

ARLbrain - Valid circular inference for fMRI analysis

Wouter Weeda, Leiden University



Universiteit
Leiden

Thanks to:

Xu Chen

Martha van Kempen

Jonathan Rosenblatt

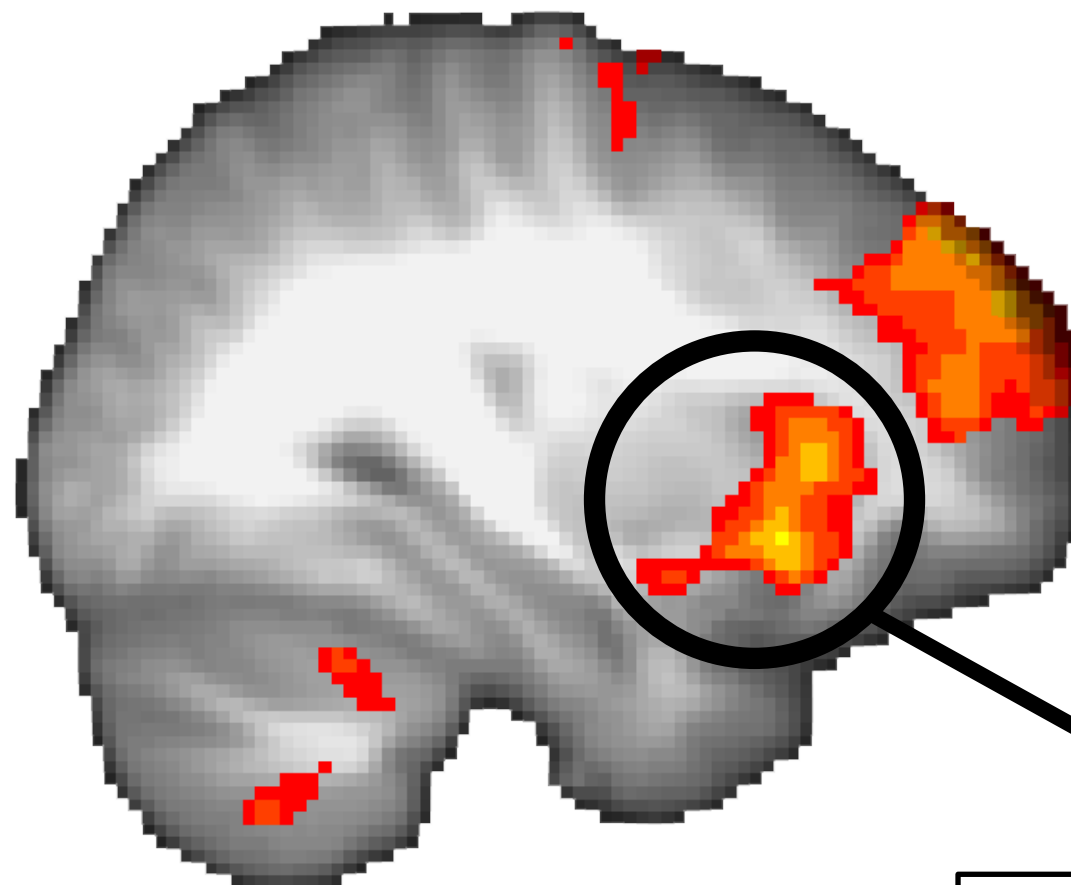
Livio Finos

Aldo Solari

Jelle Goeman

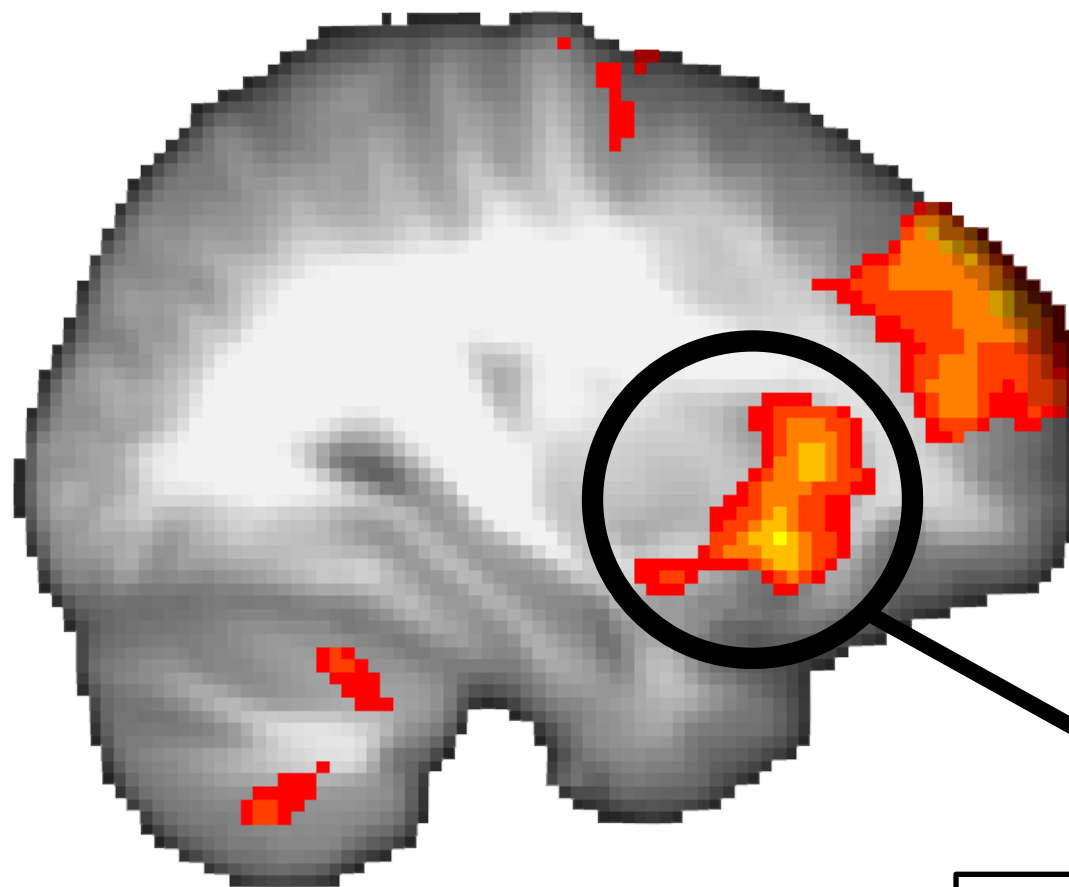


fMRI cluster inference



All voxels in this cluster are active

fMRI cluster inference



At least one voxel in
this cluster is active

Spatial specificity paradox



“The larger the cluster we find, the less we know about activity within that cluster.”

Goal

- Increase the spatial specificity of cluster inference by:
 - Estimating the number of truly active voxels of a cluster
 - For any cluster, as many times as you want.
 - With Familywise Error (FWE) control



How?

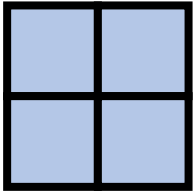
- In of stead asking:
 - Give us a dataset, and we'll tell you which regions are active.
- We ask:
 - Give us a region, and we'll tell you how much activity is in there.



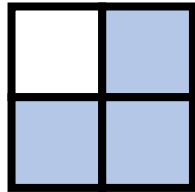
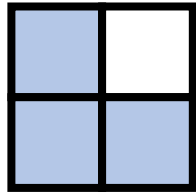
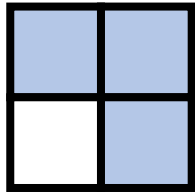
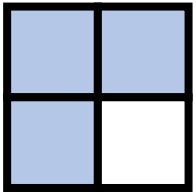
The four voxel brain

.026	.029
.031	.207

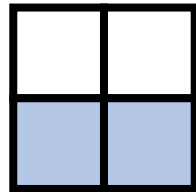
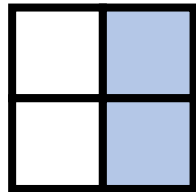
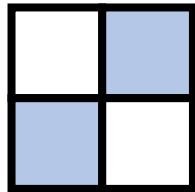
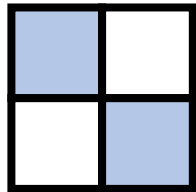
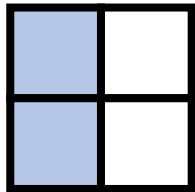
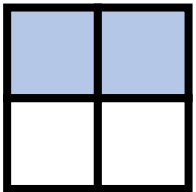
The four voxel brain - subsets



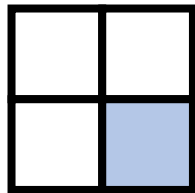
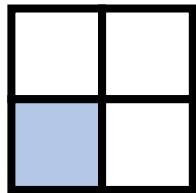
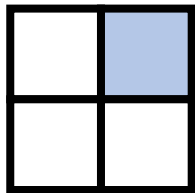
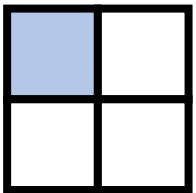
4 voxels set



3 voxels sets

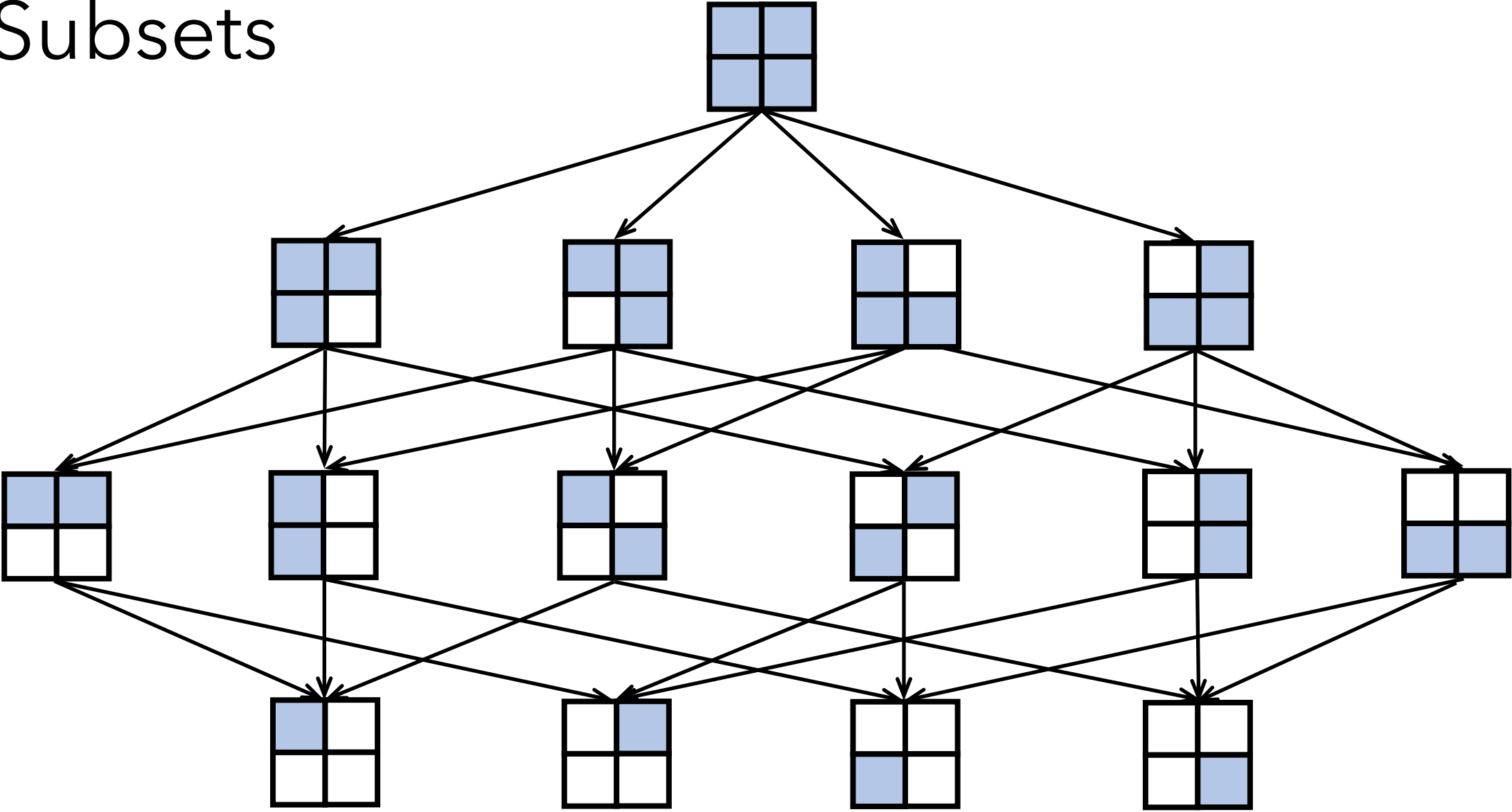


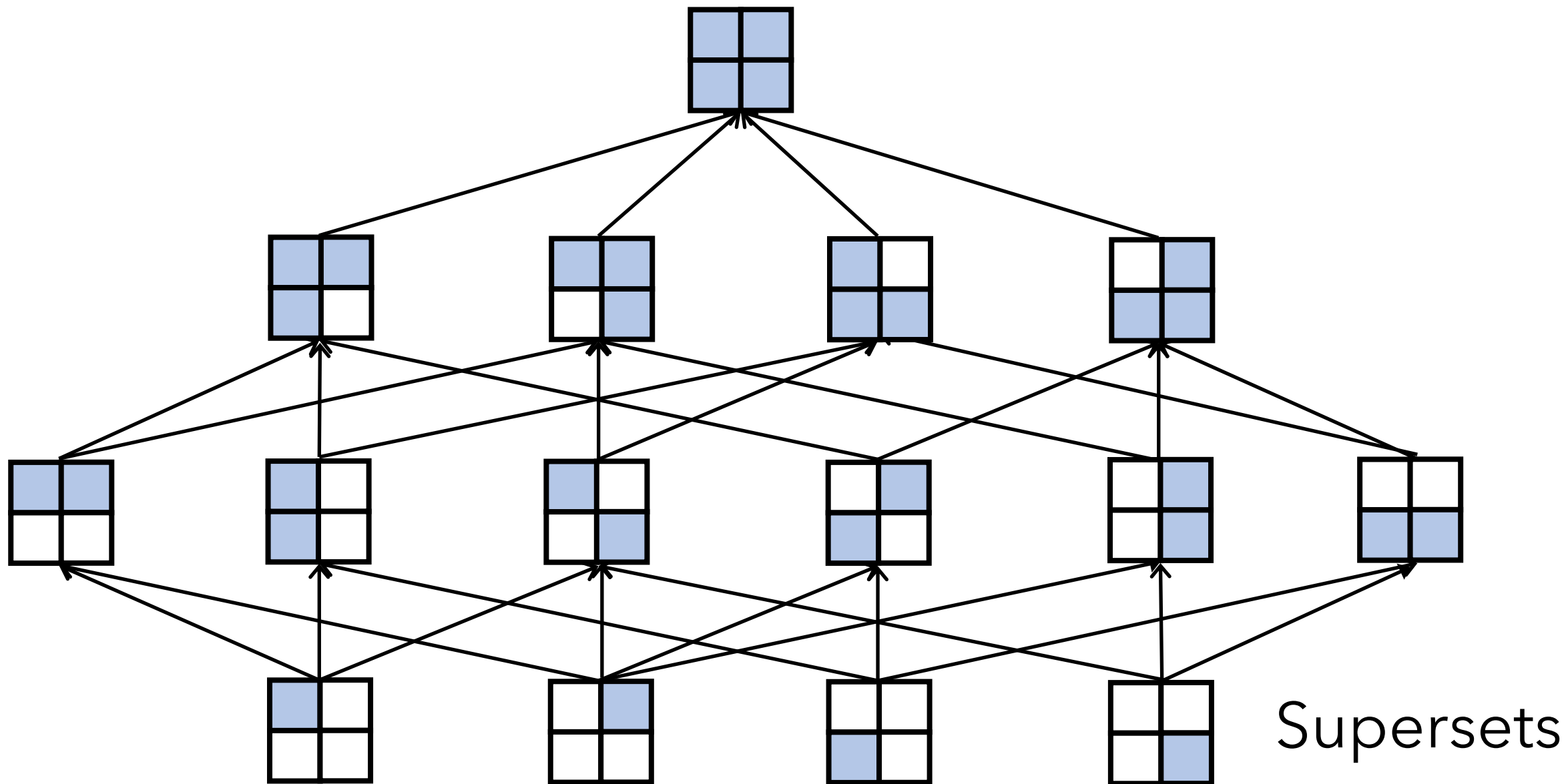
2 voxels sets



1 voxel sets

Subsets





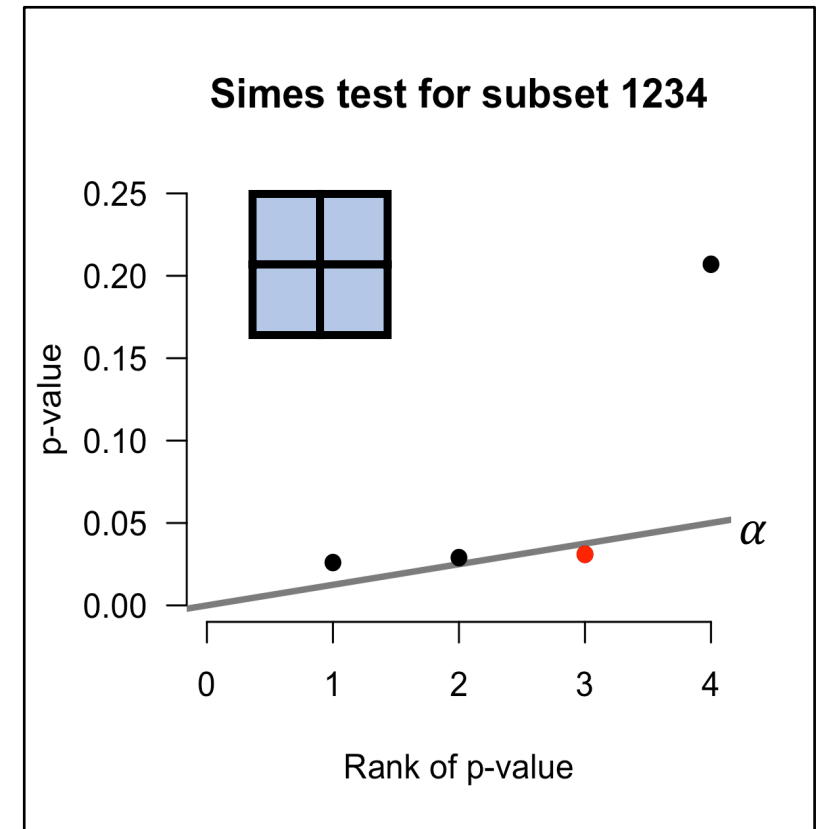
Simes test

Is there at least one active voxel in subset S ?

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

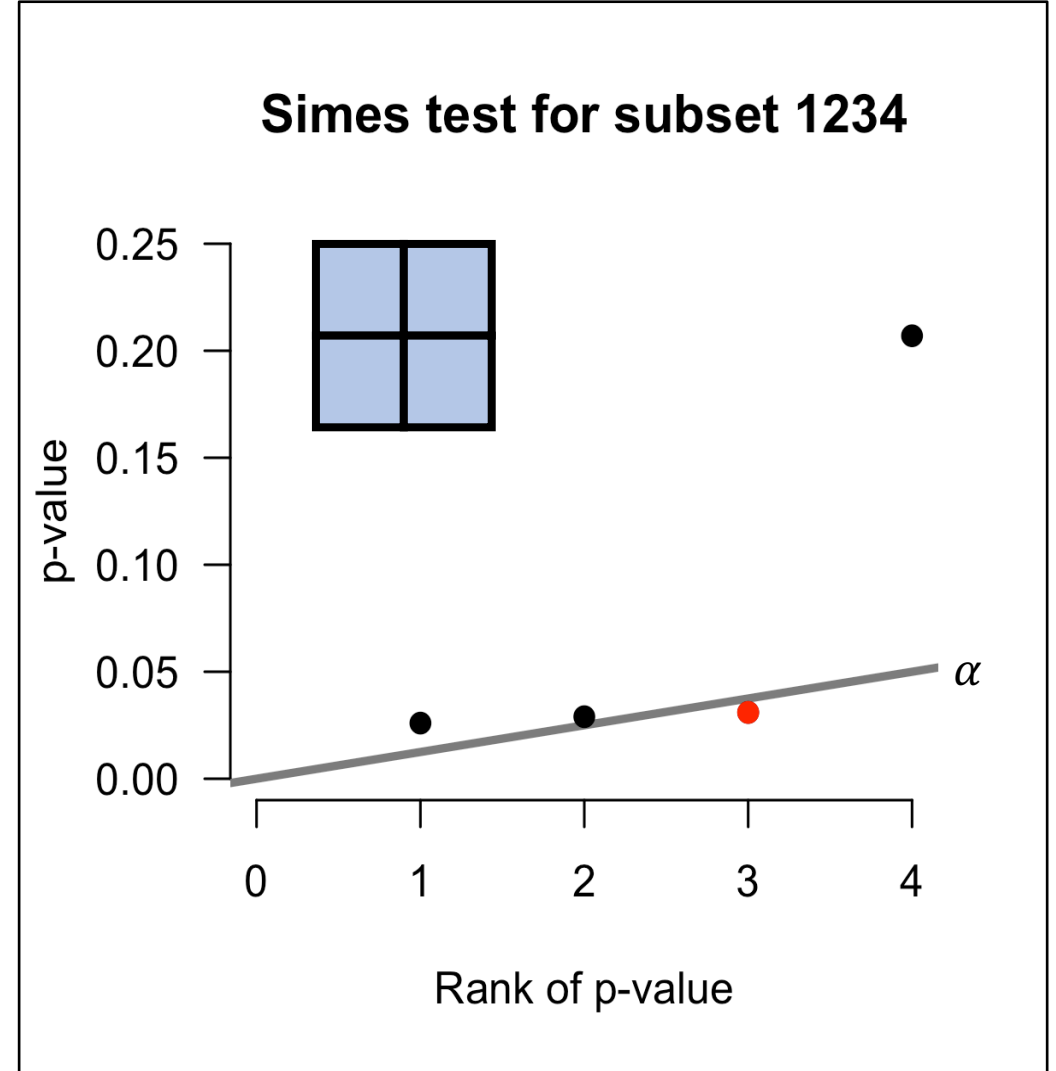
$$p_S = \min \left\{ \frac{|S|}{i} p(i:S), \text{ with } 1 \leq i \leq |S| \right\}$$

Reject H_S if $p_S \leq \alpha$.

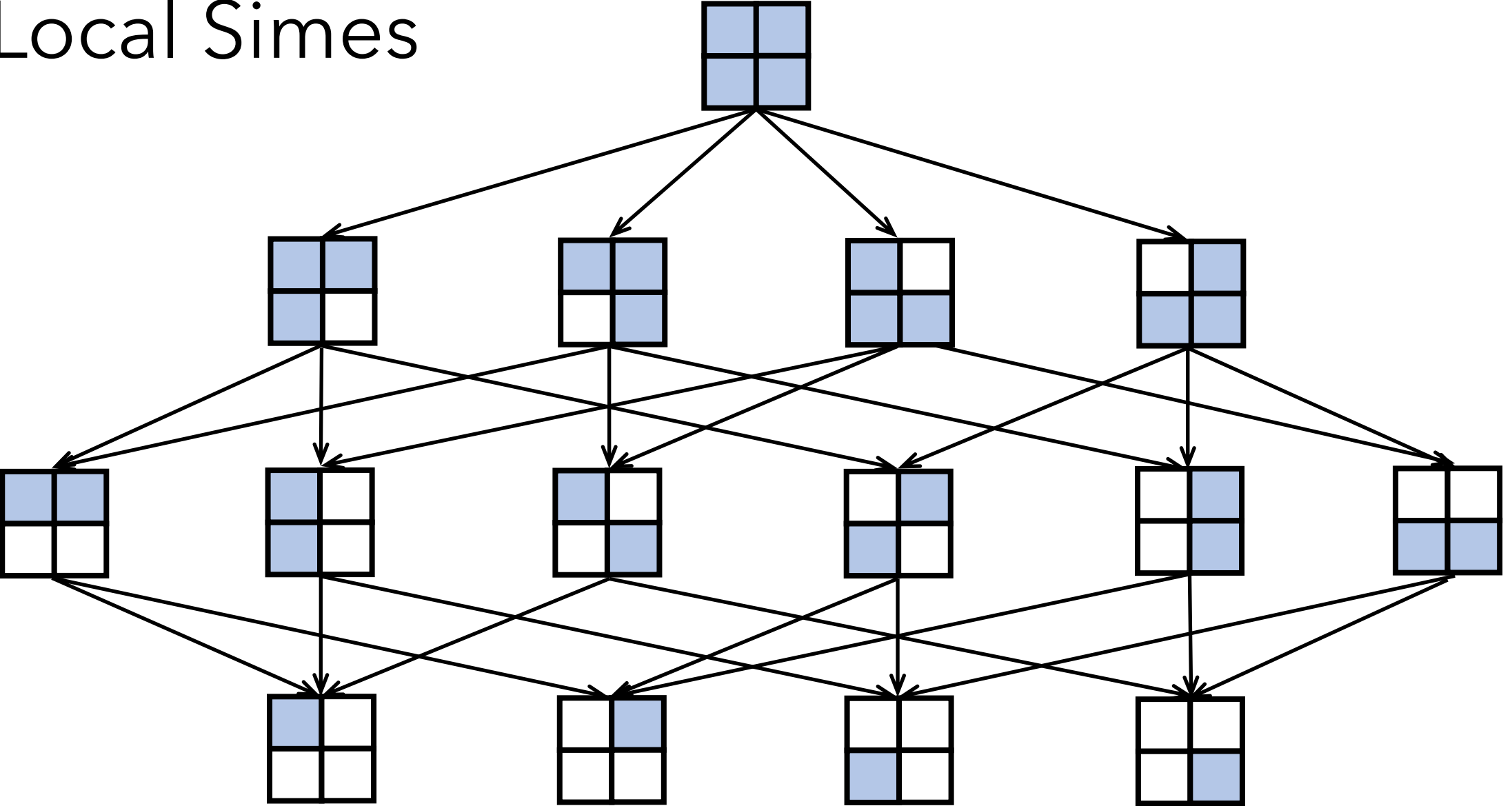


Simes test

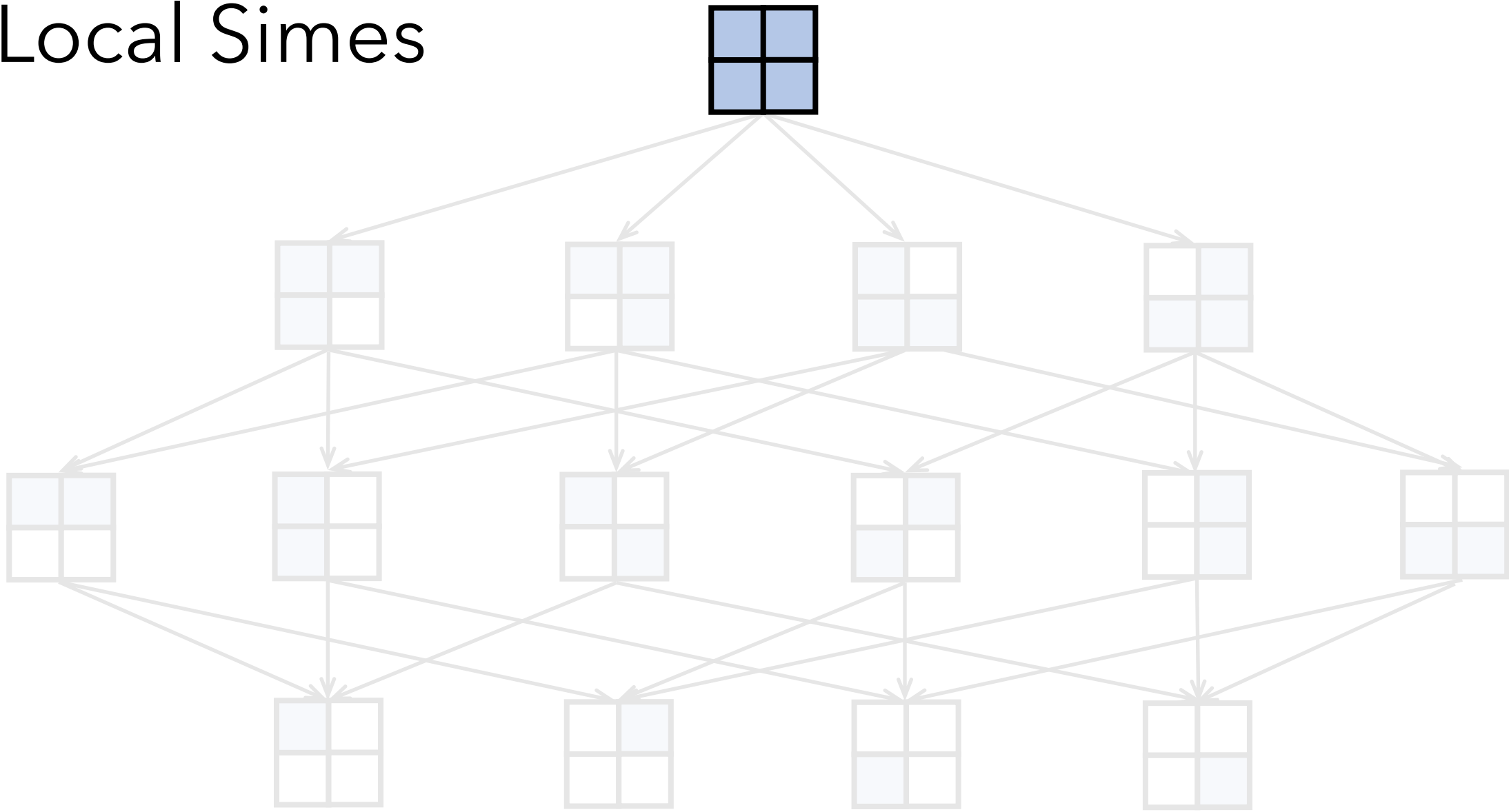
- Order p-values from smallest to largest.
- Draw a line from 0 to alpha.
- Is there any point below the line?



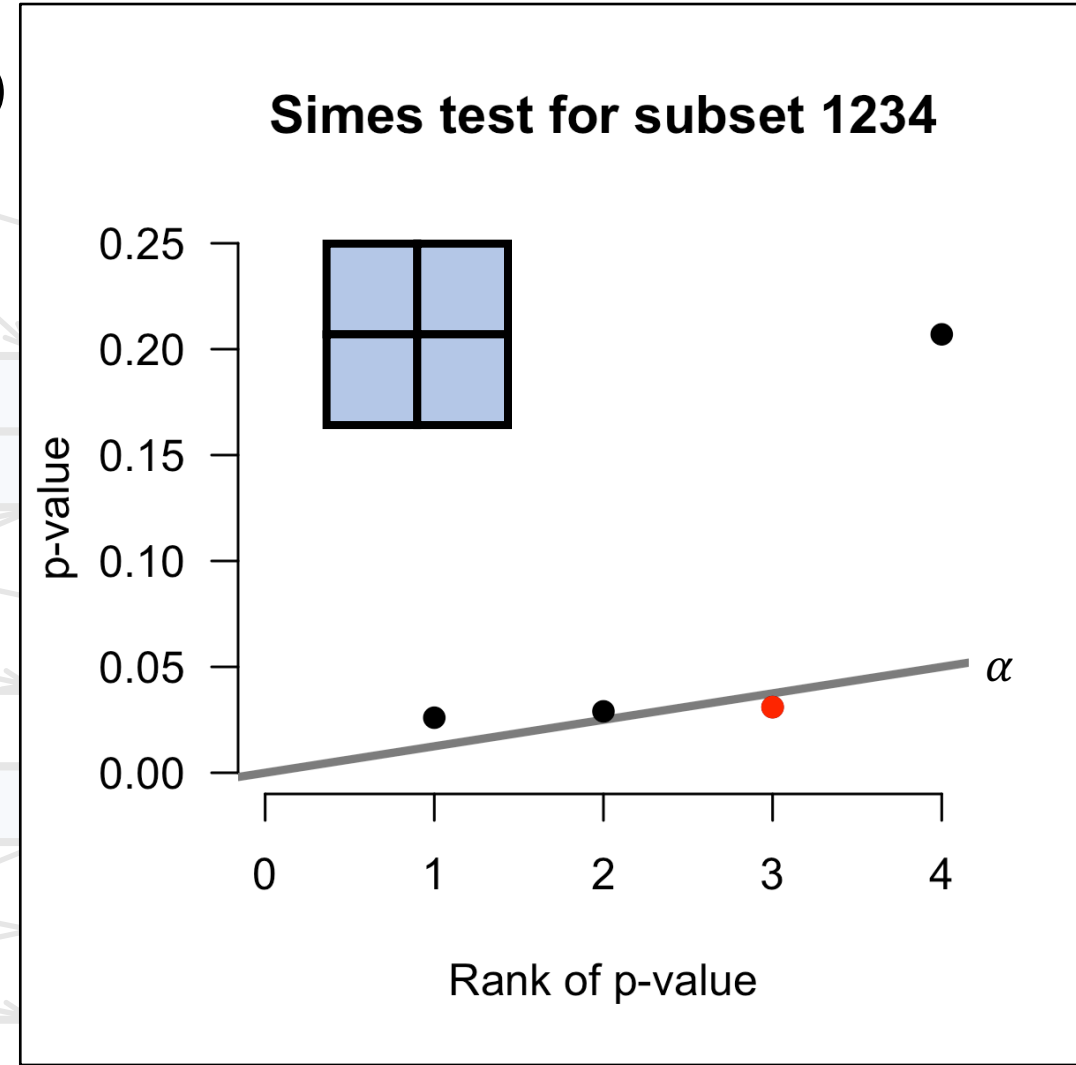
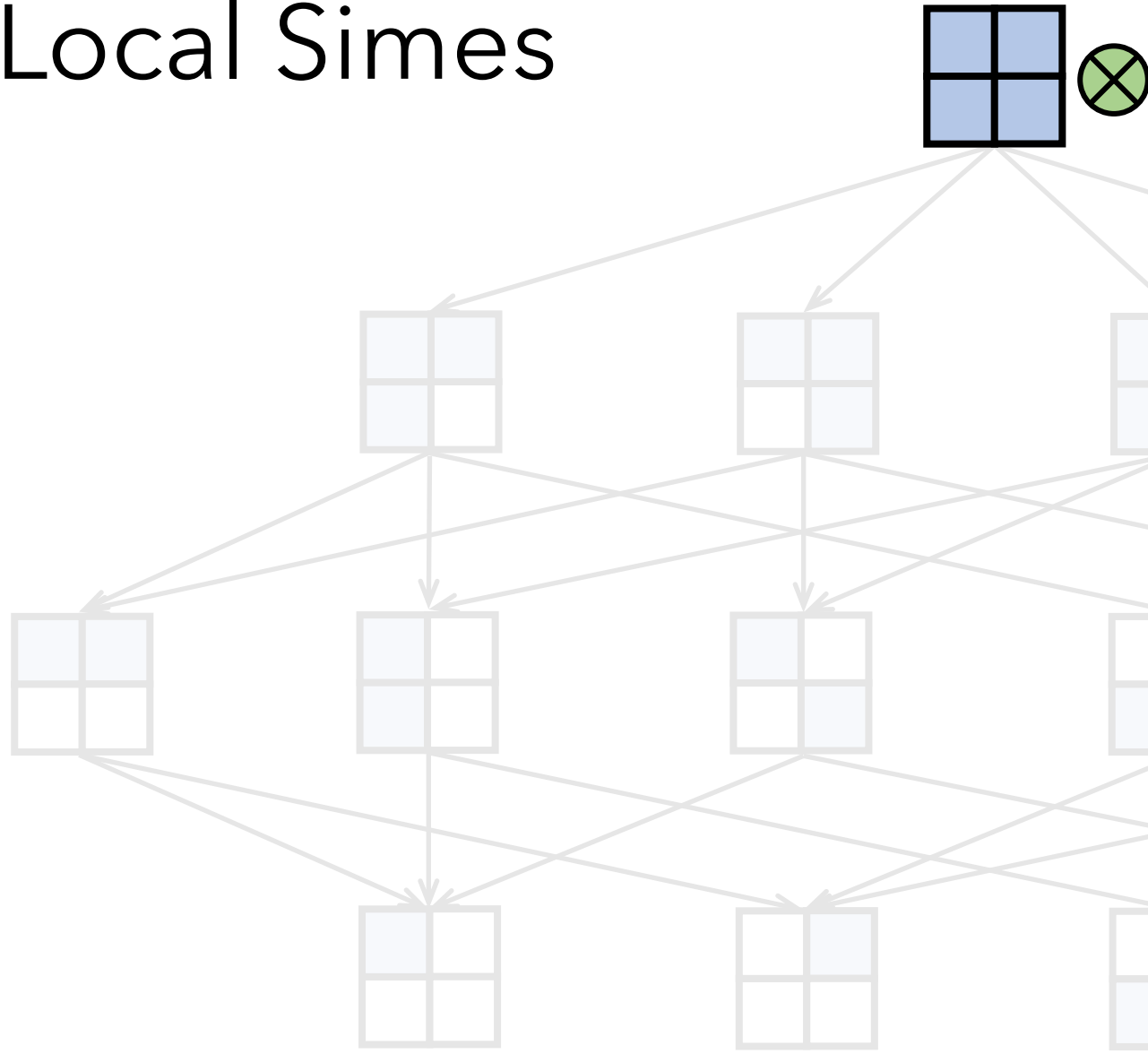
Local Simes



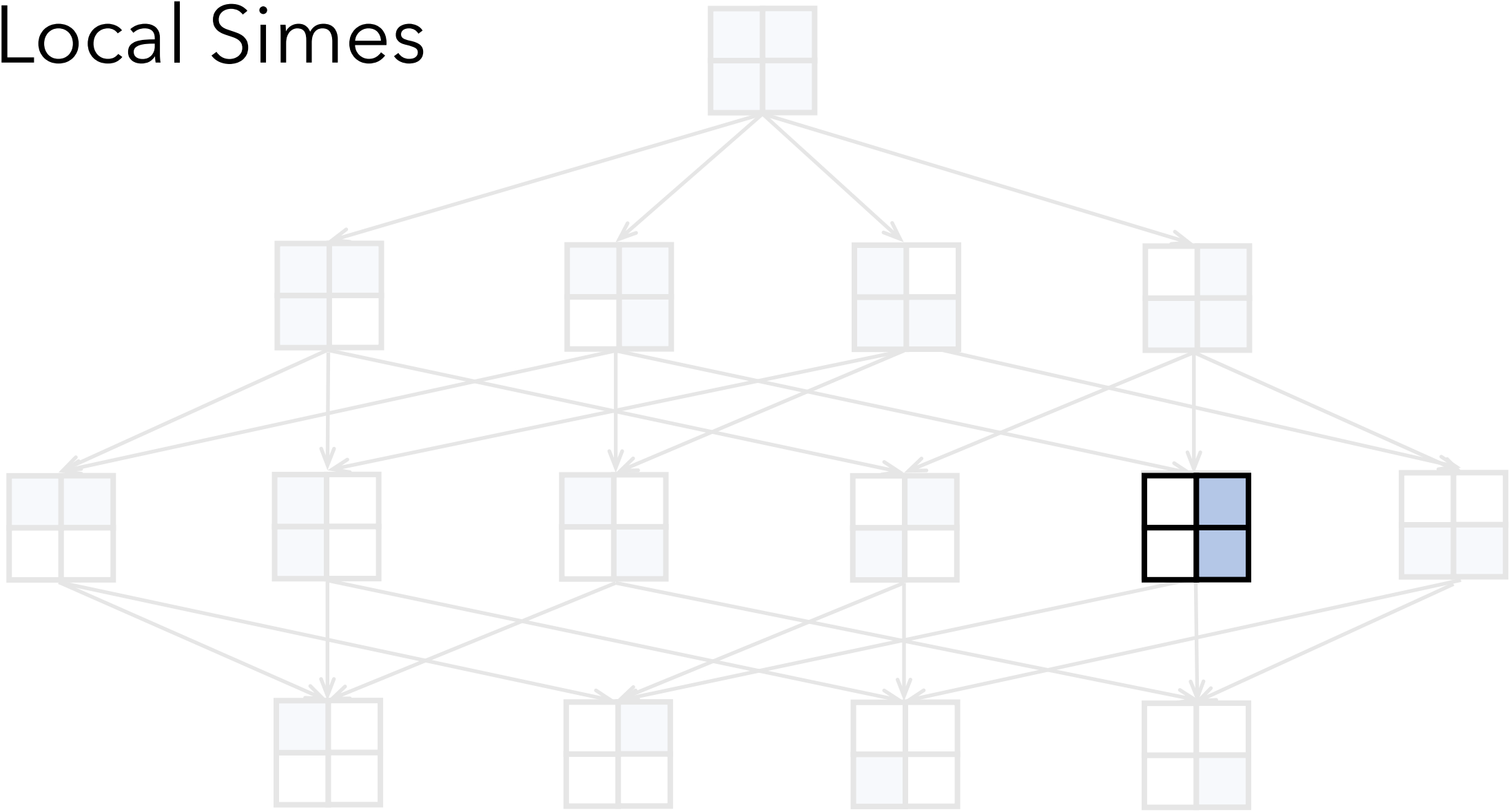
Local Simes



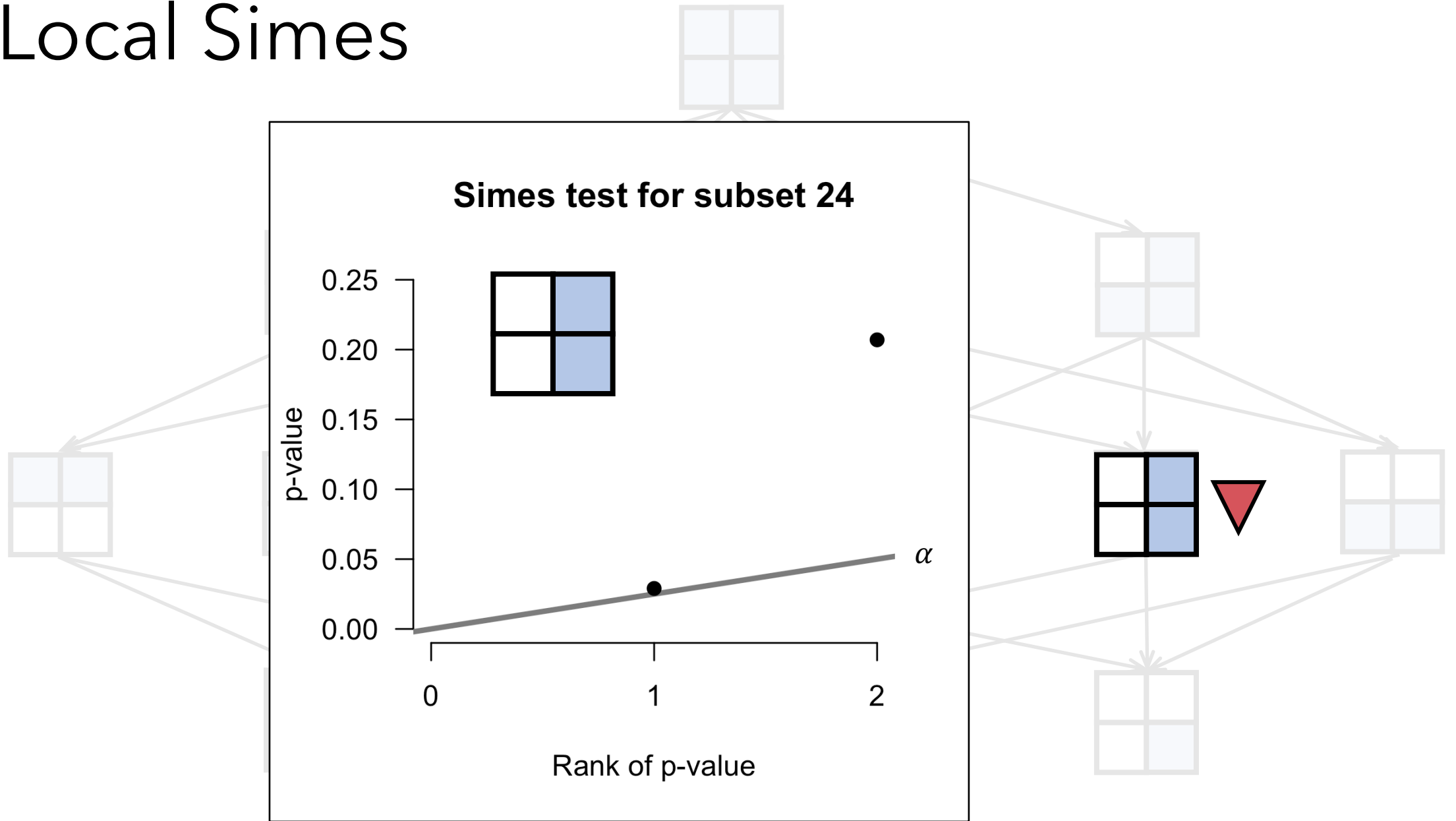
Local Simes



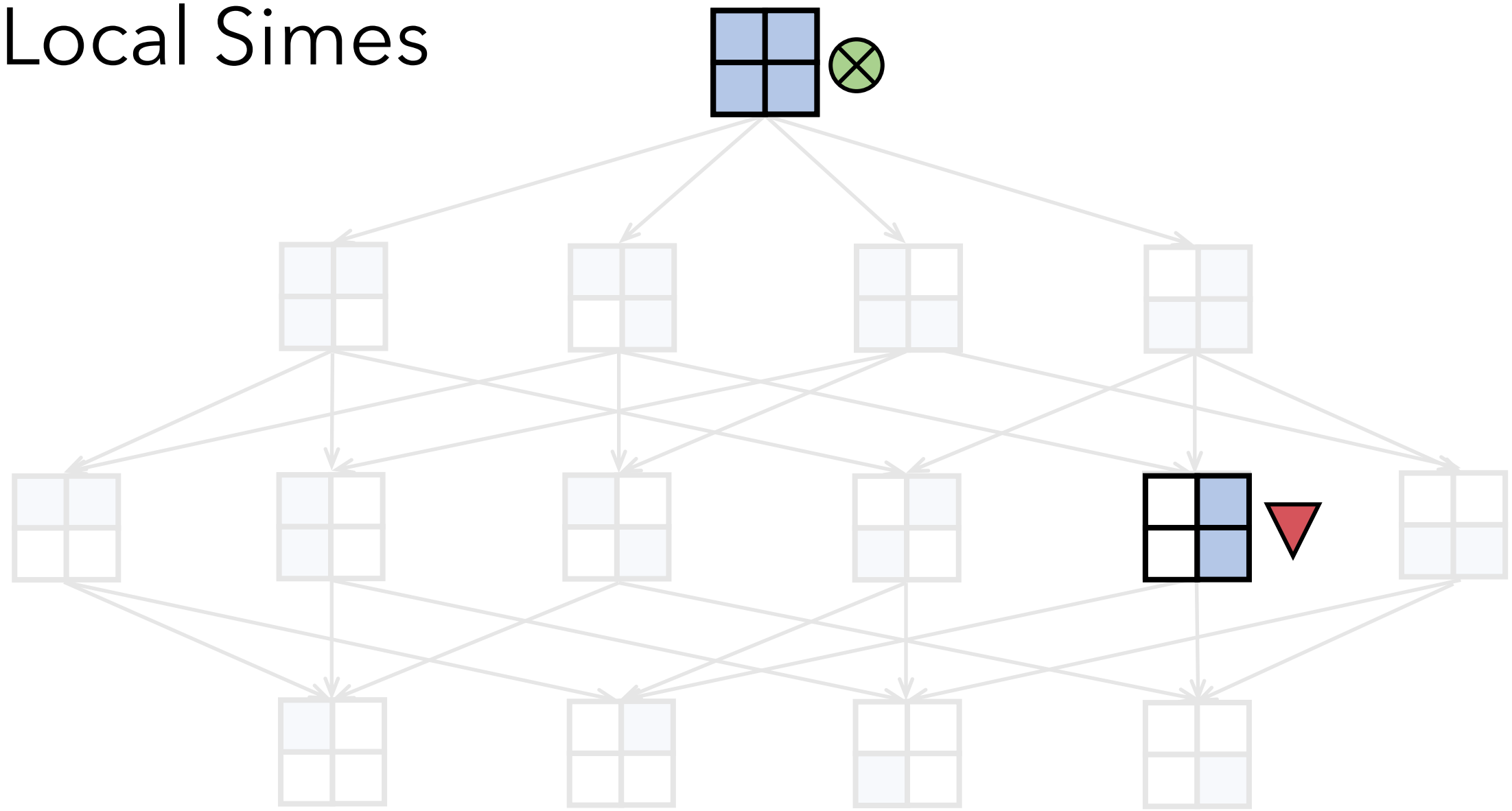
Local Simes



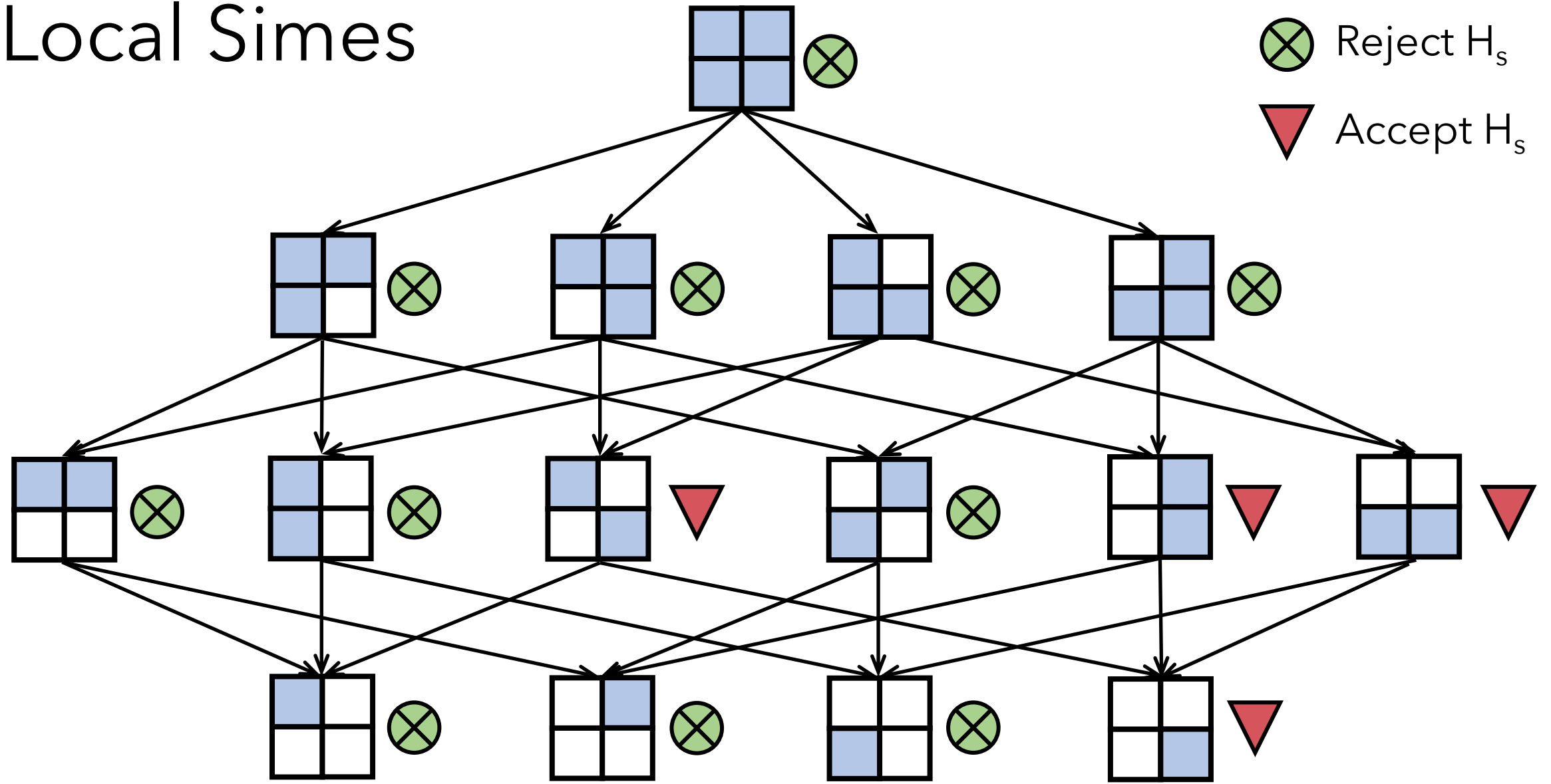
Local Simes



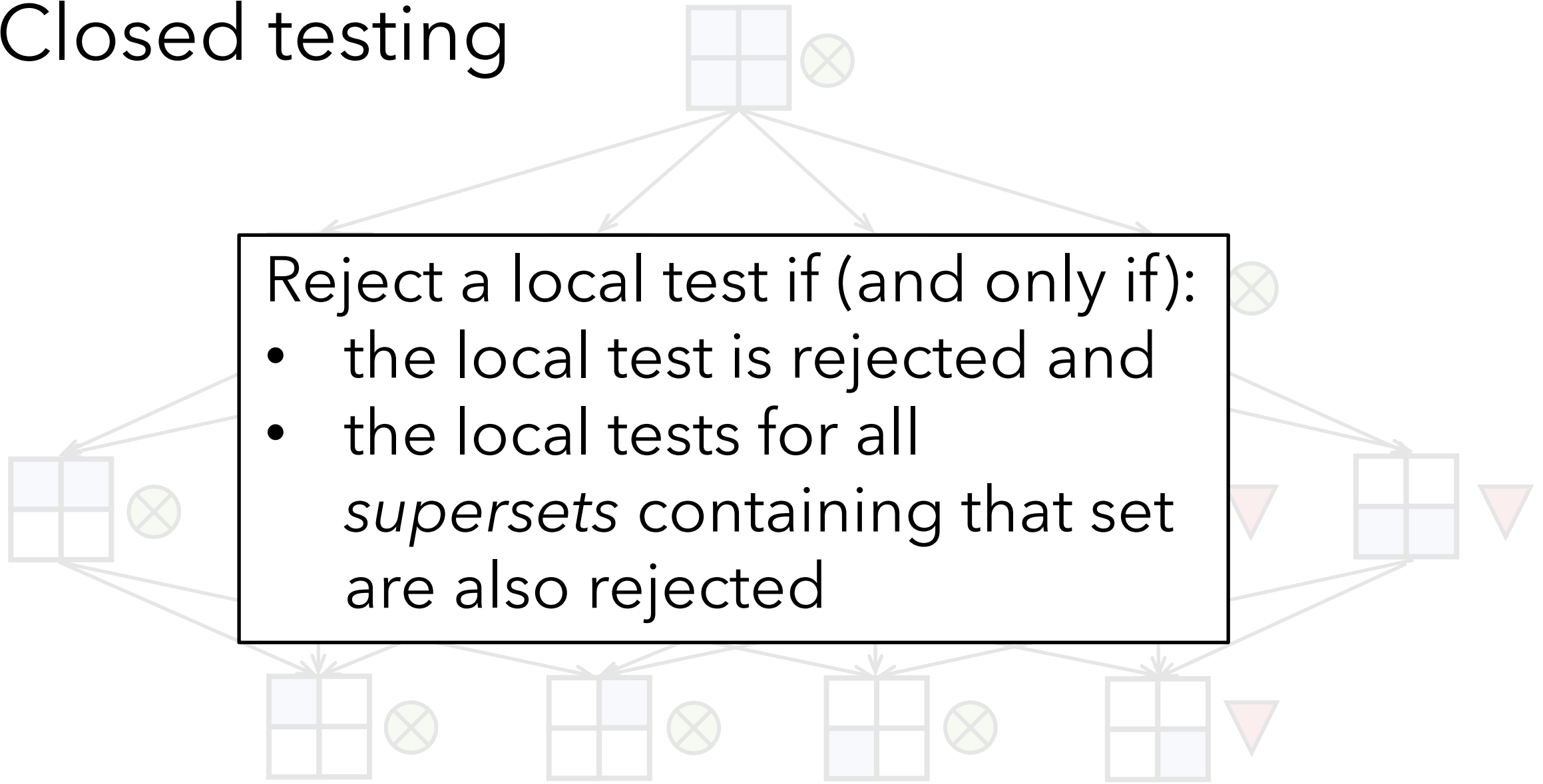
Local Simes



Local Simes



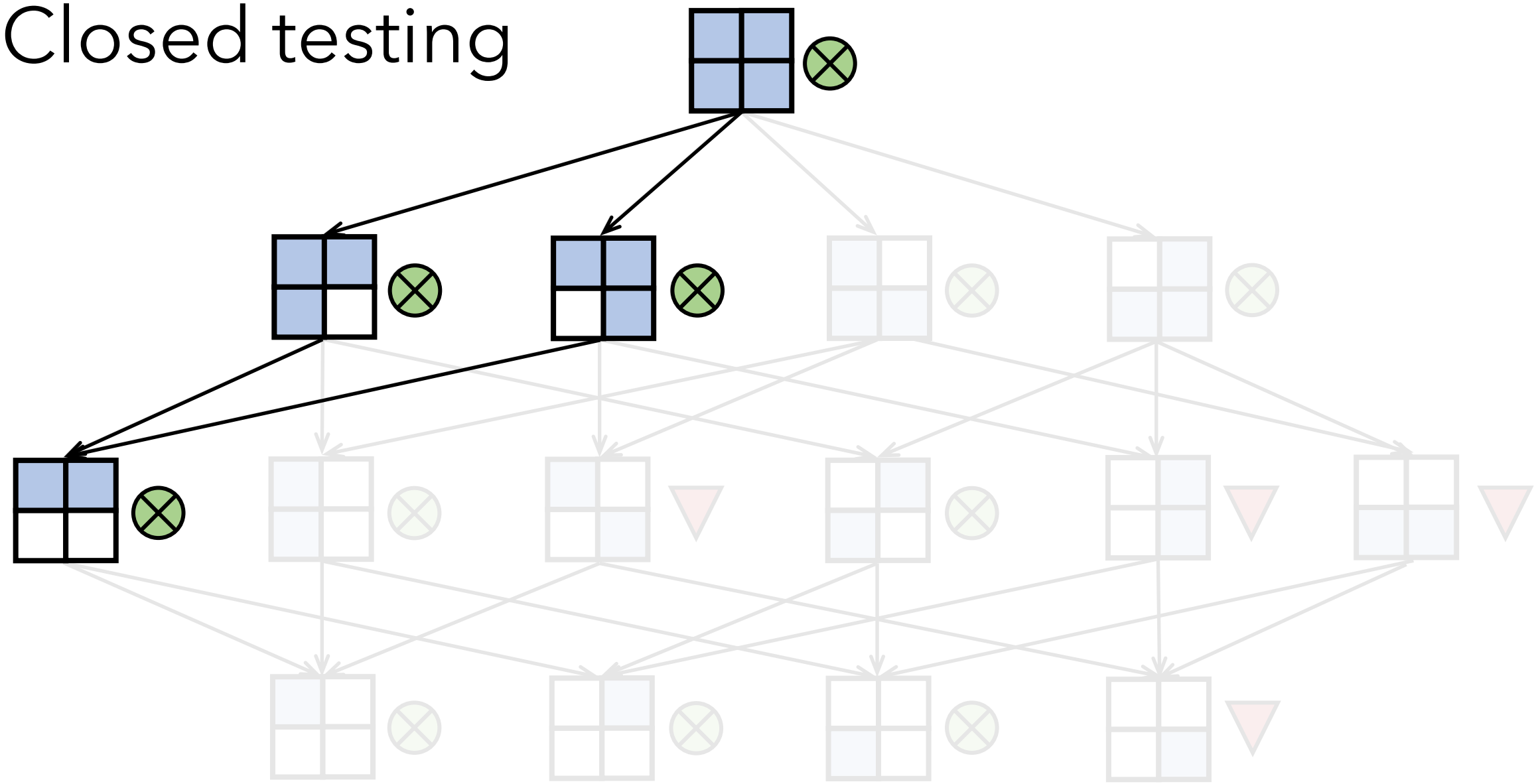
Closed testing



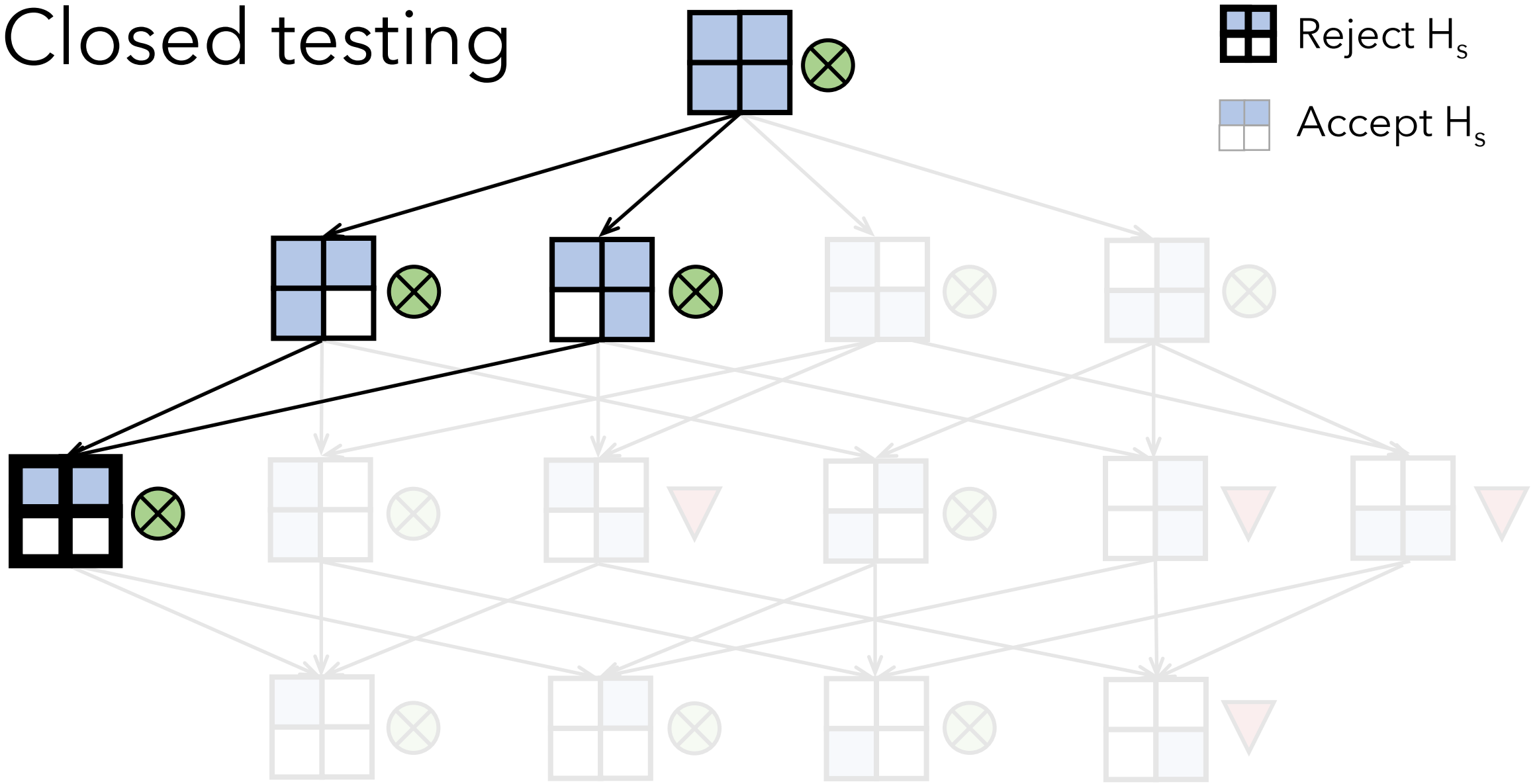
Closed testing



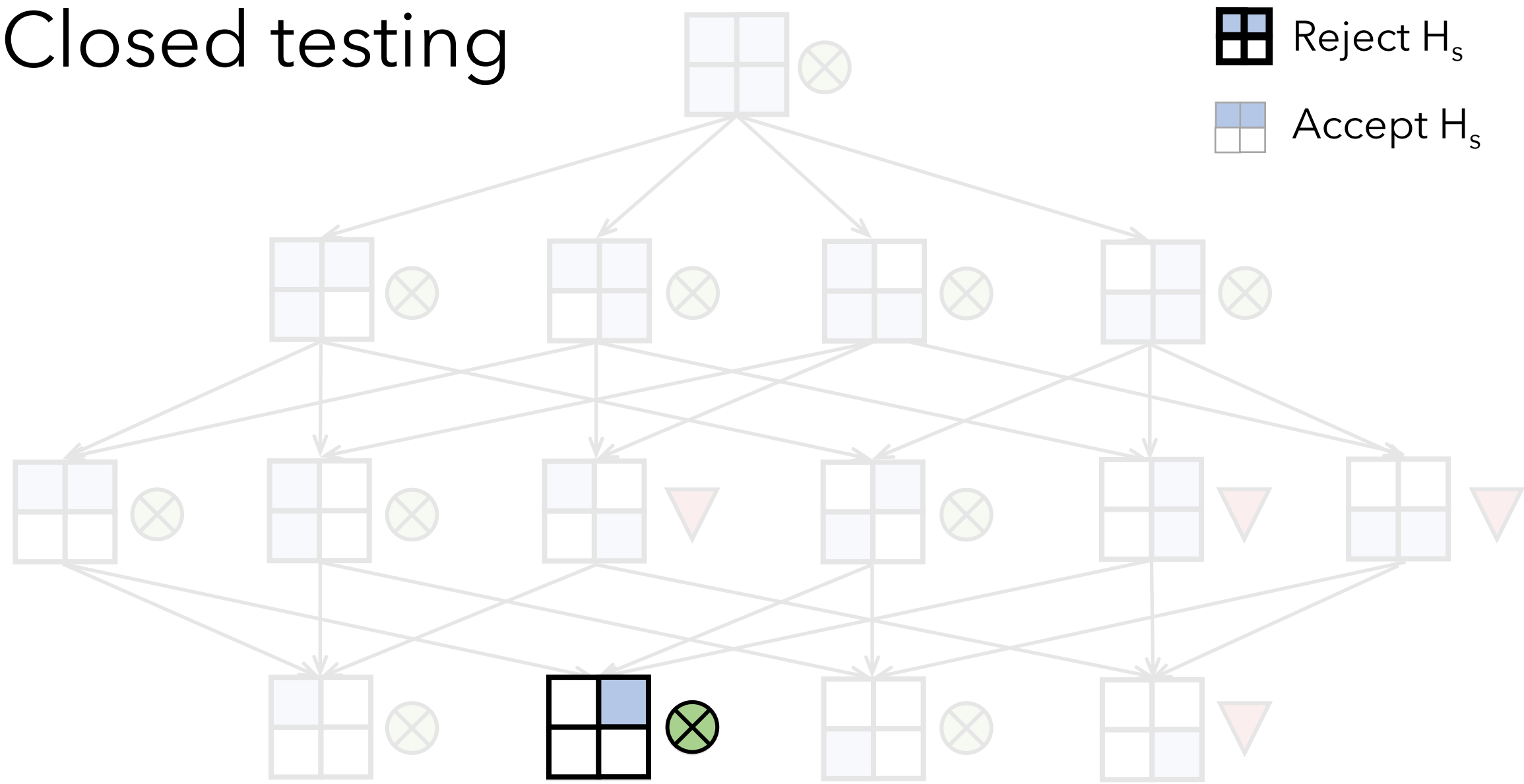
Closed testing



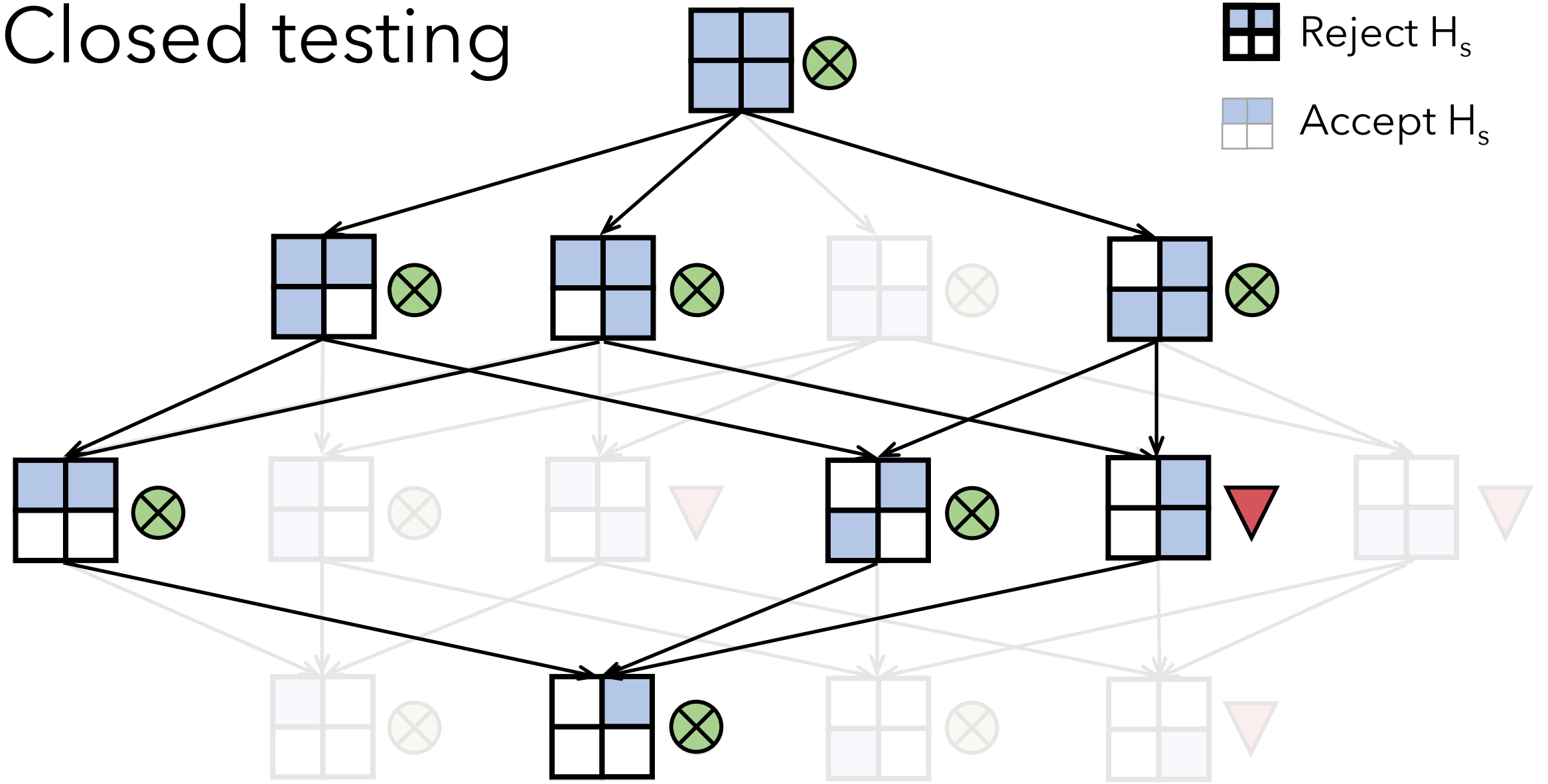
Closed testing



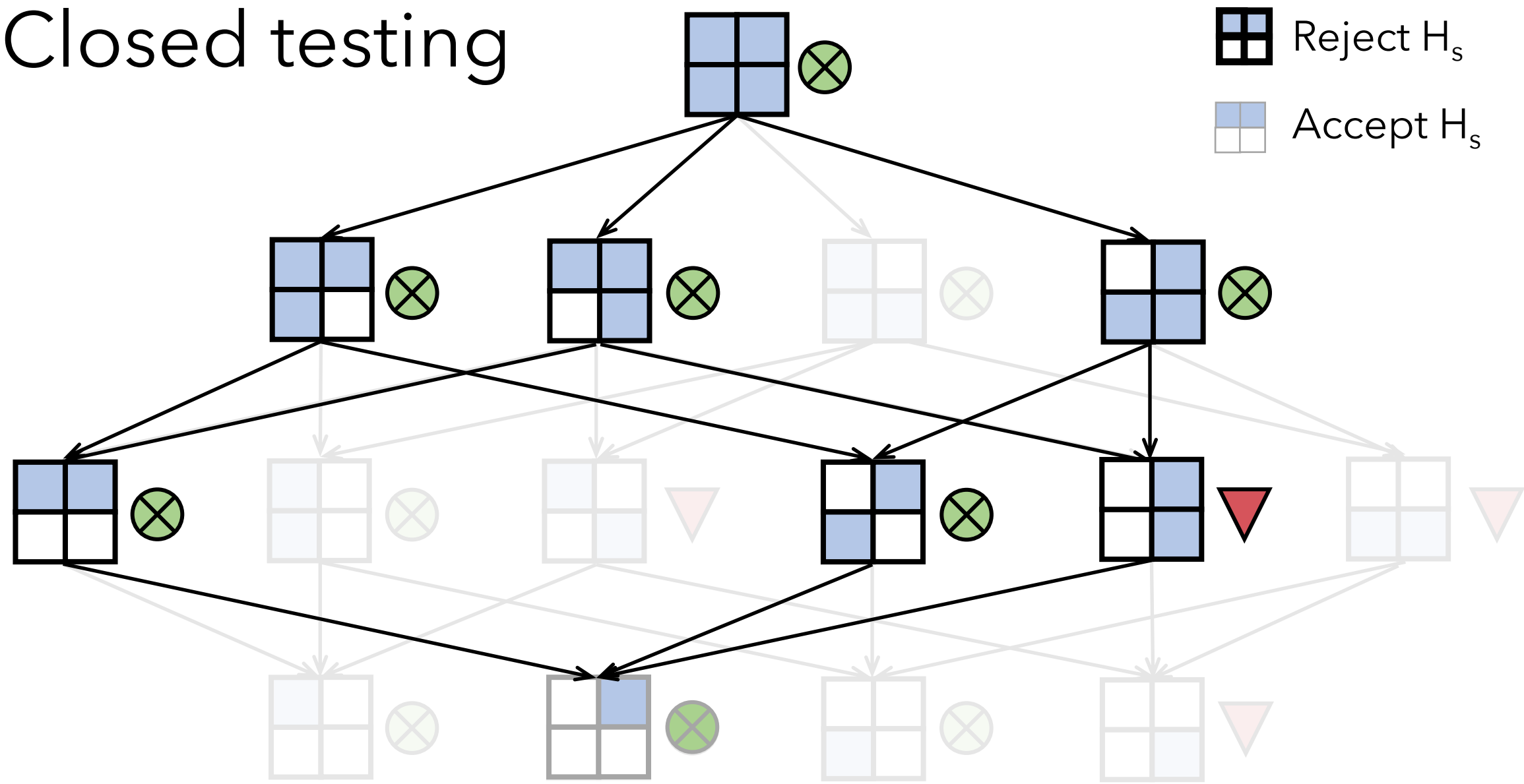
Closed testing



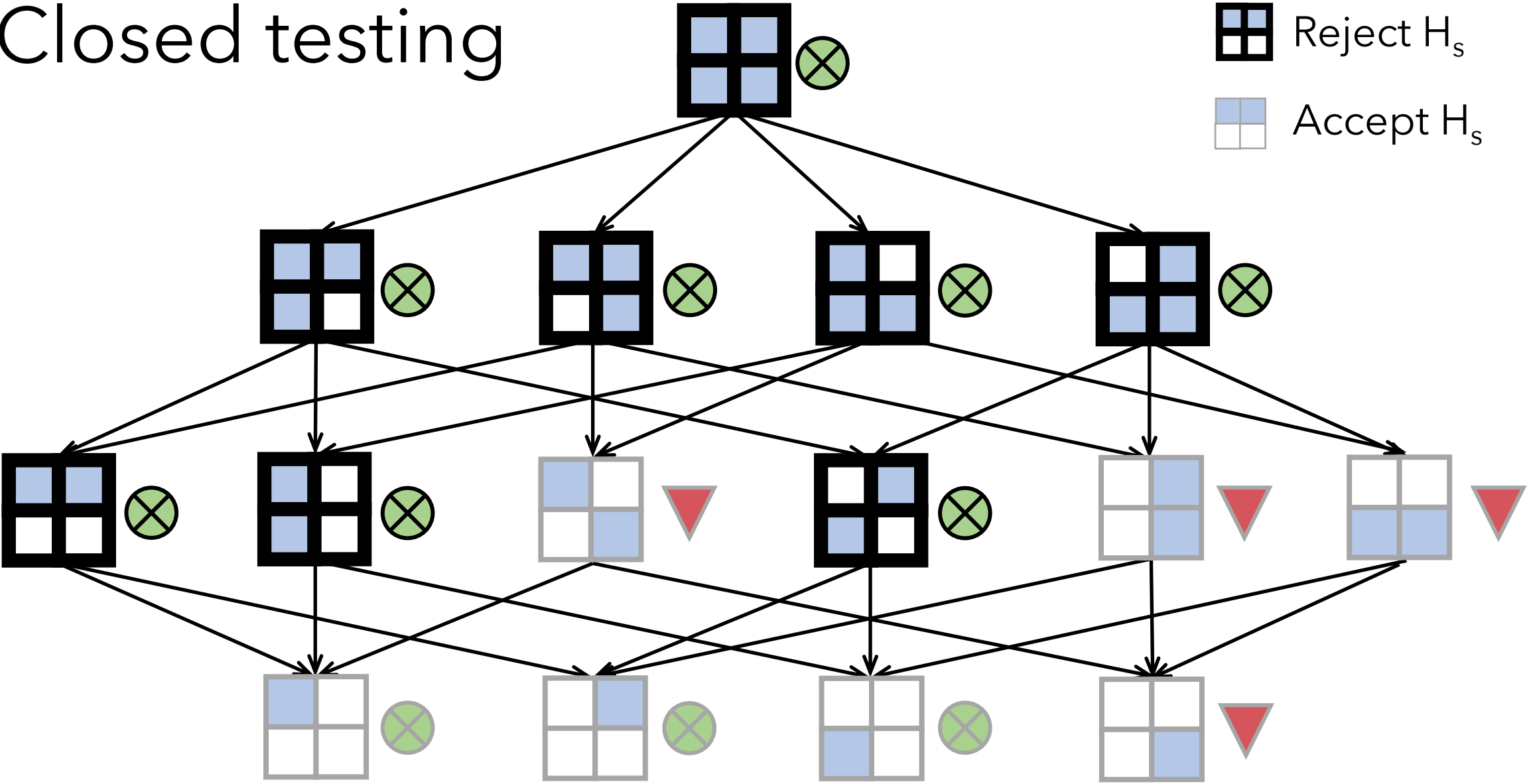
Closed testing



Closed testing



Closed testing



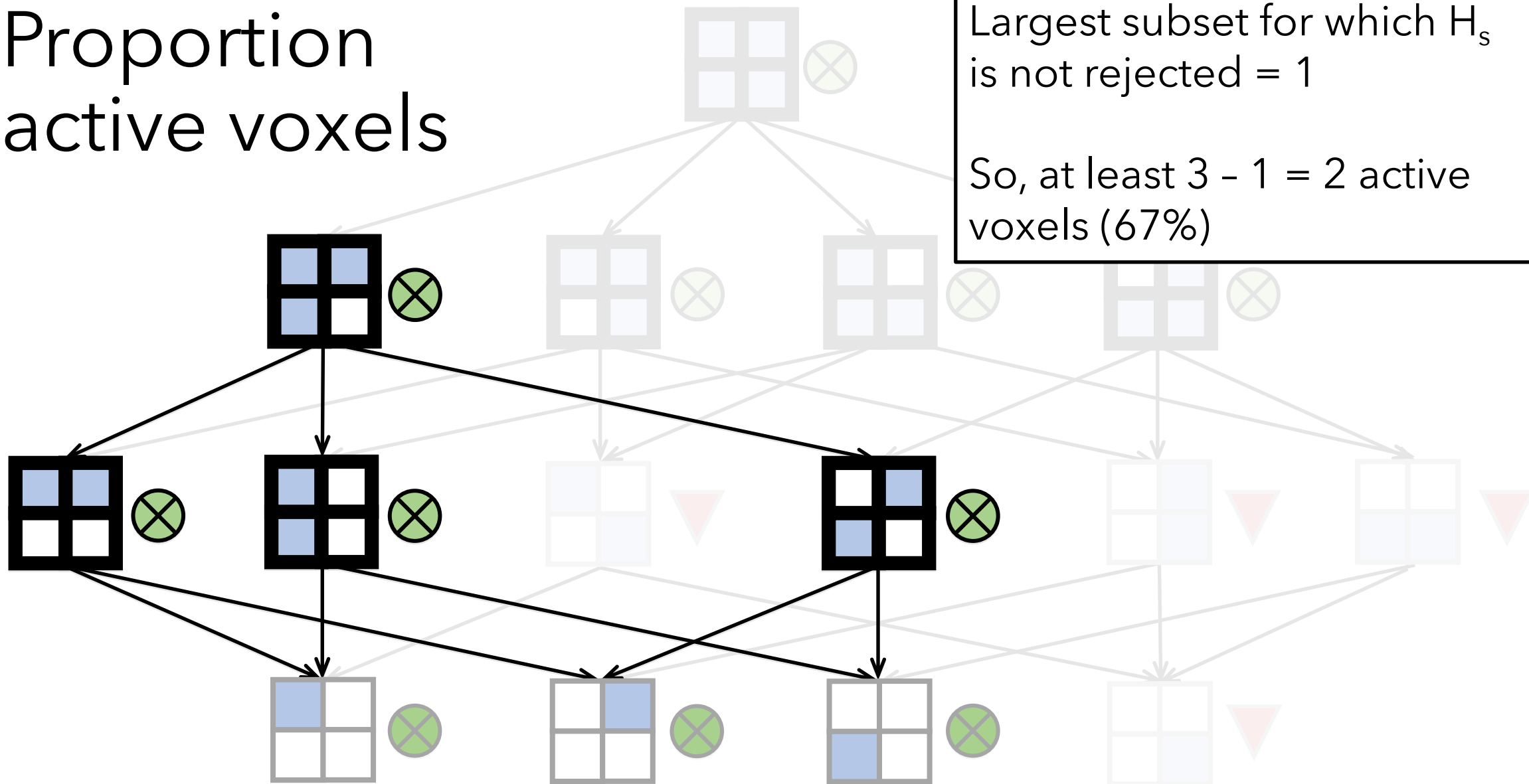
Proportion
active voxels



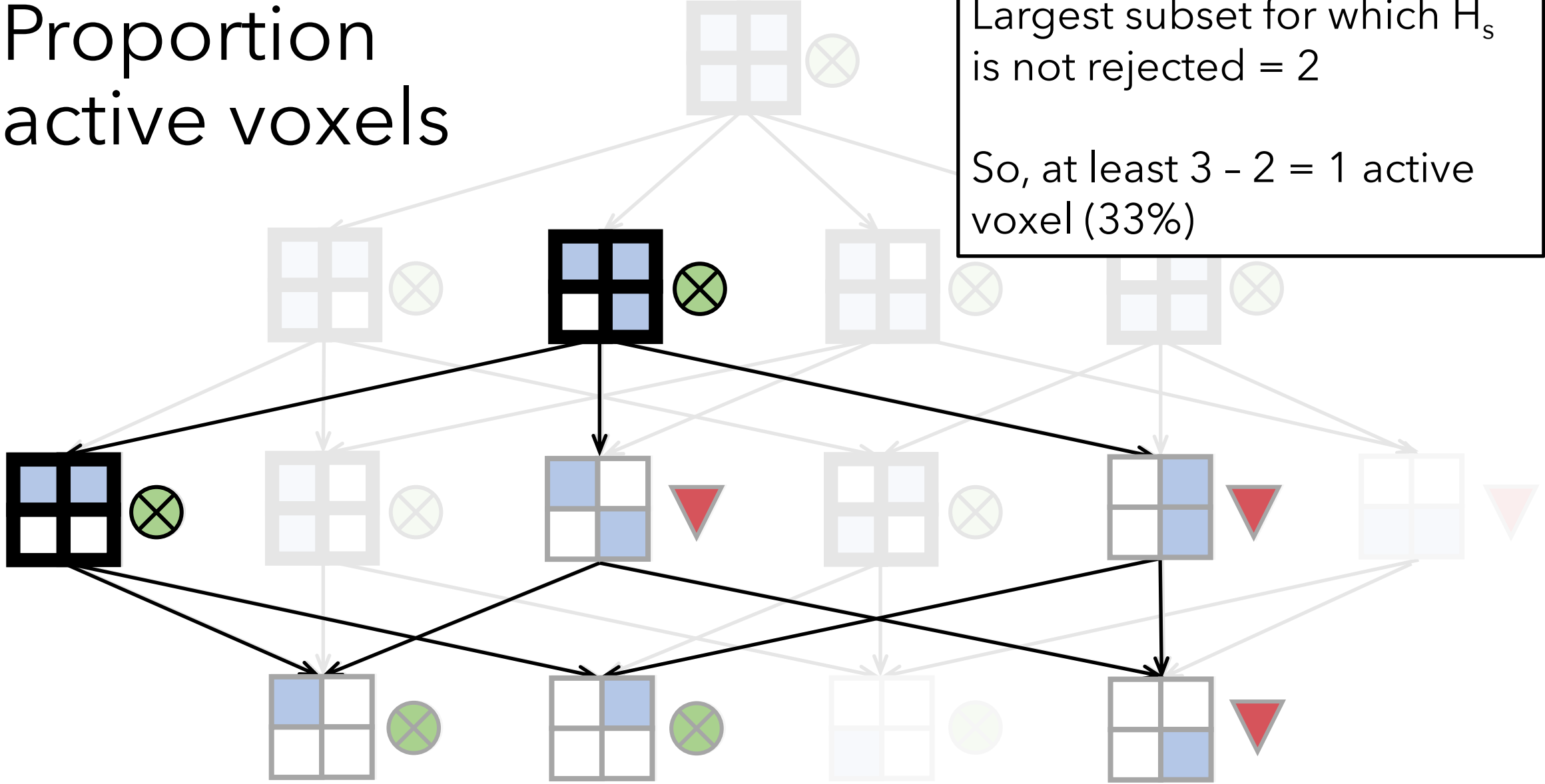
Proportion active voxels

Largest subset for which H_s
is not rejected = 1

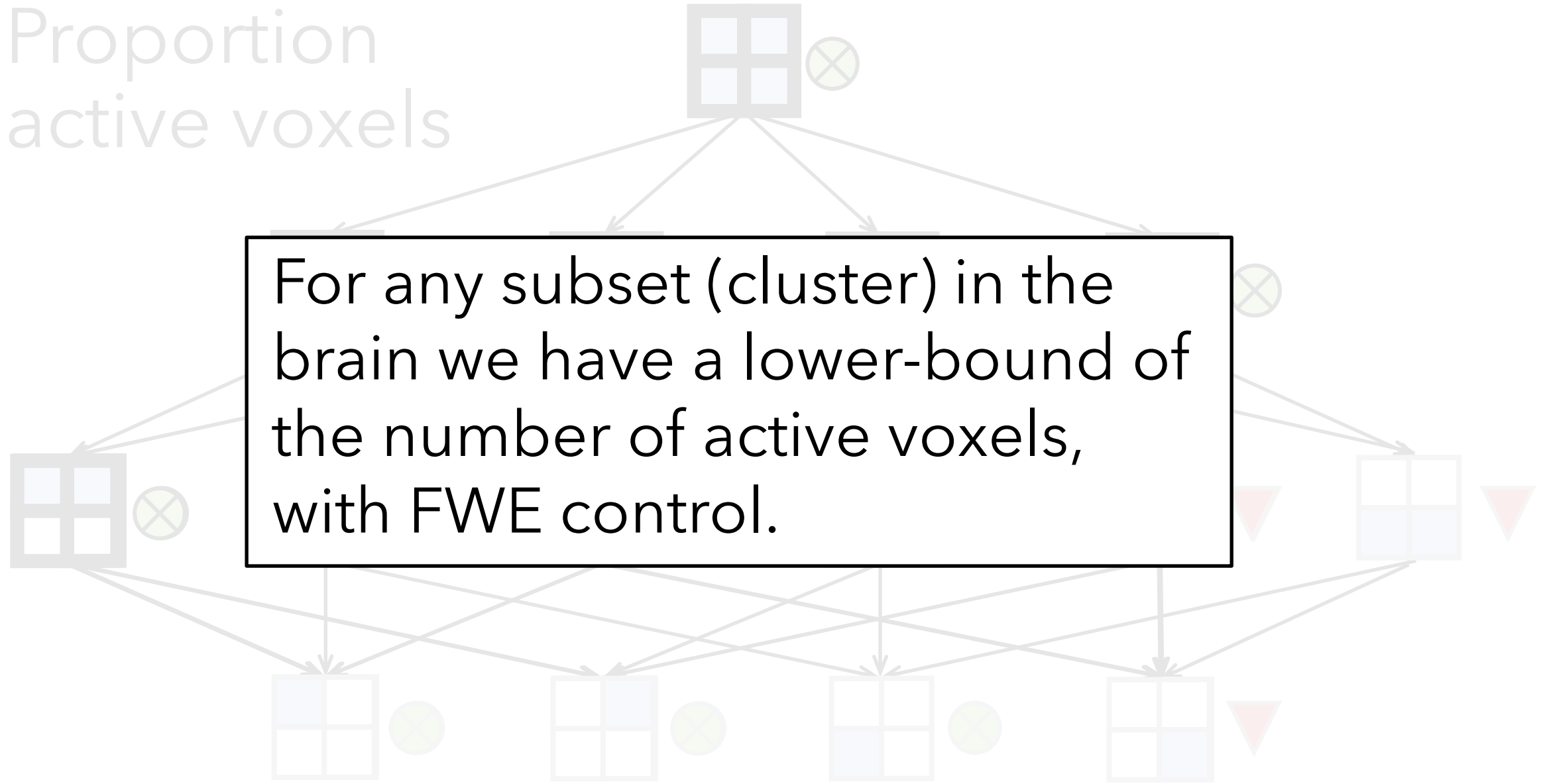
So, at least $3 - 1 = 2$ active
voxels (67%)



Proportion active voxels



Proportion
active voxels



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

- For 200,000 voxels there are $2^{200,000}$ possible subsets (a lot).
- Due to the structure of the Simes test and the closed testing procedure we can do this very efficiently.
- For 200,000 voxels: computation time within seconds.

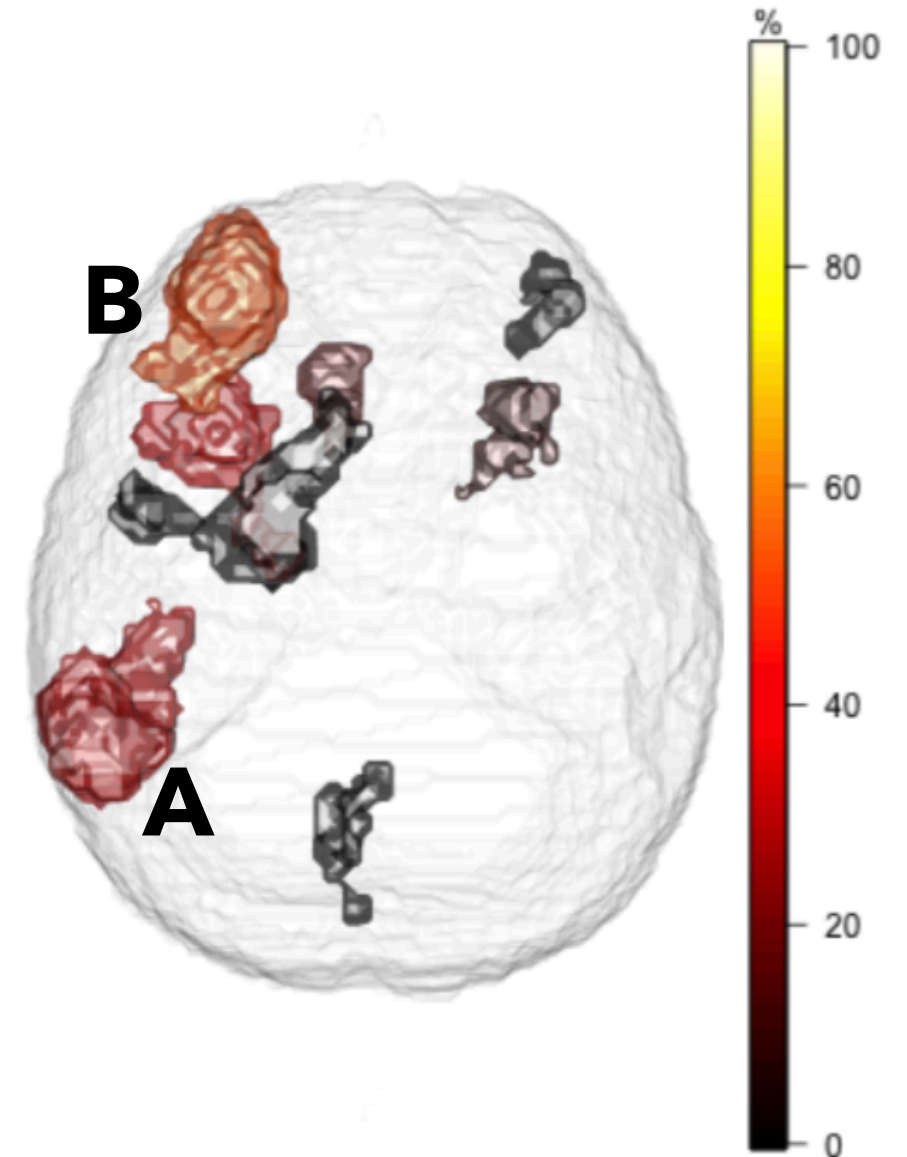
All-resolutions inference

- In exchange for your p-values and any region-of-interest you'll get:
 - Within seconds
 - A lower-bound of the number of truly active voxels of that region
 - For any region, as many times as you want.
 - With FWE control



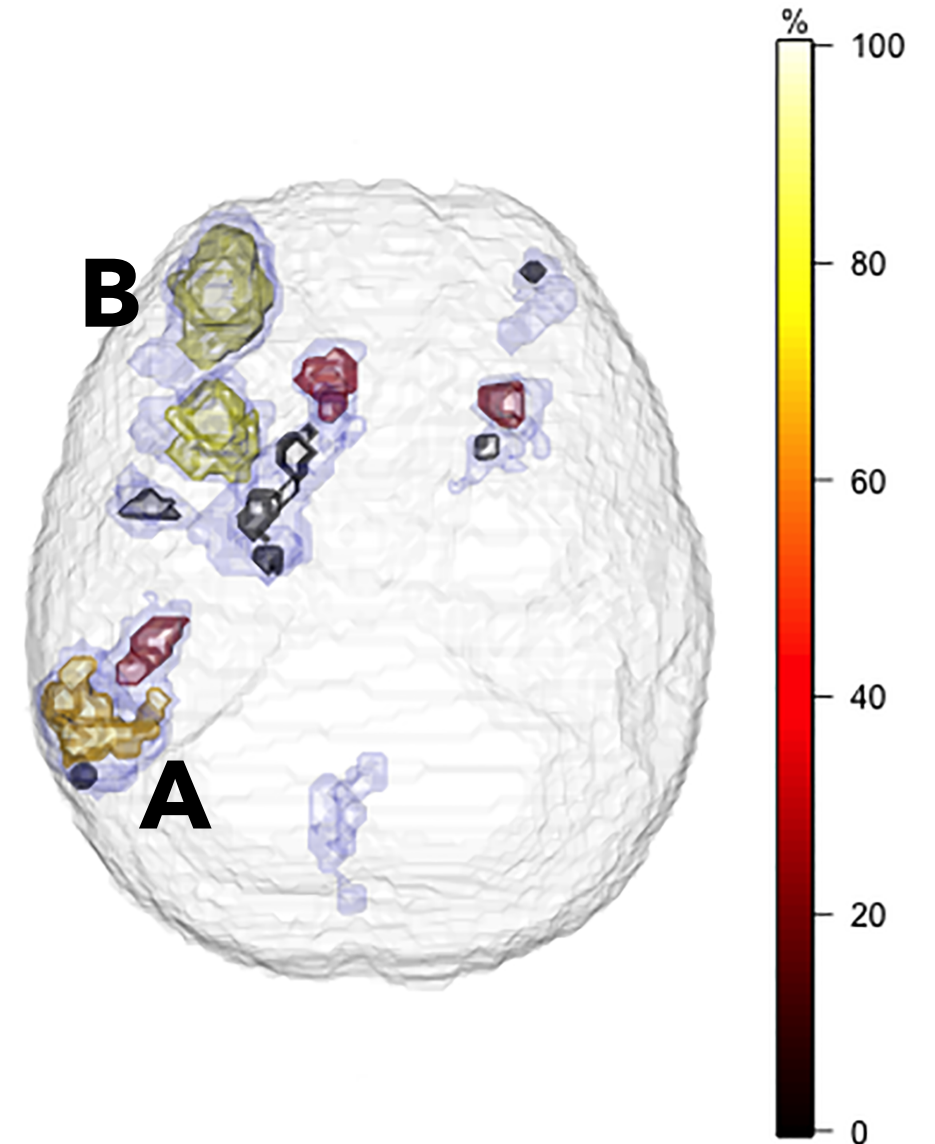
Go / No-go data

Cluster	Thres.	Size	# Active	% Active
A	$Z > 3.2$	2191	624	28.5 %
B	$Z > 3.2$	1835	847	46.2 %



Go / No-go data

Cluster	Thres.	Size	# Active	% Active
A	$Z > 3.2$	2191	624	28.5 %
1	$Z > 4$	405	267	65.9 %
2	$Z > 4$	133	31	23.3 %
3	$Z > 4$	6	0	0 %
B	$Z > 3.2$	1835	847	46.2 %
1	$Z > 4$	963	826	85.8 %



ARIBrain - R package

```
#load library
library(ARIBrain)

#run ARI
out <- ARI(Pmap = '/stats/pvalue1.nii.gz',
           clusters = '/cluster_zstat1.nii.gz',
           mask = '/mask.nii.gz',
           Statmap = '/stats/zstat1.nii.gz')
```

A hommel object for 166407 hypotheses.
Simes inequality is assumed.

With 0.95 confidence: at least 12938 discoveries.
9384 hypotheses with adjusted p-values below 0.05.

	Size	FalseNull	TrueNull	TDP	dim1	dim2	dim3	Stat
cl4	7257	6257	1000	0.86	23	56	40	10.789868
cl3	6206	5313	893	0.86	69	56	39	10.640452
cl2	694	355	339	0.51	52	51	33	6.859994
cl1	340	27	313	0.08	25	70	50	4.746142
cl0	151910	0	151910	0.00	64	79	41	4.319296

>



ARIbrain - R Shiny app

ARI Application

Dashboard

Load in Files

Perform ARI new

Zoom in on voxels

Download Axial Plot

Download Coronal Plot

Download Sagittal Plot

Download ARI Results

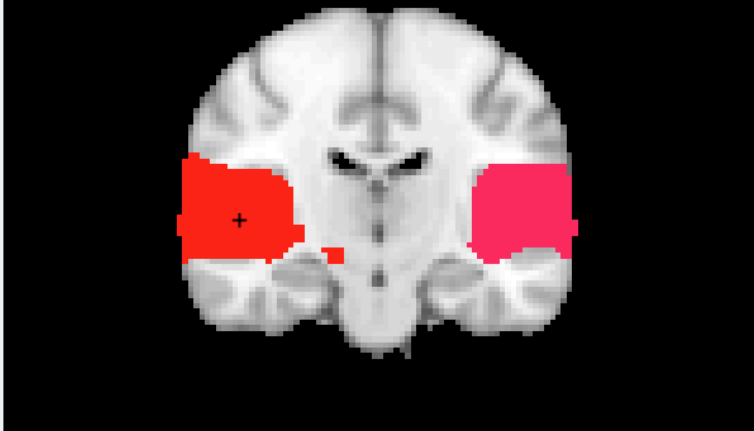

Download Zoom-in Results

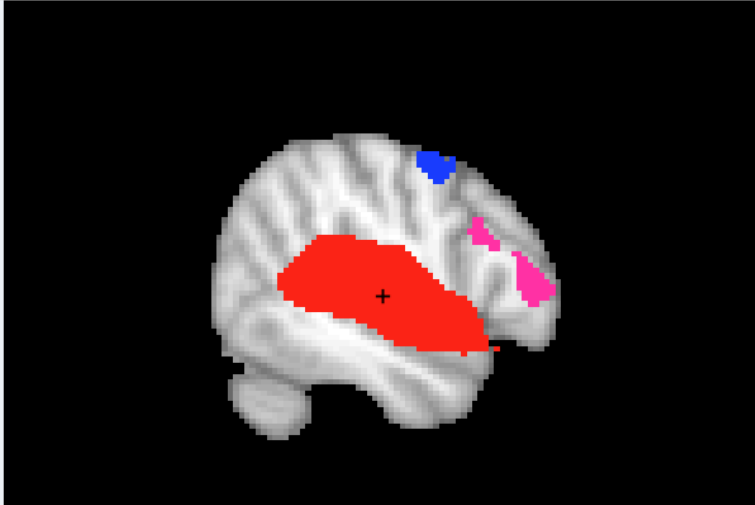
Download Cluster Map

Plots

All Clusters

Zoomed in





Step 1: Click here to load the fsl directory, statistics and mask files

x	y	z	Brain.area
20.00	55.00	38.00	No Atlas Selected

Step 2: Click here to run the All-Resolutions Inference (ARI)

cluster	Size	FalseNull	TrueNull	TDP	Statistic
cl21	9532.00	6821.00	2711.00	0.72	10.79

Step 3: Zoom in on clusters, spheres or brain areas

