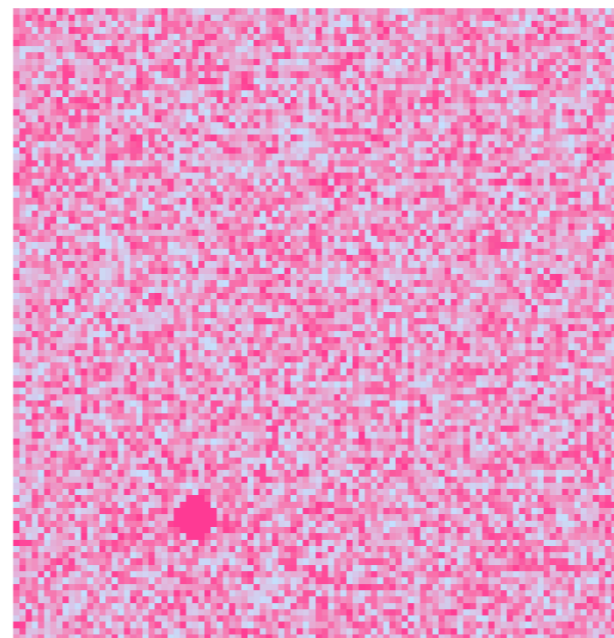
Motivating example: tumor detection in brain image



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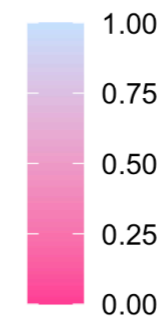


Underlying true μ_i



Observed p -values

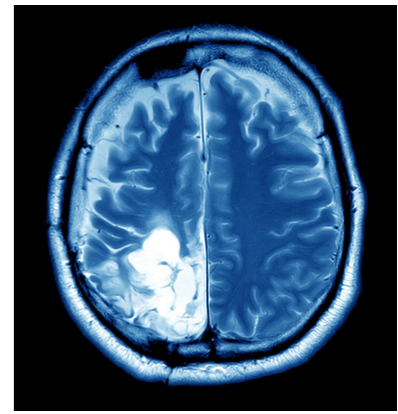
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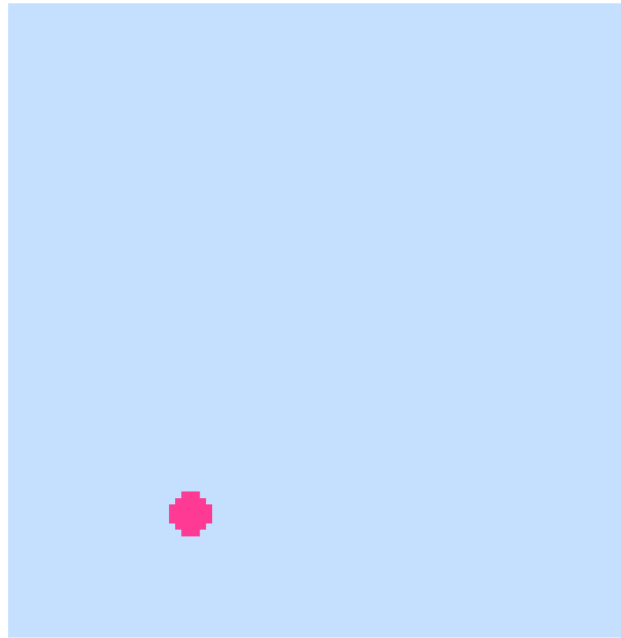
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$$\text{FWER} := \mathbb{P}(\#\text{falsely rejected nulls} \geq 1)$$

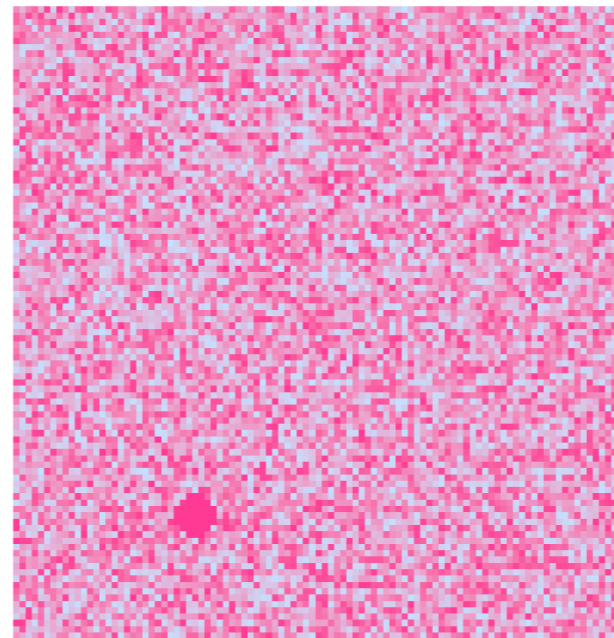
Motivating example: tumor detection in brain image



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Underlying true μ_i



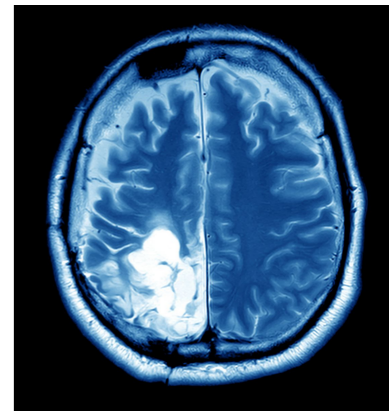
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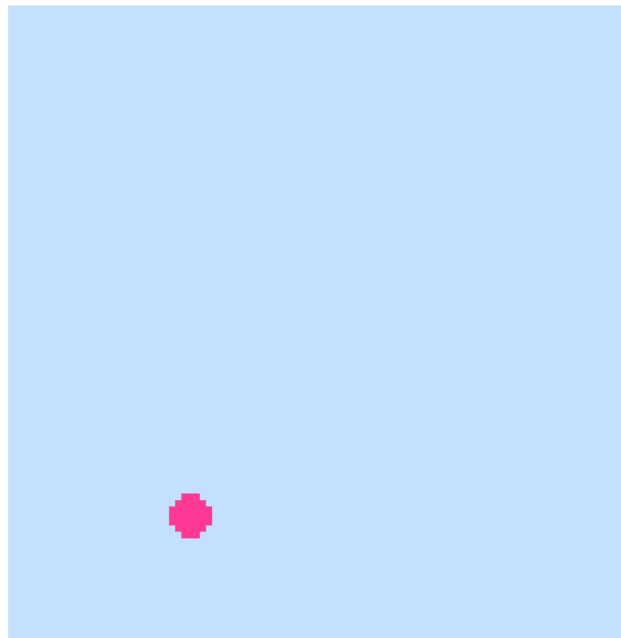
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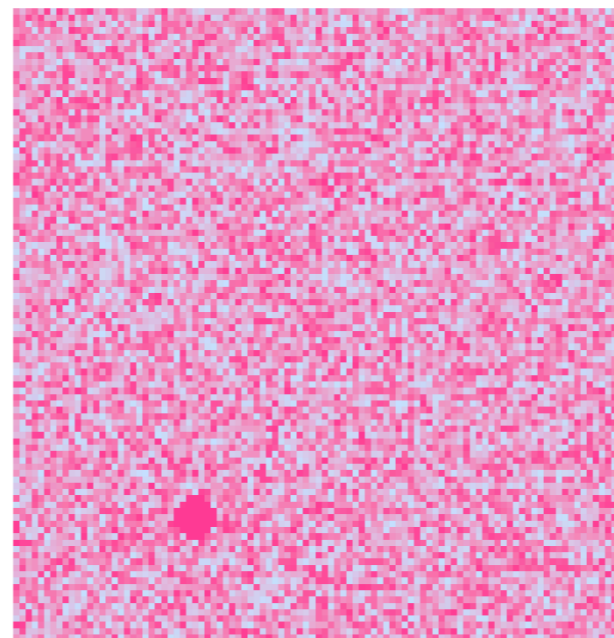
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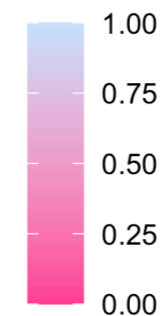


Underlying true μ_i



Observed p -values

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Task: identify non-nulls (decide whether to reject each H_i), with familywise error rate (FWER) control:

$$\text{FWER} := \mathbb{P}(\#\text{falsely rejected nulls} \geq 1) \leq \alpha \quad \text{Eg. } \alpha = 0.2$$

taking account of side information. Eg. structure, covariates...

Classical testing

Pre-fixed procedure. Eg. weighted Bonferroni: reject H_i if $P_i/w_i \leq \alpha/n$.

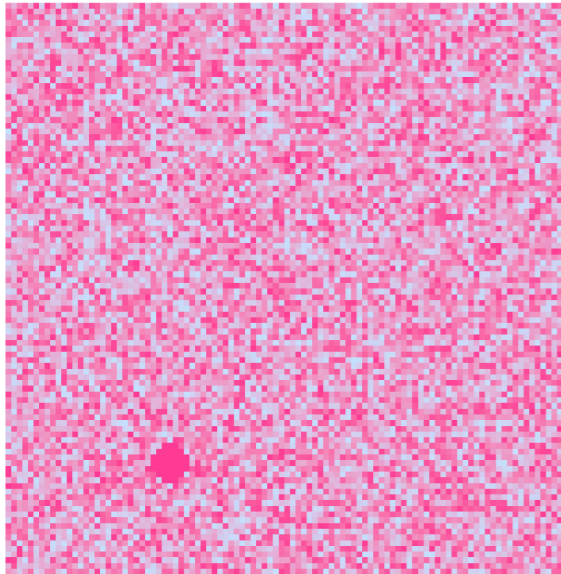
Classical testing

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Classical test
(single step)

Classical testing

Pre-fixed procedure. Eg. weighted Bonferroni: reject H_i if $P_i/w_i \leq \alpha/n$.



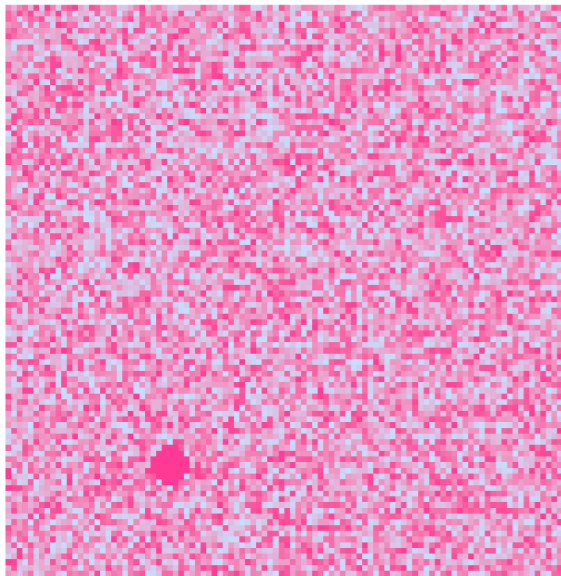
p -values

Data
(& side info)

Classical test
(single step)

Classical testing

Pre-fixed procedure. Eg. weighted Bonferroni: reject H_i if $P_i/w_i \leq \alpha/n$.



p -values

Data
(& side info)



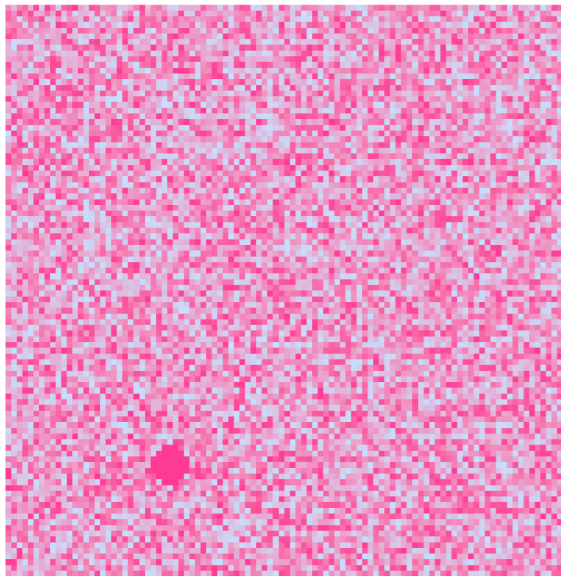
Classical test
(single step)



Rejection
set

Classical testing

Pre-fixed procedure. Eg. weighted Bonferroni: reject H_i if $P_i/w_i \leq \alpha/n$.



p -values

Data
(& side info)



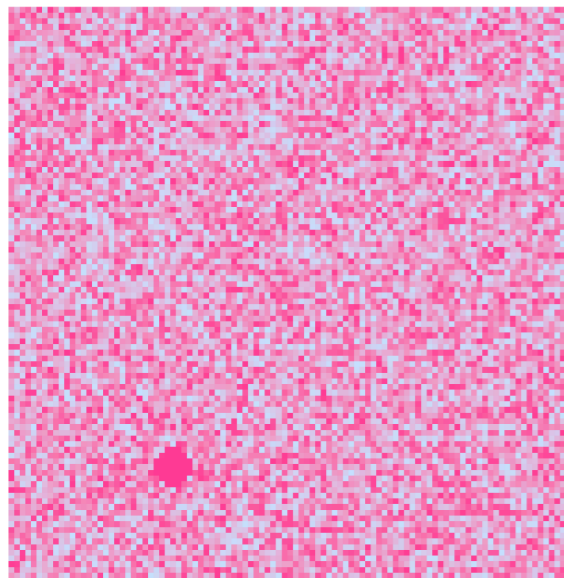
Interactive test
(multi-step)



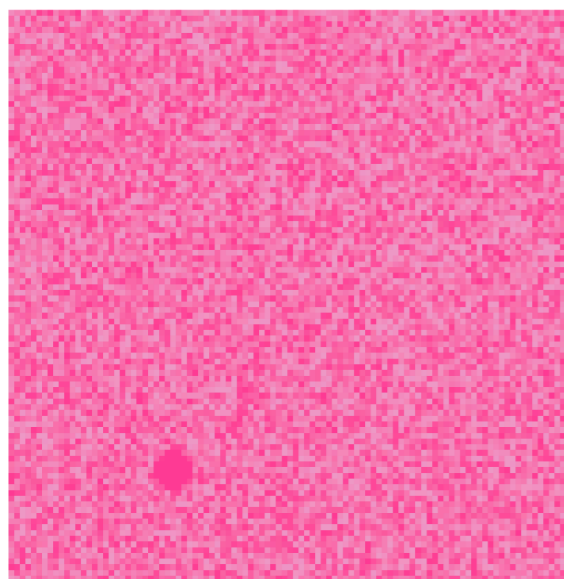
Rejection
set

Classical testing

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p -values



masked
 p -values

Masked data
(& side info)



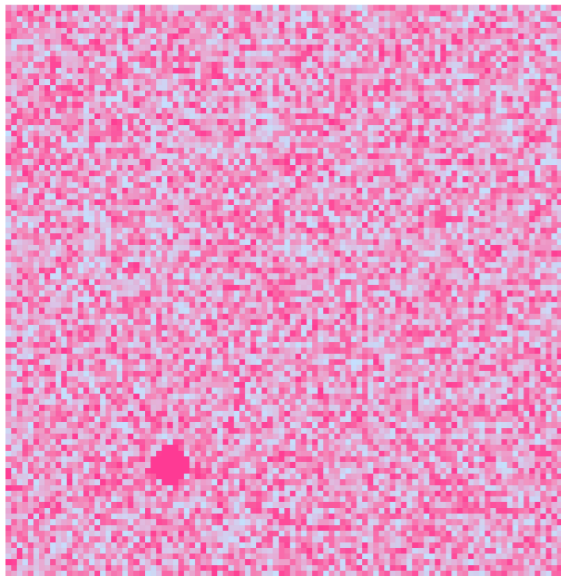
Interactive test
(multi-step)



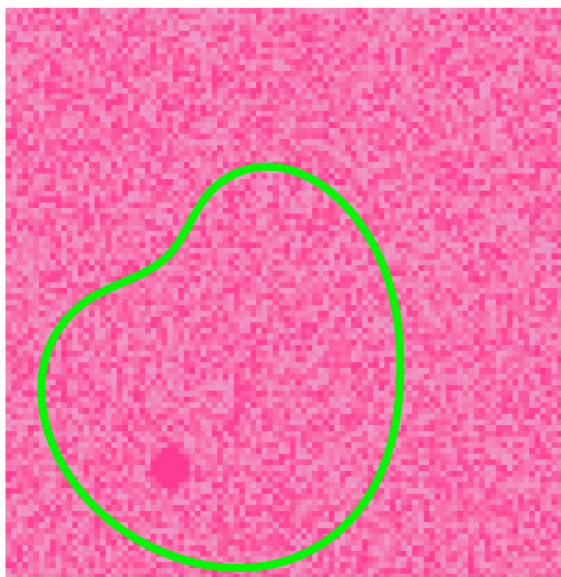
Rejection
set

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p -values



masked
 p -values

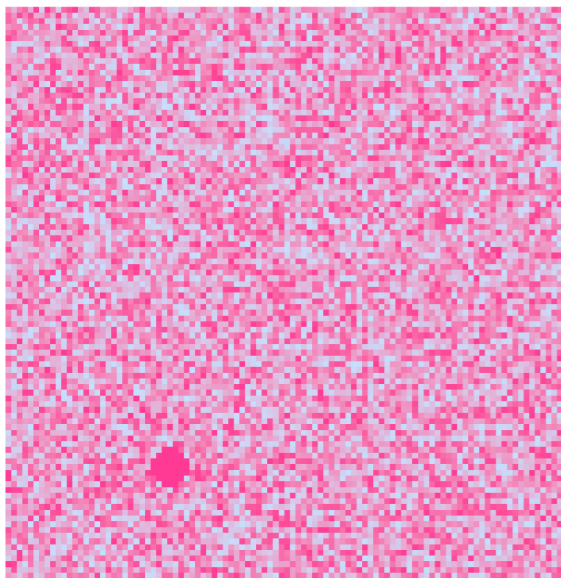
Masked data
(& side info)

Interactive test
(multi-step)

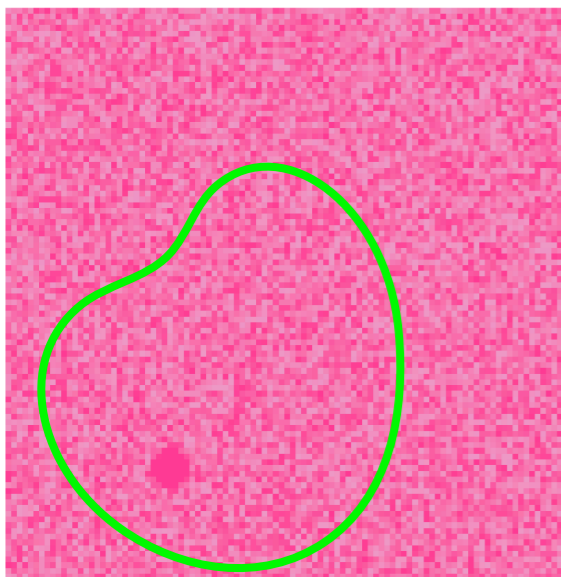
Rejection
set

Classical testing

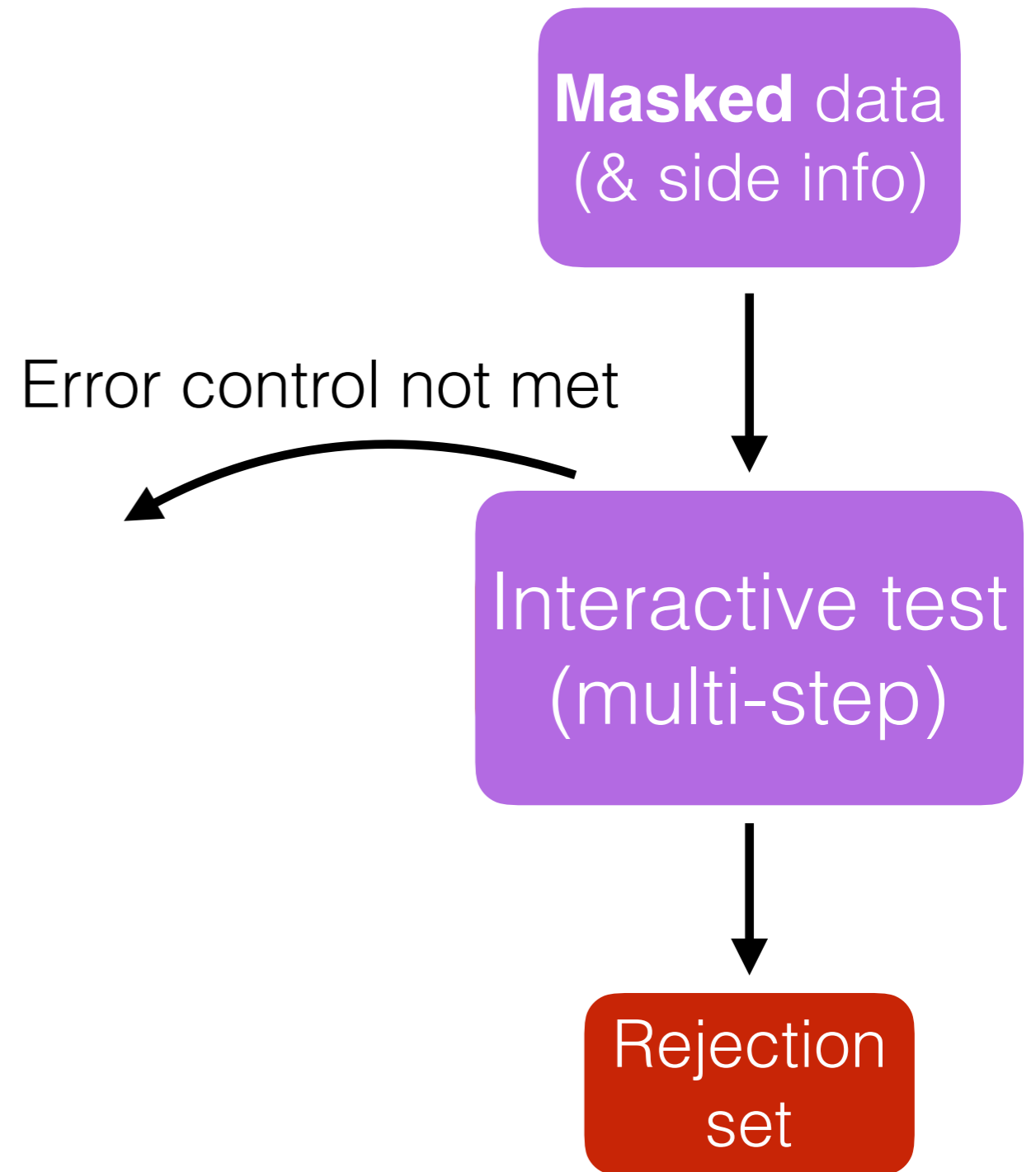
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p -values



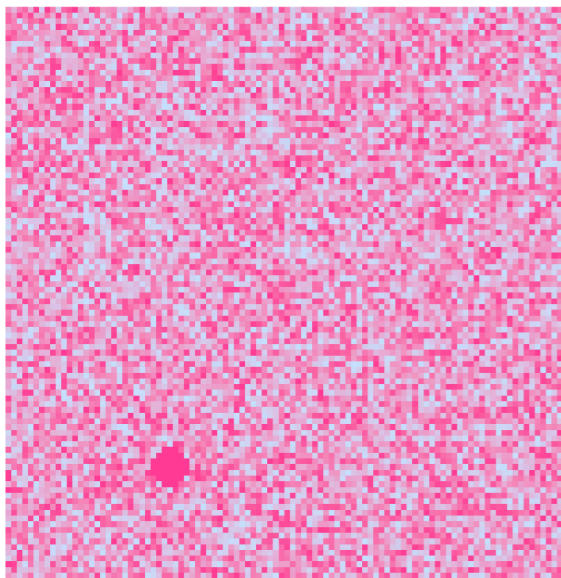
masked
 p -values



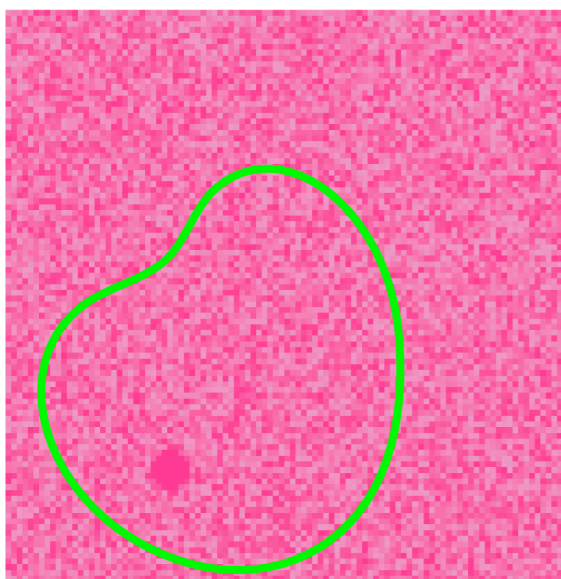
Interactive tests with FDR control:
Lei, Fithian (2018); Lei, Ramdas, Fithian (2019)

Classical testing

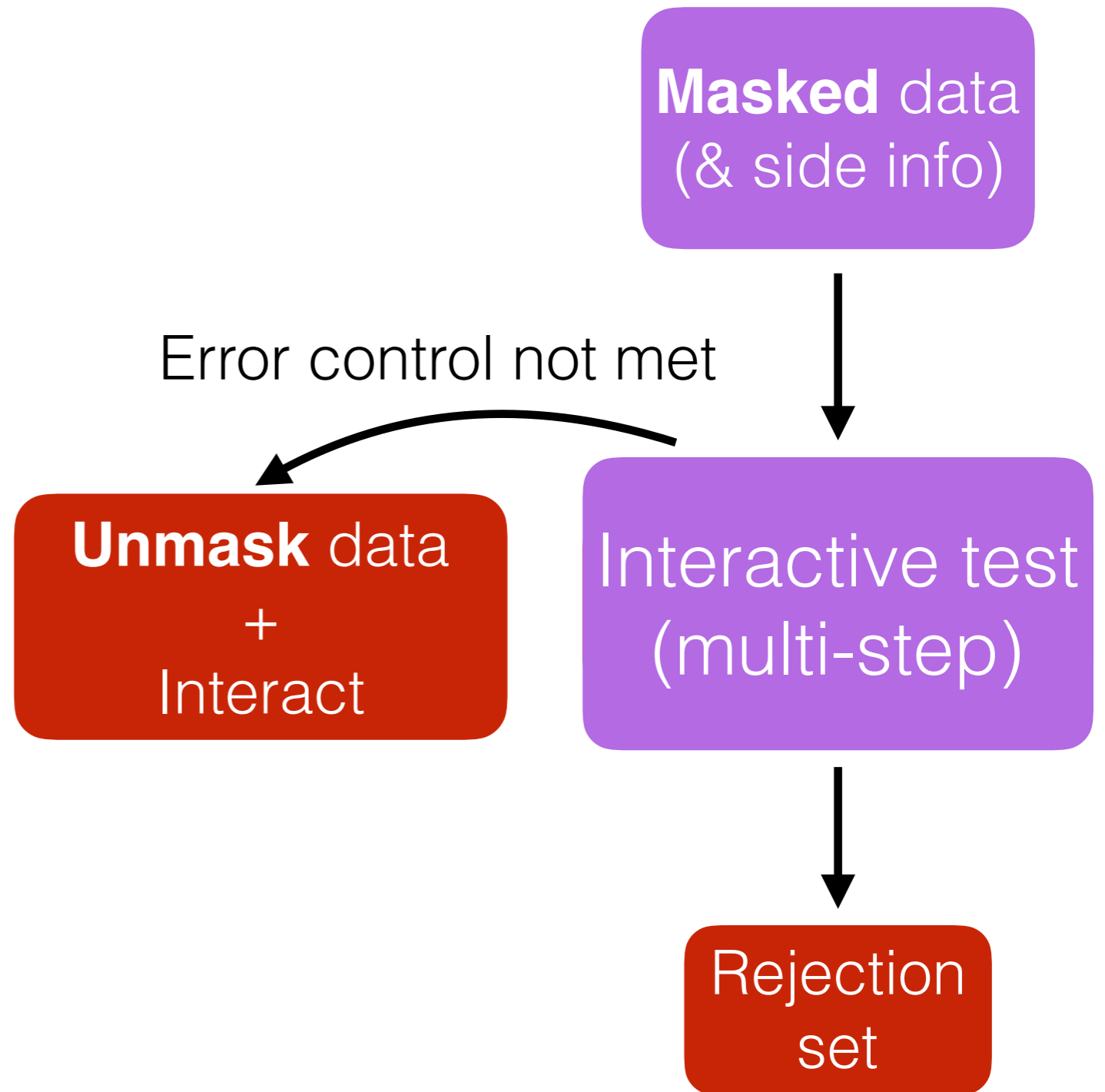
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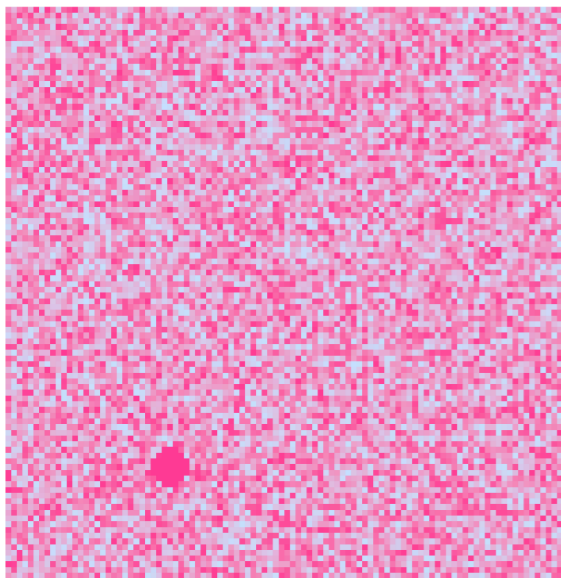
masked
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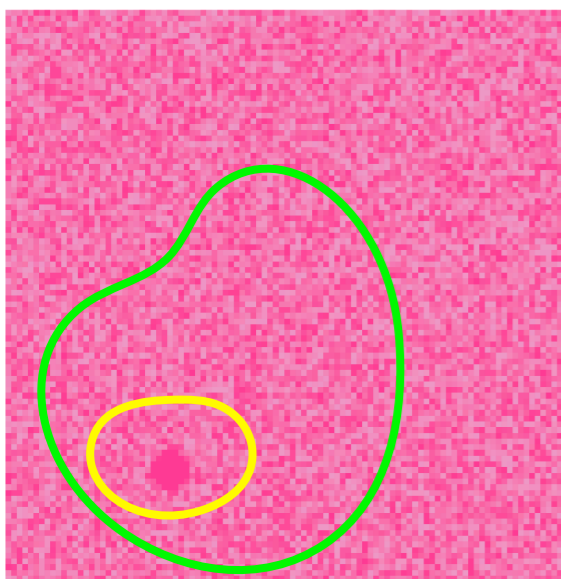
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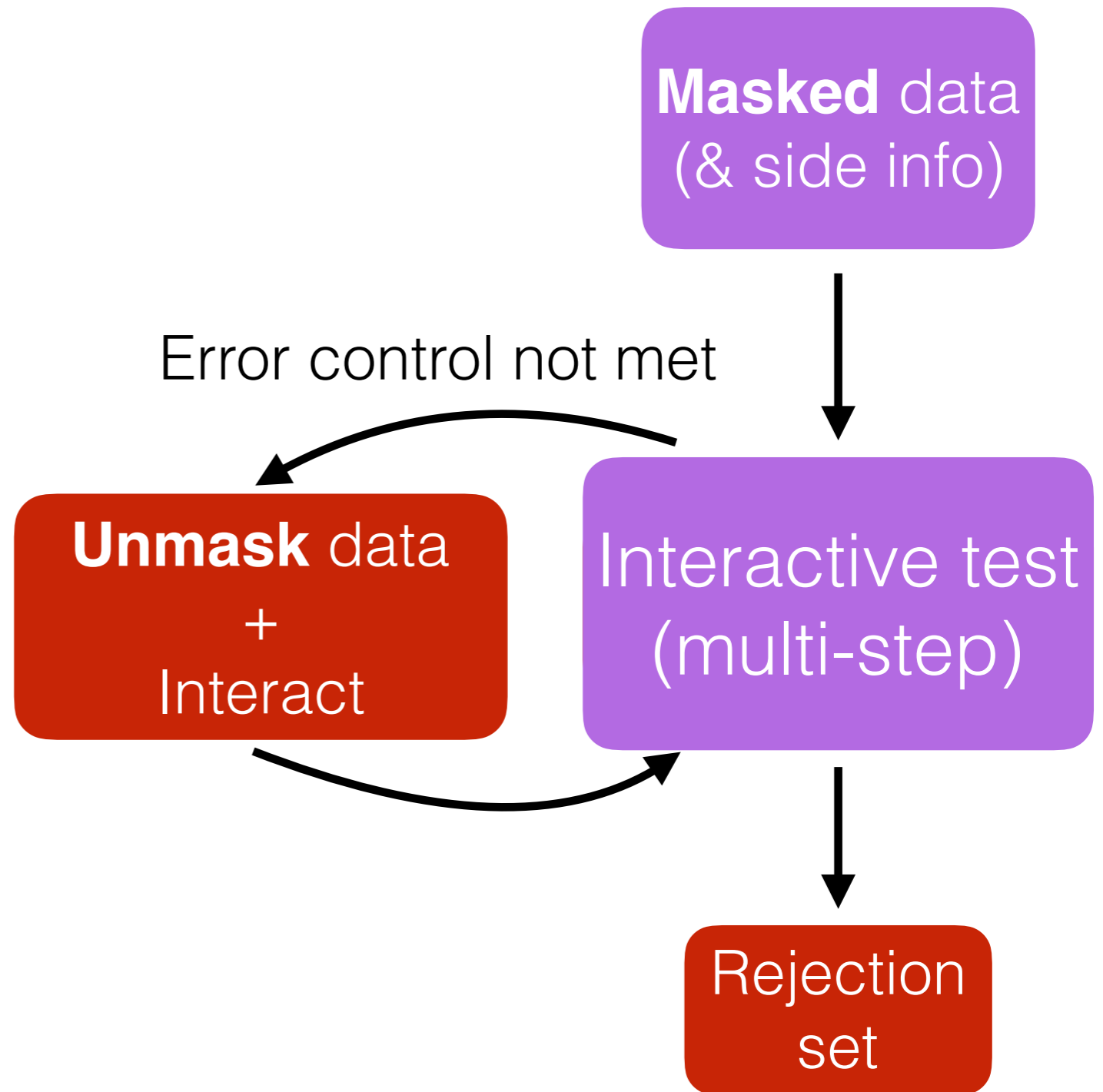
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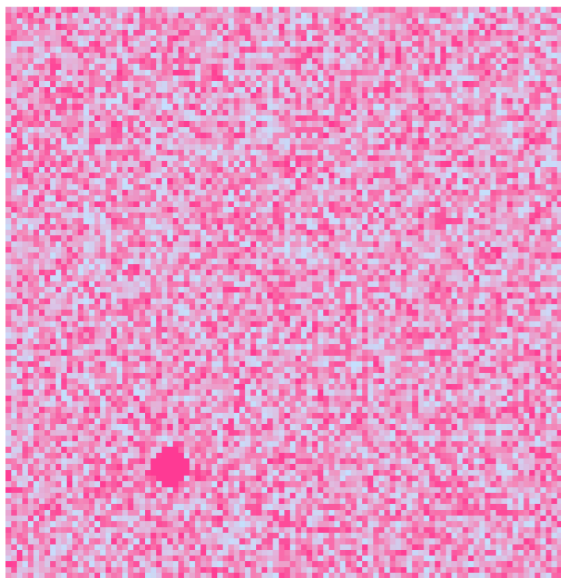
masked
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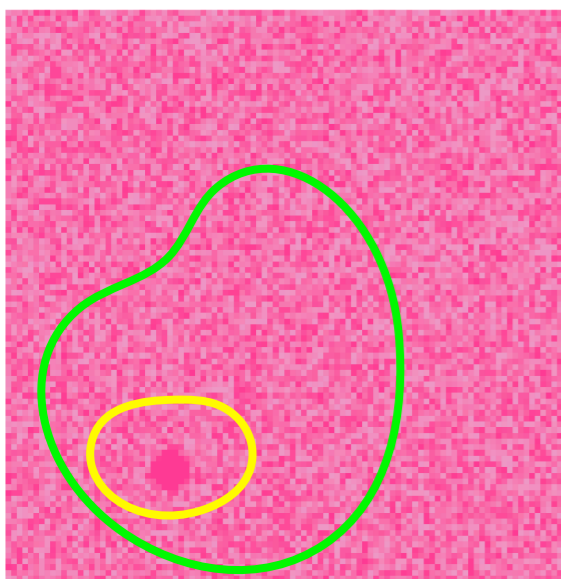
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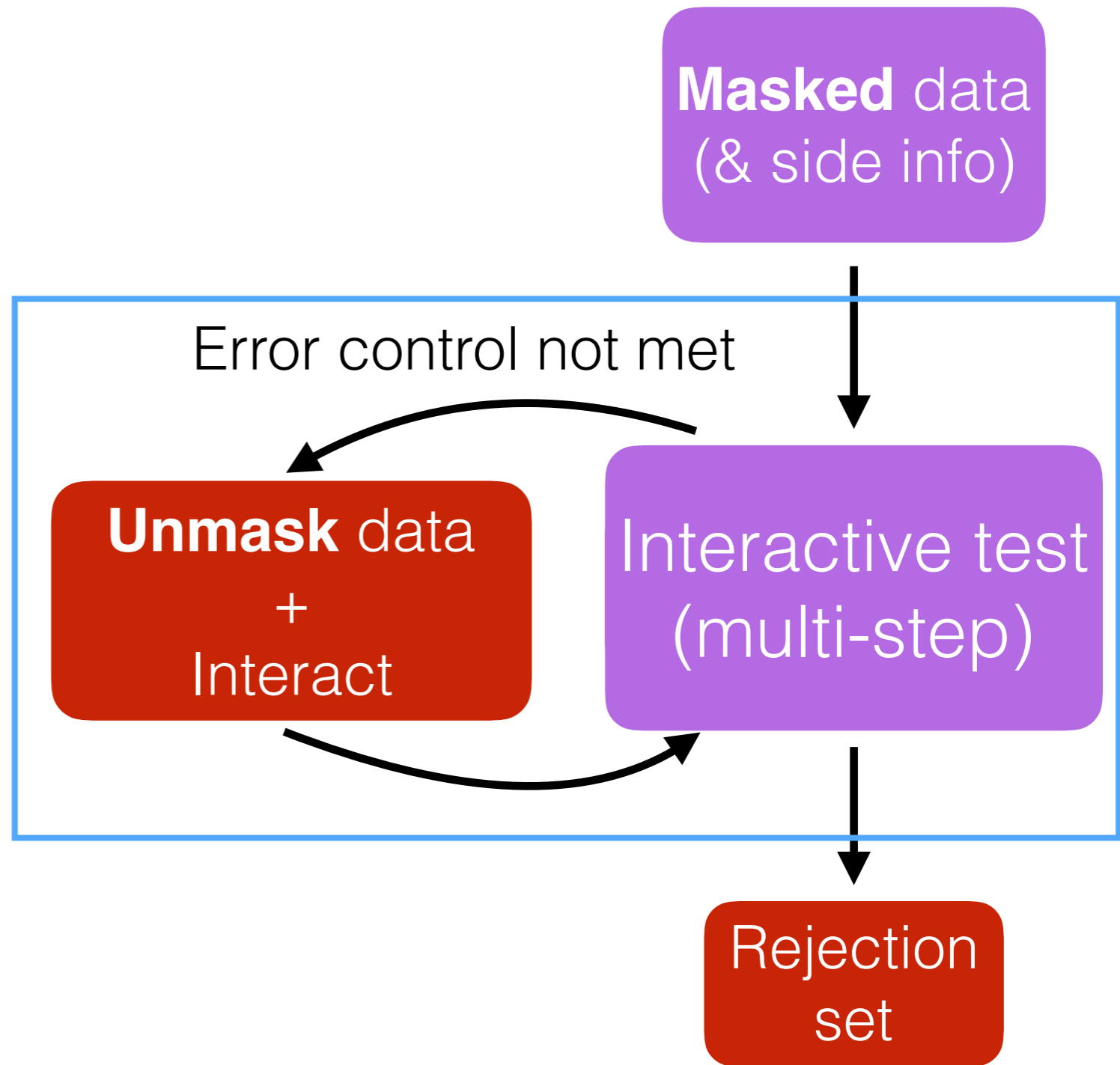
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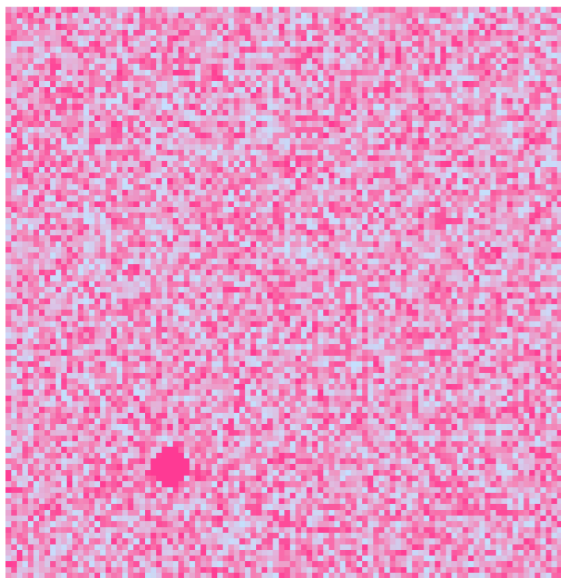
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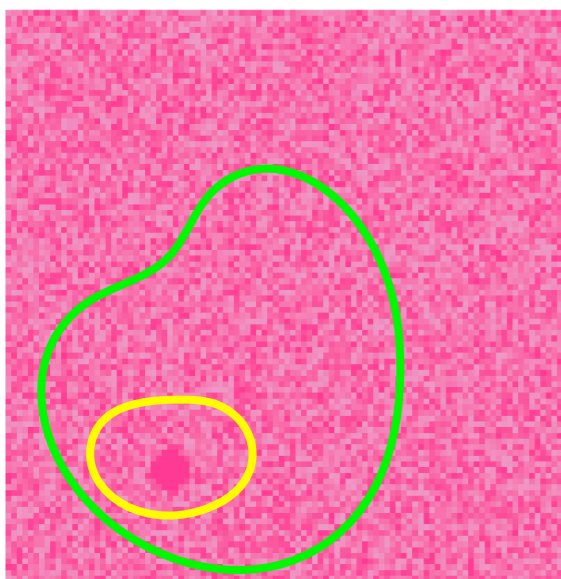
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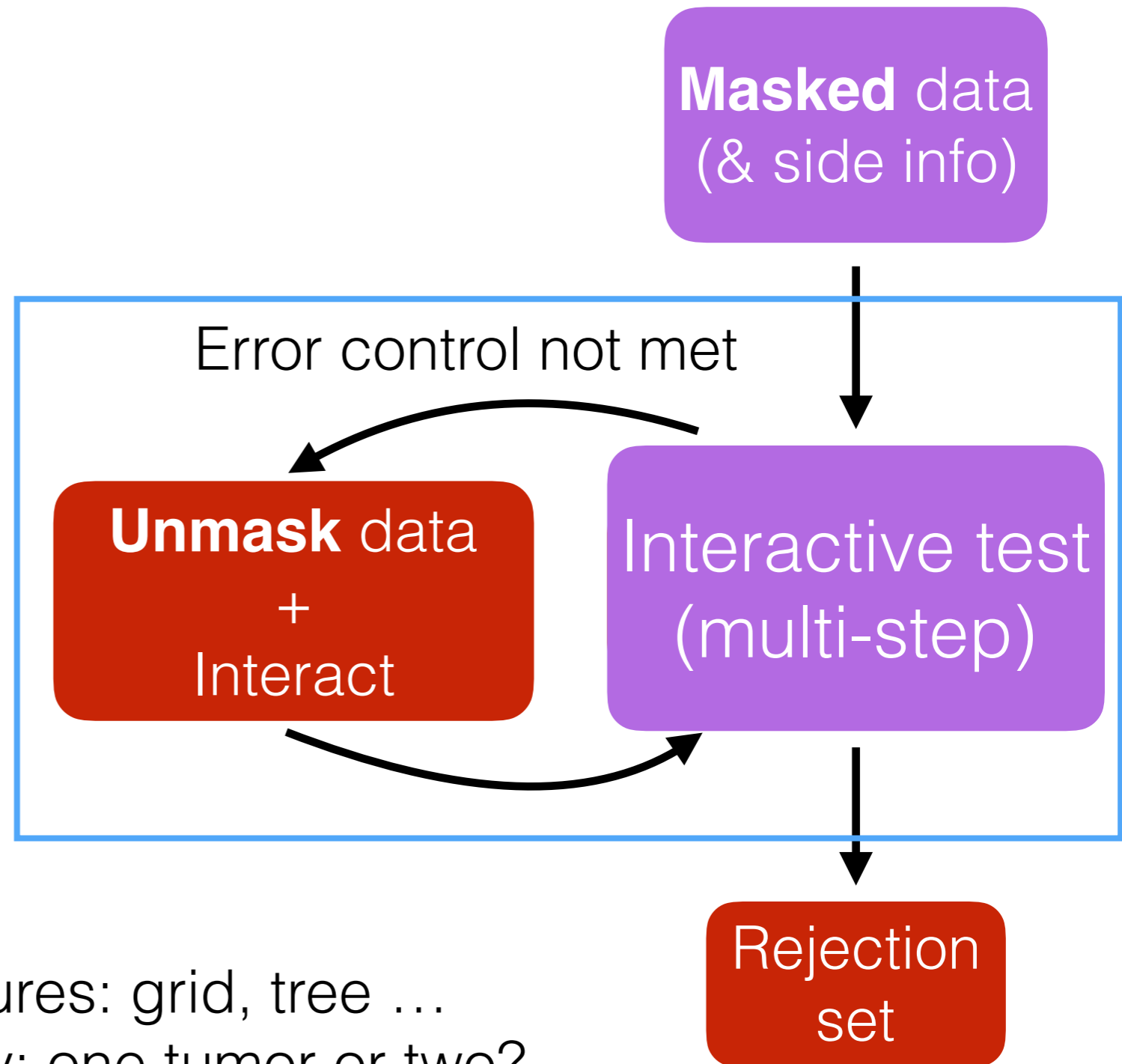
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p -values



masked
 p -values



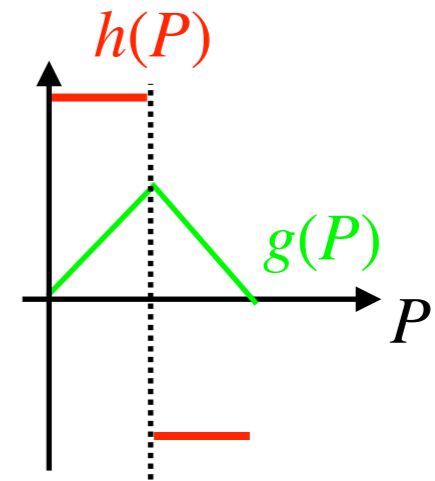
Interactive testing

- Accommodate various structures: grid, tree ...
- Be revised manually on the fly: one tumor or two?

Component 1: mask p -values

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$$P \longrightarrow \begin{cases} g(P) = \min\{P, 1 - P\} \\ h(P) = 2 \cdot \mathbb{1}\{P < 0.5\} - 1 \end{cases}$$



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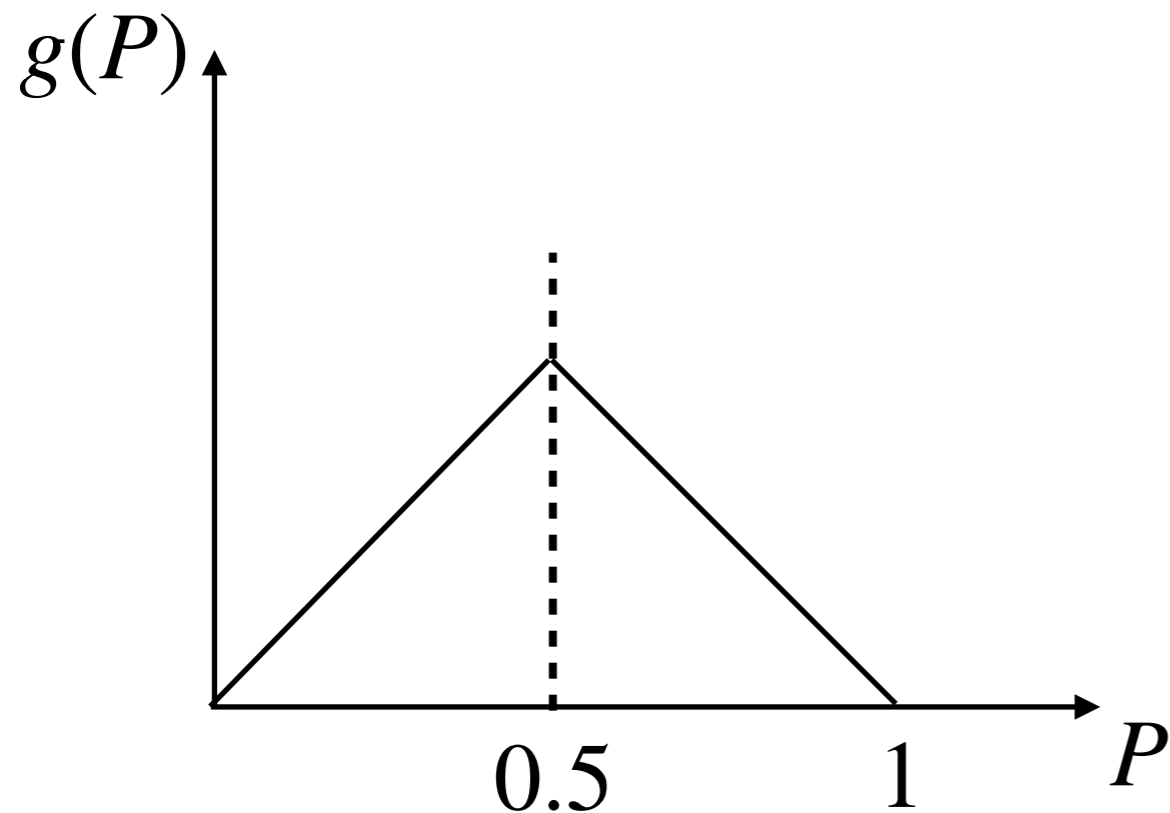
Independent
for nulls

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Independent
for nulls

Small $g(P)$ indicate non-nulls

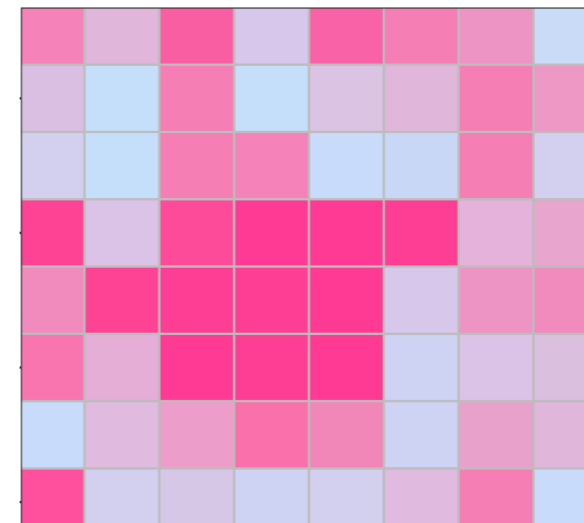
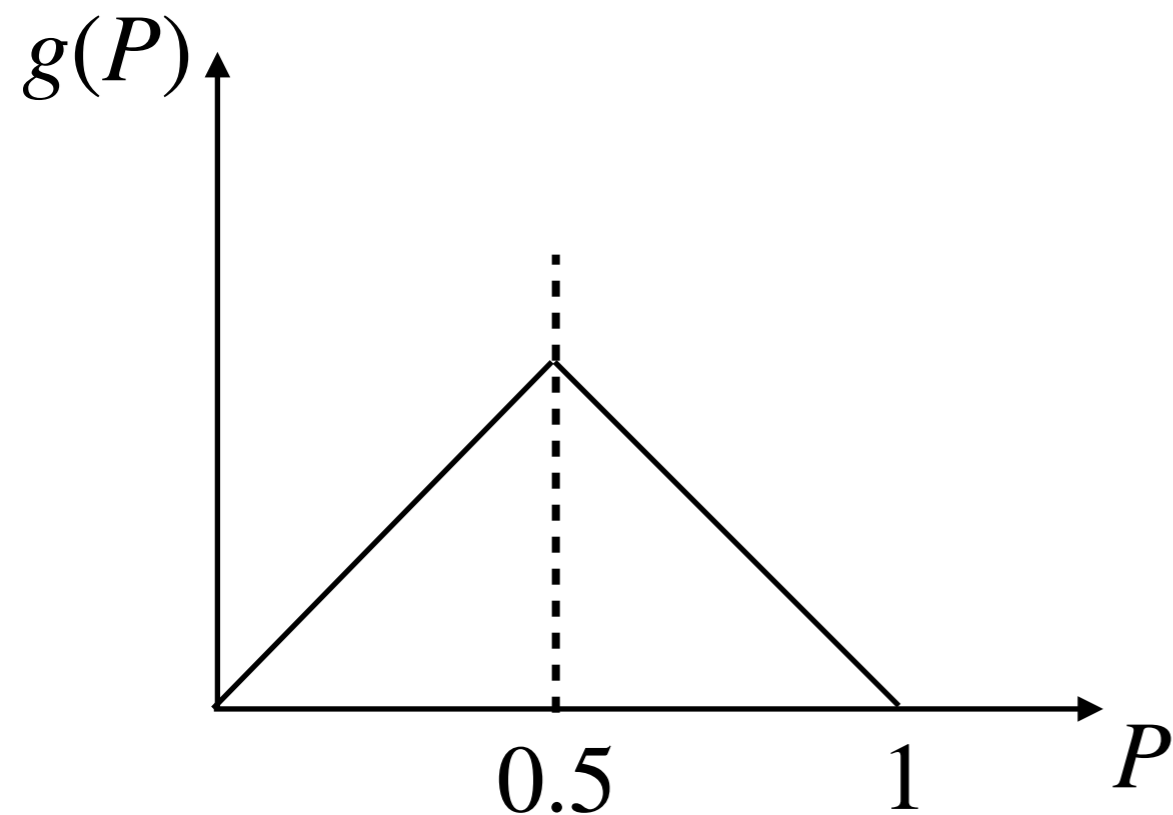


Component 1: mask p -values

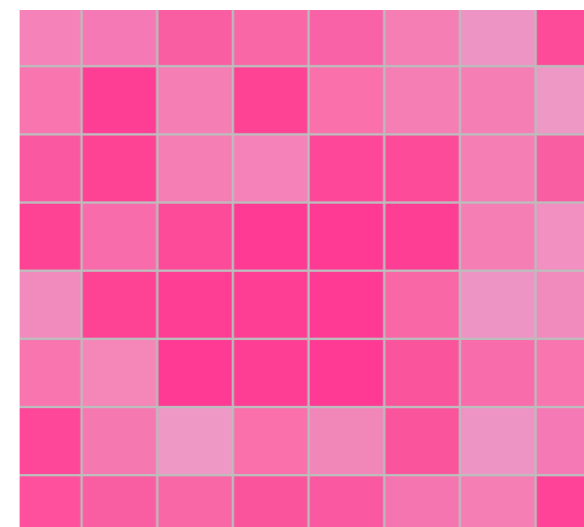
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P



$g(P)$

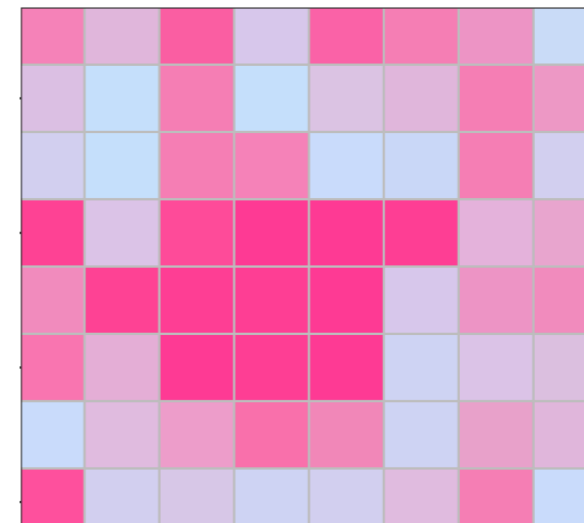
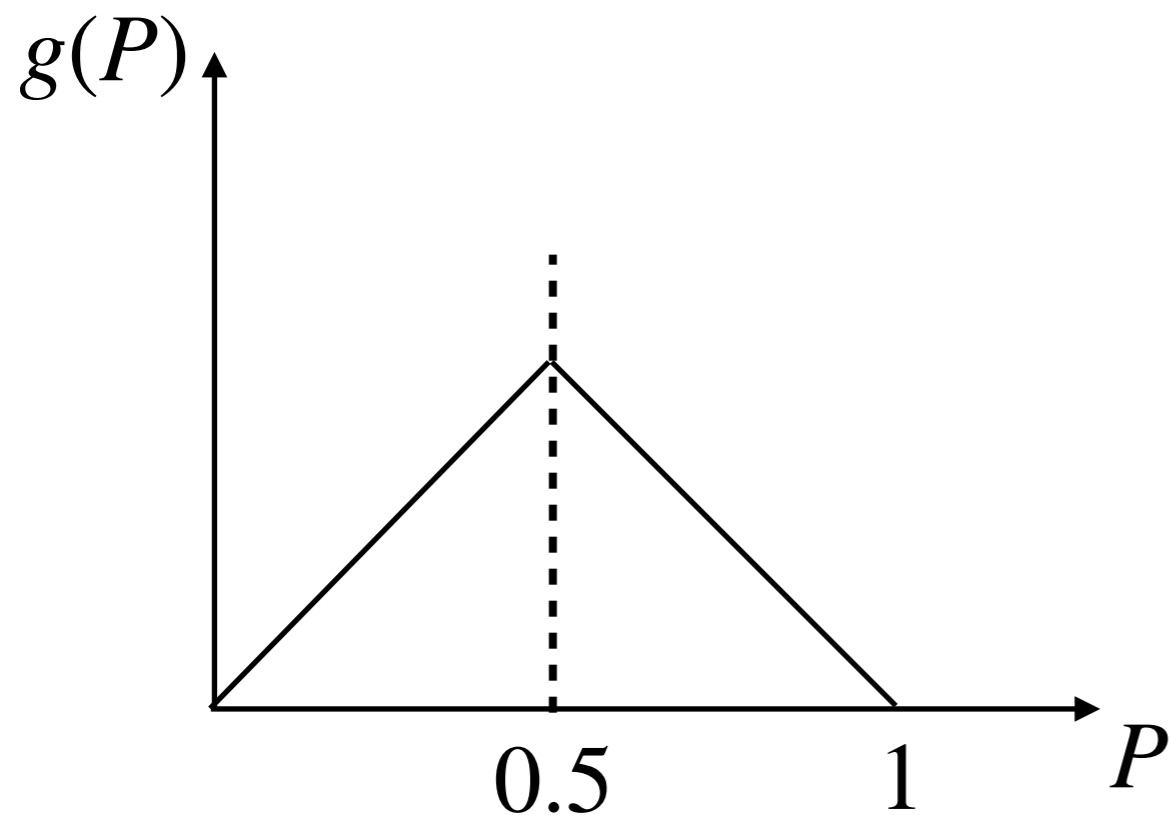
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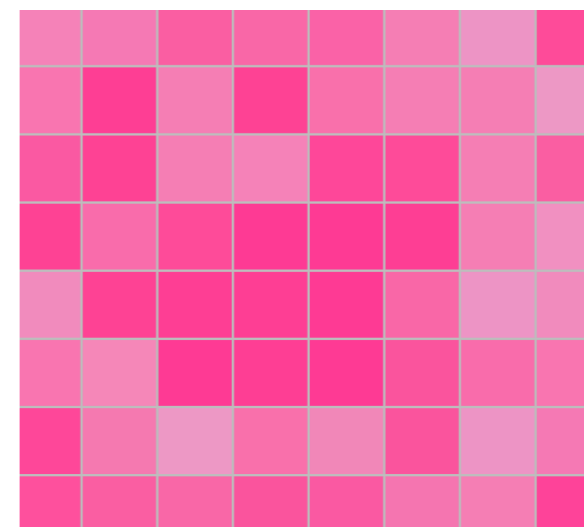
Independent
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Note: above masking works for FDR control, but not FWER control.

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P



$g(P)$

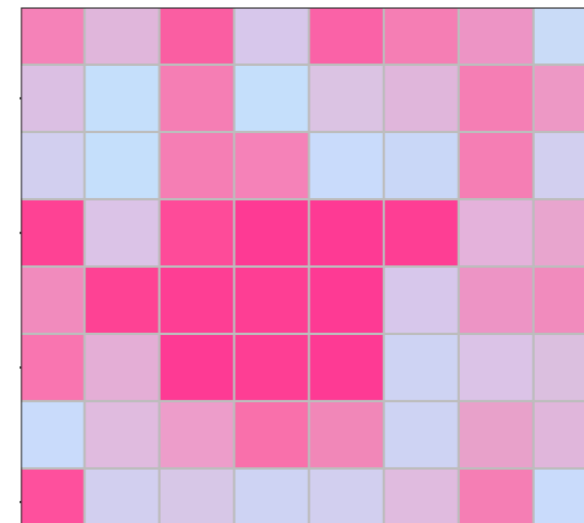
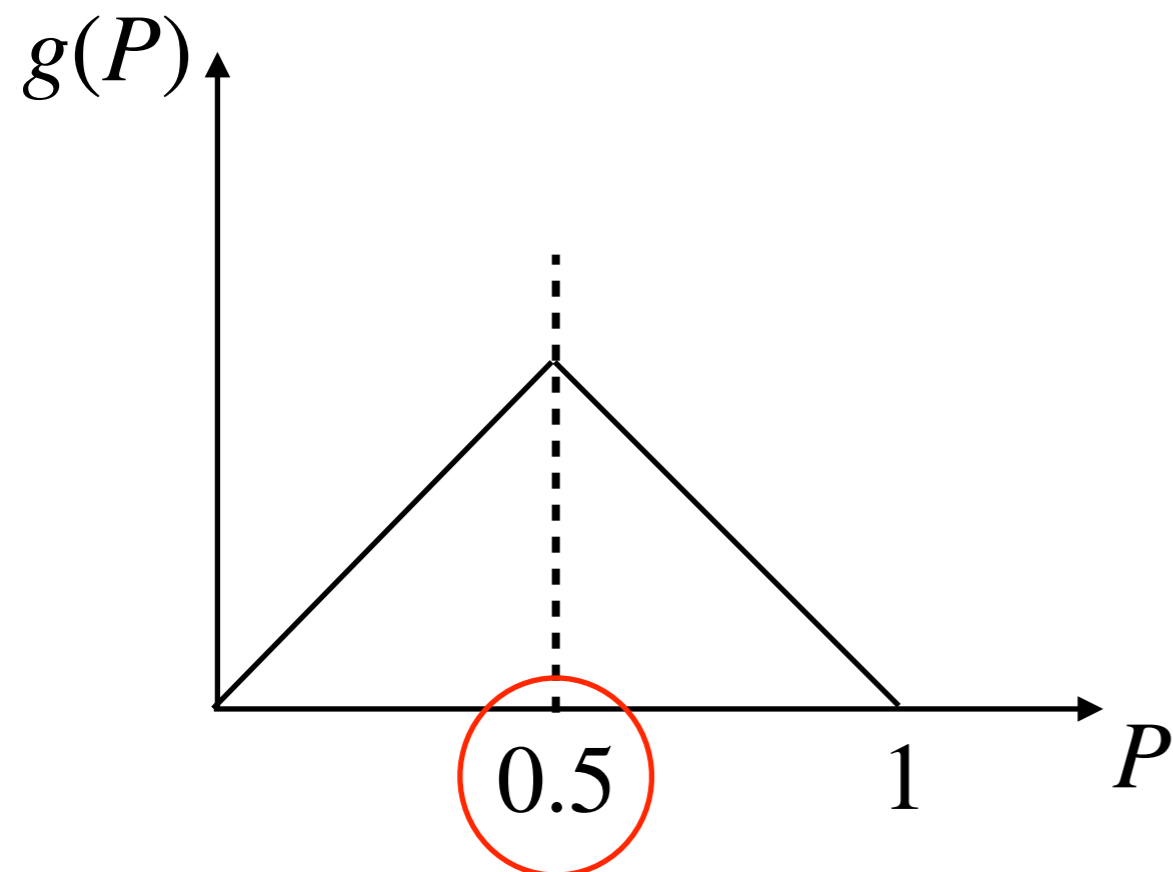
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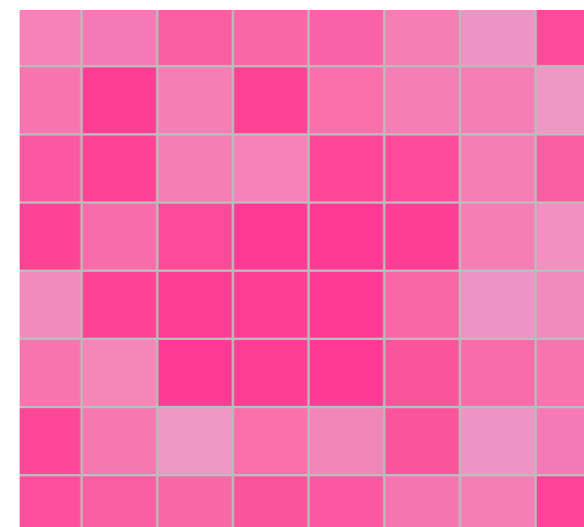
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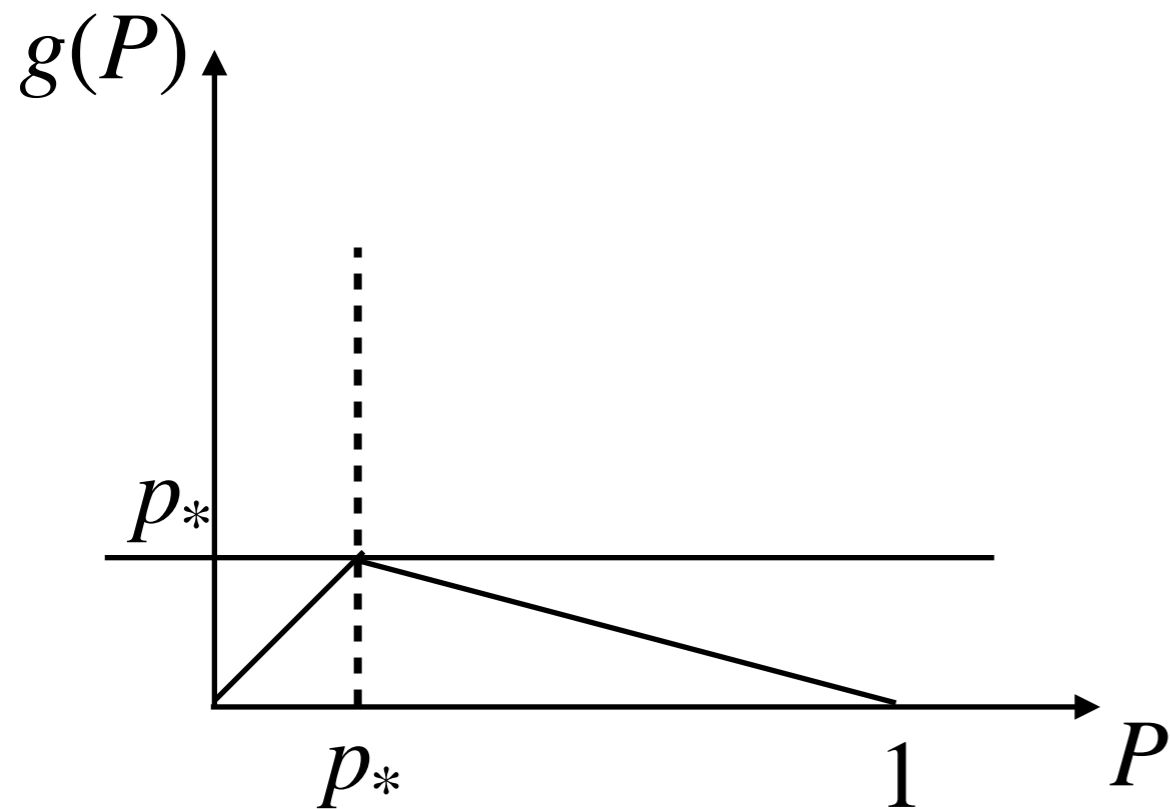
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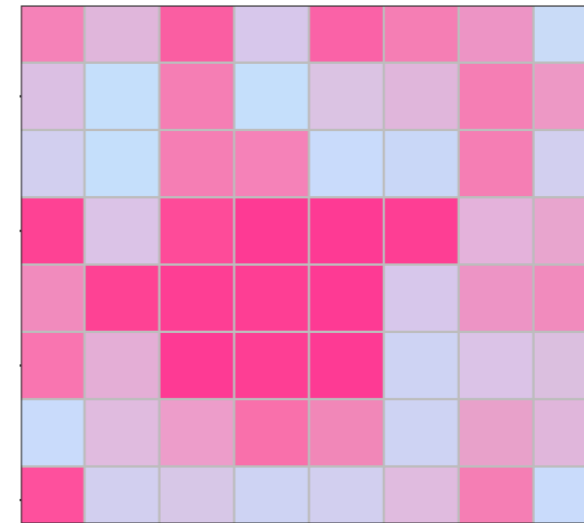
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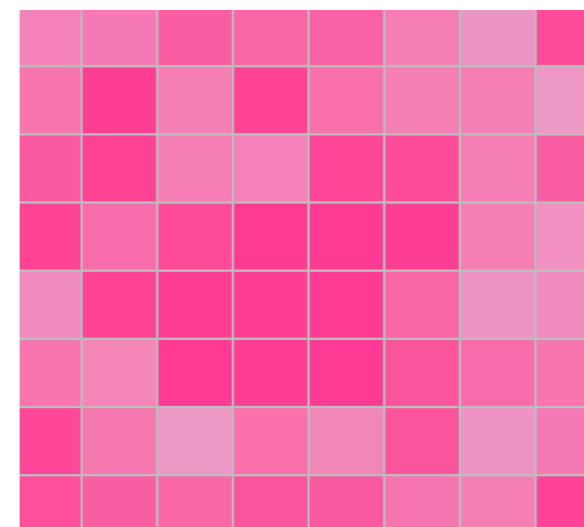
Small $g(P)$ indicate non-nulls



(Default $p_* = \alpha/2$)



P



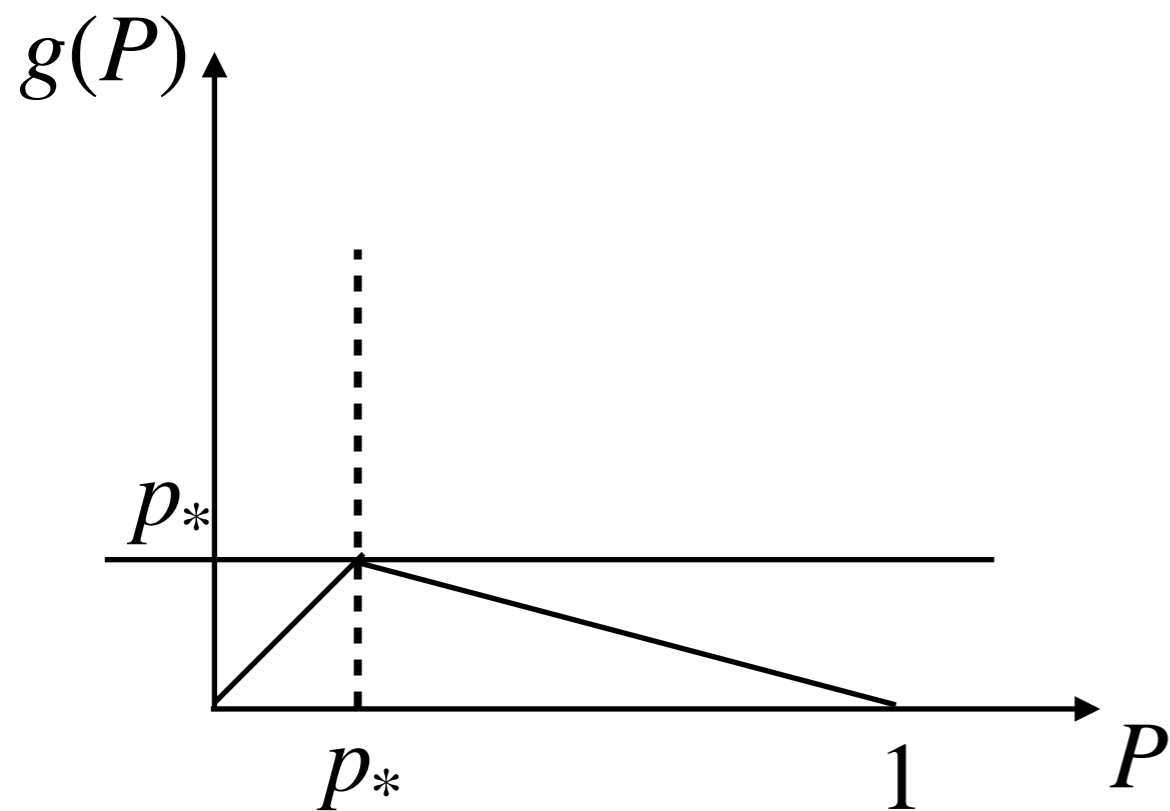
$g(P)$

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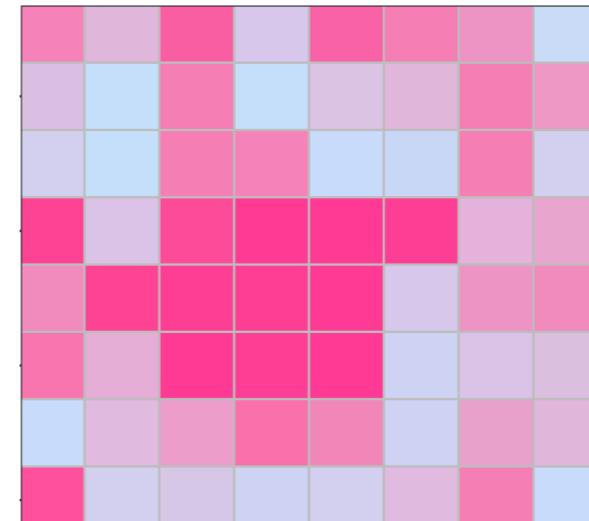
$$P \longrightarrow \begin{cases} g(P; p_*) = \min\left\{P, \frac{p_*}{1-p_*}(1-P)\right\} \\ h(P; p_*) = 2 \cdot 1\{P < p_*\} - 1 \end{cases}$$

Independent
for nulls

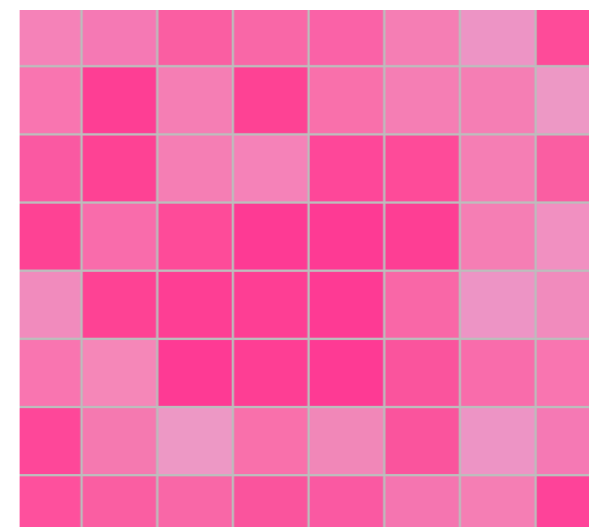
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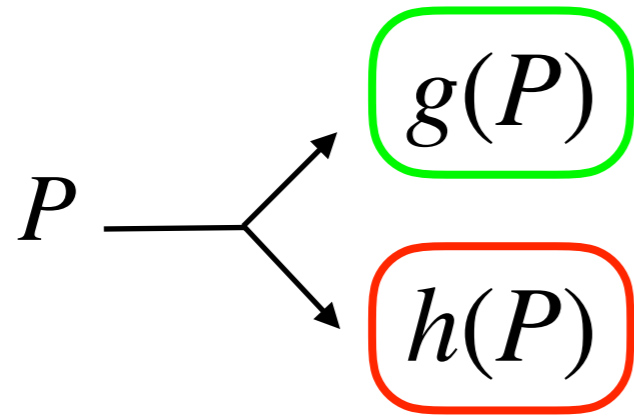


P



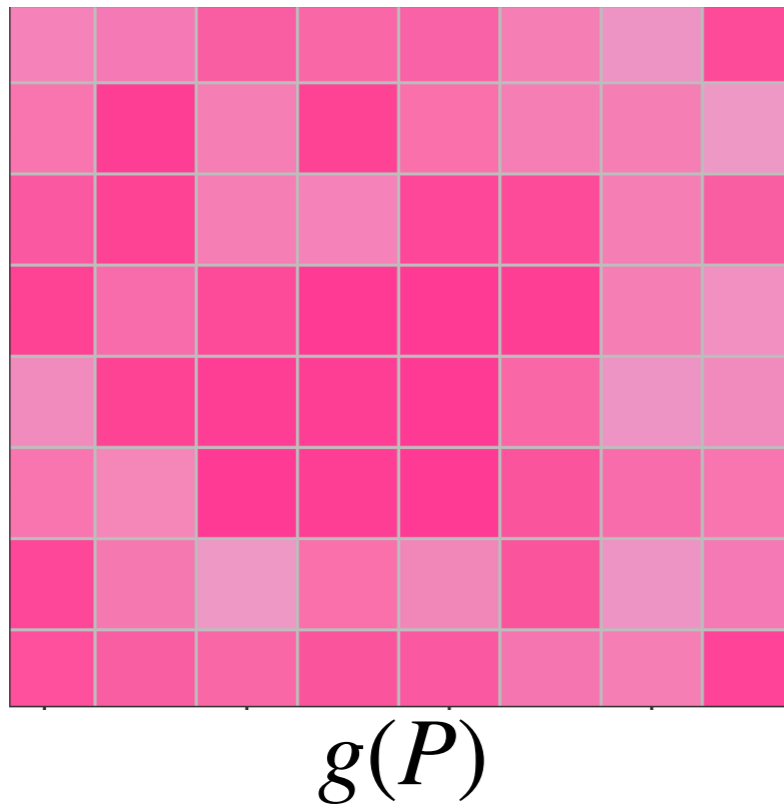
$g(P)$

Component 2: select candidate set **interactively**

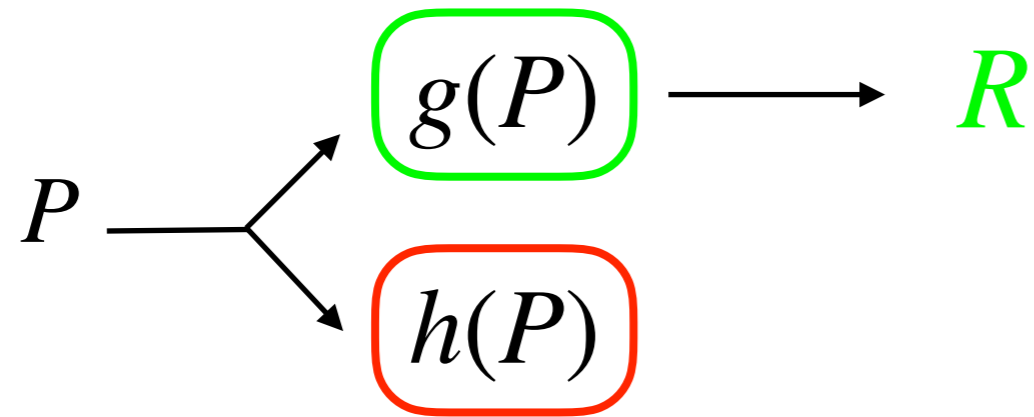


for cand. set selection

for error control

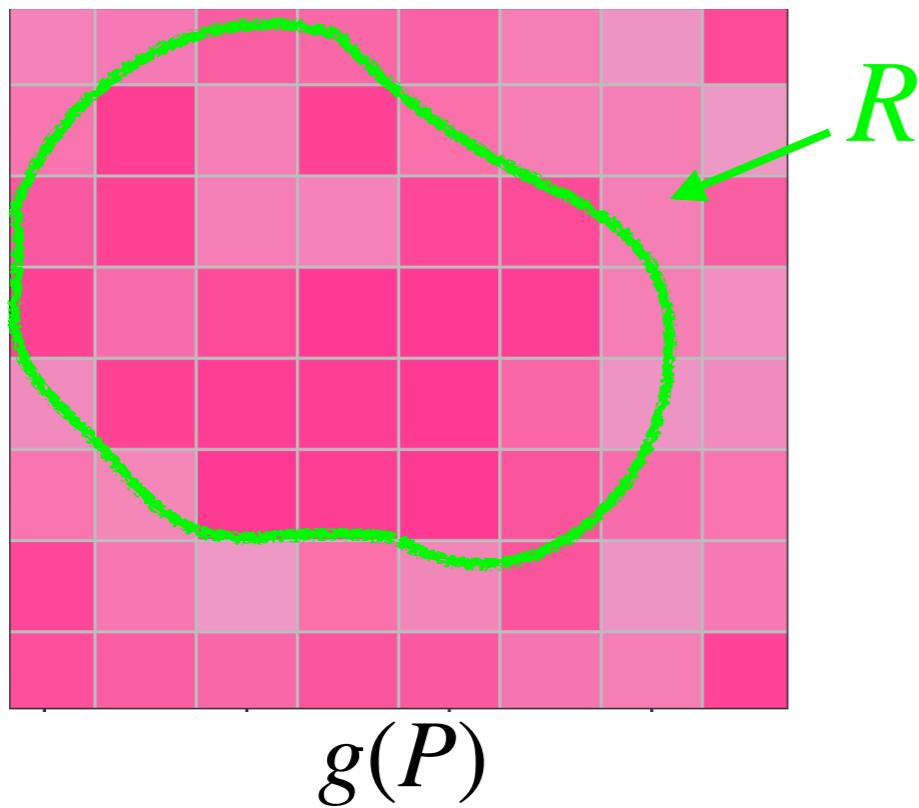


Component 2: select candidate set **interactively**

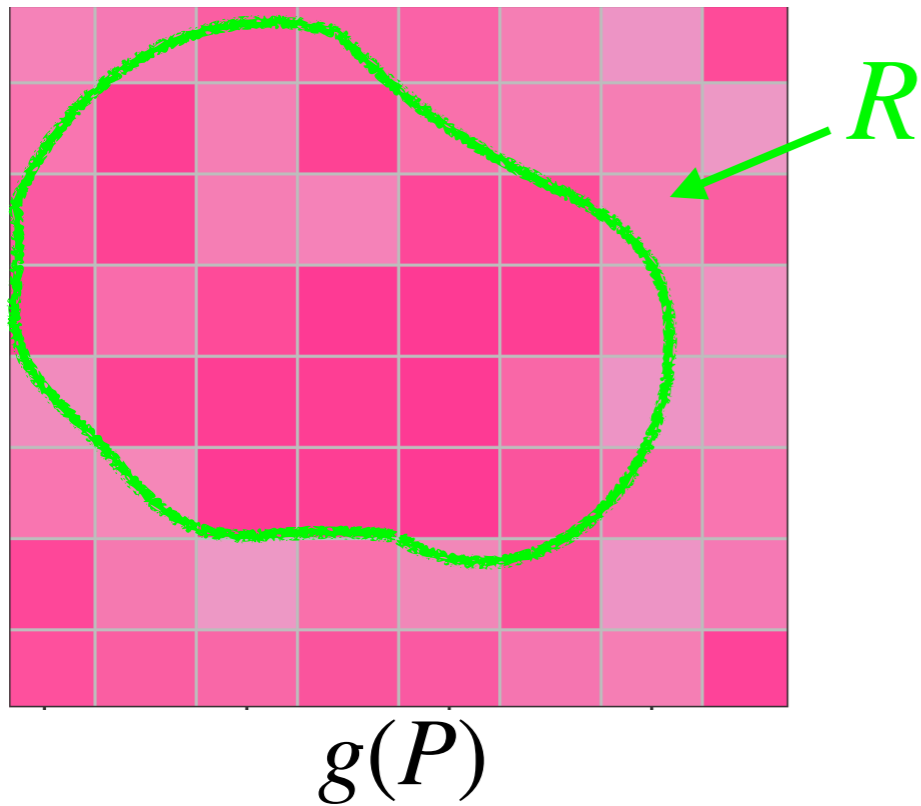
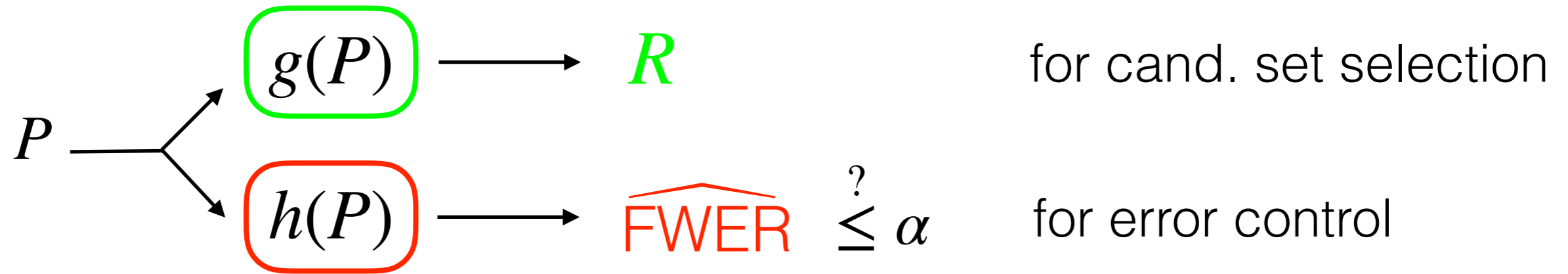


for cand. set selection

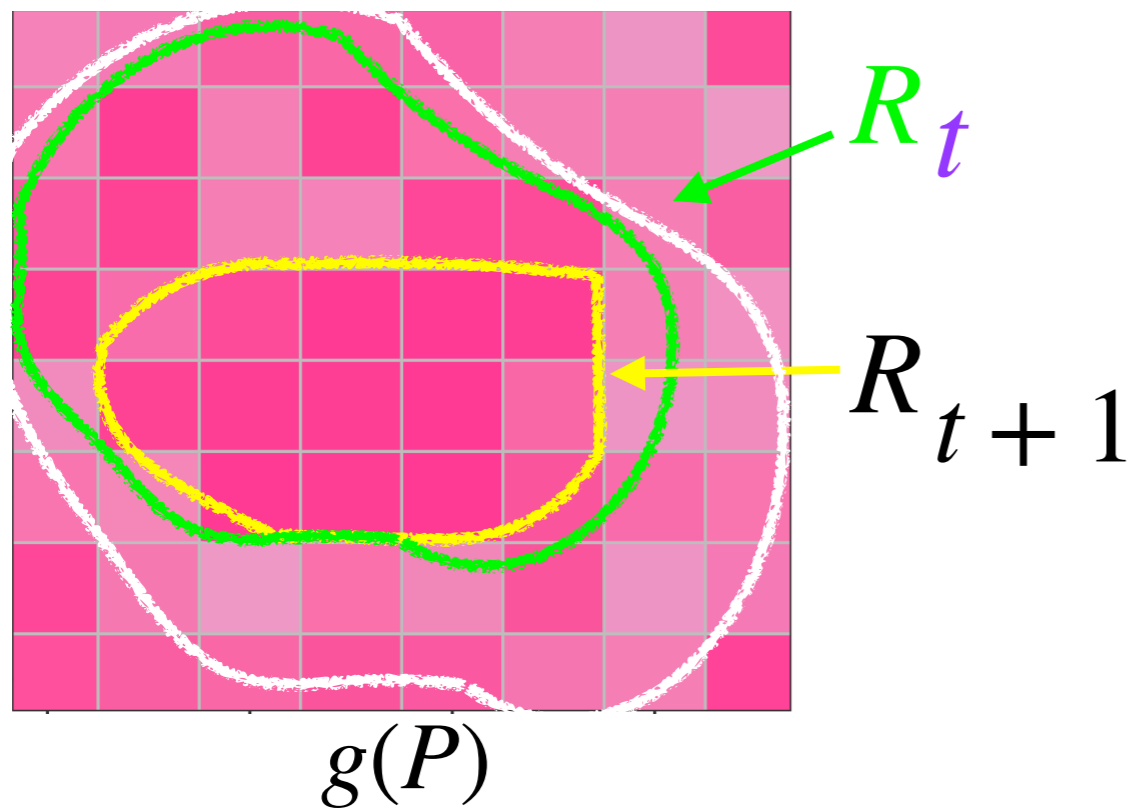
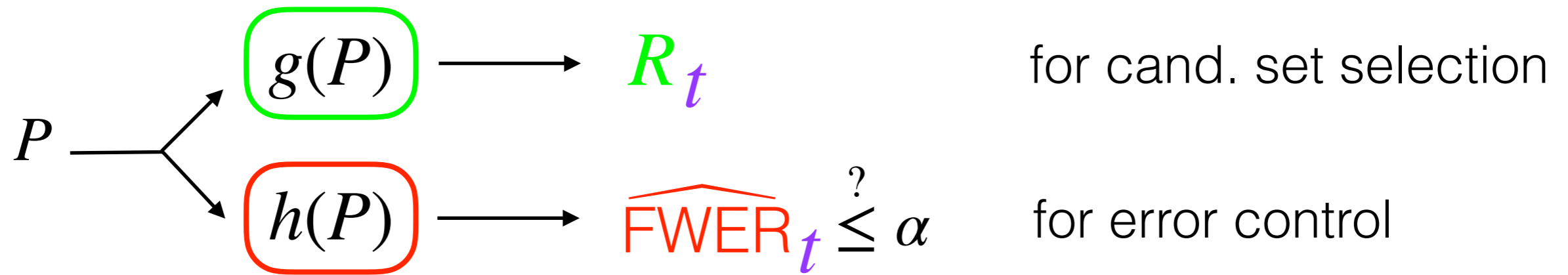
for error control



Component 2: select candidate set **interactively**



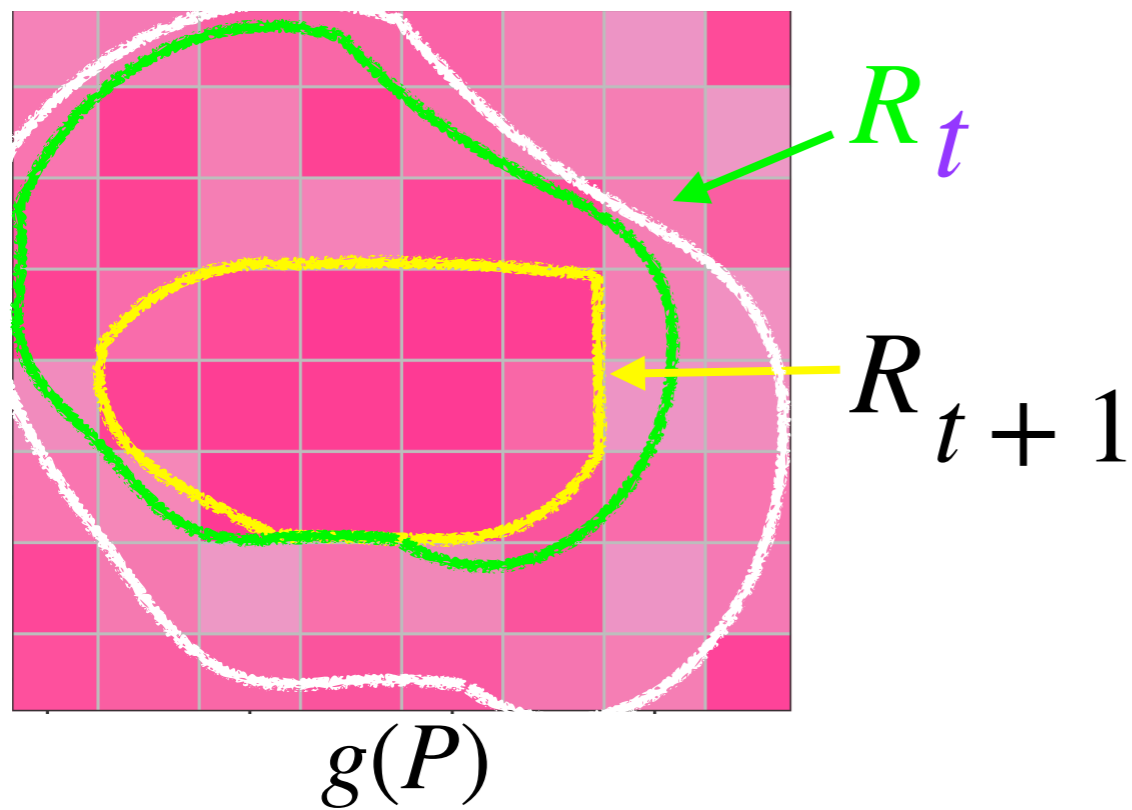
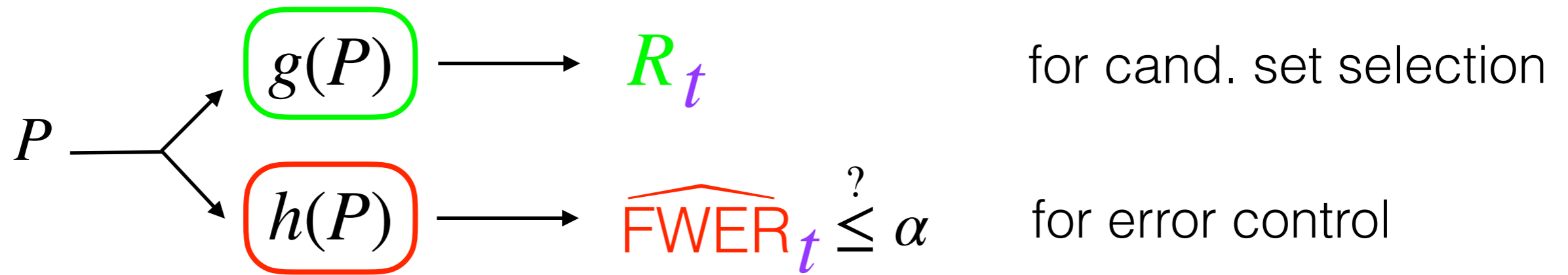
Component 2: select candidate set **interactively**



$$R_0 = \{1, \dots, n\}$$
$$R_0 \supseteq R_1 \supseteq \dots$$

progressively shrink R_t

Component 2: select candidate set **interactively**

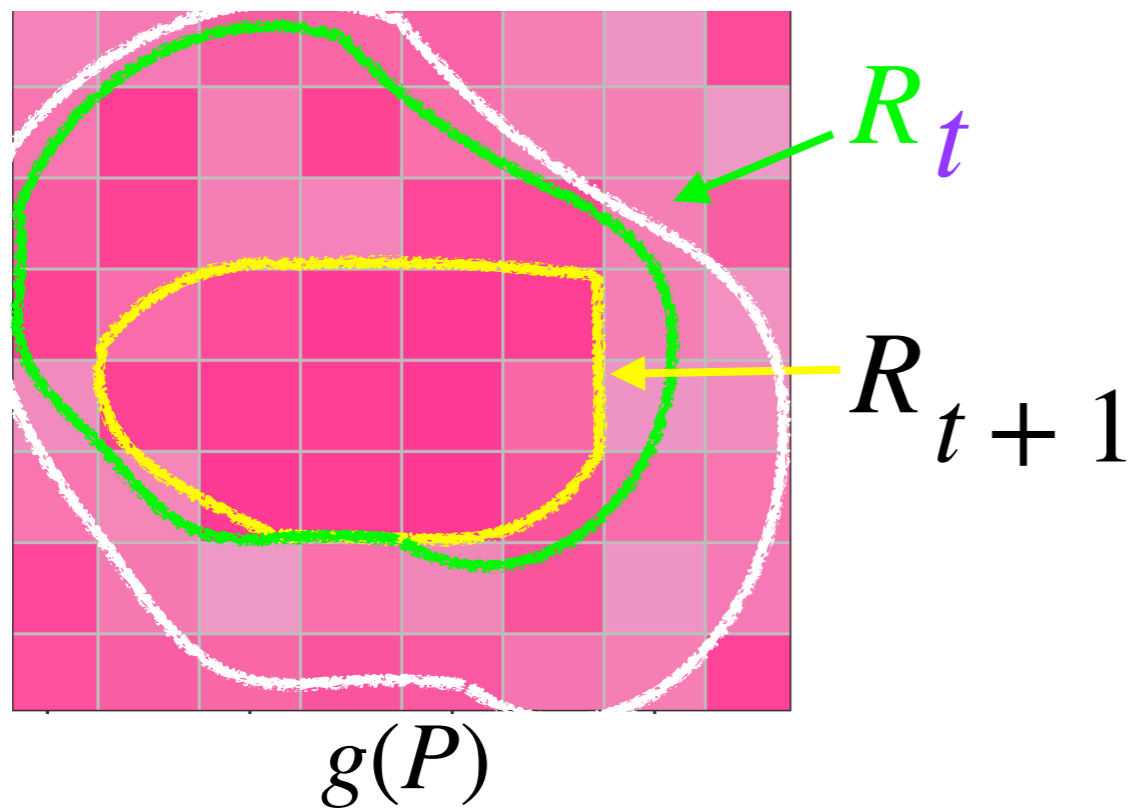
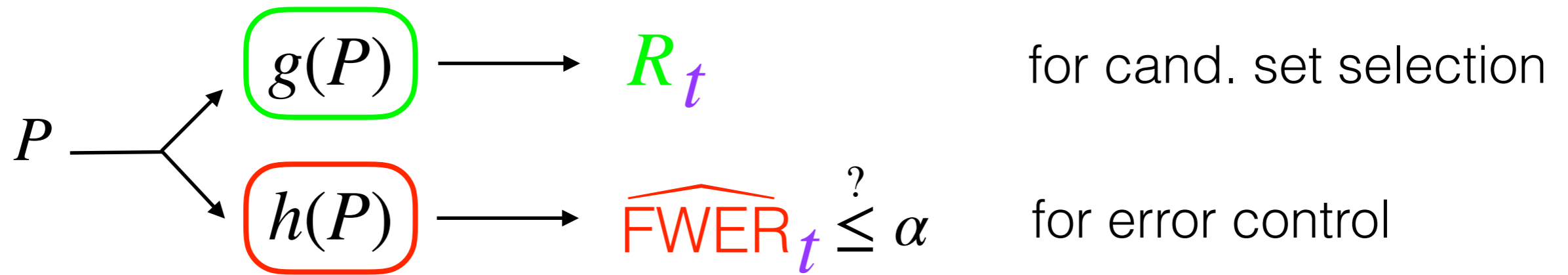


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progressively shrink R_t

$$\{g(P_i)\}_{i=1}^n$$

Component 2: select candidate set **interactively**

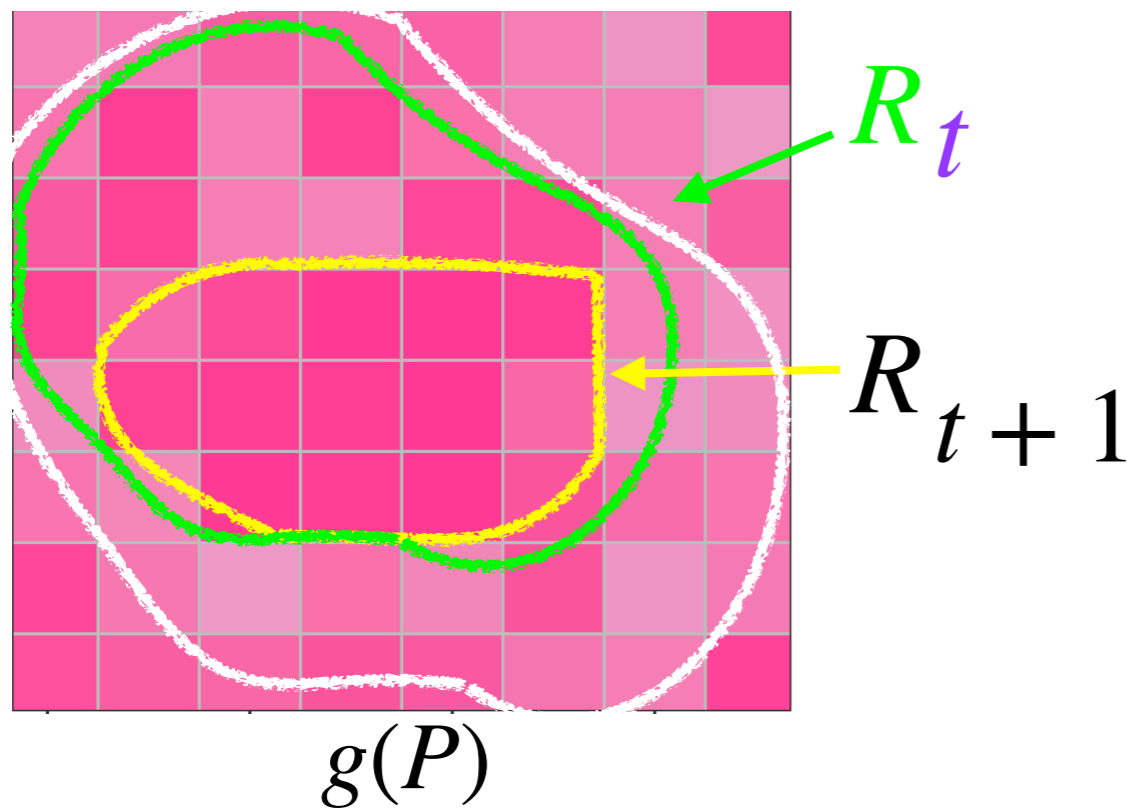
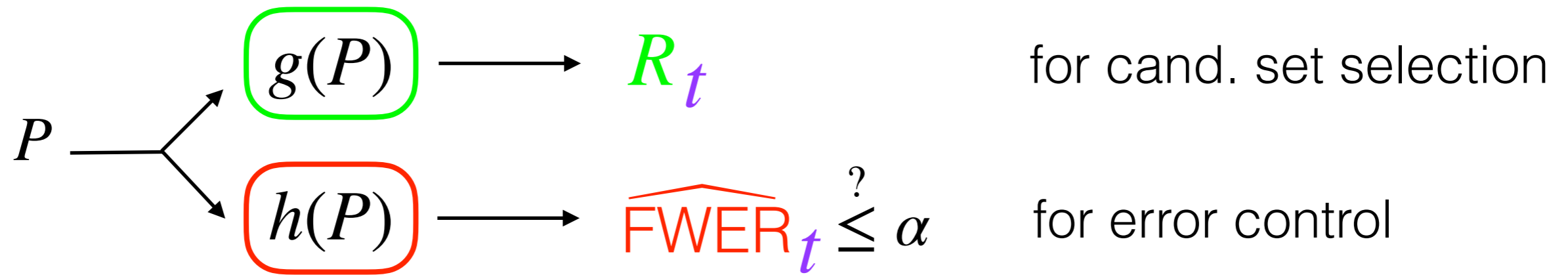


$$R_0 = \{1, \dots, n\}$$
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progressively shrink R_t

$\{g(P_i)\}_{i=1}^n$
+ coordinates

Component 2: select candidate set **interactively**

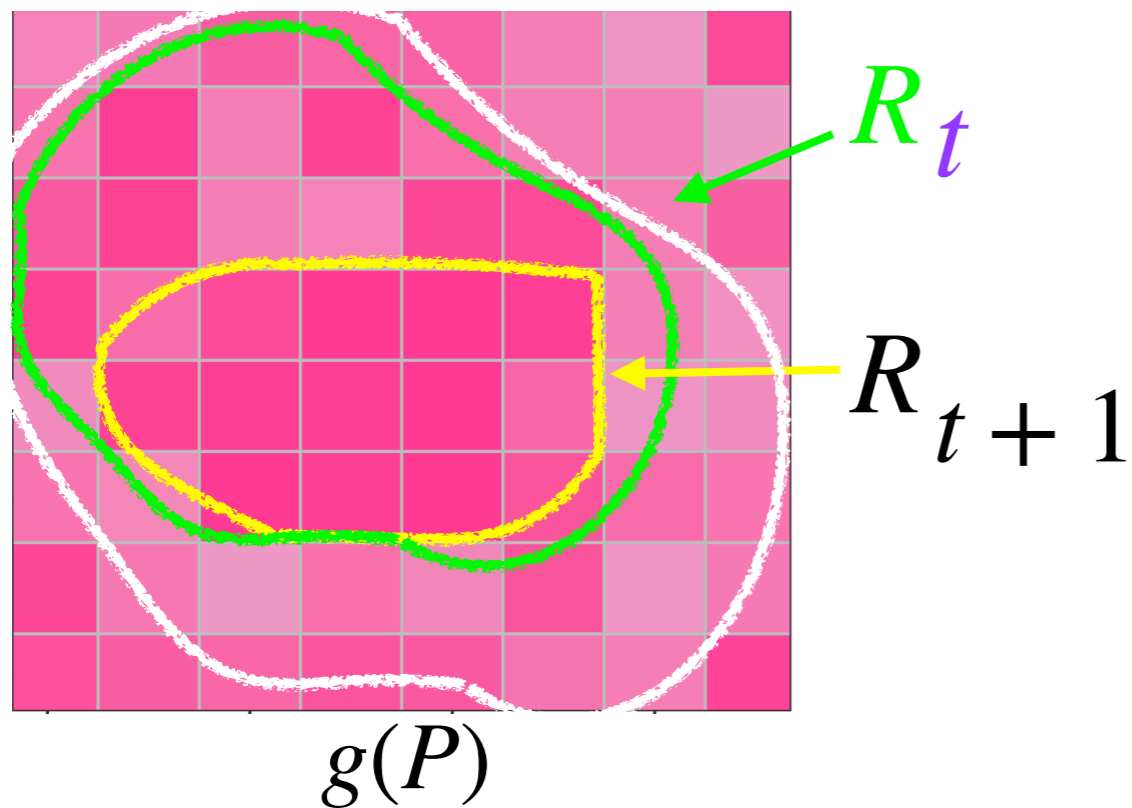
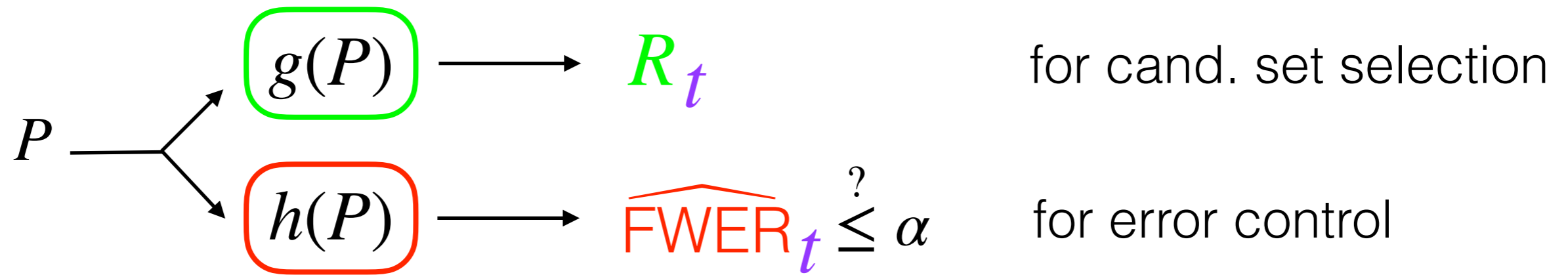


$$R_0 = \{1, \dots, n\}$$
$$R_0 \supseteq R_1 \supseteq \dots$$

progressively shrink R_t

$\{g(P_i)\}_{i=1}^n$
+ coordinates
(side information $\{x_i\}_{i=1}^n$)

Component 2: select candidate set **interactively**



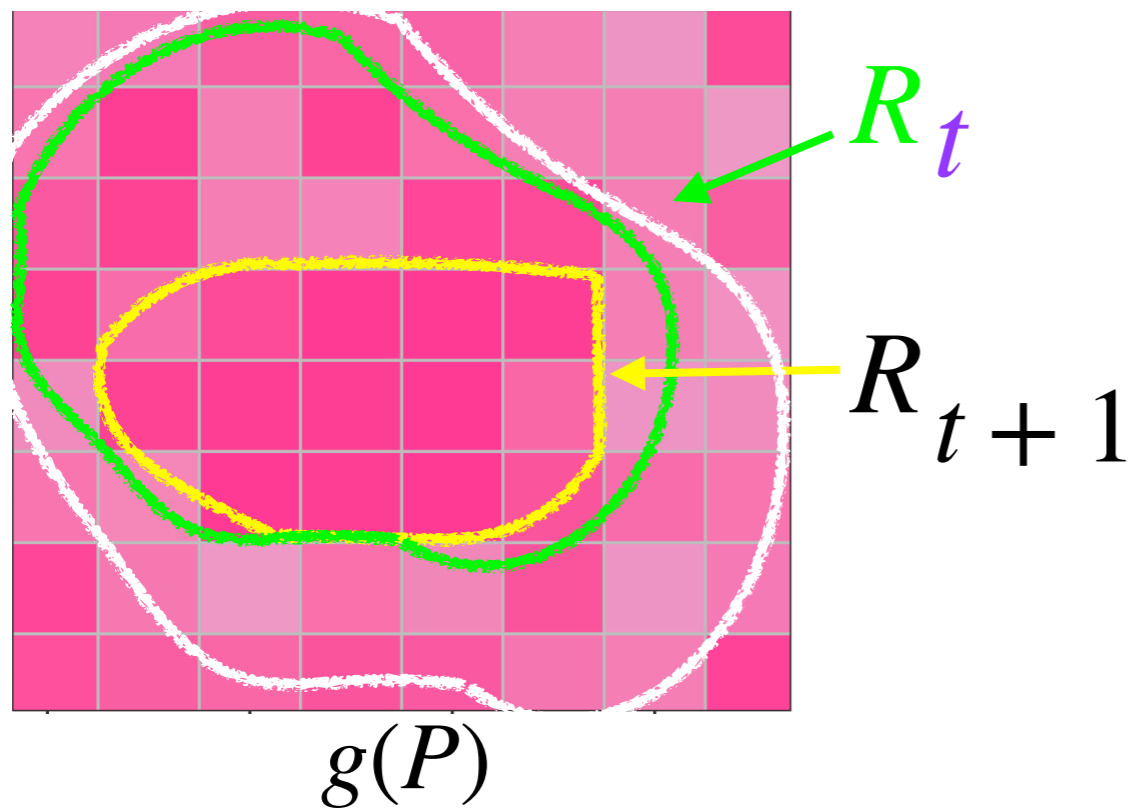
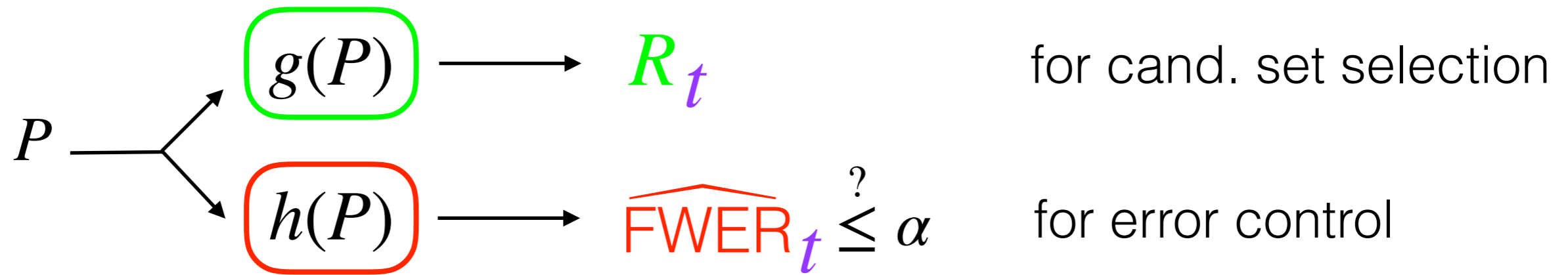
$$R_0 = \{1, \dots, n\}$$

$$R_0 \supseteq R_1 \supseteq \dots$$

progressively shrink R_t

$\{g(P_i)\}_{i=1}^n$
 + coordinates
 (side information $\{x_i\}_{i=1}^n$)
 + $\{h(P_i)\}_{i \notin R_t}$

Component 2: select candidate set **interactively**



$$R_0 = \{1, \dots, n\}$$

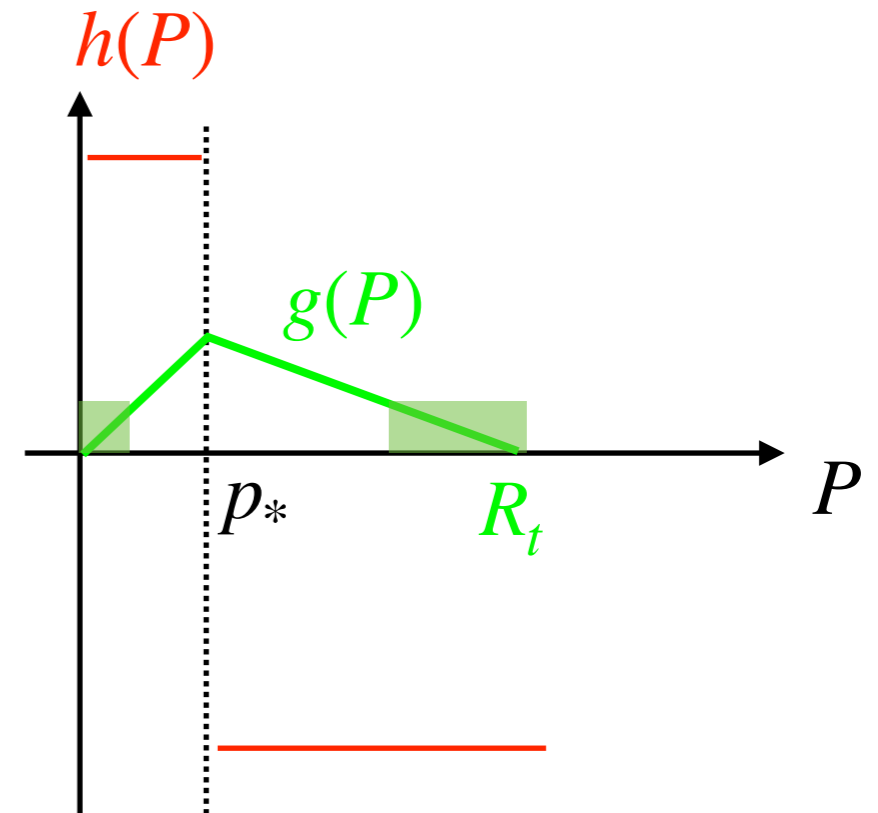
$$R_0 \supseteq R_1 \supseteq \dots$$

progressively shrink R_t
using increasing information:

$$\begin{aligned}
 & \{g(P_i)\}_{i=1}^n \\
 & + \text{coordinates} \\
 & (\text{side information } \{x_i\}_{i=1}^n) \\
 & + \{h(P_i)\}_{i \notin R_t}
 \end{aligned}$$

Component 3: error control

$$\text{FWER} := \mathbb{P}(\#\text{false rejections} \geq 1)$$

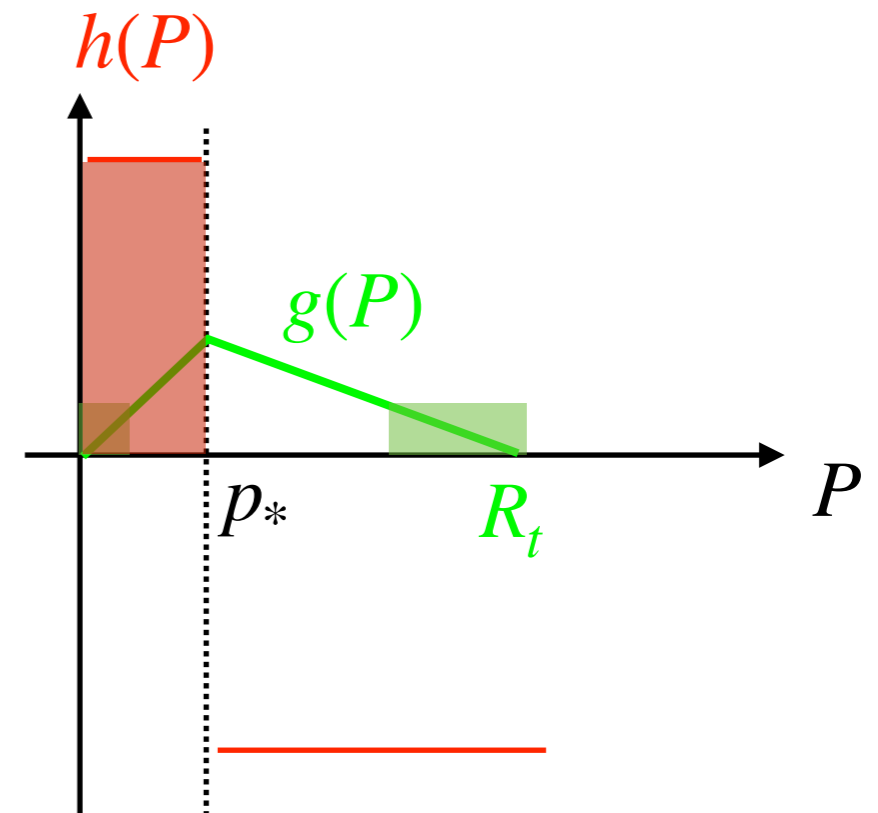


Component 3: error control

$$\text{FWER} := \mathbb{P}(\#\text{false rejections} \geq 1)$$

Reject the set

$$\{H_i \in R_t : h(P_i) = 1\}.$$



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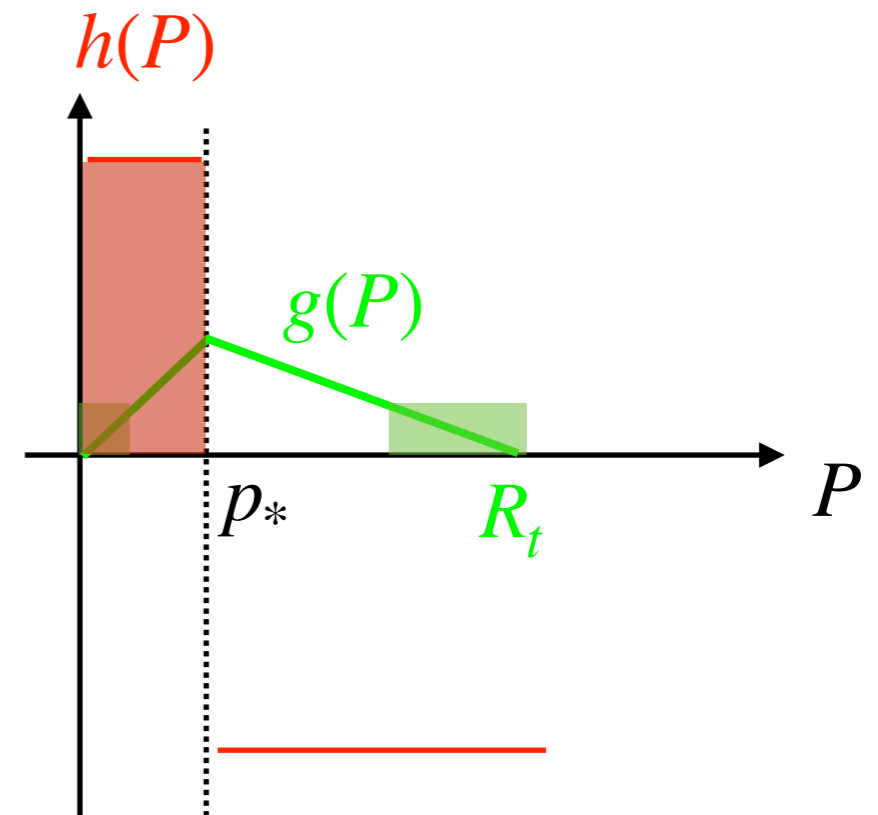
Reject the set

$$\{H_i \in R_t : h(P_i) = 1\}.$$

Count

$$R_t^- := |\{H_i \in R_t : h(P_i) = -1\}|$$

$h(P_i) \sim \text{Ber}(p_*)$ if H_i is null



FWER control using binary var.:
Janson & Su (2016)

Component 3: error control

$$\text{FWER} := \mathbb{P}(\#\text{false rejections} \geq 1)$$

Reject the set

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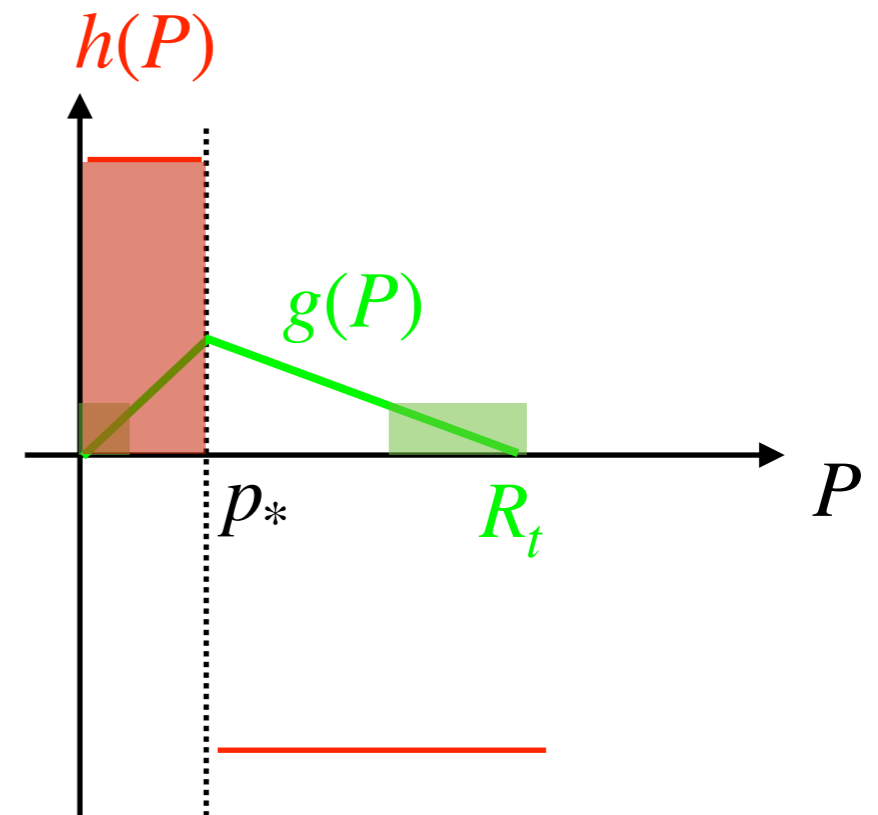
Count

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$h(P_i) \sim \text{Ber}(p_*)$ if H_i is null

Stop shrinking when

$$\widehat{\text{FWER}}_t := 1 - (1 - p_*)^{R_t^- + 1} \leq \alpha.$$



FWER control using binary var.:
Janson & Su (2016)

Component 3: error control

$$\text{FWER} := \mathbb{P}(\#\text{false rejections} \geq 1)$$

$$t = 0, \quad \#\text{rej} = 17, \quad \widehat{\text{FWER}}_t = 0.9998567$$

Reject the set

$$\{H_i \in R_t : h(P_i) = 1\}.$$

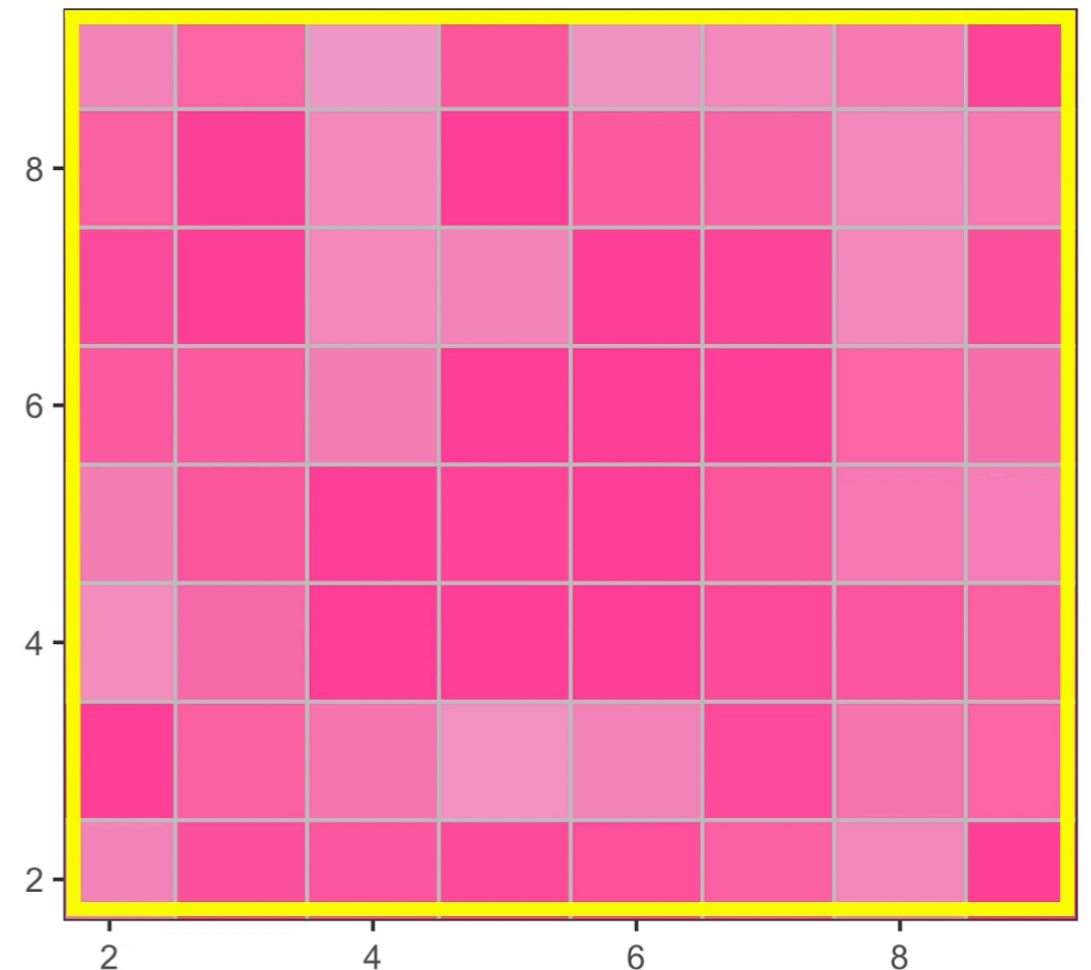
Count

$$R_t^- := |\{H_i \in R_t : h(P_i) = -1\}|$$

$$h(P_i) \sim \text{Ber}(p_*) \text{ if } H_i \text{ is null}$$

Stop shrinking when

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10 × 10 hypotheses with 9 non-nulls
($\alpha = 0.2$)

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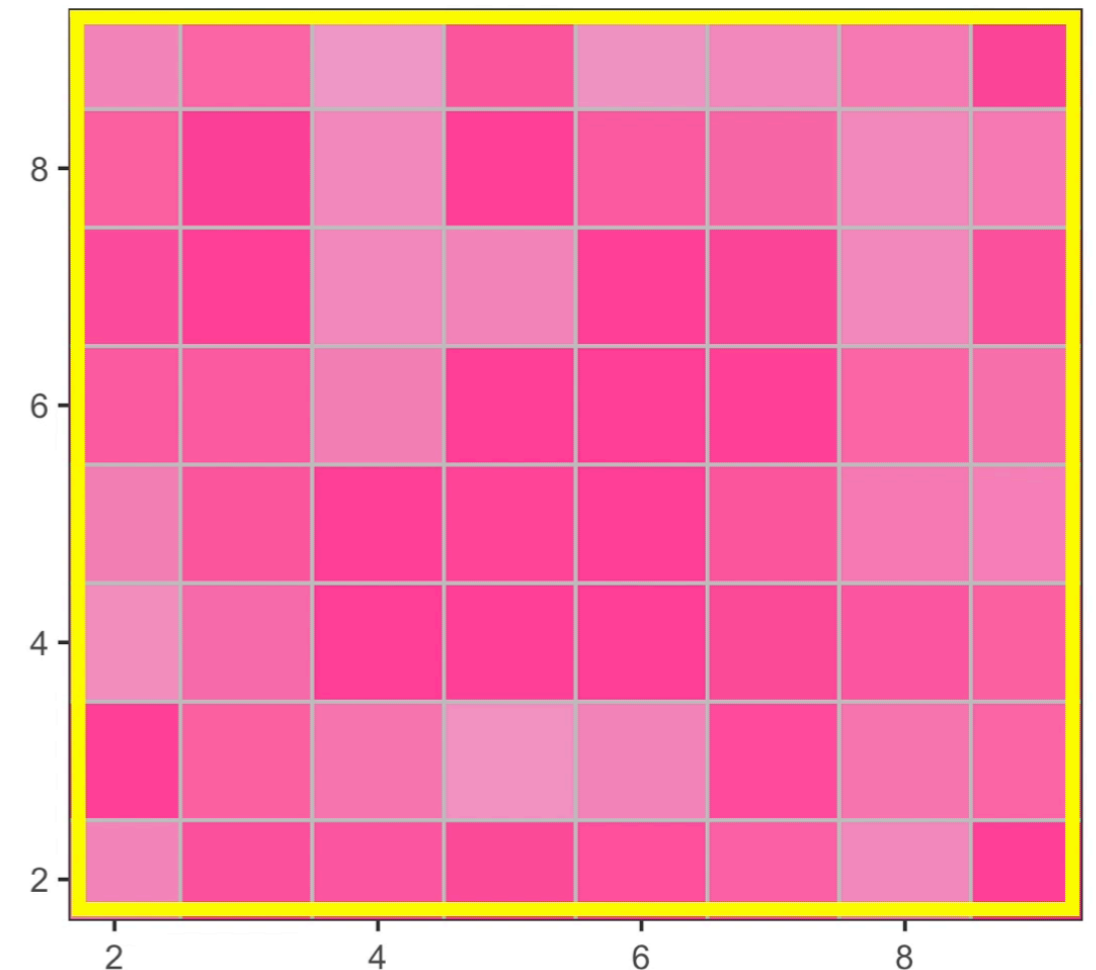
Count

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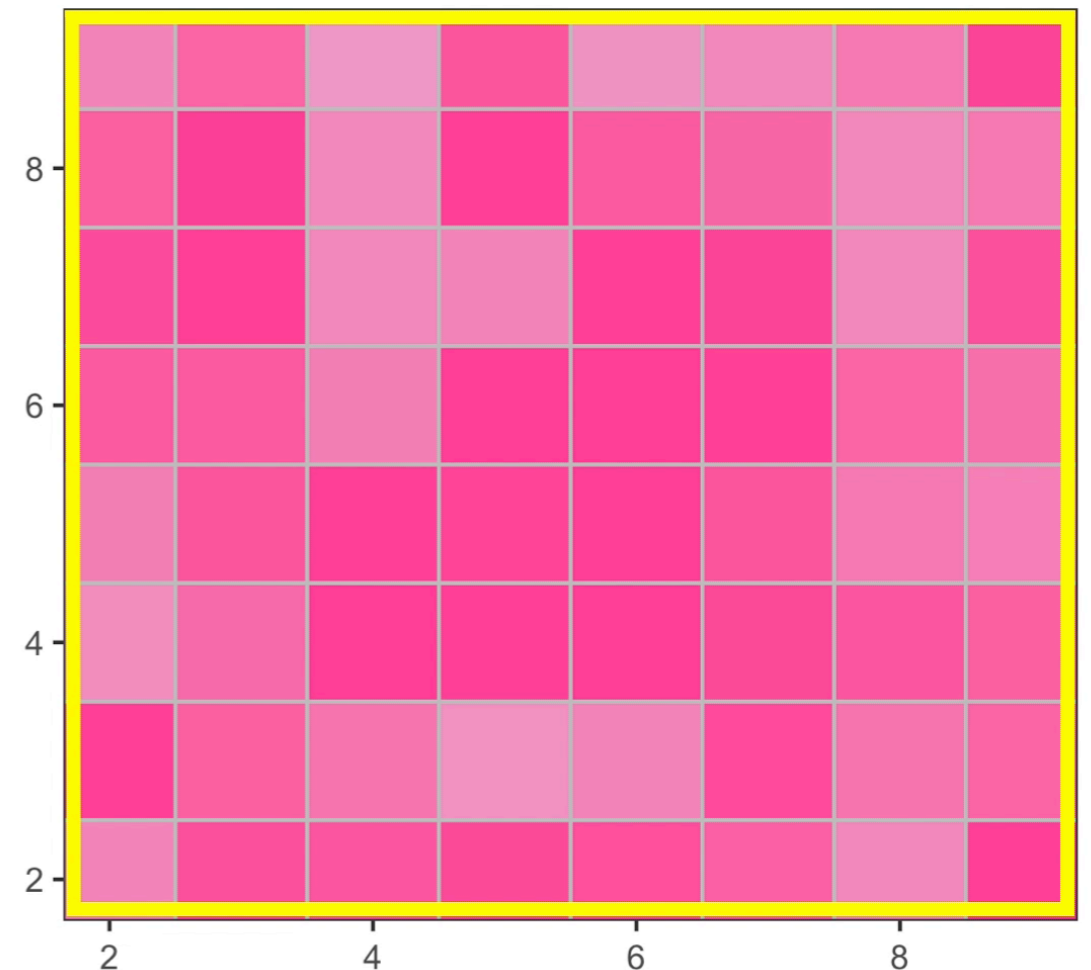
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Stop shrinking when

$$\widehat{\text{FWER}}_t := 1 - (1 - p_*)^{R_t^- + 1} \leq \alpha.$$



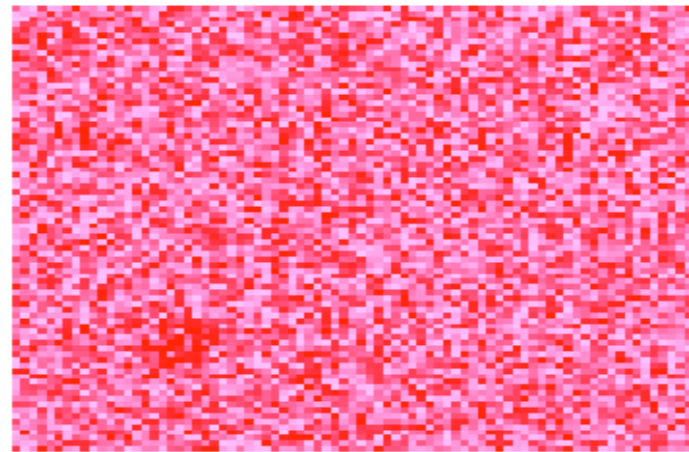
10 × 10 hypotheses with 9 non-nulls
($\alpha = 0.2$)

Theorem: the i-FWER method control FWER if null p -values are mutually independent, and are independently of the non-nulls.

FWER control using binary var.:
Janson & Su (2016)

Strategies to shrink R_t

- Estimate the probability of being non-null.



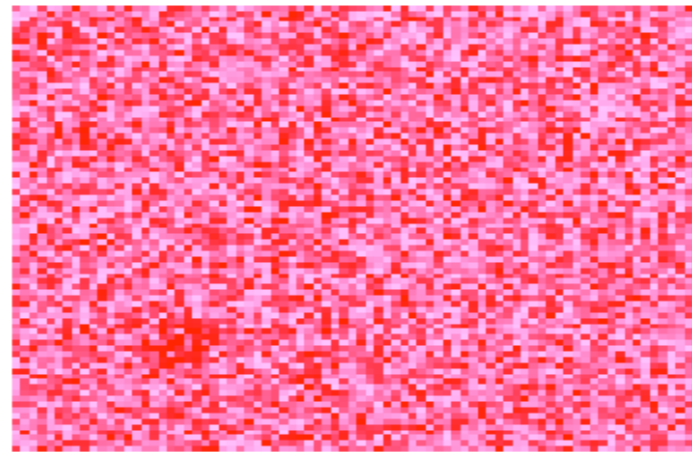
$g(P)$



smoothed
probability

Strategies to shrink R_t

- Estimate the probability of being non-null.

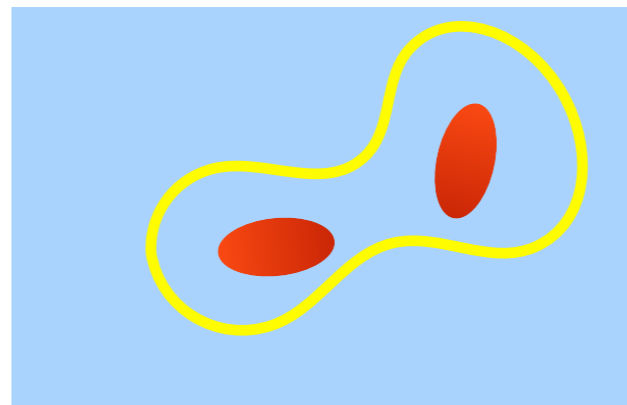


$g(P)$



smoothed
probability

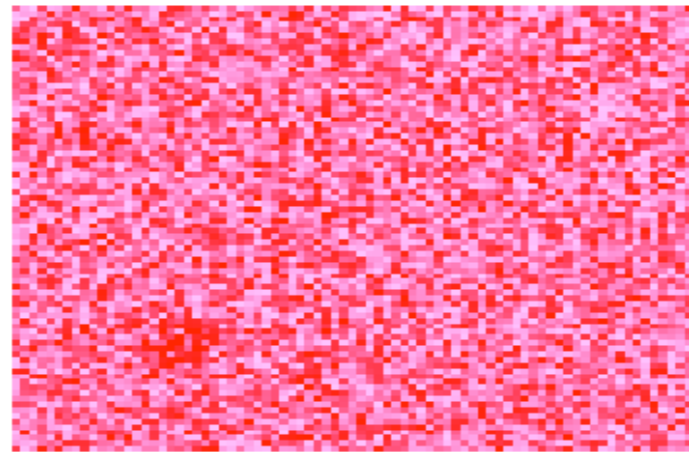
-
- Change the strategy at any step.



single center?

Strategies to shrink R_t

- Estimate the probability of being non-null.

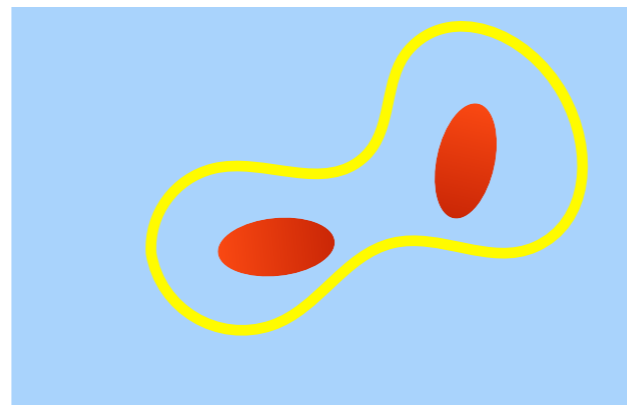


$g(P)$

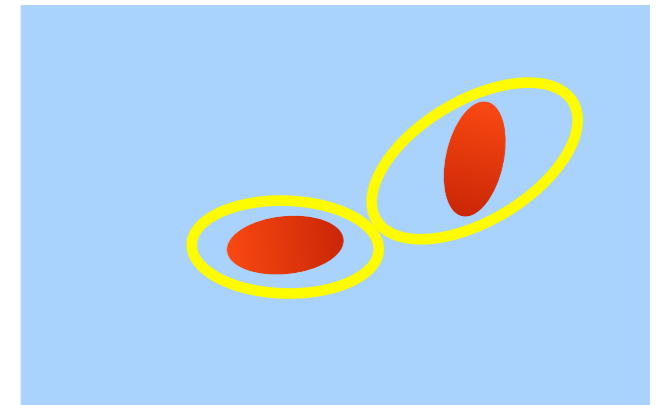


smoothed
probability

- Change the strategy at any step.



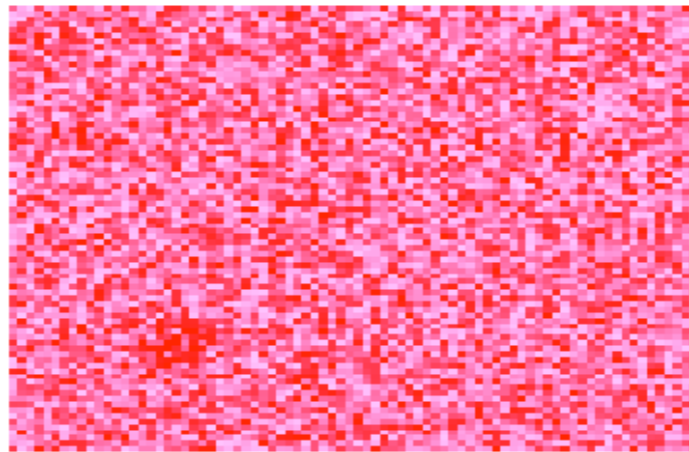
single center?



two centers

Strategies to shrink R_t

- Estimate the probability of being non-null.

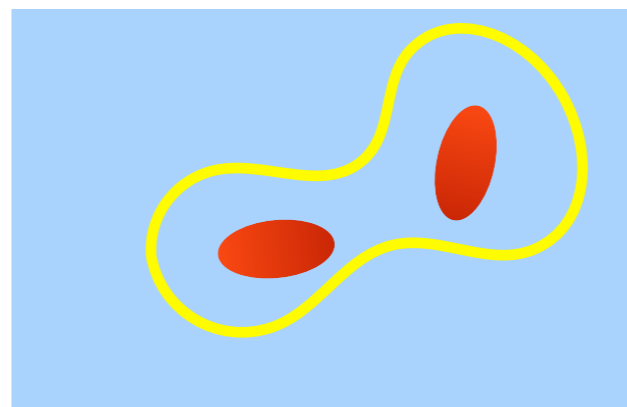


$g(P)$

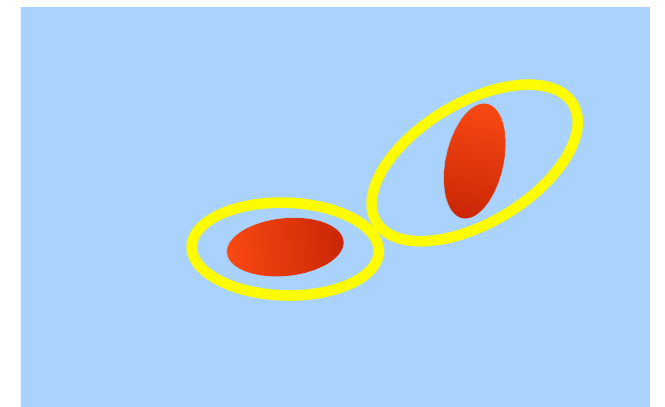


smoothed probability

- Change the strategy at any step.

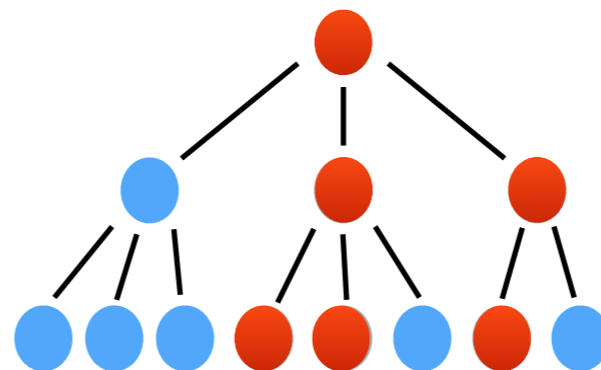


single center?



two centers

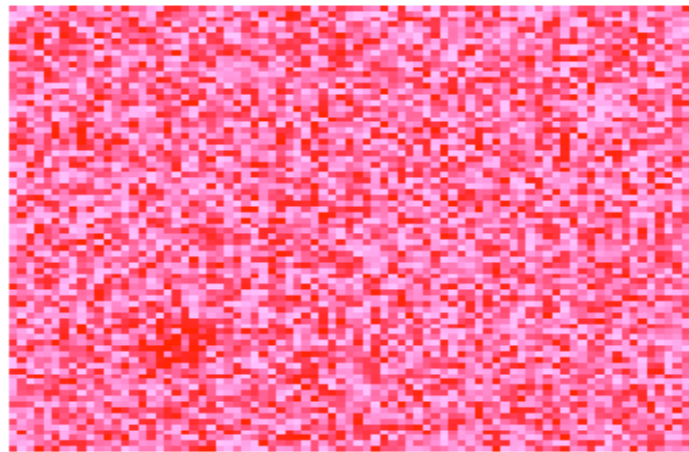
- Customize the strategy to different structures.



Hierarchical structure

Strategies to shrink R_t

- Estimate the probability of being non-null.

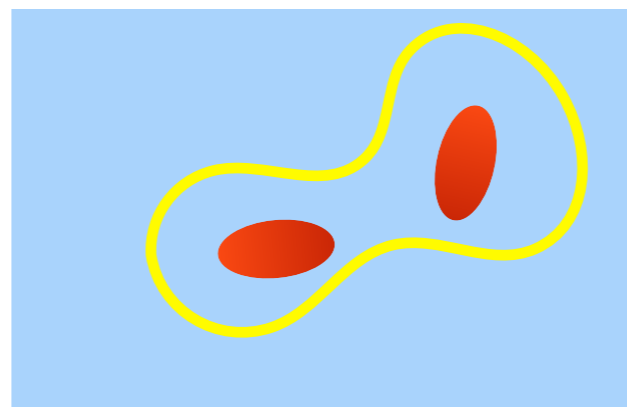


$g(P)$

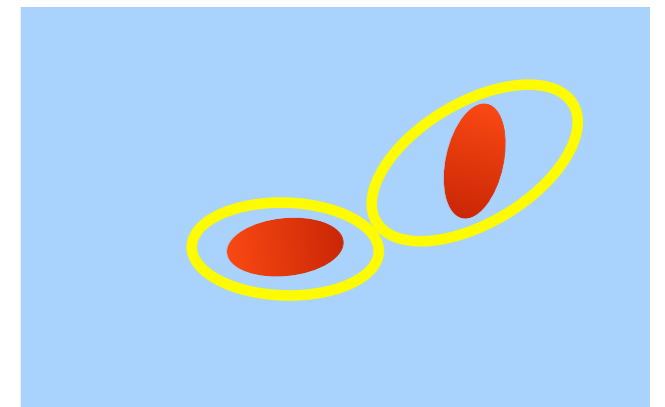


smoothed probability

- Change the strategy at any step.

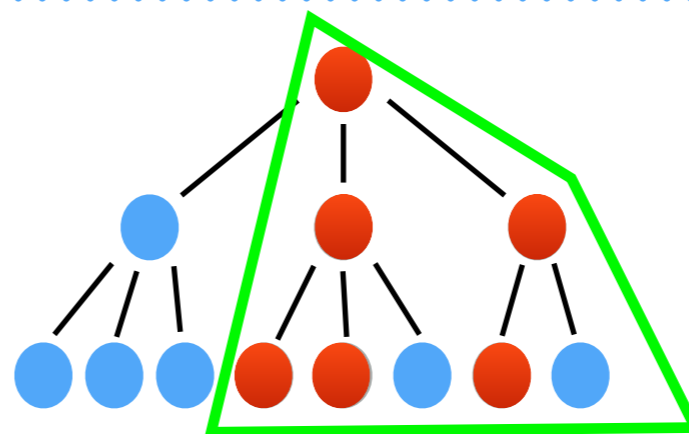


single center?



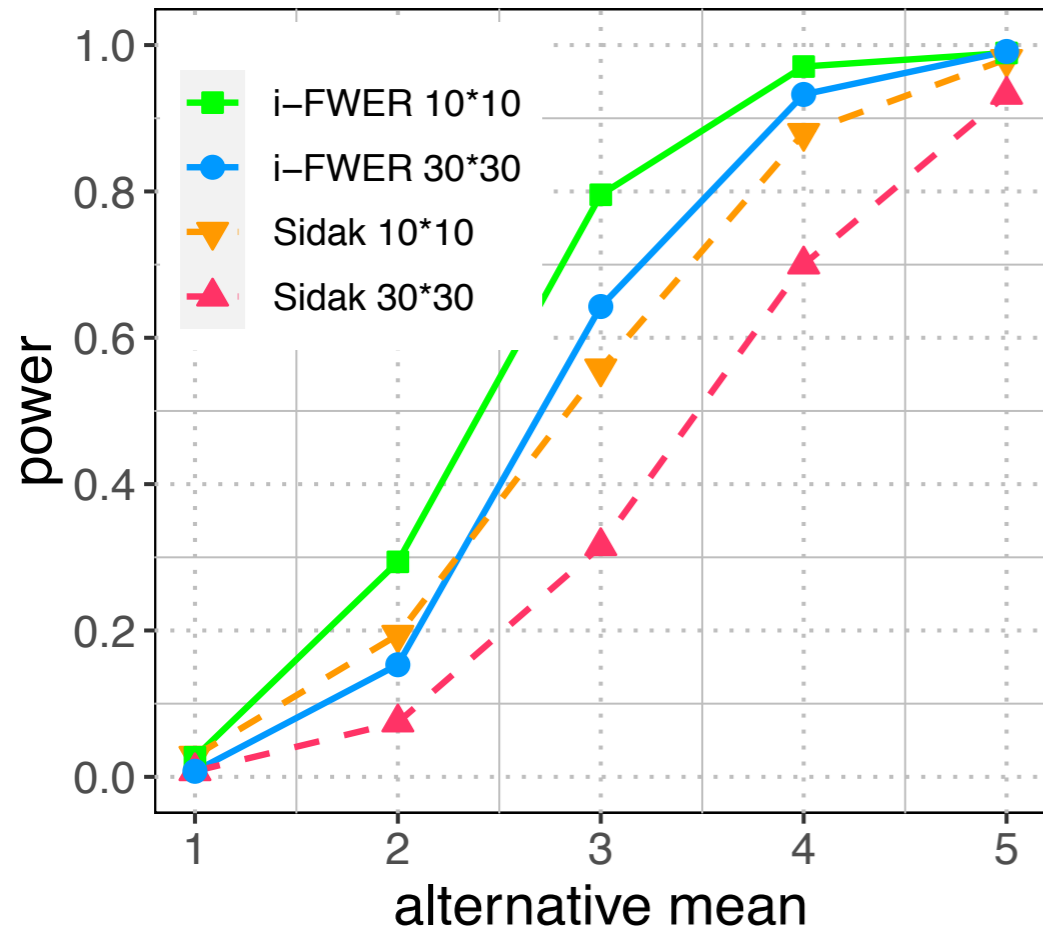
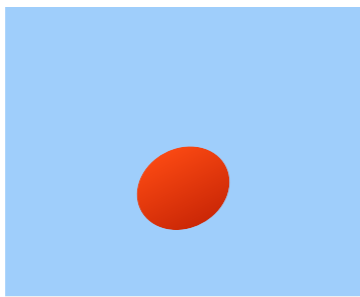
two centers

- Customize the strategy to different structures.

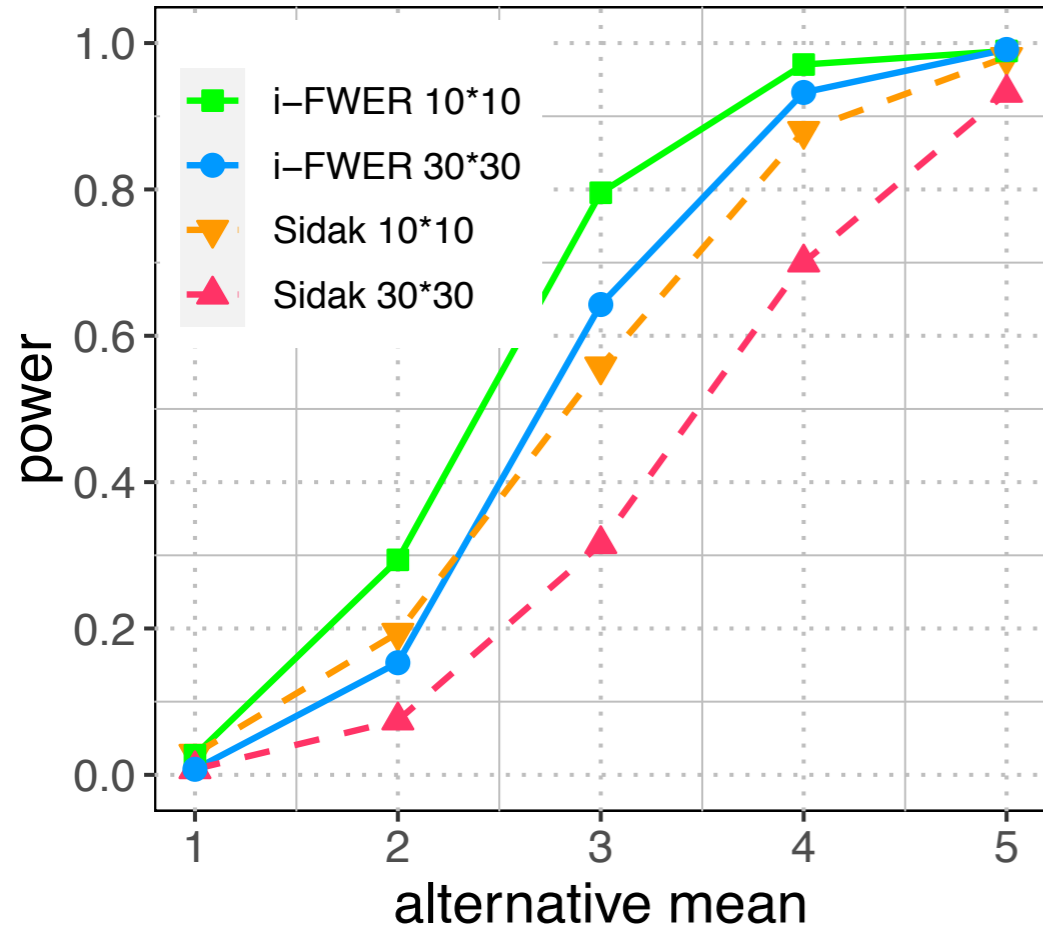
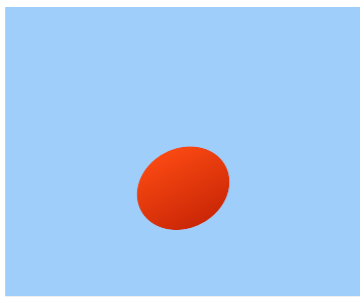


Hierarchical structure

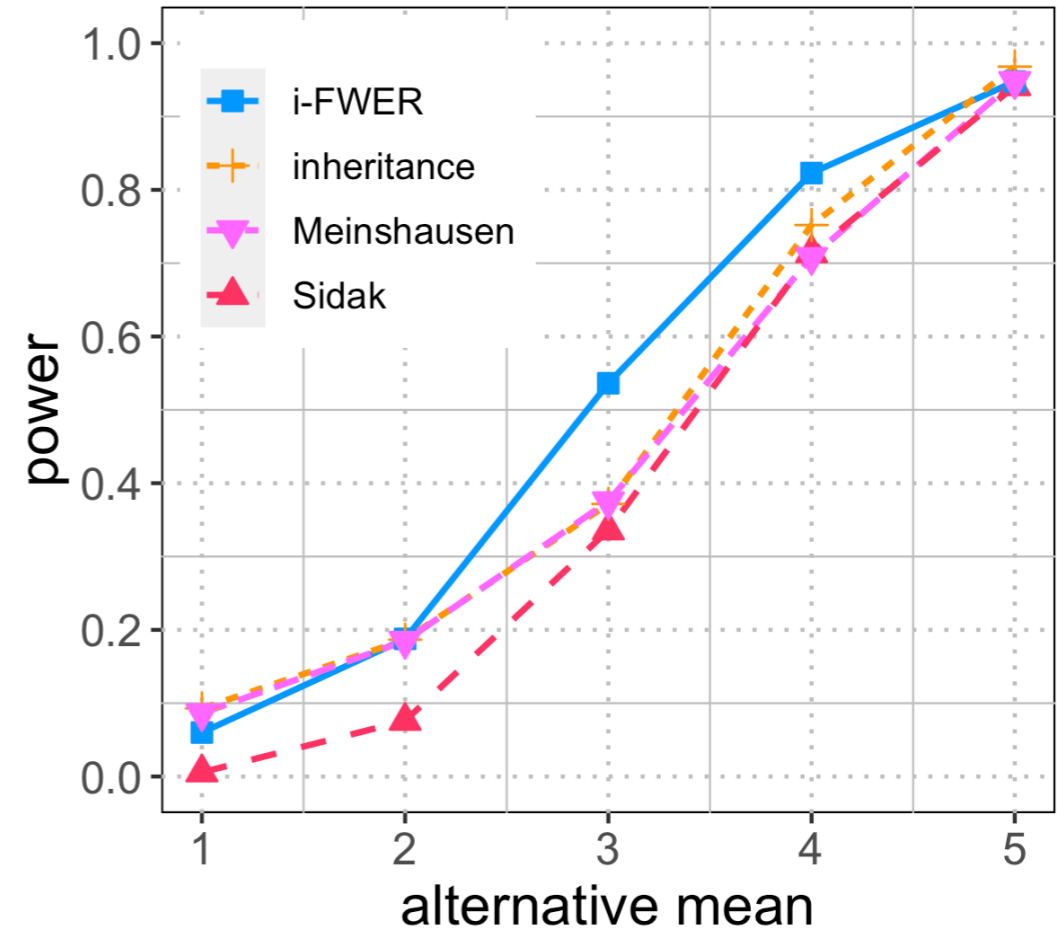
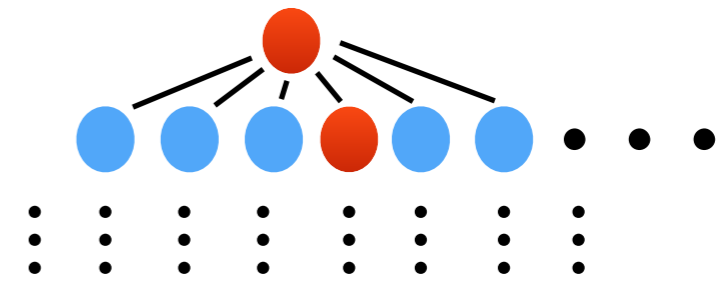
Grid



Grid

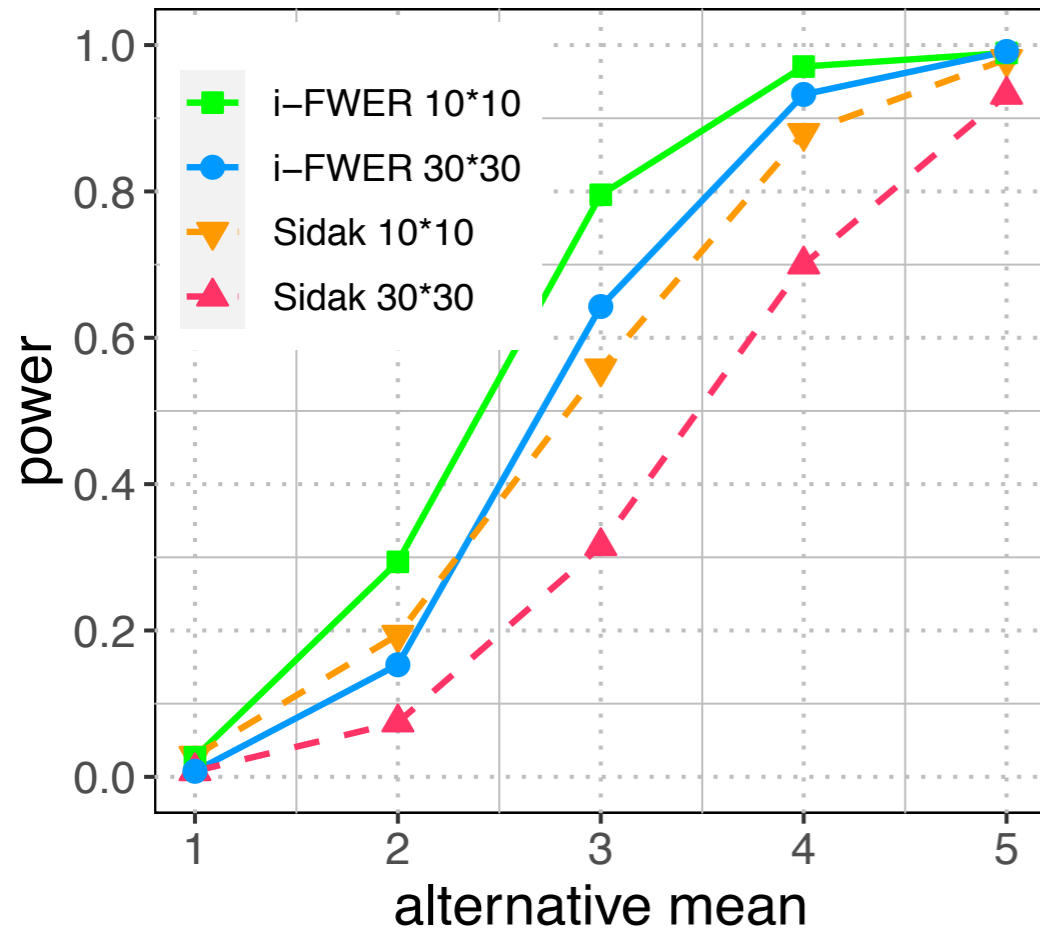
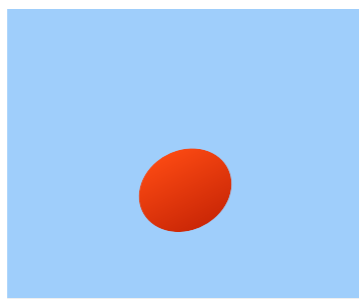


Tree

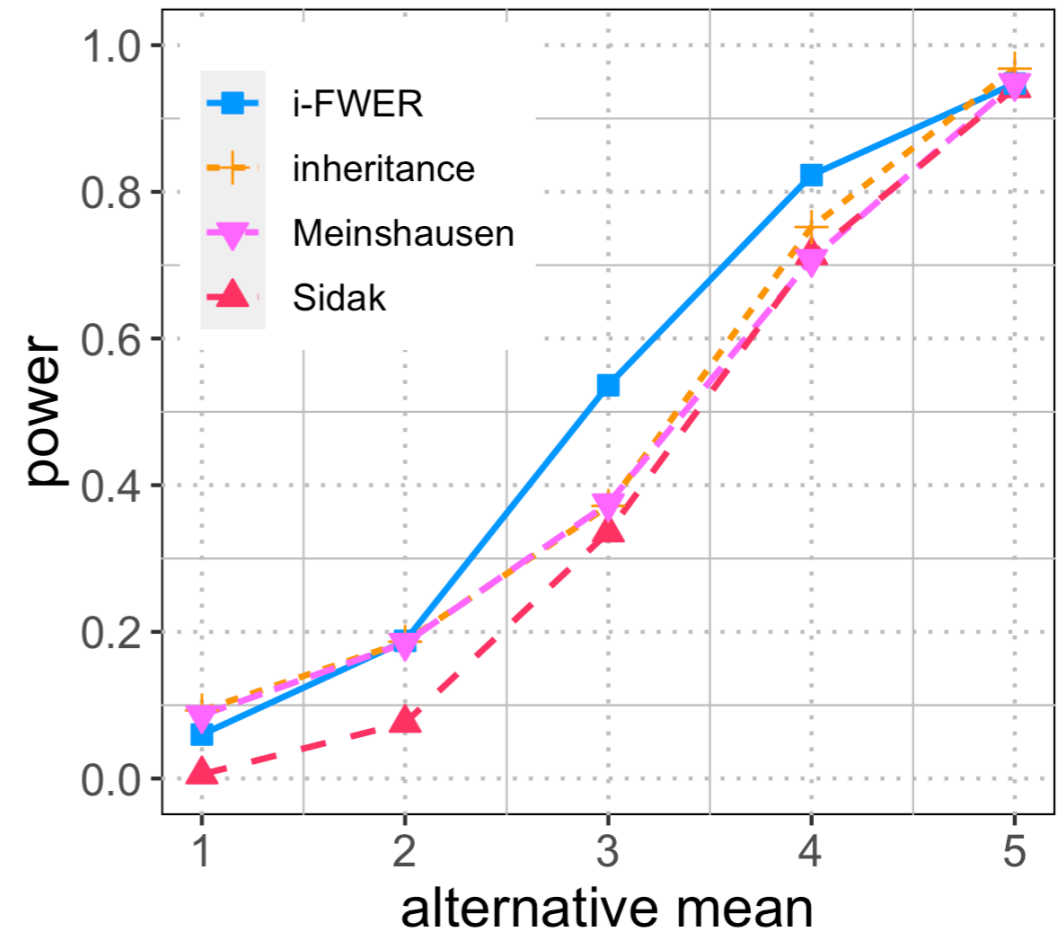
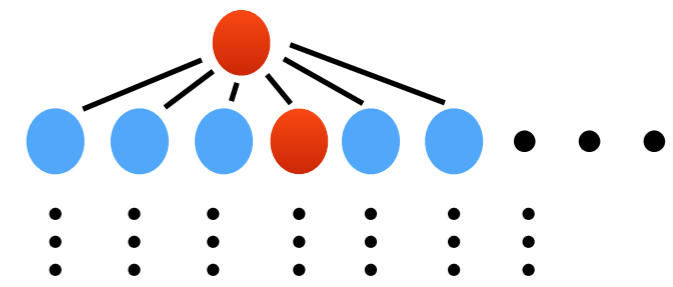


Meinshausen (2008); Goeman and Finos (2012);
Ignatiadis et al. (2016); Lei, Fithian (2018)

Grid



Tree



Real data: RNA sequence

| FWER level | IHW [★] | i-FWER |
|------------|------------------|--------|
| 0.1 | 1552 | 1613 |
| 0.2 | 1645 | 1740 |
| 0.3 | 1708 | 1844 |

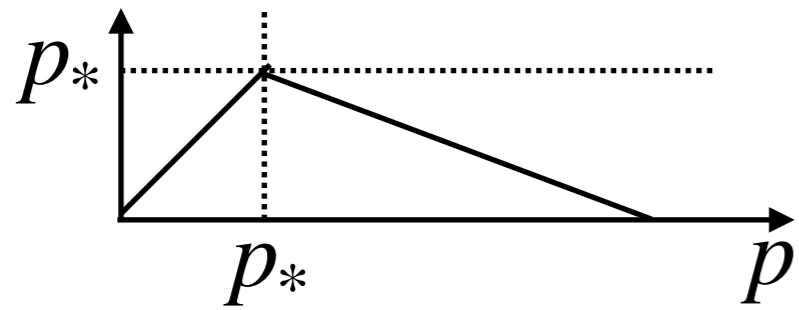
differentially expressed genes
in airway smooth muscle cell lines
in response to dexamethasone

Meinshausen (2008); Goeman and Finos (2012);
★Ignatiadis et al. (2016); Lei, Fithian (2018)

Extensions

Masking functions

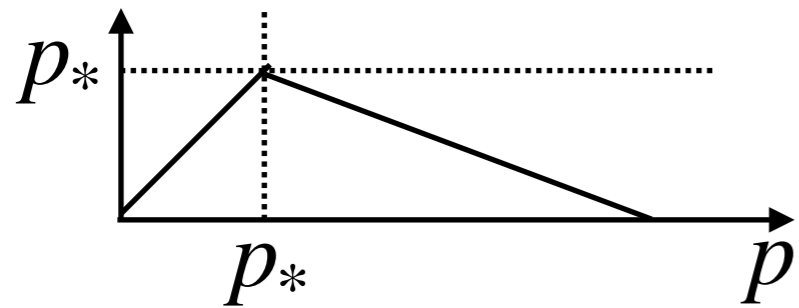
Tent function



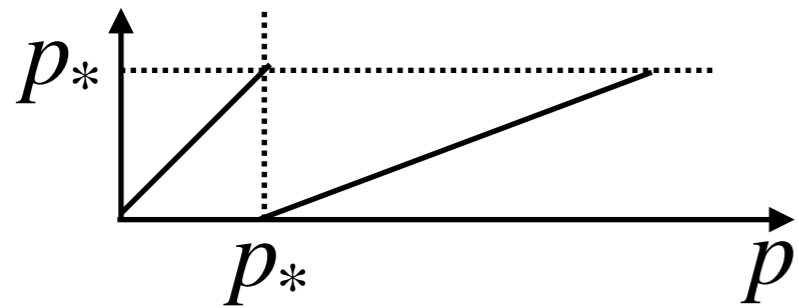
Extensions

Masking functions

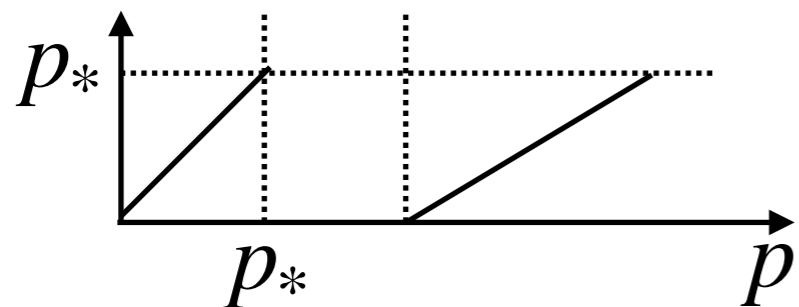
Tent function



Railway function



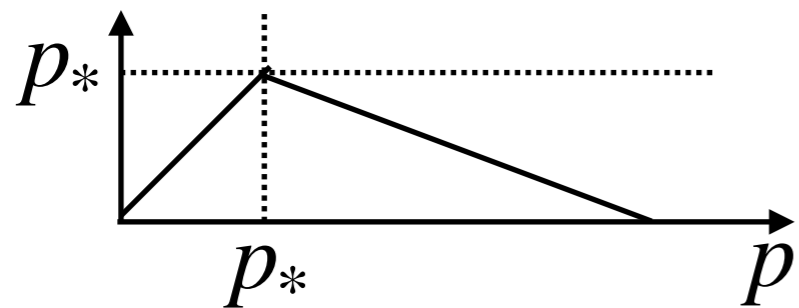
Gap function



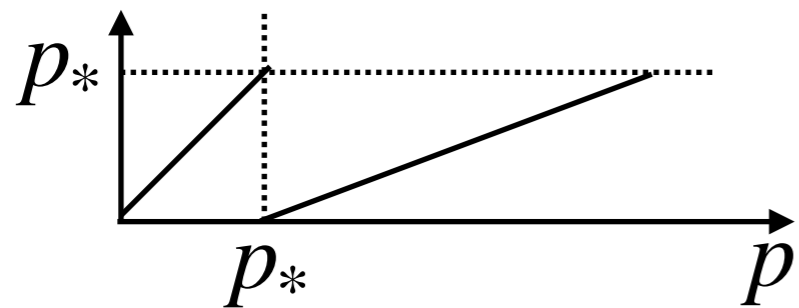
Extensions

Masking functions

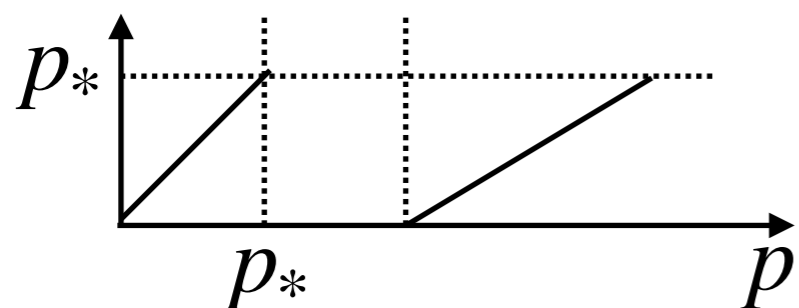
Tent function



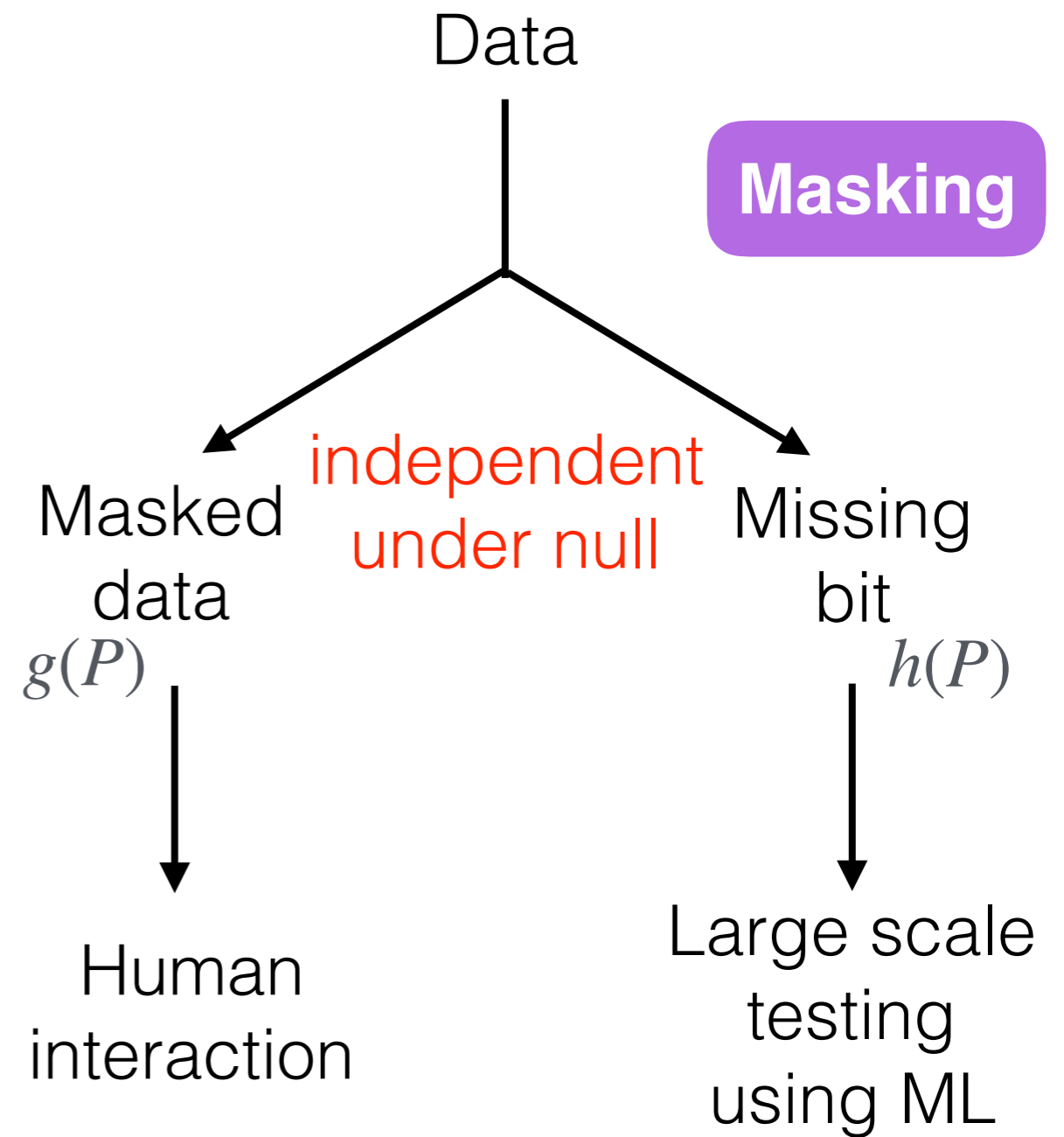
Railway function



Gap function

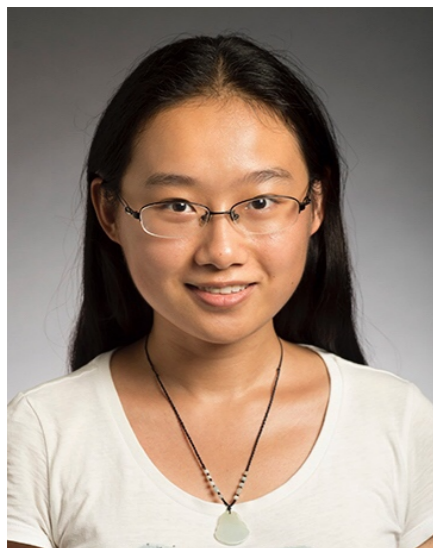


Interactive testing



Familywise error rate control by interactive unmasking

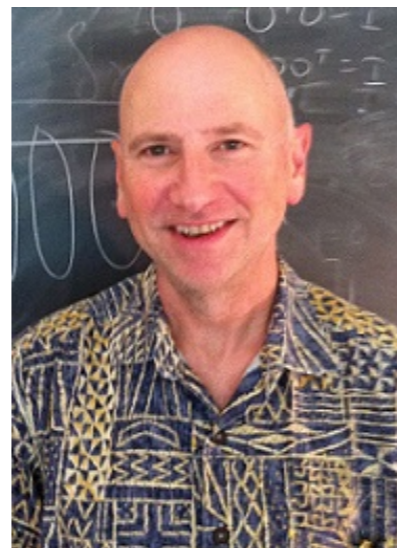
Boyan
Duan



Aaditya
Ramdas



Larry
Wasserman



Thank you!