CSC 535 HW9

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1 Introduction

All homework problems were completed. I was a little stumped on what an expected value of 24 cycles before a transition implied for the values within the transition matrix, for the cell problem. I settled on something that intuitively made sense to me, but it might not be the correct answer. Otherwise, I felt that 2.d and 2.e were very calculation heavy without much payoff in understanding. It was basically finding out how many of each probability to multiply, then multiply them all. Also, I suspect there's a high chance of human error in those problems, whereas those problems are much better suited to having a computer perform the computations.

2 A Graphical Exploration of Hidden Markov Models

2.1 A State Transition Diagram Representing Progression Through College

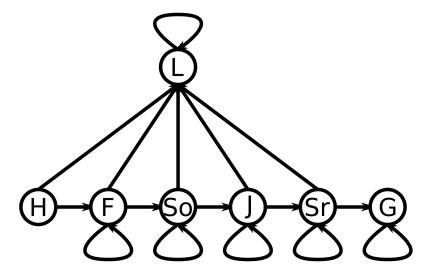


Figure 1: A state transition diagram representing progression through college. Initially, a student is in high school (H) and chooses whether to leave education (L) or continue to college (F). For each year in college, freshman (F), sophmore (So), junior (J), and senior (Sr), the student either passes his or her classes, and thus progresses to the next year, or fails some or all of his or her classes. If he or she fails classes, he or she must decide whether to continue his or her education (thus remaining in the current year) or leave academia altogether (L). If a student passes his or her senior (Sr) year, he or she graduates (G) and remains a college graduate.

2.2 A Hidden Markov Model for a Cell

This scenario is a little interesting, because it concerns taking measurements at a different time interval than the actual state transitions. In addition, the measurements are for gene expressions rather than the underlying states. We could construct the model in terms of our measurement times (one hour intervals) or in terms of the actual state transition period (24 hours). Doing things in terms of measurements times gives us finer control over the state transition as well as provides a one set of outputs (gene expressions) per cycle. Since we want the expected value of the change to be after 24 cycles, we can cook up an equation to represent the probability that the organism has not changed state after 24 hours:

$$x^{24} = 0.5$$

 $x \approx 0.97153$

I chose 0.5 here because it seemed like the most reasonable mid-point between no chance and 100% chance. I cannot explain it, but, intuitively, this seems like it would yield the correct expected value of 24 cycles. If it does not, I would love to hear in the feedback what the correct approach is. The transition matrix then becomes:

$$T = \begin{bmatrix} 0.97153 & 0.02847 \\ 0.02847 & 0.97153 \end{bmatrix}$$

The graph for the Hidden Markov Model is below. I list four observations for each cycle, representing the expression probability of each gene. I have come up with some arbitrary probabilities for those gene expressions.

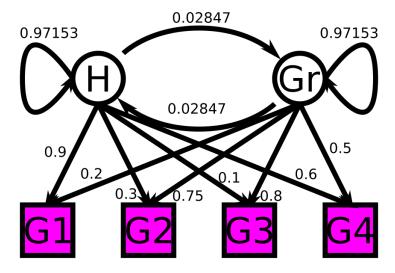


Figure 2: A graph for a HMM representing two cell states, hibernation and growth, with a state transition time of one hour. For each of those hours, a set of four genes are tested for whether they were expressed or not. The probabilities of those genes being expressed is given as the numbers next to the corresponding links.

3 Reasoning About Trails

3.1 Trail Transition Matrix

This is pretty straightforward. Each entry for an adjacent road is $\frac{1}{n}$, where n is the number of adjacent roads. Let the rows represent the current hiker's location, sorted lexicographically, and let the columns represent the probabilities for the hiker's next location, sorted lexicographically. (The value for A is in entry 1, B in 2, and so on.)

$$T = \begin{bmatrix} 0 & 1 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0.2 & 0 & 0.2 & 0.2 & 0.2 & 0 & 0.2 & 0 \\ 0 & 0.333 & 0 & 0.333 & 0 & 0 & 0.333 & 0 \\ 0 & 0.25 & 0.25 & 0 & 0 & 0.25 & 0.25 & 0 \\ 0 & 0.333 & 0 & 0 & 0 & 0.333 & 0 & 0.333 \\ 0 & 0 & 0 & 0.25 & 0.25 & 0 & 0.25 & 0.25 \\ 0 & 0.25 & 0.25 & 0.25 & 0 & 0.25 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0.5 & 0.5 & 0 & 0 \end{bmatrix}$$

3.2 Is this a Markov Process?

Markov Processes involve several key features. One is the change of state over time over a fininte number of states. States are represented by each fork in the trail. In addition, the transitions from a given state are a probability distribution. Thus, the rows in the transition matrix must sum to one. This is satisfied above. Markovian processes also assume that the only state that matters for determining the current state is the previous state. This is satisfied because the path transition probabilities are given only in terms of the current fork. One final observation is that this is a *Hidden* Markov Model, since we can only reason about the state of the traveler based on indirect observations, in this case, the sensors, which are not perfect.

3.3 Joint Probability of Observation Sequences

First, define two observation output matrices as follows, where each row is a hidden state (current trail location), and each column is probability of the sensor output being n at each location, given the hidden state. This is meant as an easy way to address a sensor output for a given hidden state.

$$O_0 = \begin{bmatrix} 0.15 & 0.75 & 0.75 & 0.75 & 0.75 & 0.75 & 0.75 \\ 0.75 & 0.15 & 0.75 & 0.75 & 0.75 & 0.75 & 0.75 & 0.75 \\ 0.75 & 0.75 & 0.15 & 0.75 & 0.75 & 0.75 & 0.75 & 0.75 \\ 0.75 & 0.75 & 0.15 & 0.75 & 0.75 & 0.75 & 0.75 & 0.75 \\ 0.75 & 0.75 & 0.75 & 0.15 & 0.75 & 0.75 & 0.75 & 0.75 \\ 0.75 & 0.75 & 0.75 & 0.75 & 0.15 & 0.75 & 0.75 & 0.75 \\ 0.75 & 0.75 & 0.75 & 0.75 & 0.15 & 0.75 & 0.75 & 0.75 \\ 0.75 & 0.75 & 0.75 & 0.75 & 0.75 & 0.15 & 0.75 & 0.75 \\ 0.75 & 0.75 & 0.75 & 0.75 & 0.75 & 0.75 & 0.15 & 0.75 \\ 0.75 & 0.75 & 0.75 & 0.75 & 0.75 & 0.75 & 0.15 & 0.75 \\ 0.25 & 0.25 & 0.25 & 0.25 & 0.25 & 0.25 & 0.25 \\ 0.25 & 0.25 &$$

You could also think of this as a tensor where $O_{i,j,k}$ represents the probability of the sensor output at fork j being k, given the hidden state of i. As we are interested in a particular sequence output, I generated a sequence of state changes and observations by rolling some dice. We could write the probability of that sequence like this:

$$\pi_1O_{1,1,1}O_{1,2,0}O_{1,3,1}O_{1,4,1}O_{1,5,0}O_{1,6,1}O_{1,7,0}O_{1,8,0} \\ T_{1,2}O_{2,1,0}O_{2,2,1}O_{2,3,0}O_{2,4,1}O_{2,5,0}O_{2,6,0}O_{2,7,0}O_{2,8,1} \\ T_{2,4}O_{4,1,0}O_{4,2,0}O_{4,3,1}O_{4,4,1}O_{4,5,0}O_{4,6,1}O_{4,7,0}O_{4,8,0}$$

And so on, and so forth. Note that, in this case, $pi_1 = 1$, since the hiker always starts at location A.

3.4 Sensor Response Sequence Probabilities

Given $R_1 = [a, b, c, d, b, c, d, b, e, f, h]$, what are the probabilities of the following sensor responses? For the following calcuations, I will make use of the tensor O defined above.

3.4.1 $[\{a\}, \{b\}, \{c\}, \{d\}, \{b\}, \{c\}, \{d\}, \{b\}, \{e\}, \{f\}, \{h\}]]$

$$P([\{a\},\{b\},\{c\},\{d\},\{b\},\{c\},\{d\},\{b\},\{c\},\{d\},\{b\},\{e\},\{f\},\{h\}])) = \\ O_{1,1,1}O_{1,2,0}O_{1,3,0}O_{1,4,0}O_{1,5,0}O_{1,6,0}O_{1,7,0}O_{1,8,0} \\ O_{1,1,0}O_{1,2,1}O_{1,3,0}O_{1,4,0}O_{1,5,0}O_{1,6,0}O_{1,7,0}O_{1,8,0} \\ O_{1,1,0}O_{1,2,0}O_{1,3,1}O_{1,4,0}O_{1,5,0}O_{1,6,0}O_{1,7,0}O_{1,8,0} \\ O_{1,1,0}O_{1,2,0}O_{1,3,0}O_{1,4,1}O_{1,5,0}O_{1,6,0}O_{1,7,0}O_{1,8,0} \\ O_{1,1,0}O_{1,2,1}O_{1,3,0}O_{1,4,0}O_{1,5,0}O_{1,6,0}O_{1,7,0}O_{1,8,0} \\ O_{1,1,0}O_{1,2,0}O_{1,3,1}O_{1,4,0}O_{1,5,0}O_{1,6,0}O_{1,7,0}O_{1,8,0} \\ O_{1,1,0}O_{1,2,0}O_{1,3,0}O_{1,4,1}O_{1,5,0}O_{1,6,0}O_{1,7,0}O_{1,8,0} \\ O_{1,1,0}O_{1,2,1}O_{1,3,0}O_{1,4,0}O_{1,5,0}O_{1,6,0}O_{1,7,0}O_{1,8,0} \\ O_{1,1,0}O_{1,2,0}O_{1,3,0}O_{1,4,0}O_{1,5,1}O_{1,6,0}O_{1,7,0}O_{1,8,0} \\ O_{1,1,0}O_{1,2,0}O_{1,3,0}O_{1,4,0}O_{1,5,0}O_{1,6,1}O_{1,7,0}O_{1,8,0} \\ O_{1,1,0}O_{1,2,0}O_{1,3,0}O_{1,4,0}O_{1,5,0}O_{1,6,1}O_{1,7,0}O_{1,8,0} \\ O_{1,1,0}O_{1,2,0}O_{1,3,0}O_{1,4,0}O_{1,5,0}O_{1,6,0}O_{1,7,0}O_{1,8,0} \\ O_{1,1,0}O_{1,2,0}O_{1,3,0}O$$

 $0.85 \times 0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75$ $0.75 \times 0.85 \times 0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75$ $0.75 \times 0.75 \times 0.85 \times 0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75$ $0.75 \times 0.75 \times 0.85 \times 0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75$ $0.75 \times 0.75 \times 0.75 \times 0.85 \times 0.75 \times 0.75 \times 0.75 \times 0.75$ $0.75 \times 0.85 \times 0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75$ $0.75 \times 0.75 \times 0.85 \times 0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75$ $0.75 \times 0.75 \times 0.85 \times 0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75$ $0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75$ $0.75 \times 0.85 \times 0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75$ $0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75$ $0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75$ $0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75$ $0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75 \times 0.75$

 $= 0.85^{11} \times 0.75^{77} \approx 4.01167 \times 10^{-11}$

We expected this to be the expected value for all sensor outputs, but, even so, the space for all of these sensor outputs after 11 states is so large that we yield an incredibly small probability.

3.4.2
$$\{a, b\}, \{b, c\}, \{c, d\}, \{d, f\}, \{b, d\}, \{c, f\}, \{d, h\}, \{b, e\}, \{e, h\}, \{f, e\}, \{f, h\}\}$$

For the following sequence, if we're clever, we can recognize that it contains every *actual* hidden state in addition to one false positive. There are thus 6 true negatives, 1 true positive, and 1 false positive for each of the 13 states. We can use this to group the exponents:

$$P([\{a,b\},\{b,c\},\{c,d\},\{d,f\},\{b,d\},\{c,f\},\{d,h\},\{b,e\},\{e,h\},\{f,e\},\{f,h\}]) = 0.75^{66} \times 0.25^{11} \times 0.85^{11} \approx 2.26460 \times 10^{-16}$$

Notice how just one false positive per state can reduce the probability by a substantial amount (by a factor of roughly a hundred thousand).

3.4.3 $\{\{\}, \{b\}, \{\}, \{f\}, \{\}, \{f\}, \{\}, \{h\}, \{\}, \{e\}, \{\}\}\}$

This sensor output sequence will probably have the lowest probability of all. There is only one true positive and several false positives. The sensors seemed to be taking a nap in this sequence.

$$P(\{\{\},\{b\},\{\},\{f\},\{\},\{f\},\{\},\{h\},\{\},\{e\},\{\}\}\}) = 0.75^{73} \times 0.25^{4} \times 0.85 \times 0.15^{10} \approx 1.45065 \times 10^{-20}$$

As expected, a very low probability. The ten false negatives really bring the probability for this sequence down.

3.5 Same Sensor Responses, Different Sequence

Now consider the hidden sequence $R_2 = [a, b, d, f, d, f, h, e, h, e, f]$ and the same sensor responses.

3.5.1 [{a}, {b}, {c}, {d}, {b}, {c}, {d}, {b}, {e}, {f}, {h}]
$$P([\{a\}, \{b\}, \{c\}, \{d\}, \{b\}, \{c\}, \{d\}, \{b\}, \{e\}, \{f\}, \{h\}]) = 0.75^{68} \times 0.25^{9} \times 0.85^{2} \times 0.15^{9} \approx 3.38287 \times 10^{-22}$$

3.5.2
$$\{a, b\}, \{b, c\}, \{c, d\}, \{d, f\}, \{b, d\}, \{c, f\}, \{d, h\}, \{b, e\}, \{e, h\}, \{f, e\}, \{f, h\}\}$$

The astute observer will note that this sequence of sensor outputs includes all of the actual hidden states of the first sequence and the second sequence. This makes the probability for the two exactly the same.

$$P([\{a,b\},\{b,c\},\{c,d\},\{d,f\},\{b,d\},\{c,f\},\{d,h\},\{b,e\},\{e,h\},\{f,e\},\{f,h\}]) = 0.75^{66} \times 0.25^{11} \times 0.85^{11} \approx 2.26460 \times 10^{-16}$$

3.5.3 [{}, {b}, {}, {f}, {}, {f}, {}, {h}, {}, {e}, {}]
$$P([{}, {b}, {}, {f}, {}, {f}, {}, {h}, {}, {e}, {}]) = 0.75^{76} \times 0.25 \times 0.85^4 \times 0.15^7 \approx 7.12706 \times 10^{-17}$$

4 HMM: Other Possible Applications?

4.1 Can a HMM Work for Baby MQLs?

For normal MQLs, movement seems very straightforward to model with a HMM. In the stationary state, they have a probability of staying in the stationary state (S), transitioning to the move left one state ("releasing the magnetic force on the left foot," what I will call ML1), or transitioning to the move right one state (MR1). According to the description, ML1 will transition to ML2 ("moving the left foot outward") with 100% probability. Then, ML2 will transition to ML3 ("activating the [left foot] magnet") with 100% probability. ML3 will transition to ML4 ("releasing the magnet on the right foot") with 100% probability. ML4 will transition to ML5 ("drawing the legs together") with 100% probability. ML5 will transition to S ("activating the right foot magnet") with 100% probability. A similar sequence can represent movement rightward. Note that some of these actions taken in isolation appear when moving right, as well, but it is important to differentiate them, as, when we take the movement as a whole, the states transition in a very different way between leftwards movement and rightwards movement.

For baby MQLs, the story is actually not much different. However, instead of two paths which are relatively deterministic, there is some chance of branching. As it was not specified in the problem, I will assume that the baby MQL will continue the movement sequence even if its arm is up. Thus, unless the baby falls, it will take the same amount of moves to complete the movement. However, for all states with a magnet detached, there will be a second state corresponding to the baby losing its balance and raising its arm. For example, the state ML1B corresponds to the state when the baby has just started its journey leftward and has raised its left arm because its left leg's magnet became detached. The state transition diagram is below:

Note that not even this figure tells the whole story, as, at each state, we observe the MQL. Thus, each state would emit an angle for each of the MQL's inner legs, inner arms, outer legs, and outer arms. We

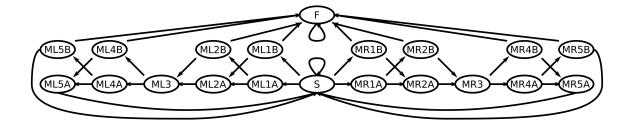


Figure 3: A state transition diagram for the movement of a baby MQL. Nodes ending in "B" represent movement when the baby is starting to lose its balance and has thus raised its corresponding arm. Note that these states can only happen when the MQL has deactivated one of its magnets. "F" represents being in the fallen state. "S" is the stationary state.

might even measure body shape, as we did in a few assignments back. Each state would have a different probability distribution for each of these emittances.

4.2 Can a HMM Work for a Malfunctioning Metronome?

A normal metronome would be easy to model with a HMM, but, in this case, the failure rate is depedent on age. This means that, as the metronome ages, its transition probabilities also change. Unfortunately, this means that we do not have a constant transition matrix. Therefore, this scenario cannot be represented by a HMM.

5 Conclusion

HMMs are powerful for modeling state transitions over time. We have seen how they can be applied to model a student's progression (or lack thereof) through college, a cell's stages of hibernation and growth, a hiker's path through a web of trails, and a baby MQL's first steps. However, HMMs have some limitations. First, they can only represent a fininte number of states. This makes them efficient to implement using vectors and matrices, but, sometimes, real world scenarios require a continuous universe of states. Second, if transitions are represented as a matrix, each row in that matrix must correspond to a probability distribution and thus sum to one. Third, a first-order HMM assumes that a state only depends on the immediately previous state and no other states, which can be a shaky assumption to make in many scenarios. Fourth, the model is hidden in the fact that the *true* state can only be inferred through a series of observations, which can have a variety of probability distributions. Last, the trasition matrix must remain constant throughout all of the transitions.