

The joys of doing a lot of t-tests; or neuroimaging statistics using closed testing

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A day in the life...

- What do you do?
 - I develop statistical methods for neuroimaging data
- Do you do SPSS all day?
 - No
- What do you do then?
 - A lot of t-tests
- Seriously, what is ~~your~~ the problem?
 - Doing a lot of t-tests increases my chance of making mistakes

The joys of doing a lot of t-tests

Part 1:

The problem

(aka: the joys of multiple comparisons)

The joys of doing a lot of t-tests

- If I do a lot of tests, the chance of me making a 'false positive' decision increase.
- Say I have 20 hypotheses, what are the chances of finding at least one false positive?
- $P(\text{at least 1 significant}) = 1 - P(\text{no test significant})$
 $= 1 - (1 - .05)^{20} = .64$

The joys of doing a lot of t-tests

- This 64% is what we call the Family-wise error rate (FWER).
- What is the chance of making at least one false-positive decision over a family of hypothesis?
- For any number of hypotheses we want to control the FWER in a sensible way (make sure we don't want to make too many wrong decisions).

The joys of doing a lot of t-tests

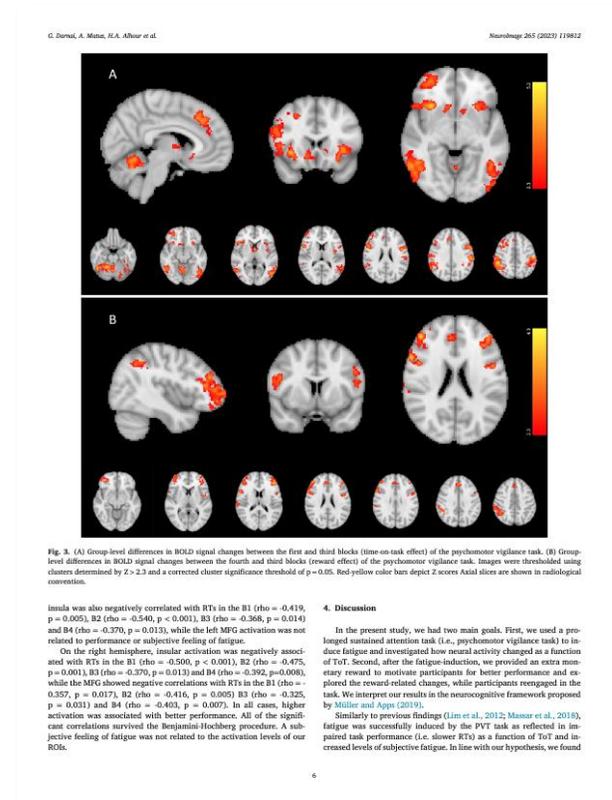
- In fMRI signal is measured within 'cubes' (voxels) in the brain over time.
- These cubes are around 3x3x3 millimeters in size.
- A typical fMRI dataset has about 200.000 voxels, measured over 300 time-points.
- Hypothesis tests are usually performed for each voxel separately.

The joys of doing a lot of t-tests

- We do over 200.000 tests, we will make some false positive decisions (declaring a voxel active while it is actually not).
- Bonferroni?
- $.05/200.000 = .00000025$ (pretty small p-value).
- Bonferroni is conservative, we will miss true activations.

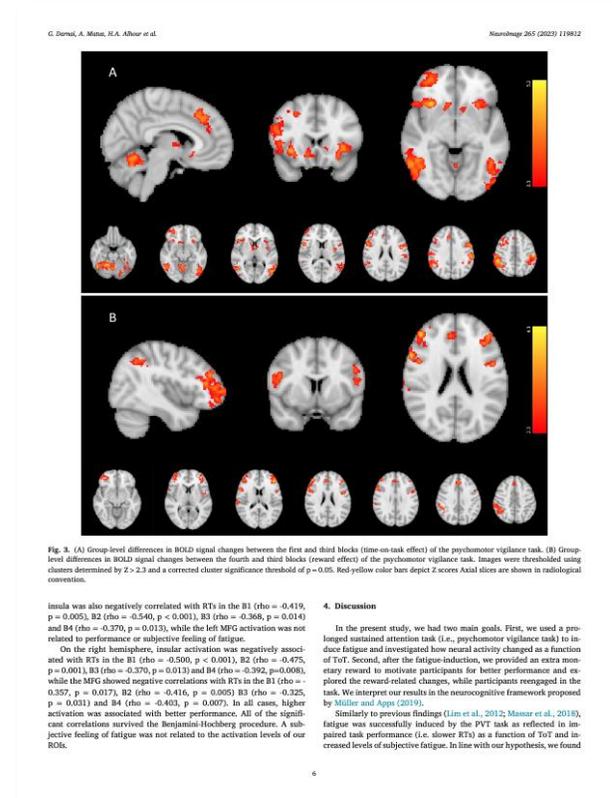
'Classical' cluster-extent analysis

- For most functional MRI studies measured signal comes from distinct locations in the brain called voxels: a 3-dimensional grid of 3x3x3 mm cubes.
- Inference in functional MRI is done on each location (voxel) separately.
- The maps that you often see are the outcomes of this inference (usually in the form of a z or t-statistic indicating significance).



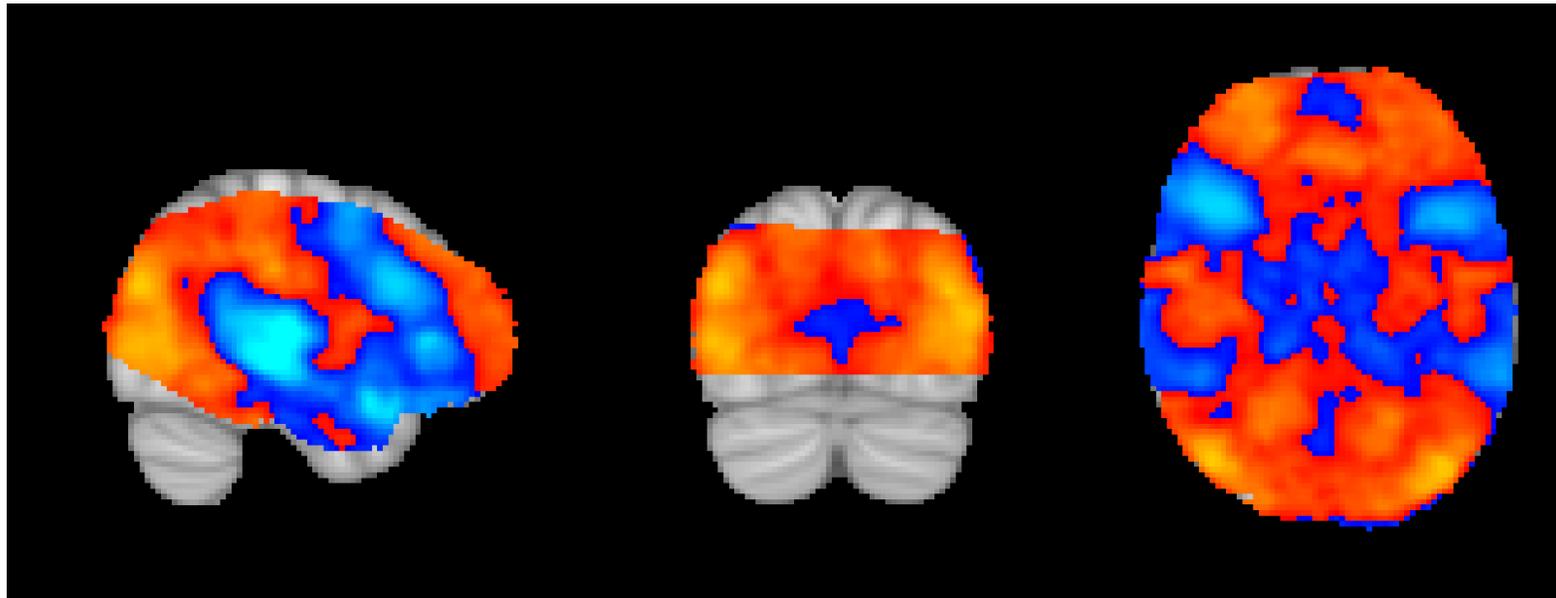
'Classical' cluster-extent analysis

- The goal of fMRI inference is to decide for each voxel whether it is active or not (using a hypothesis test).
- For each test we allow a little uncertainty of whether our decision is the right one.
- When doing multiple tests, the chances of making a wrong decision somewhere in our 'family' of tests increases dramatically.
- The family-wise error rate (FWER) of our family of tests is what we want 'controlled'.



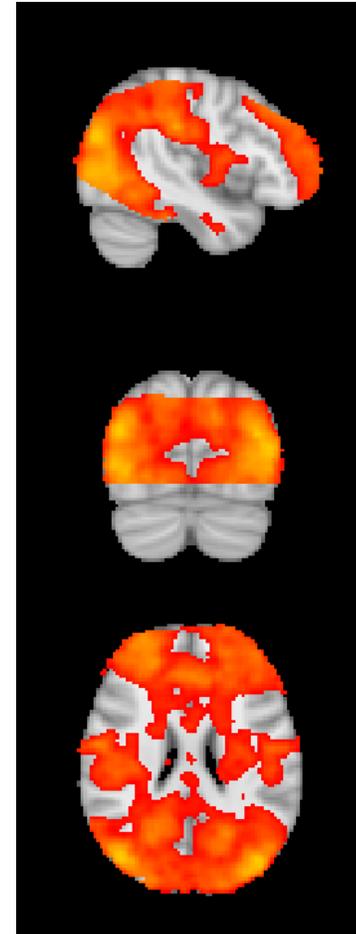
'Classical' cluster-extent analysis

Study on vocal and non-vocal sounds, Pernet et al., 2015



'Classical' cluster-extent analysis

- In total 166.407 in-mask voxels.
- Focus only on positive values for now.
- Z-statistics indicate whether a voxel is more active in the *non-vocal* condition than in the *vocal* condition.
 - H_0 = not active (z-value = 0)
 - H_1 = active (z-value > 0)

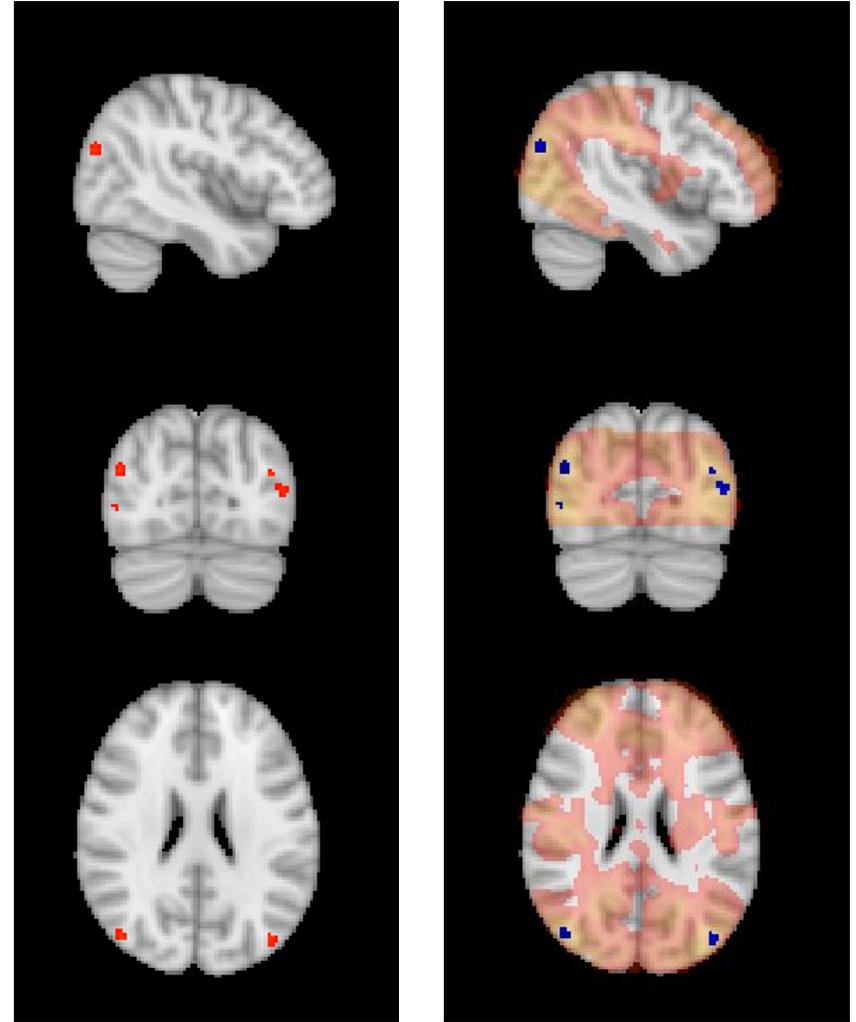


'Classical' cluster-extent analysis

- Controls the FWER over all voxels in the brain (mask). Family = all voxels.
- Easiest method to control the FWER is Bonferroni correction.
- Calculated by setting the per-voxel α to be $\alpha / \text{\#voxels}$

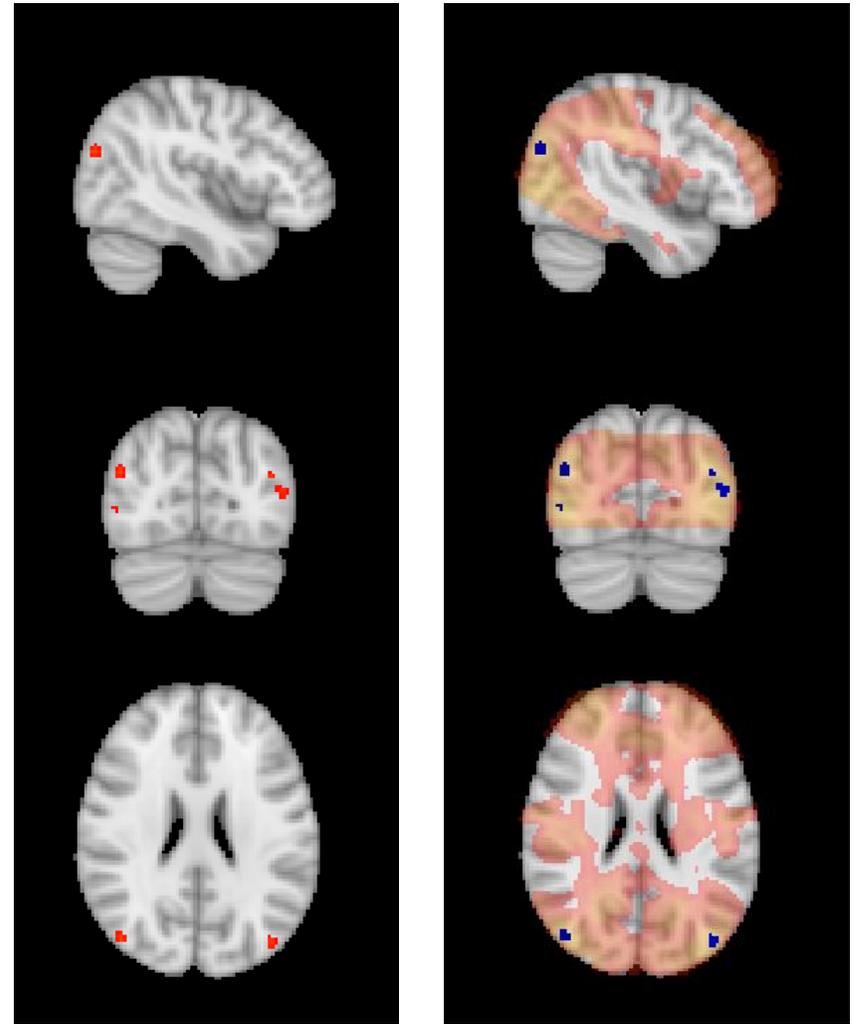
$$.05 / 166407 = .0000003$$

- Usually not very powerful.



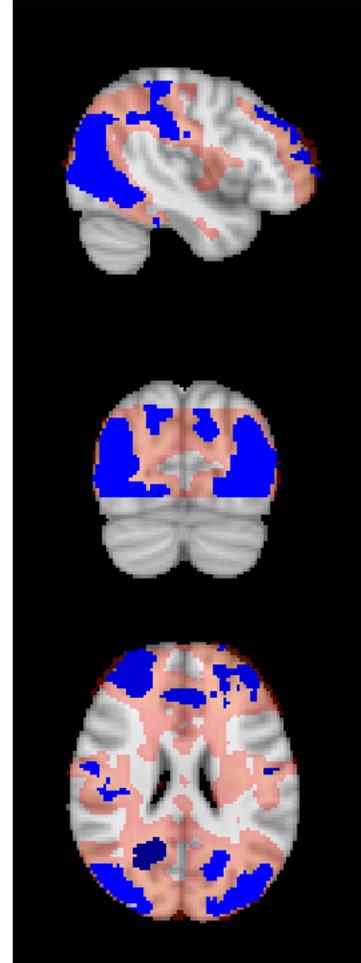
'Classical' cluster-extent analysis

- But...
- Since our family is all voxels, we know exactly where the activation is!
- In other words: we have high spatial specificity.
- (because the chance of any of these voxels being a false-positive $< 5\%$)



'Classical' cluster-extent analysis

- Usually, we are not interested in single-voxel activity per se. A more natural unit is a 'cluster' of voxels (which we will name 'blob').
- A cluster or blob is defined as a contiguous/connected set of voxels.
- We control the number of false-positive blobs (our family in FWER is thus all possible blobs, not all voxels).

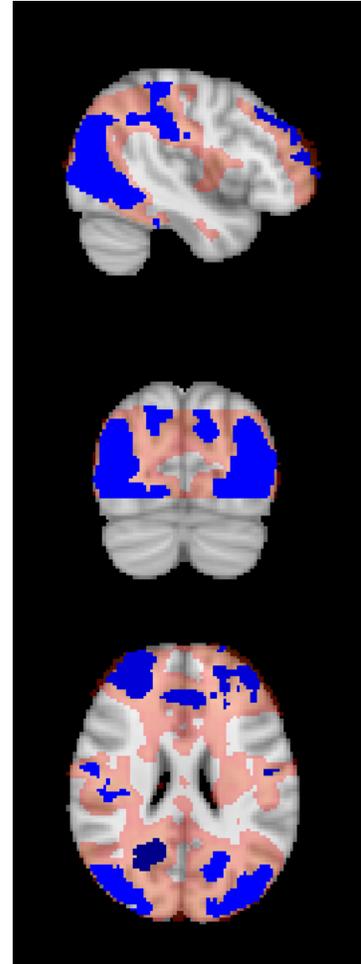


'Classical' cluster-extent analysis

- In practice, using a two-step approach:
 1. Choose a 'cluster-forming' threshold z and estimate the size of all contiguous clusters above this threshold.

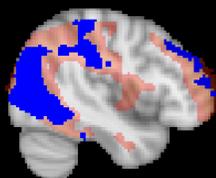
Determine the minimum cluster size k that occurs by chance under the null (95%) given the smoothness of the data and the chosen threshold z (e.g., using RFT or permutations)

2. Check which clusters are larger than k (all clusters that are larger are significant).

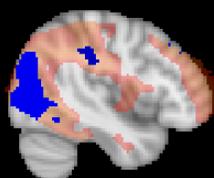


'Classical' cluster-extent analysis

$Z > 2.3$
 $k = 597$



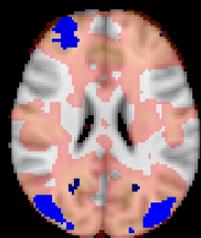
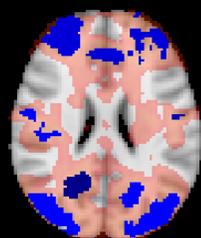
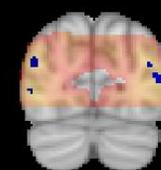
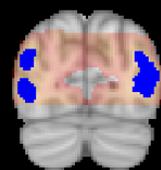
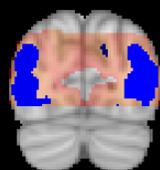
$Z > 3.1$
 $k = 160$



$Z > 4.0$
 $k = 25$

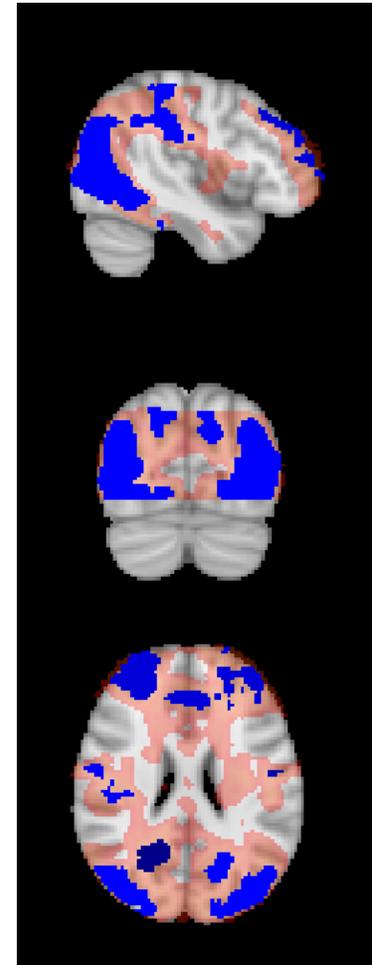


$Z > 5.0$
 $k = 1$



'Classical' cluster-extent analysis

- More powerful than voxel-wise approaches, but... more powerful in detecting activation, not in localizing it.
- Because of hypotheses being on the 'cluster' level:
 - Non-significant when cluster-extent is smaller than k
 - Significant when cluster-extent is larger than k
- No information about voxels within a cluster (clusters are large enough or not).



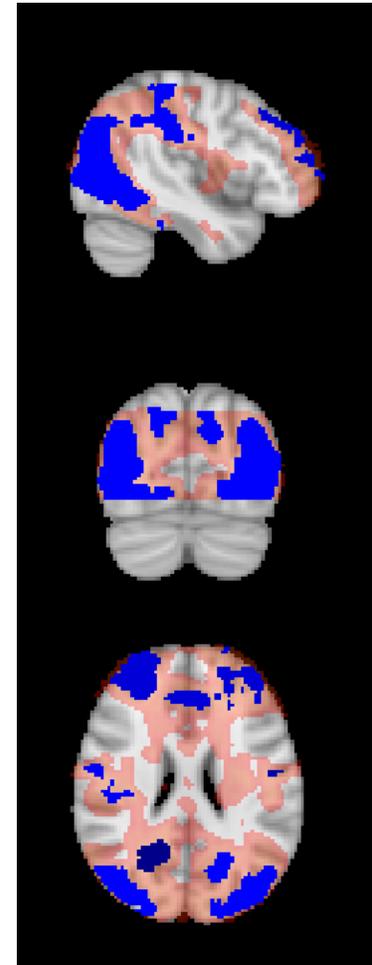
'Classical' cluster-extent analysis

- Formal way of stating this:

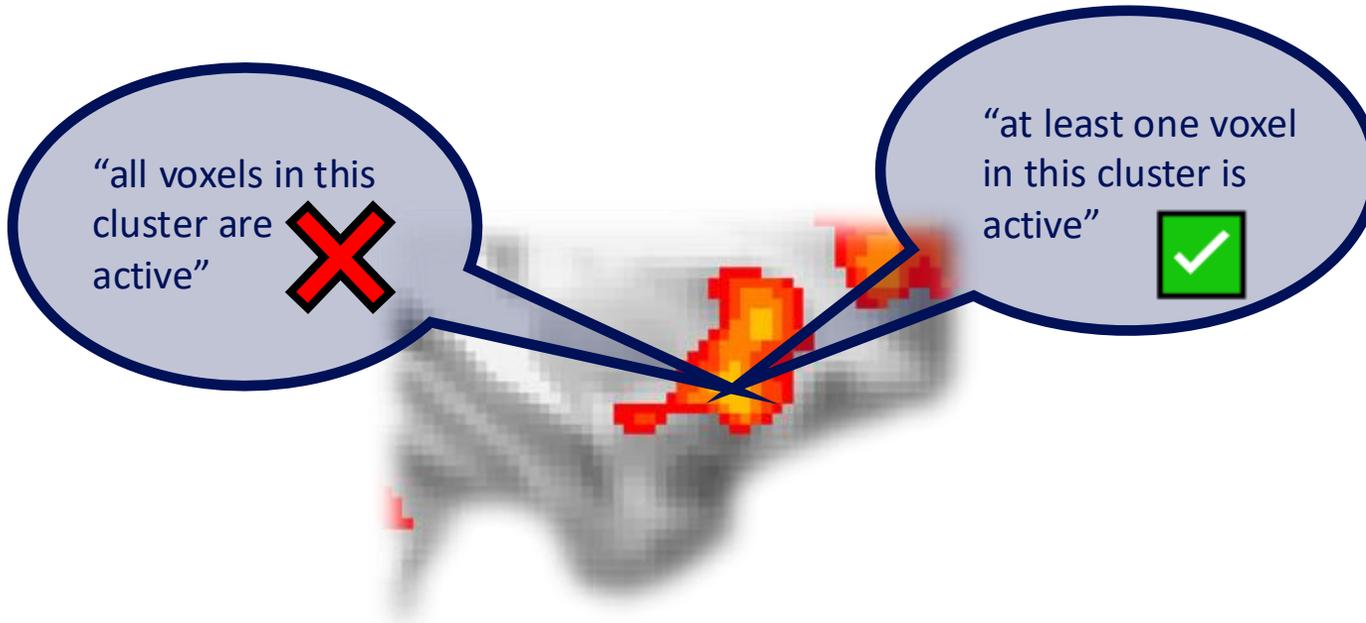
H_0 = no activation within a cluster

H_1 = at least one voxel active within a cluster

- So, the larger the cluster found, the less we know about activation within a cluster.
- This is called the Spatial specificity paradox (Woo et al., 2014, Lindquist & Mejia, 2015).



Spatial Specificity Paradox

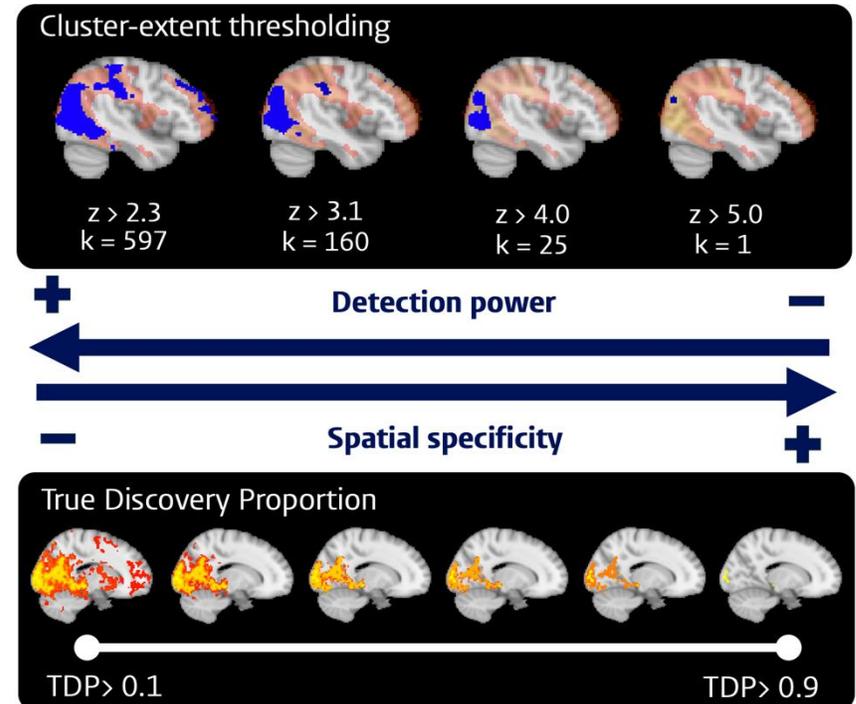


H_0 = no activation within a cluster

H_1 = at least one voxel active within a cluster

Spatial Specificity Paradox

- The statement “there is at least one voxel active” could mean that all voxels in a cluster are active. Or it could be one, we just don’t know.
- Intuition is that there is usually more than one voxel active. But cluster-extent statistics don’t allow us to test that.
- We can use TDP based methods to give us an in-depth analysis of what clusters are made of.



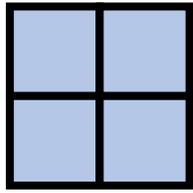
Part 2:

A crash course in closed testing

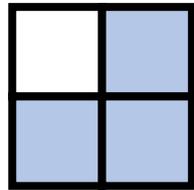
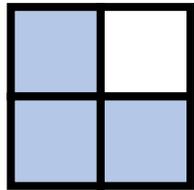
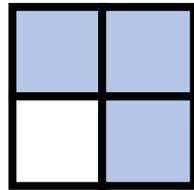
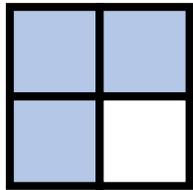
The four voxel brain

.026	.029
.031	.207

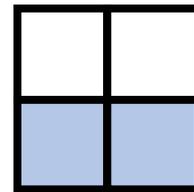
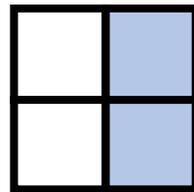
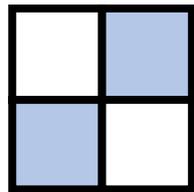
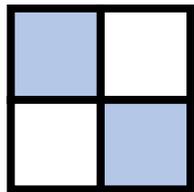
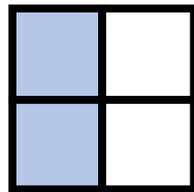
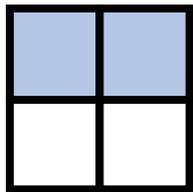
The four voxel brain – all subsets



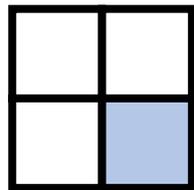
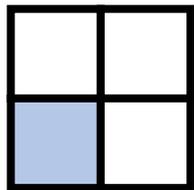
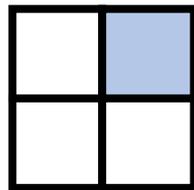
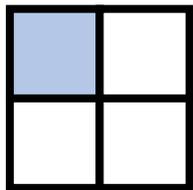
4 voxels set



3 voxels sets

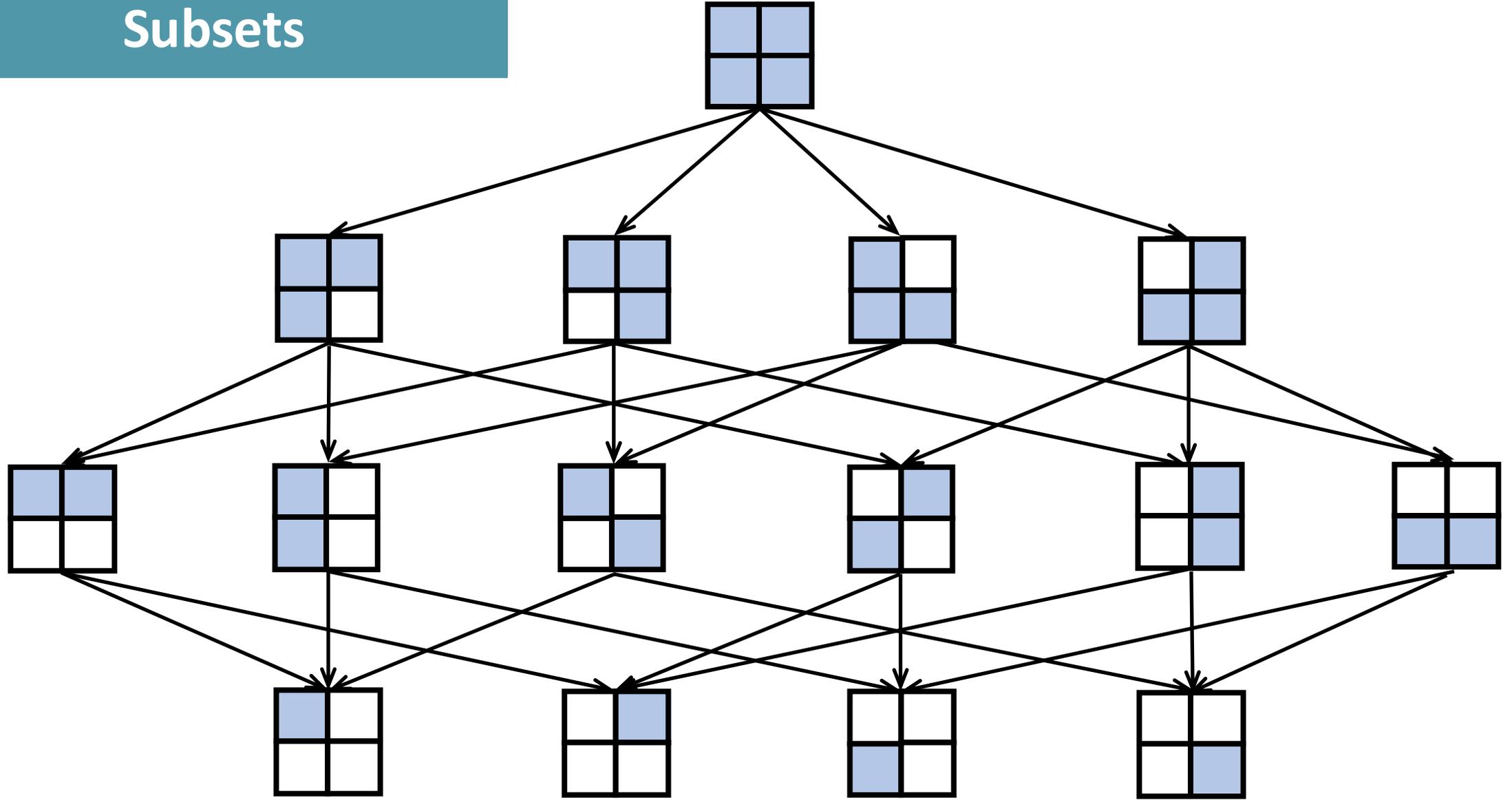


2 voxels sets

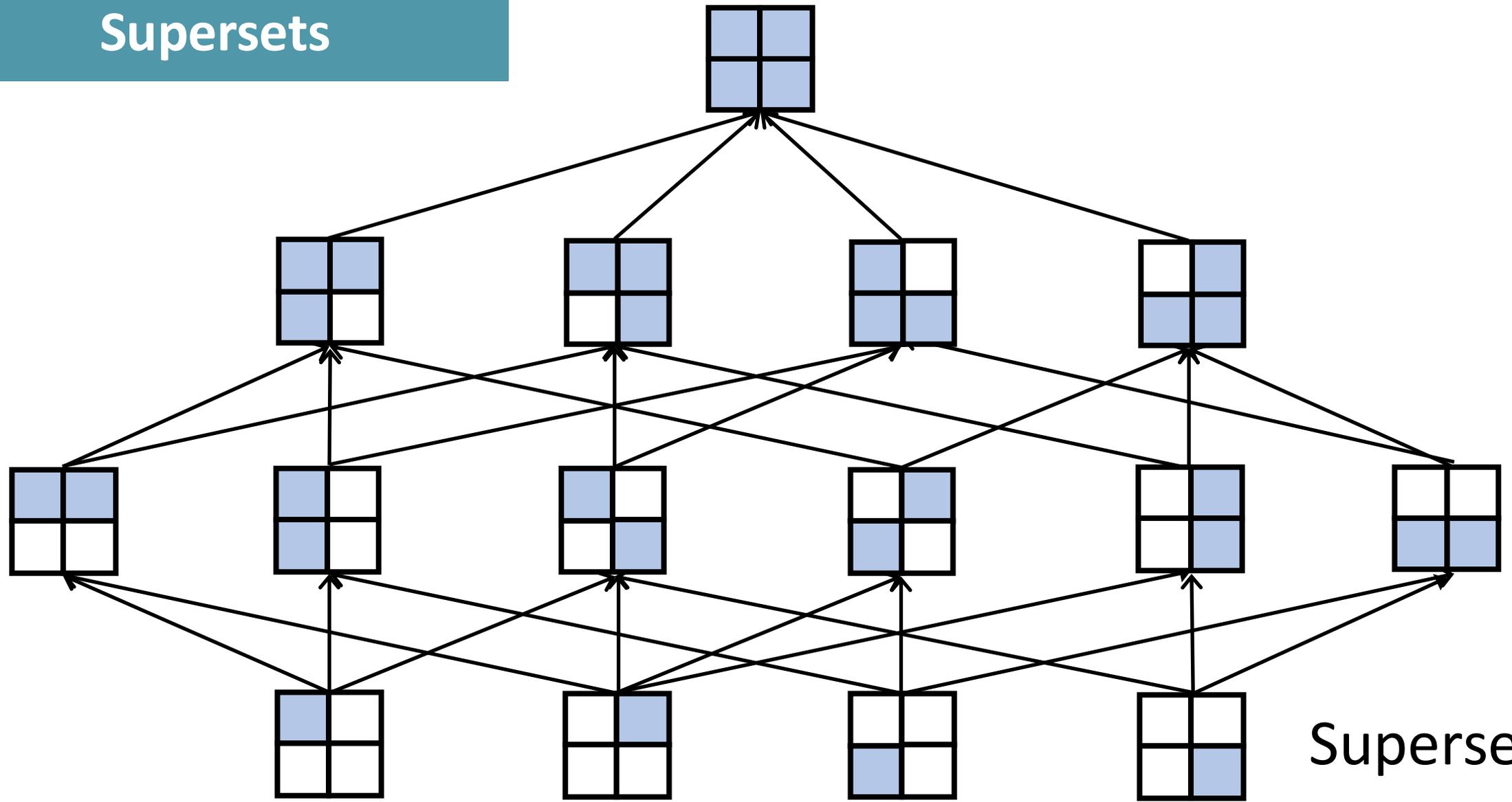


1 voxel sets

Subsets



Supersets



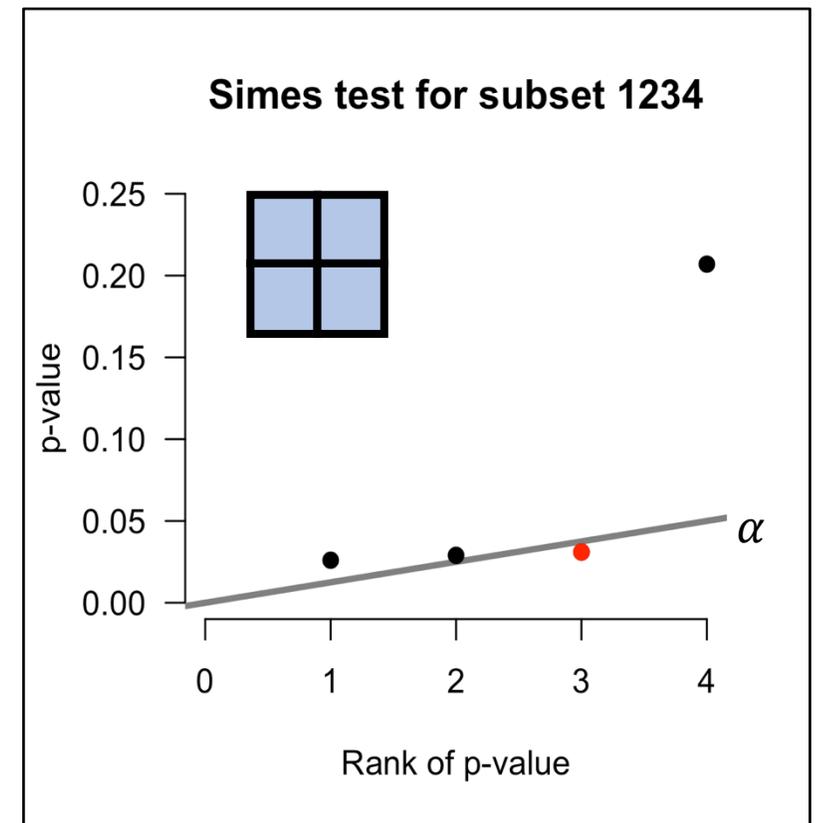
Simes test

Is there at least one active voxel in subset S ?

$$H_S: a(S) = 0$$

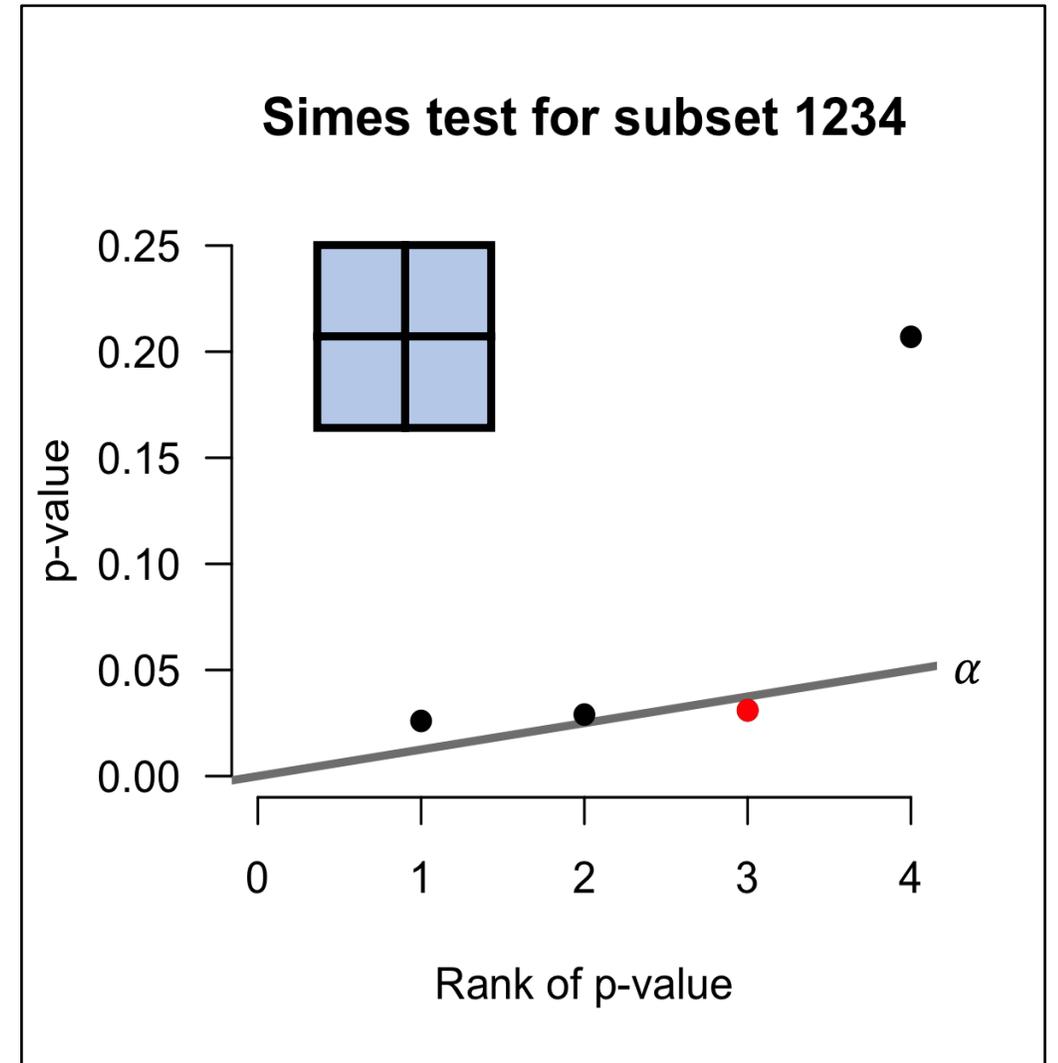
$$p_S = \min \left\{ \frac{|S|}{i} p(i:S), \text{ with } 1 \leq i \leq |S| \right\}$$

Reject H_S if $p_S \leq \alpha$.

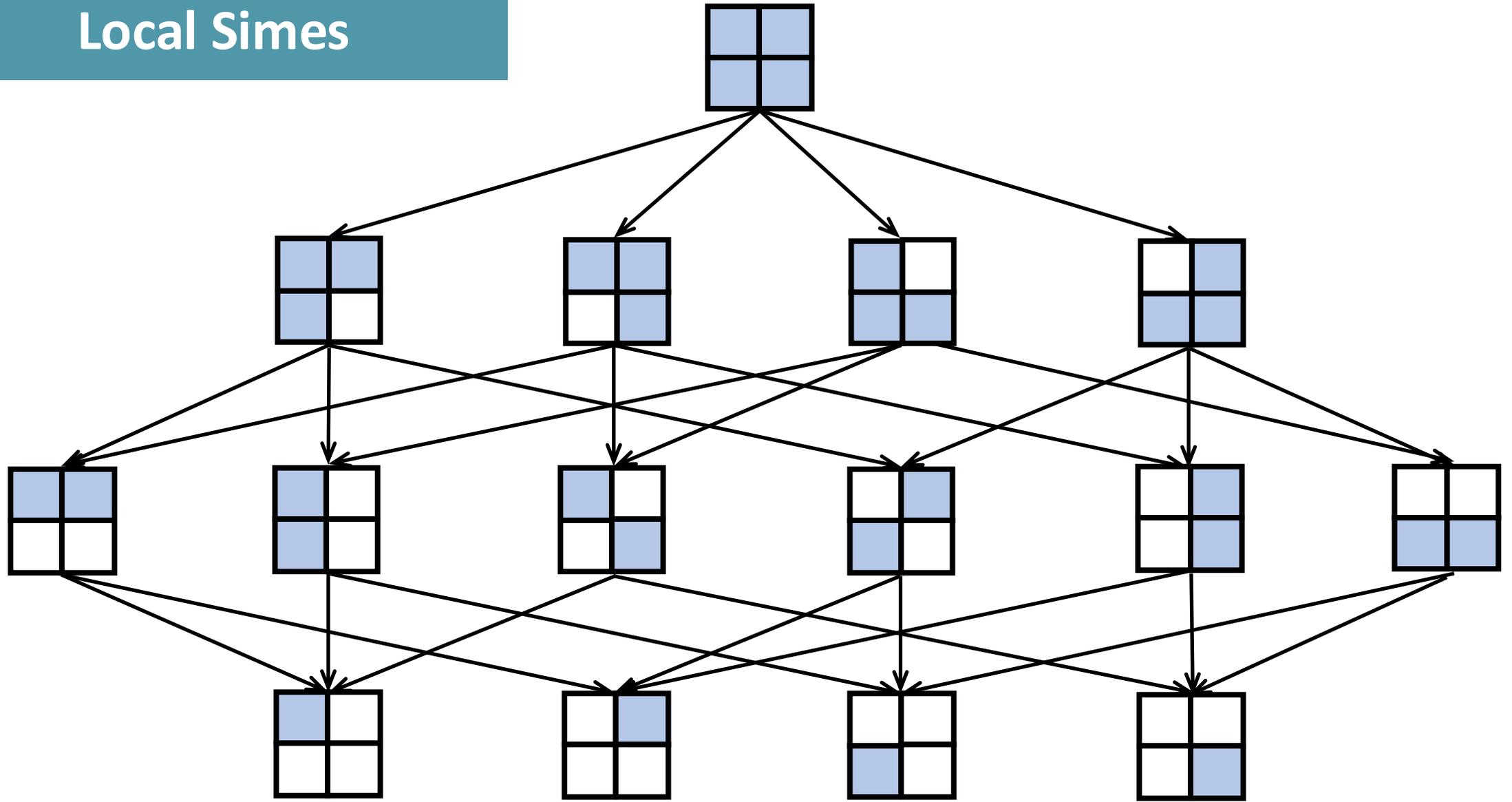


Simes test

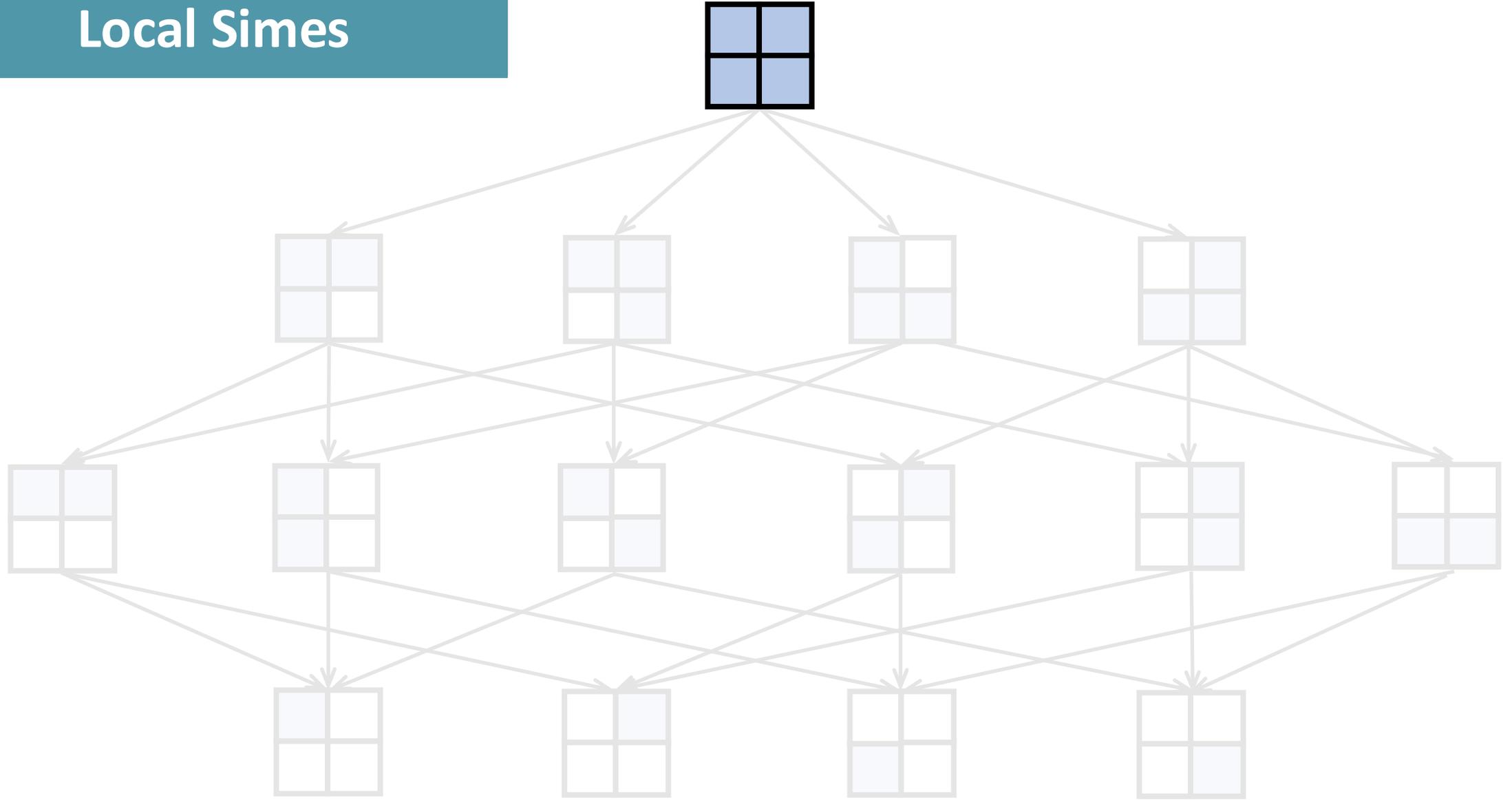
- Order p-values from smallest to largest.
- Draw a line from 0 to alpha.
- Is there any point below the line?



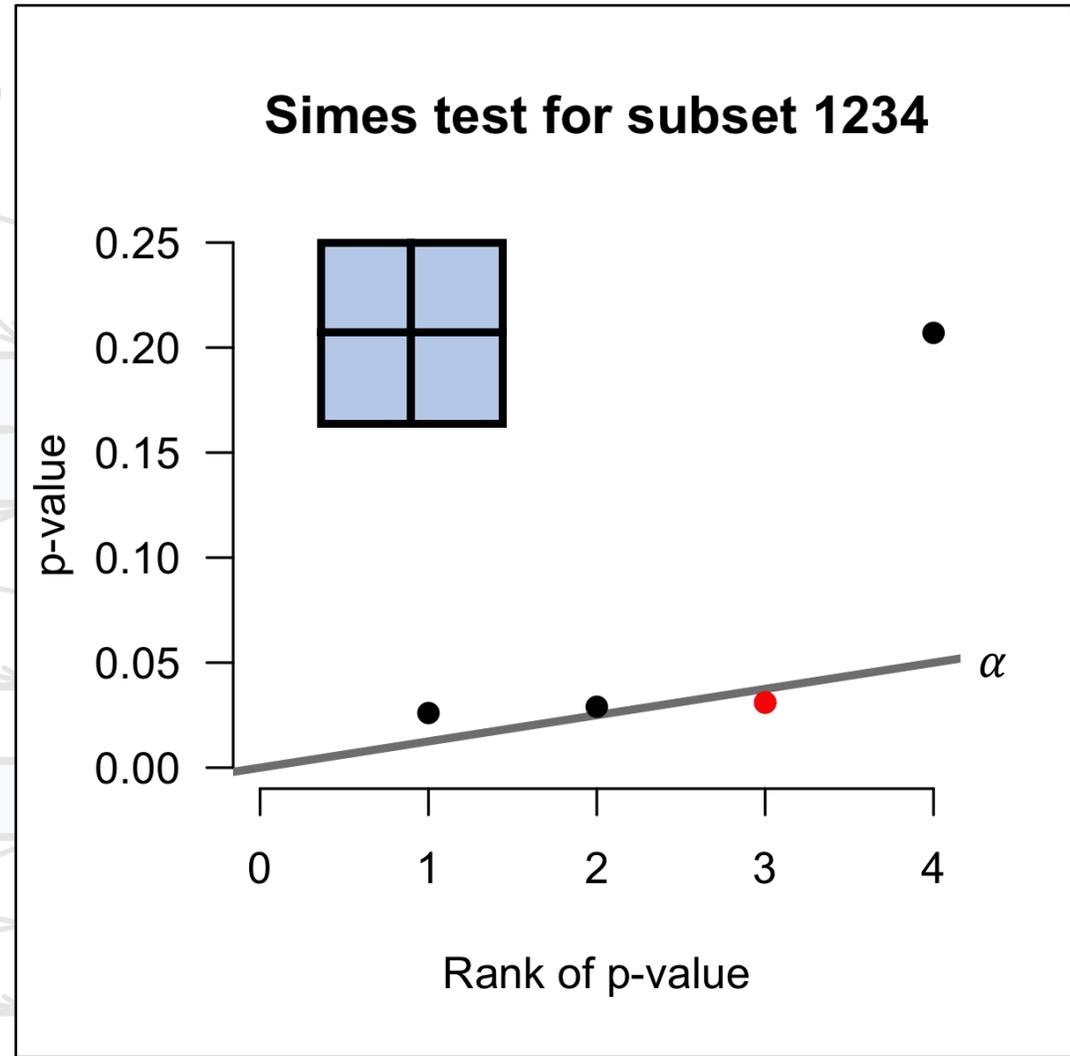
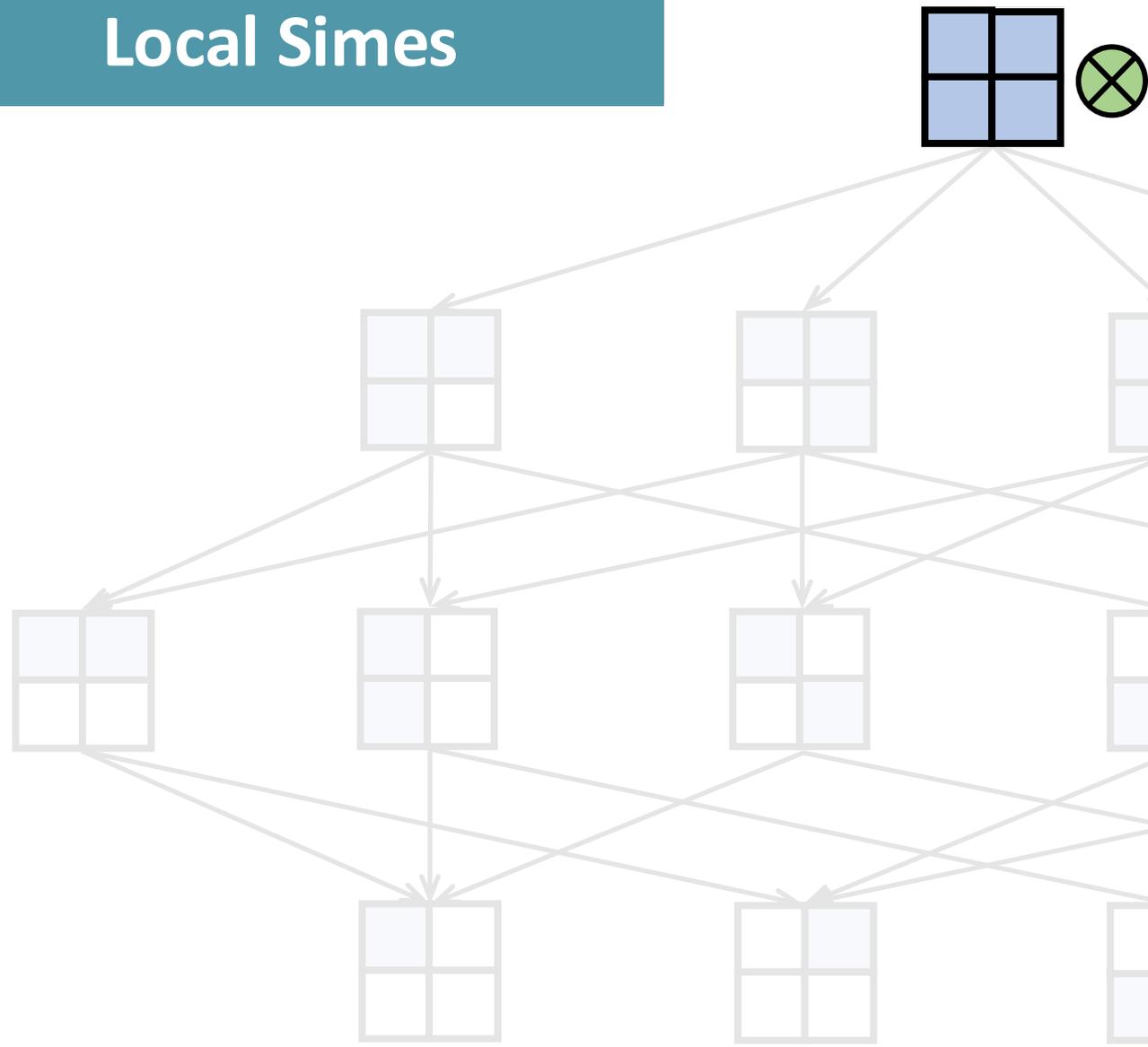
Local Simes



Local Simes



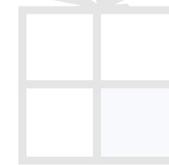
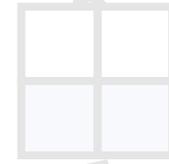
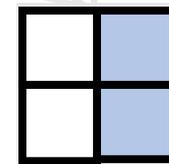
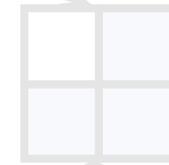
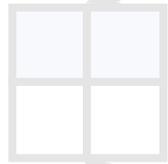
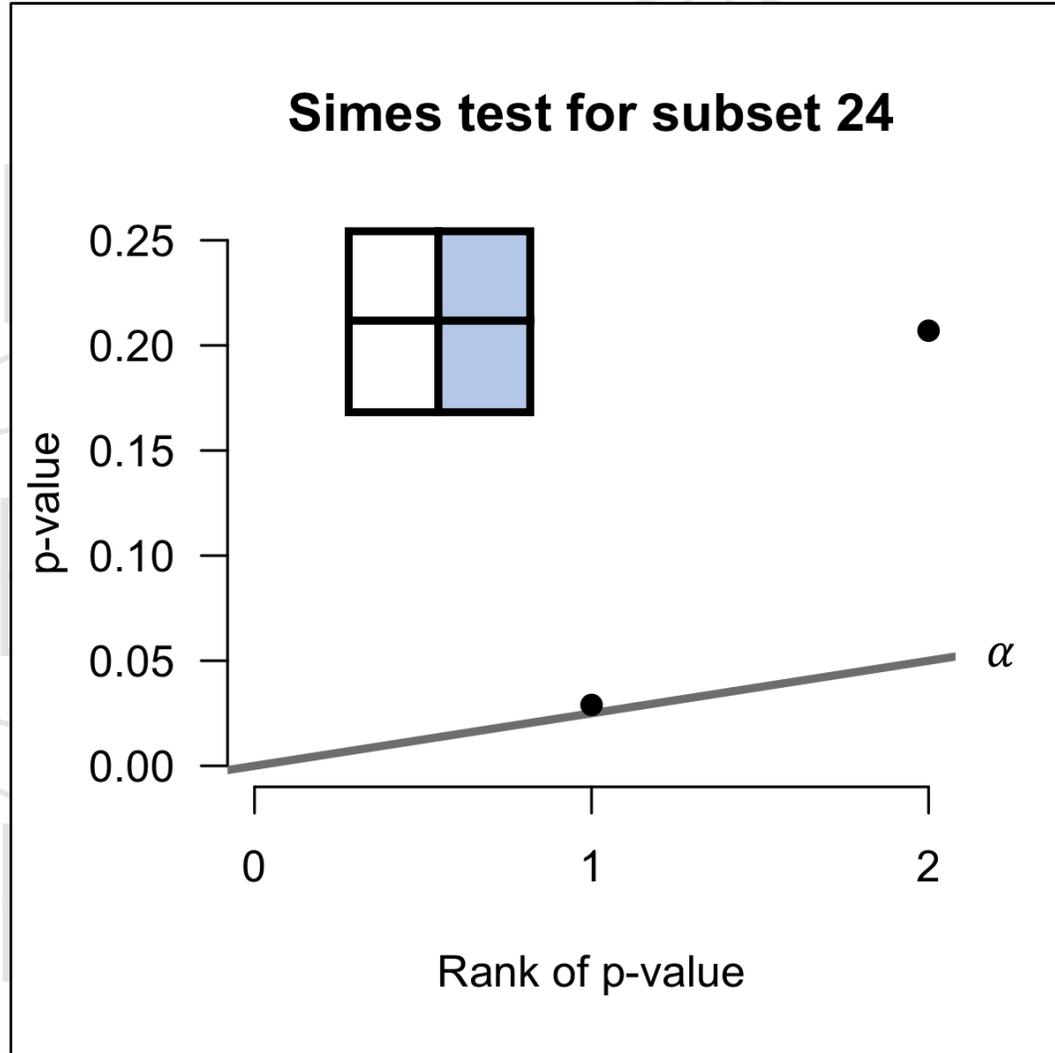
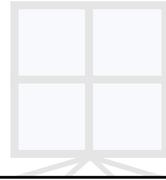
Local Simes



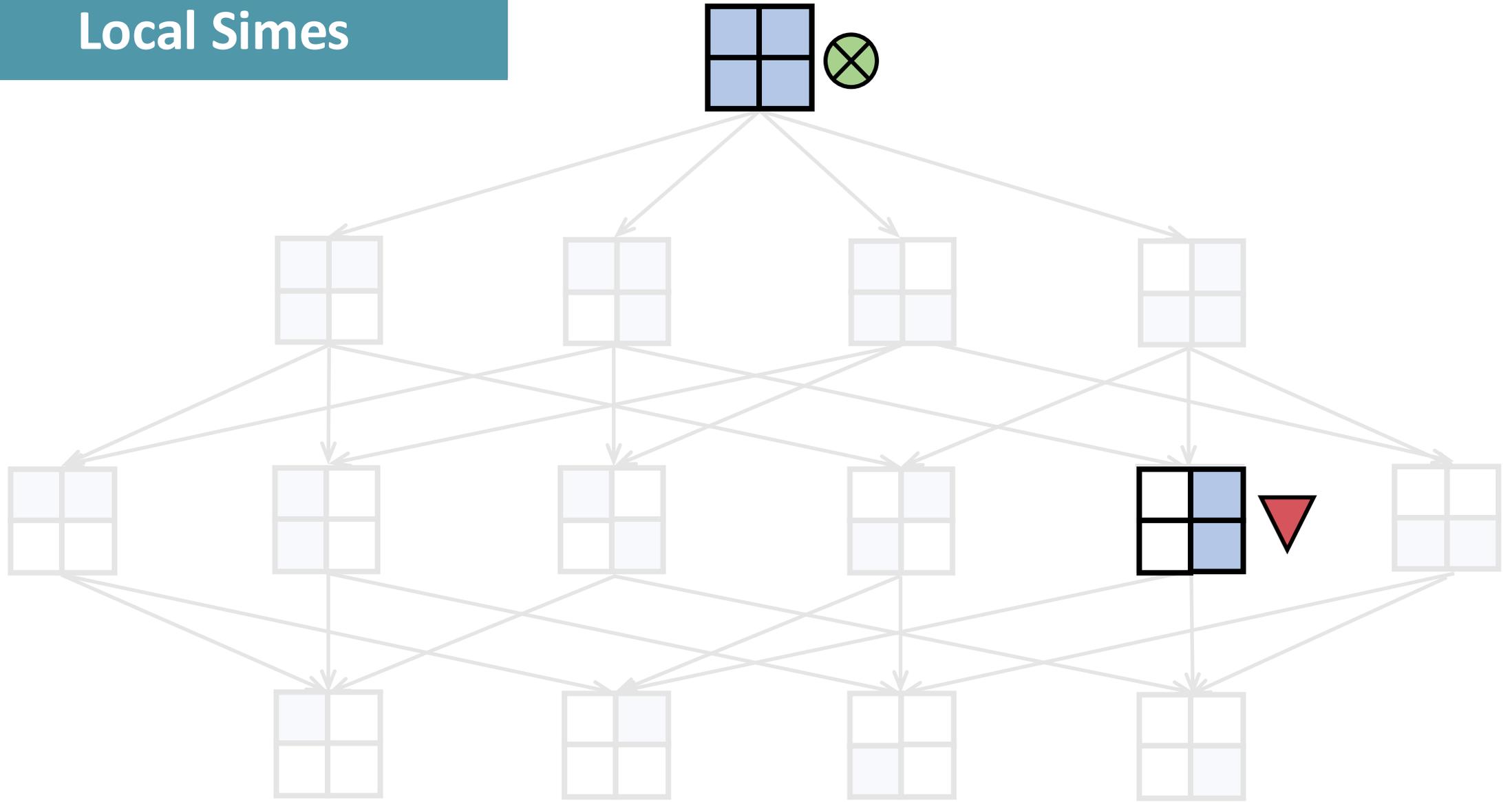
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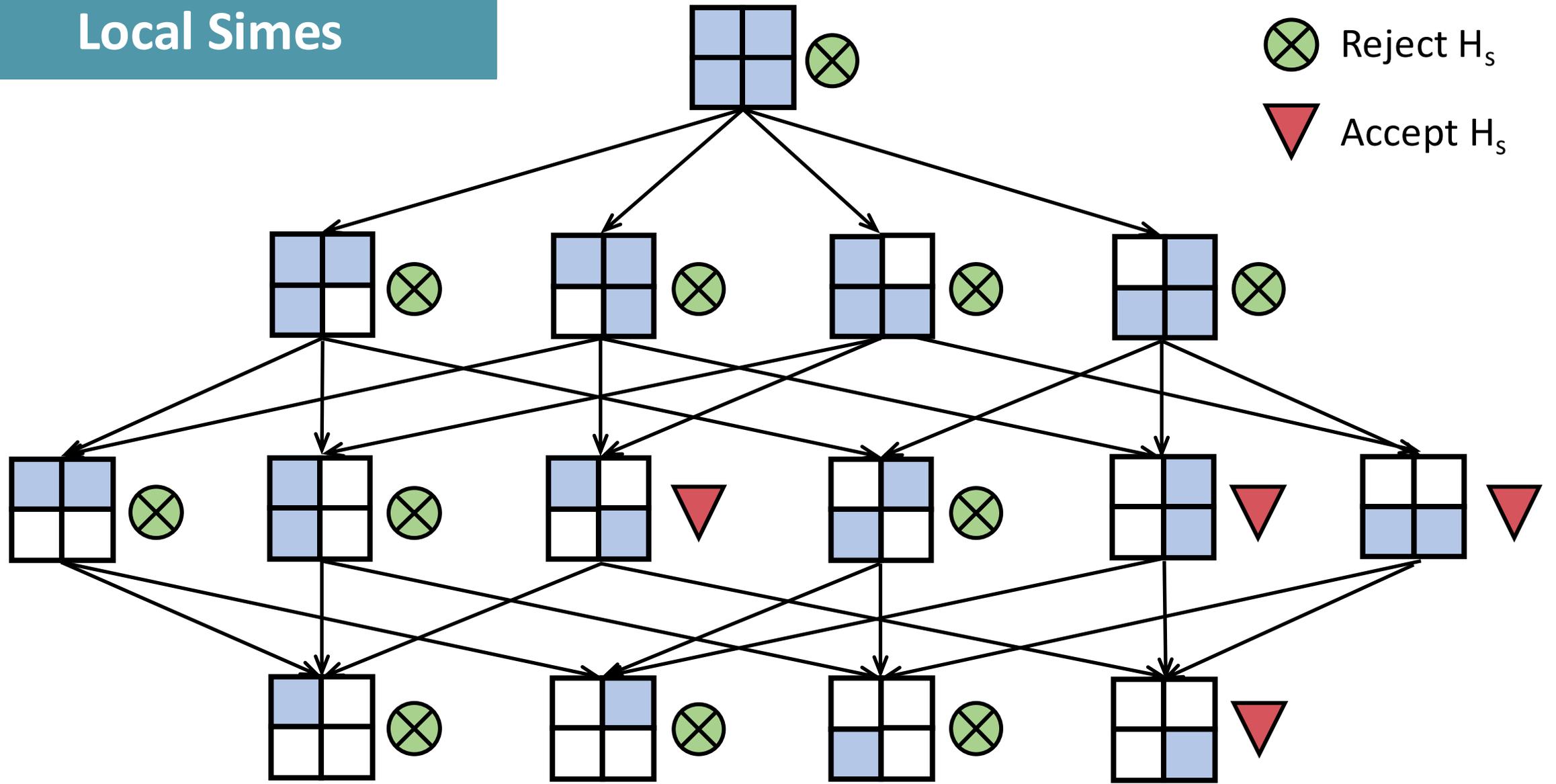
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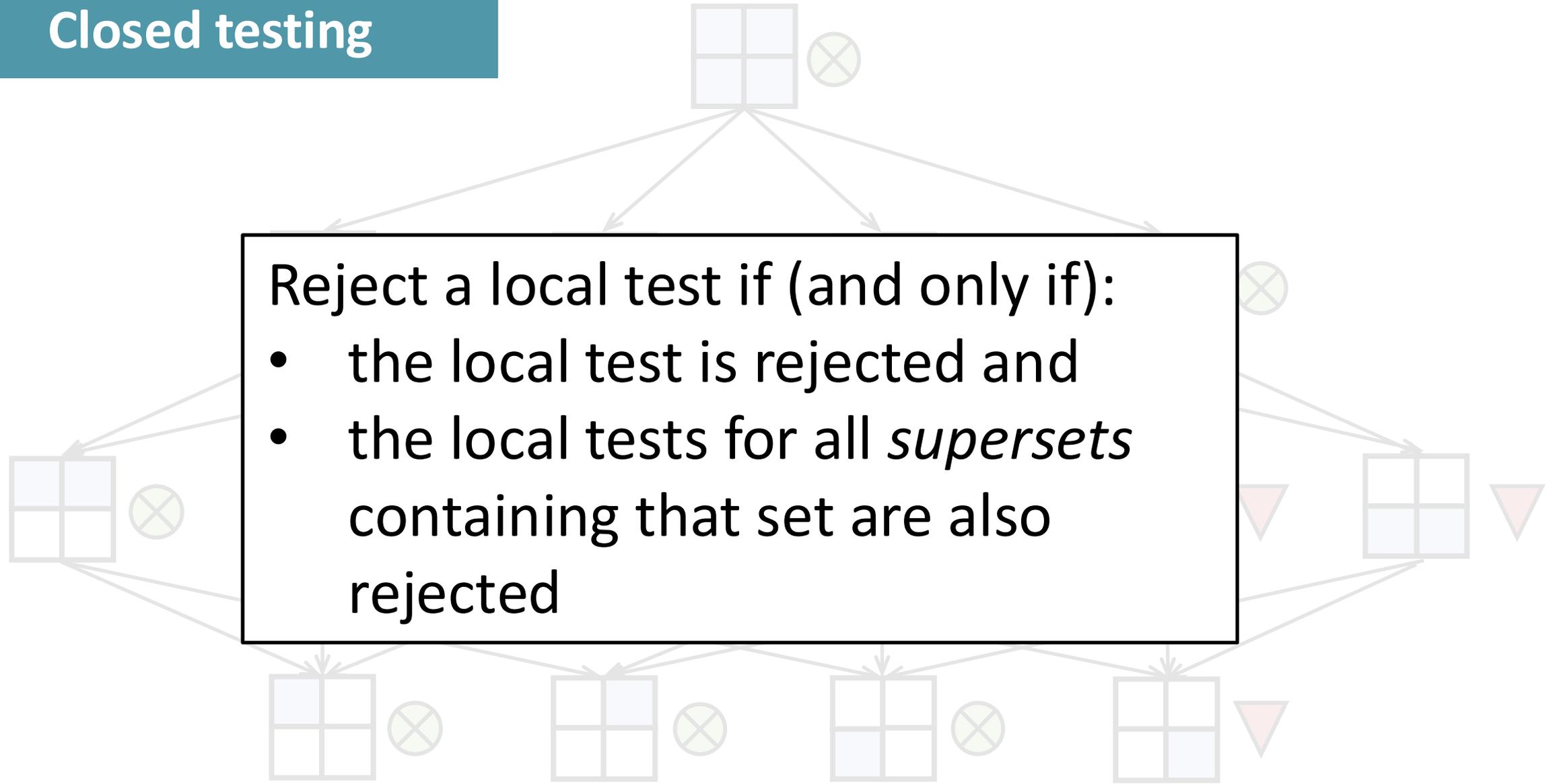
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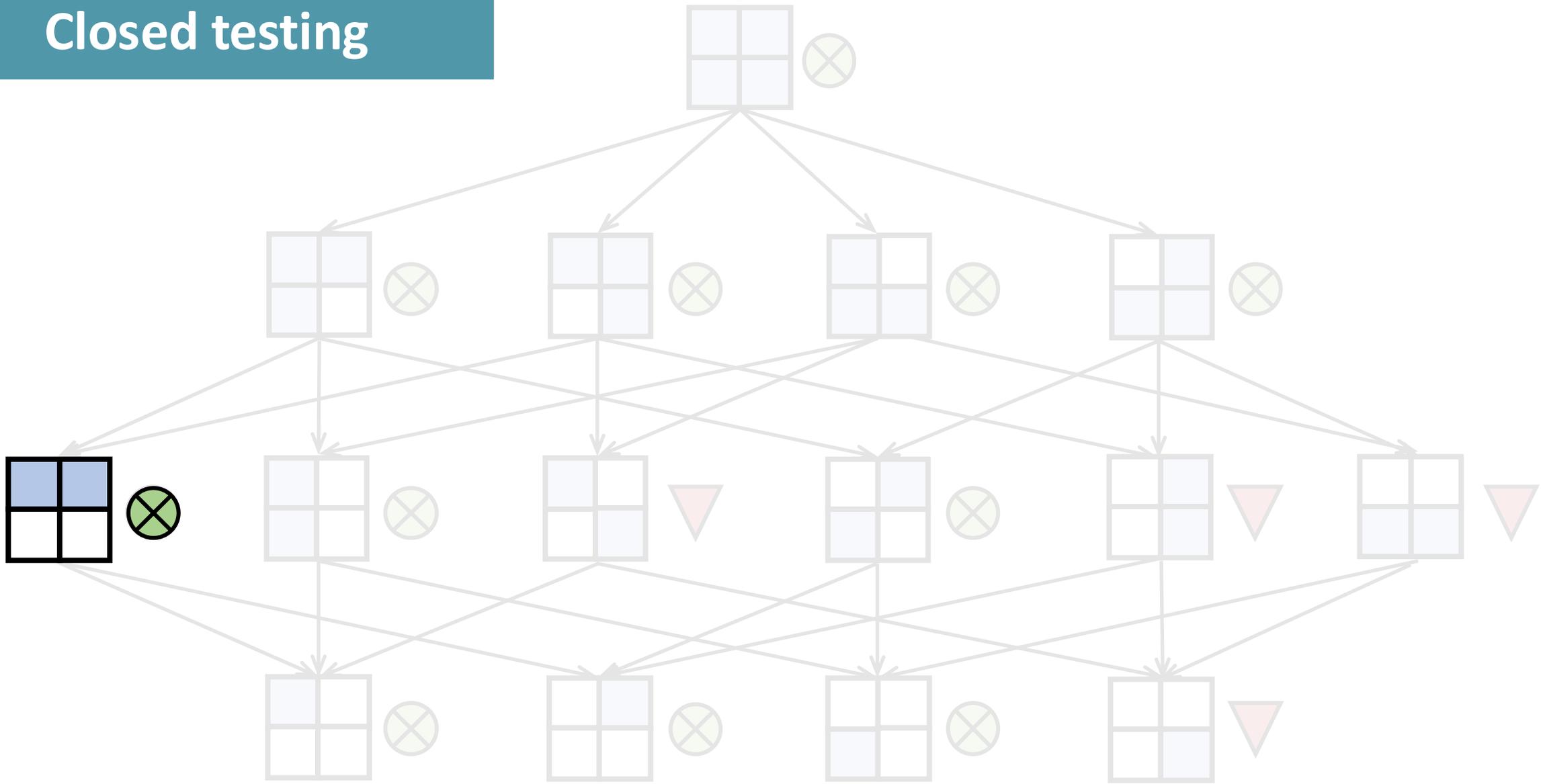
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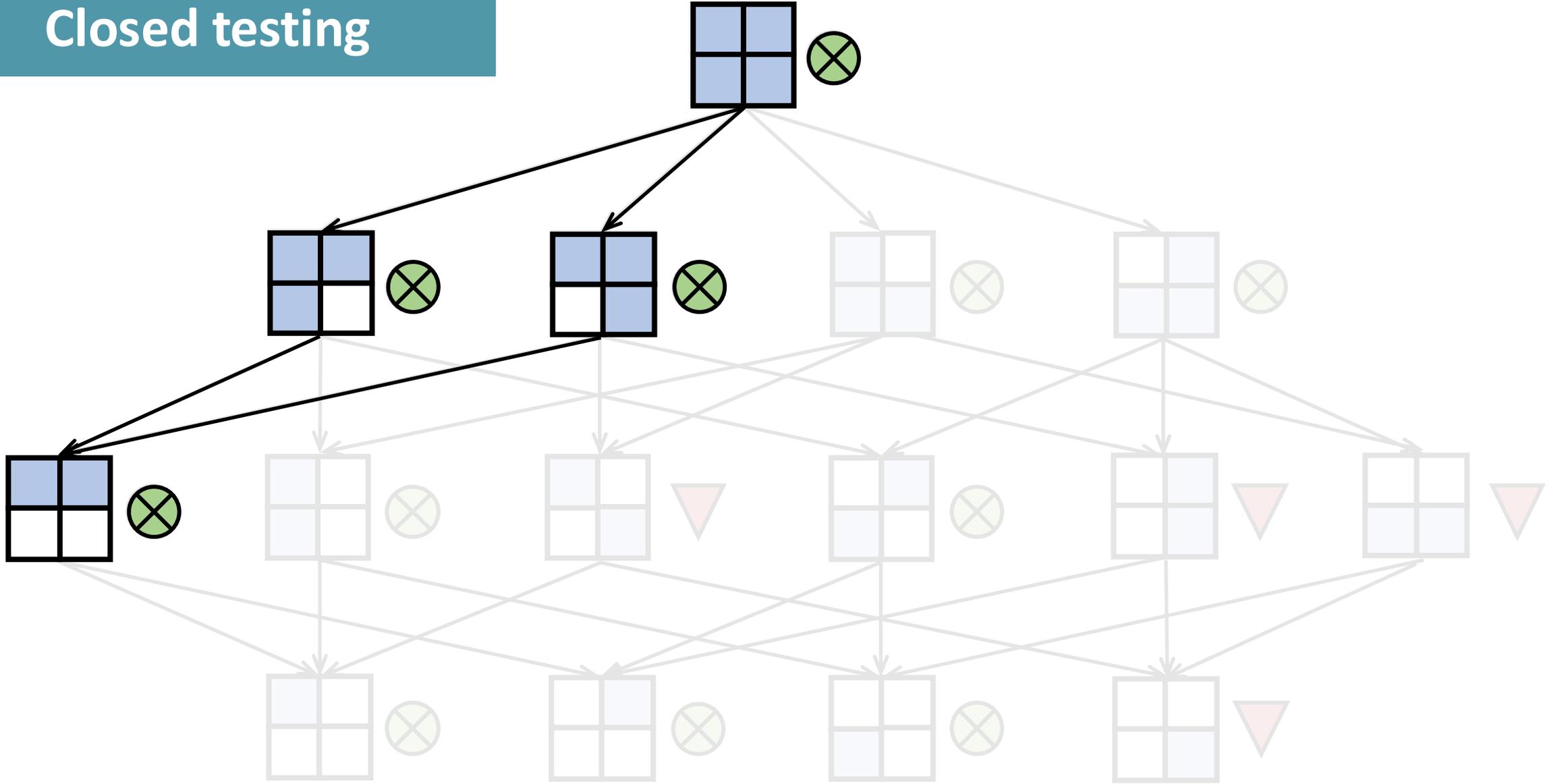
Closed testing



Closed testing

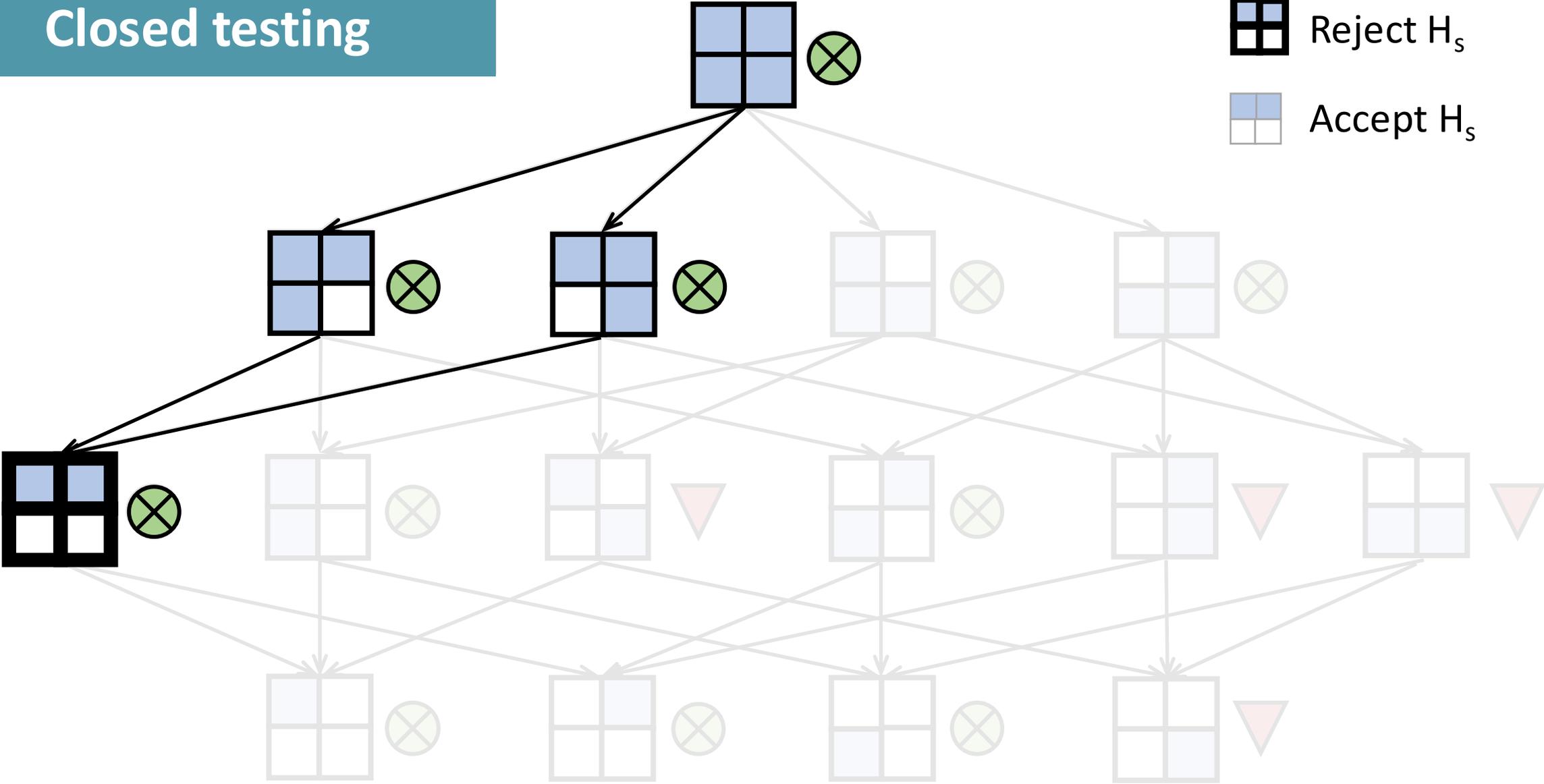


Closed testing



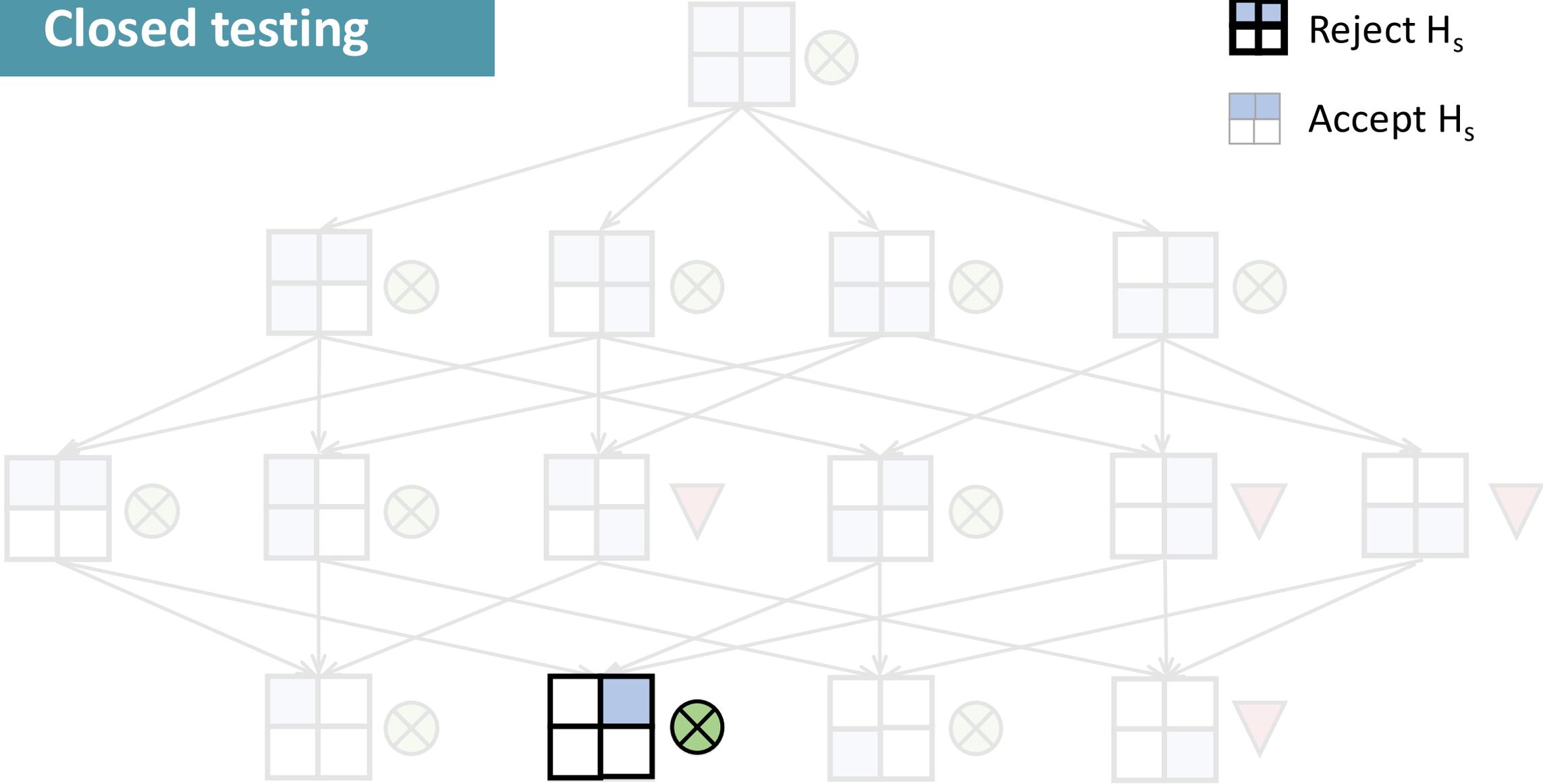
Closed testing

 Reject H_s
 Accept H_s



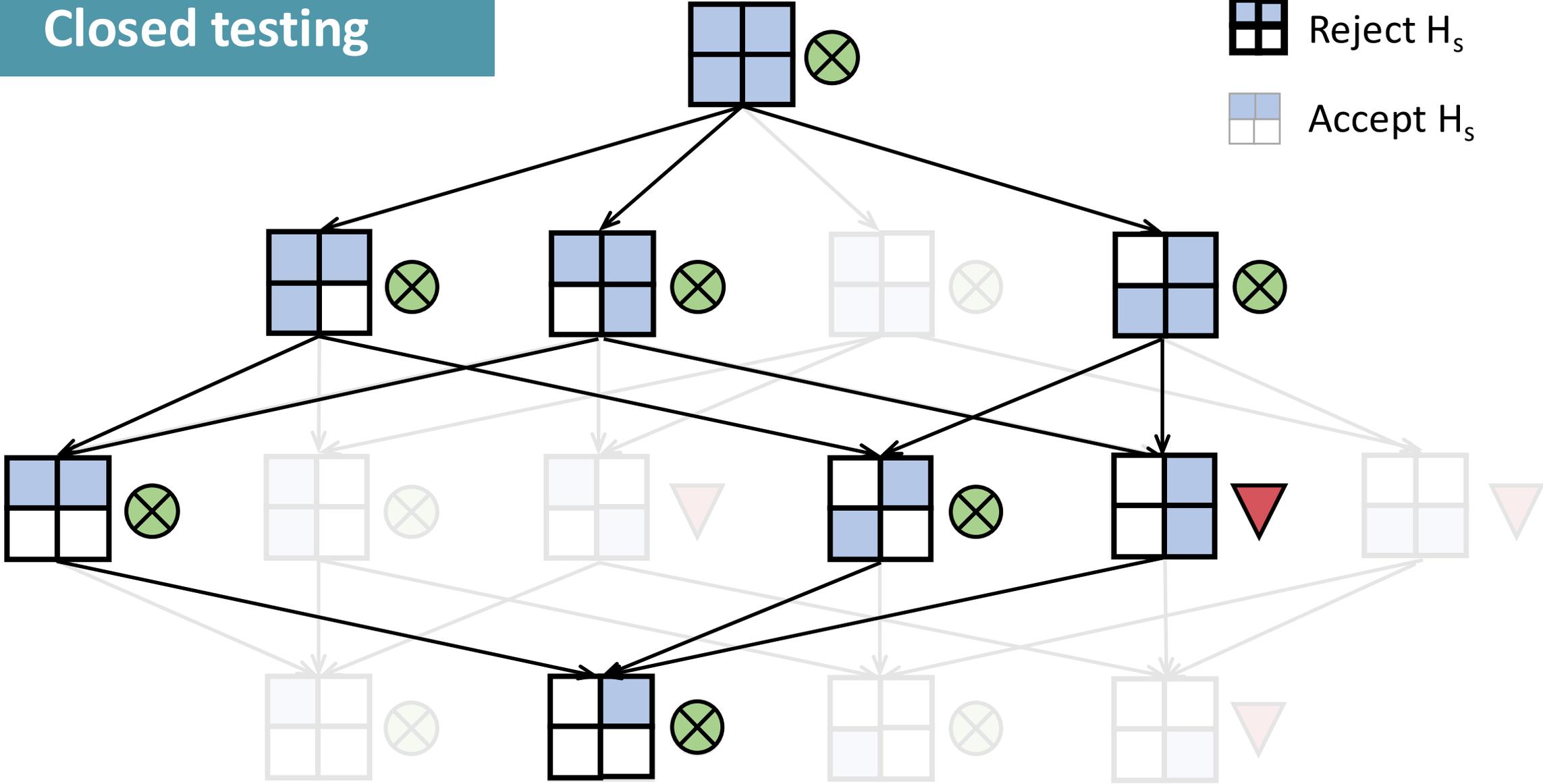
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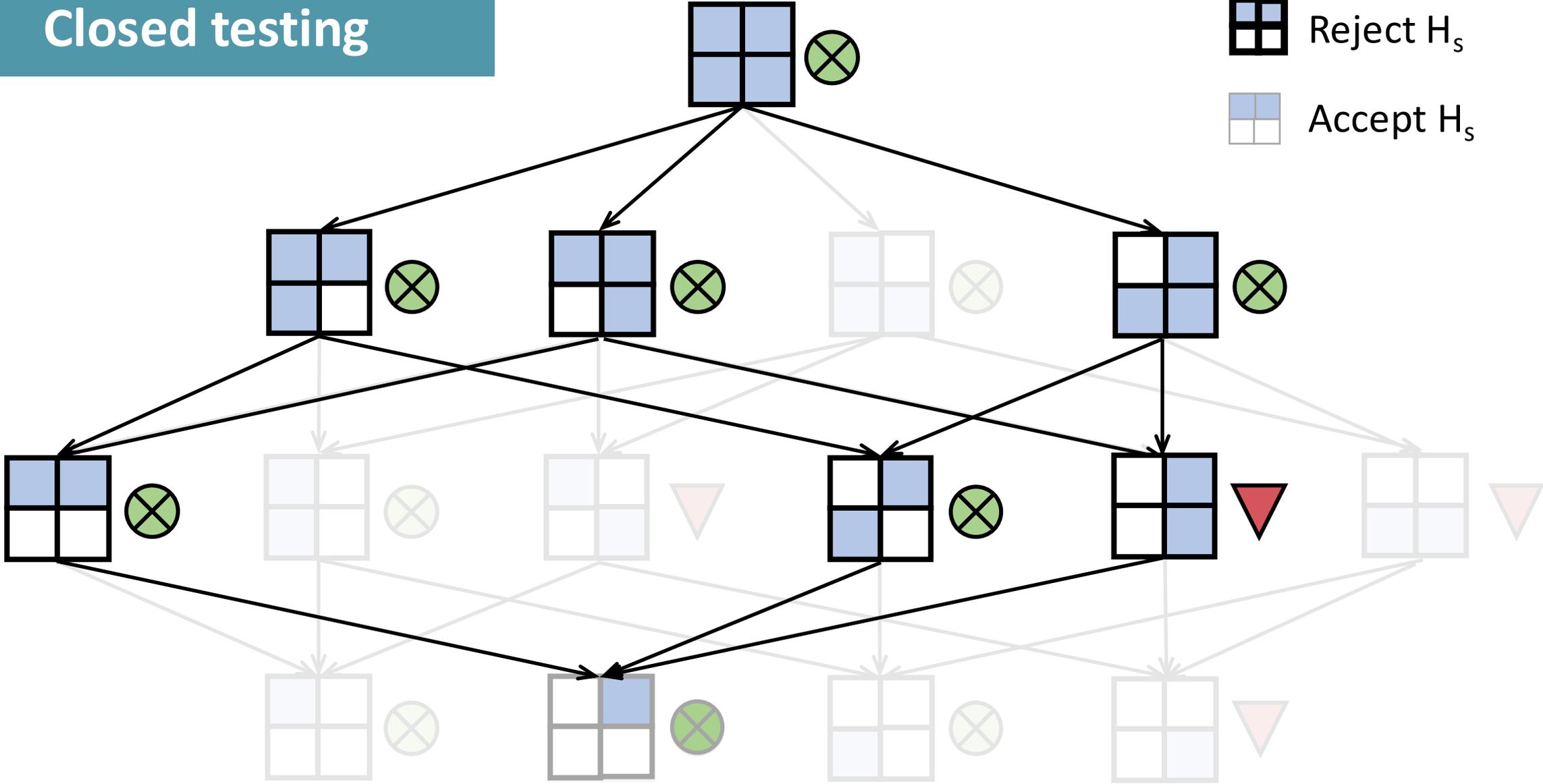
Closed testing

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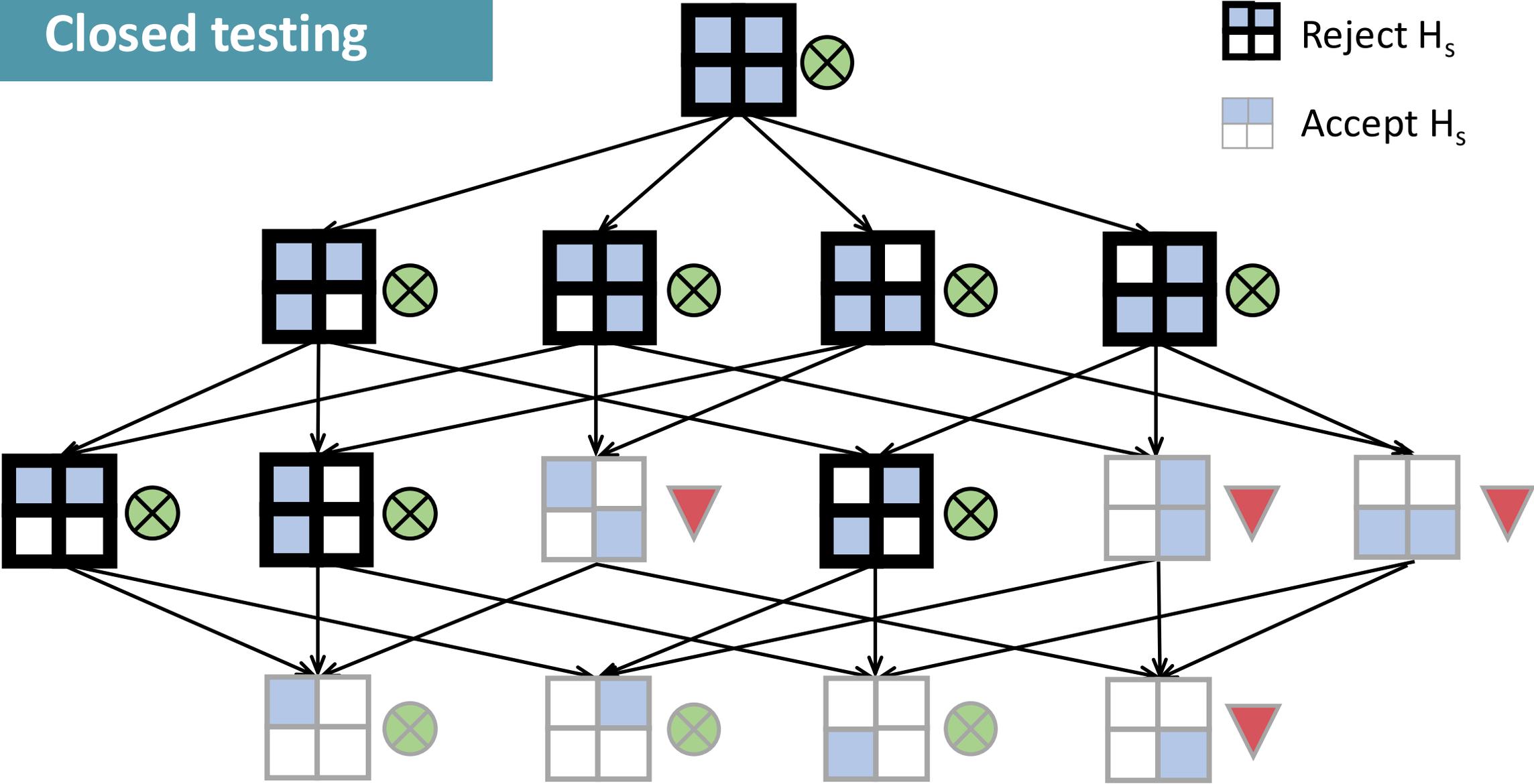
Closed testing

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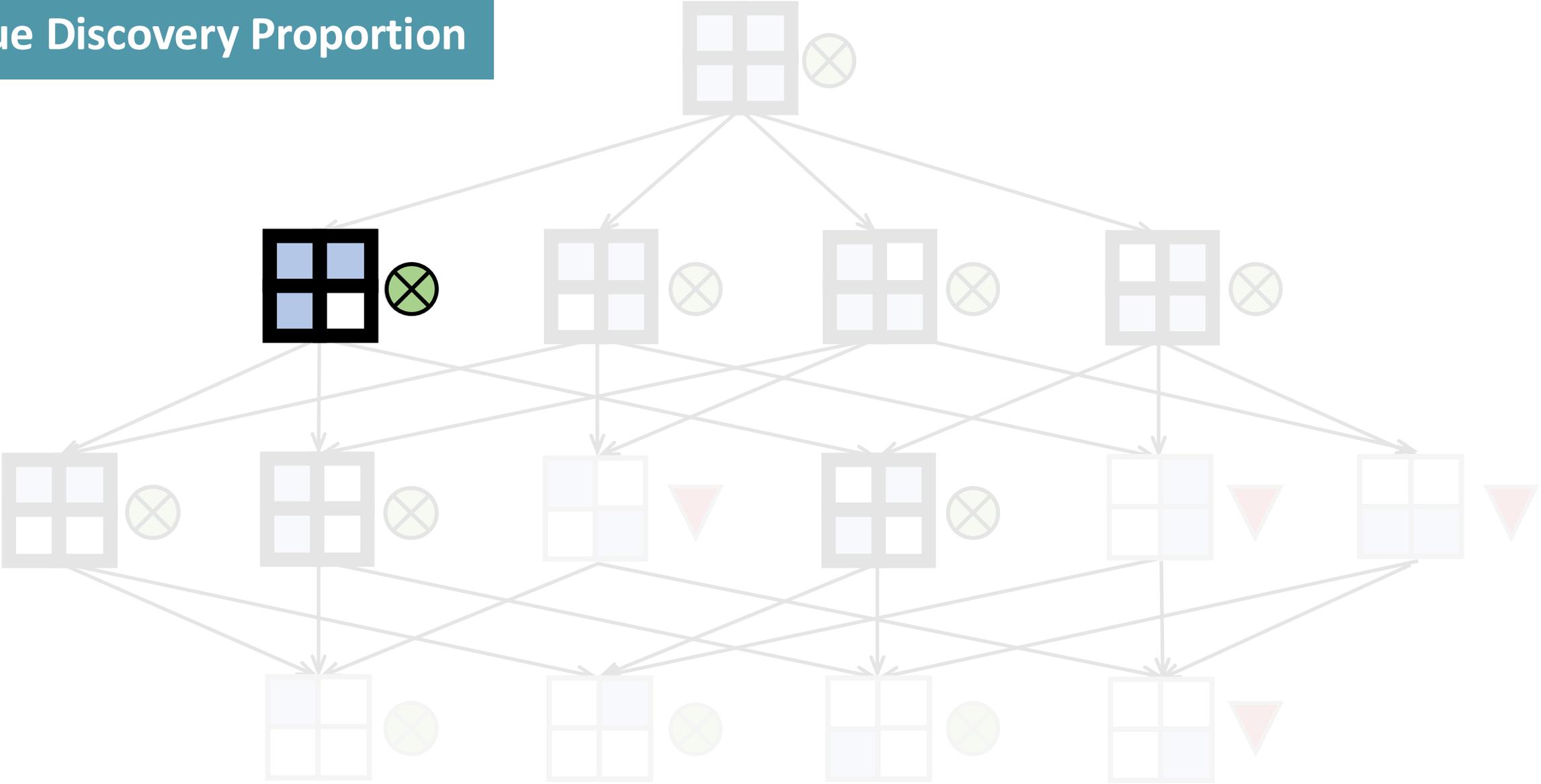


Closed testing

 Reject H_s
 Accept H_s



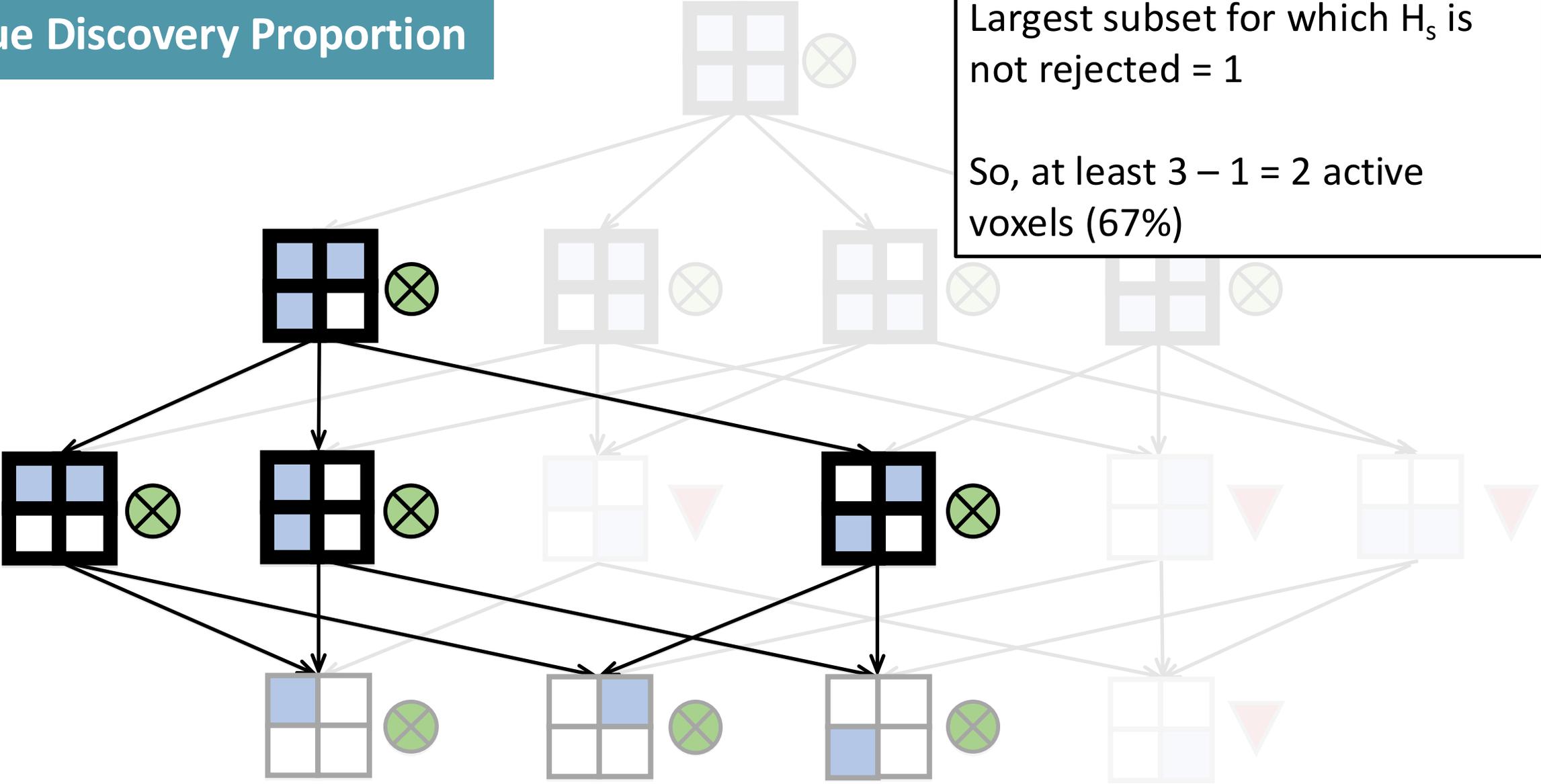
True Discovery Proportion



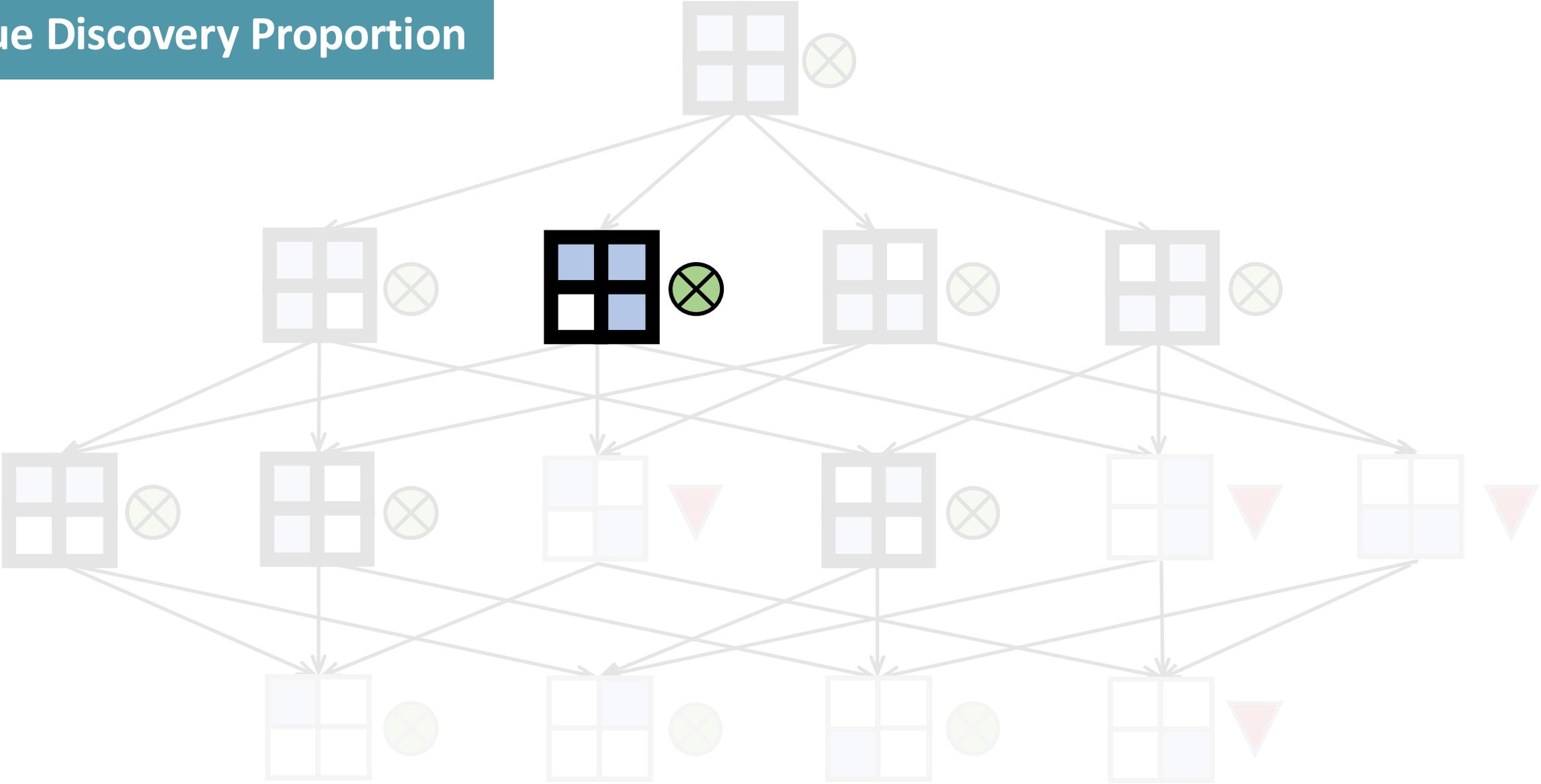
True Discovery Proportion

Largest subset for which H_s is not rejected = 1

So, at least $3 - 1 = 2$ active voxels (67%)

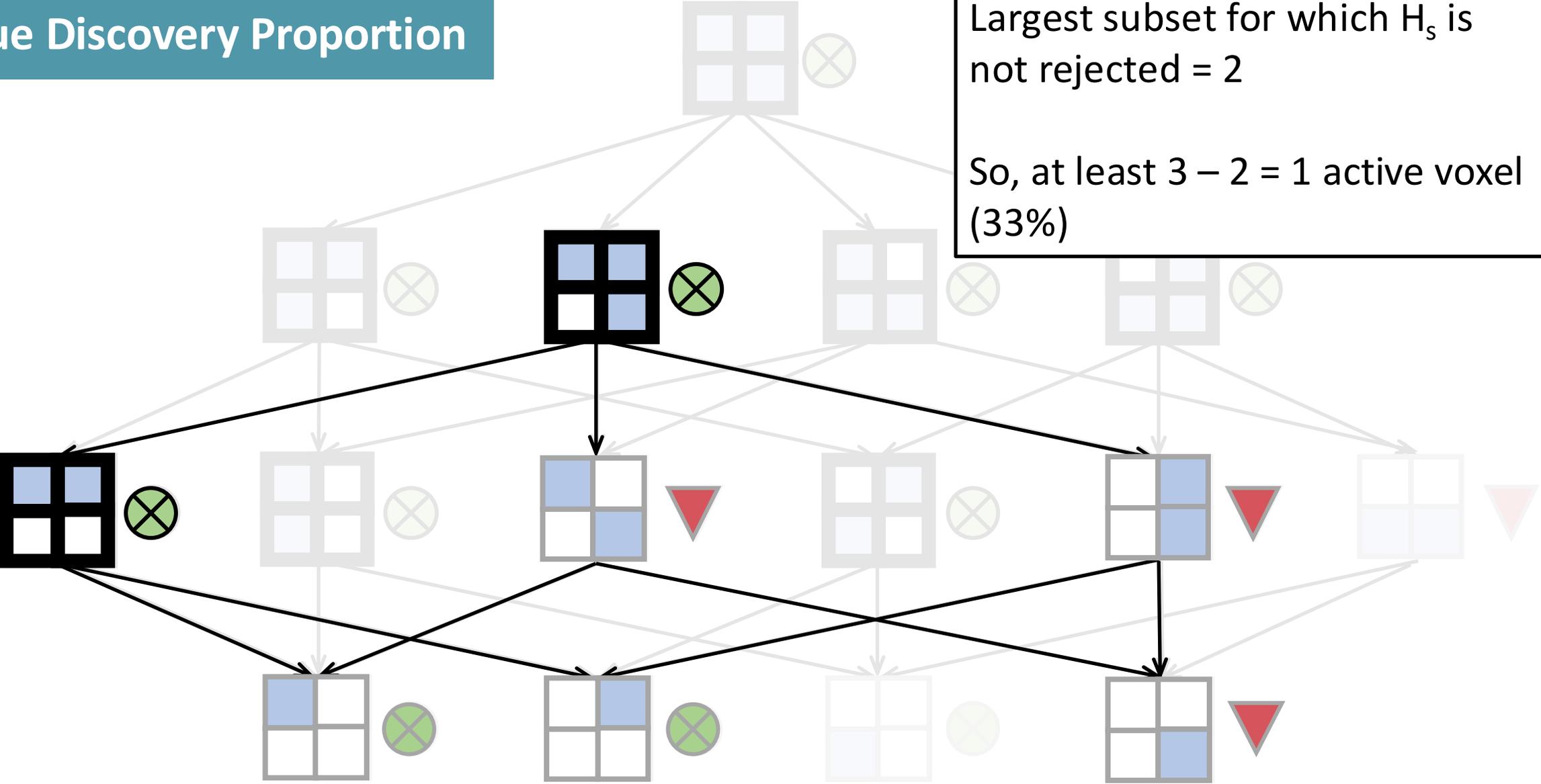


True Discovery Proportion

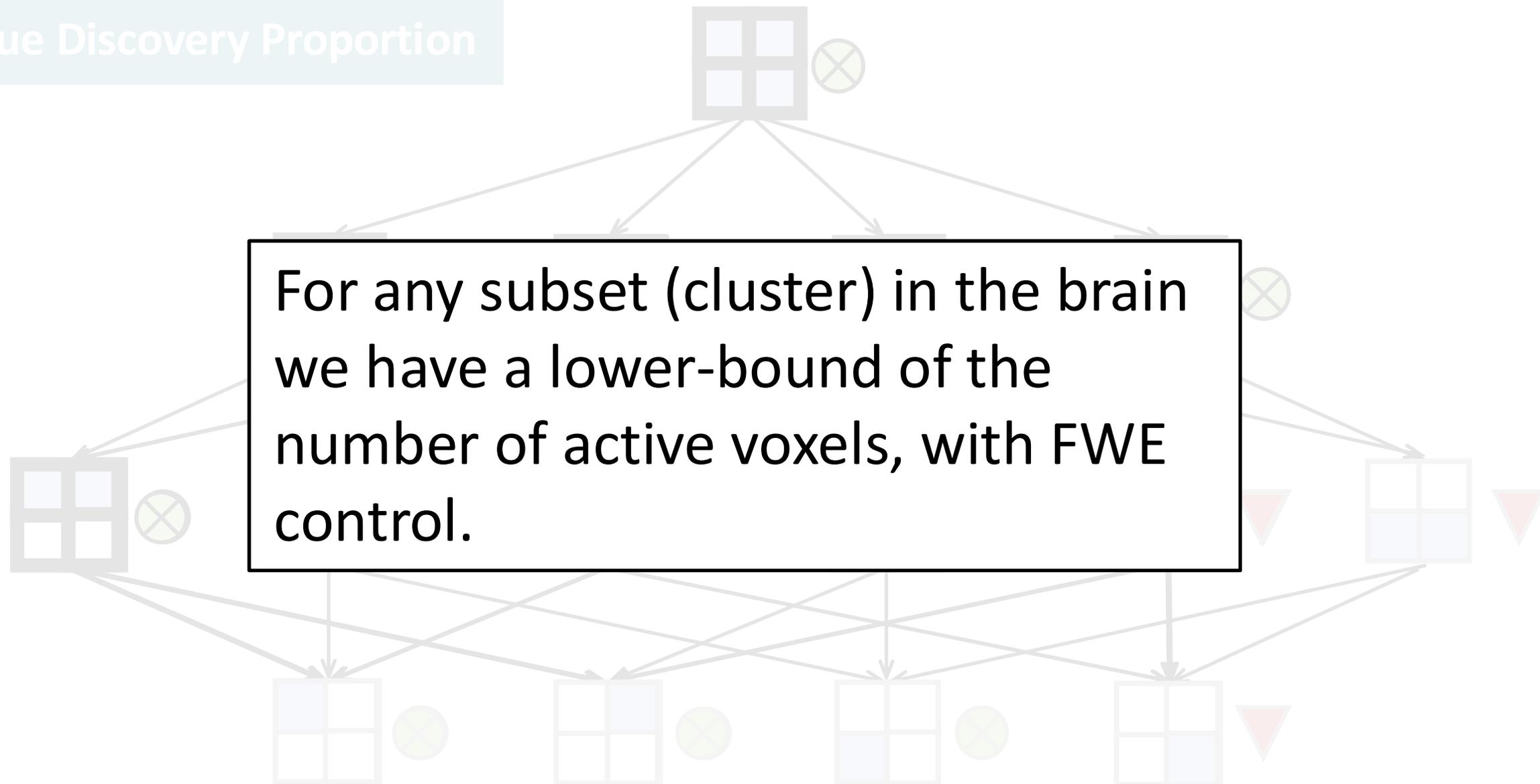


True Discovery Proportion

Largest subset for which H_s is not rejected = 2
So, at least $3 - 2 = 1$ active voxel (33%)



True Discovery Proportion



Scaling up

Ok, this works for 4 voxels, but how about 200,000 voxels?

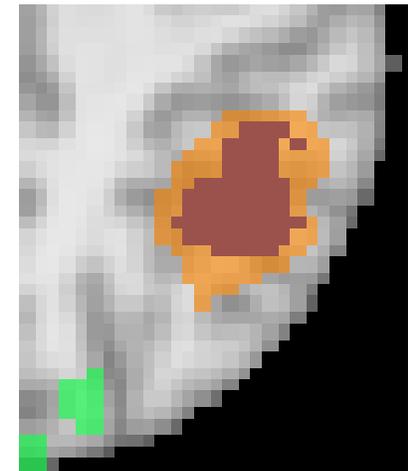
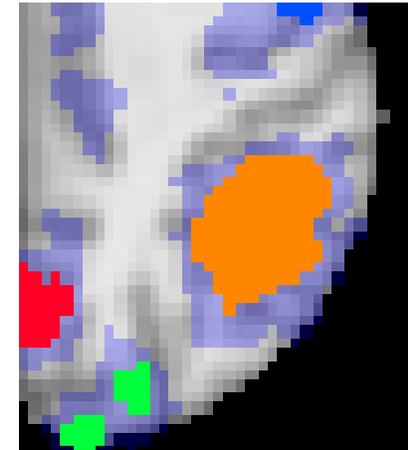
All Resolutions Inference (ARI)

Part 3: The solution

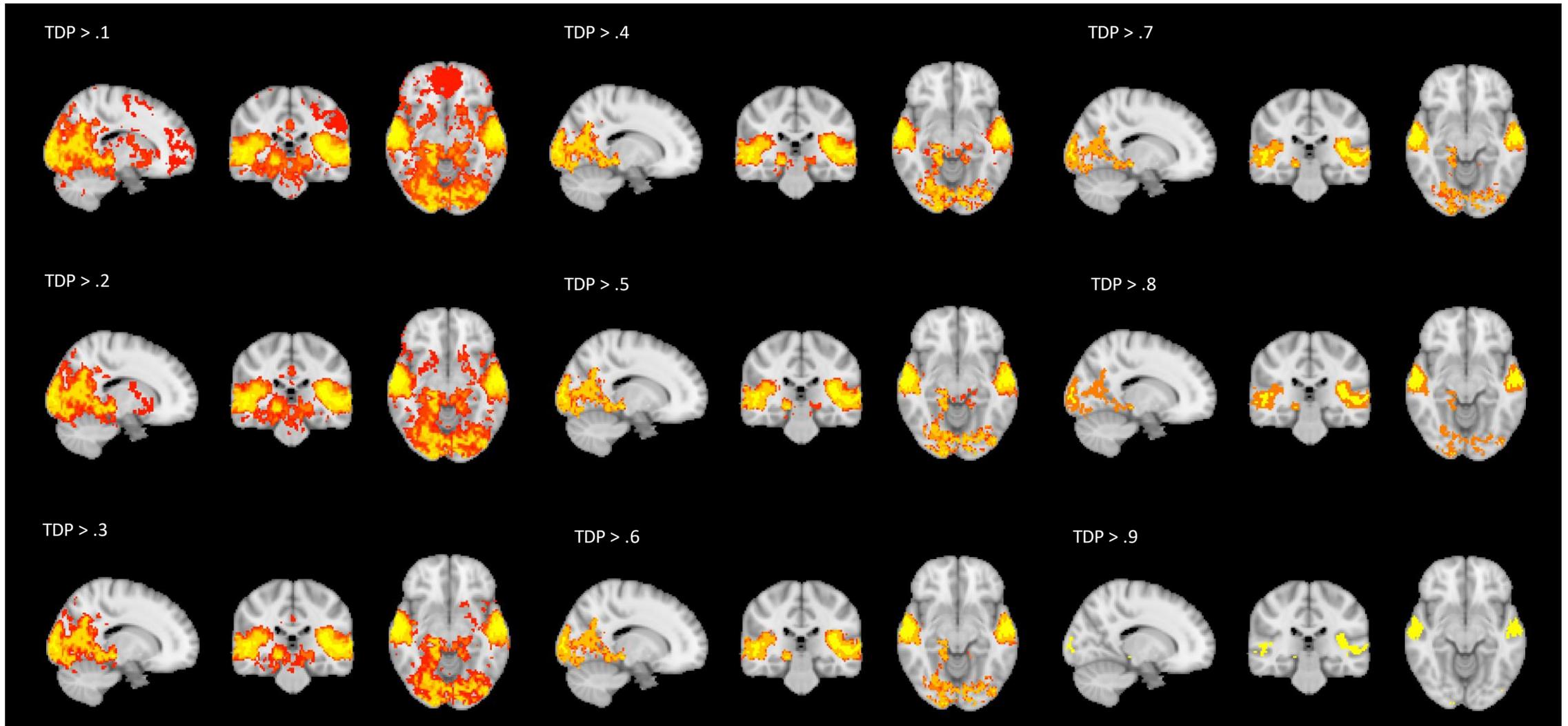
True Discovery Proportion (TDP) based methods

Rosenblatt, Finos, Weeda, Solari & Goeman (2018)

- Multiple testing correction method based on closed testing.
- True Discovery Proportion (TDP) based methods allow us to estimate the number of truly active voxels within a cluster, for all possible clusters, as many times a researcher wants, with full FWER control.
- We can also estimate clusters with at least a certain TDP. For example, “what is the largest cluster that contains at least 60% active voxels (TDP > .6)?”

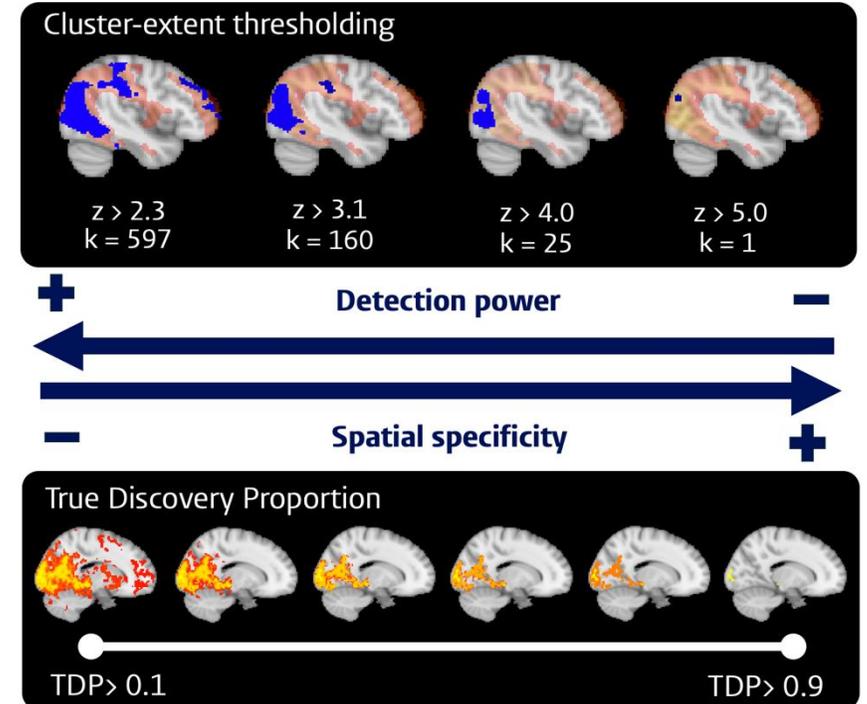


True Discovery Proportion (TDP) based methods



True Discovery Proportion (TDP) based methods

- Both methods are a trade-off between detection and localization, but:
- While cluster extent inference cannot go beyond the cluster-level
- TDP based methods can quantify this trade-off explicitly.



More information

- ARIbrain Github:

<https://github.com/aribrain/ari-core>

- References:

[1] Rosenblatt, J., et al. (2018). All-Resolutions Inference for brain imaging. *Neuroimage*, 181, 786-796.

[2] Andreella, A., et al. (2023). Permutation-based true discovery proportions for fMRI cluster analysis. *Statistics in Medicine*, 42(14): 2311-2340.

[3] Goeman, J.J., et al. (2023). Cluster extent inference revisited: quantification and localization of brain activity. *J Roy Stat Soc B*, 85(4), 1128–1153.