

DS-GA 3001.001 Special Topics in Data Science: Probabilistic Time Series Analysis
Homework 3

Due date: Oct 25, by 6pm
 YG390

Problem 1. (15p)

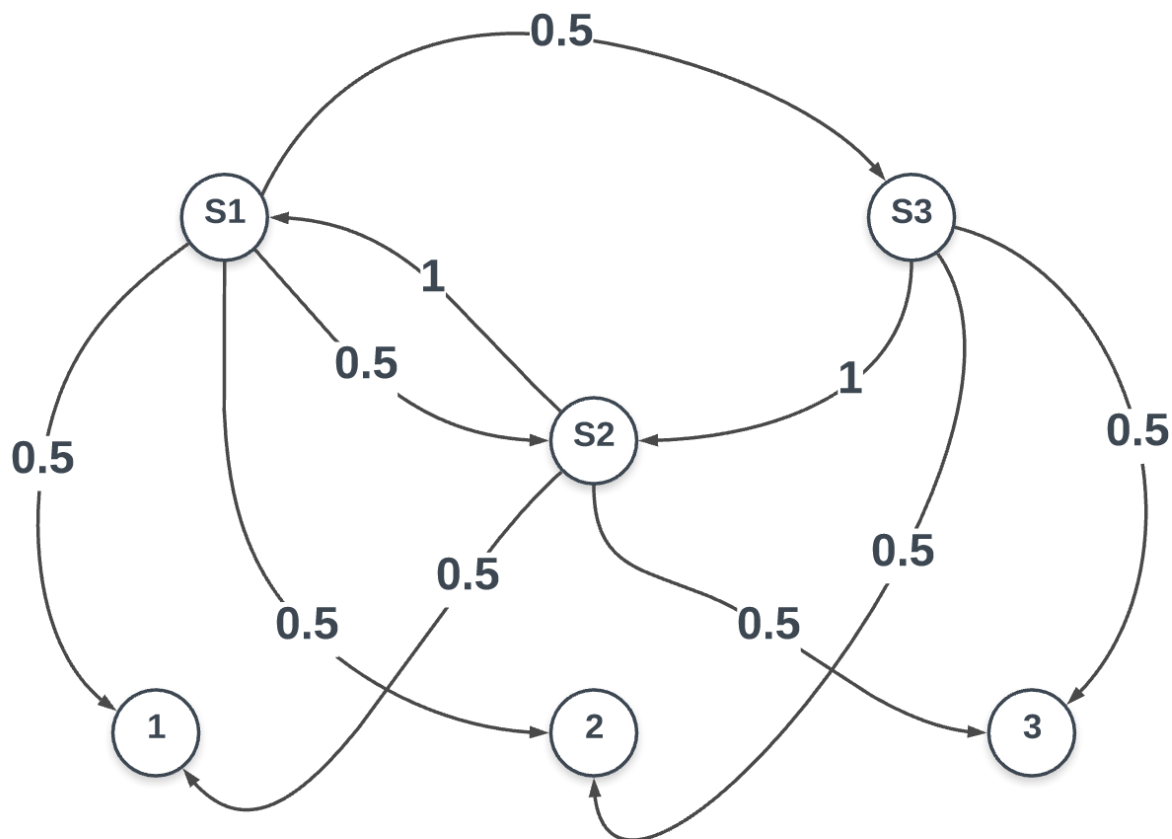
Consider the HMM with $K=3$ latent states and discrete observations $\{1, 2, 3\}$, with parameters specified by:

initial distribution $\pi = [1, 0, 0]$, transition matrix $\mathbf{A} = \begin{bmatrix} 0 & 0.5 & 0.5 \\ 1 & 0 & 0 \\ 0 & 1 & 0 \end{bmatrix}$, where $A_{ij} = P(z_{t+1} = j | z_t = i)$ and

likelihood $P(x_t | z_t)$ described by matrix entries B_{xz} : $\mathbf{B} = \begin{bmatrix} 0.5 & 0.5 & 0 \\ 0.5 & 0 & 0.5 \\ 0 & 0.5 & 0.5 \end{bmatrix}$.

Write down all possible state sequences consistent with observations a) 1, 2, 3 and b) 1, 3, 1.

Let the three latent states be $\{S_1, S_2, S_3\}$. Given the HMM with parameters $\{A, B, \pi\}$, the model can be described as:



When observing the observations 1, 2, and 3 with initial distribution π which implies we start in state S_1 then observing 2 means we have 0.5 probability of being in state S_1 or S_3 . However staying in state S_1 is not possible so the only state after seeing the sequence 1, 2 is state S_3 . Then the last observation 3 means we are in state S_2 or S_3 . but from state S_3 , the only existing transition is from S_3 to S_2 . So the sequence of observations 1, 2, 3 corresponds to the state sequence $\{S_1, S_3, S_2\}$.

When observing 1, 2, 1 with initial distribution π , seeing 2 after 1 implies we can be in state S_2 50% of the time or S_3 the rest of the time. The last observation 1 implies that the latent space is either S_1 or S_2 . Based on the transition matrix \mathbf{A} then the only possible state sequences for the sequence of observations $\{1, 2, 1\}$ are $\{S_1, S_2, S_1\}$ or $\{S_1, S_3, S_2\}$.

Problem 2. (15p)

Construct an HMM that generates the observation sequence $A^{k_1}C^{k_2}A^{k_3}C^{k_4}$ where A^{k_1} denotes k_1 repeats of symbol A and the number of repeats k_i are drawn from the set $\{1, 2, 3\}$ with equal probability.

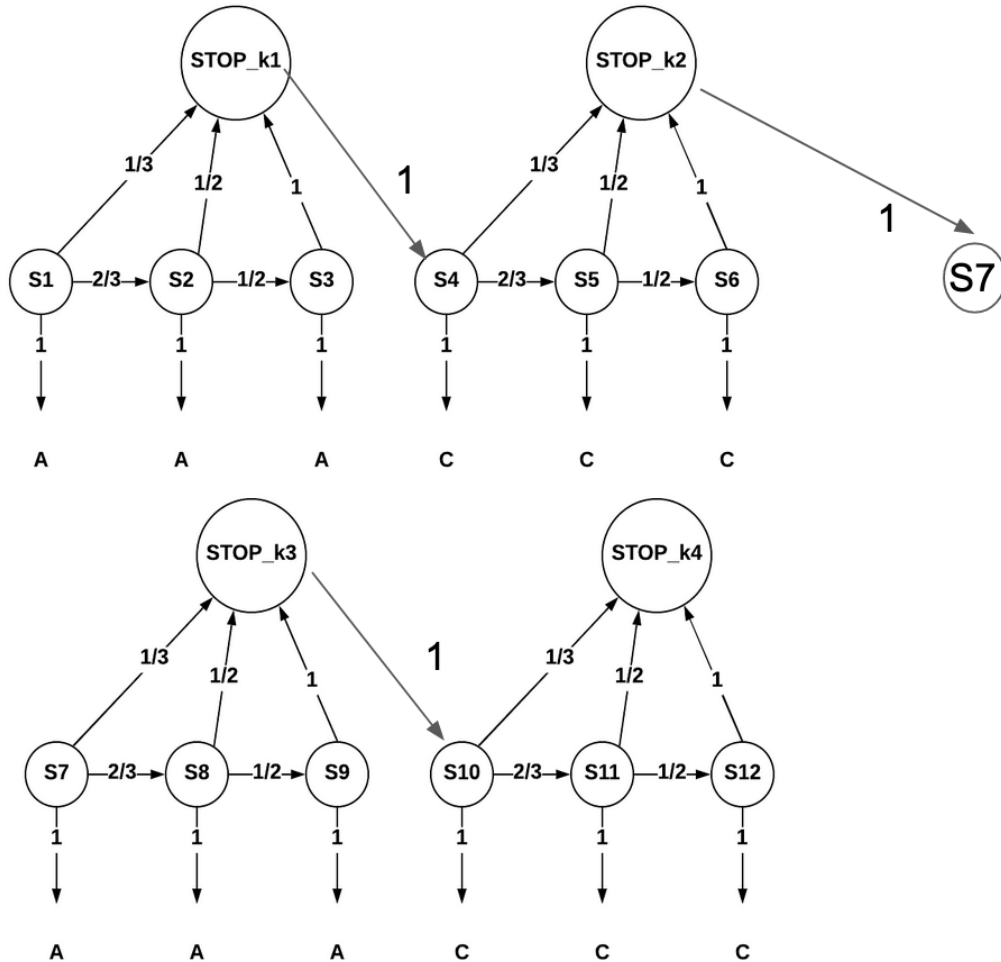
The HMM model is defined by three class of parameters $\{\mathbf{A}, \mathbf{B}, \pi\}$, and is described in the graph below. Note that there is a direct connection between the states STOP_k2 and S_7 which I broke down into two for convenience of representation. The latent states are $\{S_1, S_2, S_3, S_4, S_5, S_6, S_7, S_8, S_9, S_{10}, S_{11}, S_{12}\}$. Sequences based on k_i have equal probability $\frac{1}{3}$, meaning, that for example for k_1 : A, AA, AAA have the same probability $\frac{1}{3}$. The transition matrix has for elements the probabilities indicated on the graph and $\sum_j A_{ij} = 1$.

$$B_{ih} = \frac{\text{num of times state } i \text{ emits } h}{\text{num state } i}$$

The emission probabilities \mathbf{B} , is the identity matrix since there are as many observations repeated for A , then the number of state S_i , and the same for C and S_j .

$$\pi_i = \frac{\text{num of chains start with } i}{\text{total num of chains}}$$

All the chains start with a sequence of A in S_1 , the initial distribution is $\pi = [100000000000]^T$.



Problem 3. (20p)

Implement EM for an HMM model with K states and gaussian observations (full derivations in handout). Use this code to fit the weekly S&P 500 returns data (data/sp500w.csv) for $K = 2$ vs. $K = 3$ and compare the two results.

Hint: Use Example 6.17 from tsa4 textbook as guideline for plots and interpretation.

For the parameters initialization, we use a gaussian mixture which gives us estimates for the mean and the covariance of each components namely μ_k and Σ_k from the handout. The initial distribution of the initial state

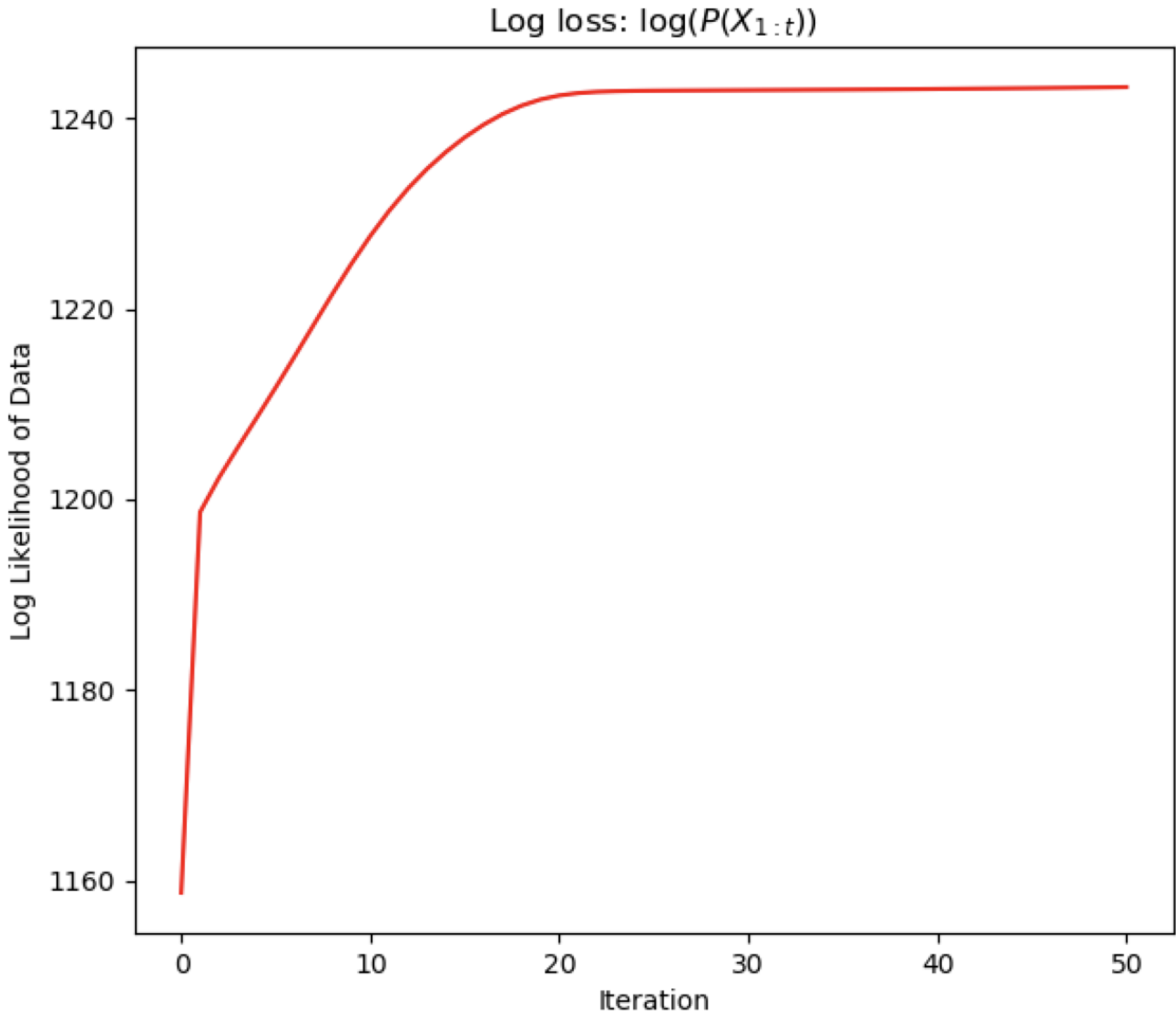
\mathbf{z}_1, π is implemented by an E-M step. The initial values of the means are provided by KMeans and updated by an expectation-maximization algorithm.

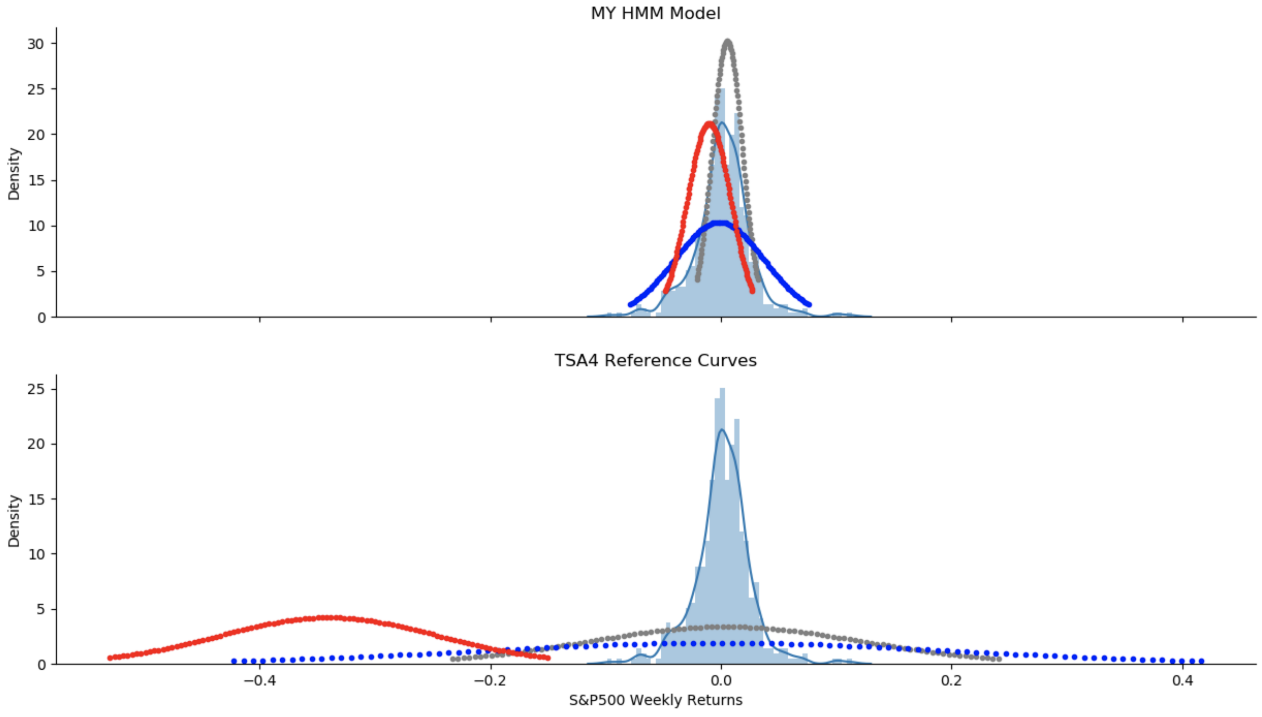
Once the initialization is completed, we alternate between the E steps, for which we run the alpha-beta algorithm with the current estimates, determining:

$$\begin{aligned}\hat{\alpha}(\mathbf{z}_i) &= P(\mathbf{z}_i | \mathbf{x}_{1:i}) = \frac{\alpha(\mathbf{z}_i)}{P(\mathbf{x}_{1:i})} \\ \hat{\beta}(\mathbf{z}_i) &= \frac{P(\mathbf{x}_{i+1:t} | \mathbf{z}_i)}{P(\mathbf{x}_{i+1:t} | \mathbf{x}_{1:i})} = \frac{\beta(\mathbf{z}_i)}{\prod_{j=i+1:t} c_j} \\ \gamma(\mathbf{z}_i) &= \hat{\alpha}(\mathbf{z}_i) \hat{\beta}(\mathbf{z}_i) \\ \xi(\mathbf{z}_{i,j}, \mathbf{z}_{i+1,k}) &= c_{i+1}^{-1} \hat{\alpha}(\mathbf{z}_i) P(\mathbf{z}_{i+1} | \mathbf{z}_i) P(\mathbf{x}_{i+1} | \mathbf{z}_{i+1}) \hat{\beta}(\mathbf{z}_{i+1})\end{aligned}$$

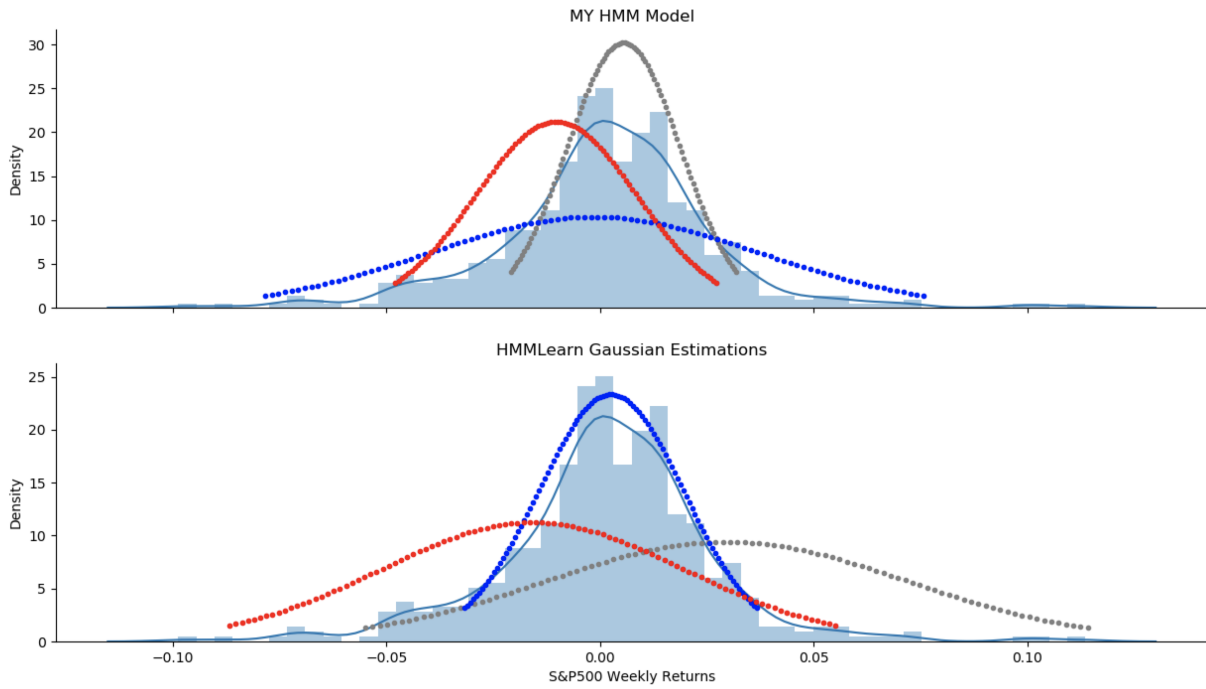
And the M step, which updates the parameters:

The result for $K = 3$, are plotted below with the top plot showing the log loss of the data, the middle plot the inference obtained with our model and the bottom the plot using the means and covariances indicated by ts4 textbook. We can see that our model to some extent captures better the distribution of the raw data.

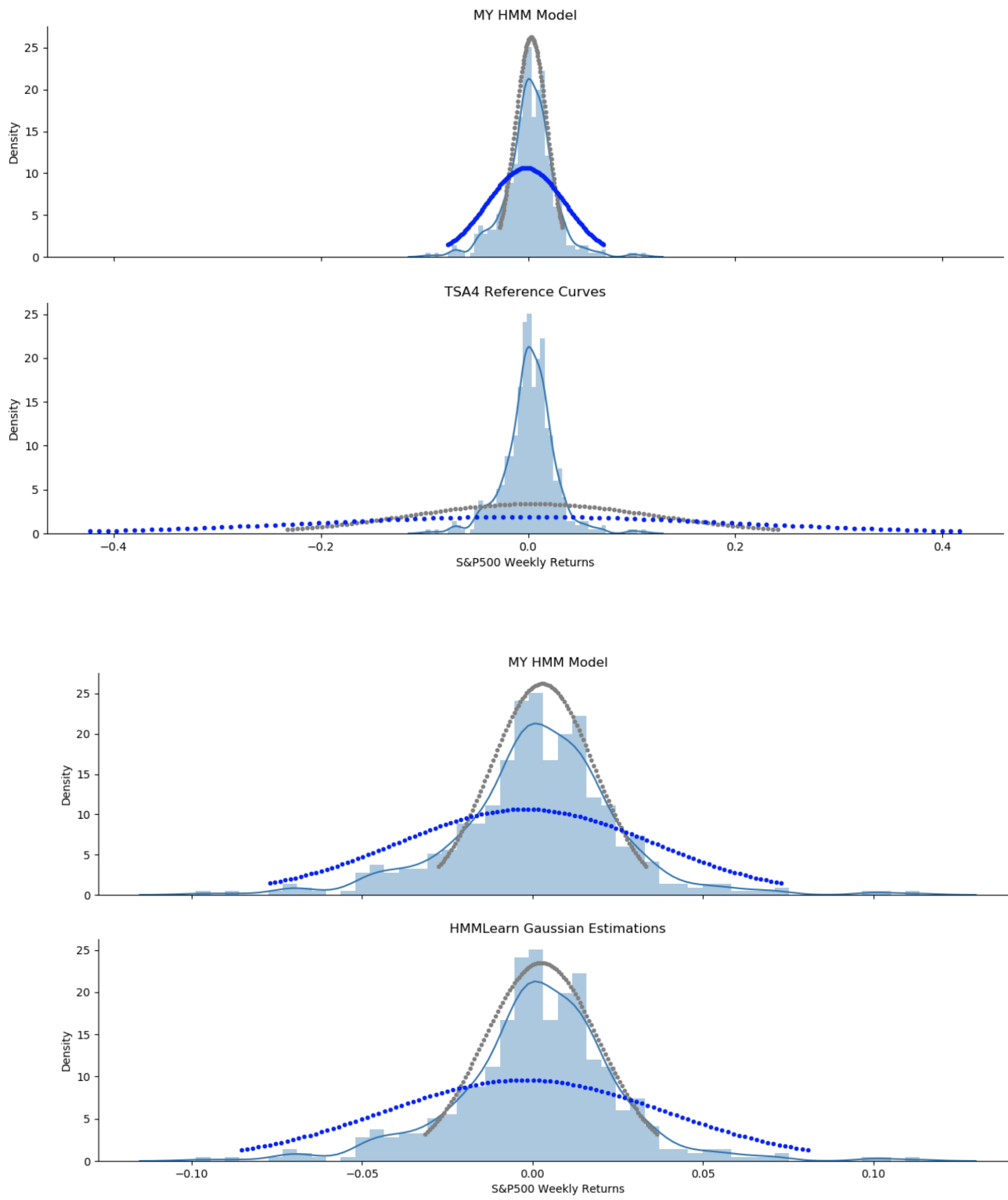




The performance of our model compared to hmmlearn Gaussian smoother, are similar, the GaussianHMM model covering more of the data along the x-axis whereas our model cover the distribution better along the y-axis.



For $K = 2$, the fit is worst but we can see that tsa4 model does not cover the distribution along the y-axis. Compared to hmmlearn gaussian model, our model performance is very similar.



Lastly, the log loss for $K = 2$:

