Conceptual and Theoretical Questions

Q1 (8 pts)

Because we are trying to estimate the parameters w in a maximum likelihood problem, the Expectation Maximization (EM) algorithm may be used to find \hat{w} . In this specific case of finding \hat{w} in an HMM problem, we can use the Baum-Welch algorithm, a special case of the EM algorithm.

First, perform the forward-backward procedure. The forward procedure is calculating the probability of observing the observed values from 1 to some t, as well as the t step of the hidden variable, given some initialized state of parameters w_i .

In the backwards-procedure, calculate the probability of seeing the same observed sequence given the hidden variable and w_i .

Finally, perform the update step of the EM algorithm, using Bayes' theorem and the probabilities calculated using the forward-backward procedures. From this, the parameters of the HMM can be updated.

Q2 (6 pts)

With \hat{f} defined by (11.2):

$$\hat{f} = rac{1}{L} \sum_l^L f(z^{(l)})$$

The expectation $E[\hat{f}\,]$ would be:

$$egin{aligned} E[\hat{f}\,] &= rac{1}{L} \sum_{l}^{L} E[f(z^{(l)})] \ &= rac{1}{L} \sum_{l}^{L} E[f] \ &= rac{1}{L} (L \cdot E[f]) \ &= E[f] \end{aligned}$$

We want to show $Var[\hat{f}\,]=rac{1}{L}E[(f-E[\hat{f}\,])^2]$ Start with the definition of variance:

$$Var[\hat{f}\,] = E[(\hat{f}\,-E[\hat{f}\,])^2] = E[\hat{f}\,]^2 - E[\hat{f}\,^2]$$

$$E[f] = E[\hat{f}\,]$$
 , so $E[f]^2 = E[\hat{f}\,]^2$.

Evaluate $E[\hat{f}^{\,2}]$:

$$egin{aligned} E[\hat{f}^{\,2}] &= \left(\sum_{l}^{L} f^{2}(z^{l})
ight)^{2} = \sum_{l}^{L} f^{2}(z^{l}) \ \hat{f}^{\,2} &= \sum_{i,j|i
eq j}^{L} f(z^{i})f(z^{j}) + \sum_{l}^{L} f^{2}(z^{l}) \ E[\hat{f}^{\,2}] &= rac{1}{L^{2}} E[f^{2}] + rac{1}{L} \sum_{i,j|i
eq j}^{L} E[f(z^{i})f(z^{j})] \ E[\hat{f}^{\,2}] &= rac{1}{L} E[f^{2}] + \sum_{i,j|i
eq j}^{L} E[f(z^{i})f(z^{j})] \ E[\hat{f}^{\,2}] &= rac{1}{L} E[f^{2}] + \left(1 - rac{1}{L}
ight) E[f]^{2} \end{aligned}$$

Combine both these terms to get:

$$Var[\hat{f}\,]=rac{1}{L}E[(f-E[\hat{f}\,])^2]$$

Q3 (6 pts)

Here, we can use Equation (11.5) to find p(y):

$$p(y) = p(z) \cdot \left| rac{dz}{dy}
ight|$$

Differentiate the inverse of the given equation of y to find $\left| \frac{dz}{dy} \right|$:

$$egin{aligned} rac{y-c}{b} &= an z \ z &= an^{-1} rac{y-c}{b} \ rac{d}{dx} an^{-1} x &= rac{1}{1+x^2} \ \left|rac{dz}{dy}
ight| &= rac{1}{1+\left(rac{y-c}{b}
ight)^2} \end{aligned}$$

As z is a uniform distributon, $p(z) = b^{-1}$, so (11.5) becomes:

$$p(y)=rac{b^{-1}}{1+\left(rac{y-c}{b}
ight)^2}$$

Which roughly follows the form given by (11.16):

$$\frac{k}{1 + \left(\frac{z-c}{h}\right)^2}$$

Application Questions

```
In []: # Imports
    import numpy as np
    from numpy import pi, sqrt, exp
    import pandas as pd
    from sklearn.gaussian_process import GaussianProcessRegressor
    from sklearn.gaussian_process.kernels import RBF
    from sklearn.linear_model import LinearRegression
    from sklearn.model_selection import cross_val_score, KFold
    from sklearn.metrics import mean_squared_error

import matplotlib.pyplot as plt
    from scipy.stats import norm
    from scipy import integrate
```

Viterbi Decoding Algorithm (20 pts)

```
In [ ]: | # Part a
                     obs_states = ["A", "C", "G", "T"]
                      hidden_states = ["H", "L"]
                      transition = np.array([
                                                              [0.5, 0.4],
                                                              [0.5, 0.6]
                                                              1)
                      emission = np.array([
                                                    [0.2, 0.3, 0.3, 0.2],
                                                    [0.3, 0.2, 0.2, 0.3]
                                                    1)
                      def viterbi_alg(obs):
                                N = len(obs)
                                h states = len(hidden states)
                                path_probs = np.zeros((h_states, N)) # 1st array is H probs, 2nd is L prob
                                for i in range(0, N):
                                          for k_index in range(len(hidden_states)): # k_index is either H (0) or
                      L (1)
                                                    nuc idx = obs states.index(obs[i])
                                                    e_l = emission[k_index][nuc_idx]
                                                    prev_p = path_probs.T[i - 1] if i > 0 else emission[:, nuc_idx]
                                                    transition_prob = transition[:, k_index] if i > 0 else np.array([
                      1.0 / h_states] * h_states)
                                                    \# k\_prob = e_l * max(prev\_p[0] * transition[0][k\_index],
                                                                                                                               prev_p[1] * transition[1][k_index])
                                                    k_{prob} = e_1 * max(prev_p[0] * transition_prob[0], prev_p[1] * transition_prob[0], prev_p[1] * transition_prob[0] * transition_prob
                      nsition_prob[1])
                                                    path_probs[k_index][i] = k_prob
                               mp_path = ""
                                for probs in path_probs.T:
                                          # mx = np.array(np.argmax(probs)).reshape(1,)
                                          # print(probs)
                                          mx = np.argwhere(probs == np.amax(probs)).reshape(1,)
                                          # print(mx)
                                          # "X" means H and L are equilikely to appear in that position
                                          mp_path += hidden_states[mx[0]] if len(mx) == 1 else "X"
                                print(mp_path)
                      viterbi alg("GGCACTGAA")
                      Because of the extra precision in my calculations, compared to the example PD
                      F, the path is slightly different (but equivalent).
```

HHHLHLHLL

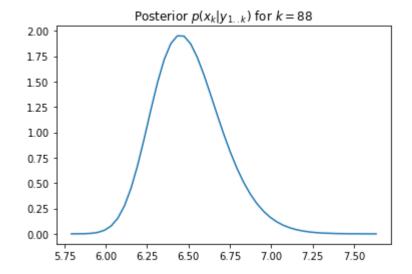
```
In [ ]: # Part b
viterbi_alg("AGTCGTA")
```

Bayesian Filtering (20 pts)

```
In [ ]: # Part a
    x_sim = [np.random.standard_normal()]
    y_sim = []
    for k in range(1, 101):
        x_sim.append(np.random.normal(0.99 * x_sim[k - 1] + 0.1, 0.1))
        y_sim.append(np.random.normal(-2 * x_sim[k] + 1, 0.4))
```

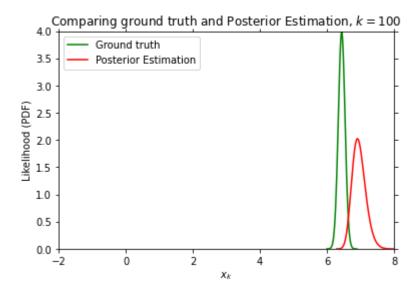
```
In [ ]: # Part b
                         post x = np.linspace(-10, 10, num=500)
                         # print(post_x)
                         def posterior prob(k, x, y=y sim):
                                     assert k >= 1, "k must be greater than or equal to 1"
                                     # measure_model = Lambda x_k: 1 / sqrt(0.4 * 2 * pi) * exp(-0.5 * (<math>y[k - 1] + pi)
                           1] - (-2 * x k + 1)) ** 2 / 0.4)
                                    measure model = 1 / sqrt(0.4 * 2 * pi) * exp(-0.5 * (y[k - 1] - (-2 * x + pi)) * exp(-0.5 * (y[k - 1] - (-2 * x + pi)) * exp(-0.5 * (y[k - 1] - (-2 * x + pi)) * exp(-0.5 * (y[k - 1] - (-2 * x + pi)) * exp(-0.5 * (y[k - 1] - (-2 * x + pi)) * exp(-0.5 * (y[k - 1] - (-2 * x + pi)) * exp(-0.5 * (y[k - 1] - (-2 * x + pi)) * exp(-0.5 * (y[k - 1] - (-2 * x + pi)) * exp(-0.5 * (y[k - 1] - (-2 * x + pi)) * exp(-0.5 * (y[k - 1] - (-2 * x + pi)) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(
                         1)) ** 2 / 0.4)
                                    def chap_kol():
                                                 \# dyn_mu = Lambda x_k_prev: (0.99 * x_k_prev) + 0.1
                                                mu = (0.99 * x) + 0.1
                                                 st_norm = norm.freeze(loc=0, scale=1)
                                                 return integrate.cumulative_trapezoid(norm.pdf(x, loc=mu, scale=0.1) *
                                                                                                                                   (posterior prob(k - 1, x, y) if k > 1 else
                         st_norm.pdf(x)),
                                                                                                                                   x, initial=0)
                                    ck = chap_kol()
                                     z_k = integrate.trapezoid(measure_model * ck, x)
                                     return (1 / z k) * measure model * ck
                         k sel = 88
                         post_y = posterior_prob(k_sel, post_x)
                         fig_post, ax_post = plt.subplots()
                         # print(integrate.trapezoid(post_y, post_x))                                  # Sanity check if you don't think
                         the AUC of the PDF is 1
                         post y[np.abs(post y) < 1e-4] = np.nan
                         ax_post.plot(post_x, post_y)
                         ax post.set title(r"Posterior p(x k|y \{1...k\}) for " + f"k=\{k sel\}")
```

Out[]: Text(0.5, 1.0, 'Posterior $p(x k|y \{1..k\})$ for k=88')



```
In [ ]: # Part c
        gt_means = np.zeros(100)
        gt vars = np.array([0.1] * 100)
        est means = np.zeros(100)
        est vars = np.zeros(100)
        fig comp, ax posterior = plt.subplots()
        ax posterior.set xlim(-1, 8)
        ax posterior.set ylim(0,4)
        def posterior vs ground truth(k, x=x sim, y=y sim):
            ax posterior.clear()
            ax_posterior.set_xlim(-2, 8)
            ax posterior.set ylim(0,4)
            x sam = np.linspace(-5, 9, num=900)
            ground truth = norm.pdf(x sam, loc=0.99 * x[k - 1] + 0.1, scale=0.1)
            ground_truth[np.abs(ground_truth) < 1e-4] = np.nan</pre>
            ax_posterior.plot(x_sam, ground_truth, label="Ground truth", color="green"
        )
            gt means[k - 1] = 0.99 * x[k - 1] + 0.1
            post est = posterior prob(k=k, x=x sam, y=y)
            est_means[k - 1] = x_sam[np.argmax(post_est)]
            est_vars[k - 1] = np.nanvar(post_est)
            post est[np.abs(post est) < 1e-4] = np.nan</pre>
            ax posterior.plot(x sam, post est, label="Posterior Estimation", color="re
        d")
            ax posterior.set title(f"Comparing ground truth and Posterior Estimation,
         k=\{k\}")
            ax posterior.set xlabel("$x k$")
            ax posterior.set ylabel("Likelihood (PDF)")
            ax posterior.legend()
        fig sum, ax sum = plt.subplots()
        for i in range(1,101):
            posterior_vs_ground_truth(i)
        k \text{ vals} = np.arange(1, 101, step=1)
        ax_sum.plot(k_vals, gt_means, label="Ground Truth Means", color="r")
        ax_sum.fill_between(k_vals, gt_means + sqrt(gt_vars), gt_means - sqrt(gt_vars
        ), color="r", alpha=0.4)
        ax sum.plot(k vals, est means, label="Estimate Means", color="b")
        ax_sum.fill_between(k_vals, est_means + sqrt(est_vars), est_means - sqrt(est_v
        ars), color="b", alpha=0.4)
        ax sum.set xlabel("$k$")
        ax_sum.set_ylabel(r"$\mu$")
        ax sum.set title(r"$\mu$ of Ground Truth and Estimate Posteriors")
        ax sum.legend()
```

Out[]: <matplotlib.legend.Legend at 0x28c395f9fd0>



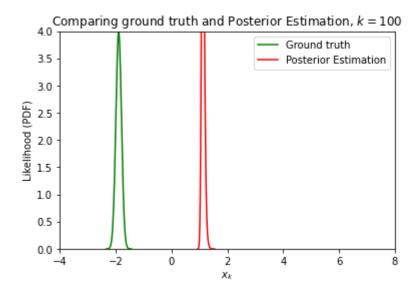
μ of Ground Truth and Estimate Posteriors Ground Truth Means Estimate Means k

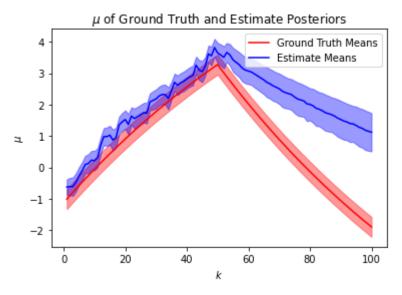
```
In [ ]: # For producing video animation of estimate vs ground truth
    import matplotlib.animation as animation
    plt.rcParams['animation.ffmpeg_path'] = "D:\\ffmpeg\\bin\\ffmpeg.exe"
    anim = animation.FuncAnimation(
        fig_comp, lambda i: posterior_vs_ground_truth(i + 1, x_sim, y_sim),
        frames=99,
        interval=1000)

writervideo = animation.FFMpegWriter(fps=5)
    anim.save("posteriors_pt_b.mp4", writer=writervideo)
```

```
In [ ]: # Part d
        df filter = pd.read excel("filter problem.xlsx", header=None, names=["index",
         "x", "y"])
        x_filter = df_filter["x"].to_numpy()
        y_filter = df_filter["y"].dropna().to_numpy()
        print(len(y filter))
        gt means d = np.zeros(100)
        gt_vars_d = np.array([0.1] * 100)
        est means d = np.zeros(100)
        est_vars_d = np.zeros(100)
        fig d, ax d = plt.subplots()
        def posterior_vs_ground_truth_d(k, x=x_filter, y=y_filter):
            ax d.clear()
            ax_d.set_xlim(-4, 8)
            ax_d.set_ylim(0,4)
            x sam = np.linspace(-5, 9, num=900)
            ground_truth = norm.pdf(x_sam, loc=0.99 * x[k - 1] + 0.1, scale=0.1)
            ground truth[np.abs(ground truth) < 1e-4] = np.nan</pre>
            ax_d.plot(x_sam, ground_truth, label="Ground truth", color="green")
            gt means d[k - 1] = 0.99 * x[k - 1] + 0.1
            post est = posterior prob(k=k, x=x sam, y=y)
            est_means_d[k - 1] = x_sam[np.argmax(post_est)]
            est vars d[k - 1] = np.nanvar(post est)
            post est[np.abs(post est) < 1e-4] = np.nan</pre>
            ax_d.plot(x_sam, post_est, label="Posterior Estimation", color="red")
            ax_d.set_title(f"Comparing ground truth and Posterior Estimation, $k={k}$"
        )
            ax_d.set_xlabel("$x_k$")
            ax_d.set_ylabel("Likelihood (PDF)")
            ax d.legend()
        # posterior vs ground truth d(99, x=x \text{ filter}, y=y \text{ filter})
        fig_sum_d, ax_sum_d = plt.subplots()
        for i in range(1,101):
            posterior_vs_ground_truth_d(i, x=x_filter, y=y_filter)
        k_vals = np.arange(1, 101, step=1)
        ax_sum_d.plot(k_vals, gt_means_d, label="Ground Truth Means", color="r")
        ax_sum_d.fill_between(k_vals, gt_means_d + sqrt(gt_vars_d), gt_means_d - sqrt(
        gt_vars_d), color="r", alpha=0.4)
        ax_sum_d.plot(k_vals, est_means_d, label="Estimate Means", color="b")
        ax sum d.fill between(k vals, est means d + sqrt(est vars d), est means d - sq
        rt(est_vars_d), color="b", alpha=0.4)
        ax sum d.set xlabel("$k$")
        ax sum d.set ylabel(r"$\mu$")
        ax sum d.set title(r"$\mu$ of Ground Truth and Estimate Posteriors")
        ax_sum_d.legend()
```

Out[]: <matplotlib.legend.Legend at 0x1ae81bd92d0>

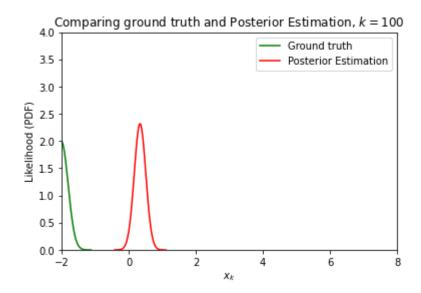


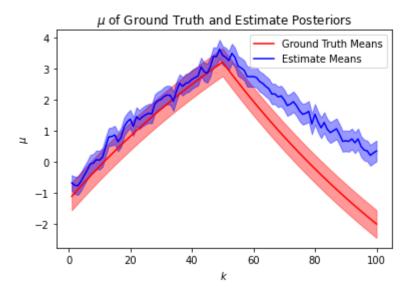


```
In [ ]: # Part e
                            gt_means_e = np.zeros(100)
                            gt vars e = np.array([0.2] * 100)
                            est_means_e = np.zeros(100)
                            est_vars_e = np.zeros(100)
                            def posterior_prob_e(k, x, y=y_filter):
                                         assert k >= 1, "k must be greater than or equal to 1"
                                         # measure_model = lambda x_k: 1 / sqrt(0.4 * 2 * pi) * exp(-0.5 * (y[k - 2 * pi) * (y[k - 2 * pi) * exp(-0.5 * (
                              1] - (-2 * x_k + 1) ** 2 / 0.4)
                                        measure_model = 1 / sqrt(0.4 * 2 * pi) * exp(-0.5 * (y[k - 1] - (-2 * x + pi)) * exp(-0.5 * (y[k - 1] - (-2 * x + pi)) * exp(-0.5 * (y[k - 1] - (-2 * x + pi)) * exp(-0.5 * (y[k - 1] - (-2 * x + pi)) * exp(-0.5 * (y[k - 1] - (-2 * x + pi)) * exp(-0.5 * (y[k - 1] - (-2 * x + pi)) * exp(-0.5 * (y[k - 1] - (-2 * x + pi)) * exp(-0.5 * (y[k - 1] - (-2 * x + pi)) * exp(-0.5 * (y[k - 1] - (-2 * x + pi)) * exp(-0.5 * (y[k - 1] - (-2 * x + pi)) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(-0.5 * (y[k - 1] - (-2 * x + pi))) * exp(
                            1)) ** 2 / 0.4)
                                         def chap kol():
                                                      st norm = norm.freeze(loc=0, scale=1)
                                                      norm funcs = np.zeros((len(x), len(x)))
                                                     for loc, i in enumerate(x):
                                                                   norm_funcs[loc] = norm.pdf(i, loc=x, scale=0.2)
                                                      rec_res = posterior_prob(k - 1, x, y) if k > 1 else st_norm.pdf(x)
                                                      return np.matmul(norm_funcs, rec_res.T)
                                                      # return integrate.cumulative_trapezoid(norm.pdf(x, loc=x, scale=0.2)
                                                                                                                                                       (posterior\_prob\_e(k - 1, x, y) if k > 1)
                               else st_norm.pdf(x),
                                                                                                                                                       x, initial=0)
                                         ck = chap kol()
                                         z_k = integrate.trapezoid(measure_model * ck, x)
                                         return (1 / z k) * measure model * ck
                            fig_e, ax_e = plt.subplots()
                            def posterior_vs_ground_truth_e(k, x=x_filter, y=y_filter):
                                         ax e.clear()
                                         ax_e.set_xlim(-2, 8)
                                         ax_e.set_ylim(0, 4)
                                         x_sam = np.linspace(-5, 9, num=900)
                                         ground_truth = norm.pdf(x_sam, loc=x[k - 1], scale=0.2)
                                         ground truth[np.abs(ground truth) < 1e-4] = np.nan</pre>
                                         ax_e.plot(x_sam, ground_truth, label="Ground truth", color="green")
                                         gt_means_e[k - 1] = x[k - 1]
                                         post_est = posterior_prob_e(k=k, x=x_sam, y=y)
                                         est_means_e[k - 1] = x_sam[np.argmax(post_est)]
                                         est vars e[k - 1] = np.nanvar(post est)
                                         post_est[np.abs(post_est) < 1e-4] = np.nan</pre>
                                         ax_e.plot(x_sam, post_est, label="Posterior Estimation", color="red")
                                        ax e.set title(f"Comparing ground truth and Posterior Estimation, $k={k}$"
                                         ax e.set xlabel("$x k$")
                                         ax e.set ylabel("Likelihood (PDF)")
                                         ax_e.legend()
                            posterior_prob_e(k=1, x=np.linspace(-10,10,num=900))
```

```
fig_sum_, ax_sum = plt.subplots()
for i in range(1,101):
    posterior_vs_ground_truth_e(i, x=x_filter, y=y_filter)
k_vals = np.arange(1, 101, step=1)
ax_sum.plot(k_vals, gt_means_e, label="Ground Truth Means", color="r")
ax_sum.fill_between(k_vals, gt_means_e + sqrt(gt_vars_e), gt_means_e - sqrt(gt_vars_e), color="r", alpha=0.4)
ax_sum.plot(k_vals, est_means_e, label="Estimate Means", color="b")
ax_sum.fill_between(k_vals, est_means_e + sqrt(est_vars_e), est_means_e - sqrt(est_vars_e), color="b", alpha=0.4)
ax_sum.set_xlabel("$k$")
ax_sum.set_ylabel(r"$\mu$")
ax_sum.set_title(r"$\mu$ of Ground Truth and Estimate Posteriors")
ax_sum.legend()
```

Out[]: <matplotlib.legend.Legend at 0x1aea24a51b0>





Part f

In part c, the estimated posterior distributions closely matched the simulated data produced by the state and measurement models, although they did consistently overestimate by a small amount. This can also be seen in the video "posteriors_pt_b.mp4".

In part d, the model begins by closely estimating the ground truth data, but the posteriors do not translate leftwards as fast as the data does.

In part e, the same pattern occurs, but with overall higher accuracy.

Sequential MCMC (20 pts)

```
In [ ]: | # Part a
        df filter = pd.read excel("filter problem.xlsx", header=None, names=["index",
        "x", "y"])
        x filter = df filter["x"]
        y_filter = df_filter["y"].dropna().to_numpy()
        df filter.head()
        gt means = np.zeros(100)
        gt vars = np.array([0.1] * 100)
        def sis(particles, k max):
            w = np.full((particles, k_max), 1.0 / particles) # weights, initialized to
        1/particles
            x = np.zeros((particles, k max))
            x[:, 0] = np.random.standard normal(size=(particles,)) # Particles, initia
         lized to x1
            means = np.zeros(k max)
            vars = np.zeros(k_max)
            for k in range(1, k max):
                # Note that our proposal distribution is equal to our dynamical model
                imp_particles = np.random.choice(x[:, k - 1], size=particles, p=w[:, k
        - 1])
                for i in range(particles):
                     x[i, k] = np.random.normal(loc=imp_particles[i], scale=0.2)
                     state model = norm.pdf(x[i, k], x[i, k - 1], scale=0.2)
                     measure\_model = norm.pdf(y\_filter[k - 1], loc=-2 * x[i][k - 1] + 1
         , scale=0.4)
                     w[i, k] = measure model * state model
                w[:, k] /= sum(w[:, k])
                means[k - 1] = np.mean(x[:, k])
                vars[k - 1] = np.var(x[:, k])
            return means, vars
        def compare_sis(ax, particles):
            x_sis = np.arange(1, 101, step=1)
            gt means = x filter[0:100]
            gt vars = [0.2] * 100
            ax.plot(x_sis, gt_means, color="green", label="Ground truth means")
            ax.fill_between(x_sis, gt_means - sqrt(gt_vars), gt_means + sqrt(gt_vars),
        color="green", alpha=0.4)
            means sis, vars sis = sis(particles=particles, k max=100)
            ax.plot(x_sis, means_sis, color="red", label="Estimated means")
            ax.set title(f"Estimated vs ground truth means using SIS, particles={parti
        cles}")
            ax.fill between(x sis, means sis + sqrt(vars sis), means sis - sqrt(vars s
        is), color="red", alpha=0.4)
            ax.legend()
        fig_sis, ax_sis = plt.subplots()
        compare sis(ax sis, 100)
```

```
# For some reason, my plots for a and b of this question do not appear in my Jupyter Notebook export, so I have attached them separately in Canvas
```

```
In [ ]: # Part b
fig_sis_b, ax_sis_b = plt.subplots()
compare_sis(ax_sis_b, 1000)
```

GP Linear Regression (20 pts)

```
In [ ]: | # Part a
        df_sm = pd.read_csv("synchronous machine.csv",
                            delimiter=";",
                            decimal=",")
        print(df_sm.head())
        x_sm = df_sm.drop(columns=["If", "dIf"]).values
        y_sm = df_sm["If"].values
        def model mse(model):
            train_mse = []
            test mse = []
            for train_loc, test_loc in KFold(n_splits=10, shuffle=True).split(x_sm, y_
        sm):
                train x, test x = x sm[train loc], x sm[test loc]
                train_y, test_y = y_sm[train_loc], y_sm[test_loc]
                model_tmp = None
                if model == GaussianProcessRegressor:
                    model tmp = model(random state=0)
                else:
                    model_tmp = model()
                model tmp.fit(train x, train y)
                train_mse.append(mean_squared_error(model_tmp.predict(train_x), train_
        y))
                test mse.append(mean squared error(model tmp.predict(test x), test y))
            return train_mse, test_mse
        gp_train_mse, gp_test_mse = model_mse(GaussianProcessRegressor)
            Ιy
                  ΡF
                         e
                              dIf
                                      Ιf
        0 3.0 0.66 0.34 0.383 1.563
        1 3.0 0.68 0.32 0.372 1.552
        2 3.0 0.70 0.30 0.360 1.540
        3 3.0 0.72 0.28 0.338 1.518
        4 3.0 0.74 0.26 0.317 1.497
In [ ]: | # Part b
        lin_model = LinearRegression()
```

lin_train_mse, lin_test_mse = model_mse(LinearRegression)

print(lin train mse)

```
In [ ]: # Part c
        err x = np.arange(1, 11, step=1)
        fig comp, ax comp = plt.subplots()
        ax_comp.plot(err_x, gp_train_mse, label="GP MSE")
        ax_comp.plot(err_x, lin_train_mse, label="Linear Regression MSE")
        ax_comp.set_title("Comparing MSE of GP and Linear Regression Models (Train)")
        ax comp.set xlabel("Cross validation folds")
        ax comp.set ylabel("MSE")
        ax_comp.legend()
        fig_comp, ax_comp = plt.subplots()
        ax_comp.plot(err_x, gp_test_mse, label="GP MSE")
        ax_comp.plot(err_x, lin_test_mse, label="Linear Regression MSE")
        ax comp.set title("Comparing MSE of GP and Linear Regression Models (Test)")
        ax_comp.set_xlabel("Cross validation folds")
        ax comp.set ylabel("MSE")
        ax_comp.legend()
         ="""
        Evidently, the MSE in both the train and test datasets is lower for the Gaussi
        an Process model, and higher in the Linear Regression model
```

