

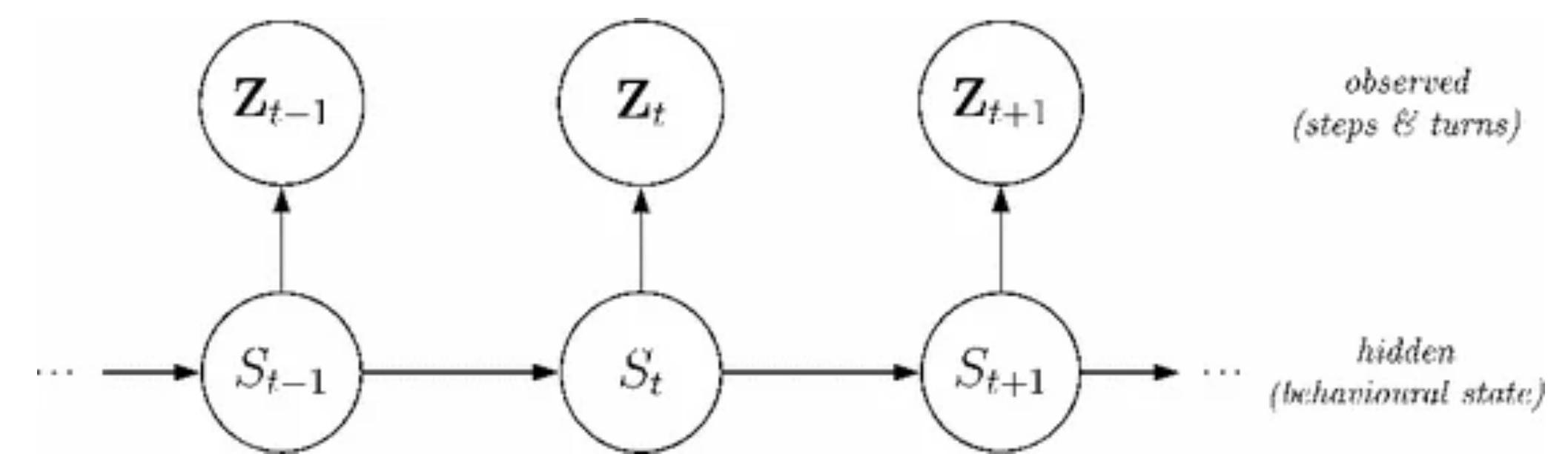
HIDDEN MARKOV MODELS FOR MOVEMENT ECOLOGY



Josh Cullen
September 8, 2022

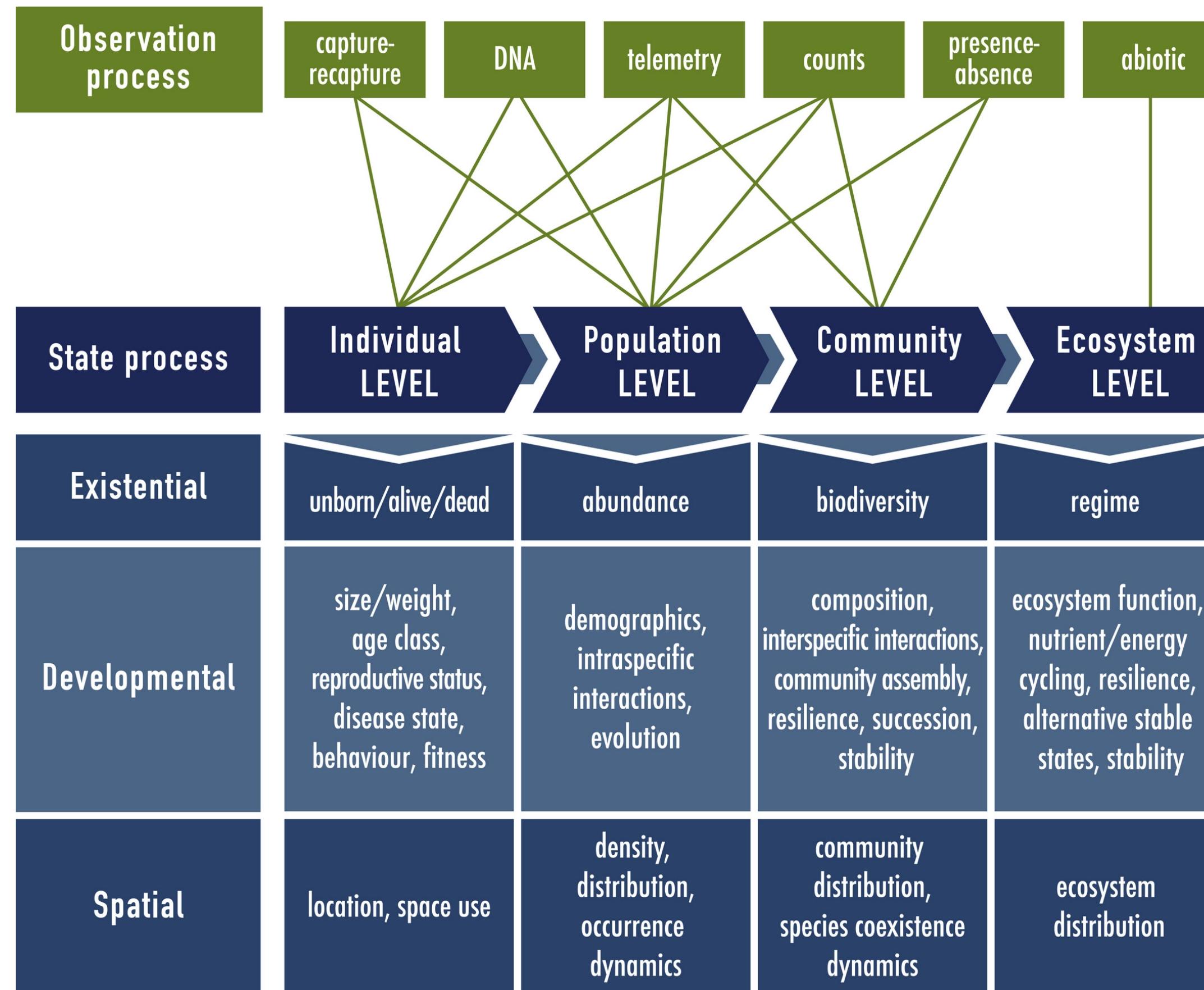
What is a hidden Markov model (HMM)?

- **McClintock et al. (2020)**: “A special class of state-space model with a finite number of hidden states that typically assumes some form of Markov property and the conditional independence property.”
- **Langrock et al. (2012)**: “Discrete-time hidden Markov models (HMMs) achieve considerable computational gains by focusing on observations that are regularly spaced in time, and for which the measurement error is negligible. These conditions are often met, in particular for data related to terrestrial animals, so that a likelihood-based HMM approach is feasible.”
- **Patterson et al. (2009)**: “...HMMs provide a statistically rigorous framework for incorporating covariates, for allowing for the autocorrelation commonly encountered in [animal telemetry] data, and for making inferences about behavioural states.”
- **Patterson et al. (2017)**: For HMMs, “one typically considers bivariate time series comprising step lengths and turning angles, regularly spaced in time and assumed to be observed with no or only negligible error.”



Patterson et al. 2017

What is a hidden Markov model (HMM)?

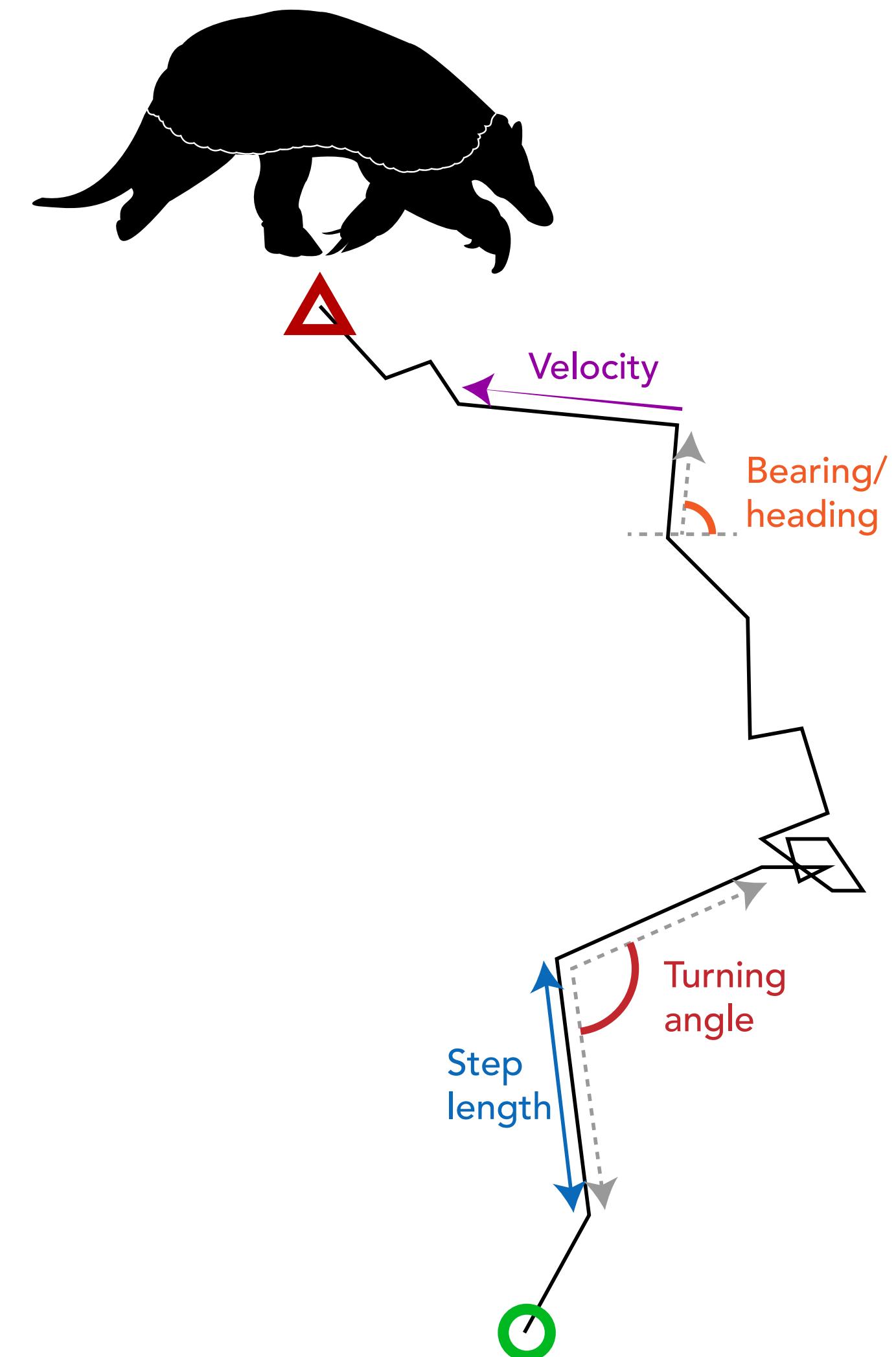


Latent variable models

	State space	
	Continuous	Discrete
Temporal dependence	State-space model	Hidden Markov model
Temporal independence	Continuous mixture model	Finite mixture model

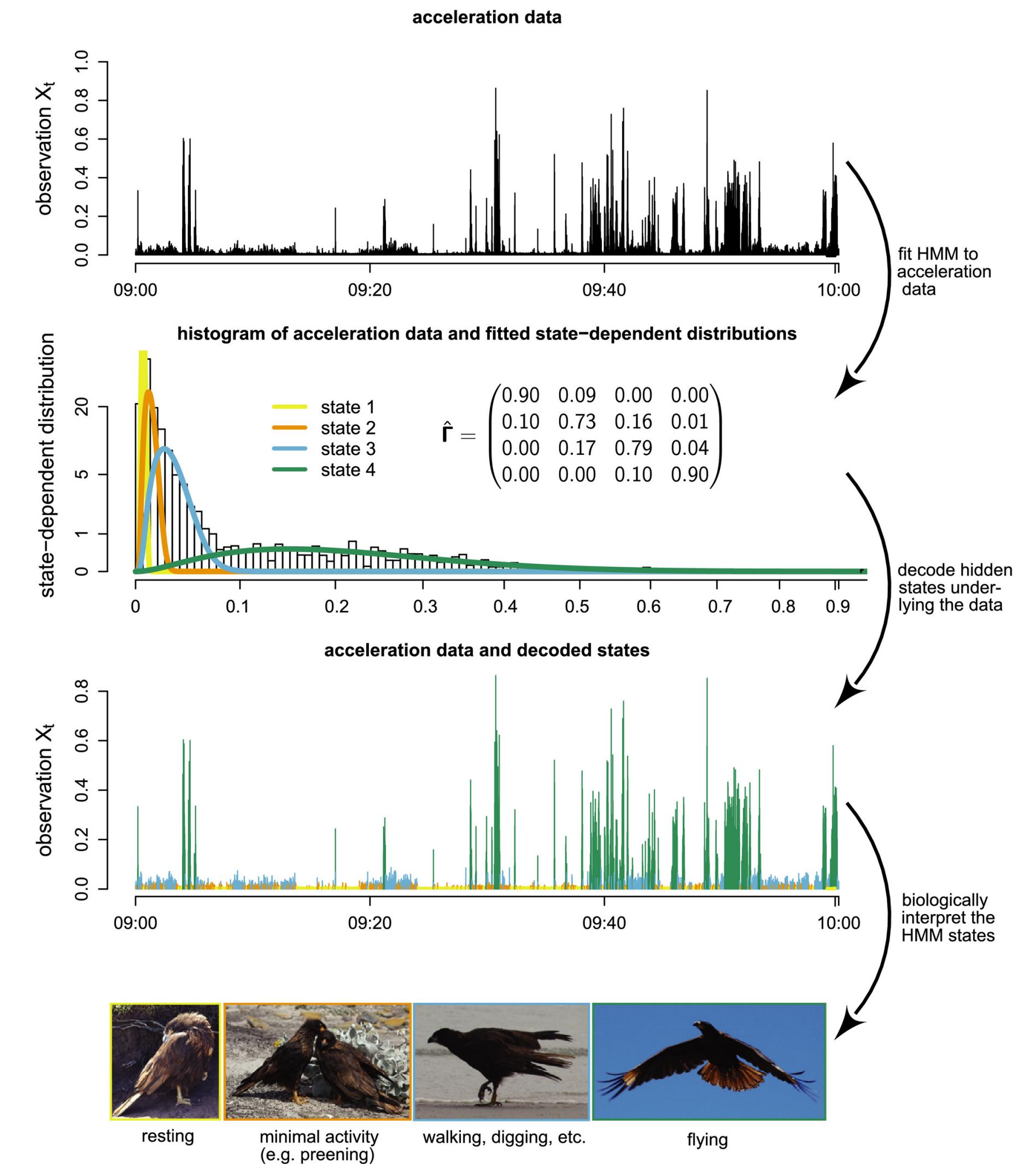
Movement metrics analyzed by HMMs

- Bivariate positions or their increments
- Distance between successive observations
- Compass direction (i.e., heading/bearing)
- Changes in direction between successive relocations



Which process models are used for HMMs?

- Almost always use discrete-time models w/ CRW
- Some options starting to become available for biased RW (BRW), biased-CRW (BCRW), and continuous-time
 - But infrequently used



Components of an HMM

1. Initial state distribution (δ)

- Defines the starting probabilities for each of the N states

2. State transition probability matrix (t.p.m.; Γ)

- Defines the probabilities of remaining in (or changing) behavioral states

3. State-dependent distributions ($f_i(\mathbf{z}_t)$)

- Defines the distributions of the data streams that characterize each state

$$\begin{aligned}\mathcal{L}^{\text{HMM}}(\theta) &= f(\mathbf{z}_1, \dots, \mathbf{z}_T) \\ &= \sum_{s_1=1}^N \dots \sum_{s_T=1}^N f(\mathbf{z}_1, \dots, \mathbf{z}_T | s_1, \dots, s_T) f(s_1, \dots, s_T) \\ &= \sum_{s_1=1}^N \dots \sum_{s_T=1}^N \delta_{s_1} \prod_{t=1}^T f(l_t | s_t) f(\phi_t | s_t) \prod_{t=2}^T \gamma_{s_{t-1}, s_t},\end{aligned}$$

Model parameters
Realizations of state-dependent process for $t \in [1, T]$
Probability of being in behavioral state s at time t

Patterson et al. 2017

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Example of 2-state HMM

Behavioral state
at time $t = 1$

$$\delta = [Pr(S_1 = 1), Pr(S_1 = 2)]$$

Behavioral state
index for all N states

$$\sum_{i=1}^N \delta_i = 1$$

Probabilities must sum to 1

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Example of 2-state HMM

Probability of switching from state j to i

$$\Gamma = (\gamma_{ij}) = \begin{bmatrix} \gamma_{11} & \gamma_{12} \\ \gamma_{21} & \gamma_{22} \end{bmatrix}$$

Stay in state 1

Switch from state 2 to 1

Switch from state 1 to 2

Stay in state 2

$$\Gamma^{(t)} = (\gamma_{ij}^{(t)}) = \frac{\exp(\eta_{ij}^{(t)})}{\sum_{k=1}^N \exp(\eta_{ik}^{(t)})}$$

Multinomial logit link function

η_{ij} = linear model

Incorporate effect of covariates on state switching probability

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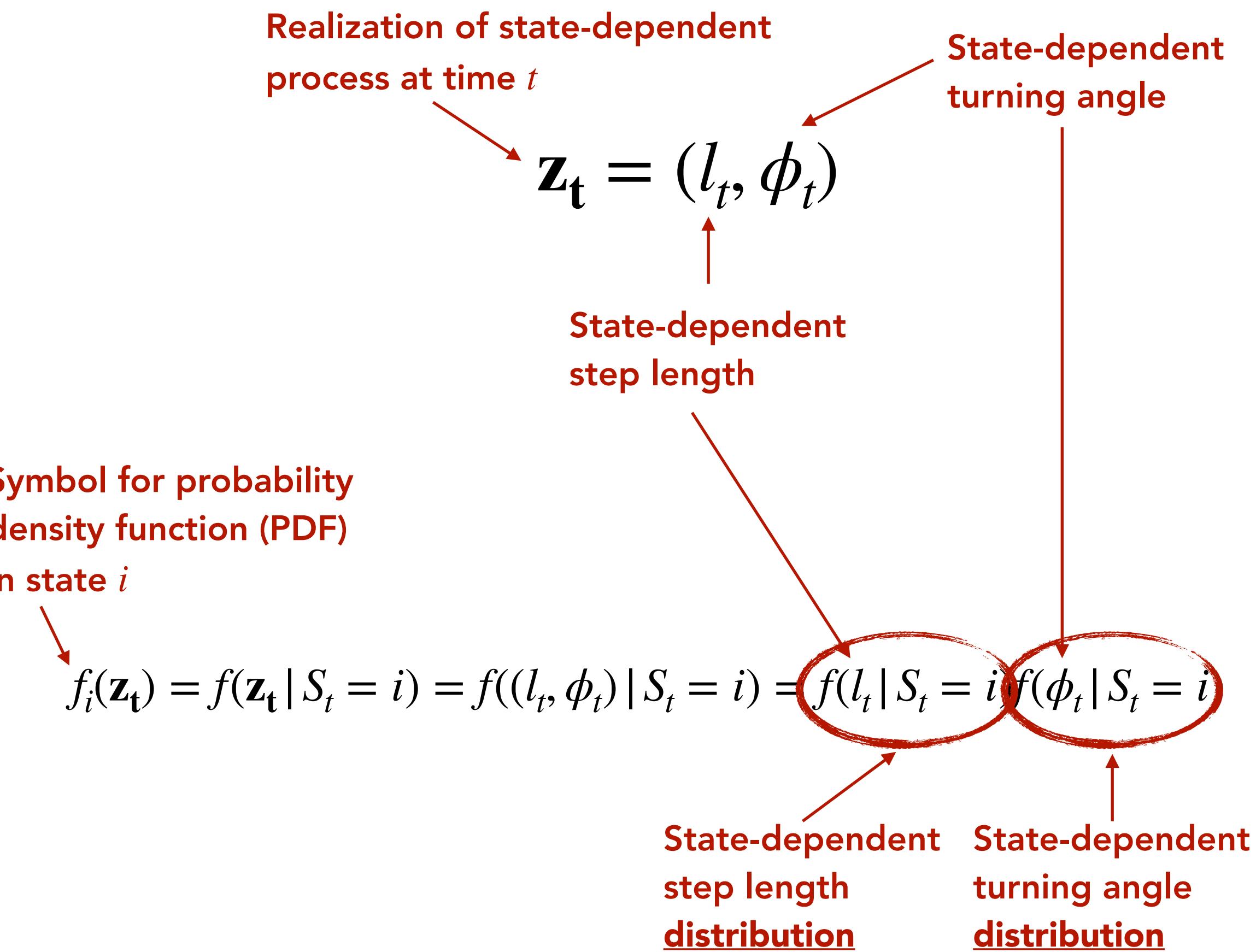
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Example of 2-state HMM

(Including only step lengths and turning angles)



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Patterson et al. 2017

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The diagram illustrates the components of the HMM likelihood function. Red arrows point from labels to specific terms in the equation:

- An arrow points from "Initial state distribution" to δ_{s_1} .
- An arrow points from "State-dependent turning angle distribution" to $f(\phi_t | s_t)$.
- An arrow points from "State-dependent step length distribution" to $f(l_t | s_t)$.
- An arrow points from "Probability of transition from state j to i " to γ_{s_{t-1}, s_t} .

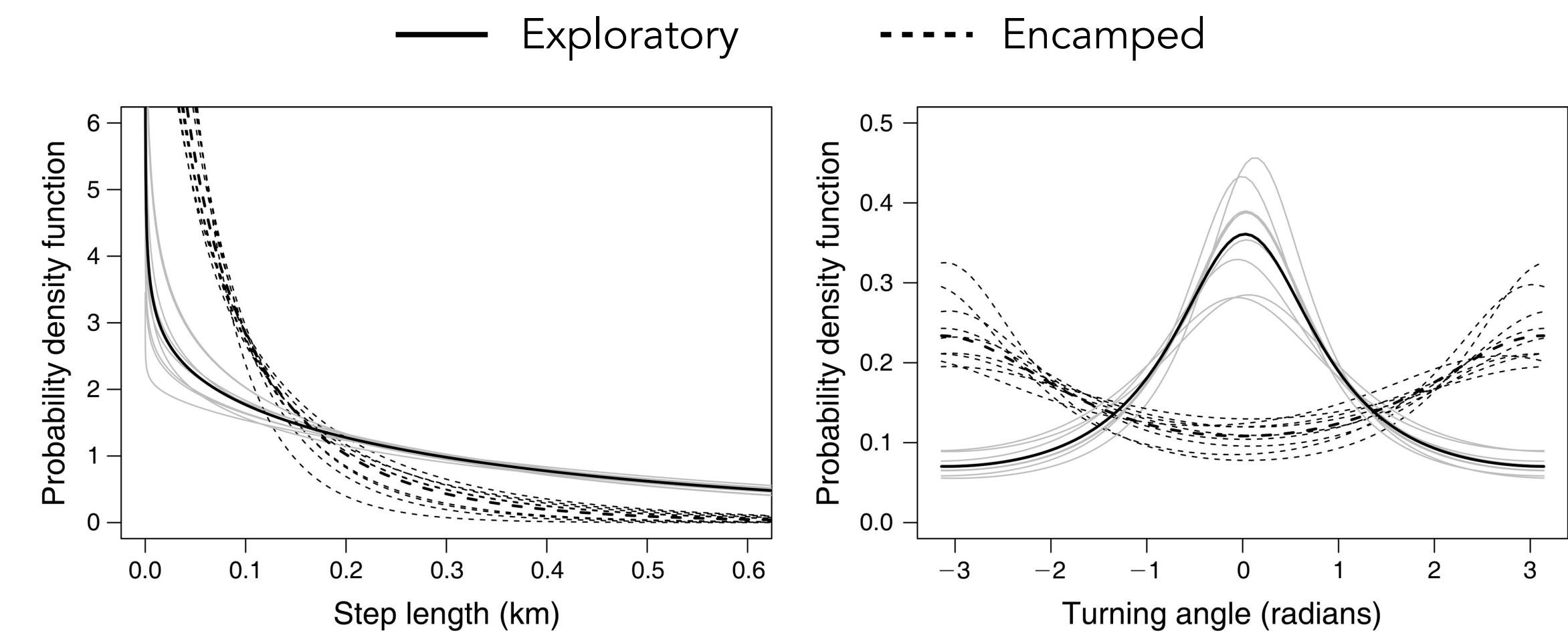
Common PDFs for movement variables

Step length

- Real, positive values that often exhibit a right-skewed distribution
- Gamma, Weibull

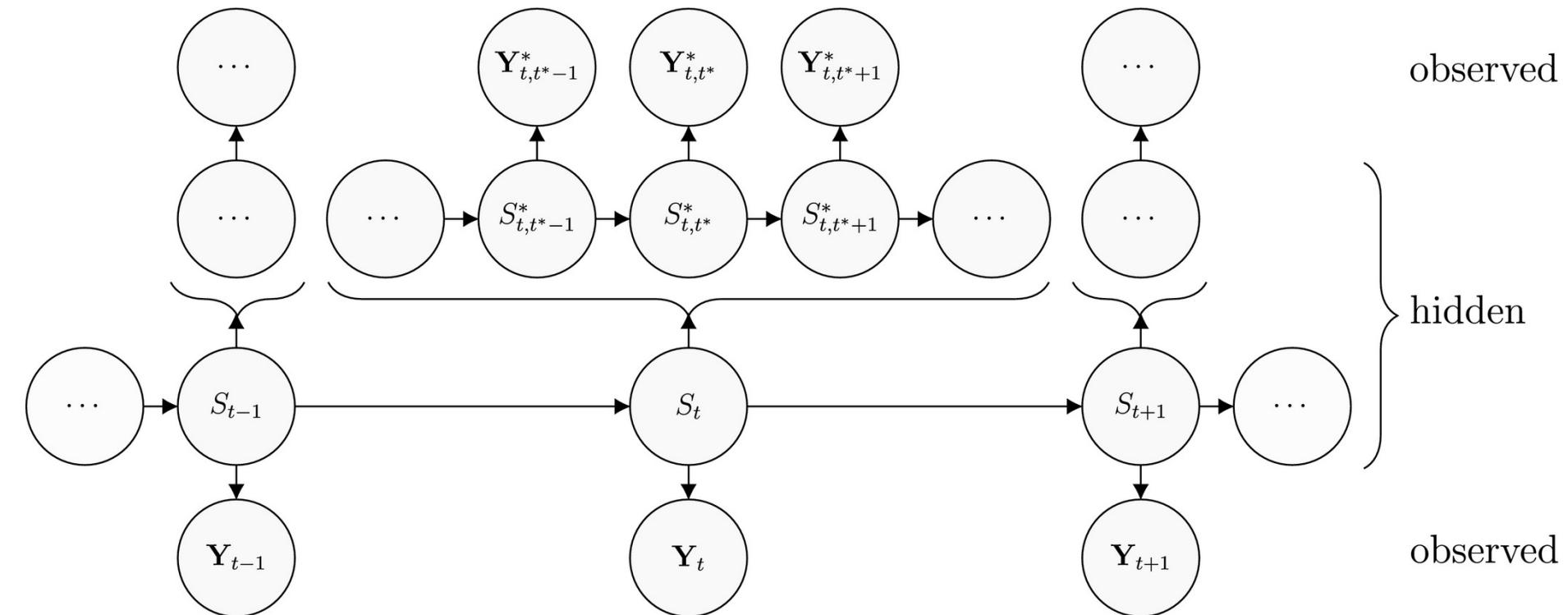
Turning angle

- Real values constrained from $-\pi$ to $+\pi$
- von Mises, wrapped Cauchy

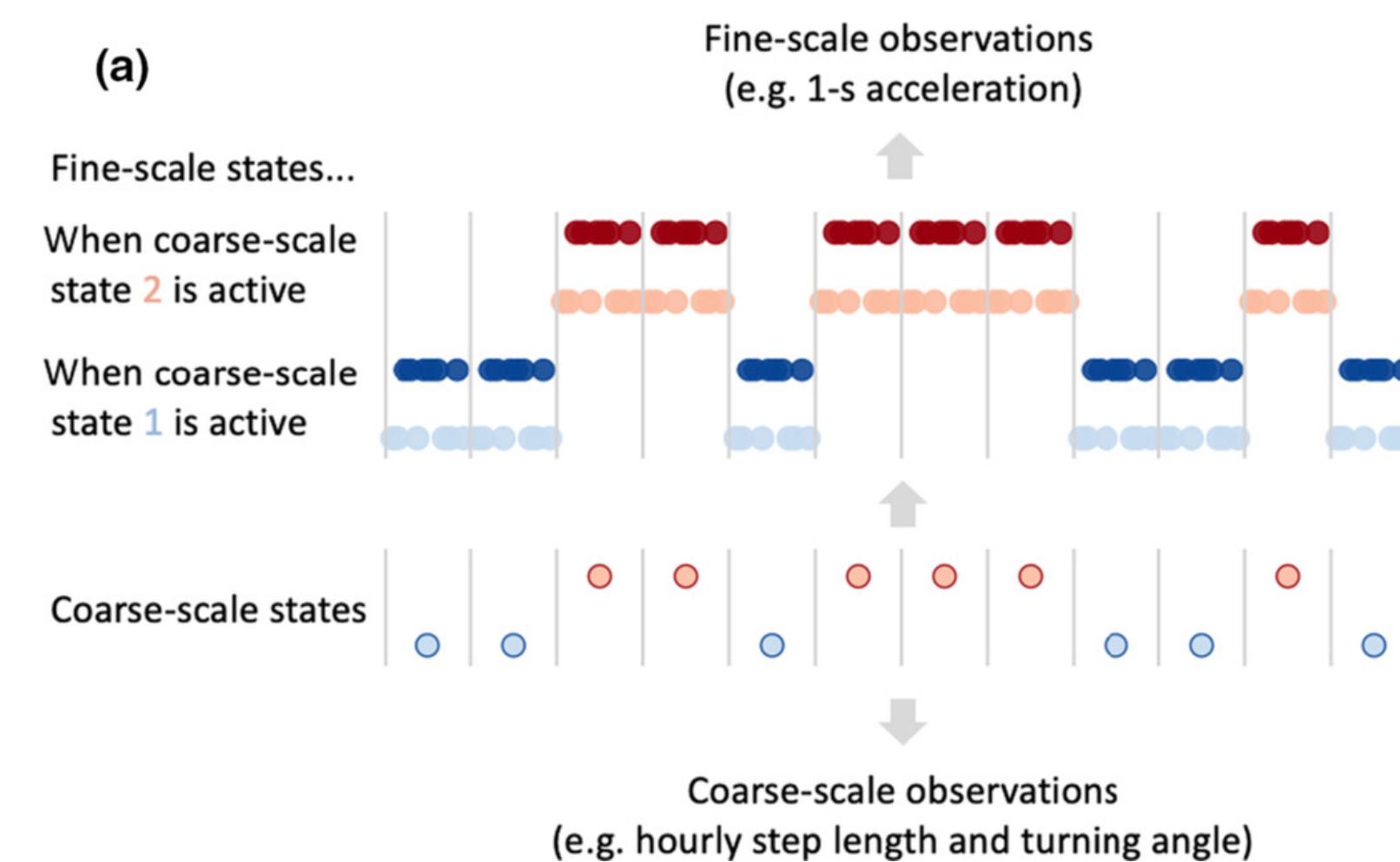


Hierarchical HMMs

Fine-scale

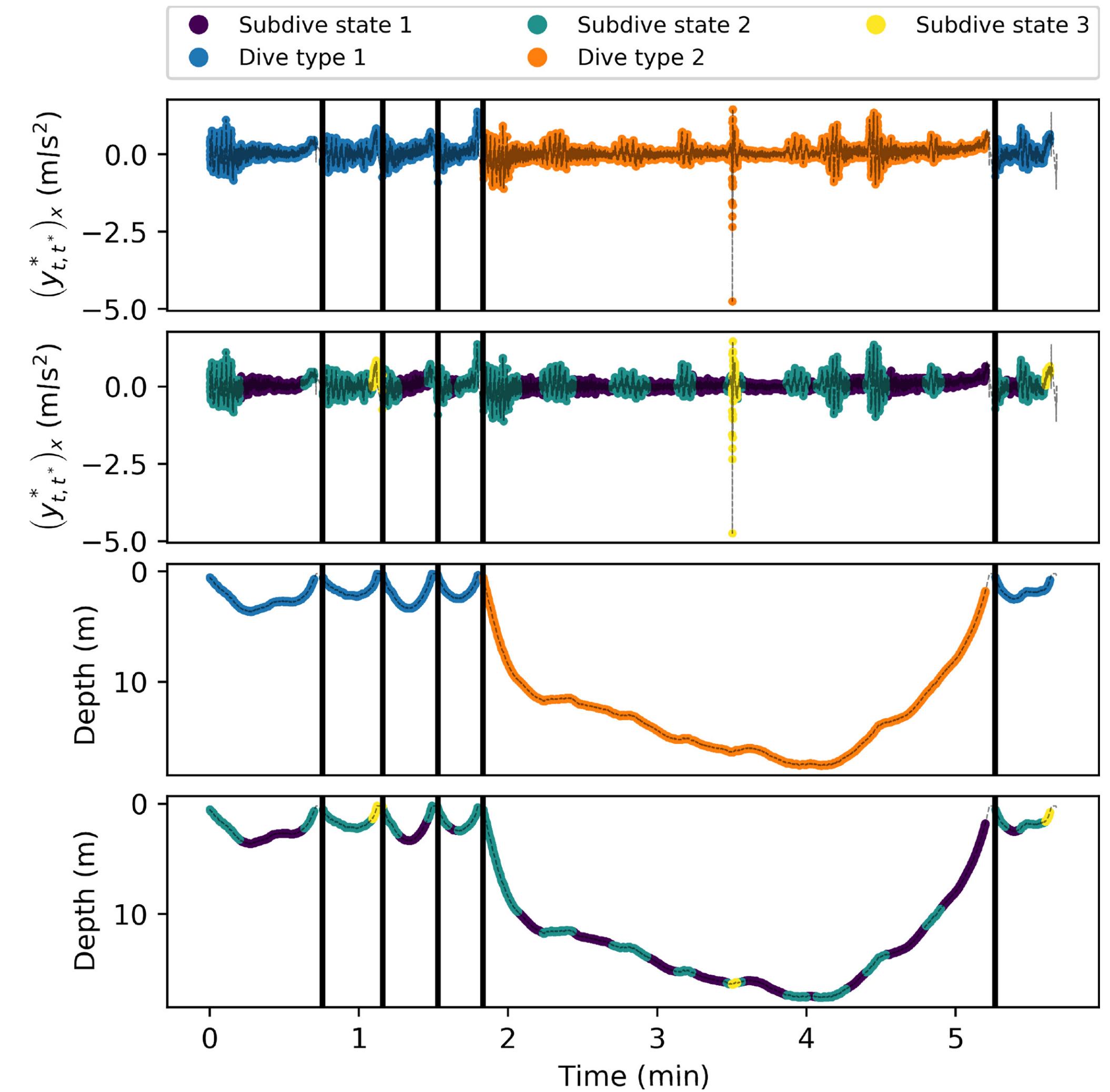


Coarse-scale



Glennie et al. 2022

Northern resident killer whale dive sequence

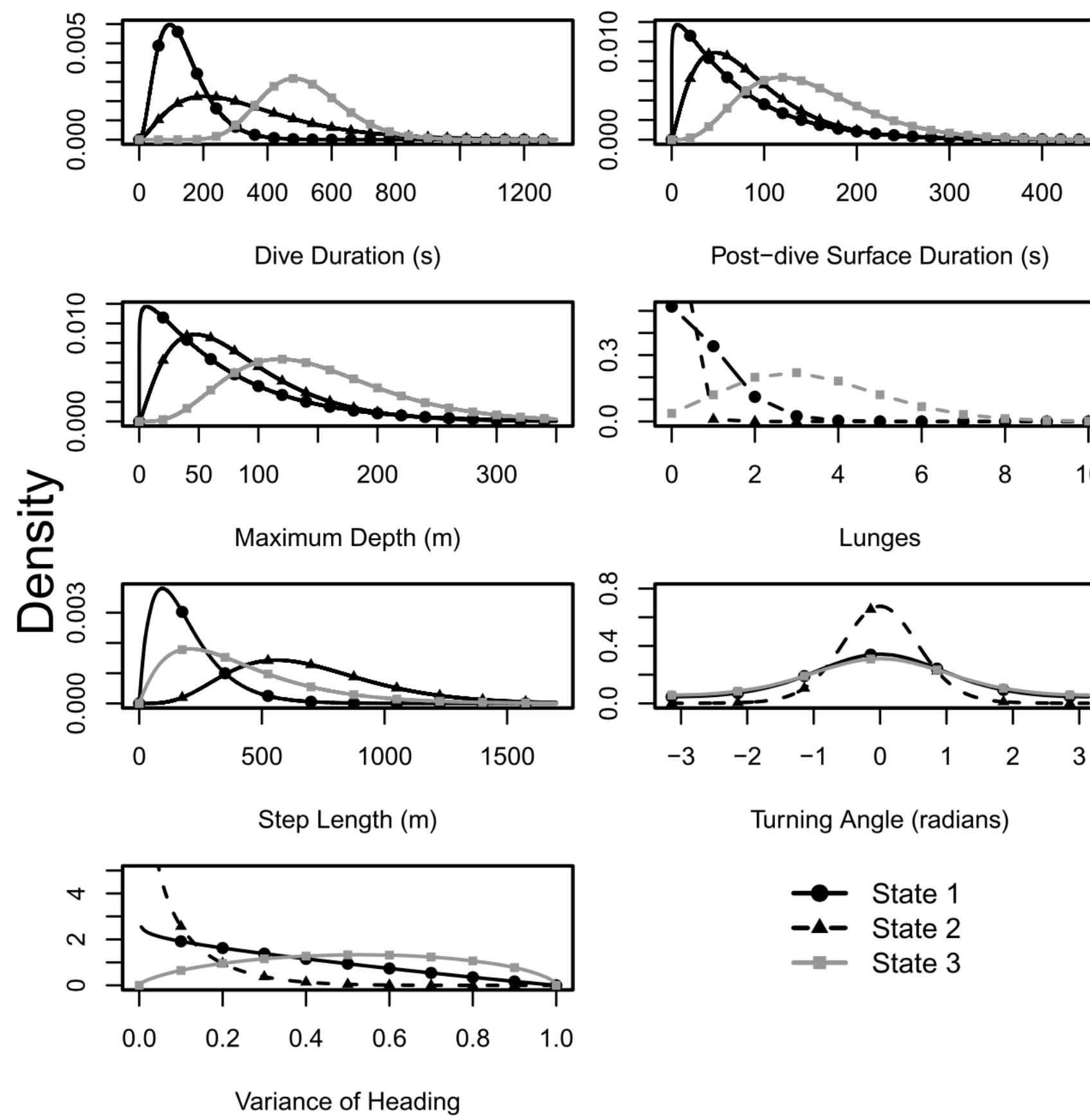


Sidrow et al. 2022

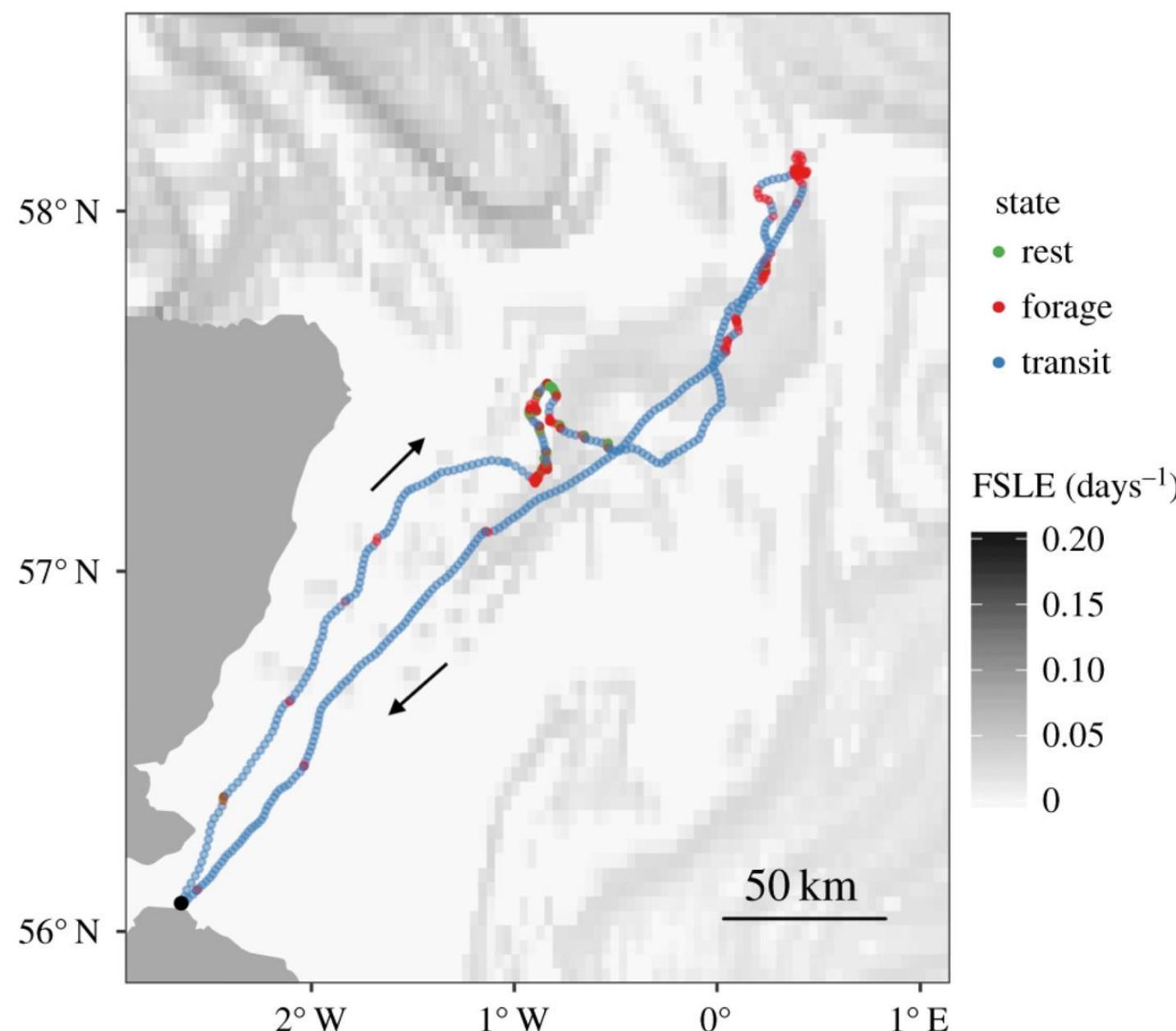
Methods to fit HMM

- Maximum likelihood estimation (MLE) is most common (for behavioral state estimation)
- Bayesian methods (via MCMC, HMC, etc) also possible, but less common
- Readily available functions to fit animal movement models using the moveHMM or momentuHMM R packages
 - hmmTMB is very new package that uses Laplace approximation via TMB package (similar to foieGras)
- If interested in comparison among available R packages, look at Table 2 of McClintock et al. (2020) for a comparison

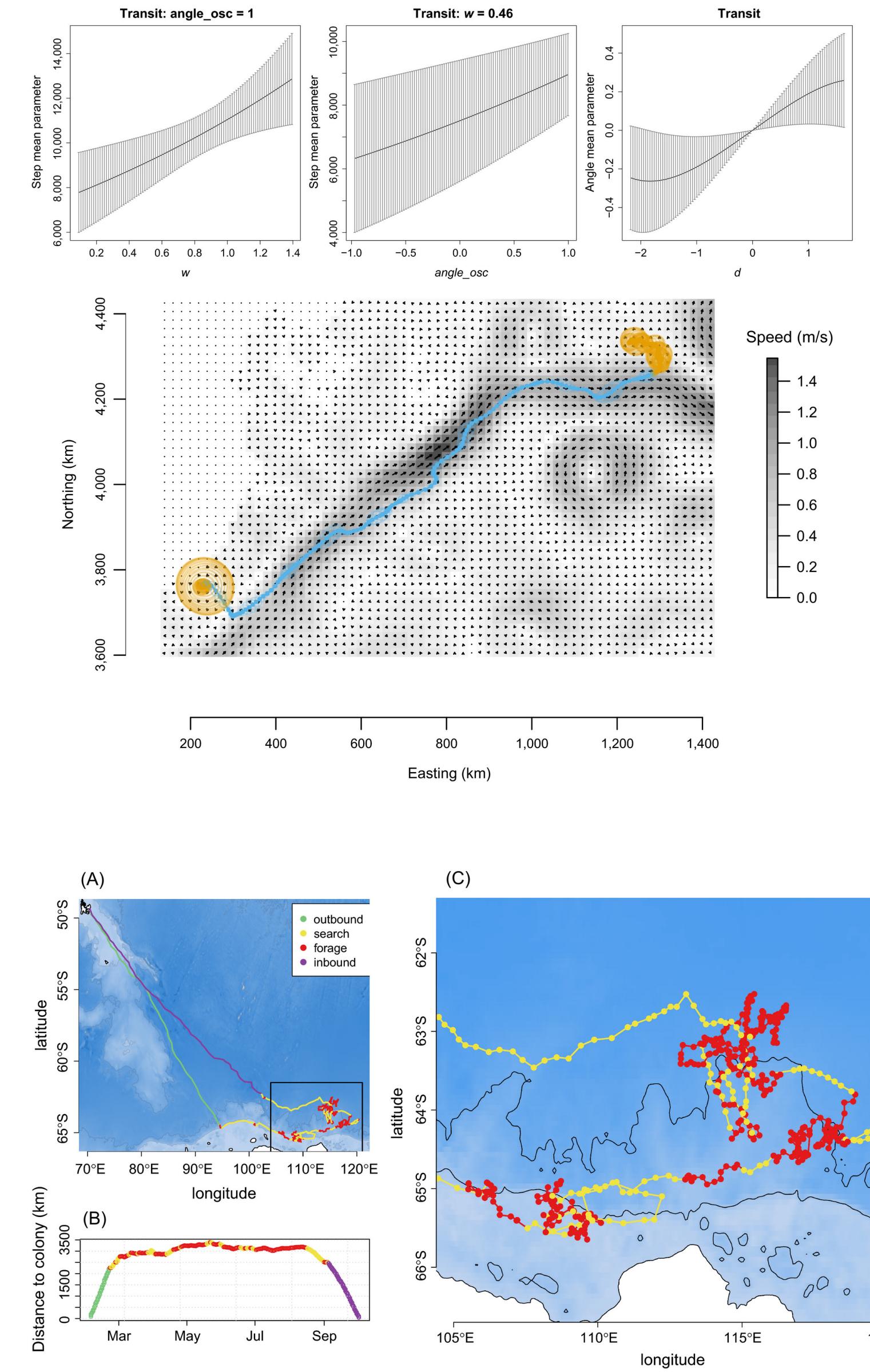
Motivating examples



Deruiter et al. 2017



Grecian et al. 2018



Michelot et al. 2017

Let's do some modeling!



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