## Homework9

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### 1 Homework

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```
[]: import pandas as pd
import numpy as np
from sklearn.preprocessing import OneHotEncoder
from IPython.display import display_markdown
import platform
import os
```

### 1.1 Problem 1

```
file_path = os.getcwd() + "/"
if platform.system() == "Windows":
    file_path = file_path.replace("/", "\\")

pb1 = pd.read_csv(file_path + "hmm_pb1.csv", header=None)
pb1 = np.array(pb1).squeeze()
```

```
[]: transition_probs = np.array([[0.95, 0.05], [0.10, 0.90]]) # a values
emission_probs = np.array([[1/6, 1/6, 1/6, 1/6, 1/6, 1/6], [1/10, 1/10, 1/10, 1/10, 1/10, 1/10]) # b values
initial_probs = np.array([0.5, 0.5]) # pi values

# # Observed sequence (dice rolls)
# obs_seq = [1, 3, 5, 2, 4, 6]
```

```
[]: # Define number of hidden states (fair, loaded)
NUM_STATES = 2

# Define the number of time steps (observations) on the shape of pb1
num_time_steps = pb1.shape[0]

# Initialize arrays for storing values
log_vit_probs = np.zeros((NUM_STATES, num_time_steps))
maximizing_states = np.zeros((NUM_STATES, num_time_steps))
most_likely_seq = -1 * np.ones_like(pb1)
```

```
# Calculate intial log probs for each state
     initial_log_probs = np.log(emission_probs[:, pb1[0] - 1] * initial_probs)
     log_vit_probs[:, 0] = initial_log_probs
[]: # Iterative step of the Viterbi algorithm
     for t in range(1, pb1.shape[0]):
         # Calculate log probs of transitioning form previous states to current_{\sqcup}
         transition_log_probs = np.log(transition_probs) + log_vit_probs[:, t-1].
      \hookrightarrowreshape(-1,1)
         # Calculate log probs of emitting the current observations from each state
         emission_log_probs = np.log(emission_probs[:, pb1[t] - 1])
         # Compute total log probs for each state at current time step t
         total_log_probs = emission_log_probs + np.max(transition_log_probs, axis=0)
         # Update Viterbi matrix with total log probs
         log_vit_probs[:, t] = total_log_probs
         # Store maximizing state for this t
         maximizing_states[:, t] = np.argmax(transition_log_probs, axis=0)
     # Find most likely sequence of hidden states
     # Initialize last stae in most liekly sequence
     most_likely_seq[-1] = np.argmax(log_vit_probs[:, -1])
     # Iterate backward through time steps to determine the most likely sequence
     for t in range(pb1.shape[0] - 1, 0, -1):
         # Determine the state at the current time step based on the pointer from
      → the next time step
         most_likely_seq[t - 1] = maximizing_states[most_likely_seq[t], t]
     # Display the most likely sequence of hidden states
     display_markdown("The most probable sequence of the hidden state is:", raw=True)
     print(most_likely_seq + 1) # Add 1 to each state index for human-readable_
      \hookrightarrow output
```

The most probable sequence of the hidden state is:

[]: # Initialization of the Viterbi algorithm

## 

```
[]: | # Define a function that implements the forward-backward algorithms
     def forward backward(dataset, initial probs, transition probs, emission probs):
        log_alpha = np.zeros((2, dataset.shape[0]))
        log_beta = np.zeros((2, dataset.shape[0]))
        # Compute forward probs
        log_alpha[:, 0] = np.log(emission_probs[:, dataset[0] - 1] * initial_probs)
        log_alpha[:, 0] -= log_alpha[:, 0].sum()
        for t in range(1, dataset.shape[0]):
             alpha_numerator = np.log(emission_probs[:, dataset[t] - 1]) + np.log(np.
      sum(transition_probs * np.exp(log_alpha[:, t - 1].reshape(-1, 1)), axis=0))
             alpha_denominator = np.log(np.sum(np.exp(alpha_numerator)))
             log_alpha[:, t] = alpha_numerator - alpha_denominator
         # Compute backward probs
        log_beta[:, -1] = 0.5
        for t in range(dataset.shape[0] - 2, -1, -1):
             beta_numerator = np.log(np.sum(transition_probs * (np.exp(log_beta[:, t_
      4+ 1]) * emission_probs[:, dataset[t + 1] - 1]), axis=1))
             beta_denominator = np.log(np.sum(np.exp(beta_numerator)))
             log_beta[:, t] = beta_numerator - beta_denominator
        return log_alpha, log_beta
```

# 

The following shows  $\alpha_{138}^1/\alpha_{138}^2$ .

### 12.487100265980972

The following shows  $\beta_{138}^1/\beta_{138}^2$ .

### 3.540995425670584

### 1.2 Problem 2

```
[]: pb2 = pd.read csv(file path + "hmm pb2.csv", header=None)
     pb2 = np.array(pb1).squeeze()
[]: # Define a function that implements the forward-backward algorithms
     def forward_backward(dataset, initial_probs, transition_probs, emission_probs):
         log_alpha = np.zeros((2, dataset.shape[0]))
         log_beta = np.zeros((2, dataset.shape[0]))
         epsilon = 1e-10  # Small epsilon value to handle zero probabilities and
      →avoid division by zero
         # Compute forward probs
         emission_probs_safe = np.maximum(emission_probs[:, dataset[0] - 1], epsilon)
         log_alpha[:, 0] = np.log(emission_probs_safe * initial_probs)
         log_alpha[:, 0] -= log_alpha[:, 0].sum()
         for t in range(1, dataset.shape[0]):
             # Handle zero or negative emission probabilities by adding a small_{\sqcup}
      ⇔epsilon value
             emission probs safe = np.maximum(emission probs[:, dataset[t] - 1],
      ⇔epsilon)
             alpha_numerator = np.log(emission_probs_safe) + np.log(np.
      sum(transition_probs * np.exp(log_alpha[:, t - 1].reshape(-1, 1)), axis=0))
             alpha_denominator = np.log(np.sum(np.exp(alpha_numerator)))
             log_alpha[:, t] = alpha_numerator - alpha_denominator
         # Compute backward probs
         log_beta[:, -1] = 0.5
         for t in range(dataset.shape[0] - 2, -1, -1):
             # Handle zero or negative emission and transition probabilities by \Box
      ⇔adding a small epsilon value
             emission_probs_safe = np.maximum(emission_probs[:, dataset[t + 1] - 1],_
      ⇔epsilon)
             transition_probs_safe = np.maximum(transition_probs, epsilon)
             beta_numerator = np.log(np.sum(transition_probs_safe * (np.

exp(log_beta[:, t + 1]) * emission_probs_safe), axis=1))

             beta_denominator = np.log(np.sum(np.exp(beta_numerator)))
             log_beta[:, t] = beta_numerator - beta_denominator
         return log_alpha, log_beta
```

```
[]: def baum_welch_algorithm(dataset, initial_probs, transition_probs,_u
      →emission_probs, num_iter=10):
         # Initialization
         learned initial probs = initial probs # learned pi
         learned_transition_probs = transition_probs # learned a
         learned_emission_probs = emission_probs # learned b
         epsilon = 1e-10
         # Create one hot encoded dataset for calculation
         encoder = OneHotEncoder(categories = [[1,2,3,4,5,6]], sparse_output=False)
         encoded_dataset = np.transpose(encoder.fit_transform(dataset.reshape(-1,1)))
         for i in range(num_iter):
             # Compute forward and backward probabilities
             alpha, beta = forward_backward(dataset, learned_initial_probs,__
      Glearned_transition_probs, learned_emission_probs)
             ## E-step
             # Calculate b x(t+1)^{\hat{i}}
             next_emission_probs = learned_emission_probs[:, np.roll(dataset - 1,__
      →-1)]
             # Handle zero or negative emission probabilities by adding a small_{f \sqcup}
      ⇔epsilon value
             emission_probs_safe = np.maximum(next_emission_probs, epsilon)
             # Compute xi
             xi_numerator = alpha.reshape(2, 1, -1) * learned_transition_probs.
      \Rightarrowreshape(2, 2, 1) * np.roll(
                 beta.reshape(1, 2, -1), shift=-1, axis=-1) * emission_probs_safe.
      \rightarrowreshape(1, 2, -1)
             xi_denominator = np.sum(xi_numerator, axis=(0, 1)).reshape(1, 1, -1)
             xi_denominator[xi_denominator == 0] = epsilon # Ensure we do not divide_
      ⇔by zero
             xi = xi_numerator / xi_denominator
             # Check for division by zero in xi_denominator
             assert xi_denominator.min() != 0
             # Calculate gamma (expected number of visits to each state)
             gamma_numerator = alpha * beta
             gamma_denominator = np.sum(gamma_numerator, axis=0).reshape(1, -1)
             gamma_denominator[gamma_denominator == 0] = epsilon # ensure we don't_\( \)
      ⇔divide by zero
             gamma = gamma_numerator / gamma_denominator
```

```
# Check for division by zero in gamma_denominator
             assert gamma_denominator.min() != 0
             ## M-step
             # Update parameters based on the expected counts
             learned_initial_probs = gamma[:, 0]
             learned_transition_probs = np.sum(xi[:, :, :-1], axis=2) / np.
      \Rightarrowsum(gamma[:, :-1], axis=1).reshape(-1, 1)
             learned_emission_probs = np.sum(gamma.reshape(2, 1, -1) *__
      -encoded dataset.reshape(1, 6, -1), axis=2) / np.sum(gamma, axis=1).
      \rightarrowreshape(-1, 1)
         return learned_initial_probs, learned_transition_probs,_
      →learned_emission_probs
[]: # Run Baum-Welch algorithm
     learned_initial_probs, learned_transition_probs, learned_emission_probs = __
      →baum_welch_algorithm(pb2, initial_probs, transition_probs, emission_probs,
      ⇒2000)
[]: # Display the learned parameters
     display_markdown("The learned initial probability is:", raw=True)
     print(learned_initial_probs)
     display_markdown("The learned transition probability is:", raw=True)
     print(learned_transition_probs)
     display_markdown("The learned emission probability is:", raw=True)
     print(learned_emission_probs)
    The learned initial probability is:
    [0.28806428 0.71193572]
    The learned transition probability is:
    [[0.46355313 0.53577229]
     [0.59087941 0.40972353]]
    The learned emission probability is:
    [[2.40936207e-01 8.08309542e-12 8.08184746e-12 6.05410064e-12
      7.51223751e-12 7.59063793e-01]
     [3.37287677e-12 2.26580671e-01 2.26580671e-01 2.76931932e-01
      2.64344117e-01 5.56260842e-03]]
```