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VETERINARY
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BAYESIAN NETWORKS MEET OBSERVATIONAL DATA



Outline of the talk

- ▶ Mixed models – correction for grouped data
- ▶ Heuristic search
- ▶ MCMC over structures

- ▶ *Other advanced methods/features:*
 - ▶ *Scoring system*
 - ▶ *Tunable parameter prior*
 - ▶ *Structural prior*
 - ▶ *Data separation*
 - ▶ *Covariate adjustment*
 - ▶ *Likelihood contribution*

CORRECTION FOR CLUSTERING

Correction for grouped data

- The way the data were collected has a clear **grouping aspect**
- Then potential for **non-independence** between data points
- Lead to analyses which are **over-optimistic**
- As the **true level of variation** in the data is **under-estimated**
- Could impact study result ... or not!
- **Good practice to check!**

In practice:

- Random effect
- **GLM** -> **GLMM** for each node
- Fit the DAG and check the posterior distribution (**widening**)
- If needed one can **incorporate random effect** in the **scoring** scheme

Pitfalls:

- High computational complexity!

Find maximum a posteriori score

✓ Exact search

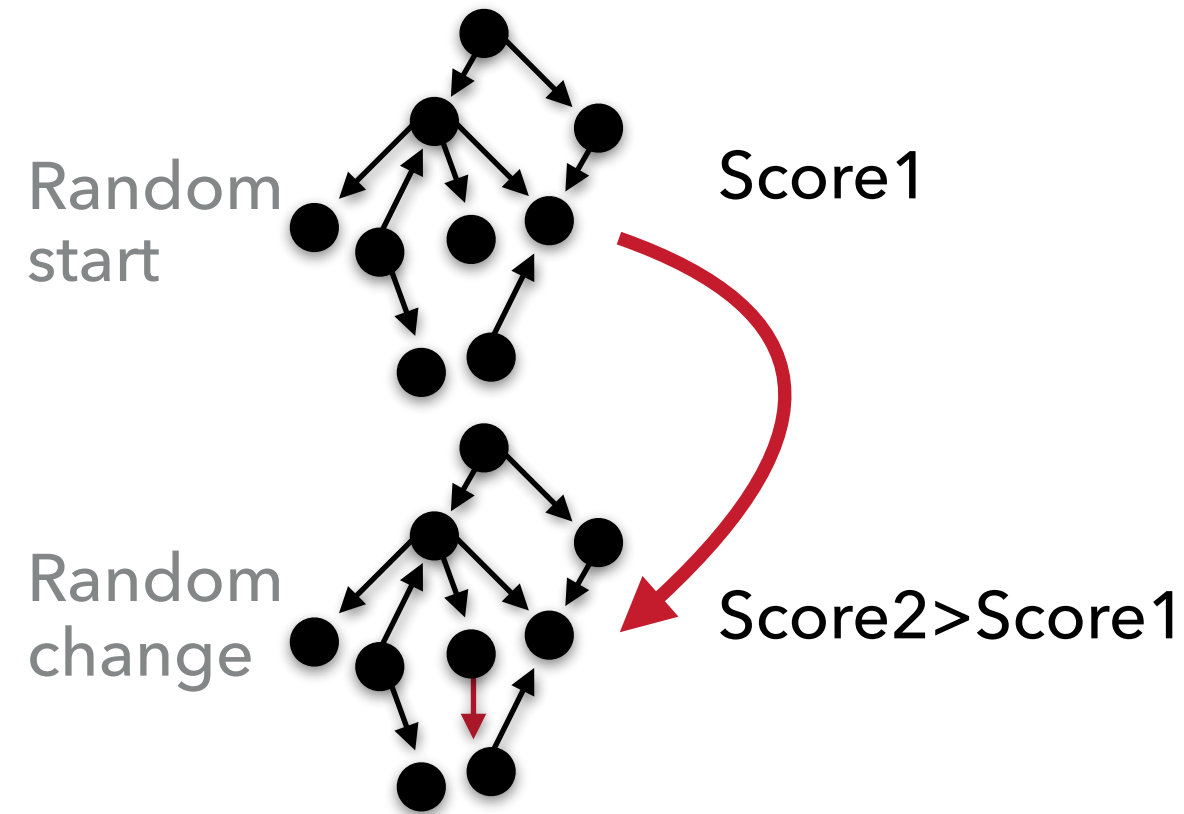
▶ Heuristic search

▶ MCMC over structures

HEURISTIC SEARCH

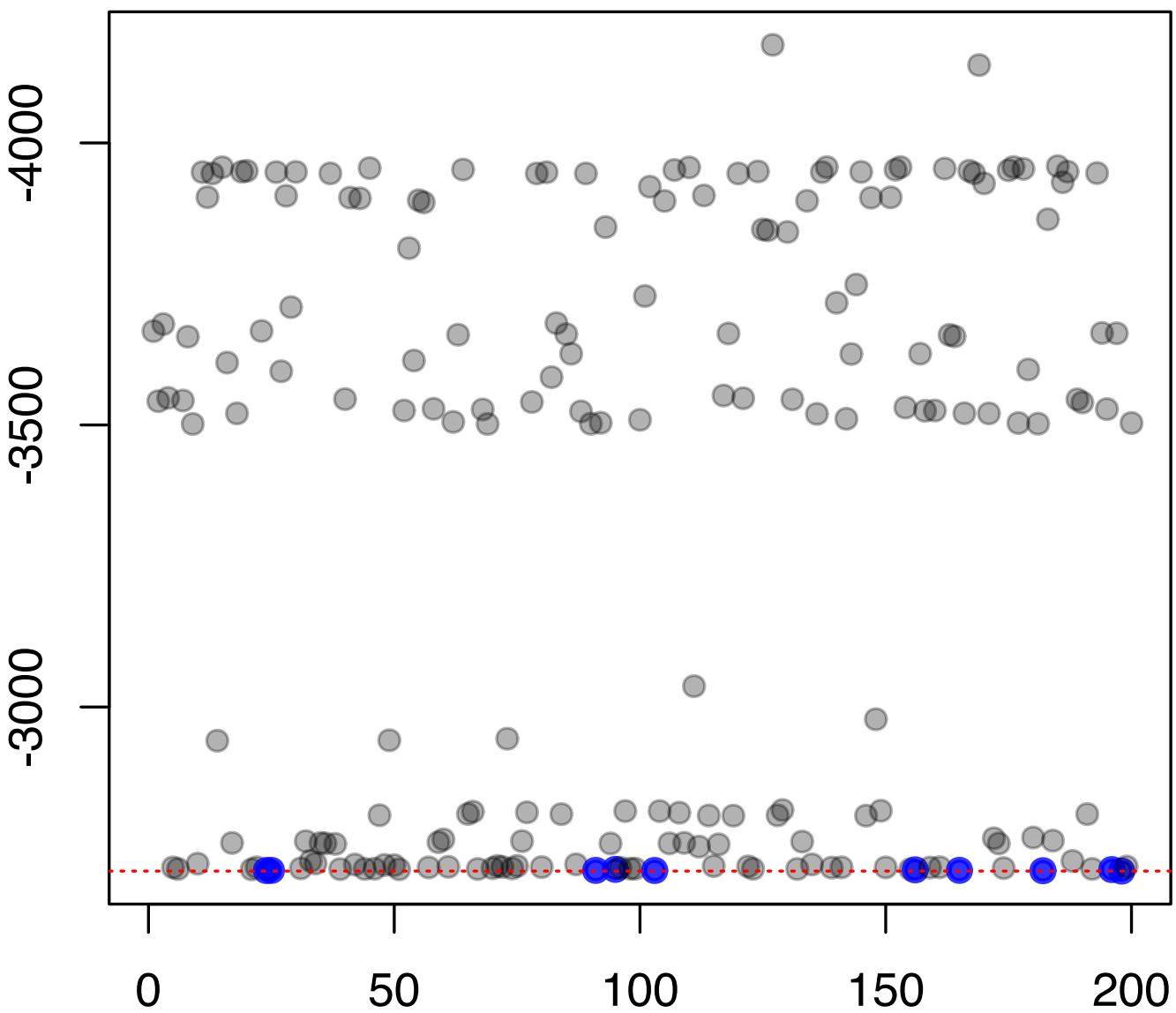
Heuristic search: Greedy Hill-Climbing

- ▶ Simplest heuristic local search
 - ▶ Start with a given network
 - ▶ empty network
 - ▶ best tree
 - ▶ a random network
 - ▶ At each iteration
 - ▶ Evaluate all possible changes
 - ▶ Apply change that leads to best improvement in score
 - ▶ Reiterate
 - ▶ Stop when no modification improves score
- ▶ *Pitfalls:*
 - ▶ Local Maxima
 - ▶ Plateaus
- ▶ *Solution:*
 - ▶ Tabu
 - ▶ Random restart
 - ▶ Simulated annealing

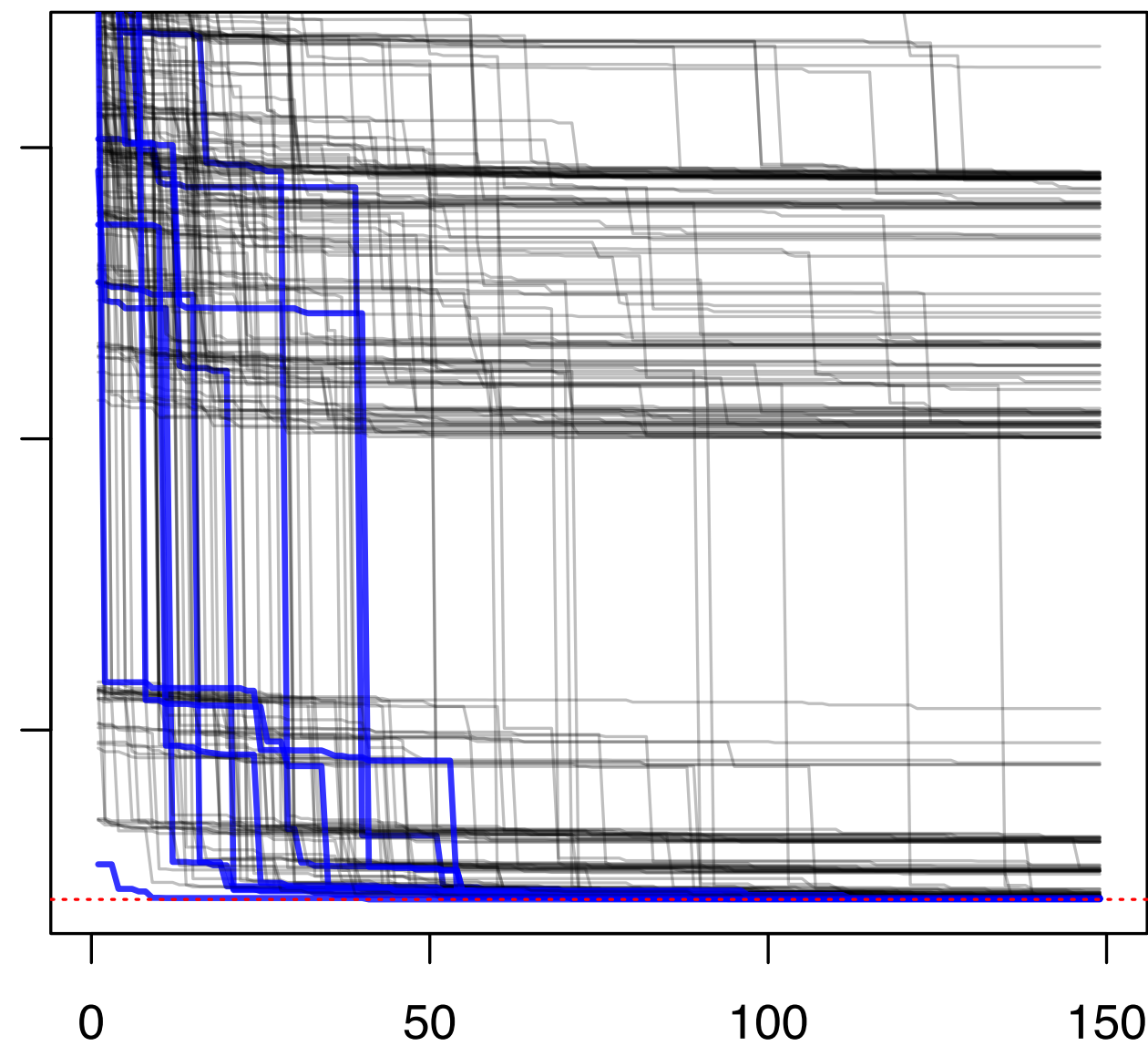


HEURISTIC SEARCH

```
num.searches <- 200
max.steps <- 150
heur.res <- quiet(search.heuristic(score.cache = mycache,
                                   score = "mlik",
                                   data.dists = dist,
                                   max.parents = 4,
                                   start.dag = "random",
                                   num.searches = num.searches,
                                   max.steps = max.steps,
                                   seed = 3213,
                                   verbose = TRUE,
                                   algo = "hc"))
```

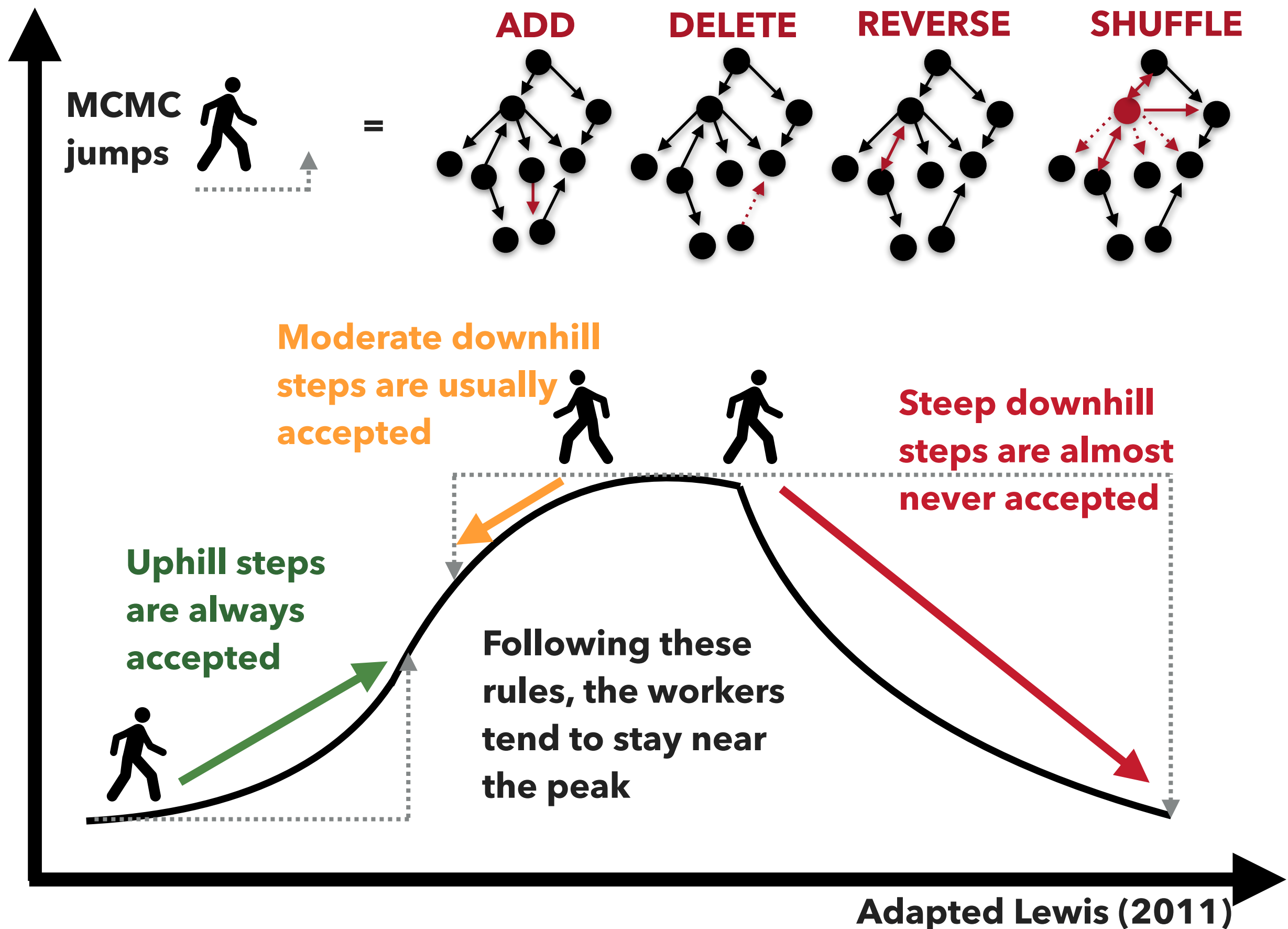


Index of heuristic search



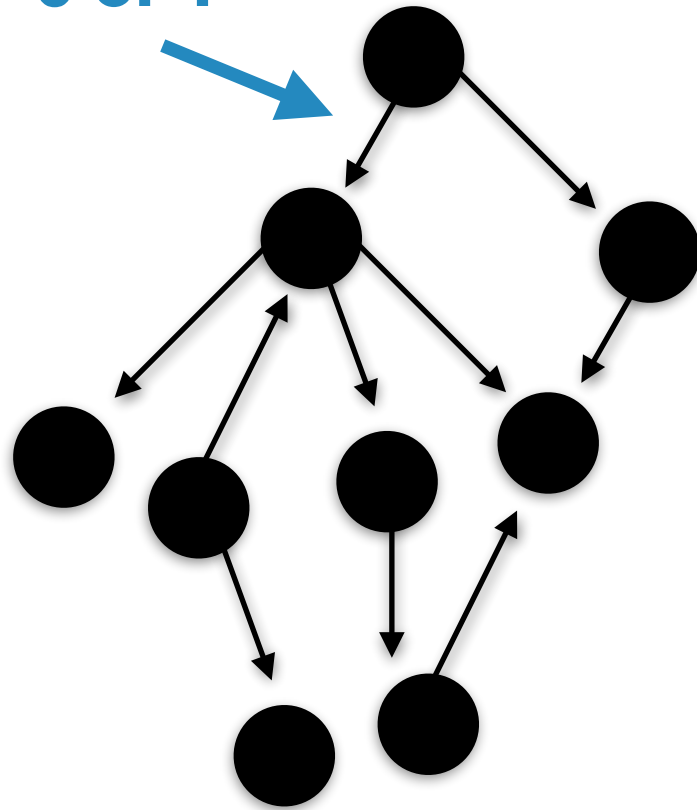
Number of Steps

MCMC OVER STRUCTURES

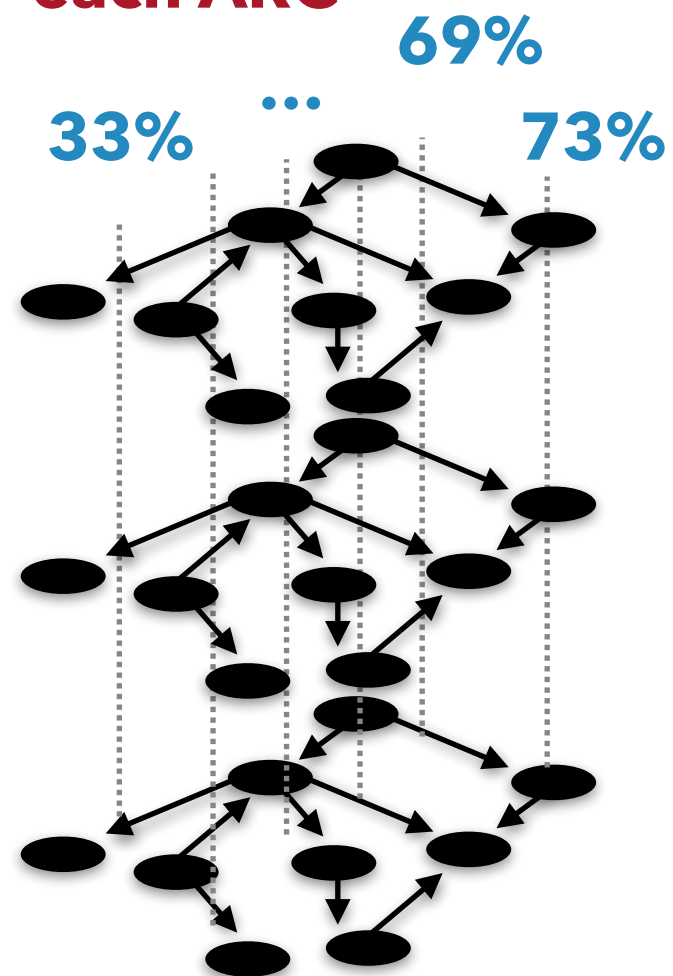


Best Unique Bayesian Network

0 or 1



Counting prevalence of each ARC



MCMC OVER STRUCTURES

MCMC over structures

- ▶ Selecting the most probable structure
- ▶ Controlling for overfitting
- ▶ Sampling the landscape of high scoring structures
 - ▶ In applied perspective avoid reducing the richness of BN modelling to only **one** structure
 - ▶ Quantify the marginal impact of relationships by marginalising out over structures



HANDS-ON EXERCICES