

Task1

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```
[ ]: import numpy as np
import matplotlib.pyplot as plt
from IPython.display import display, Markdown, Math
import pandas as pd

plt.style.use("default")
```

7 Task A

7.1 The probability density function $p(x|\theta)$ for this model is the sum of the two Gaussian pdf with the factors π_1, π_2 with $\pi_1 + \pi_2 = 1$ so the integral over $p(x|\theta)$ is still 1.

$$8 \quad p(x|\theta) = \pi_1 \frac{1}{\sqrt{2\pi\sigma_{c_1}^2}} \exp\left(-\frac{1}{2\sigma_{c_1}^2}(x - \mu_{c_1})^2\right) + \pi_2 \frac{1}{\sqrt{2\pi\sigma_{c_2}^2}} \exp\left(-\frac{1}{2\sigma_{c_2}^2}(x - \mu_{c_2})^2\right)$$

```
[ ]: # We want to have random samples. Therefore draw N numbers of uniform
      ↳ distribution and if the
      # value is below 0.4 we draw a value from the first Gaussian and above from the
      ↳ second Gaussian

def gen_samples(N, pi, mu, sigma):
```

```

    assert all([len(l) == 2 for l in [pi, mu, sigma]]), "Lists need to have the
↳length 2."

    randoms = np.random.random(N)
    samples = []
    for r in randoms:
        if r <= pi[0]:
            samples.append(np.random.normal(mu[0], sigma[0], 1))
        else:
            samples.append(np.random.normal(mu[1], sigma[1], 1))

    return np.array(samples).flatten()

# Just the normal distribution
def gaussian_pdf(x, mu, sigma):

    return 1 / np.sqrt(2 * np.pi * sigma ** 2) * np.exp(- (x - mu) ** 2 / (2 *
↳sigma ** 2))

# Combined Gaussian distribution
def pdf_combined(x, pi, mu, sigma):
    length = len(pi)
    assert all([len(l) == length for l in [pi, mu, sigma]]), "Lists need to
↳have the same length."

    prob = 0
    for p, m, s in zip(pi, mu, sigma):
        prob += p * gaussian_pdf(x, m, s)

    return prob

```

9 Task B

```

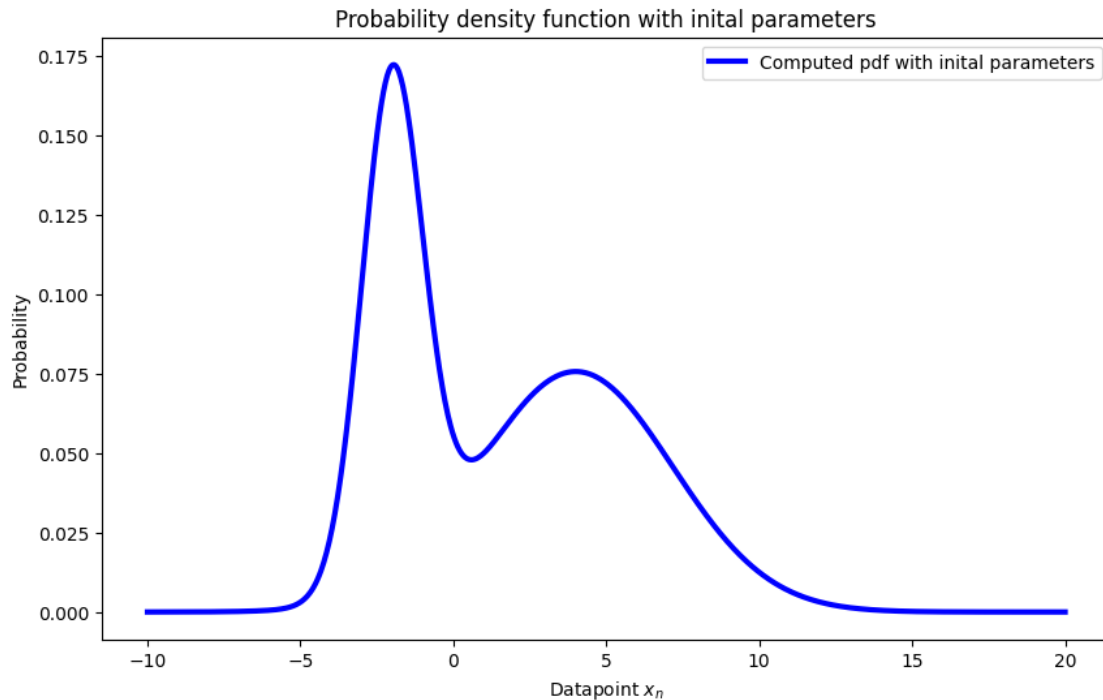
[ ]: fig, ax = plt.subplots(1,1, figsize = (10,6))

# Parameters and x-parameters for line plot
x = np.linspace(-10, 20, 1000)
pi = [0.4, 0.6]
mu = [-2, 4]
sigma = [1, np.sqrt(10)]

# Compute pdf
pdf_task_b = pdf_combined(x, pi, mu, sigma)
ax.plot(x, pdf_task_b, c="b", lw=3, label="Computed pdf with inital parameters")
ax.set_xlabel("Datapoint $x_n$")
ax.set_ylabel("Probability")

```

```
ax.set_title("Probability density function with inital parameters")
ax.legend()
plt.show()
```



10 Task C + D

10.1 Scatter Plot and histograms of sampled data

```
[ ]: pi = [0.4, 0.6]
mu = [-2, 4]
sigma = [1, np.sqrt(10)]

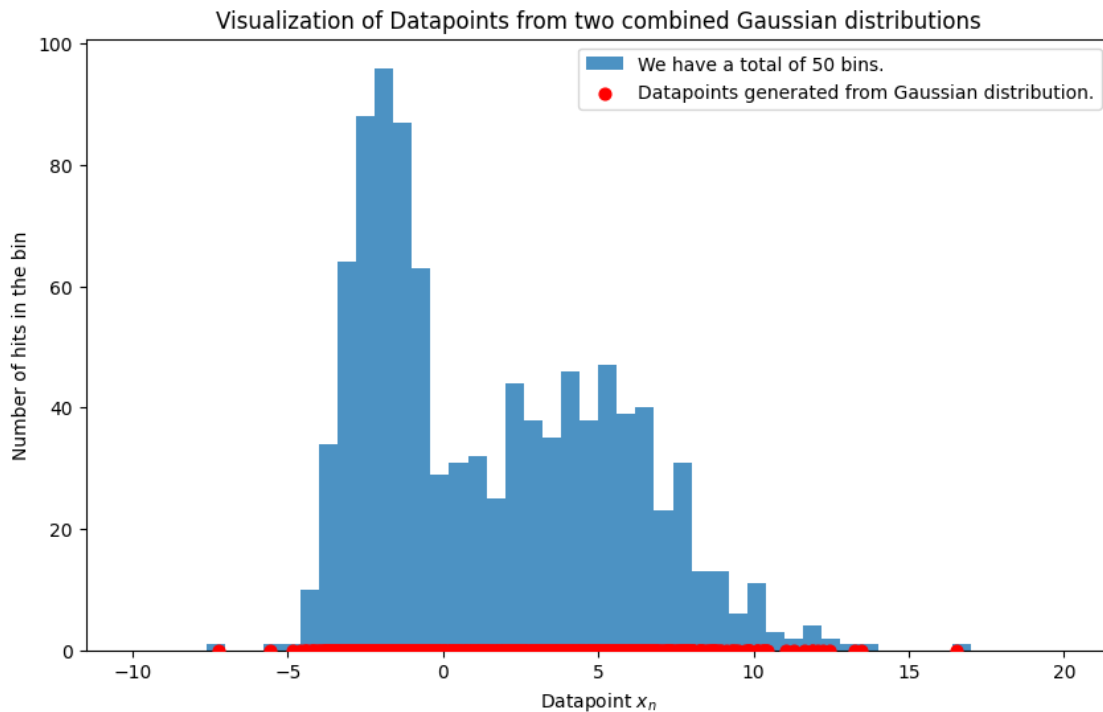
fig, ax = plt.subplots(1,1, figsize = (10,6))
gaussian_combined_samples = gen_samples(1000, pi, mu, sigma)
y=np.zeros(len(gaussian_combined_samples))

# Create histogram of the distributed datapoints
bins = np.linspace(-10, 20, 51)
ax.scatter(gaussian_combined_samples, y, c="r", zorder=10, s=40,
    ↳label="Datapoints generated from Gaussian distribution.")
ax.hist(gaussian_combined_samples, bins, alpha = 0.8, label=f"We have a total
    ↳of {len(bins)-1} bins.")
```

```

ax.set_xlabel("Datapoint  $x_n$ ")
ax.set_ylabel("Number of hits in the bin")
ax.set_title("Visualization of Datapoints from two combined Gaussian_
↳distributions")
ax.legend()
ax.set_ylim(bottom=-0.01)
plt.show()

```



11 Task E

11.1 The same as Task D but with normalized height of histogram

```

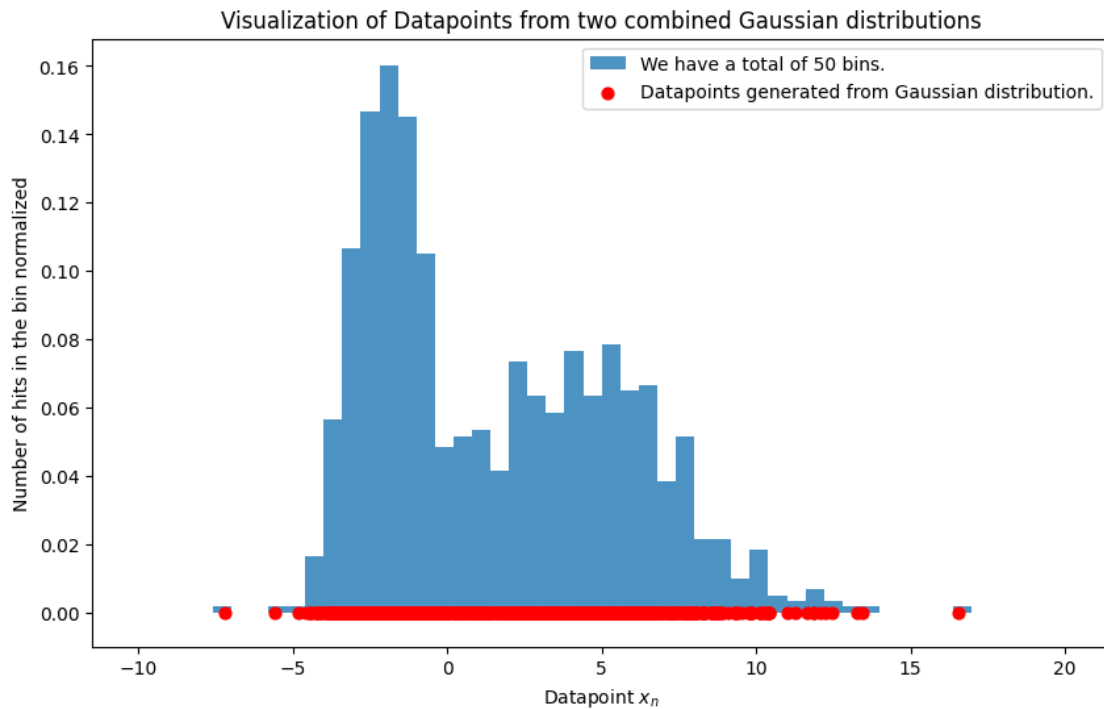
[ ]: fig, ax = plt.subplots(1,1, figsize = (10,6))

# Create histogram of the distributed datapoints
ax.scatter(gaussian_combined_samples, y, c="r", zorder=10, s=40,
↳label="Datapoints generated from Gaussian distribution.")
ax.hist(gaussian_combined_samples, bins, alpha = 0.8, density=True, label=f"We_
↳have a total of {len(bins)-1} bins.")

ax.set_xlabel("Datapoint  $x_n$ ")
ax.set_ylabel("Number of hits in the bin normalized")

```

```
ax.set_title("Visualization of Datapoints from two combined Gaussian_
↳distributions")
ax.legend()
ax.set_ylim(bottom=-0.01)
plt.show()
```



12 Task F

12.1 The same as Task E but also with the pdf used in Task B

12.2 The density approximation is better when you use more bins and more data points. With more bins you have a finer division of the x axis and more datapoints increases the possible accuracy in general

```
[ ]: bins_num = [30, 60, 90, 120]
N_num = [100, 1000, 10000, 100_000]

fig, ax = plt.subplots(len(bins_num), len(N_num), sharex=True, sharey=True,
↳figsize = (20, 15))

for i, b in enumerate(bins_num):
    for j, N in enumerate(N_num):
        # Create histogram of the distributed datapoints
        bins = np.linspace(-10, 20, b+1)
```

```

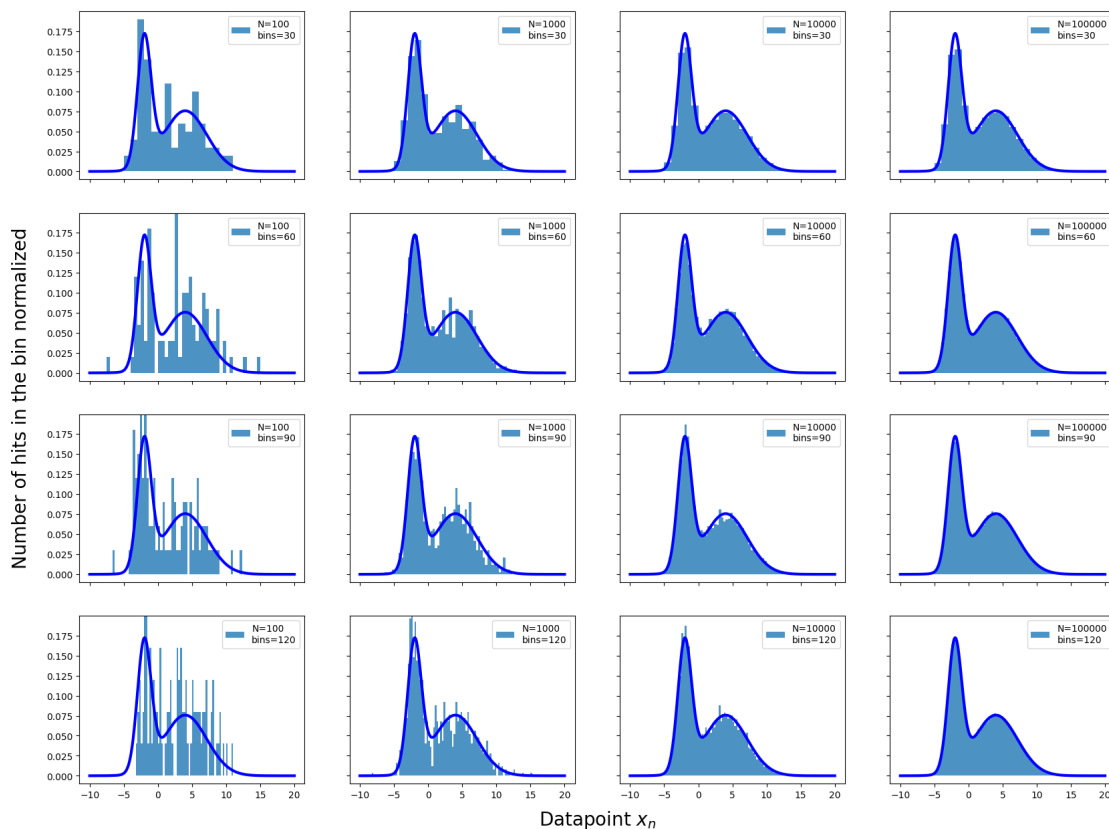
gaussian_combined_samples = gen_samples(N, pi, mu, sigma)
ax[i,j].hist(gaussian_combined_samples, bins, alpha = 0.8,
→density=True, label=f"N={N}\nbins={b}")
ax[i,j].plot(x, pdf_task_b, c="b", lw=3)

ax[i,j].legend()
ax[i,j].set_ylim(bottom=-0.01)
plt.suptitle("Histogram of different N datapoints and bin sizes", fontsize=20)

fig.add_subplot(111, frameon=False)
plt.tick_params(labelcolor='none', which='both', top=False, bottom=False,
→left=False, right=False)
plt.xlabel("Datapoint $x_n$", fontsize=20, labelpad=10)
plt.ylabel("Number of hits in the bin normalized", fontsize=20, labelpad=30)
plt.show()

```

Histogram of different N datapoints and bin sizes



13 Task G

14 The general equation for the log likelihood

$$15 \quad \log L(\theta) = \mathcal{L}(\theta) = \sum_{n=1}^N \log[p(x_n|\theta)] = \sum_{n=1}^N \sum_{c=1}^C \log[p(c|\theta) p(x|c, \theta)]$$

$$16 \quad \mathcal{L}(\theta) = \sum_{n=1}^N \log\left[\pi_1 \frac{1}{\sqrt{2\pi\sigma_{c_1}^2}} \exp\left(-\frac{1}{2\sigma_{c_1}^2}(x - \mu_{c_1})^2\right) + \pi_2 \frac{1}{\sqrt{2\pi\sigma_{c_2}^2}} \exp\left(-\frac{1}{2\sigma_{c_2}^2}(x - \mu_{c_2})^2\right)\right]$$

17 Task H

```
[ ]: def log_likelihood(pdf_func, x, **kwargs):  
  
    return np.log(pdf_func(x, **kwargs)).sum()
```

```
[ ]: # Optimal parameters  
pi_correct = [0.4, 0.6]  
mu_correct = [-2, 4]  
sigma_correct = [1, np.sqrt(10)]  
  
# Correct parameters log likelihood  
dfs = []  
  
for N in [100, 1000, 10000]:  
    results = []  
    gaussian_combined_samples = gen_samples(N, pi_correct, mu_correct,   
→sigma_correct)  
    log_likely = log_likelihood(pdf_combined, gaussian_combined_samples,   
→pi=pi_correct, mu=mu_correct, sigma=sigma_correct)  
    results.append({f"N={N}": log_likelihood": np.round(log_likely,0), "Pi":   
→str(np.round(pi_correct,2)), "Mu": str(np.round(mu_correct,2)), "Sigma":   
→str(np.round(sigma_correct,2))})  
  
    # Other parameters with some random noise  
    random_pi = np.random.uniform(size=15)  
    pi_vec = [[p, 1 - p] for p in random_pi]  
    mu_vec = zip(np.random.uniform(-1, 1, 15) - 2, np.random.uniform(-1, 1, 15)   
→+ 4)  
    sigma_vec = zip(np.random.uniform(-1, 1, 15) + 1, np.random.uniform(-1, 1,   
→15) + np.sqrt(10))  
  
    # Compute the log likelihood  
    for pi, mu, sigma in zip(pi_vec, mu_vec, sigma_vec):  
        log_likely = log_likelihood(pdf_combined, gaussian_combined_samples,   
→mu=mu, pi=pi, sigma=sigma)
```

```

        results.append({f"N={N}": log_likelihood": np.round(log_likely,0), "Pi":
↪str(np.round(pi,2)), "Mu": str(np.round(mu,2)), "Sigma": str(np.
↪round(sigma,2))})

df_results = pd.DataFrame(results)
df_results = df_results.set_index(f"N={N}": log_likelihood").
↪sort_index(ascending=False)
dfs.append(df_results)

```

18 Print out dataframes

18.1 As expected: The correct parameters have the best (highest) log likelihood and parameters close to these correct ones give a better log likelihood. Furthermore, the log likelihood increases if the number of datapoints increases because we sum up more values. If we increase the number of datapoints by 10 the log likelihood also increases around a factor of 10.

```

[ ]: for df in dfs:
      display(df)

```

	Pi	Mu	Sigma
N=100: log likelihood			
-252.0	[0.4 0.6]	[-2 4]	[1. 3.16]
-253.0	[0.44 0.56]	[-1.73 4.8]	[0.74 2.84]
-258.0	[0.27 0.73]	[-1.51 3.35]	[1.08 2.78]
-260.0	[0.18 0.82]	[-1.92 3.25]	[1.24 3.35]
-264.0	[0.59 0.41]	[-1.97 4.85]	[1.72 2.81]
-267.0	[0.2 0.8]	[-1.58 3.05]	[1.83 3.08]
-269.0	[0.38 0.62]	[-1.04 4.64]	[0.84 2.88]
-270.0	[0.29 0.71]	[-2.63 4.11]	[1.9 2.61]
-272.0	[0.13 0.87]	[-1.93 3.49]	[0.19 3.53]
-275.0	[0.39 0.61]	[-2.91 4.47]	[0.6 3.65]
-277.0	[0.07 0.93]	[-1.42 3.49]	[1.91 3.56]
-286.0	[0.53 0.47]	[-1.45 4.08]	[0.43 2.88]
-310.0	[0.09 0.91]	[-1.23 4.42]	[0.33 2.82]
-315.0	[0.87 0.13]	[-2.31 3.54]	[0.78 2.54]
-316.0	[0.9 0.1]	[-1.59 4.24]	[0.99 2.88]
-568.0	[1. 0.]	[-2.08 3.5]	[0.72 3.68]
	Pi	Mu	Sigma
N=1000: log likelihood			
-2583.0	[0.4 0.6]	[-2 4]	[1. 3.16]
-2687.0	[0.29 0.71]	[-1.86 4.63]	[1.62 3.26]
-2698.0	[0.58 0.42]	[-1.77 3.73]	[1.91 3.15]
-2704.0	[0.67 0.33]	[-1.69 4.85]	[1.48 3.66]
-2723.0	[0.42 0.58]	[-2.8 4.45]	[1.71 3.55]
-2740.0	[0.41 0.59]	[-1.63 3.23]	[0.42 3.39]

-2763.0	[0.7 0.3]	[-1.98 4.33]	[1.47 3.93]
-2763.0	[0.4 0.6]	[-1.22 3.59]	[1.75 2.31]
-2781.0	[0.45 0.55]	[-1.04 3.39]	[1.07 2.45]
-2782.0	[0.64 0.36]	[-1.67 3.58]	[0.67 2.77]
-2904.0	[0.28 0.72]	[-1.94 4.85]	[0.46 2.44]
-2966.0	[0.88 0.12]	[-1.12 4.22]	[1.85 2.94]
-3021.0	[0.19 0.81]	[-1.4 3.25]	[0.01 3.92]
-3250.0	[0.92 0.08]	[-2.34 4.43]	[1.77 3.6]
-3738.0	[0.52 0.48]	[-2.4 3.79]	[0.05 2.8]
-5446.0	[0.94 0.06]	[-2.39 4.07]	[0.1 2.57]

	Pi	Mu	Sigma
N=10000: log likelyhood			
-26071.0	[0.4 0.6]	[-2 4]	[1. 3.16]
-26241.0	[0.49 0.51]	[-1.79 3.87]	[1.16 2.95]
-27702.0	[0.63 0.37]	[-2.47 4.67]	[0.92 3.63]
-27721.0	[0.21 0.79]	[-2.42 4.35]	[1.96 4.02]
-27966.0	[0.26 0.74]	[-2.89 3.09]	[0.38 3.37]
-28115.0	[0.11 0.89]	[-2.73 4.49]	[0.64 3.89]
-28359.0	[0.7 0.3]	[-2.54 4.73]	[0.99 2.56]
-31199.0	[0.03 0.97]	[-2.89 3.67]	[1. 2.54]
-31256.0	[0.65 0.35]	[-1.44 3.72]	[0.58 2.17]
-31302.0	[0.91 0.09]	[-1.11 3.76]	[1.66 2.78]
-32186.0	[0.48 0.52]	[-2.1 4.87]	[0.24 3.02]
-34182.0	[0.91 0.09]	[-2.94 3.64]	[1.46 2.37]
-35879.0	[0.8 0.2]	[-2.22 3.01]	[0.32 2.44]
-36053.0	[0.75 0.25]	[-2.6 4.63]	[0.26 3.52]
-37181.0	[0.95 0.05]	[-2.39 4.56]	[0.79 3.95]
-38405.0	[0.98 0.02]	[-1.82 3.57]	[1.21 3.1]