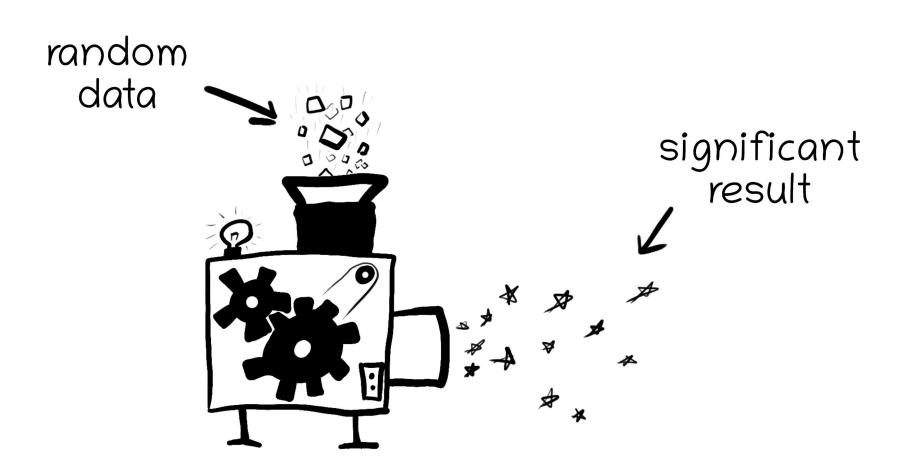
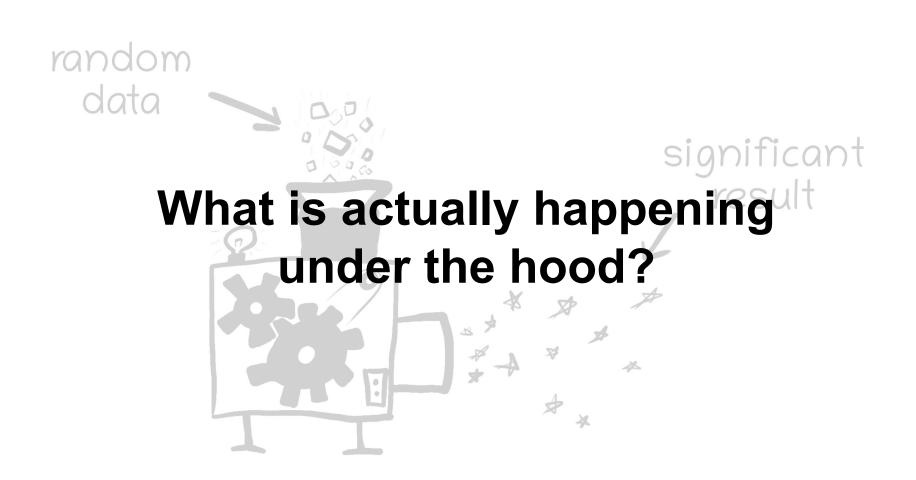
#### Hidden States SSE models

CIDs - Character Independent models

## The crisis with SSE models:



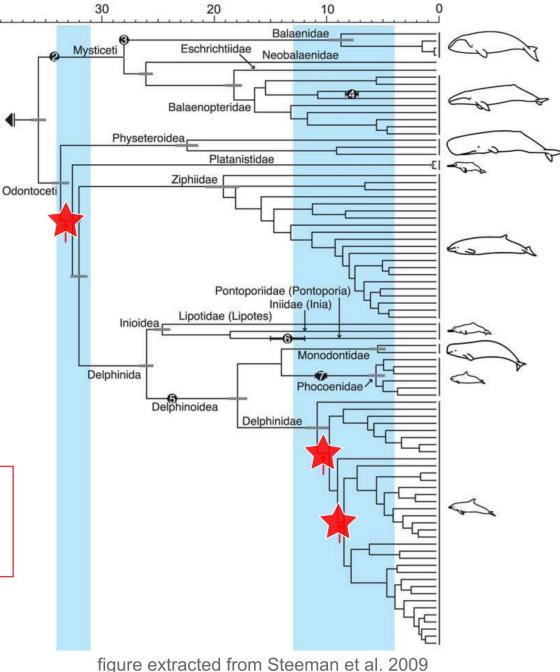
### The crisis with SSE models:

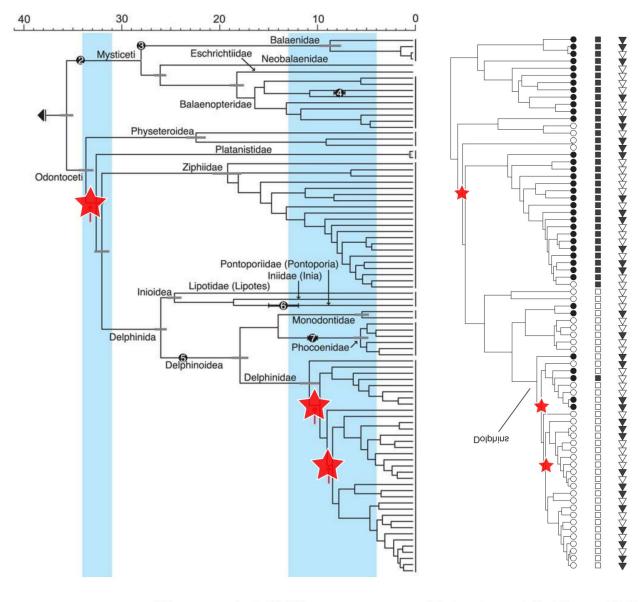


# whales & dolphins

(Steeman et al. 2009)



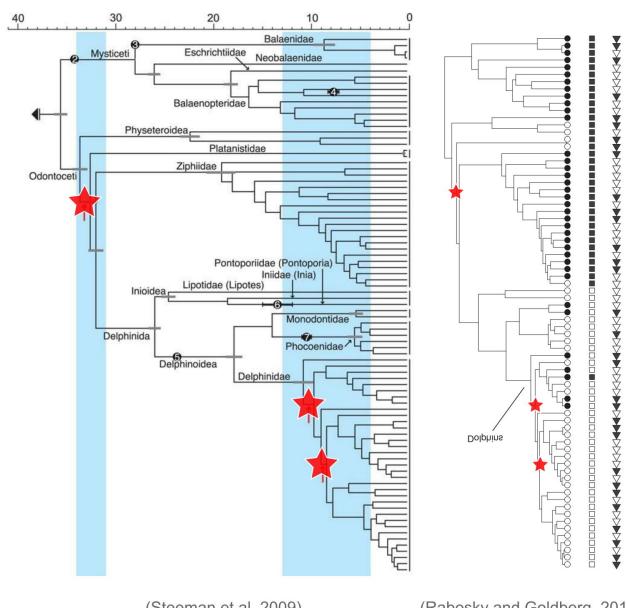




How useful is to reject a model that assumes a single rate of diversification?

(Steeman et al. 2009)

(Rabosky and Goldberg, 2015)



How useful is to reject a model that assumes a single rate of diversification?

State-dependent diversification? or Multiple rates of diversification?

(Steeman et al. 2009)

(Rabosky and Goldberg, 2015)

# Using empirical phylogenies and random traits we get a very high rate of false positives with standard BiSSE models

TABLE 1. Type I error rates for neutral character simulations on 186 empirical phylogenies of birds, squamates, amphibians, and fishes

State frequency	N	P < 0.05	P < 0.001
$0.10 \le x < 0.15$ $0.15 \le x < 0.20$ $0.20 \le x < 0.25$ $0.25 \le x < 0.30$ $0.30 \le x < 0.35$ $0.35 \le x < 0.40$ $0.40 \le x < 0.45$ $0.45 < x < 0.50$	1562 713 444 241 291 171 163 128	0.404 0.456 0.617 0.631 0.632 0.608 0.669 0.656	0.164 0.212 0.387 0.402 0.423 0.298 0.423 0.414
and you do consideration and the same of t			

Notes: Results are binned by frequency of the rarer character state and are pooled across transition rates and clades. Data are identical to those presented in Figure 5. The second column gives the number of simulated realizations. The third and fourth columns report the proportion of realizations for which state-independent diversification is rejected.

## Hidden rates SSE models

## state-dependent diversification

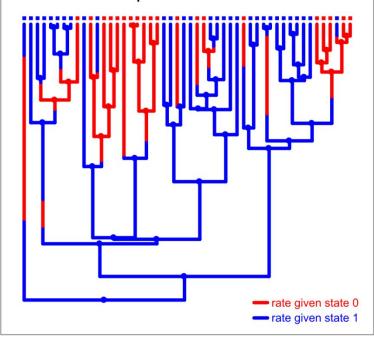
$$\lambda_{0A} = \lambda_{0B} \quad \mu_{0A} = \mu_{0B}$$

$$\mu_{0A} = \mu_{0B}$$

$$\lambda_{1A} = \lambda_{1B} \quad \mu_{1A} = \mu_{1B}$$

$$\mu_{1A} = \mu_{1B}$$

4 free diversification parameters



### state-independent diversification

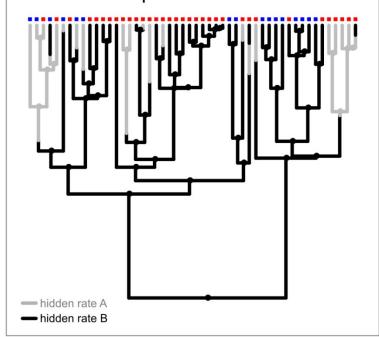
$$\lambda_{0A} = \lambda_{1A}$$

$$\lambda_{0A} = \lambda_{1A}$$
  $\mu_{0A} = \mu_{1A}$ 

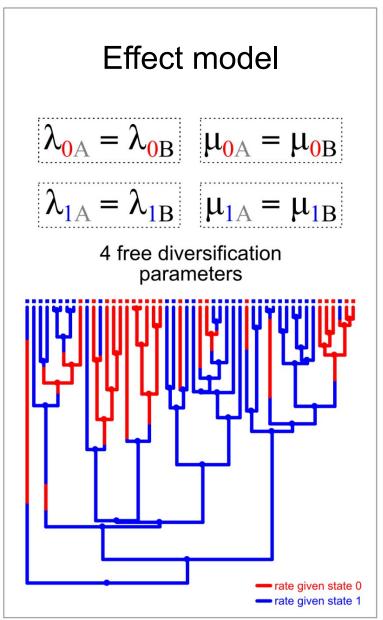
$$\lambda_{0B} = \lambda_{1B}$$

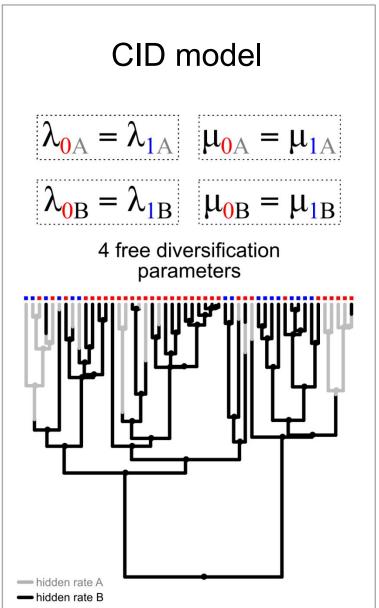
$$\lambda_{0B} = \lambda_{1B} \quad \mu_{0B} = \mu_{1B}$$

4 free diversification parameters



## Hidden rates SSE models



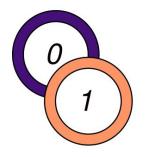


# **GeoSSE**Geographic SSE models

 $d_0$  - dispersion

 $S_0$  - speciation

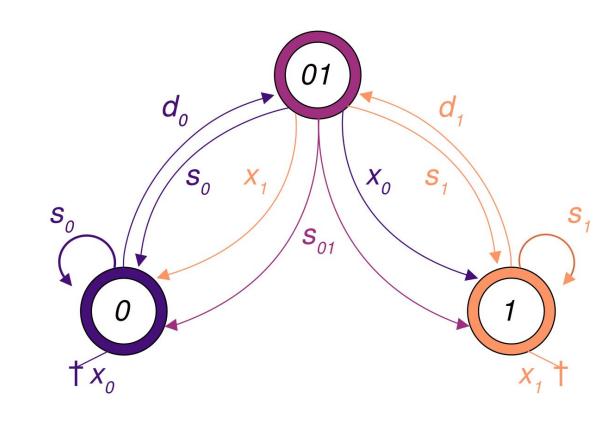
 $X_0$  - extinction

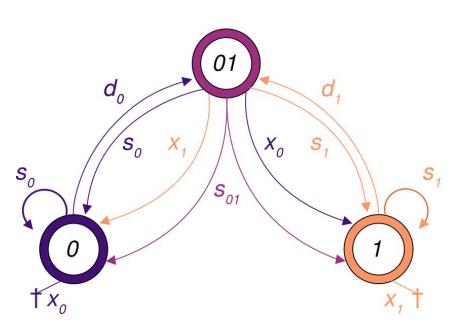


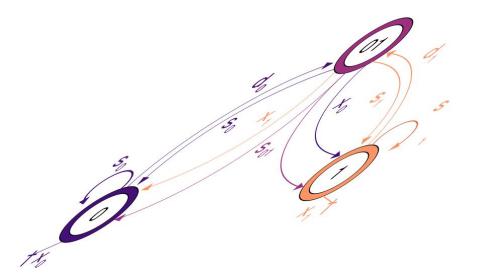
endemic areas

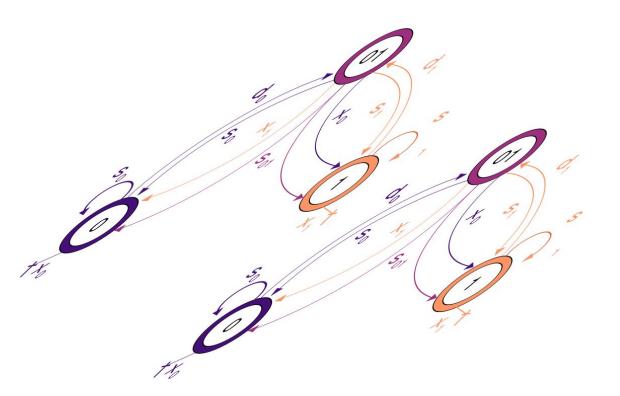


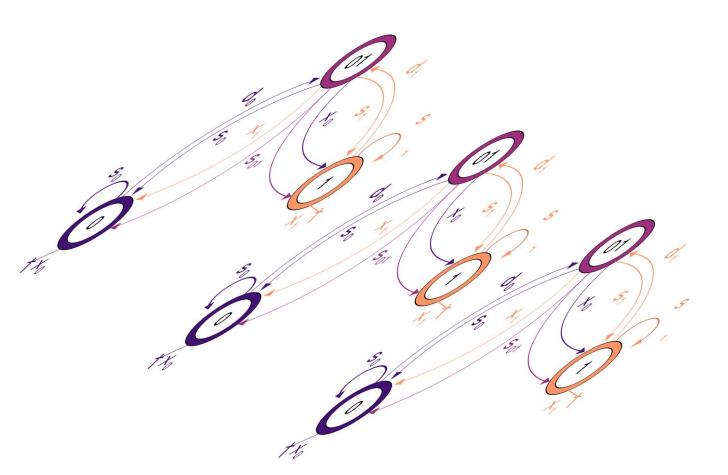
widespread areas

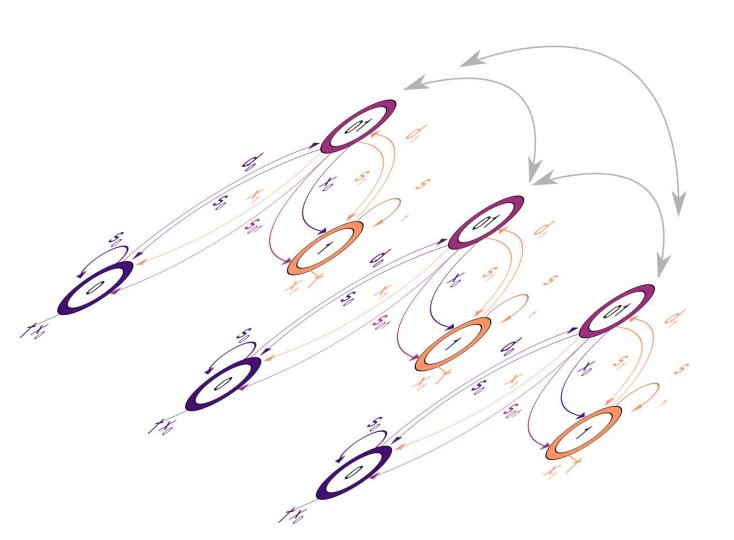


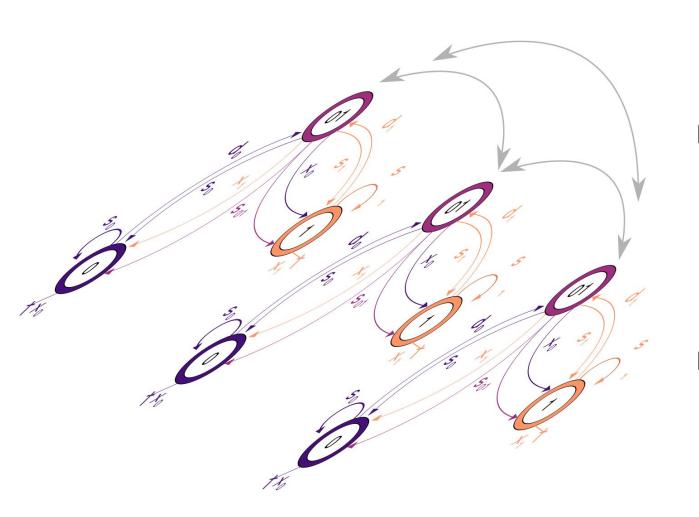












# Area-dependent diversification

partition diversification shifts between areas

# Area-independent diversification

partition diversification shifts within areas

Review of how hidden states work and what do they mean:



# Hidden state models improve state-dependent diversification approaches, including biogeographical models

Daniel S. Caetano, 1,2 D Brian C. O'Meara, and Jeremy M. Beaulieu 1

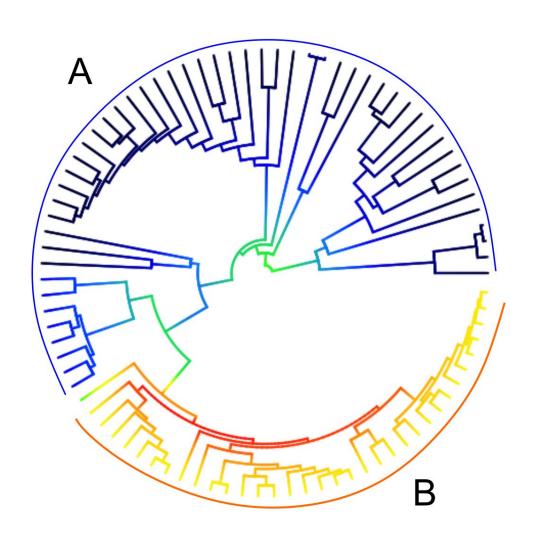
<sup>1</sup>Department of Biological Sciences, University of Arkansas, Fayetteville, Arkansas 72701

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<sup>3</sup>Department of Ecology and Evolutionary Biology, University of Tennessee, Knoxville, Tennessee 37996-1610

# Common questions when using Hidden states SSE models

# "... hidden state model was favored, thus there are unobserved traits influencing diversification."



there are *always* multiple traits influencing diversification

hidden states are a tool to accomodate variation in rates across the tree

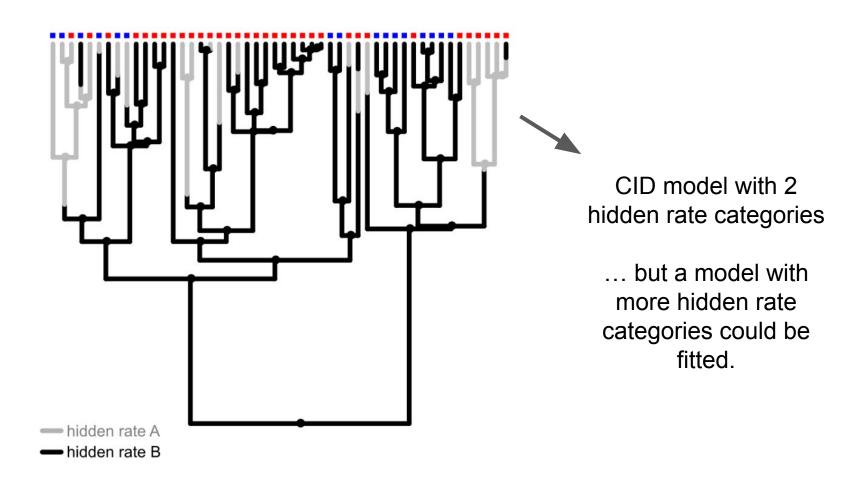
hidden states *simplify* the real variation on rates

# "Too many models to work with!" or

"Model A and model B have the very similar AIC values."

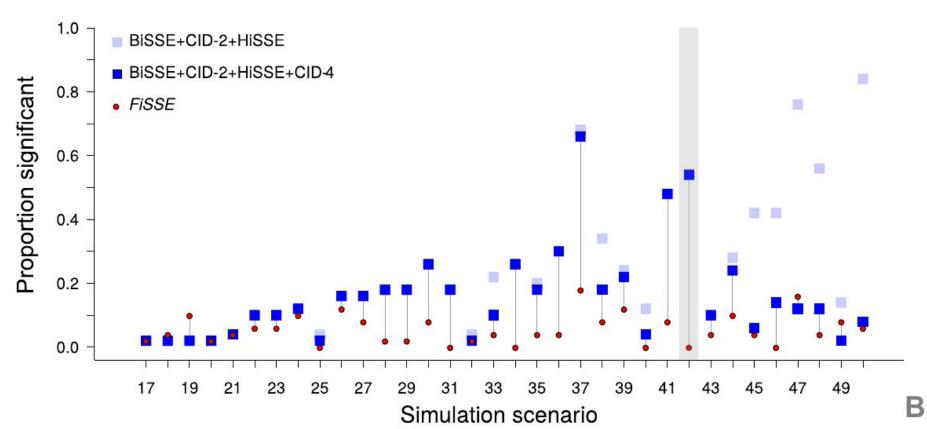
Model averaging allows to take into account multiple models that explain well the observed data.

"How many hidden rate categories do we need to use?"



## The number of **hidden rate categories** needs to match the **rate categories of the state-dependent model**

#### State-independent scenarios







### State 0

hidden rate A



hidden rate B



## State 1

hidden rate A



hidden rate B



### State 0

hidden rate A



hidden rate B



## State 1

hidden rate A



hidden rate B





Hidden states **do not** occur with the same frequency across the tree.

The weight of a hidden rate is proportional to the frequency of that hidden state in the data.

We can use equilibrium frequencies to compute the expected frequency of each hidden rate among tips and nodes.

### State 0

hidden rate A



hidden rate B



### State 1

hidden rate A



hidden rate B

