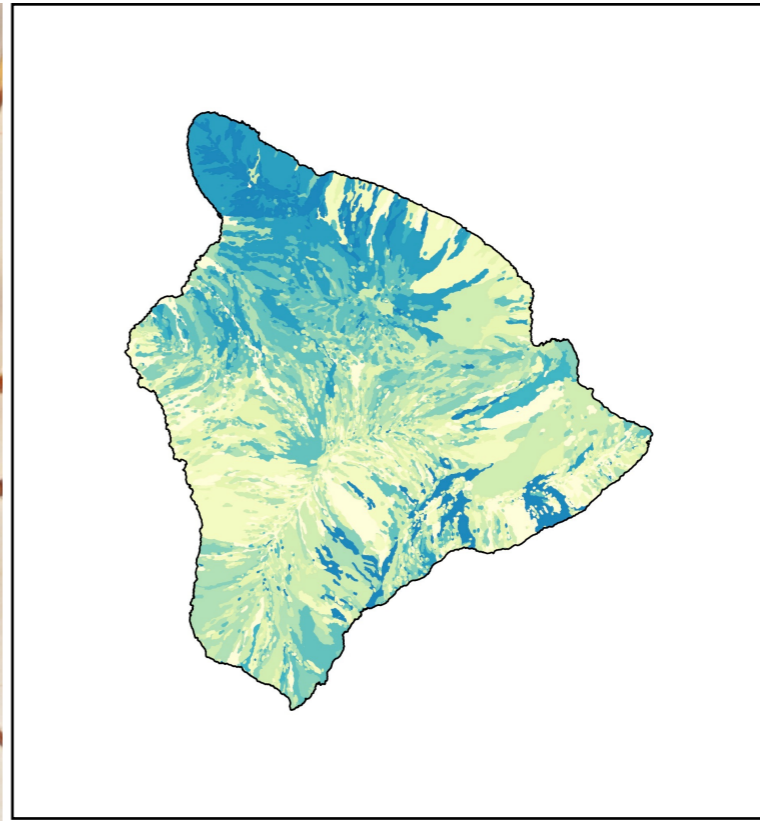


Biocollections, hierarchical models & islands



$$\int_{\theta_S} \frac{\mathbb{P}(\hat{D} | S, M) \mathbb{P}(S, M)}{\mathbb{P}(\hat{D})} d\theta_S$$



Andy Rominger

nature.berkeley.edu/~rominger

Island Biology • 19 July 2016

Thanks!

David Ackerly, Lucy Chang, Margaret Evans, Rosie Gillespie, Rob Guralnick, Jun Ying Lim, Cory Merow, Gio Rapacciuolo, Ethan White, Adam Wilson



Berkeley Initiative in
UNIVERSITY OF CALIFORNIA
Global Change Biology

GORDON AND BETTY
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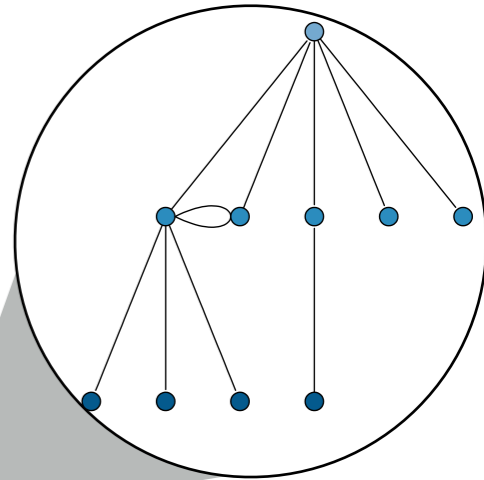
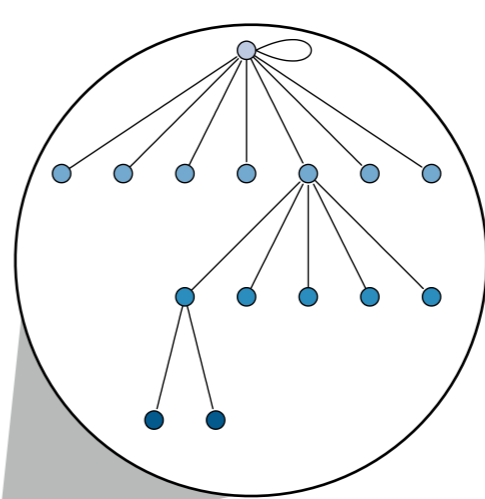
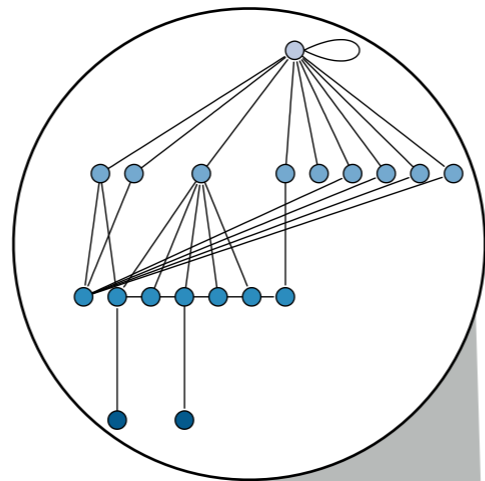
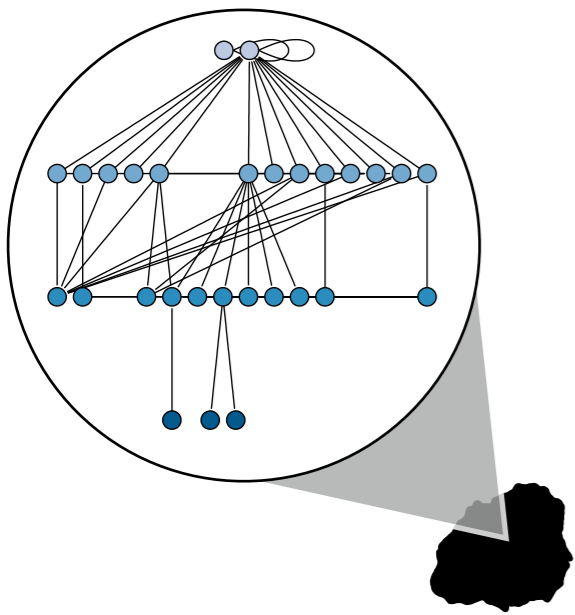
NIMBioS

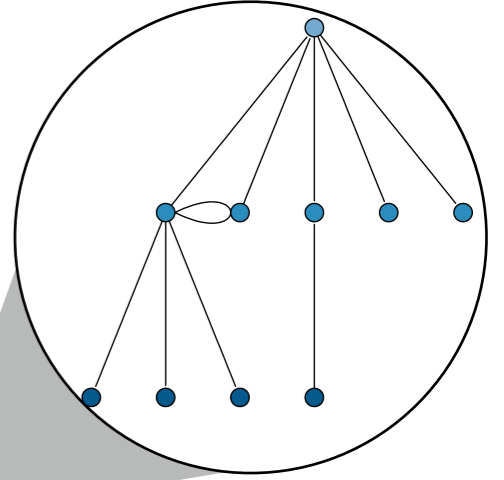
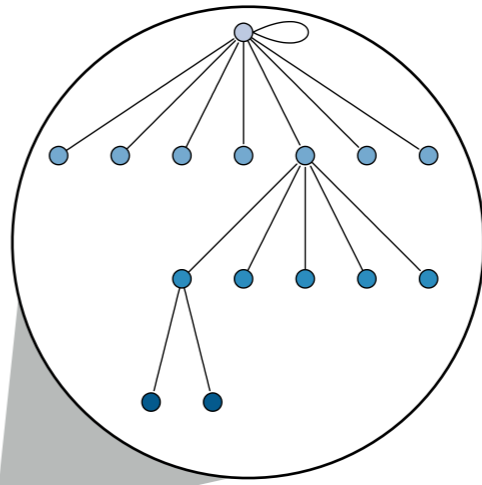
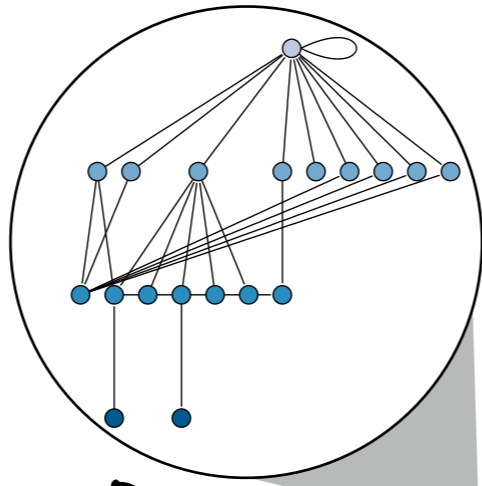
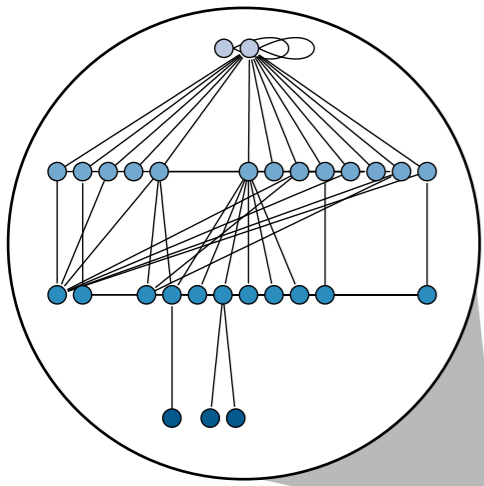


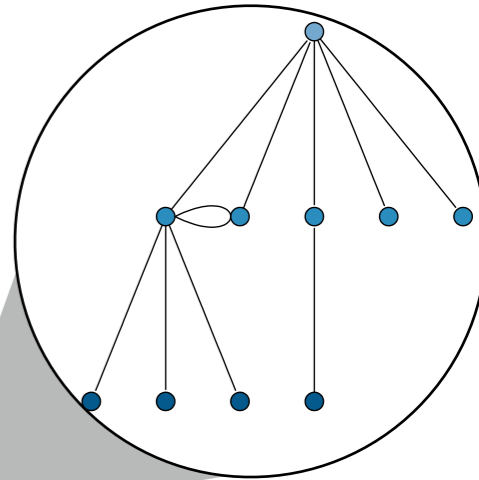
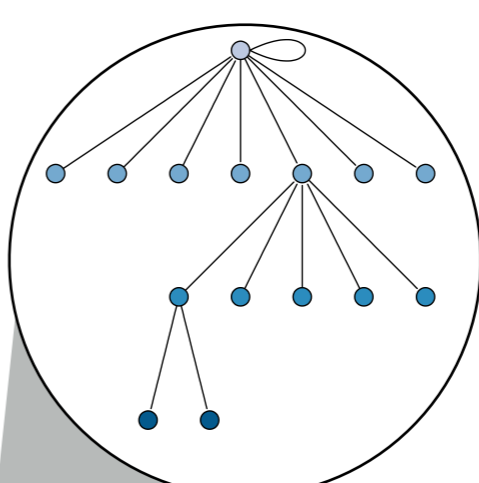
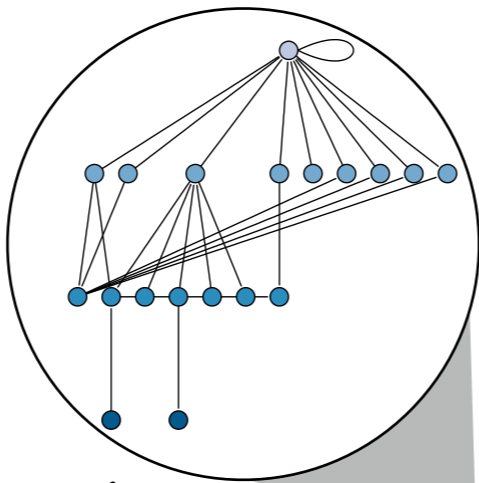
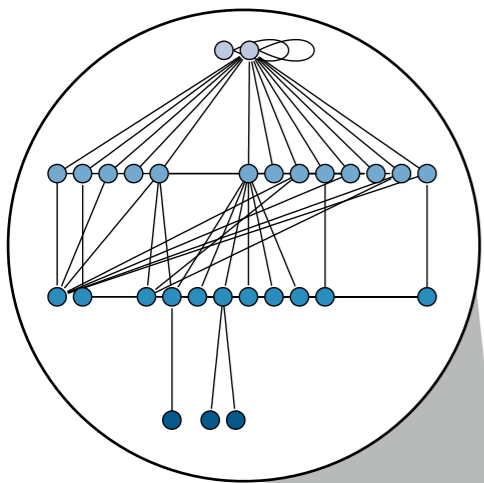
iDigBio

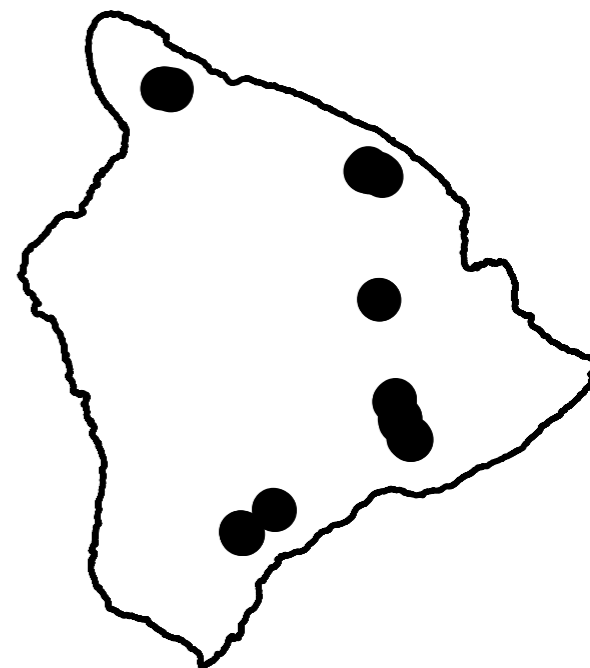
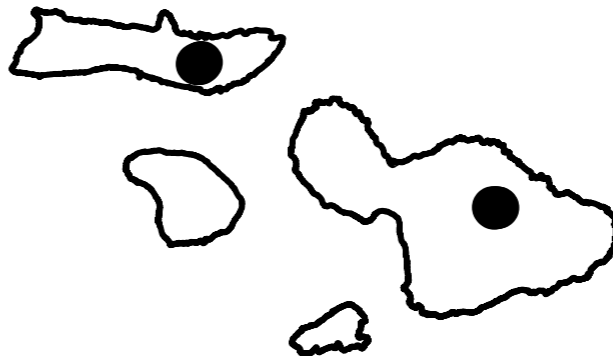
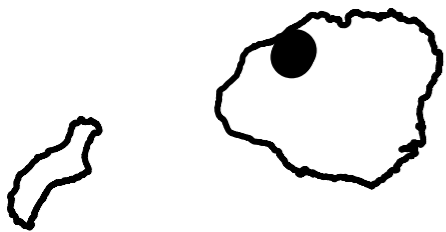
Integrated Digitized Biocollections

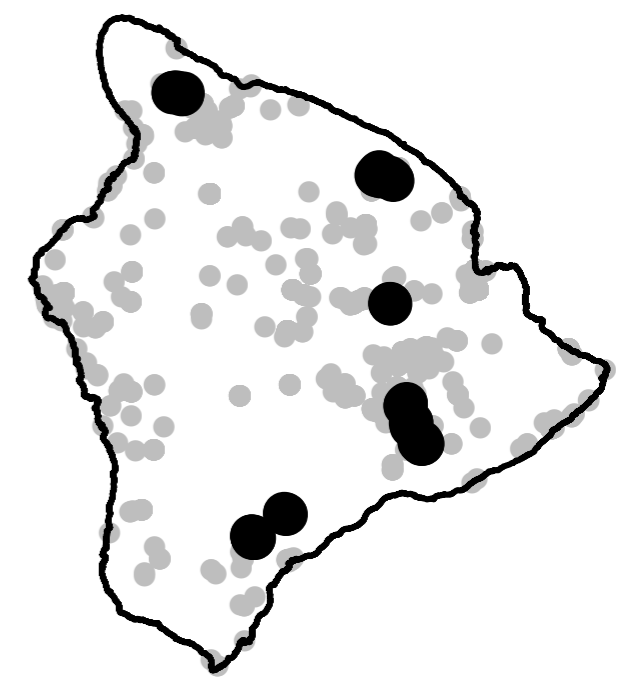
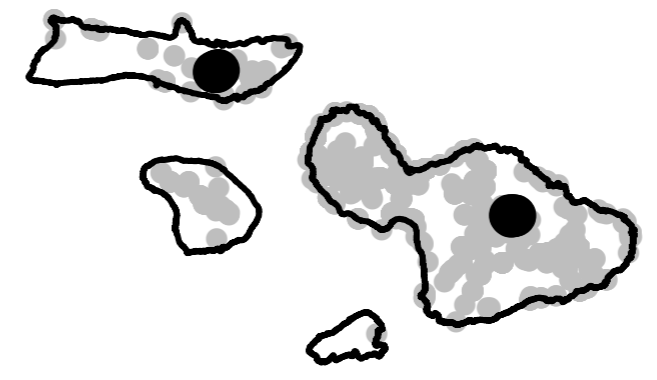
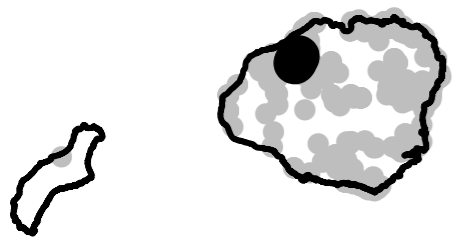






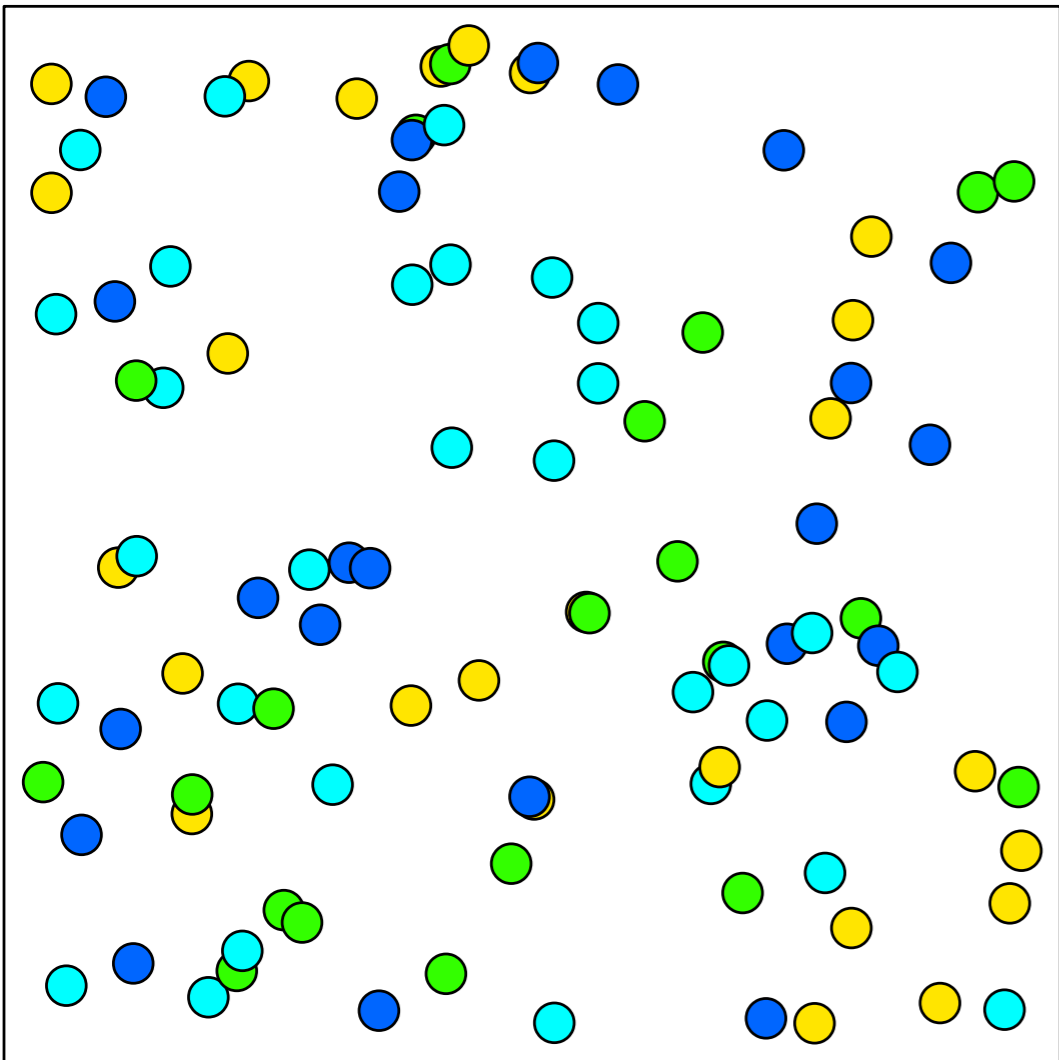




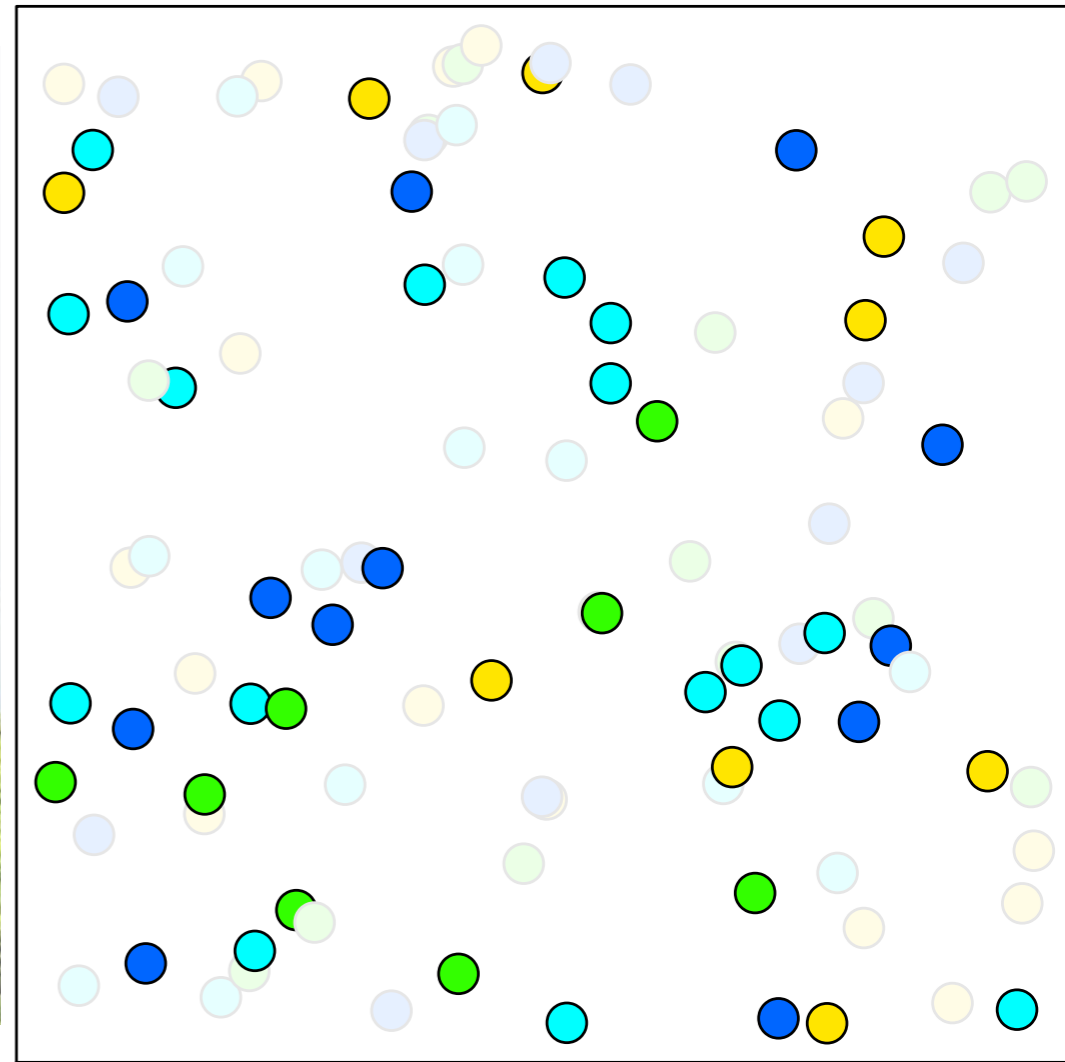
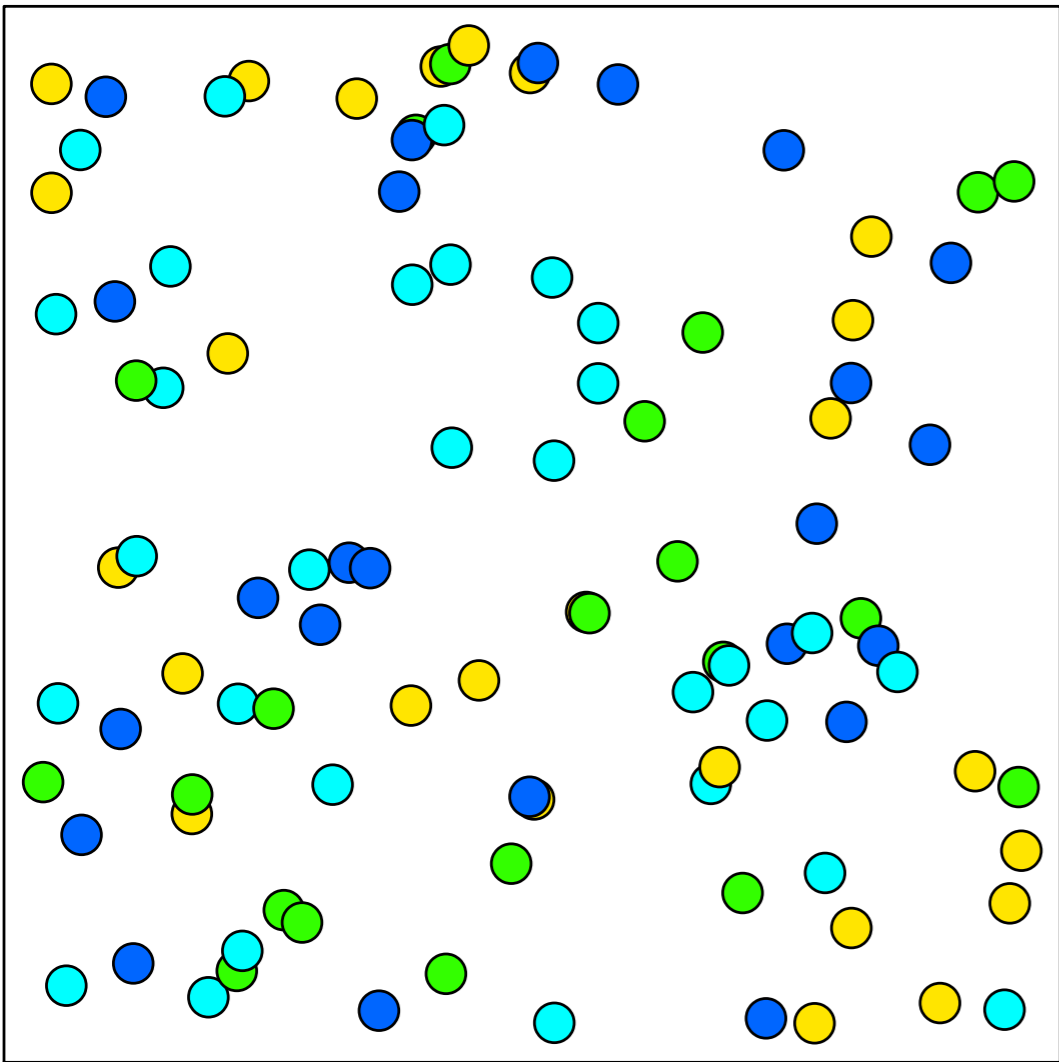


Sampling problems

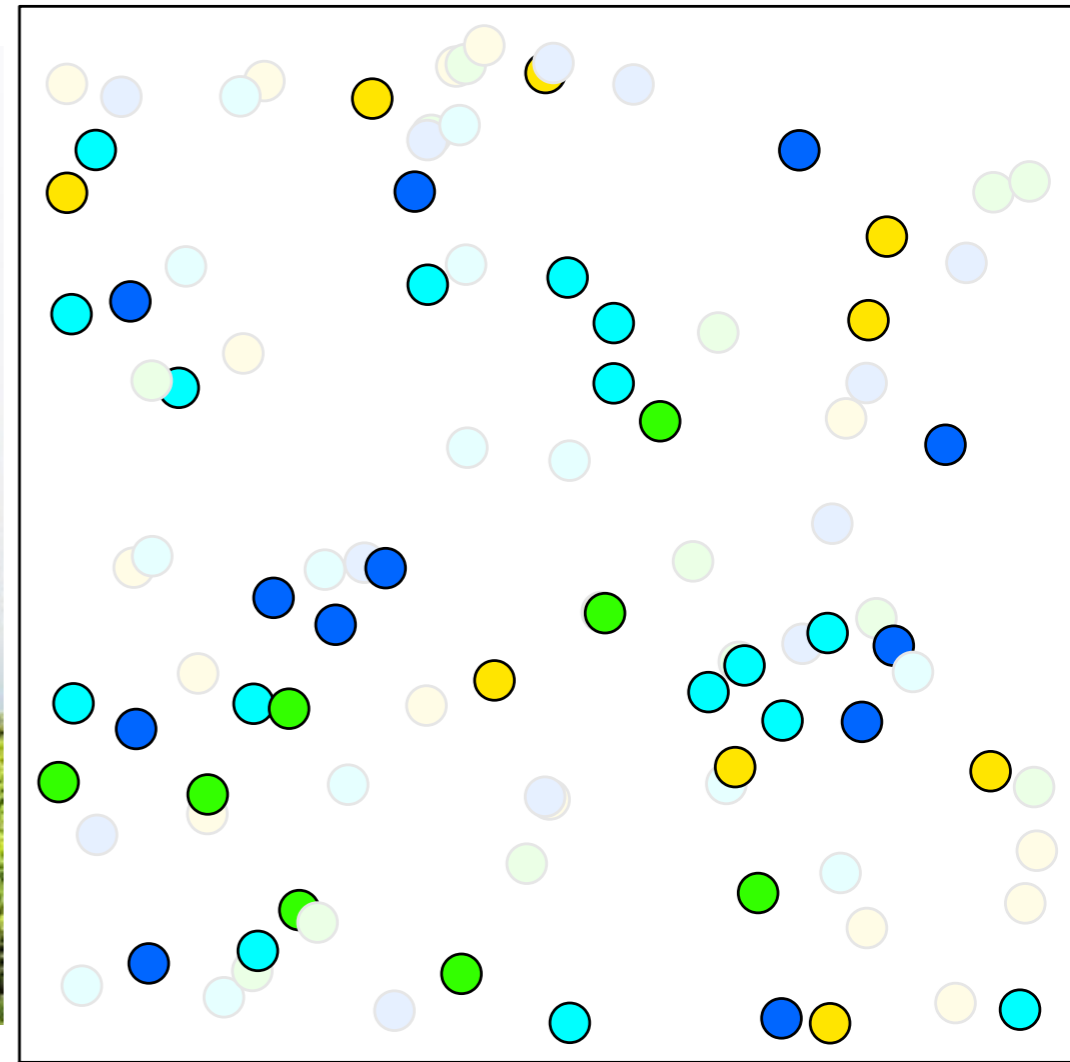
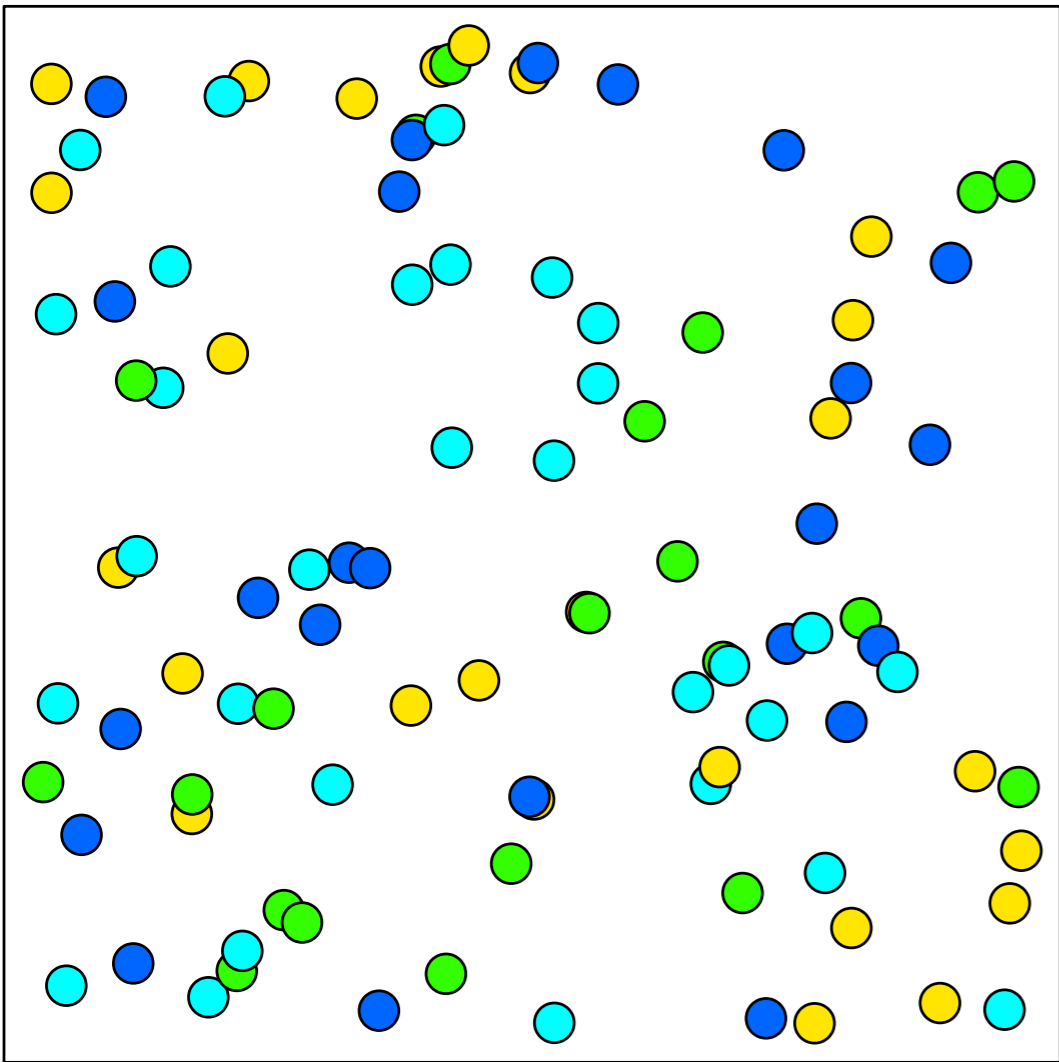
Sampling problems



Sampling problems



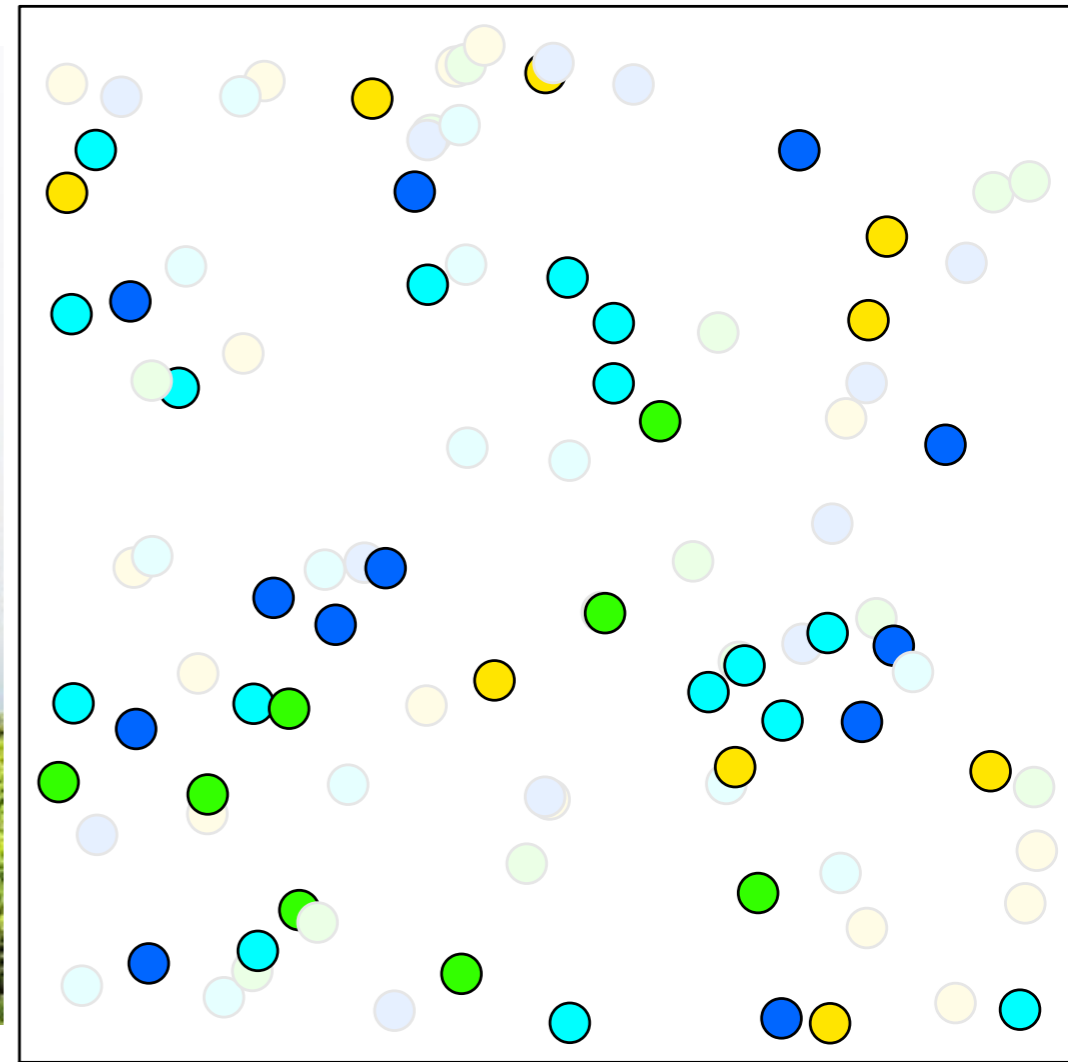
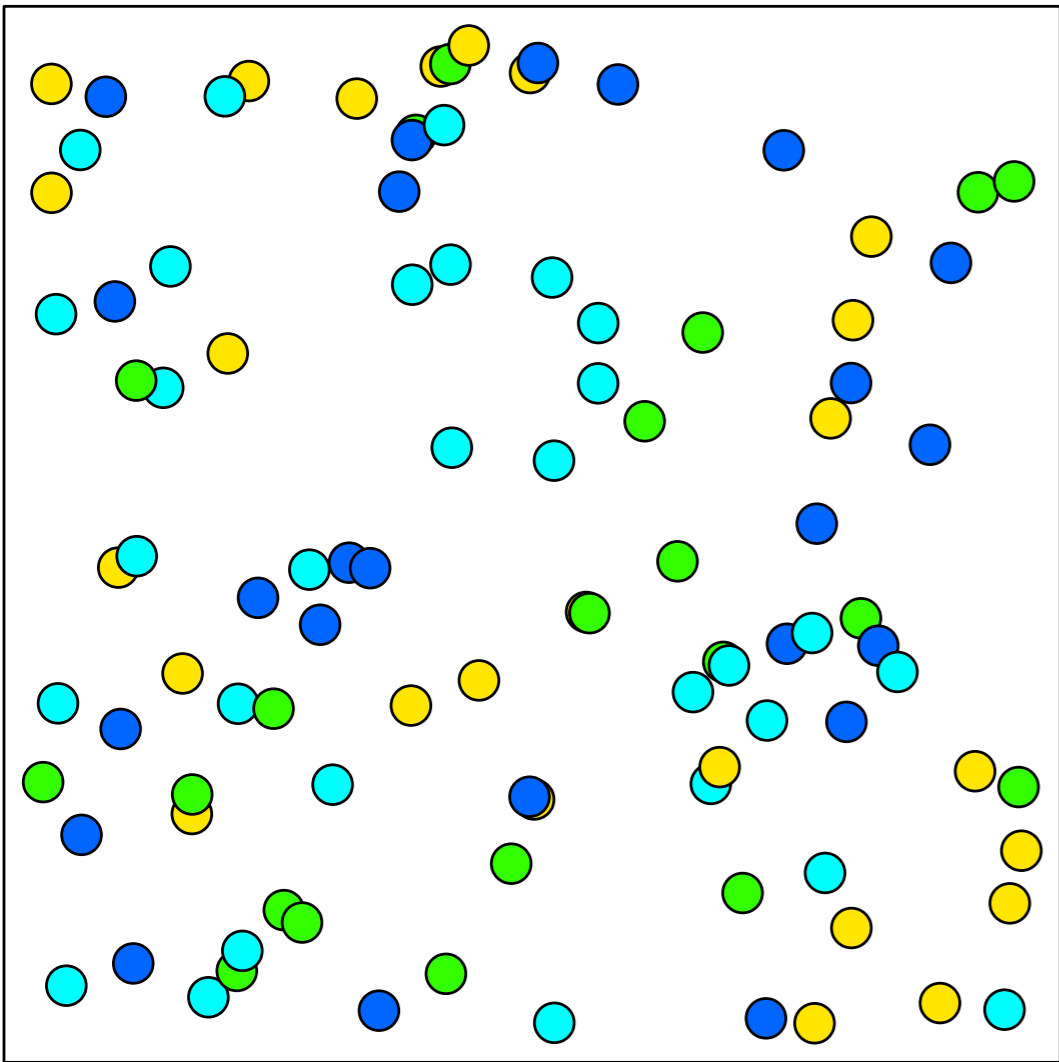
Hierarchical models



Hierarchical models

latent data

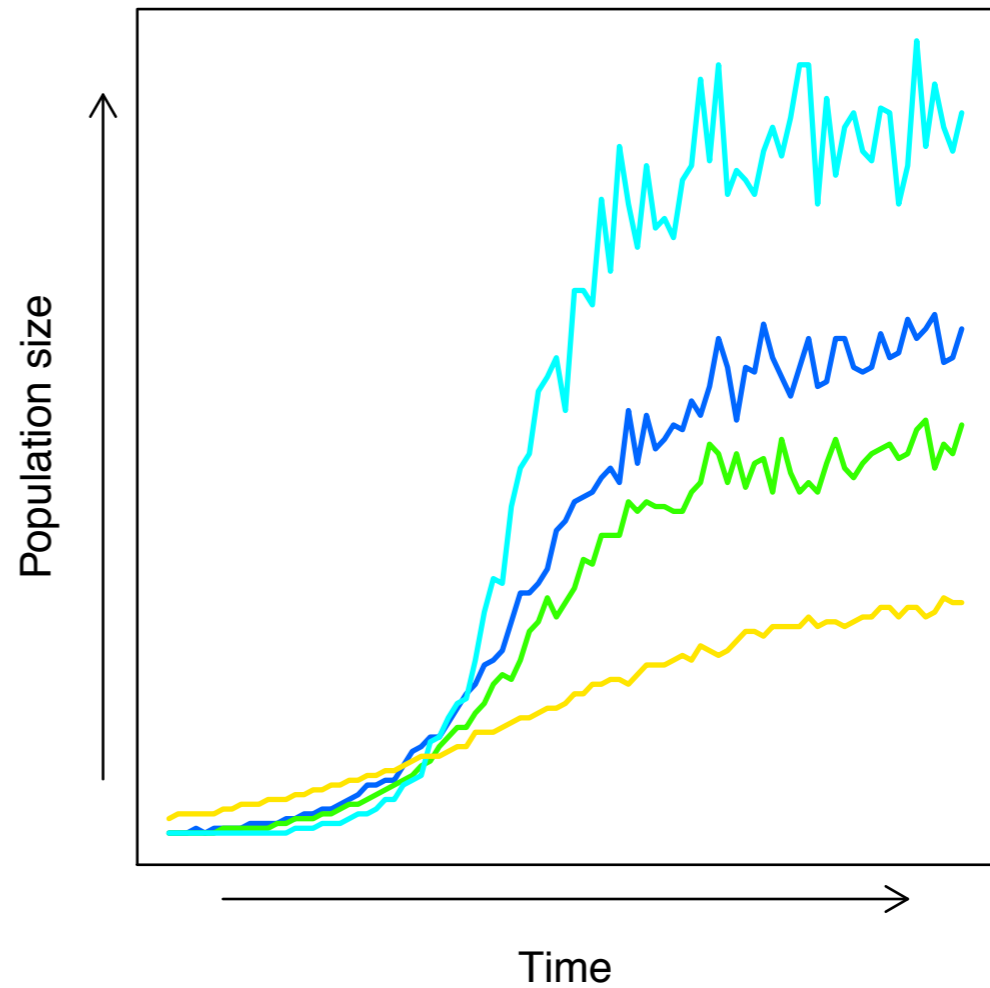
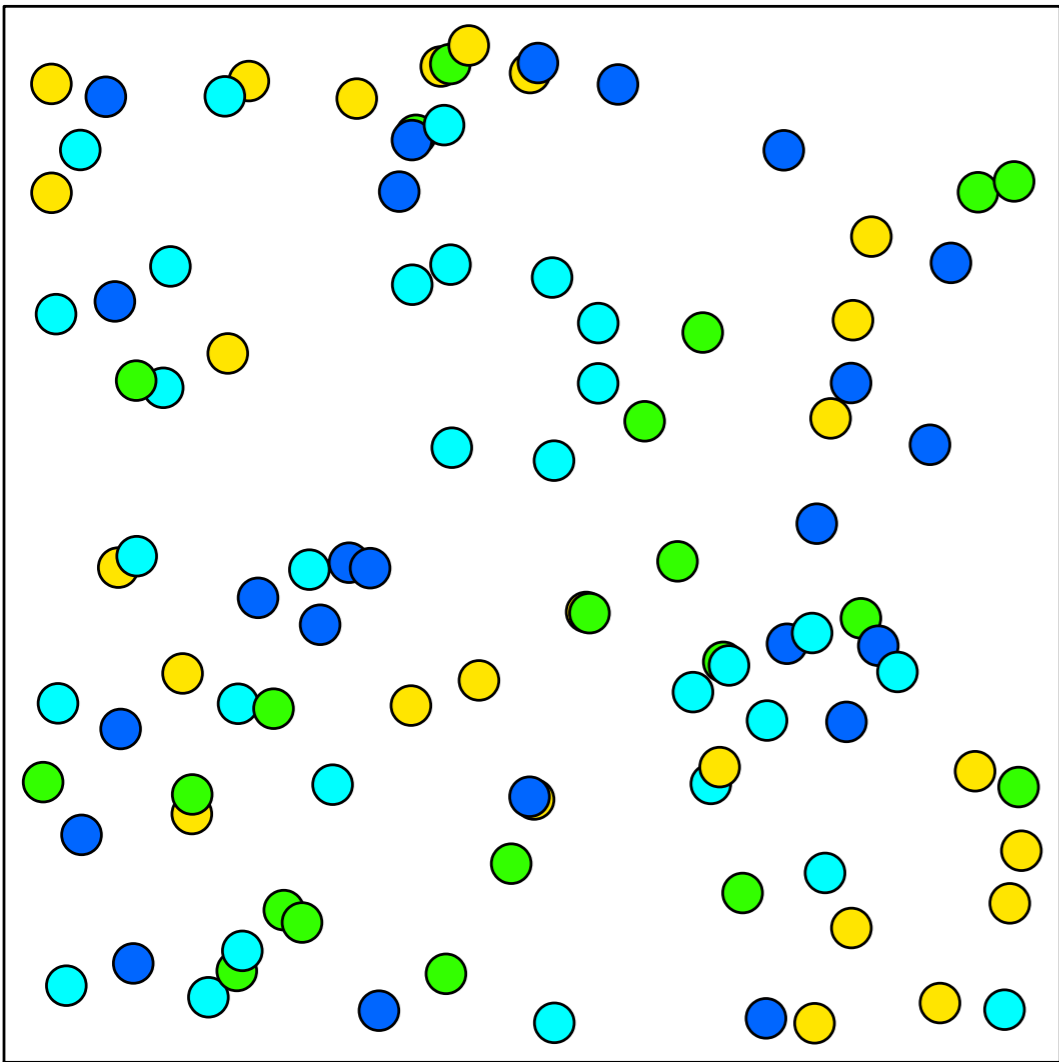
real data



$$\mathbb{P}(\text{real data} \mid \text{sampling, latent data})$$

Hierarchical models

latent data



$\mathbb{P}(\text{latent data}|\text{model})$

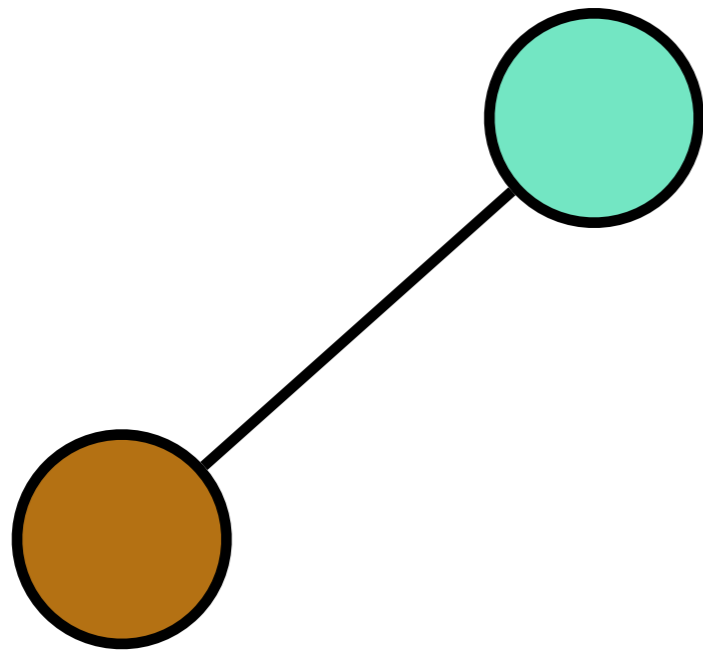
Hierarchical models



$$\mathbb{P}(\text{model} \mid \text{real data}) = \int_{\theta_S} \frac{\mathbb{P}(\text{real data} \mid \text{sampling, latent data}) \mathbb{P}(\text{latent data} \mid \text{model}) \mathbb{P}(\text{sampling, model})}{\mathbb{P}(\text{real data})} d\theta_S$$

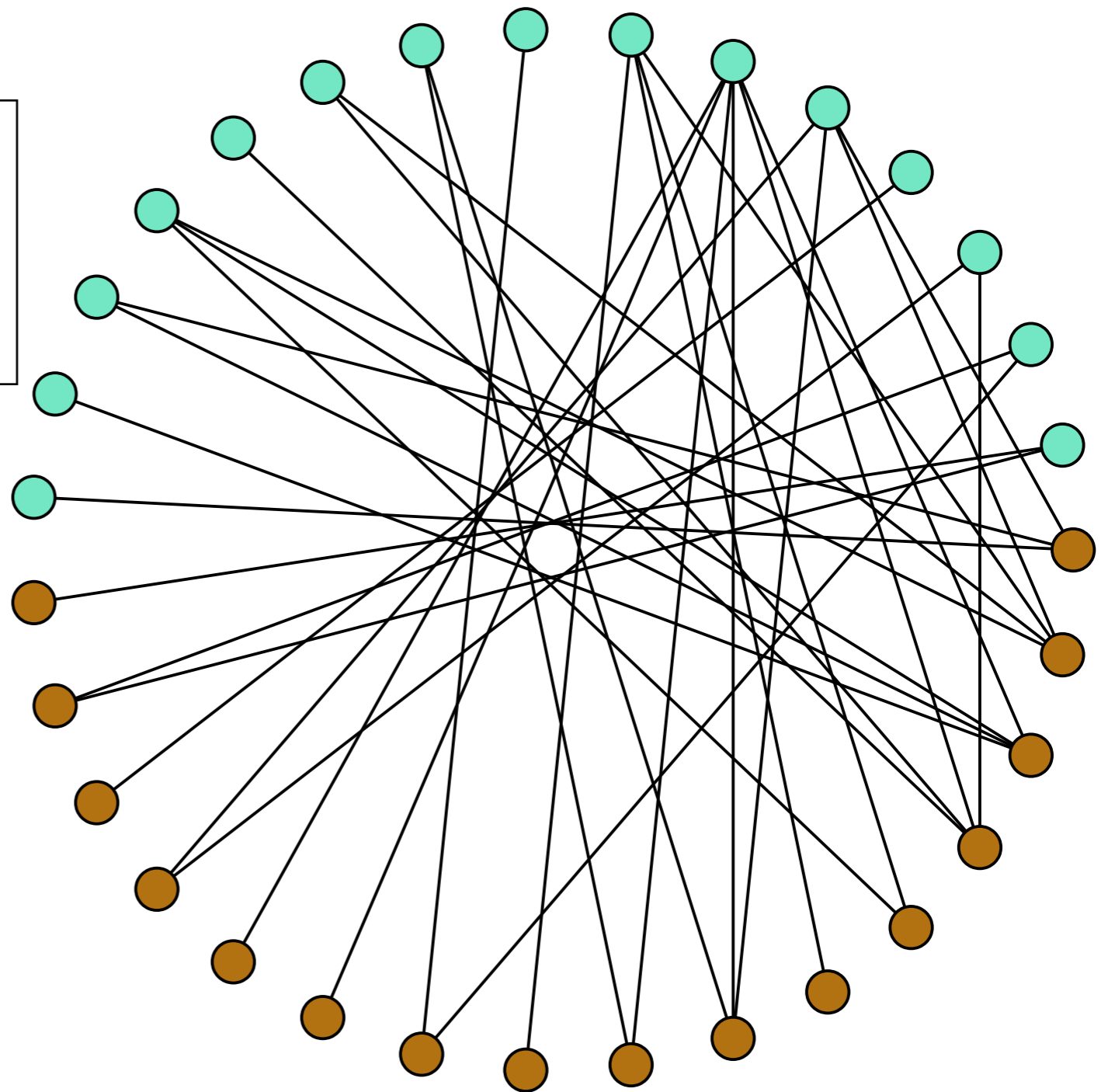
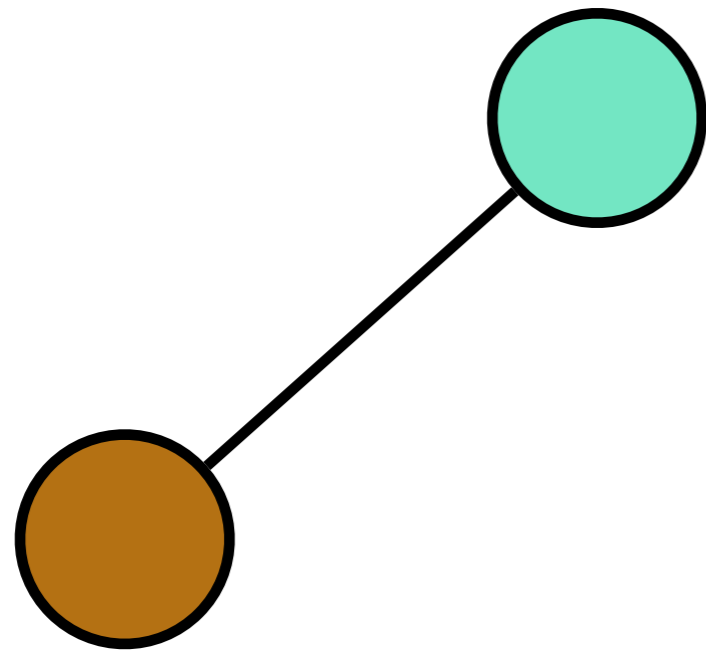
Use case: Networks

HDIM6643; laupLSAG_10
beating Melicope 65 sec.
09:25–10:08; 2015-oct-1
B. Cote coll.



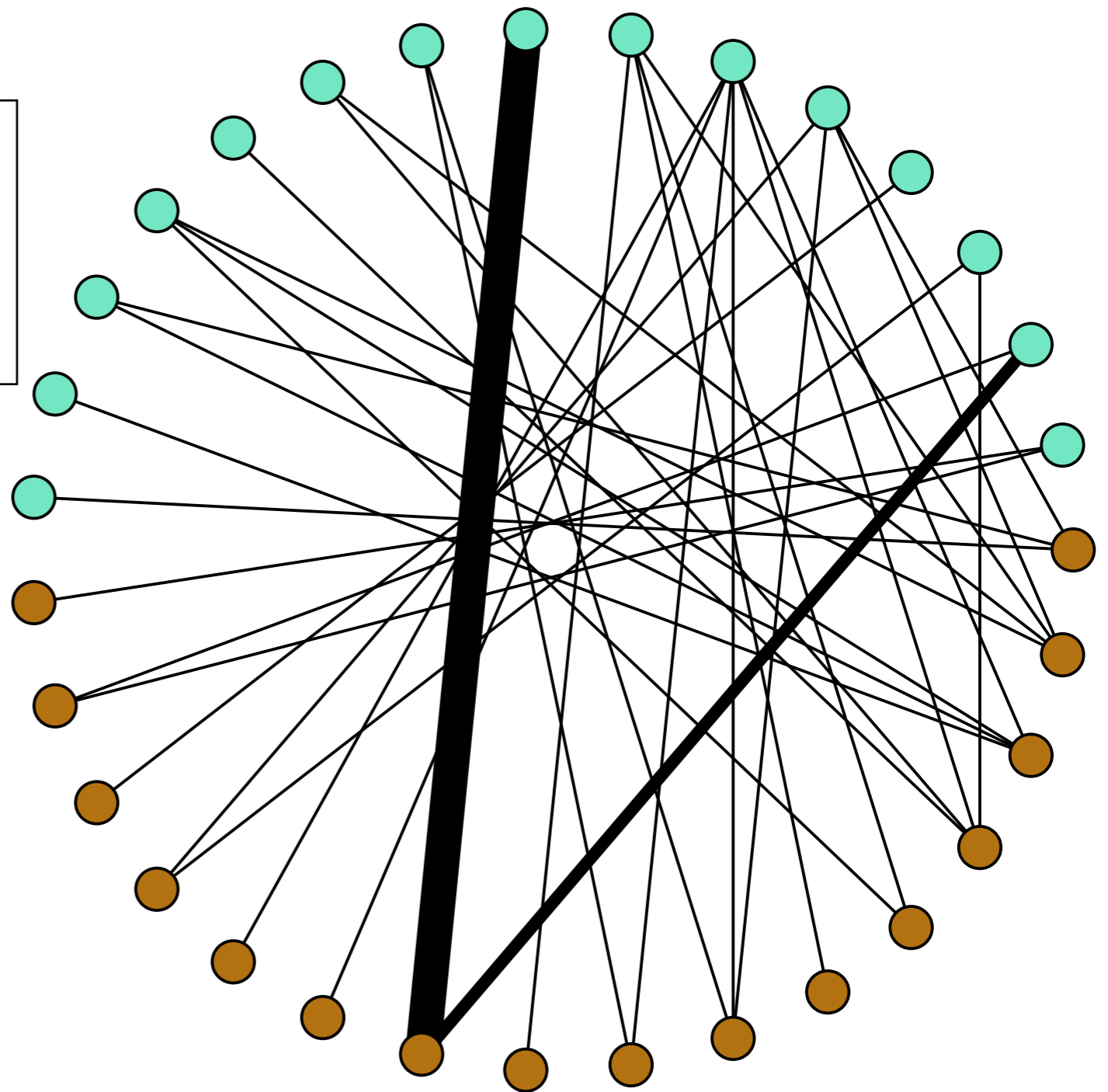
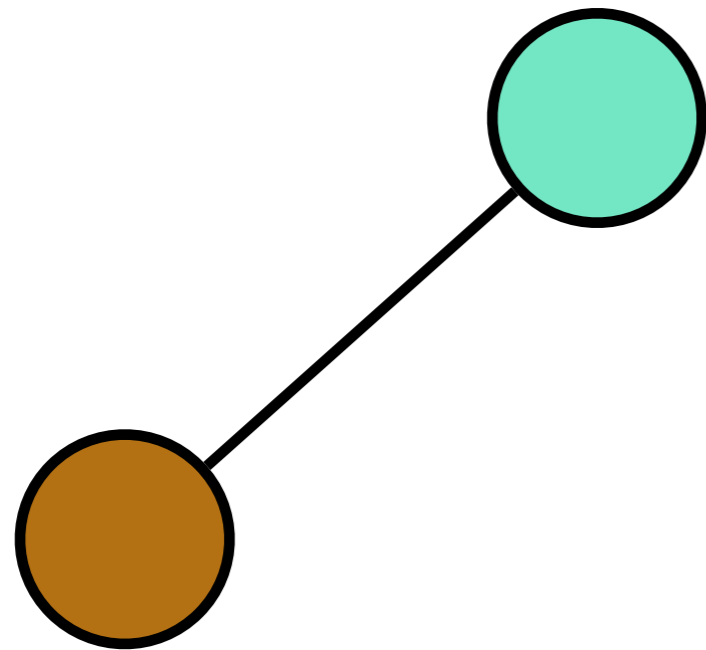
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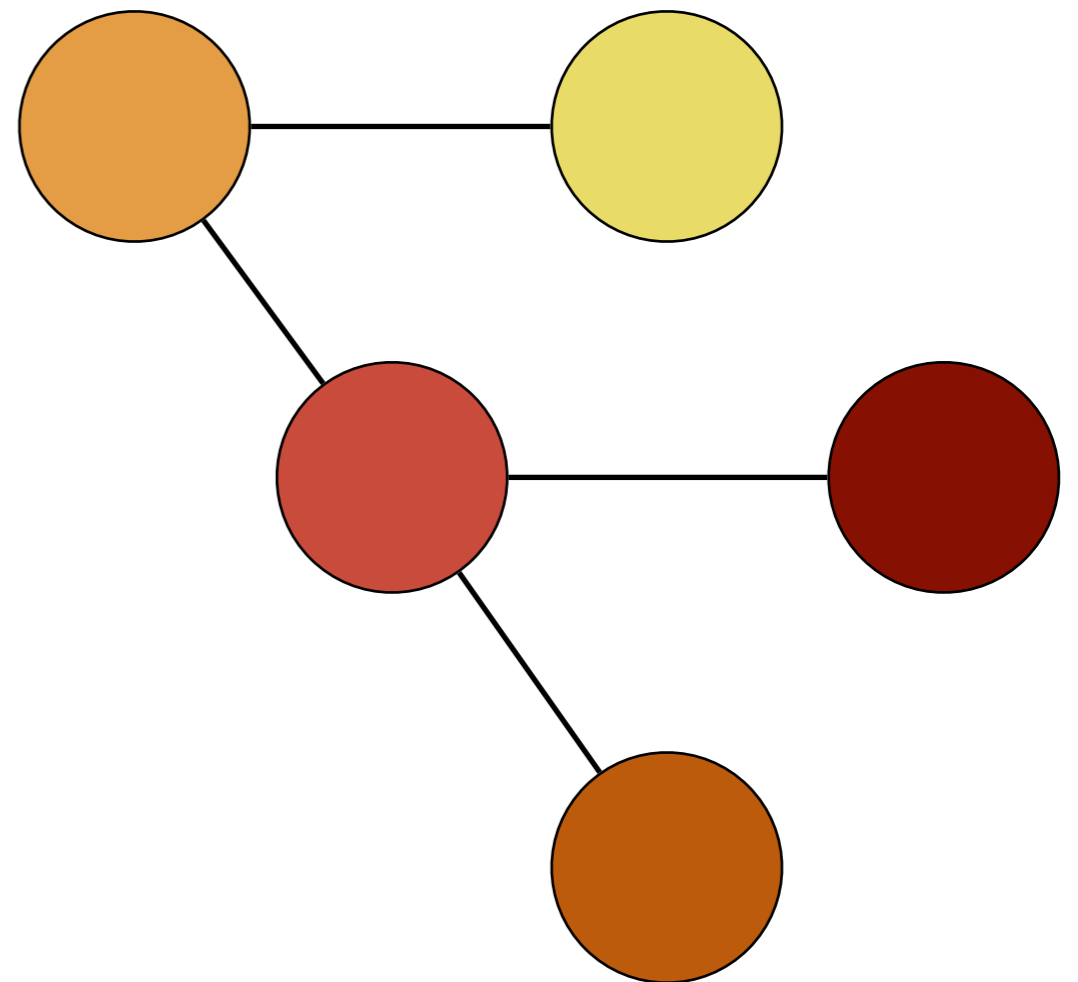
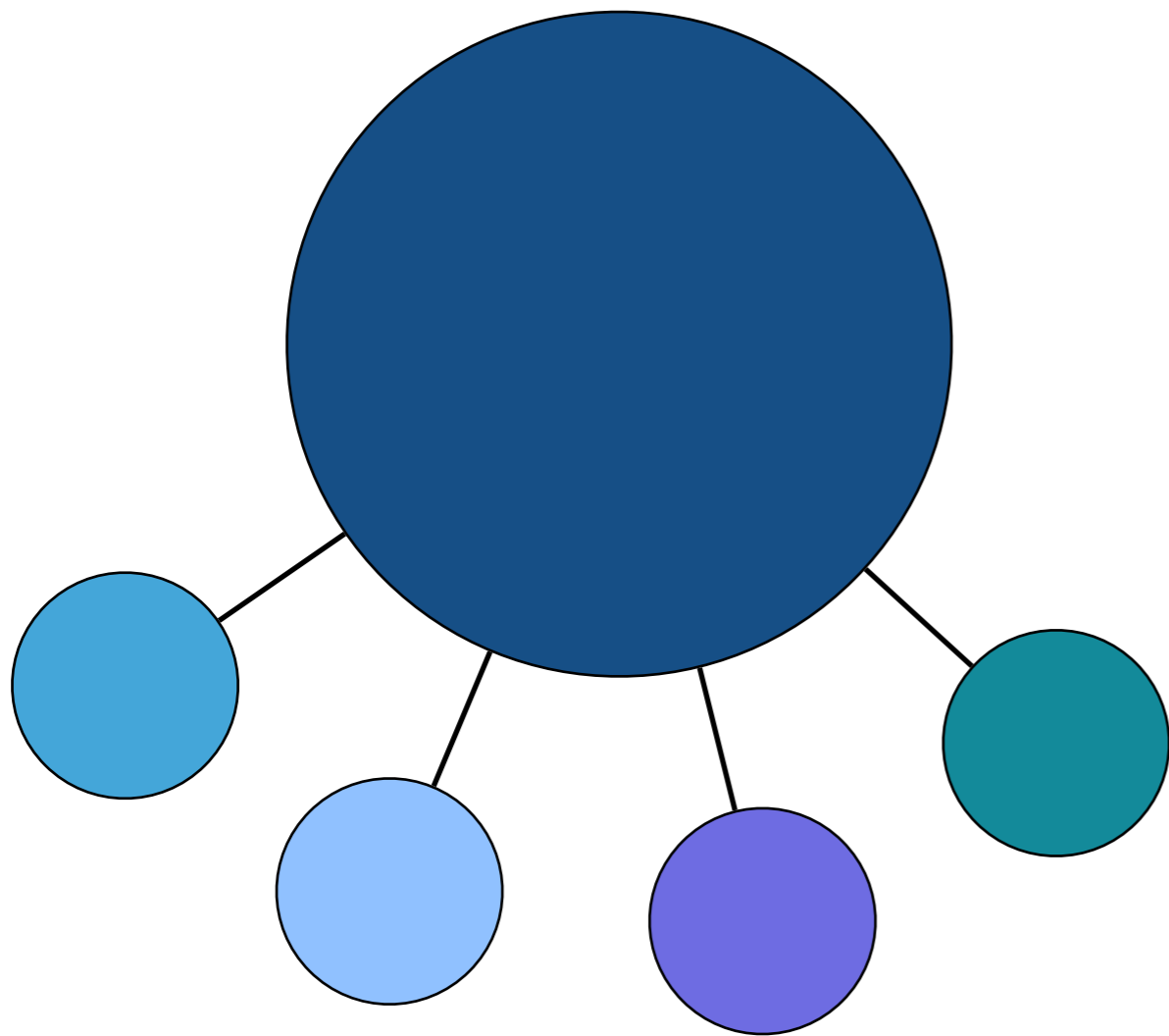
HDIM6643; laupLSAG_10
beating Melicope 65 sec.
09:25–10:08; 2015-oct-1
B. Cote coll.

CHALLENGE

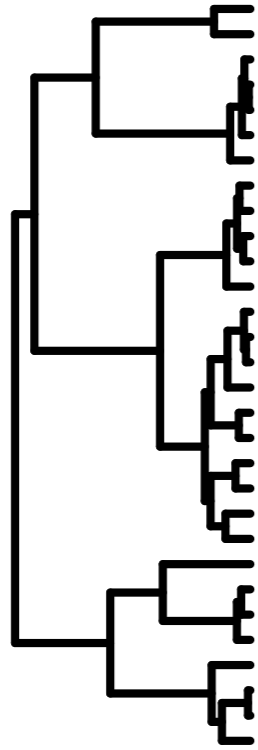


Use case: Macro eco-evo

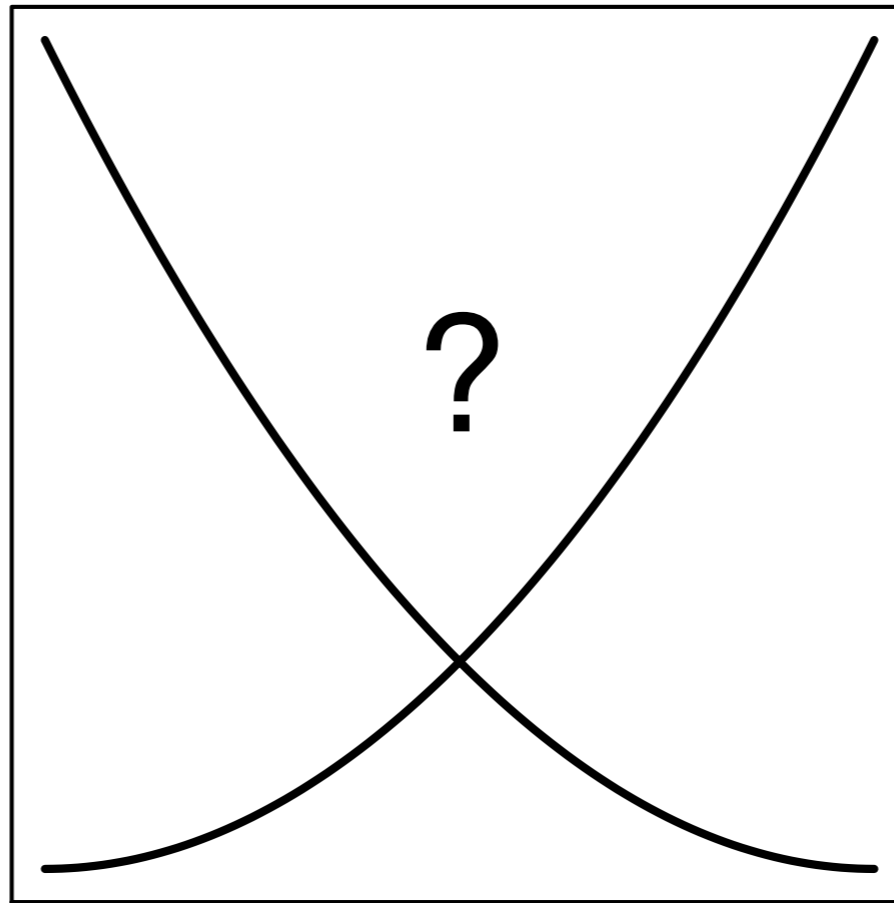
Do widespread species distributions promote or inhibit diversification?



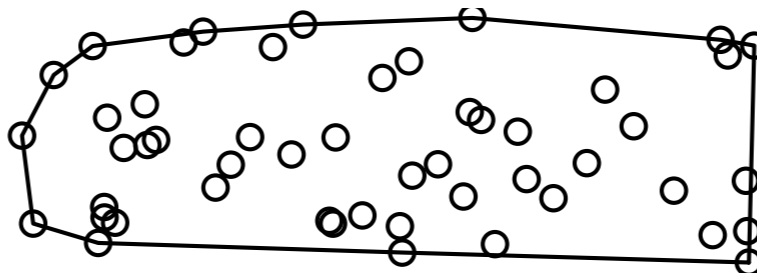
Use case: Macro eco-evo



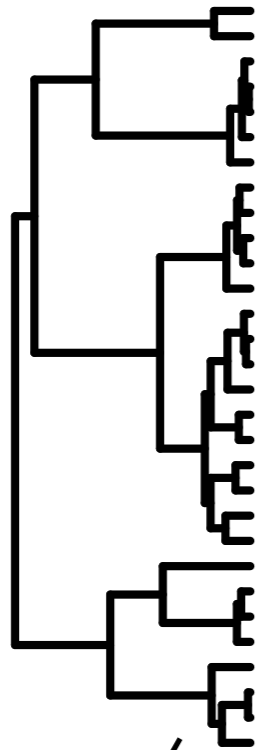
Clade speciation rate



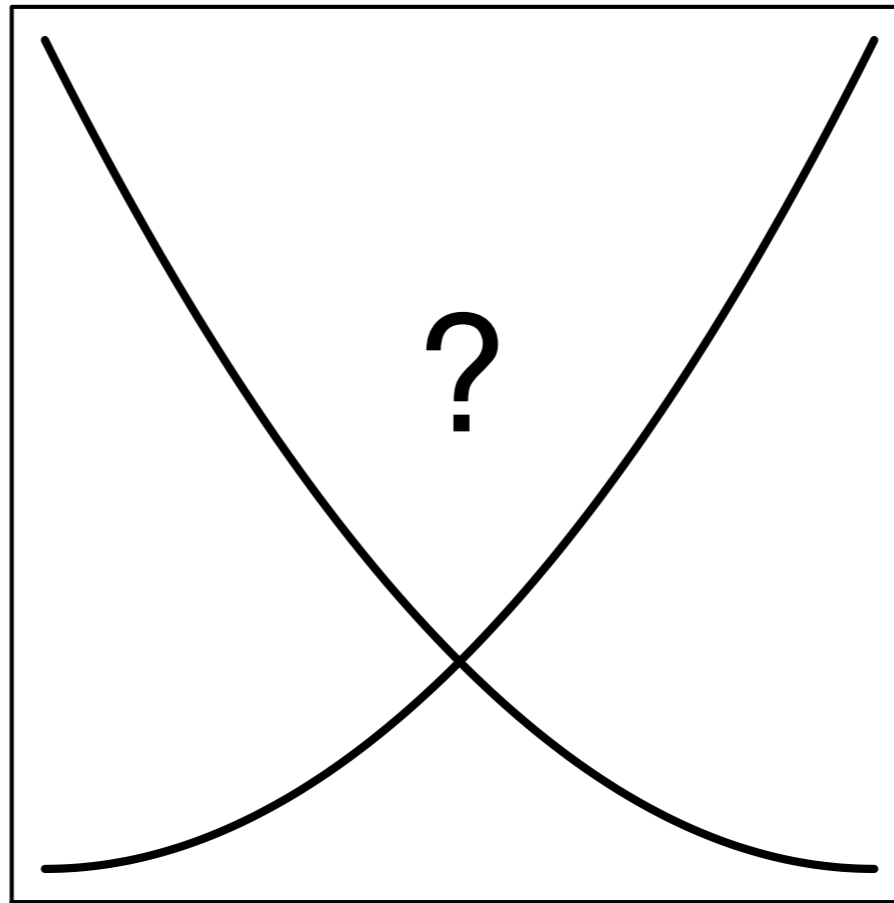
Clade range size



Use case: Macro eco-evo

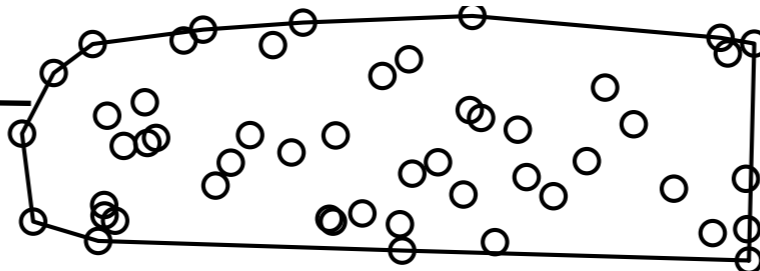


Clade speciation rate

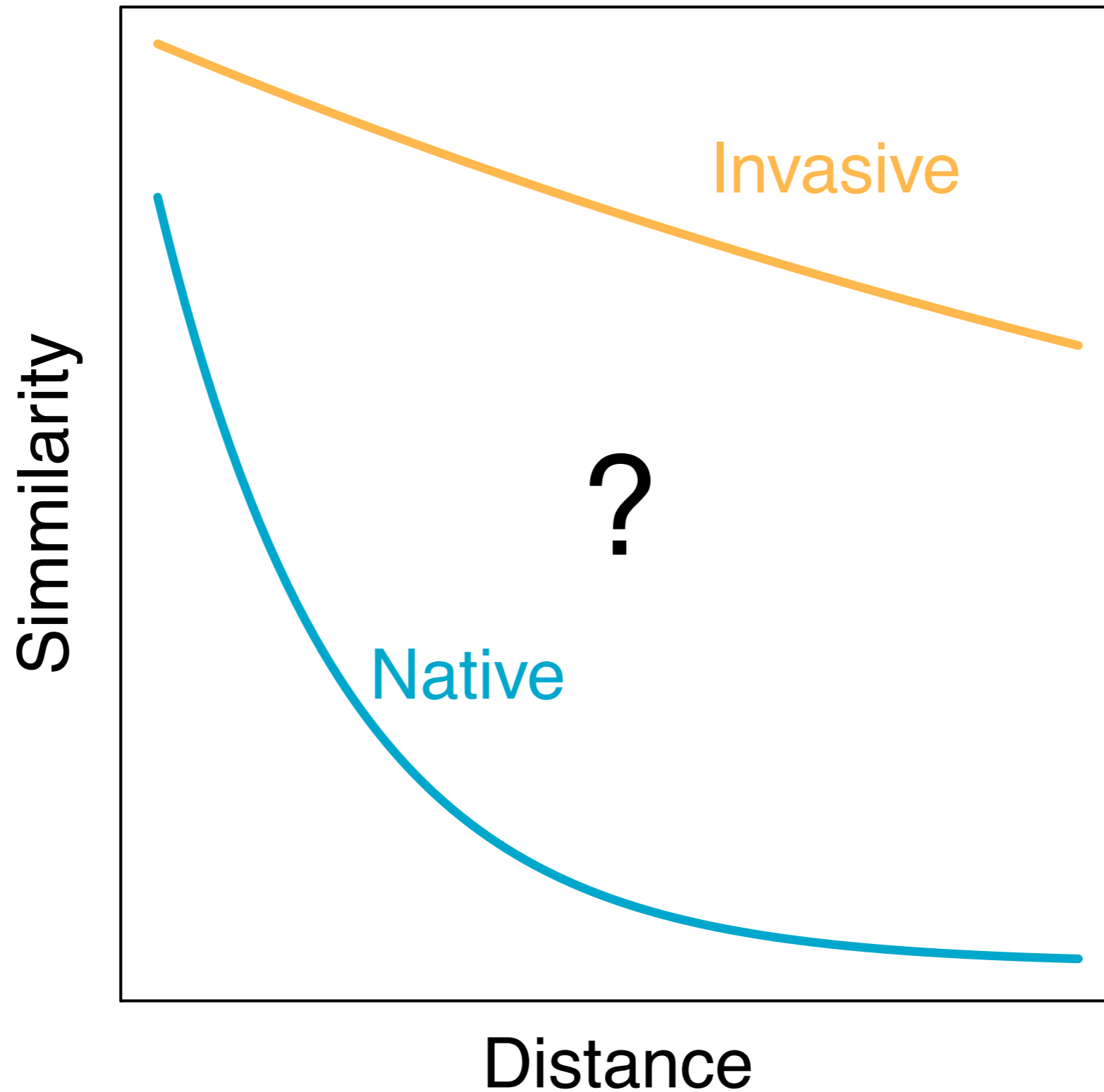


Clade range size

CHALLENGE



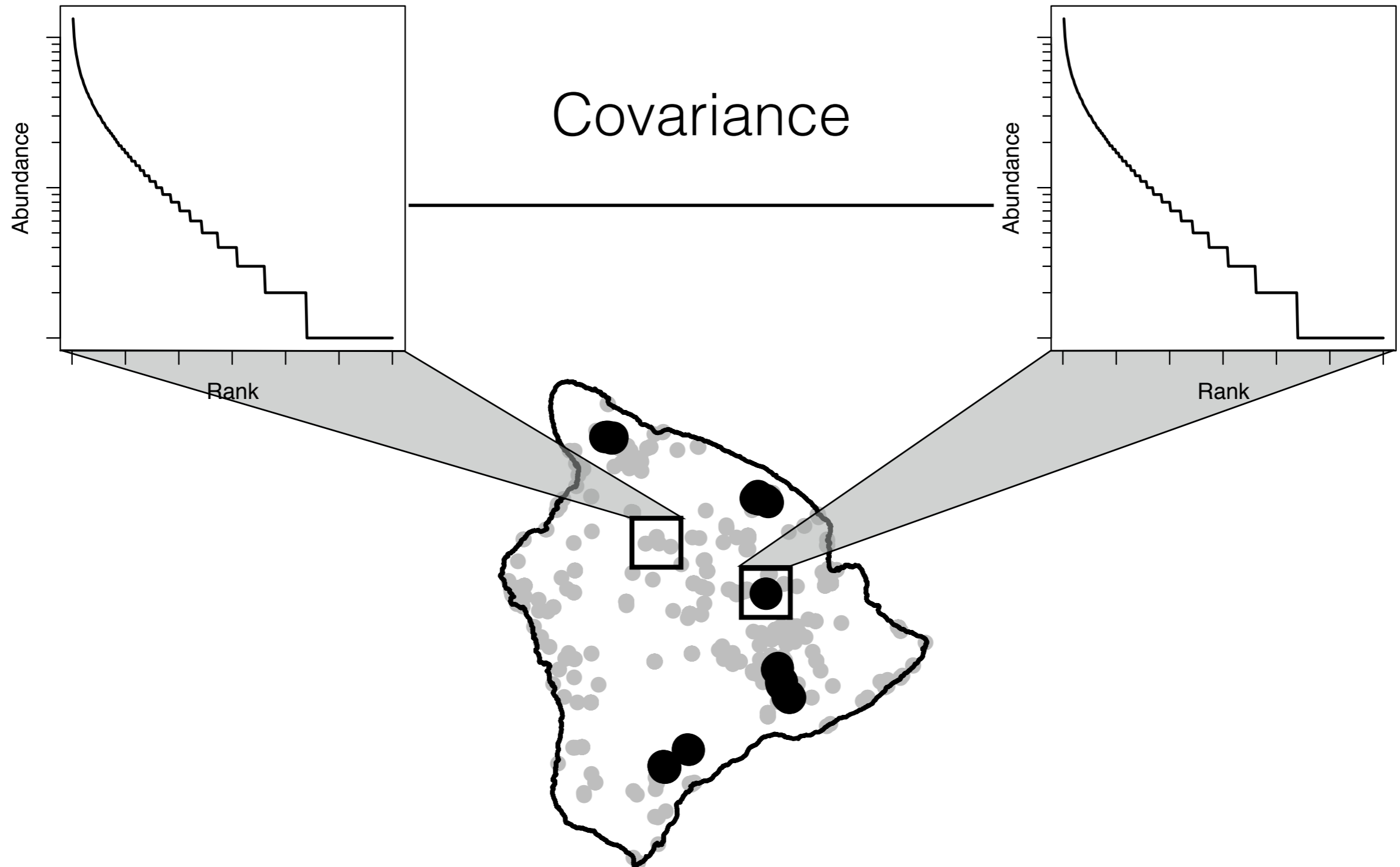
Use case: Invasion β



Use case: Invasion β

CHALLENGE

Use case: Invasion β

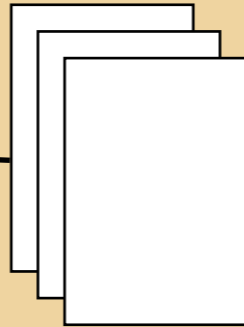


Digitization

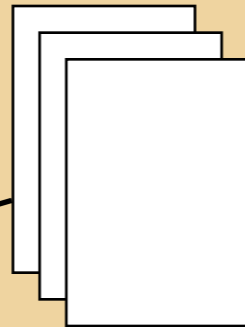


Databases/API

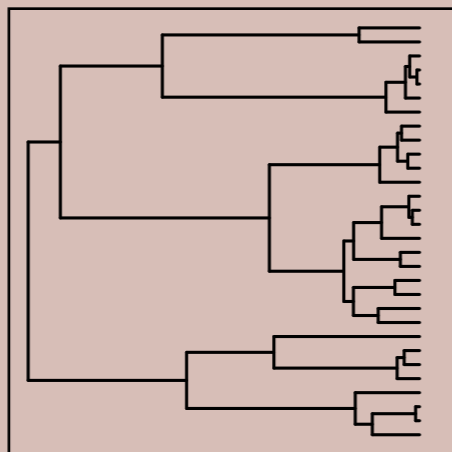
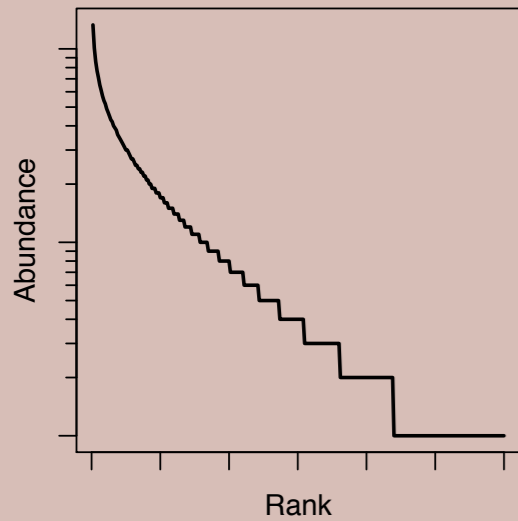
e.g. iDigBio



e.g. Dryad



Published studies



e.g. TreeBase

R packages

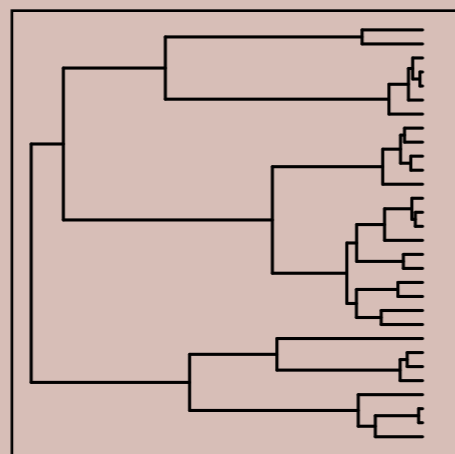
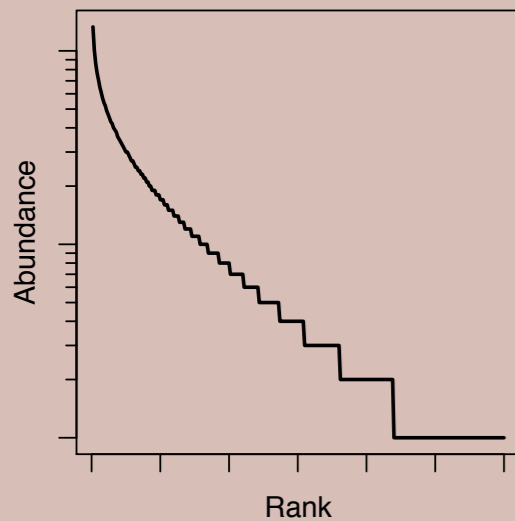
e.g. ridigbio

Query

Digitization

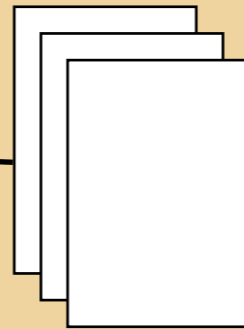


Published studies

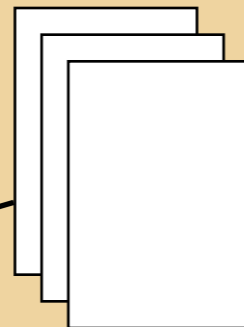


Databases/API

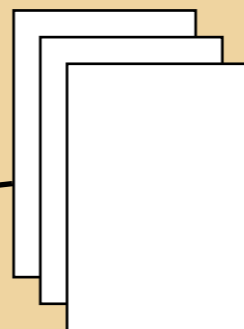
e.g. iDigBio



e.g. Dryad



e.g. TreeBase



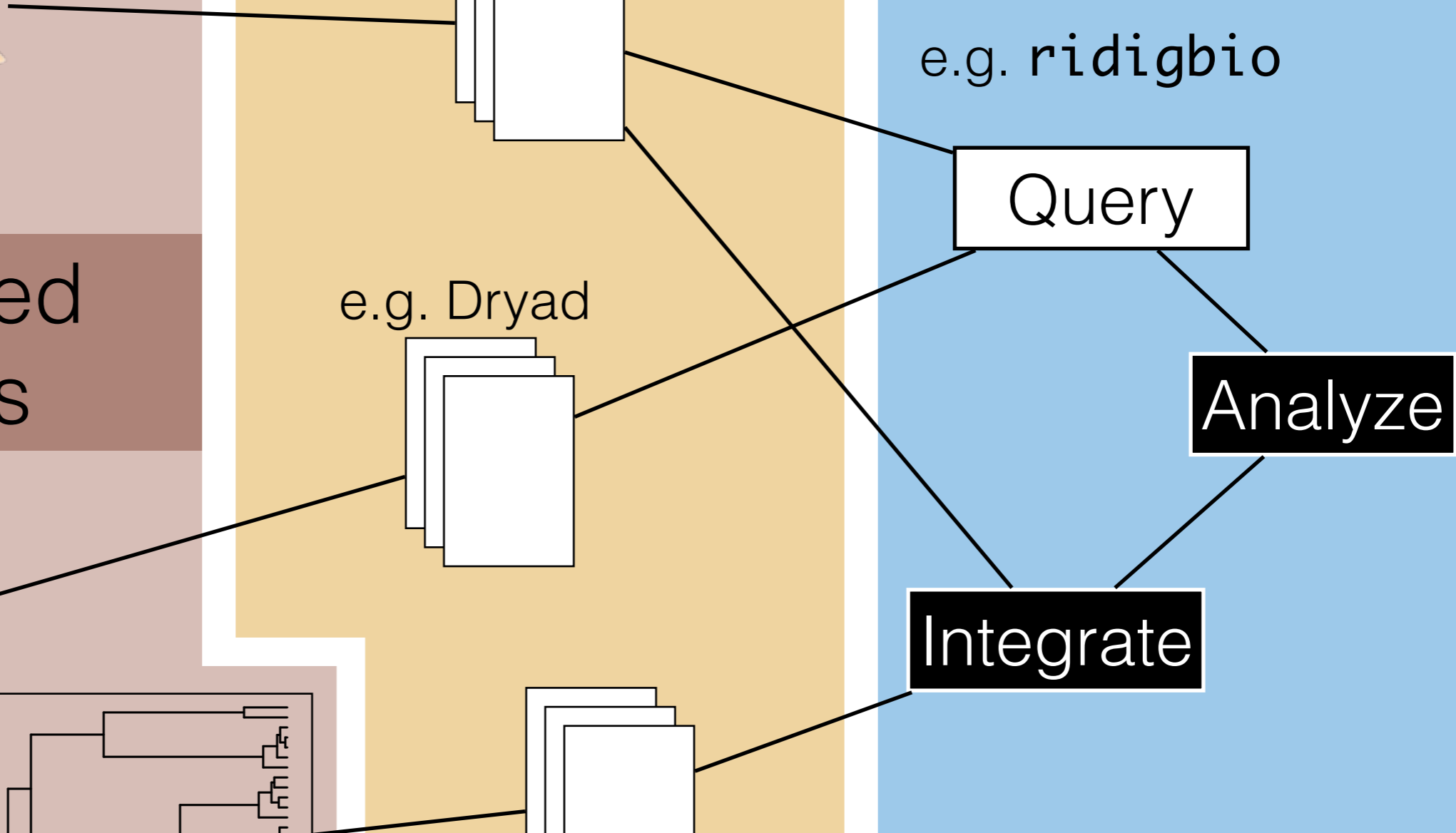
R packages

e.g. ridigbio

Query

Analyze

Integrate

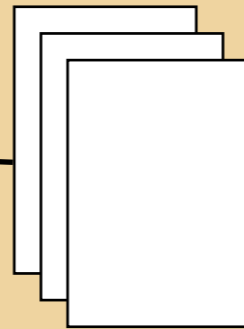


Digitization

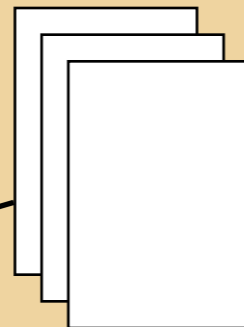


Databases/API

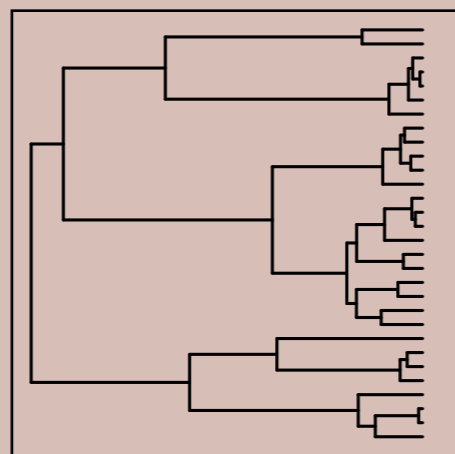
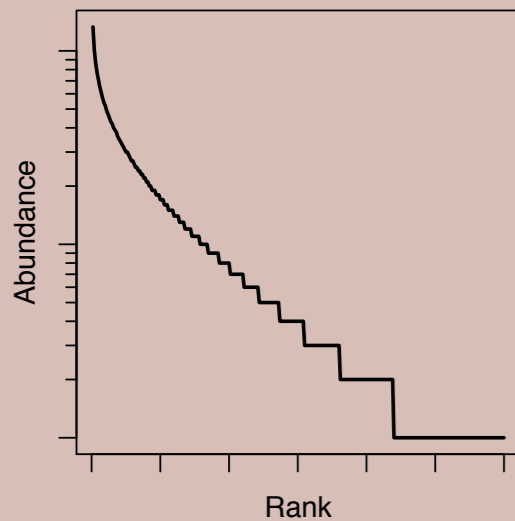
e.g. iDigBio



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