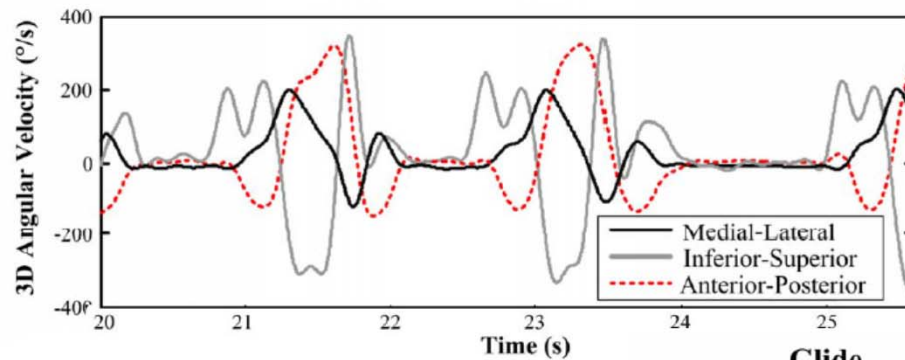


TIME SERIES DATA AND HIDDEN MARKOV MODELS

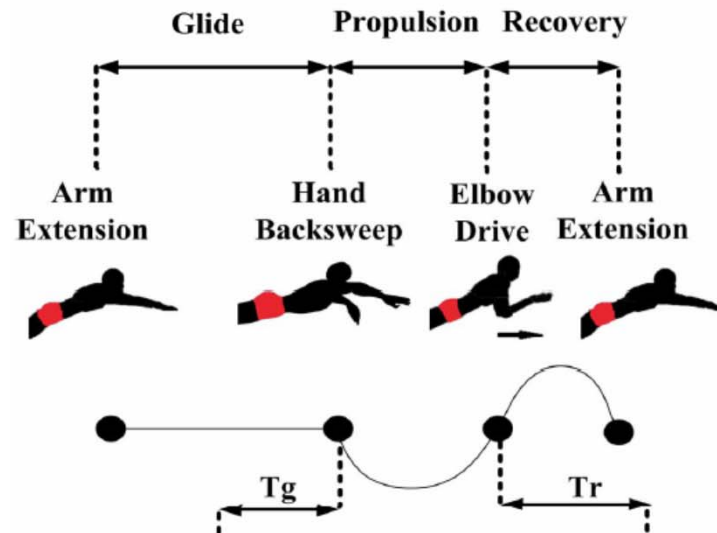
Paul Wu

Time Series Analysis: Breaststroke

2

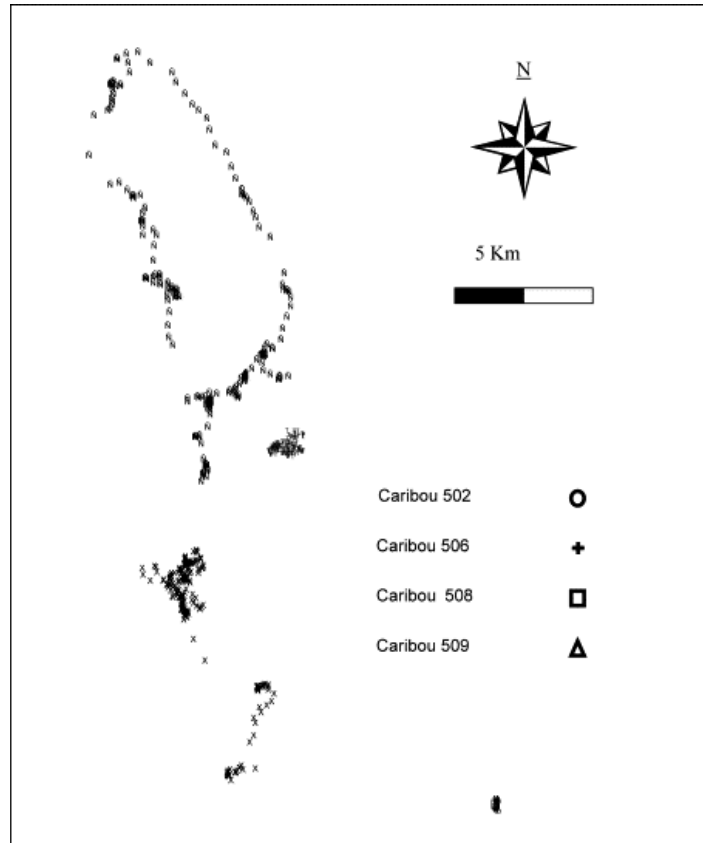


(Dadashi, 2013)



Spatio-Temporal Analysis

3



Observations: distance travelled and turn angle

3 behavioural states (bedding, foraging and relocating)

(Franke, 2004)

Hidden Markov Models (HMMs)

4

- What characterises these problems?
 - ▣ Multivariate time series
 - ▣ Data at each point in time is not independent
 - ▣ Data distribution may change over time
 - ▣ Data sequences not of the same length
- Why HMM?
 - ▣ Want to understand some process of a dynamical system
 - ▣ Unobserved/latent variables
 - ▣ Simple and computational efficiency



Where are HMMs used?

5

- Widely used (Zucchini, 2016):
 - ▣ Recognition (face, handwriting)
 - ▣ Bioinformatics
 - ▣ Environment (rainfall, earthquakes)
 - ▣ Finance
 - ▣ Ecology (animal movements)



Basics: Markov Chains

6

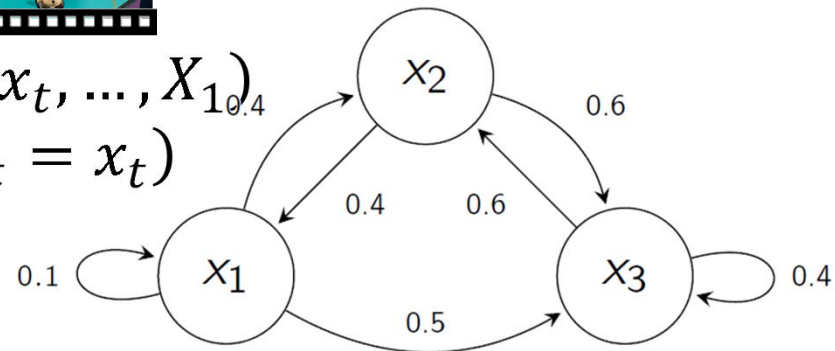
Markov chain = sequence of random variables

X_1, X_2, \dots, X_t where:



<https://fixhepc.com/support/faq/analysis/1250/what-s-simple.html>
<https://bopnetlimited.com/trapping-notes/>

$$P(X_{t+1} = x_{t+1} | X_t = x_t, \dots, X_1 = x_1) \\ = P(X_{t+1} = x_{t+1} | X_t = x_t)$$



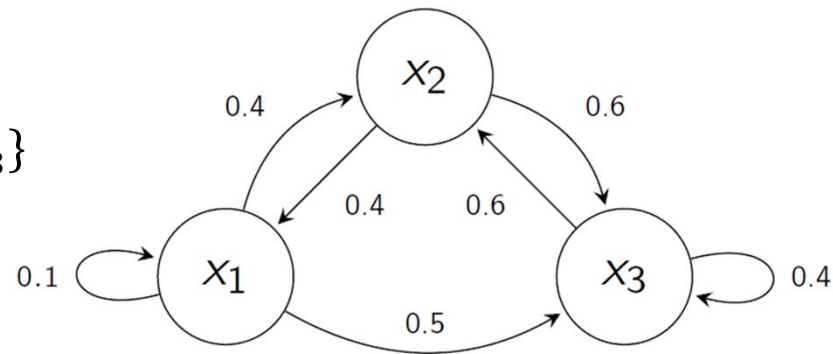
(Frazzoli, 2010)

Transition Matrix

7

$$T = P(x_j|x_i) = \begin{pmatrix} \text{from} \backslash \text{to} & x_1 & x_2 & x_3 \\ x_1 & 0.1 & 0.4 & 0.5 \\ x_2 & 0.4 & 0 & 0.6 \\ x_3 & 0 & 0.6 & 0.4 \end{pmatrix}$$

Finite set $X = \{x_1, x_2, x_3\}$



Transition Matrix

8

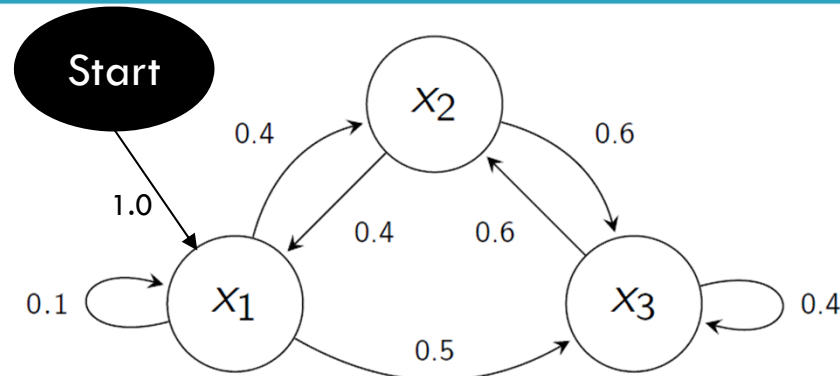
- If we know the state at time step 1 is $X_1 = x_1$, then $p_1 = \pi = (1,0,0)$.

- $p_2 = T'p_1 = (0.1,0.4,0.5) = (0.1 \times 1, 0.4 \times 1, 0.5 \times 1)$

- $p_3 = (T')^2 p_1 = (0.17,0.34,0.49) = (0.1 \times 0.1 + 0.4 \times 0.4, 0.1 \times 0.1 + 0.6 \times 0.5, 0.1 \times 0.5 + 0.6 \times 0.4 + 0.4 \times 0.5)$

- $p_t = (T')^t p_1 = (0.15,0.36,0.49)$

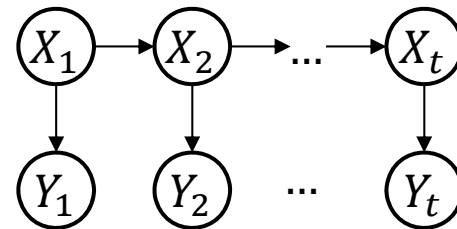
- For some systems, stationary distribution $p^\infty = T'p^\infty$



Hidden Markov Model

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- States X (finite set)
- Observations Y (finite set)
- T (transition probabilities)
- Q (emission=measurement probabilities)
- π (initial/prior state probability distribution)



Moles at Night: HMM

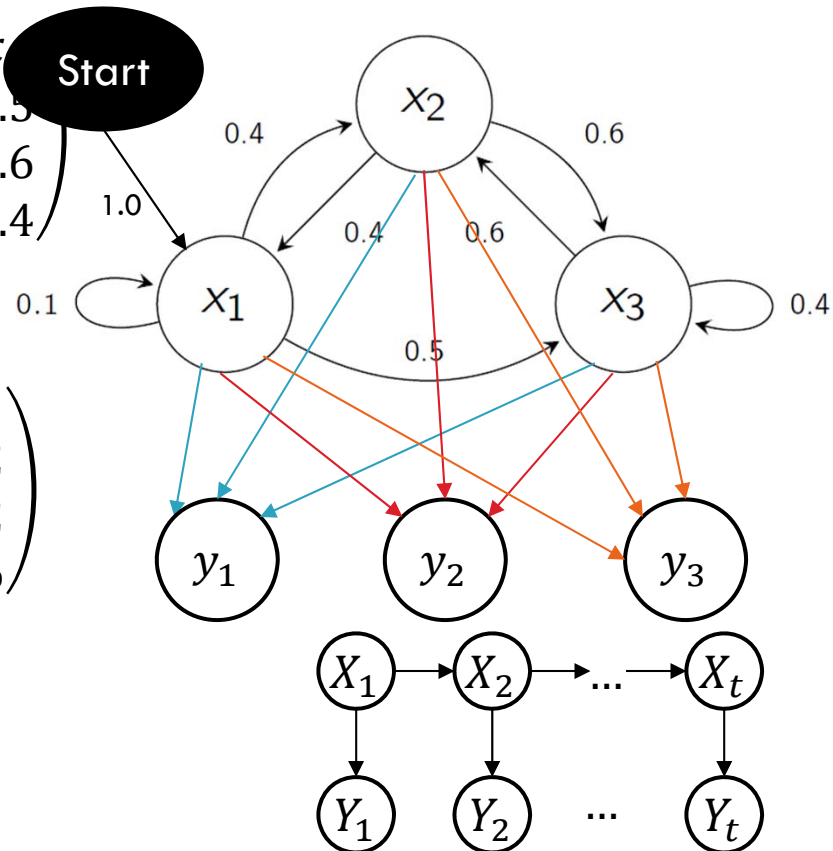
10

$$T = \begin{pmatrix} \text{from} \backslash \text{to} & x_1 & x_2 & x_3 \\ x_1 & 0.1 & 0.4 & 0.5 \\ x_2 & 0.4 & 0 & 0.6 \\ x_3 & 0 & 0.6 & 0.4 \end{pmatrix}$$

$$Q = P(y_j | x_i)$$

$$= \begin{pmatrix} \text{from} \backslash \text{to} & y_1 & y_2 & y_3 \\ x_1 & 0.6 & 0.2 & 0.2 \\ x_2 & 0.2 & 0.6 & 0.2 \\ x_3 & 0.2 & 0.2 & 0.6 \end{pmatrix}$$

$$p_1 = \pi = (1, 0, 0)$$



HMM: What Can It Do?

11

- Given observations y up to time t , we can do:
 - ▣ Filtering – what is the current state distribution $X_t = p_t$
 - ▣ Smoothing – hindsight, what is $X_u, u < t$
 - ▣ Prediction – forecast, what is $X_u, u > t$
 - ▣ Decoding – find the most likely state history $X_{1,\dots,t}$.



HMM Types

12

- Discrete time
 - ▣ Discrete observations (genetics: observe letters A, G, C, T)
 - ▣ Continuous observations - q_{x_i} has a distribution (e.g. Gaussian, Poisson)
 - ▣ Covariates – prior, transition
- Continuous time



Approach for Today

13

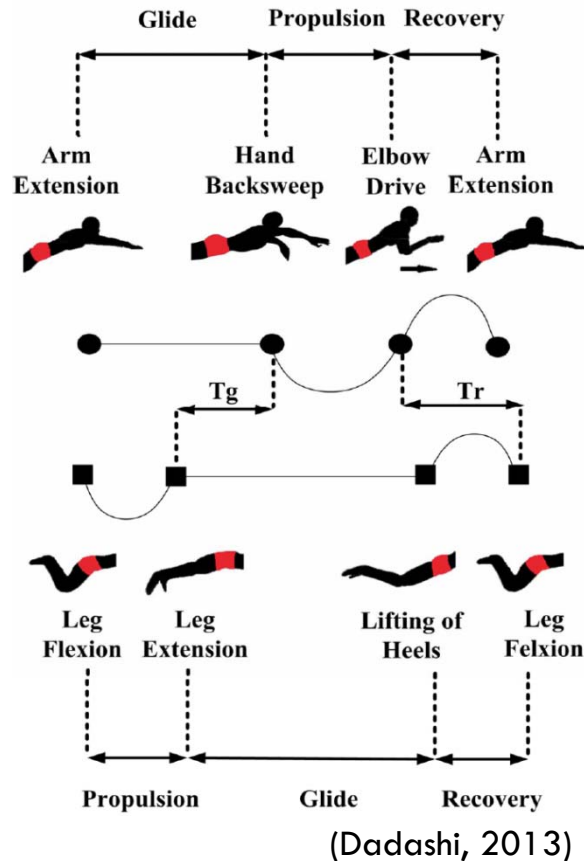
- Case study 1: swimming (Dadashi, 2013)
- Case study 2: animal movement
(Franke, 2004)
- Practical: response time (Dutilh, 2011)



Case Study 1: Swimming

14

- 7 swimmers, 2 100Hz IMUs (arm + leg)
- 3x200m trials with target speeds
- Temporal phases of locomotion – arm-leg motor organisation affects speed fluctuations and energy
- Total time gap = $T_g + T_r$



Case Study 1: Data

15

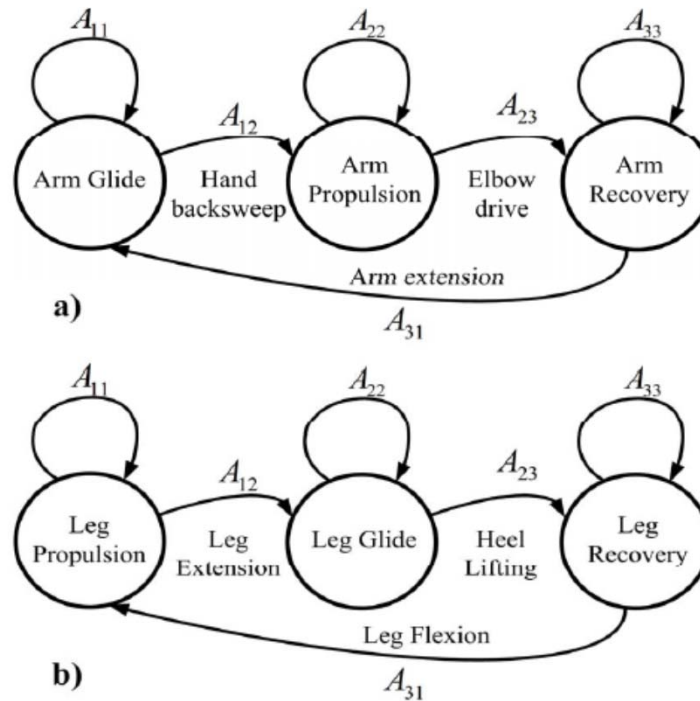
- Observations: IMU sensors – medial-lateral, inferior-superior, anterior-posterior angular velocities for arm and leg
- States and thus prior and transitions: expert annotated video



Case Study 1: Goal

16

Find T_g via the most probable sequence of states



(Dadashi, 2013)

Case Study 1: Estimating Parameters

17

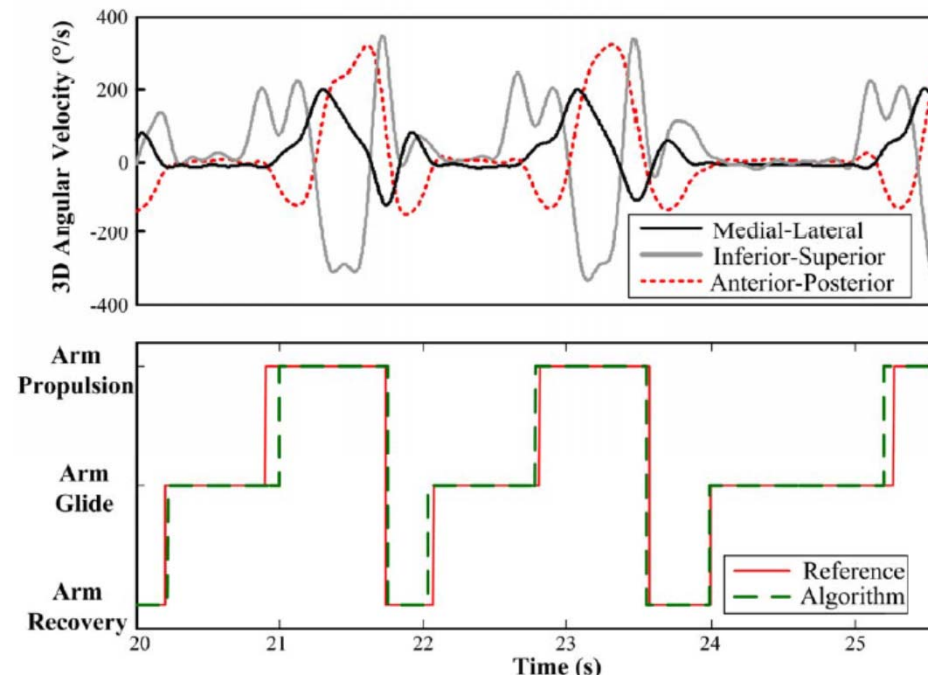
- π : Prior estimated as frequency of states in training set
- T : Transition probabilities estimated as $\frac{\text{\#samples in training set with transition from state } i \text{ to } j}{\text{total \#samples labelled } j}$.
- Q : Emissions modelled as multi-variate Gaussian mixtures.
 - ▣ If Y is discrete, q is estimated $\frac{\text{\#outputs } i \text{ from state } j}{\text{total \#samples of state } j}$.
 - ▣ mclust, mixtools



Supervised learning

Results

18



(Dadashi, 2013)

Results and Learnings

19

- Error in estimating the state/phase:
 - ▣ Sensitivity 93.5% for arm, 94.4% for leg; specificity of 96.2% and 97.2%, respectively
 - ▣ Boosted model – use output of classifiers (linear + non-parametric estimator, Gaussian kernel) as input to HMM
- Error in estimating TG: $-11 \pm 52\text{ms}$



Case Study 2: Caribou Movements

20

- Developing a set of interpretable states for caribou behaviours
 - ▣ State and transitions ('strategy') not known/defined in advance!
 - ▣ Postulate 3 hidden states – bedding, feeding and relocating
- 12 caribou over 10 days with GPS tracking
- Goal: interpret duration of behaviour and transition to different behaviours



(Franke, 2004)

Case Study 2: Data

21

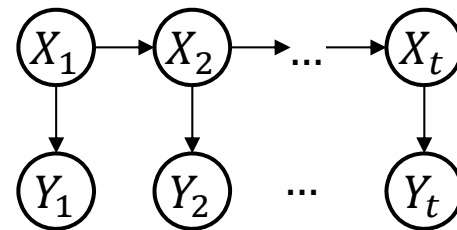
- Distance moved and turn angle
 - ▣ Distance discretised into zero $< 20\text{m}$, short 20-100m, medium 100-250m, long $> 250\text{m}$
 - ▣ Angles discretised: ahead $316-45^\circ$, right $46-135^\circ$, back $136-225^\circ$, left $226-315^\circ$.



Case Study 2: Estimating Parameters

22

- Expectation maximisation (EM)
 - ▣ “Missing data” are the states
 - ▣ E: estimate probability of the state sequence given observations
 - ▣ M: fit new model parameters
 - ▣ R – depmixS4



Case Study 2: Results

Table 4
Multiple-observation HMM for caribou 506

	State 1 _(t+1) (<i>B</i>)	State 2 _(t+1) (<i>F</i>)	State 3 _(t+1) (<i>R</i>)
A: State transitions			
State 1 _(t) (<i>B</i>)	0.87	0.037	0.09
State 2 _(t) (<i>F</i>)	0.90	0.04	0.06
State 3 _(t) (<i>R</i>)	0.14	0.02	0.84

Table 5
Multiple-observation HMM for caribou 508

	State 1 _(t+1) (<i>B</i>)	State 2 _(t+1) (<i>F</i>)	State 3 _(t+1) (<i>R</i>)
A: State transitions			
State 1 _(t) (<i>B</i>)	0.68	0.29	0.02
State 2 _(t) (<i>F</i>)	0.69	0.28	0.03
State 3 _(t) (<i>R</i>)	0.21	0.10	0.69



(Franke, 2004)

Case Study 2: Results

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Table 5
Multiple-observation HMM for caribou 508

	State 1 _(t+1) (<i>B</i>)	State 2 _(t+1) (<i>F</i>)	State 3 _(t+1) (<i>R</i>)		
A: State transitions					
State 1 _(t) (<i>B</i>)	0.68	0.29	0.02		
State 2 _(t) (<i>F</i>)	0.69	0.28	0.03		
State 3 _(t) (<i>R</i>)	0.21	0.10	0.69		
	Stationary	Short	Medium	Long	
B1: Distance between locations					
State 1 _(t) (<i>B</i>)	0.99	0.01	0.00	0.00	
State 2 _(t) (<i>F</i>)	0.91	0.09	0.00	0.00	
State 3 _(t) (<i>R</i>)	0.00	0.85	0.11	0.04	
	Stationary	Ahead	Right	Left	Backward
B2: Turn angle					
State 1 _(t) (<i>B</i>)	1.00	0.00	0.00	0.00	0.00
State 2 _(t) (<i>F</i>)	1.00	0.00	0.00	0.00	0.00
State 3 _(t) (<i>R</i>)	0.00	0.50	0.11	0.17	0.22



Results and Learnings

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- Relating behaviour to habitat and conservation
- Challenges
 - ▣ Potential for insertion and deletion of states
 - ▣ Initialisation and convergence to local maximum
 - ▣ Constraints: challenging for EM and result in incorrect parameter estimates
 - depmixs4: direct optimisation



Results and Learnings

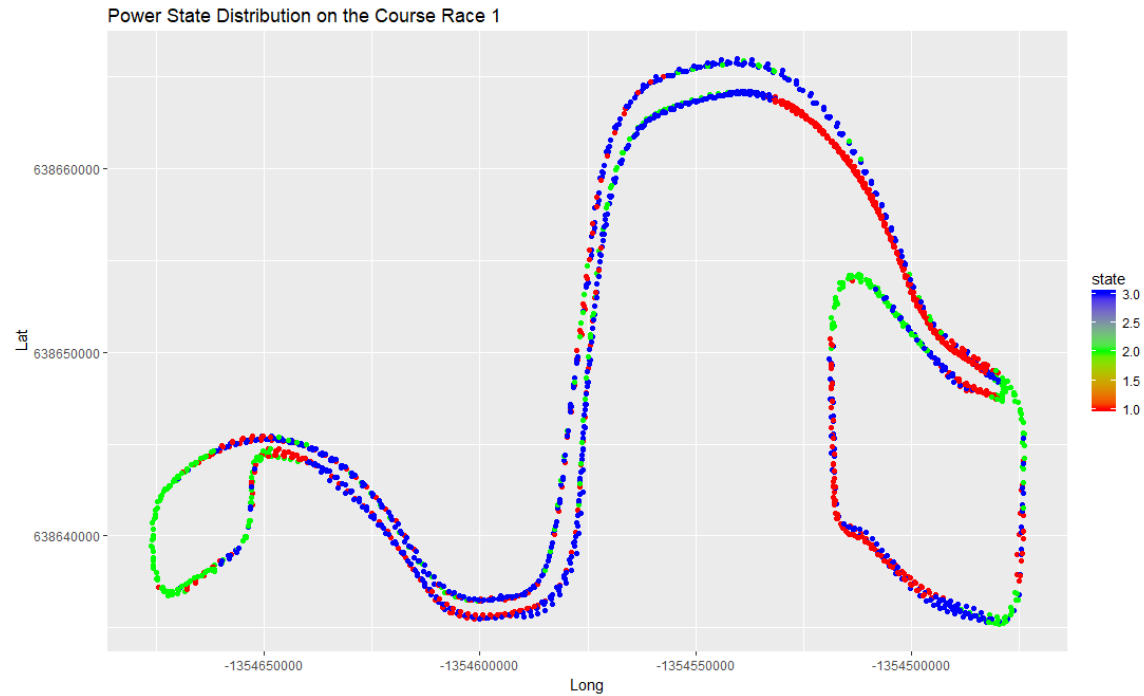
26

- Covariates – affect prior or transition
 - ▣ Time of day
 - ▣ Season
 - ▣ Spatial – terrain and vegetation



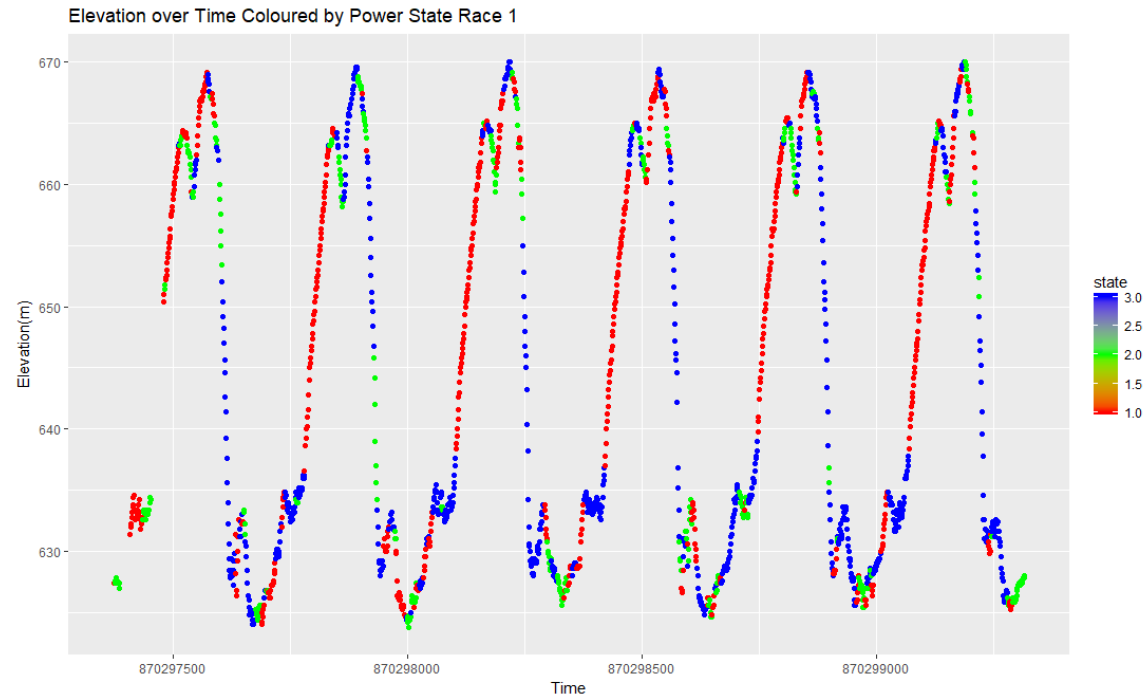
QAS Example of HMM

27



QAS Example for HMM

28



Reflection

29

- How might HMMs be used for your work?



Prac: Response Time

30

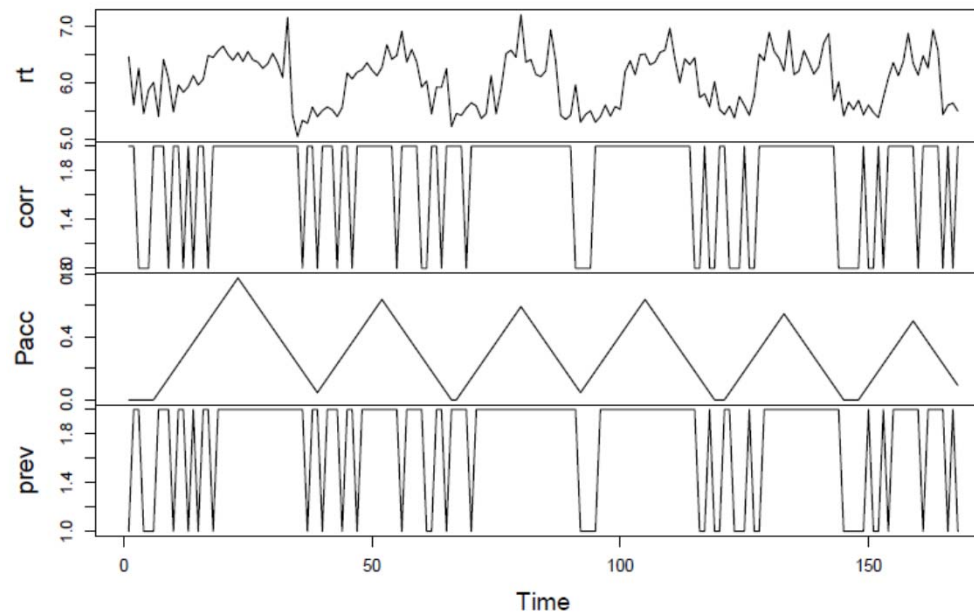
Dutilh, Gilles, et al. "A phase transition model for the speed-accuracy trade-off in response time experiments." *Cognitive Science* 35.2 (2011): 211-250.



Prac: Response Time

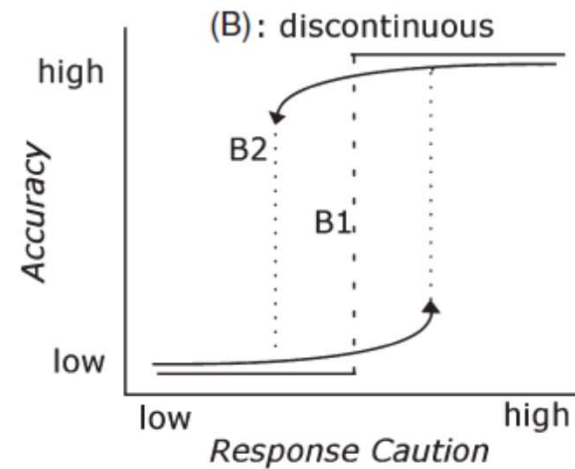
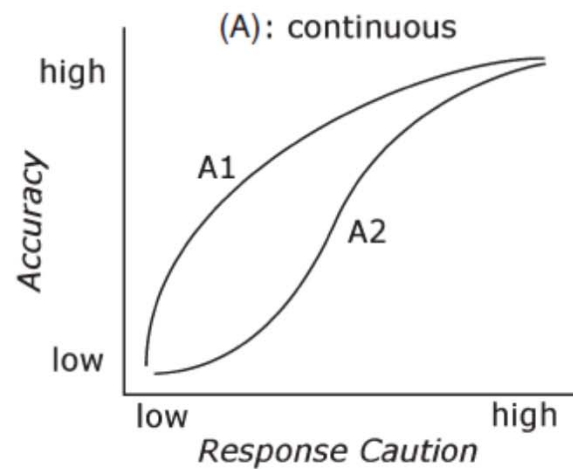
31

Speed-accuracy trade-off



Prac: Response Time

32



Prac: Load Data

33

- ☐ # load the library and data
- ☐ `install.packages('depmixS4')`
- ☐ `library(depmixS4)`
- ☐ `data(speed)`
- ☐ # View the data
- ☐ `View(speed)`



Prac: Visualise the Data

34

```
library(ggplot2)
library(reshape2)
speed = cbind(speed,list(t=1:nrow(speed)))
mspeed = melt(speed,measure.vars=c('rt','corr','Pacc'))
ggplot(mspeed,aes(x=t,y=value,group=variable)) +
  geom_line() +
  facet_wrap(~variable,scales='free',ncol=1)
```



Prac: Fit a HMM

35

```
> modsimple =  
depmix(response=rt~1,data=speed,nstates=2,trstart=runif(4))  
  
> fmsimple = fit(modsimple)  
  
iteration 0 logLik: -305.3318  
iteration 5 logLik: -305.3198  
iteration 10 logLik: -305.3066  
iteration 15 logLik: -305.2831  
iteration 40 logLik: -88.92333  
iteration 45 logLik: -88.71571  
iteration 50 logLik: -88.71502  
converged at iteration 52 with logLik: -88.71502
```



Prac: Fit a HMM

36

```
> summary(fmsimple)
```

Initial state probabilities model

pr1 pr2

1 0

Transition matrix

toS1 toS2

fromS1 0.916 0.084

fromS2 0.117 0.883

Response parameters

Resp 1 : gaussian

Re1.(Intercept) Re1.sd

St1 6.385 0.244

St2 5.510 0.192



Prac: Results

37

```
> fmsimple
```

Convergence info: Log
likelihood converged to within
tol. (relative change)

'log Lik.' -88.71502 (df=7)

AIC: 191.43

BIC: 220.0215

```
> head(fmsimple@posterior)
```

	state	S1	S2
1	1	1.000000000	0.000000000
2	2	0.052835996	0.94716400
3	1	0.993992010	0.00600799
4	2	0.005894012	0.99410599
5	2	0.063159278	0.93684072
6	2	0.457223228	0.54277677



Filtering, smoothing, prediction, forecasting

Prac: Questions

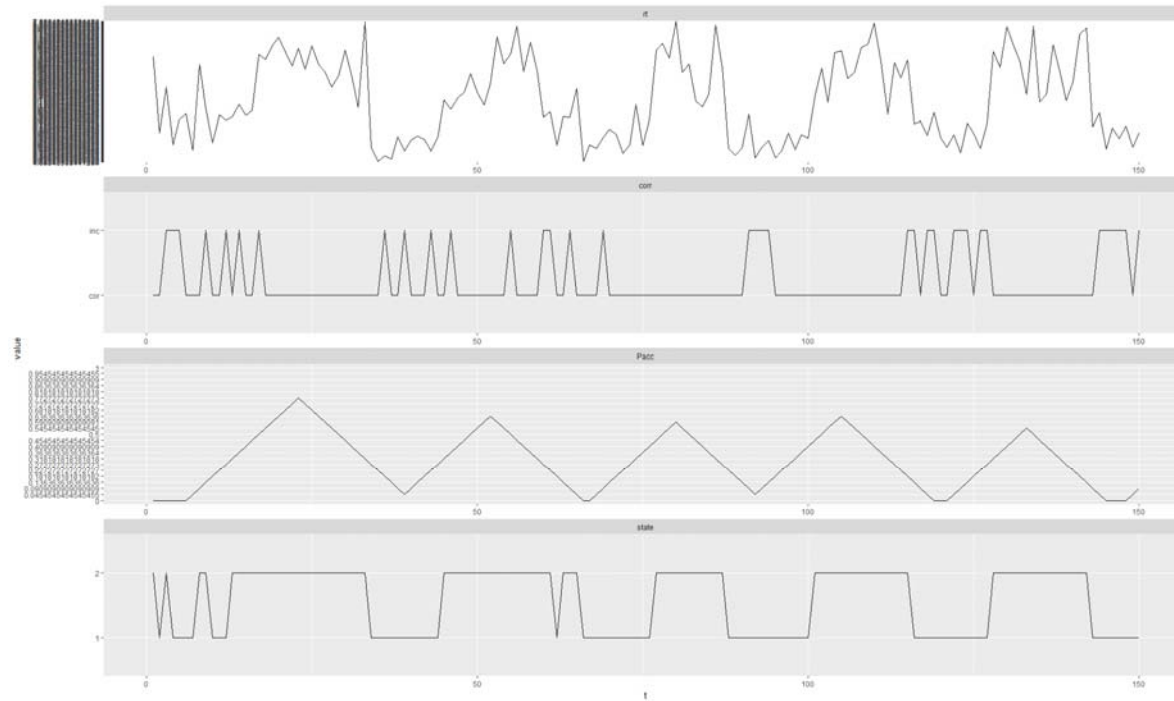
38

- Q1: Explain your results
- Q2: What did we learn from this?



Prac: Questions

39



Extension: Fit with Covariates

40

```
modtrans =  
depmix(response=rt~1,transition=~Pacc,data=speed,nsta  
tes=2,instart=runif(2))  
fmtrans = fit(modtrans)
```

- ☐ Explain your results?
- ☐ What is different?
- ☐ What additional information can we learn?



Extension : Fit with Covariates

41

```
> summary(fmtrans)
```

Initial state probabilities model

```
pr1 pr2
```

```
0 1
```

Transition model for state (component) 1

Model of type multinomial (mlogit), formula:

```
~Pacc
```

Coefficients:

```
St1 St2
```

```
(Intercept) 0 -3.128895
```

```
Pacc 0 5.971186
```

Probabilities at zero values of the covariates.

```
0.958069 0.04193098
```

Transition model for state (component) 2

Model of type multinomial (mlogit), formula:

```
~Pacc
```

Coefficients:

```
St1 St2
```

```
(Intercept) 0 -3.126696
```

```
Pacc 0 15.354371
```

Probabilities at zero values of the covariates.

```
0.9579806 0.04201939
```

Response parameters

Resp 1 : gaussian

```
Re1.(Intercept) Re1.sd
```

```
St1 5.507 0.187
```

```
St2 6.386 0.242
```



Extension : Fit with Covariates

42

```
> fmtrans
```

Convergence info: Log
likelihood converged to within
tol. (relative change)

'log Lik.' -44.19949 (df=9)

AIC: 106.399

BIC: 143.1595



Extension2: Multivariate HMM

43

```
modmulti = depmix(list(rt~1,corr~1),data=speed,nstates=2,  
family=list(gaussian(),multinomial('identity')),transition=~scale(Pa  
cc), instart=runif(2))
```

```
fmmulti = fit(modmulti)
```



Extension2: Multivariate HMM

44

Initial state probabilities model

pr1 pr2

0 1

Transition model for state (component) 1

Model of type multinomial (mlogit), formula: $\sim \text{scale}(\text{Pacc})$

Coefficients:

St1 St2

(Intercept) 0 -0.9265658

scale(Pacc) 0 1.5984641

Probabilities at zero values of the covariates.

0.716378 0.283622

Transition model for state (component) 2

Model of type multinomial (mlogit), formula:

$\sim \text{scale}(\text{Pacc})$

Coefficients:

St1 St2

(Intercept) 0 2.401846

scale(Pacc) 0 3.722318

Probabilities at zero values of the covariates.

0.08303203 0.916968

Response parameters

Resp 1 : gaussian

Resp 2 : multinomial

Re1.(Intercept) Re1.sd Re2.inc Re2.cor

St1 5.517 0.197 0.475 **0.525**

St2 6.391 0.239 0.098 **0.902**



Extension2: Multivariate HMM

45

Convergence info: Log
likelihood converged to within
tol. (relative change)

'log Lik.' -255.5337 (df=11)

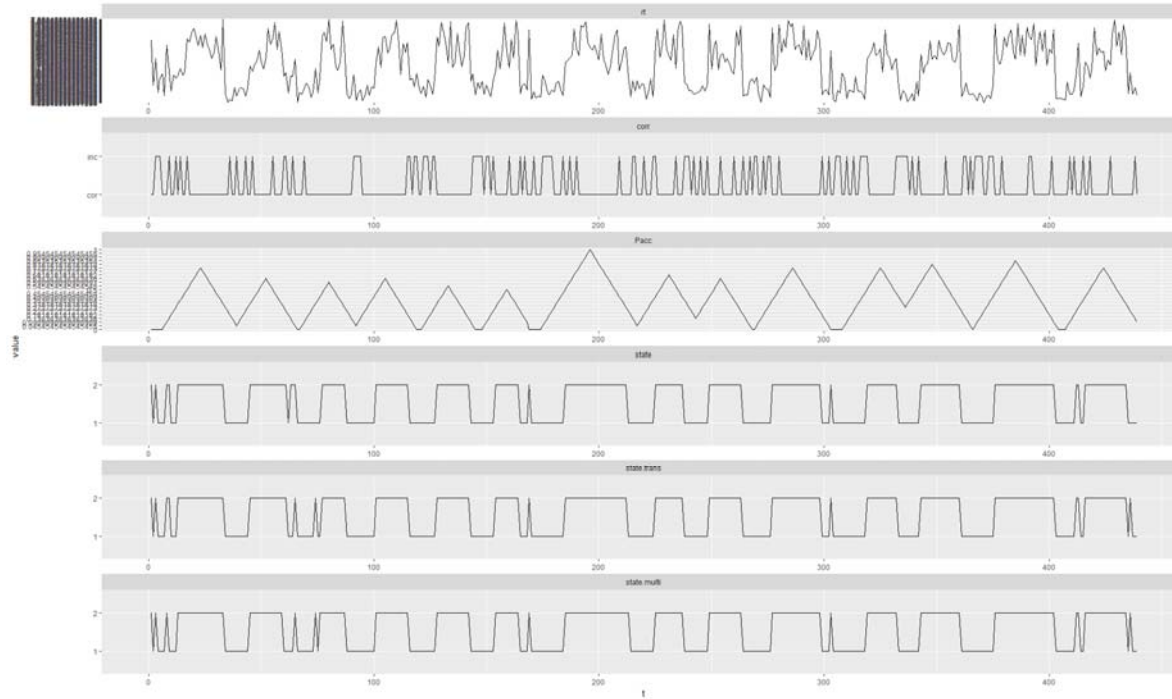
AIC: 533.0674

BIC: 577.9969



Extension2: Multivariate HMM

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Questions

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Paul Wu

No moles were harmed in the making of this presentation



References

48

- Dadashi, F., A. Arami, F. Crettenand, G. P. Millet, J. Komar, L. Seifert and K. Aminian (2013). A Hidden Markov Model of the breaststroke swimming temporal phases using wearable inertial measurement units. 2013 IEEE International Conference on Body Sensor Networks.
- Franke, Alastair, Terry Caelli, and Robert J. Hudson. "Analysis of movements and behavior of caribou (*Rangifer tarandus*) using hidden Markov models." *Ecological Modelling* 173.2-3 (2004): 259-270.



References

49

- Zucchini, Walter, Iain L. MacDonald, and Roland Langrock. *Hidden Markov models for time series: an introduction using R*. CRC press, 2016.
- Roohi, N. “HMM: Viterbi Algorithm – a toy example”, University of Pennsylvania. <http://www.cis.upenn.edu/~cis262/notes/Example-Viterbi-DNA.pdf>.
- Rabusseau, G. and Islam, R. “Lecture 9: Hidden Markov Models.” McGill University. <https://rllabmcgill.github.io/COMP-652/lectures.html>.
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- Langrock, Roland, et al. "Flexible and practical modeling of animal telemetry data: hidden Markov models and extensions." *Ecology* 93.11 (2012): 2336-2342.





Prac Case Study: Cycling Power

52

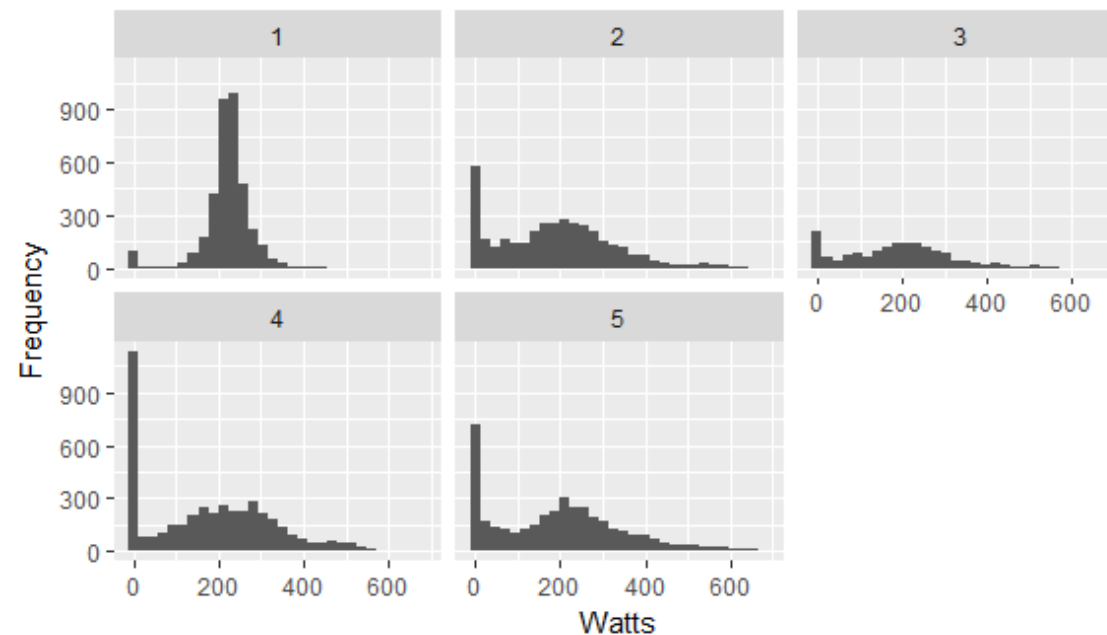
- Goal: understand power variability and physiological states in cycling
- Data:
 - ▣ GPS: latitude, longitude, elevation
 - ▣ Power
 - ▣ Speed
 - ▣ Cadence



Prac Case Study: Power

53

Power Histogram for Each Race



Prac Case Study: Model

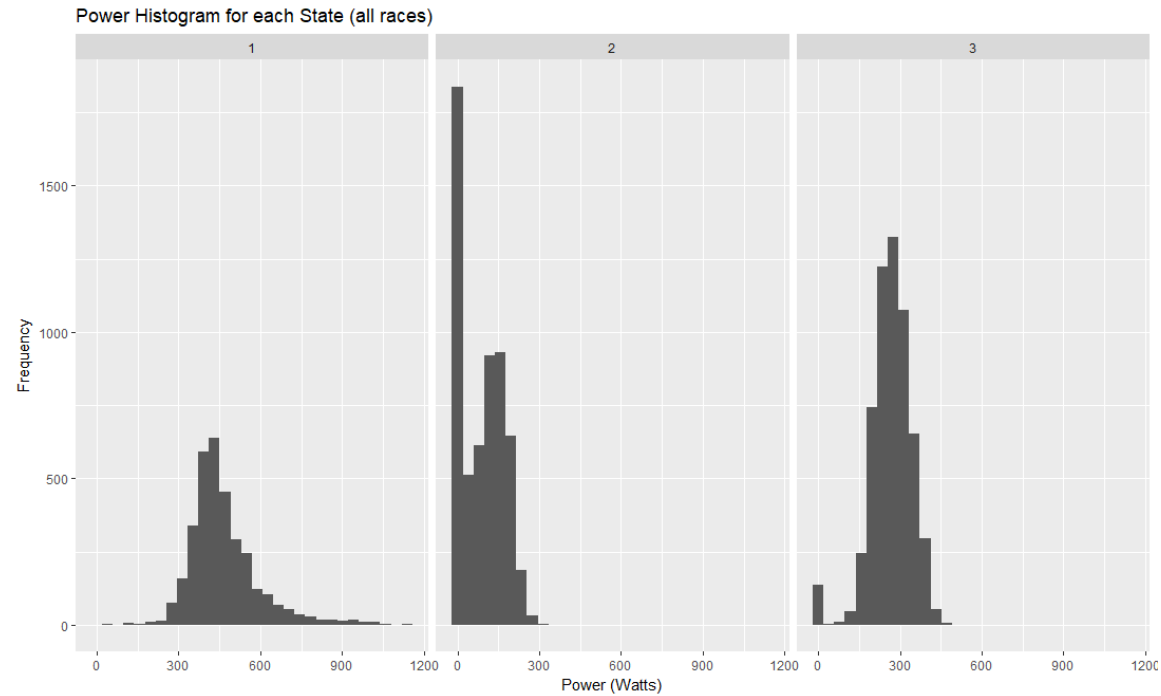
54

```
mod <- depmix(response = Power ~ 1 + Speed  
+ Cadence + Elevation, data = R, nstates =  
3)  
fm <- fit(mod)
```



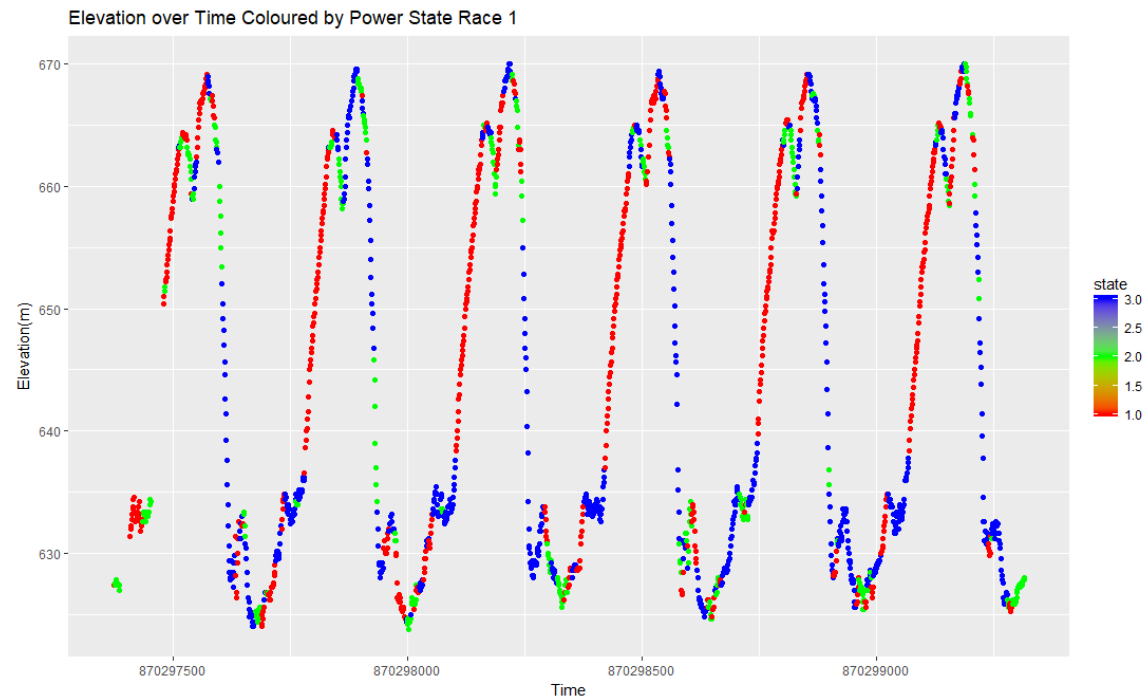
Prac Case Study: Results

55



Prac Case Study: Results

56



Power State Distribution on the Course Race 1

