COMP0086 Summative Assignment

Nov 14, 2022

Question 1

- (a) Our sample space for images is $\{0,1\}^D$, where each of our D dimensions can only take binary values (D being the number of pixels in the image). The exponential family best suited on this sample space is the D-dimensional multivariate Bernoulli distribution because it shares the same sample space. On the other hand, a D-dimensional multivariate Gaussian has the sample space \mathbb{R}^D , which does not match the sample space of our data. It is not immediately clear how the likelihood of an image of binary (discrete) values would be calculated under the continuous distribution of a multivariate Gaussian. Thus it would be inappropriate to model this dataset of images with a multivariate Gaussian.
- (b) For $\{\mathbf{x}^{(n)}\}_{n=1}^N$, a data set of N images, the joint likelihood (assuming images are independently and identically distributed) is the product of N, D-dimensional multivariate Bernoulli distributions:

$$P(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|\mathbf{p}) = \prod_{n=1}^{N} P(\mathbf{x}^{(n)}|\mathbf{p})$$

Substituting the D-dimensional multivariate Bernoulli:

$$P(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|\mathbf{p}) = \prod_{n=1}^{N} \prod_{d=1}^{D} p_d^{x_d^{(n)}} (1 - p_d)^{1 - x_d^{(n)}}$$

Taking the logarithm, we get the log likelihood:

$$\mathcal{L}(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|\mathbf{p}) = \sum_{n=1}^{N} \sum_{d=1}^{D} [x_d^{(n)} \log(p_d) + (1 - x_d^{(n)}) \log(1 - p_d)]$$

Note that since the logarithm is a monotonically increasing function on \mathbb{R}_+ , the maximisers and minimisers of the likelihood do not change. Thus, to solve for the maximum likelihood estimate, \hat{p}_d , we can take the derivative of $\mathcal{L}(\{\mathbf{x}^{(n)}\}_{n=1}^N|\mathbf{p})$ with respect to p_d , the d^{th} element of \mathbf{p} :

$$\frac{\partial \mathcal{L}(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|\mathbf{p})}{\partial p_{d}} = \sum_{n=1}^{N} \left(\frac{x_{d}^{(n)}}{p_{d}} - \frac{1 - x_{d}^{(n)}}{1 - p_{d}}\right)$$
$$\frac{\partial \mathcal{L}(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|\mathbf{p})}{\partial p_{d}} = \frac{\sum_{n=1}^{N} x_{d}^{(n)}}{p_{d}} - \frac{\sum_{n=1}^{N} (1 - x_{d}^{(n)})}{1 - p_{d}}$$

and set the derivative to zero to solve for \hat{p}_d :

$$\frac{\sum_{n=1}^{N} x_d^{(n)}}{\hat{p}_d} - \frac{\sum_{n=1}^{N} (1 - x_d^{(n)})}{1 - \hat{p}_d} = 0$$

$$\sum_{n=1}^{N} x_d^{(n)} - \hat{p}_d \sum_{n=1}^{N} x_d^{(n)} - \hat{p}_d \cdot N + \hat{p}_d \sum_{n=1}^{N} x_d^{(n)} = 0$$

$$\hat{p}_d = \frac{1}{N} \sum_{n=1}^{N} x_d^{(n)}$$

Because we assume that each pixel is independent (we are taking the product of D one dimensional Bernoulli distributions), we can express the maximum likelihood for \mathbf{p} in vectorised form as $\hat{\mathbf{p}}^{MLE}$:

$$\hat{\mathbf{p}}^{MLE} = \frac{1}{N} \sum_{n=1}^{N} \mathbf{x}^{(n)}$$

(c) From Bayes' Theorem:

$$P(\mathbf{p}|\{\mathbf{x}^{(n)}\}_{n=1}^{N}) = \frac{P(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|\mathbf{p})P(\mathbf{p})}{P(\{\mathbf{x}^{(n)}\}_{n=1}^{N})}$$

Taking the logarithm:

$$\mathcal{L}(\mathbf{p}|\{\mathbf{x}^{(n)}\}_{n=1}^{N}) = \mathcal{L}(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|\mathbf{p}) + \mathcal{L}(\mathbf{p}) - \mathcal{L}(\{\mathbf{x}^{(n)}\}_{n=1}^{N})$$

Taking the derivative with respect to p_d :

$$\frac{\partial \mathcal{L}(\mathbf{p}|\{\mathbf{x}^{(n)}\}_{n=1}^{N})}{\partial p_{d}} = \frac{\partial \mathcal{L}(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|\mathbf{p})}{\partial p_{d}} + \frac{\partial \mathcal{L}(\mathbf{p})}{\partial p_{d}}$$

where $\frac{\partial \mathcal{L}(\{\mathbf{X}^{(n)}\}_{n=1}^{N})}{\partial p_d} = 0$ because it doesn't depend on p_d .

We know from (b):

$$\frac{\partial \mathcal{L}(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|\mathbf{p})}{\partial p_d} = \frac{\sum_{n=1}^{N} x_d^{(n)}}{p_d} - \frac{\sum_{n=1}^{N} (1 - x_d^{(n)})}{1 - p_d}$$

For the second term $\frac{\partial \mathcal{L}(\mathbf{p})}{\partial p_d}$, we start with $P(\mathbf{p})$, assuming each pixel to have an independent prior:

$$P(\mathbf{p}) = \prod_{d=1}^{D} P(p_d)$$

and assuming a Beta prior on each p_d :

$$P(\mathbf{p}) = \prod_{d=1}^{D} \frac{1}{B(\alpha, \beta)} p_d^{\alpha - 1} (1 - p_d)^{\beta - 1}$$

Taking the logarithm:

$$\mathcal{L}(\mathbf{p}) = \sum_{d=1}^{D} -\log(B(\alpha, \beta)) + (\alpha - 1)\log p_d + (\beta - 1)\log(1 - p_d)$$

Taking the derivative with respect to p_d :

$$\frac{\partial \mathcal{L}(\mathbf{p})}{\partial p_d} = \frac{(\alpha - 1)}{p_d} - \frac{(\beta - 1)}{1 - p_d}$$

Since we are only concerned with p_d , we are only left with a single element of the summation pertaining to p_d .

Combining, we have an expression for $\frac{\partial \mathcal{L}(\mathbf{p}|\{\mathbf{X}^{(n)}\}_{n=1}^{N})}{\partial p_d}$:

$$\frac{\partial \mathcal{L}(\mathbf{p}|\{\mathbf{x}^{(n)}\}_{n=1}^{N})}{\partial p_{d}} = \frac{\sum_{n=1}^{N} x_{d}^{(n)}}{p_{d}} - \frac{\sum_{n=1}^{N} (1 - x_{d}^{(n)})}{1 - p_{d}} + \frac{(\alpha - 1)}{p_{d}} - \frac{(\beta - 1)}{1 - p_{d}}$$
$$\frac{\partial \mathcal{L}(\mathbf{p}|\{\mathbf{x}^{(n)}\}_{n=1}^{N})}{\partial n_{d}} = \frac{(\alpha - 1) + \sum_{n=1}^{N} x_{d}^{(n)}}{n_{d}} - \frac{(\beta - 1) + \sum_{n=1}^{N} (1 - x_{d}^{(n)})}{1 - n_{d}}$$

To find the maximum a posteriori (MAP) estimate $\hat{p_d}$ set $\frac{\partial \mathcal{L}(\mathbf{p}|\{\mathbf{X}^{(n)}\}_{n=1}^N)}{\partial p_d} = 0$ and solve:

$$0 = \frac{(\alpha - 1) + \sum_{n=1}^{N} x_d^{(n)}}{\hat{p_d}} - \frac{(\beta - 1) + \sum_{n=1}^{N} (1 - x_d^{(n)})}{1 - \hat{p_d}}$$

$$0 = (1 - \hat{p_d})(\alpha - 1) + (1 - \hat{p_d}) \left(\sum_{n=1}^{N} x_d^{(n)}\right) - \hat{p_d}(\beta - 1) - \hat{p_d} \left(\sum_{n=1}^{N} (1 - x_d^{(n)})\right)$$

$$0 = (\alpha - \alpha \hat{p_d} + \hat{p_d} - 1) + \left(\sum_{n=1}^{N} x_d^{(n)} - \hat{p_d} \sum_{n=1}^{N} x_d^{(n)}\right) - (\hat{p_d}\beta - \hat{p_d}) - \left(\hat{p_d} \cdot N - \hat{p_d} \sum_{n=1}^{N} x_d^{(n)}\right)$$

Cancelling the $\hat{p}_d \sum_{n=1}^N x_d^{(n)}$ terms:

$$0 = \alpha - \alpha \hat{p_d} + \hat{p_d} - 1 + \sum_{n=1}^{N} x_d^{(n)} - \hat{p_d}\beta + \hat{p_d} - \hat{p_d} \cdot N$$
$$0 = \hat{p_d}(2 - \alpha - \beta - N) + \alpha - 1 + \sum_{n=1}^{N} x_d^{(n)}$$
$$\hat{p_d} = \frac{\alpha - 1 + \sum_{n=1}^{N} x_d^{(n)}}{(N + \alpha + \beta - 2)}$$

Due to independence of our likelihood and priors for each dimension, we can express the maximum a priori for \mathbf{p} in vectorised form as $\hat{\mathbf{p}}^{MAP}$:

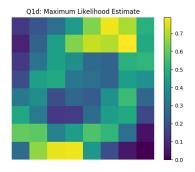
$$\hat{\mathbf{p}}^{MAP} = \frac{\alpha - 1 + \sum_{n=1}^{N} \mathbf{x}^n}{(N + \alpha + \beta - 2)}$$

(d&e) The Python code for MLE and MAP:

```
import matplotlib.pyplot as plt
import numpy as np
3
     {\tt def\_compute\_maximum\_likelihood\_estimate(x: np.ndarray)} \ -\!\!\!> \ np.ndarray:
6
           :param x: numpy array of shape (N, D) :return: MLE estimate """
10
           return np.mean(x, axis=0)
     def _compute_maximum_a_priori_estimate(
    x: np.ndarray, alpha: float, beta: float
14
     ) -> np.ndarray:
           Calculates MAP estimate of images
           :param x: numpy array of shape (N, D)
:param alpha: param of prior distribution
:param beta: param of prior distribution
:return: MAP estimate
"""
19
20
22
23
24
25
           n, = x.shape
           return (alpha - 1 + np.sum(x, axis=0)) / (n + alpha + beta - 2)
     def d(x: np.ndarray, figure_path: str, figure_title: str) -> None:
30
           Produces answers for question 1d :param x: numpy array of shape (N, D)
31
33
34
           :param figure_path: path to store figure
:param figure_title: figure title
35
36
           maximum_likelihood = _compute_maximum_likelihood_estimate(x)
38
           plt.figure()
            plt.imshow(
                 np.reshape(maximum_likelihood, (8, 8)),
41
                 interpolation="None",
42
           plt.colorbar()
           plt.axis("off")
plt.title(figure_title)
44
45
           plt.savefig(figure_path)
49
          x: np.ndarray, alpha: float, beta: float, figure_path: str, figure_title: str
50
51
     ) -> None:
           Produces answers for question 1e:param x: numpy array of shape (N, D):param alpha: param of prior distribution
55
56
           :param beta: param of prior distribution
:param figure_path: path to store figure
:param figure_title: figure title
58
59
           :return:
60
61
           maximum\_a\_priori = \_compute\_maximum\_a\_priori\_estimate(x, alpha, beta)
62
            plt.figure()
            plt.imshow(
64
                 np.reshape(maximum_a_priori, (8, 8)),
65
                 interpolation="None",
           plt.colorbar()
    resis("off")
67
           plt.axis("off")
plt.title(figure_title)
plt.savefig(f"{figure_path}.png")
69
70
71
72
73
74
75
76
77
78
79
            maximum\_likelihood = \_compute\_maximum\_likelihood\_estimate(x)
            plt.figure()
                 {\tt np.reshape(maximum\_a\_priori-maximum\_likelihood,\ (8\,,\ 8))}\,,
                 interpolation="None",
           plt.colorbar()
           plt.axis("off")
plt.title(f"MAP vs MLE")
80
            plt.savefig(f"{figure_path}-mle-vs-map.png")
```

src/solutions/q1.py

Displaying the learned parameters:



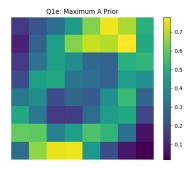


Figure 1: ML parameters

Figure 2: MAP parameters

Comparing the equations:

$$\hat{\mathbf{p}}^{MLE} = \frac{1}{N} \sum_{n=1}^{N} \mathbf{x}^{(n)}$$

and

$$\hat{\mathbf{p}}^{MAP} = \frac{\alpha - 1 + \sum_{n=1}^{N} \mathbf{x}^n}{(N + \alpha + \beta - 2)}$$

As the number of data points increases, $\hat{\mathbf{p}}^{MAP}$ approaches $\frac{1}{N}\sum_{n=1}^{N}\mathbf{x}^{(n)}$, the $\hat{\mathbf{p}}^{MLE}$. This makes sense because as our data set gets bigger, the effect of the prior diminishes. However, if a specific pixel in all of the images of our data set are white or all black, the MLE for that pixel would either be 1 or 0. This may not be representative of our intuitions about images, as there should be some non-zero probability of a pixel being black or white. By introducing an appropriate prior we can ensure that the probability of that pixel will never be exactly zero or one. In our case, with a Beta(3,3) prior on each pixel, our parameter values are biased to be closer to 0.5 and to never be at the extremities 0 and 1. We can see this in Figure 2 where the range of our parameters is smaller than the range of Figure 1 and doesn't include zero. Figure 3 visualises $\hat{\mathbf{p}}^{MAP} - \hat{\mathbf{p}}^{MLE}$ and we can see that for likelihoods greater than 0.5 in the MLE, the MAP has a lower value and for likelihoods less than 0.5, the MAP has a higher value, confirming our intuitions.

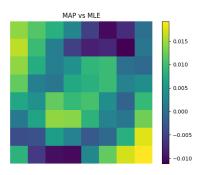


Figure 3: $\hat{\mathbf{p}}^{MAP} - \hat{\mathbf{p}}^{MLE}$

Priors can also help ensure numerical stability during calculations. The logarithm of zero is negative infinity, so having if the MLE is zero it can be problematic for log-likelihood calculations whereas MAP can ensure non-zero probabilities. Interestingly, when $\alpha = \beta = 1$, $\hat{\mathbf{p}}^{MLE} = \hat{\mathbf{p}}^{MAP}$. This is when the prior is a uniform distribution and so there is uniform bias on the location of \mathbf{p} and we recover the MLE.

On the other hand, a mis-specified prior can be problematic, as the estimated parameters might be skewed by the prior and not properly represent the underlying data generating process, this can result in parameter estimates that are 'worse' than using the MLE if our data set is limited in size.

Question 2

When all D components are generated from a Bernoulli distribution with $p_d = 0.5$, we have the likelihood function for model M_1 :

$$P(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|\mathbf{p}^{(1)} = [0.5, 0.5, ..., 0.5]^{T}, M_{1}) = \prod_{n=1}^{N} \prod_{d=1}^{D} (0.5)^{x_{d}^{(n)}} (0.5)^{1-x_{d}^{(n)}}$$

Knowing that either $x_d^{(n)}$ or $1 - x_d^{(n)}$ will be 1 and the other zero, we can simplify $(0.5)^{x_d^{(n)}}(1 - 0.5)^{1-x_d^{(n)}}$ to 0.5:

$$P(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|\mathbf{p}^{(1)} = [0.5, 0.5, ..., 0.5]^{T}, M_{1}) = \prod_{n=1}^{N} \prod_{d=1}^{D} (0.5)$$

$$P(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|\mathbf{p}^{(1)} = [0.5, 0.5, ..., 0.5]^{T}, M_1) = 0.5^{N \cdot D}$$

When all D components are generated from Bernoulli distributions with unknown, but identical, p_d , we have the likelihood function for model M_2 :

$$P(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|\mathbf{p}^{(2)} = [p_d, p_d, ..., p_d]^T, M_2) = \prod_{n=1}^{N} \prod_{d'=1}^{D} p_d^{x_{d'}^{(n)}} (1 - p_d)^{1 - x_{d'}^{(n)}}$$

When each component is Bernoulli distributed with separate, unknown p_d , we have the likelihood function for model M_3 :

$$P(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|\mathbf{p}^{(3)} = [p_1, p_2, ..., p_D]^T, M_3) = \prod_{n=1}^{N} \prod_{d=1}^{D} p_d^{x_d^{(n)}} (1 - p_d)^{1 - x_d^{(n)}}$$

For each model M_i , we can marginalise out $\mathbf{p}^{(i)}$ to get $P(\{\mathbf{x}^{(n)}\}_{n=1}^N | M_i)$:

$$P(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|M_i) = \int_0^1 \dots \int_0^1 P(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|\mathbf{p}^{(i)}, M_i) P(\mathbf{p}^{(i)}|M_i) dp_1 \dots dp_D$$

Given that the prior of any unknown probabilities is uniform, i.e. $P(\mathbf{p}^{(i)}|M_i) = 1$. We can simplify:

$$P(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|M_i) = \int_0^1 \dots \int_0^1 P(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|\mathbf{p}^{(i)}, M_i) dp_1 \dots dp_D$$

For M_1 :

$$P(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|M_1) = \int_0^1 \dots \int_0^1 0.5^{N \cdot D} d\theta_1 \dots d\theta_D$$

We can remove the integrals:

$$P(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|M_1) = 0.5^{N \cdot D}$$

For M_2 , we have that all pixels share some probability p_d so we only need to integrate over a single variable p_d :

$$P(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|M_2) = \int_0^1 \prod_{n=1}^{N} \prod_{d'=1}^{D} p_d^{x_{d'}^{(n)}} (1 - p_d)^{1 - x_{d'}^{(n)}} dp_d$$

Changing the products to sums:

$$P(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|M_2) = \int_0^1 p_d^{\sum_{n=1}^{N} \sum_{d'=1}^{D} x_{d'}^{(n)}} (1 - p_d)^{\sum_{n=1}^{N} \sum_{d'=1}^{D} 1 - x_{d'}^{(n)}} dp_d$$

Rewriting:

$$P(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|M_2) = \int_0^1 (p_d)^K (1 - p_{d=1})^{N \cdot D - K} dp_d$$

where $K = \sum_{n=1}^{N} \sum_{d'=1}^{D} x_{d'}^{(n)}$.

This integral is the beta function:

$$P(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|M_2) = \frac{K!(N \cdot D - K)!}{(N \cdot D + 1)!}$$

For M_3 , we need an integral for each p_d :

$$P(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|M_3) = \int_0^1 \dots \int_0^1 \prod_{n=1}^N \prod_{d=1}^D p_d^{x_d^{(n)}} (1 - p_d)^{1 - x_d^{(n)}} dp_1 \dots dp_D$$

We can separate the integrals to only contain the relevant p_d :

$$P(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|M_3) = \prod_{d=1}^{D} \left(\int_0^1 \prod_{n=1}^{N} p_d^{x_d^{(n)}} (1 - p_d)^{1 - x_d^{(n)}} dp_d \right)$$

Changing the products to sums:

$$P(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|M_3) = \prod_{d=1}^{D} \left(\int_0^1 p_d^{\sum_{n=1}^{N} x_d^{(n)}} (1 - p_d)^{\sum_{n=1}^{N} 1 - x_d^{(n)}} dp_d \right)$$

In this case, we have the product of integrals where each evaluates to a beta function:

$$P(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|M_3) = \prod_{d=1}^{D} \frac{K_d!(N-K_d)!}{(N+1)!}$$

where $K_d = \sum_{n=1}^{N} x_d^{(n)}$.

The posterior probability of a model M_i can be expressed:

$$P(M_i | \{\mathbf{x}^{(n)}\}_{n=1}^N) = \frac{P(\{\mathbf{x}^{(n)}\}_{n=1}^N | M_i) P(M_i)}{P(\{\mathbf{x}^{(n)}\}_{n=1}^N)}$$

We only have three models, so in this case the normalisation $P(\{\mathbf{x}^{(n)}\}_{n=1}^N)$ can be expressed as a sum:

$$P(M_i|\{\mathbf{x}^{(n)}\}_{n=1}^N) = \frac{P(\{\mathbf{x}^{(n)}\}_{n=1}^N|M_i)P(M_i)}{\sum_{i\in\{1,2,3\}}P(\{\mathbf{x}^{(n)}\}_{n=1}^N|M_i)P(M_i)}$$

Given that $P(M_i) = \frac{1}{3}$ for all $i \in \{1, 2, 3\}$:

$$P(M_i | \{\mathbf{x}^{(n)}\}_{n=1}^N) = \frac{P(\{\mathbf{x}^{(n)}\}_{n=1}^N | M_i)}{\sum_{i \in \{1,2,3\}} P(\{\mathbf{x}^{(n)}\}_{n=1}^N | M_i)}$$

Calculating the posterior probabilities of each of the three models having generated the data in binarydigits.txt using Python, we can show the values in the Table 1.

i	$P(M_i \{\mathbf{x}^{(n)}\}_{n=1}^N)$
1	1E-1924
2	1E-1858
3	1-(1E-1924)-(1E-1858)

Table 1: Posterior Probabilities

We can see that for models specified to have the same parameter value for all pixels, like M_1 , are very unlikely with the given data set. This makes sense because it is specifying models where the image is essentially a uniform shade, which is not reflective of our digit images. Moreover, M_1 specifies a specific value of 0.5 for all the parameters whereas M_2 specifies any value for all the parameters as long as it's the same. So the model M_1 is just one possible model specified in M_2 and we can see this reflected in our probabilities when $P(M_2|\{\mathbf{x}^{(n)}\}_{n=1}^N) > P(M_1|\{\mathbf{x}^{(n)}\}_{n=1}^N)$.

The Python code for calculating the posterior probabilities of the three models:

```
import pandas as pd
      from scipy.special import betaln, logsumexp
 6
7
8
      \begin{array}{lll} \textbf{def} & \texttt{-log-p-d-given-m1} \, (\, x \colon \; \texttt{np.ndarray} \,) \; -\!\!\!> \; \textbf{float} : \end{array}
            Calculates log likelihood of model 1: param x: numpy array of shape (N, D): return: log likelihood """
10
11
            n, d = x.shape
13
14
            return n * d * np.log(0.5)
16
17
      def _log_p_d_given_m2(x: np.ndarray):
18
19
            Calculates log likelihood of model 2
            :param x: numpy array of shape (N, D)
:return: log likelihood
"""
20
21
22
            \begin{array}{ll} n\,,\;\;d=\,x\,.\,shape \\ k\,=\,np\,.\,sum(\,x\,)\,.\,astype\,(\,in\,t\,) \\ return\;\;betaln\,(\,k\,+\,1\,,\;n\,*\,d\,-\,k\,+\,1\,) \end{array}
23
24
25
26
27
28
      def _log_p_d_given_m3(x: np.ndarray):
            :param x: numpy array of shape (N, D) return: log likelihood
29
30
31
33
34
            n, _ = x.shape
k_d = np.sum(x, axis=0).astype(int)
            return logsumexp(betaln(k_d + 1, n - k_d + 1))
36
38
      def _log_p_model_given_data(x) -> np.ndarray:
            Calculates posterior log likelihood of models given image data
40
            :param x: numpy array of shape (N, D)
:return: posterior log likelihood
"""
41
42
44
            log_p_d_given_m = np.array(
45
                         -\log_{p}_{d}_{given_{m}}1(x),
                         log_p_d_given_m2(x),
log_p_d_given_m3(x),
47
48
49
50
51
            log_p_m_given_data = log_p_d_given_m - logsumexp(log_p_d_given_m)
            return log_p_m_given_data
55
      \begin{array}{lll} \texttt{def} & \texttt{c(x: np.ndarray, table\_path: str)} \ -\!\!\!> \ None: \end{array}
56
            Produces answers for question 2c
            :param x: numpy array of shape (N,\ D) :param table_path: path to store table posterior likelihoods
58
59
61
62
            log_p_m_given_data = _log_p_model_given_data(x)
            df = pd.DataFrame(
                   data=np.array(
                              np.arange(len(log-p-m-given_data)).astype(int) + 1,
[f"1E{int(x/np.log(10))}" for x in log-p-m-given_data[:-1]]
+ [
65
67
                                     f"1-\{'-'.join([f'(1E\{int(x/np.log(10))\})'|for x in log_p_m_given_data[:-1]])\}"
70
71
72
73
74
                  ).T.
                   columns=["Model", "P(M_i |D)"],
            df.set_index("Model", inplace=True)
df.to_csv(table_path)
```

src/solutions/q2.py

Question 3

(a) The likelihood for a model consisting of a mixture of K multivariate Bernoulli distributions can be expressed as the product across N data points:

$$P(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|\theta) = \prod_{i=1}^{N} P(\mathbf{x}^{(n)}|\theta)$$

where $\{\mathbf{x}^{(n)}\}_{n=1}^{N}$ is our data set with $\mathbf{x}^{(n)} \in \mathbb{R}^{D \times 1}$ and $\theta = \{\pi, \mathbf{P}\}$ are our parameters, $\pi = [\pi_1, ..., \pi_K] \in \mathbb{R}^{K \times 1}$ our K mixing proportions $(0 \le \pi_k \le 1; \sum_k \pi_k = 1)$ and $\mathbf{P} \in \mathbb{R}^{D \times K}$ the K Bernoulli parameter vectors with elements p_{kd} denoting the probability that pixel d takes value 1 given mixture component k. We also assume the images are iid and that the pixels are independent of each other within each component distribution.

For each $P(\mathbf{x}^{(n)}|\theta)$:

$$P(\mathbf{x}^{(n)}|\theta) = \sum_{k=1}^{K} \pi_k \prod_{d=1}^{D} (p_{kd})^{x_d^{(n)}} (1 - p_{kd})^{1 - x_d^{(n)}}$$

The log-likelihood $\mathcal{L}(\mathbf{x}^{(n)}|\theta)$ can be expressed in vector form:

$$\mathcal{L}(\mathbf{x}^{(n)}|\theta) = \log \sum_{k=1}^{K} \pi_k \exp\left((\mathbf{x}^{(n)})^T \log(\mathbf{P}_k) + (1 - \mathbf{x}^{(n)})^T \log(1 - \mathbf{P}_k)\right)$$

where \mathbf{P}_k is the k^{th} column of \mathbf{P} . This can can be further vectorised using Python scipy's logsumexp operation.

Moreover, the log-likelihood $\mathcal{L}(\{\mathbf{x}^{(n)}\}_{n=1}^N | \theta)$ can be expressed:

$$\mathcal{L}(\{\mathbf{x}^{(n)}\}_{n=1}^{N}|\theta) = \sum_{k=1}^{N} \left(\log \sum_{k=1}^{K} \pi_k \exp\left((\mathbf{x}^{(n)})^T \log(\mathbf{P}_k) + (1-\mathbf{x}^{(n)})^T \log(1-\mathbf{P}_k)\right)\right)$$

(b) We know that:

$$P(A|B) \propto P(B|A)P(A)$$

Thus,

$$P(s^{(n)} = k | \mathbf{x}^{(n)}, \pi, \mathbf{P}) \propto P(\mathbf{x}^{(n)} | s^{(n)} = k, \pi, \mathbf{P}) P(s^{(n)} = k | \pi, \mathbf{P})$$

where $s^{(n)} \in \{1, ..., K\}$ a discrete hidden variable with $P(s^{(n)} = k | \pi) = \pi_k$. Note that $P(s^{(n)} = k | \pi, \mathbf{P}) = P(s^{(n)} = k | \pi)$ as $s^{(n)} = k$ isn't dependent on \mathbf{P} .

Let \tilde{r}_{nk} be the unnormalised responsibility $P(\mathbf{x}^{(n)}|s^{(n)}=k,\pi,\mathbf{P})P(s^{(n)}=k|\pi,\mathbf{P})$. Using the mixture for component k, π_k and the likelihood function of component k:

$$\tilde{r}_{nk} = \pi_k \prod_{l=1}^{D} (p_{kd})^{x_d^{(n)}} (1 - p_{kd})^{1 - x_d^{(n)}}$$

Normalising across the components:

$$r_{nk} = \frac{\tilde{r}_{nk}}{\sum_{j=1}^{K} \tilde{r}_{nj}}$$

we have calculated $P(s^{(n)} = k | \mathbf{x}^{(n)}, \pi, \mathbf{P})$ for the E step of an EM algorithm. Moreover,

$$\log \tilde{r}_{nk} = \log \pi_k + \sum_{d=1}^{D} \left(x_d^{(n)} \log(p_{kd}) + (1 - x_d^{(n)}) \log(1 - \exp(\log(p_{kd}))) \right)$$

and

$$\log r_{nk} = \log \tilde{r}_{nk} - \log \sum_{i=1}^{K} \exp(\log \tilde{r}_{nj})$$

which can be vectorised as $\log \mathbf{r}$ calculated with $\log \pi$ and $\log \mathbf{P}$ using Python scipy's logsum exp operation.

(c) We know that the expectation log joint can be expressed:

$$\left\langle \sum_{n} \log P(\mathbf{x}^{(n)}, s^{(n)} | \pi, \mathbf{P}) \right\rangle_{q(\{s^{(n)}\})} = \sum_{n=1}^{N} q(s^{(n)}) \log P(\mathbf{x}^{(n)}, s^{(n)} | \pi, \mathbf{P})$$

Let this quantity be E. For each term of the summation in E:

$$q(s^{(n)}) = \mathbf{r}_n^T$$

and

$$\log P(\mathbf{x}^{(n)}, s^{(n)} | \pi, \mathbf{P}) = \log[P(\mathbf{x}^{(n)} | s^{(n)}, \pi, \mathbf{P})P(s^{(n)} | \pi, \mathbf{P})]$$

which is the vectorised version of $\log \tilde{r}_{nk}$ from part (b) so:

$$\log P(\mathbf{x}^{(n)}, s^{(n)} | \pi, \mathbf{P}) = \log(\pi) + \log(\mathbf{P})^T \mathbf{x}^{(n)} + \log(1 - \mathbf{P})^T (1 - \mathbf{x}^{(n)})$$

Combining:

$$E = \sum_{n} \mathbf{r}_{n}^{T} [\log(\pi) + \log(\mathbf{P})^{T} \mathbf{x}^{(n)} + \log(1 - \mathbf{P})^{T} (1 - \mathbf{x}^{(n)})]$$

To maximise with respect to π and \mathbf{P} for the M step, we want to take the derivative, set to zero, and solve for $\hat{\pi}$ and $\hat{\mathbf{P}}$.

For the k^{th} element of π :

$$\frac{\partial E}{\partial \pi_k} = \sum_{n} r_{nk} \frac{1}{\pi_k}$$

We can calculate the maximiser with:

$$\frac{\partial E}{\partial \pi_k} + \lambda = 0$$

where λ is a Lagrange multiplier ensuring that the mixing proportions sum to unity.

Thus,

$$\hat{\pi}_k = \frac{\sum_n r_{nk}}{N}$$

For the dk^{th} element of **P**:

$$\frac{\partial E}{\partial \mathbf{P}_{dk}} = \sum_{n} r_{nk} \frac{\partial}{\partial \mathbf{P}_{dk}} [x_d^{(n)} \log \mathbf{P}_{dk} + (1 - x_d^{(n)}) \log(1 - \mathbf{P}_{dk})]$$

Simplifying:

$$\frac{\partial E}{\partial \mathbf{P}_{dk}} = \sum_{n} r_{nk} \left(\frac{x_d^{(n)}}{\mathbf{P}_{dk}} - \frac{1 - x_d^{(n)}}{1 - \mathbf{P}_{dk}} \right)$$

Setting the derivative to zero:

$$\frac{\sum_{n} x_{d}^{(n)} r_{nk}}{\hat{\mathbf{P}}_{dk}} - \frac{\sum_{n} r_{nk} - \sum_{n} x_{d}^{(n)} r_{nk}}{1 - \hat{\mathbf{P}}_{dk}} = 0$$

Solving for $\hat{\mathbf{P}}_{dk}$:

$$\hat{\mathbf{P}}_{dk} \sum_{n} r_{nk} - \hat{\mathbf{P}}_{dk} \sum_{n} x_d^{(n)} r_{nk} = \sum_{n} x_d^{(n)} r_{nk} - \hat{\mathbf{P}}_{dk} \sum_{n} x_d^{(n)} r_{nk}$$

Thus,

$$\hat{\mathbf{P}}_{dk} = \frac{\sum_{n} x_d^{(n)} r_{nk}}{\sum_{n} r_{nk}}$$

We have the maximizing parameters for the expected log-joint

$$\arg \max_{\pi, \mathbf{P}} \left\langle \sum_{n} \log P(\mathbf{x}^{(n)}, s^{(n)} | \pi, \mathbf{P}) \right\rangle_{q(\{s^{(n)}\})}$$

thus obtaining an iterative update for the parameters π and **P** in the M-step of EM.

For numerical stability, we can compute the maximisation step for the MAP of \mathbf{P} , by solving for $\hat{\mathbf{P}}_{dk}^{MAP}$ with:

$$\frac{\partial E'}{\partial \mathbf{P}_{dk}} = 0$$

where

$$E' = \sum_{n=1}^{N} q(s^{(n)}) \log P(\mathbf{P}|\pi, \mathbf{x}^{(n)}, s^{(n)})$$

and from Bayes':

$$\log P(\mathbf{P}|\pi, \mathbf{x}^{(n)}, s^{(n)}) = \log P(\mathbf{x}^{(n)}, s^{(n)}|\pi, \mathbf{P}) + \log P(\mathbf{P}) - \log P(\mathbf{x}^{(n)}, s^{(n)}|\pi)$$

Assuming an independent Beta prior on each pixel of each component:

$$\log P(\mathbf{P}) = \sum_{k=1}^{K} \sum_{d=1}^{D} -\log(B(\alpha, \beta)) + (\alpha - 1)\log \mathbf{P}_{dk} + (\beta - 1)\log(1 - \mathbf{P}_{dk})$$

and

$$\frac{\partial \log P(\mathbf{P})}{\partial \mathbf{P}_{dk}} = \frac{(\alpha - 1)}{\mathbf{P}_{dk}} - \frac{(\beta - 1)}{1 - \mathbf{P}_{dk}}$$

Thus, the derivative can be expressed as:

$$\frac{\partial E'}{\partial \mathbf{P}_{dk}} = \sum_{n} \left(r_{nk} \left(\frac{\partial \log P(\mathbf{x}^{(n)}, s^{(n)} | \pi, \mathbf{P})}{\partial \mathbf{P}_{dk}} + \frac{\partial \log P(\mathbf{P})}{\partial \mathbf{P}_{dk}} \right) \right)$$

Substituting the appropriate expressions:

$$\frac{\partial E'}{\partial \mathbf{P}_{dk}} = \sum_{n} \left(r_{nk} \left(\frac{x_d^{(n)}}{\mathbf{P}_{dk}} - \frac{1 - x_d^{(n)}}{1 - \mathbf{P}_{dk}} + \frac{(\alpha - 1)}{\mathbf{P}_{dk}} - \frac{(\beta - 1)}{1 - \mathbf{P}_{dk}} \right) \right)$$

Simplifying:

$$\frac{\partial E'}{\partial \mathbf{P}_{dk}} = \frac{\sum_{n} r_{nk} (\alpha - 1 + x_d^{(n)})}{\mathbf{P}_{dk}} - \frac{\sum_{n} r_{nk} (\beta - x_d^{(n)})}{1 - \mathbf{P}_{dk}}$$

Setting $\frac{\partial E'}{\partial \mathbf{P}_{dk}} = 0$ we can calculate $\hat{\mathbf{P}}_{dk}^{MAP}$:

$$\sum_{n} r_{nk}(\alpha - 1 + x_d^{(n)}) - \hat{\mathbf{P}}_{dk} \sum_{n} r_{nk}(\alpha - 1 + x_d^{(n)}) = \hat{\mathbf{P}}_{dk} \sum_{n} r_{nk}(\beta - x_d^{(n)})$$

$$\hat{\mathbf{P}}_{dk}^{MAP} = \frac{\sum_{n} r_{nk} (x_d^{(n)} + \alpha - 1)}{(\alpha + \beta - 1)(\sum_{n} r_{nk})}$$

As a sense check, we can see when setting $\alpha = 1$ and $\beta = 1$ we recover $\hat{\mathbf{P}}_{dk}^{MLE}$ as we would expect. For the following parts, a very weak Beta(1+1e-5, 1+1e-5) prior was used.

(d) Plotting the unnormalised posterior likelihood as a function of the iteration number for different k values:

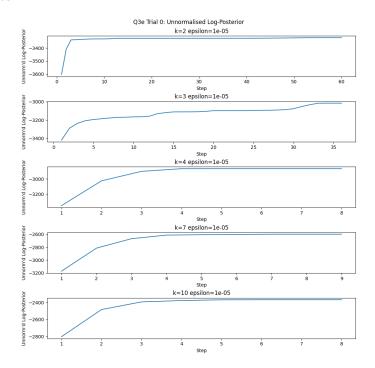


Figure 4: Unnormalised Log Posterior vs Iteration Number

where *epsilon* is the stopping condition for when the unnormalised log posterior converges sufficiently. Note that the normalisation constant for the log posterior $\log P(\mathbf{x}^{(n)}, s^{(n)}|\pi)$ is intractable and so only the unnormalised portion $\log P(\mathbf{x}^{(n)}, s^{(n)}|\pi, \mathbf{P}) + \log P(\mathbf{P})$ was computed and reported.

Displaying the parameters found for $K \in \{2, 3, 4, 7, 10\}$:

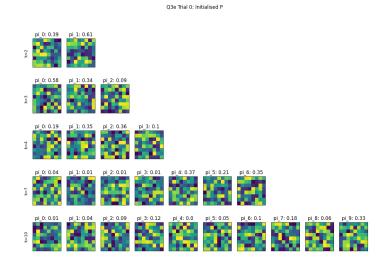


Figure 5: Randomly initialised parameters

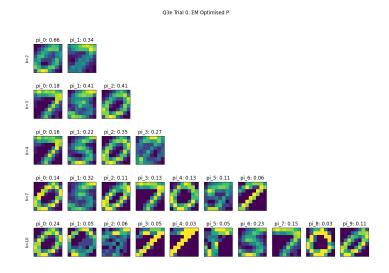


Figure 6: EM optimised parameters

The Python code for the EM algorithm:

```
from dataclasses import dataclass
from typing import List, Tuple
 3
      import matplotlib.pyplot as plt
      import numpy as np
import pandas as pd
from scipy.special import betaln, logsumexp
      from sklearn.manifold import TSNE
10
      from src.constants import DEFAULT_SEED
13
14
      @\,d\,a\,t\,a\,c\,l\,a\,s\,s
      class Theta:
           Data class containing the model parameters log_pi: the logarithm of the mixing proportions (1, k) log_p_matrix: the logarithm of the probability where the (i,j)th element is the probability that pixel j takes value 1 under mixture component i (d, k)
16
17
19
20
22
           log_pi: np.ndarray
log_p_matrix: np.ndarray
24
25
            @property
           def pi(self) -> np.ndarray:
26
27
                 Calculates the mixing proportions :return: vector of mixing proportions (1, k)
30
31
                 return np.exp(self.log_pi)
33
34
           @property
def p_matrix(self) -> np.ndarray:
35
                 Calculates the Bernoulli parameters : return: matrix Bernoulli parameters (d, k)
36
38
                 d, k = self.log_p_matrix.shape
                  \begin{array}{ll} image\_dimension = int (np.sqrt(d)) \\ return \ np.exp(self.log\_p\_matrix).reshape(image\_dimension, image\_dimension, -1) \\ \end{array} 
42
44
            def log_one_minus_p_matrix(self) -> np.ndarray:
45
                 Compute log(1-P) where P=exp(log_p_matrix)
                 :return: an array of the same shape as log_p_matrix\ (d, k)
49
                 log_of_one = np.zeros(self.log_p_matrix.shape)
                 50
51
52
53
54
                 return np.array(logsumexp(stacked_sum, b=weights, axis=0))
55
56
57
            def log_pi_repeated(self, n: int) -> np.ndarray:
                 Repeats the \log_{-p}i vector n times along axis 0
                 :param n: number of repetitions
:return: an array of shape (n, k)
58
59
60
61
                 \begin{array}{lll} \textbf{return} & \texttt{np.repeat} \, (\, \, \texttt{self.log.pi} \, \, , \, \, \, \texttt{n} \, , \, \, \, \texttt{axis} \! = \! 0) \end{array}
62
63
      \label{eq:def_init_params} \begin{array}{lll} def & \verb"init_params" (k: int), & d: int) & -> & Theta: \end{array}
64
65
66
           Random initialisation of theta parameters (log_pi and log_p_matrix)
            :param k: Number of components
:param d: Image dimension (number of pixels in a single image)
:return: theta: the parameters of the model
67
70
71
72
73
74
                 log_pi=np.log(np.random.dirichlet(np.ones(k), size=1)), log_p=matrix=np.log(np.random.uniform(low=0, high=1, size=(d, k))),
      def _compute_log_component_p_x_i_given_theta(x: np.ndarray, theta: Theta) -> np.ndarray:
            Compute the unweighted probability of each mixing component for each image
            :param x: the image data (n, d) :param theta: the parameters of the model :return: an array of the unweighted probabilities (n, k)
80
81
83
84
            return x @ theta.log_p_matrix + (1 - x) @ theta.log_one_minus_p_matrix
86
      def _compute_log_p_x_i_given_theta(x: np.ndarray, theta: Theta) -> np.ndarray:
            Computes the log likelihood of each image in the dataset x
89
            :param x: the image data (n, d) :param theta: the parameters of the model
90
91
            return: log_p_x_i_given_theta: a log likelihood array containing the log likelihood of each image (n
92
03
```

```
94
                        n, = x.shape
                        log_component_probabilities = _compute_log_component_p_x_i_given_theta(
 95
                               x, theta
# (n, k)
 96
  97
 98
                        return np.array(
                                 logsumexp(
 99
100
                                            log_component_probabilities
                                                   theta.log_pi_repeated(n), # scale each component by component probability
                                             axis=1.
                       )
107
             \label{log_likelihood} \begin{array}{ll} def & \verb|compute_log_likelihood(x: np.ndarray, theta: Theta) -> float: \end{array}
108
109
                        Computes the log likelihood of all images in the dataset x
                        :param x: the image data (n, d) :param theta: the parameters of the model
                        return: log_p_x_given_theta: the log likelihood array across all images
                        return np.sum(_compute_log_p_x_i_given_theta(x, theta)).item()
114
             def _compute_log_prior(
                       theta: Theta, alpha_parameter: float, beta_parameter: float
118
             ) -> float:
119
                       Compute the prior \log probability of the P matrix under a Beta prior :param theta: the parameters of the model
                        :param alpha_parameter: alpha parameter of the beta prior
:param beta_parameter: beta parameter of the beta prior
:return: log_p_of_p_matrix
124
126
                        return np.sum(
                                -betaln(alpha_parameter, beta_parameter)
+ (alpha_parameter - 1) * theta.log_p_matrix
+ (beta_parameter - 1) * theta.log_one_minus_p_matrix
130
                        ).item()
132
            x: np.ndarray, theta: Theta, alpha_parameter: float, beta_parameter: float) -> float:
136
137
                       Compute the unnormalised posterior log probability of the P matrix :param x: the image data (n,\ d) :param theta: the parameters of the model
138
139
140
                        :param alpha-parameter: alpha parameter of the beta prior
:param beta-parameter: beta parameter of the beta prior
141
                        :return: log_p_of_p_matrix
143
144
                        log_likelihood = _compute_log_likelihood(x, theta)
                        log_prior = _compute_log_prior(theta, alpha_parameter, beta_parameter)
return log_likelihood + log_prior
146
147
149
             def _compute_log_e_step(x: np.ndarray, theta: Theta) -> np.ndarray:
                       Compute the e step of expectation maximisation :param x: the image data (n, d) :param theta: the parameters of the model :return: an array of the log responsibilities of k mixture components for each image (n, k) """
154
                       \label{log_runnormalised} \begin{array}{l} \text{log_rr_unnormalised} = \text{\_compute\_log\_component\_p\_x\_i\_given\_theta}(x, \text{ theta}) \\ \text{log\_rr_normaliser} = \text{logsumexp}(\text{log\_r_unnormalised}, \text{ axis=1}) \\ \text{log\_responsibility} = \text{log\_rr_unnormalised} - \text{log\_rr_normaliser}[:, \text{ np.newaxis}] \\ \text{return log\_responsibility} \end{array}
157
160
161
             def _compute_log_pi_hat(log_responsibility: np.ndarray) -> np.ndarray:
164
165
                       (n, k)
                        :return: an array of the maximised log mixing proportions (1, k)
167
168
                        n, _ = log_responsibility.shape
                        return (logsumexp(log_responsibility, axis=0) - np.log(n)).reshape(1, -1)
170
\frac{173}{174}
             def _compute_log_p_matrix_hat(
                        x: np.ndarray,
                        log_responsibility: np.ndarray,
\begin{array}{c} 176 \\ 177 \end{array}
                       alpha_parameter: float, beta_parameter: float,
178
             ) -> np.ndarray:
180
                        Compute the log of the maximised pixel probabilities
                        compute the log t the image data (n, d) :param x: the image data (n, d) :param \log_{-1} \exp_{-1} \exp_{-1}
181
182
                        :param alpha_parameter: alpha parameter of the beta prior
:param beta_parameter: beta parameter of the beta prior
:return: an array of the maximised pixel probabilities for each component (d, k)
"""
183
184
186
                      n, d = x.shape
187
```

```
188
            _, k = log_responsibility.shape
189
            190
191
192
                  log_responsibility[:, np.newaxis, :], d, axis=1
            ) # (n, d, k)
194
195
             log\_p\_matrix\_unnormalised\_posterior \, = \, logsumexp\,(
            log_responsibility_repeated , b=(x_repeated + alpha_parameter - 1) , axis=0 ) # (d, k)
196
198
            \label{log_pmatrix_normaliser_posterior} \begin{array}{l} log\_nmatrix\_normaliser\_posterior = logsumexp(\\ log\_responsibility\_repeated \;, \; b=(alpha\_parameter \; + \; beta\_parameter \; - \; 1) \;, \; axis=0 \end{array}
199
200
            ) # (d, k)
201
202
203
             log_p_matrix_normalised_posterior = (
204
                  log_p_matrix_unnormalised_posterior - log_p_matrix_normaliser_posterior
             ) # (d,
205
                         k)
206
             return log_p_matrix_normalised_posterior
207
208
209
      def _compute_log_m_step(
210
            x: np.ndarray
             log_responsibility: np.ndarray,
                                     float,
            alpha_parameter: float
beta_parameter: float,
213
       ) -> Theta:
216
            Compute the m step of expectation maximisation :param x: the image data (n, d) \,
218
             :param log_responsibility: an array of the log responsibilities of k mixture components for each image
             (n, k)
219
             :param alpha-parameter: alpha parameter of the beta prior
            :param beta-parameter: beta parameter of the beta prior :return: thetas optimised after maximisation step
220
222
            return Theta(
                  log_pi=_compute_log_pi_hat(log_responsibility),
                  log-p-matrix=_compute_log-p-matrix-hat(
    x, log_responsibility, alpha_parameter, beta_parameter
226
228
229
230
      {\tt def\_run\_expectation\_maximisation} \, (
232
            x: np.ndarray,
             theta: Theta,
233
            alpha_parameter: float,
beta_parameter: float,
      236
237
            Run the expectation maximisation algorithm
240
             :param x:
                          the image data (n, d)
            :param theta: initial theta parameters
:param alpha_parameter: alpha parameter of the beta prior
242
243
244
             :param beta_parameter: beta parameter of the beta prior
            :param max_number_of_steps: the maximum number of steps to run the algorithm
:param epsilon: the minimum required change in log posterior, otherwise the algorithm stops early
:return: a tuple containing the optimised thetas, the log responsibilities,

and the log log_posteriors at each step of the algorithm
245
247
248
            log_responsibility = None
250
            log_responsibility = None
log_posteriors = []
for _ in range(max_number_of_steps):
    log_responsibility = _compute_log_e_step(x, theta)
    theta = _compute_log_m_step(
253
254
                      x, log_responsibility, alpha_parameter, beta_parameter
256
258
                  \log_{-posteriors}.append(
                       _compute_unnormalised_log_posterior_likelihood(
259
260
                           x, theta, alpha-parameter, beta-parameter
261
                  )
262
                  # check for early stopping
if len(log-posteriors) > 1:
   if (log-posteriors[-1] - log-posteriors[-2]) < epsilon:</pre>
264
265
266
267
                             break
268
             return theta, log_responsibility, log_posteriors
269
      def _visualise_p_matrix(
    thetas: List[Theta], ks: List[int], figure_title: str, figure_path: str
      ) -> None:
273
            Visualises the P matrix for different thetas and ks:param thetas: list of Theta instances:param ks: list of k values used for each Theta:param figure_title: name of figure
:param figure_path: path to store figure
276
278
279
            :return:
281
           n = len(ks)
282
```

```
m = np.max(ks)
283
              fig = plt.figure()
fig.set_figwidth(15)
fig.set_figheight(10)
284
285
286
              for i, k in enumerate(ks):
for j in range(k):
ax = plt.subplot(n, m, m * i + j + 1)
287
288
289
                           ax.imshow(
thetas[i].p_matrix[:, :, j],
interpolation="None",
200
291
203
                           ax.tick_params (
294
                                 axis="x",
which="both",
295
296
                                  bottom=False ,
298
                                  top=False,
                           ax.tick_params(
300
                                 axis="y",
which="both",
301
302
                                  left=False,
303
304
                                  right=False,
305
                           ax.xaxis.set_ticklabels([])
ax.yaxis.set_ticklabels([])
ax.set_title(f"pi_{j}: {np.round(thetas[i].pi[0, j], 2)}")
306
307
308
309
                           if j == 0:
              ax.set_ylabel(f"{k=}")
fig.suptitle(figure_title)
310
311
              plt.savefig(figure_path)
313
314
315
       def _visualise_responsibility_clusters (
              log_responsibilities: List[np.ndarray],
ks: List[int],
figure_title: str,
316
318
       figure-path: str ,
) -> None:
319
320
321
              Visualise responsibility vectors of images using TSNE for different k values
322
              :param log_responsibilities: list of log responsibilities for different ks:param ks: list of k values used for each Theta
:param figure_title: name of figure
324
325
              :param figure_path: path to store figure
              :return:
327
328
              n = len(ks)
fig = plt.figure()
fig.set_figwidth(5
330
331
332
              fig.set_figheight(5)
              for i, k in enumerate(ks):
    if k > 2:
        # use TSNE when we have more than 2 dimensions
        embedding = TSNE(
335
                                 n_components=2,
                                 learning_rate="auto",
init="random",
338
339
                                  perplexity = 10,
340
                                  random_state=DEFAULT_SEED.
341
                           ). fit_transform (log_responsibilities[i])
                    else:
# otherwise we can visualise responsibility vectors without dimensionality reduction
embedding = np.exp(log_responsibilities[i])
ax = plt.subplot(1, n, i + 1)
ax.scatter(embedding[:, 0], embedding[:, 1])
344
346
                    ax.set_title(f"{k=}
              fig.suptitle(figure_title)
349
350
              plt.savefig(figure_path, bbox_inches="tight")
352
353
       def _plot_log_posteriors (
    log_posteriors: List[List[float]],
354
              ks: List[int],
epsilon: float,
figure_title: str,
355
356
357
              figure_path: str,
       ) -> None:
360
              Plot log posteriors as a function of EM iteration for different ks:
param log_posteriors: list of vectors, each representing the log posterior during EM for a specific k:
param ks: list of k values used for each Theta
param epsilon: value used for early stopping of EM
361
363
364
365
               :param figure_title: name of figure
366
               :param figure-path: path to store figure
367
              :return:
368
              fig \ , \ ax = plt.subplots(len(ks), 1, constrained\_layout=True)
369
              fig.set_figwidth(10)
370
              fig.set_figheight(10)
              fig.set_lignelgit(lt)
for i, k in enumerate(ks):
    ax[i].plot(np.arange(1, len(log_posteriors[i]) + 1), log_posteriors[i])
    ax[i].set_xlabel("Step")
    ax[i].set_ylabel(f"Unnorm'd Log_Posterior")
    ax[i].set_title(f"{k=} {epsilon=}")
373
374
375
              plt.suptitle(figure_title)
```

```
379
              plt.savefig(figure_path)
381
       ks: List[int], log_posteriors: List[List[float]], i: int, n: int, d: int) -> pd.DataFrame:
382
383
384
385
              Compute the compress rate, not taking into account the cost of storing model parameters :param ks: k values to use for each trial :param log_posteriors: list of vectors, each representing the log_posterior during EM for a specific k
386
387
               :param i: trial number
:param n: number of data points
:param d: number of dimensions per data point
380
390
391
               return: dataframe containing the compression rate for this trial
392
393
394
              df = pd.DataFrame(
395
                     data = [
396
                                  397
399
400
                     columns=ks.
401
               ) . T
402
              df = df.reset_index()
df.columns = ["k value", f"Trial {i}"]
return df.set_index("k value")
403
404
405
406
407
408
        def _compute_total_compression_ratio(
       ks: List[int], log-posteriors: List[List[float]], i: int, n: int, d: int) -> pd.DataFrame:
409
410
412
              Compute the total compress ratio, taking into account the cost of storing model parameters (assuming
               float64)
413
               :param ks: k values to use for each trial
              :param log-posteriors: list of vectors, each representing the log posterior during EM for a specific k :param i: trial number :param n: number of data points :param d: number of dimensions per data point
414
416
417
               return: dataframe containing the total compression ratios for this trial
418
419
              df = pd.DataFrame(
420
421
                     data = [
422
423
                                  np.round(
                                        (-log_{-posterior}[-1] + (64 * ks[j] * (d + 1))) / (np.log(2) * n * d), 2,
424
425
426
427
                                   for j, log_posterior in enumerate(log_posteriors)
428
                           ]
430
                     columns=ks,
              ) . T
431
              df = df.reset_index()
432
              dr = dr.reset_index()
df.columns = ["k value", f"Trial {i}"]
return df.set_index("k value")
433
434
435
436
437
              x: np.ndarray,
438
               alpha_parameter: float,
439
               beta_parameter: float,
441
               number_of_trials: int,
              ks: List[int],
epsilon: float,
442
443
444
               max_number_of_steps: int ,
              figure_path: str,
figure_title: str
445
447
              \verb|compression_csv_path|: & str|,
448
       ) -> None:
449
              Produces answers for question 3e:
param x: numpy array of shape (N, D)
:param alpha_parameter: alpha parameter of the beta prior
:param beta_parameter: beta parameter of the beta prior
:param number_of_trials: number of trails to run EM
450
451
452
453
454
               :param ks: k values to use for each trial
:param epsilon: value used for early stopping of EM
:param max_number_of_steps: maximum number of steps during EM
455
456
458
               :param figure_title: base name of figures
:param figure_path: base paths to store figure
459
460
               :param compression_csv_path: path to store bits data
              :return:
461
462
463
             n, d = x.shape
np.random.seed(DEFAULT_SEED)
df.compression_list: List[pd.DataFrame] = []
df_total_compression_list: List[pd.DataFrame] = []
for i in range(number_of_trials):
   init_thetas: List[Theta] = []
   em_thetas: List[Theta] = []
   log_posteriors: List[List[float]] = []
   log_responsibilities: List[np.ndarray] = []
   for j, k in enumerate(ks):
        init_theta = _init_params(k, d)
              n, d = x.shape
464
465
466
467
468
469
470
```

```
em_theta, log_responsibility, log_posterior = _run_expectation_maximisation(
475
                                theta=init_theta ,
alpha_parameter=alpha_parameter ,
476
477
478
                                 beta_parameter=beta_parameter,
                                 epsilon=epsilon,
479
480
                                max_number_of_steps=max_number_of_steps,
481
                          init_thetas.append(init_theta)
482
                          em_thetas.append(em_theta)
                          log_responsibilities.append(log_responsibility)
log_posteriors.append(log_posterior)
484
485
486
487
                    _visualise_p_matrix( init_thetas,
488
489
                          figure_title=f"{figure_title} Trial {i}: Initialised P",
figure_path=f"{figure_path}-{i}-initialised-p.png",
490
492
                    _visualise_p_matrix(
em_thetas,
493
                          figure_title=f"{figure_title} Trial {i}: EM Optimised P", figure_path=f"{figure_path}-{i}-optimised-p.png",
495
496
498
                    _visualise_responsibility_clusters (
499
500
                          log_responsibilities,
                          ks, figure_title=f"{figure_title} Trial {i}: TSNE Responsibility Visualisation", figure_path=f"{figure_path}-{i}-tsne.png",
501
502
503
504
505
                    _plot_log_posteriors (
506
                          log_posteriors,
                          ks,
epsilon,
507
                          figure_title=f"{figure_title} Trial {i}: Unnormalised Log-Posterior", figure_path=f"{figure_path}-{i}-log-pos.png",
509
                    df_compression_list.append(
513
                          _compute_compression_rate(ks, log_posteriors, i, n, d)
                    df_total_compression_list.append(
    _compute_total_compression_ratio(ks, log_posteriors, i, n, d)
             \label{eq:pd.concat} $$ \frac{1}{concat}(df_{compression\_list}, axis=1).to_{csv}(f^{*}\{compression_{csv\_path}\}.csv^{*}) $$ pd.concat(df_{total\_compression\_list}, axis=1).to_{csv}(f^{*}\{compression_{csv\_path}\}-total.csv^{*}) $$ f^{*}\{compression_{csv\_path}\}-total.csv^{*}\}$
518
520
```

src/solutions/q3.py

(e) Running the algorithm a few times starting from randomly chosen initial conditions and visualising the parameters:

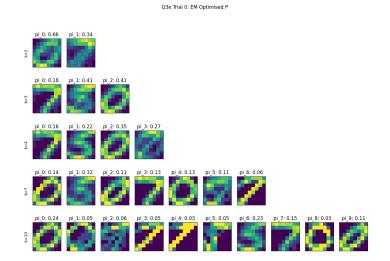


Figure 7: EM optimised parameters: Trial 0

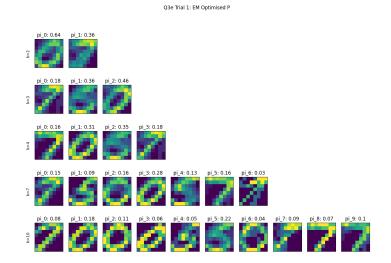


Figure 8: EM optimised parameters: Trial 1

Q3e Trial 2: EM Optimised P

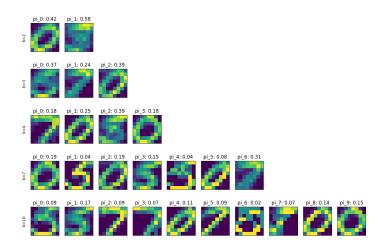


Figure 9: EM optimised parameters: Trial 2

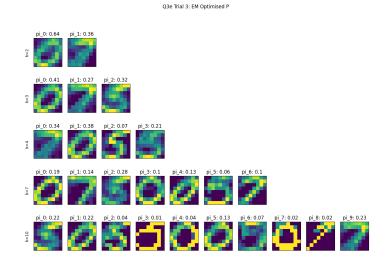


Figure 10: EM optimised parameters: Trial 3

For smaller k, we can visually see that we obtain very similar solutions (a seven and a zero for k = 2, although for trial 2, it looks more like zero and five mixed with seven). For k = 3, we get one each of zero, seven, and five, but in different permutations for different trials. However for higher k, we see that this may not always be the case. For Trial 1 of k = 10, we have two 5's whereas in Trial 3 we have four 5's. Interestingly, different clusters of the same digits can be different, representing different variants of the written digit (i.e. a slanted zero, a slightly slanted zero, and a symmetric zero).

Moreover, looking at the responsibilities of each mixture component, we can see that when k is relatively small they are relatively evenly distributed. However for k=7 and especially k=10, we can see some components have very small probability (i.e. π_3 of trial 0 and k=10). It will be unlikely for those components to represent very distinct clusters (i.e. the parameters for π_3 and π_4 are very similar in trial 0 and k=10). Moreover, for these small probabilities, we see that the parameters are almost like binary images. This shows that there are not many images represented in these components. This can be verified when we perform a TSNE visualisation of the responsibility vector for each of the images (Note that for k=2, just the responsibility vector is plotted because it is two dimensional). We can see that for large k, qualitatively the number of clusters no longer matches the k value, indicating that some mixtures are redundant. For example for k=7 and k=10 we can only qualitatively see three or four clusters with TSNE.

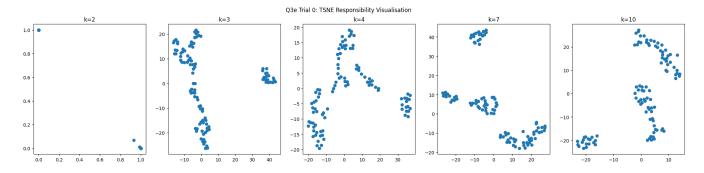


Figure 11: TSNE Visualisation of Image responsibilities: Trial 0

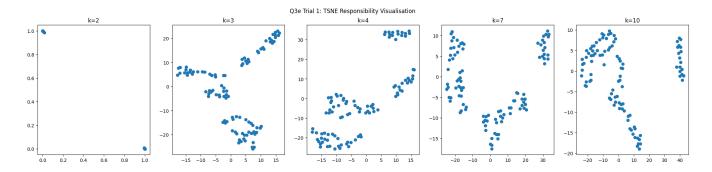


Figure 12: TSNE Visualisation of Image responsibilities: Trial 1

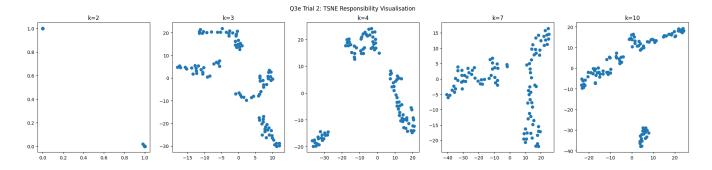


Figure 13: TSNE Visualisation of Image responsibilities: Trial 2

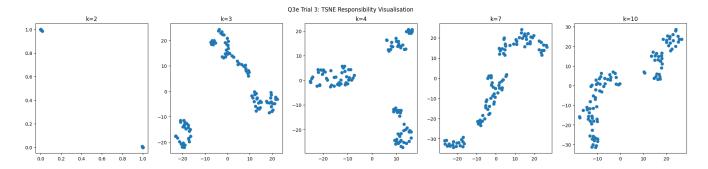


Figure 14: TSNE Visualisation of Image responsibilities: Trial 3

Improvements to the model could include searching for an optimal k by maximising the log posterior with regularisation on the magnitude of k to balance maximising log posterior with minimising model complexity. Additionally, adding a prior on the responsibility components can be helpful to ensure a more even distribution across mixture components unlike the components visualised here. This could help promote more meaningful clusters as k increases. Moreover, more experimentation for choosing better priors can be helpful to find better separation between mixtures. Increasing the size of our data set (i.e. more images) and resolution of our images (i.e. more pixels) can help the model better understand the distinguishing nuances of different mixtures and provide better clustering, although the number of images and the resolution should scale together to ensure that the model doesn't learn the noise in the higher resolution images. This is assuming we are able to scale our computing resources. Finally, given that we know that there are ten digits and our current data set only includes a subset of these digits, we can also expand our data set to include all ten digits. Hopefully, for k = 10, we will then be able to achieve a unique digit for each mixing component, rather than variations of repeated digits as we see now.

(f) The log-likelihood in bits can be expressed as:

$$\log_2(P(\{\mathbf{x}^{(n)}\}_{n=1}^N | \theta))$$

The length of the naive encoding of these binary data is $N \cdot D$, the number of pixels in $\{\mathbf{x}^{(n)}\}_{n=1}^{N}$. This is because the images are binary so each pixel can be represented with a single bit. We can compute compression rate with respect to the ratio of log-likelihood bits to the length of the naive encoding for each k:

$$rate = 1 - \frac{\log_2(P(\{\mathbf{x}^{(n)}\}_{n=1}^N | \theta))}{N \cdot D}$$

Presenting the compression rates for different trials and k values:

k value	Trial 0	Trial 1	Trial 2	Trial 3
2	0.25	0.25	0.26	0.25
3	0.32	0.28	0.29	0.31
4	0.35	0.36	0.36	0.32
7	0.41	0.42	0.38	0.43
10	0.47	0.47	0.49	0.43

Table 2: Compression Rates

As k increases, we can see that our compression rate gets better. This is intuitive because with higher k we are specifying a more complex and expressive model (i.e. with more parameters) and thus we are able to capture more of the structure of the data in the model. Thus, the bit rate, or information provided by a sample decreases with respect to the complexity of the model and thus our compression rate increases. From the source coding theorem, lossless compression algorithms are lower bounded by the entropy of the underlying data generating distribution $P(\mathbf{x})$. This is $-\sum_{\mathbf{x}\in\mathcal{X}}P(\mathbf{x})\log P(\mathbf{x})$. On the other hand, with EM we are maximising $\langle \log P(Z, X|\theta) \rangle_{q(X)} + H[q]$ or minimising the negative of this. When our proposal distribution P matches q, we recover H[P] and we get the optimal model, the data generating model for encoding our data. This makes sense because we would be able to compress our data with the best possible distribution to represent the data. This matches the lower bound of the source coding theorem. However, because it is very unlikely that our proposal q will actually match P, our compression rate will always be worse than the optimal compression rate of H[P]. On the other hand, a compression algorithm would compressions on a per image basis, independent of the other images. And thus, it is able to attain a better compression rate for that image and is much closer to the source coding theorem lower bound. Depending on the data (i.e. H[P] of the data), the compression rate of gzip can range from 60% to 88% (https://web.dev/optimizing-content-efficiency-optimizeencoding-and-transfer/text-compression-with-gzip), much higher than that of our models, as we expected.

(g) The total cost of encoding with model parameters and data is:

$$\log_2(P(\{\mathbf{x}^{(n)}\}_{n=1}^N | \theta)) + M \cdot K \cdot D + M \cdot K$$

Where M is the cost of storing a single float value, in our case we used float64 so 64 bits. The first term is the log-likelihood as expressed in part (f), the second term is the cost of storing \mathbf{P} , and the last term is the cost of storing π . The latter two terms scale with the value of k. This means that as k increases, our compression rate deteriorates. Looking at the total compression ratio $\frac{\log_2(P(\{\mathbf{X}^{(n)}\}_{n=1}^N|\theta))+M\cdot K\cdot D+M\cdot K}{N\cdot D}$ in a table:

k value	Trial 0	Trial 1	Trial 2	Trial 3
2	2.62	2.63	2.62	2.63
3	3.49	3.54	3.52	3.5
4	4.4	4.39	4.39	4.43
7	7.15	7.14	7.18	7.14
10	9.91	9.91	9.89	9.95

Table 3: Total Compression Ratios

We can see that the ratio is greater than one, meaning that this is actually worse than the naive encoding. This is due to the high cost of storing each value in the model parameters being 64 bits. However, because this remains constant with respect to our data set size, as N increases these ratios will approach $\frac{\log_2(P(\{\mathbf{X}^{(n)}\}_{n=1}^N|\theta)}{N\cdot D}$ and we'll recover our compression rates from part (f). By increasing k, we see that the ratio increases, and thus our compression rate is worse. This makes sense because we are essentially slowly storing the data into the model. In the extreme example of k=N we can have the parameters for each mixture model as an image in the data. Although in this case, we could store our parameters as binary values instead of 64 bit floats. Our mixture component is uniform because each mixture is equally likely (all images are equally likely) so there is no need to store any values for π . Thus, we would recover the cost of naive encoding our data. Taking into account the model parameters further verifies that the cost of encoding with this approach is much higher than the cost of gzip.

Question 5

(a) The formulae for the ML estimates of $P(s_i = \alpha | s_{i-1} = \beta) = \Psi(\alpha, \beta)$:

$$\Psi(\alpha,\beta) = \frac{N_{\alpha,\beta}}{N_{\beta}}$$

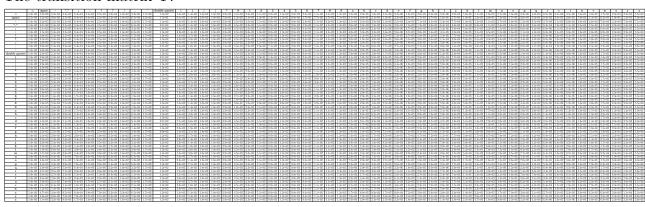
where $N_{\alpha,\beta}$ is the count of the number of occurrences of the pair (α,β) , where β is before α in the text and N_{β} is the number of occurrences of β . Moreover to ensure ergodicity, a one was added to each $N_{\alpha,\beta}$. This was also taken into account for the normaliser N_{β} .

Moreover, the stationary distribution ϕ can be calculated using the power method:

- (i) Initialise any $\phi^{(0)} \in \mathbb{R}^{53 \times 1}$ and $\sum_i \phi_i^{(0)} = 1$
- (ii) Repeat $\phi^{(i+1)} = \Psi \phi^{(i)}$
- (iii) Terminate when $\phi^{(i+1)} \phi^{(i)} < \epsilon$

where $\Psi \in \mathbf{R}^{53 \times 53}$ containing the transition probabilities, $\Psi_{i,j} = P(s^j | s^i)$ where s^i is the i^{th} symbol and s^j is the j^{th} symbol with respect to the indices of symbols in the Ψ matrix, and ϵ is some small number indicating sufficient convergence of the distribution to be considered stationary. The function $\phi(\gamma)$ is simply the index of symbol γ in the vector ϕ .

The transition matrix Ψ :



(Apologies for the tiny font, latex was being difficult)

The invariant distribution ϕ :

Symbol	Probability
=	1.7e-05
space	1.7e-01
-	6.1e-04
	1.2e-02
;	3.9e-04
	2.9e-04
<u> </u>	6.0e-04
?	4.7e-04
/	1.9e-05
	7.7e-03
,	1.9e-05
double quotes	2.4e-05
(2.3e-04
)	2.2e-04
	1.7e-05
1	1.7e-05
*	1.1e-04
0	6.9e-05
1	1.4e-04
2	6.0e-05
3	3.4e-05
4	2.3e-05
5	3.2e-05
6	3.2e-05
7	2.8e-05
8	7.6e-05
9	2.6e-05
a	6.6e-02
b	1.1e-02
С	2.0e-02
d	3.8e-02
e	1.0e-01
f	1.8e-02
g	1.6e-02
h	5.4e-02
i	5.6e-02
j	8.5e-04
k	6.4e-03
1	3.1e-02
m	2.0e-02
n	5.9e-02
0	6.2e-02
р	1.5e-02
q	7.7e-04
r	4.7e-02
S	5.2e-02
t	7.2e-02
u	2.1e-02
v	8.5e-03
w	1.9e-02
x	1.4e-03
У	1.5e-02
Z	7.4e-04
	!

(b) The latent variables $\sigma(s)$ for different symbols s are not independent. This is because by choosing an encoding for one symbol $e = \sigma(s)$, the encoding for a second symbol $\sigma(s')$ cannot be e. We have 53 symbols but only 52 degrees of freedom, because once we have defined the encoding for 52 symbols, the encoding for the 53^{rd} symbol cannot be chosen. Thus, there exists a dependence between $\sigma(s)$ for different symbols s.

The joint probability of the encrypted text $e_1e_2\cdots e_n$ given σ :

$$P(e_1, e_2, ..., e_n | \sigma) = \phi(\gamma = \sigma^{-1}(e_1)) \prod_{i=2}^n \psi(\alpha = \sigma^{-1}(e_i), \beta = \sigma^{-1}(e_{i-1}))$$

because σ is the encoding function, mapping a symbol s into the encoded symbol e, we require σ^{-1} the decoding function mapping the encoded symbol e back to s.

(c) The proposal probability $S(\sigma \to \sigma')$ depends on the permutations of σ and σ' . Our proposal generating process restricts us to choose a proposal σ' that differs from σ only at two spots:

$$\sigma'(s^i) = \sigma(s^j)$$

$$\sigma'(s^j) = \sigma(s^i)$$

for any two symbols s^i and s^j of the 53 possible symbols $(s^i \neq s^j)$.

Therefore, if the above doesn't hold for σ' , $S(\sigma \to \sigma') = 0$. From σ there are $\binom{53}{2}$ possible proposal σ' 's with the above property. Because we are assuming a uniform prior distribution over σ 's, the transition probability of a σ' that satisfies the above property is $S(\sigma \to \sigma') = \frac{1}{\binom{53}{2}}$.

The MH acceptance probability is given as:

$$A(\sigma \to \sigma'|\mathcal{D}) = \min\{1, \frac{S(\sigma' \to \sigma)P(\sigma'|\mathcal{D})}{S(\sigma \to \sigma')P(\sigma|\mathcal{D})})\}$$

where $S(\sigma \to \sigma')$ is the conditional transition probability of σ' given σ and \mathcal{D} is our encrypted text $e_1, e_2, ..., e_n$.

 $S(\sigma \to \sigma') = S(\sigma' \to \sigma)$ for all σ and σ' that differ only at two spots because the probability in this case will always be $\frac{1}{\binom{53}{2}}$, so we can simplify:

$$A(\sigma \to \sigma' | \mathcal{D}) = \min\{1, \frac{P(\sigma' | \mathcal{D})}{P(\sigma | \mathcal{D})}\}$$

From Bayes' Theorem:

$$P(\sigma|\mathcal{D}) = \frac{P(\mathcal{D}|\sigma)P(\sigma)}{\sum_{\sigma'} P(\mathcal{D}|\sigma')P(\sigma')}$$

We are assuming a uniform prior for σ , so $P(\sigma)$ is a constant and we can simplify further:

$$A(\sigma \to \sigma'|\mathcal{D}) = \min\{1, \frac{P(\mathcal{D}|\sigma')}{P(\mathcal{D}|\sigma)}\}$$

This is the acceptance probability for a given proposal σ' . The expression for $P(\mathcal{D}|\sigma)$ is $P(e_1, e_2, ..., e_n|\sigma)$ described in the previous part.

(d) Reporting the current decryption of the first 60 symbols after every 100 iterations:

MH Iteration	Current Decryption 6m p2 2namr'=)mk pn=' batm'=)3t' 2')=q p2 8)*9'= r)b' p' qn
100	er pl losrua= drk po=a bstra=dita lad=n pl -df:a= udba pa no
200	er nl loiruah drw noha bitrahdsta ladhp nl xdymah udba na po
300 400	er nl loiruav srw nova bitravsdta lasvp nl xsymav usba na po er vd dsir,an orw vsna bitranolta daony vd uophan ,oba va ys
500	er c, ,sirdan or. csna bitranolta ,aony c, uophan doba ca ys
700	en ck kyindar on. cyra bitnarolta kaors ck uophar doba ca sy en pk klindar on. plra bitnaroyta kaors pk uochar doba pa sl
800	en p, ,londar in. plra botnariyta ,airs p, fichar diba pa sl en pu ulondar in. plra botnariyta uairs pu fichar diba pa sl
900	en pu ulondar in. plra botnariyta uairs pu fichar diba pa sl
1100	en pl luondar in. pura botnariyta lairs pl fighar diba pa su en pl luondar in. pura cotnarixta lairs pl fighar dica pa su
1200	en pk kuondar inl pura comnarixma kairs pk fighar dica pa su
1300	en ck kuondar inl cura pomnarixma kairs ck fighar dipa ca su en ck koundar inl cora pumnarixma kairs ck fighar dipa ca so
1500	en ck koundar inl cora vumnarixma kairs ck fithar diva ca so
1600 1700	en ck koundar inl cora vumnarixma kairs ck fithar diva ca so
1800	an ck kounder inl core vumnerixme keirs ck fither dive ce so an ck kounler ind core vumnerixme keirs ck fither live ce so
1900	an ck kounler ind core vumnerixme keirs ck fither live ce so
2000	an ck kounler ind core vumnerixme keirs ck fither live ce so an ck kounler ind core vumnerixme keirs ck fither live ce so
2200	an ck kounler ind core vunnerixme keirs ck fither live ce so
2300	an ck kounger ind core vumnerixme keirs ck fither give ce so
2400 2500	an ck kounger ind core vulnerixle keirs ck fither give ce so an mk kounger ind more vulnerixle keirs mk fither give me so
2600	an mk kounger ind more vulneriple keirs mk fither give me so
2700 2800	an mk kounger ind more vulneriple keirs mk fither give me so an mk kounger ind more vulneriple keirs mk fither give me so
2900	an mk kounger ind more vulneriple keirs mk fither give me so
3000 3100	an mk kounger ind more vulneriple keirs mk fither give me so
3200	an mf founger ind more vulneriple feirs mf kither give me so an mf founger ind more vulneriple feirs mf kither give me so
3300	an mf founger ind more vulneriple feirs mf kither give me so
3400 3500	an mf founger ind more vulneriple feirs mf kither give me so an mf founger ind more vulneriple feirs mf kither give me so
3600	an mf founger ind more vulneriple feirs mf kither give me so
3700 3800	an mf founger ind more vulneriple feirs mf kither give me so an mf founger ind more vulneriple feirs mf kither give me so
3900	in mf founger and more vulneraple fears mf kather gave me so
4000	in mf founger and more vulneraple fears mf kather gave me so
4100 4200	in mf founger and more vulneraple fears mf kather gave me so in mf founger and more vulnerable fears mf kather gave me so
4300	in mf founger and more vulnerable fears mf kather gave me so
4400 4500	in mf founger and more vulnerable fears mf yather gave me so in mf founger and more vulnerable fears mf yather gave me so
4600	in mf founger and more vulnerable fears mf yather gave me so
4700 4800	in mf founger and more vulnerable fears mf yather gave me so
4900	in mf founger and more vulnerable fears mf yather gave me so in mf founger and more vulnerable fears mf yather gave me so
5000	in mf founger and more vulnerable fears mf yather gave me so
5100 5200	in mf founger and more vulnerable fears mf yather gave me so in mf founger and more vulnerable fears mf yather gave me so
5300	in my younger and more vulnerable years my father gave me so
5400 5500	in my younger and more vulnerable years my father gave me so in my younger and more vulnerable years my father gave me so in my younger and more vulnerable years my father gave me so
5600	in my younger and more vulnerable years my father gave me so
5700 5800	in my younger and more vulnerable years my father gave me so in my younger and more vulnerable years my father gave me so
5900	in my younger and more vulnerable years my father gave me so
6100	in my younger and more vulnerable years my father gave me so in my younger and more vulnerable years my father gave me so
6200	in my younger and more vulnerable years my father gave me so
6300 6400	in my younger and more vulnerable years my father gave me so in my younger and more vulnerable years my father gave me so
6500	in my younger and more vulnerable years my father gave me so
6600	in my younger and more vulnerable years my father gave me so
6700 6800	in my younger and more vulnerable years my father gave me so in my younger and more vulnerable years my father gave me so
6900	in my younger and more vulnerable years my father gave me so
7000 7100	in my younger and more vulnerable years my father gave me so in my younger and more vulnerable years my father gave me so
7200	in my younger and more vulnerable years my father gave me so
7300 7400	in my younger and more vulnerable years my father gave me so
7500	in my younger and more vulnerable years my father gave me so in my younger and more vulnerable years my father gave me so
7600	in my younger and more vulnerable years my father gave me so
7700 7800	in my younger and more vulnerable years my father gave me so in my younger and more vulnerable years my father gave me so
7900	in my younger and more vulnerable years my father gave me so
8000 8100	in my younger and more vulnerable years my father gave me so in my younger and more vulnerable years my father gave me so
8200	in my younger and more vulnerable years my father gave me so
8300 8400	in my younger and more vulnerable years my father gave me so
8500	in my younger and more vulnerable years my father gave me so in my younger and more vulnerable years my father gave me so
8600	in my younger and more vulnerable years my father gave me so
8700 8800	in my younger and more vulnerable years my father gave me so in my younger and more vulnerable years my father gave me so
8900	in my younger and more vulnerable years my father gave me so
9000	in my younger and more vulnerable years my father gave me so
9200	in my younger and more vulnerable years my father gave me so in my younger and more vulnerable years my father gave me so
9300	in my younger and more vulnerable years my father gave me so
9400 9500	in my younger and more vulnerable years my father gave me so in my younger and more vulnerable years my father gave me so
9600	in my younger and more vulnerable years my father gave me so
9700 9800	in my younger and more vulnerable years my father gave me so in my younger and more vulnerable years my father gave me so
9900	in my younger and more vulnerable years my father gave me so
10000	in my younger and more vulnerable years my father gave me so

The corresponding σ :

s	$\sigma(s)$
	,
	l l
space	x h
	, 1
;	n
: !	r
?	e
	f
	b
· ,	3
double quetes	5
double quotes	4
	9
	i
<u>l</u>	0
*	1
0	
1	z m
2	c
3	
4	;
5	
6	*
7	k
8	:
9	q
a)
<u></u>	2
- c	-
d	7
e	,
f	0
g	s
h	!
i	j
j	(
k	8
1	y
m	v
n	d
0	=
P	space
q	6
r	g
s	t
t	double quotes
u	p
v	j
w	a
x	u
У	?
z	w
	1

To help with chain initialisation, 10000 different σ 's were first randomly and independently sampled. The σ with the best log-likelihood was chosen as the starting point for the MH chain and the algorithm was then run for 10000 iterations. Moreover, ten different trials of this was performed, where the trial with the best log-likelihood was displayed. The decrypted message for each of the ten trials:

Trial	Decryption
-0	itedcecoutl featpedof eyunt fa.n ec afredcevas, felay ed ero
1	in my younger and more vulnerable years my father gave me so
2	in cy yomnker and core vmlnerable years cy father kave ce so
- 3	is hy ytoswer asd htre volseraule yearm hy fanger wave he mt
4	in my younger and more vulnerable years my father gave me so
5	"5407""0""][4)81094307]180(['4819*'80""891207""0:96=810)9(807802]"
6	"542)(2(]94""18234=2)]812:9'4183*'12(13862)(2[307182""3:12)126]"
7	ioadcaclyon earowadle agy.o erk. ac retadcafrsu eanrg ad atl
- 8	in my younker and more vulnerable years my father kave me so
- 0	in my younker and more vulnerable years my father lave me so

The Python code for the MH sampler:

```
from typing import Dict, List, Tuple
3
     import numpy as np
import pandas as pd
     from sklearn.preprocessing import normalize
     from src.constants import DEFAULT_SEED
     def _convert_to_scientific_notation(x: float) -> str:
10
           Convert value to string in scientific notation
           :param x: value to convert
:return: string of x in scientific notation
"""
13
14
           return "\{:.1e\}".format(float(x))
19
     class Decrypter:
           \begin{array}{lll} def & \_\_init\_\_(self \;,\; decryption\_dict \colon \; Dict[\,str \;,\; \,str \,]\,) \; -\!\!> \; None \colon \\ \end{array}
20
                Decrypter containing the mapping a symbol to its encrypted symbol :param decryption_dict:
22
24
25
                 self.decryption_dict = decryption_dict
26
           def decrypt(self, encrypted_message: str) -> str:
                Decrypts an encrypted message using the decryption dictionary
30
                 : param\ encrypted\_message:\ the\ encrypted\ message\ to\ decrypt
                 :return: decrypted message
33
34
                return \ "".join([self.decryption\_dict[x] \ for \ x \ in \ encrypted\_message])
35
           def table(self) -> pd.DataFrame:
36
                Generate table containing symbol decryptions :return: pandas table of decryptions """
38
                decrpyter_table = pd.DataFrame(
    self.decryption_dict.items(), columns=["s", "sigma(s)"]
42
                decrpyter_table [decrpyter_table == ""] = "space"
decrpyter_table [decrpyter_table == '"'] = "double quotes"
return decrpyter_table.set_index("s")
44
45
     class Statistics:
           def __init__(
    self ,
50
51
                training_text: str,
symbols: List[str],
invariant_stopping_epsilon: float = 5e-20,
55
56
57
           ) -> None:
                Statistics for text
                :param training_text: training text for calculating transition and invariant probability :param symbols: symbols in the training text :param invariant_stopping_epsilon: stopping condition for constructing the invariant distribution
58
59
60
61
62
                self.training_text = training_text
                 self.symbols = symbols
                self.num.symbols = len(symbols)
self.symbols_dict = self._construct_symbols_dictionary(symbols)
self.transition_matrix = self._construct_transition_matrix(
64
67
                      {\tt training\_text}\ ,\ {\tt self.symbols\_dict}
69
70
71
72
73
74
75
76
                 self.invariant_distribution = self._approximate_invariant_distribution(
                      invariant_stopping_epsilon
                 self.log_transition_matrix = np.log(self.transition_matrix)
                 self.log_invariant_distribution = np.log(self.invariant_distribution)
           def list_of_symbols_for_df(self) -> List[str]:
                Replace certain symbols to prepare for dataframe :return: list of symbols with some replacements ""
                x = self.symbols.copy()
                83
84
                return x
86
           @property
           def transition_table(self) -> pd.DataFrame:
                Generate a table containing transition probabilities :return: transition probabilities
89
91
                df_transitions = pd.DataFrame(
92
                      data=self.transition_matrix
94
                      columns=self.list_of_symbols_for_df,
```

```
95
96
                 df_transitions.index = self.list_of_symbols_for_df
                 return df_transitions.applymap(_convert_to_scientific_notation)
97
98
aa
           def invariant_distribution_table(self) -> pd.DataFrame:
100
                Generate a table containing invariant distribution probabilities return: invariant distribution probabilities
104
                df =
                     pd . DataFrame (
106
                           data = self.invariant_distribution.reshape(1, -1),
107
108
                           columns = self.list_of_symbols_for_df,
                      .applymap(_convert_to_scientific_notation)
110
                      .transpose()
                      .reset_index()
                df.columns = ["Symbol", "Probability"]
return df.set_index("Symbol")
114
           @staticmethod
           \label{lem:def_construct_symbols_dictionary(symbols: List[str]) -> Dict[str, int]:} \\
                 Construct a dictionary mapping each symbol to an integer to index the transition matrix
                and the invariant distribution :param symbols: list of symbols to map :return: symbol to integer mapping """
124
                return \ \{k\colon \ v \ for \ v \,, \ k \ in \ enumerate (\, symbols \,) \,\}
126
           def _construct_transition_matrix(
           self, text: str, symbols_dict: Dict[str, int]
128
130
                Constructs the transition matrix for a given text :param text: string to calculate transition matrix with
                 :param symbols_dict: dictionary mapping symbol to a dictionary
134
                :return:
                # initialise with ones to ensure ergodicity
transition_matrix = np.ones((self.num_symbols, self.num_symbols))
for i in range(1, len(text));
136
138
                140
142
144
                \textcolor{return}{\texttt{return}} \hspace{0.2cm} \texttt{transition\_matrix}
145
           def _approximate_invariant_distribution (
147
                self, invariant_stopping_epsilon: float
           ) -> np.ndarray:
148
                Approximate the invariant distribution with the power method :param invariant_stopping_epsilon: stopping condition for constructing the invariant distribution :return: the invariant distribution as a vector (number of symbols, 1)
150
                {\tt invariant\_distribution} \ = \ {\tt np.zeros} \, (\, (\, {\tt self.num\_symbols} \, , \ 1) \, )
                previous_invariant_distribution = invariant_distribution.copy()
156
                # make sure it's a proper distribution that sums to one
158
                invariant_distribution [0] = 1
                while (
    np.linalg.norm(invariant_distribution - previous_invariant_distribution)
161
                     > invariant_stopping_epsilon
163
                      previous_invariant_distribution = invariant_distribution.copy()
164
                invariant_distribution = self.transition_matrix @ invariant_distribution return invariant_distribution
165
166
168
           def log_transition_probability(self, alpha: str, beta: str) -> float:
169
                Look up the log probability of the transition from symbol alpha to beta :param alpha: symbol that is being transitioned from :param beta: symbol that is being transitioned to :return: probability of transition """
172
                return self.log_transition_matrix[
    self.symbols_dict[beta], self.symbols_dict[alpha]
176
178
179
           def log_invariant_probability(self, gamma: str) -> float:
180
                Look up the log probability of a symbol with respect to the invariant distribution
181
                :param gamma: symbol to query
:return: log probability of the symbol
182
183
184
185
                return self.log_invariant_distribution[self.symbols_dict[gamma]].item()
186
           def compute_log_probability(self, text: str) -> float:
187
                Compute the log probability of a given text containing symbols
189
                :param text: text to compute log probability for
190
```

```
191
                   :return: log probability of the text
192
                   \begin{split} \log_{\texttt{probability}} &= \texttt{self.log\_invariant\_probability} \, (\texttt{text} \, [0]) \\ \text{for } i & \text{in } \texttt{range} \, (1, \, \, \texttt{len} \, (\texttt{text})) \colon \\ & \text{log\_probability} \, +\! = \, \texttt{self.log\_transition\_probability} \, (\texttt{text} \, [\, i \, ] \, , \, \, \texttt{text} \, [\, i \, - \, 1]) \end{split}
194
195
                   return log_probability
196
197
198
       class MetropolisHastingsDecryption:
    def __init__(self , symbols: List[str]):
    """
199
200
201
                   Metropolis Hastings MCMC for Decryption :param symbols: set of symbols to decrypt
202
203
204
205
                   self.symbols = symbols
206
207
             def generate_random_decrypter(self) -> Decrypter:
208
209
                   Generates a random decrypter
                   :return: a Decrypter instantiation
210
212
                   return Decrypter (
213
                                self.symbols[i]: self.symbols[x]
                                for i, x in enumerate
                                     np.random.permutation(np.arange(len(self.symbols)))
216
218
219
                   )
220
             @staticmethod
             \begin{array}{ll} \textbf{def} & \texttt{generate\_proposal\_decryption} \, (\, \texttt{decrypter} \, \colon \, \, \texttt{Decrypter} \, ) \, \, \, \to \, \, \texttt{Decrypter} \, : \\ \end{array}
222
223
                   Generate a proposal decrypter by randomly swapping two of the decryption mappings : param decrypter: the decrypter used to generate the proposal
224
                    :return: a proposal decrypter
226
                   x1 = np.random.choice(list(decrypter.decryption_dict.keys()))
                   x2 = np.random.choice(list(decrypter.decryption_dict.keys()))
proposal_decryption = decrypter.decryption_dict.copy()
proposal_decryption[x2], proposal_decryption[x1] = (
230
                          decrypter.decryption_dict[x1], decrypter.decryption_dict[x2],
233
234
                    return Decrypter (proposal_decryption)
236
237
              @staticmethod
             def _choose_decrypter(
    statistics: Statistics,
238
                    encrypted_message: str,
current_decrypter: Decrypter,
240
241
                   proposal_decrypter: Decrypter,
243
             ) -> Decrypter:
244
                   Choose between the current and proposal decrypter
                   :param statistics: Statistics instantiation for calculating log probabilities :param encrypted_message: the encrypted message
246
248
                    :param current_decrypter: the current decrypter
249
                    :param proposal_decrypter: the proposal decrypter
                   :return:
251
                   # calculate log probabilities
current_log_probability = statistics.compute_log_probability(
254
                         text=current_decrypter.decrypt(encrypted_message),
                   proposal_log_probability = statistics.compute_log_probability(
    text=proposal_decrypter.decrypt(encrypted_message),
257
258
                   )
                   # calculate acceptance probability
acceptance_probability = np.min(
260
261
262
                          [1, np.exp(proposal_log_probability - current_log_probability)]
263
264
                   # choose decrypter using the acceptance probability
                   return np.random.choice(
265
                         [current_decrypter, proposal_decrypter],
p=[1 - acceptance_probability, acceptance_probability],
266
268
269
270
              def _find_good_starting_decrypter(
271
                   self, statistics: Statistics,
                    encrypted_message: str
274
                   number\_start\_attempts: int,
             ) -> Decrypter:
276
                    Find a good starting decrypter for the sampler by choosing the one with the best log likelihood: param statistics: Statistics instantiation for calculating log probabilities \\
277
                   :param number_start.attempts: number of possible starting decrypters to check :return: the best starting decrypter for the sampler
279
280
282
                    best_log_likelihood = -np.float("inf")
283
                    best_decrypter = None
285
                    for _ in range(number_start_attempts):
                          decrypter = self.generate_random_decrypter()
286
```

```
288
                               statistics.compute_log_probability(
                                    text=decrypter.decrypt(encrypted_message)
289
290
291
                              > best_log_likelihood
292
                        ):
293
                              best_decrypter = decrypter
204
                  return best_decrypter
295
             def run (
297
                   self,
                   encrypted_message: str.
298
                   statistics: Statistics,
299
300
                   number_of_mh_loops: int
                   number_start_attempts: int,
301
302
                   log_decryption_interval: int,
303
             log_decryption_size: int ,
) -> Tuple[Decrypter , List[str]]:
304
305
                  Run the sampler with two steps:
306
                        1. find a good starting decrypter for the sampler 2. run the sampler
307
308
                   :param encrypted_message: the encrypted message
                  :param encrypted_message: the encrypted message
:param statistics: Statistics instantiation for calculating log probabilities
:param number_of_mh_loops: number of loops to run the metropolis hastings sampler
:param number_start_attempts: number of possible starting decrypters to check
:param log_decryption_interval: number of samples between logging the decrypted message
:param log_decryption_size: number of symbols to decrypt when logging the decrypted message
:return: a tuple containing the decrypter found from the sampler and the logged decryption message
"""
309
310
311
312
314
315
317
                  decrypter = self._find_good_starting_decrypter(
    statistics, encrypted_message, number_start_attempts
318
319
                  logged_decryption_message = [
    decrypter.decrypt(encrypted_message)[:log_decryption_size]
320
321
322
                   for i in range(1, number_of_mh_loops + 1):
    if (i + 1) % log_decryption_interval == 0:
323
                              logged_decryption_message.append(
    decrypter.decrypt(encrypted_message)[:log_decryption_size]
325
326
                         proposal_decrypter = self.generate_proposal_decryption(decrypter)
decrypter = self._choose_decrypter(
328
329
330
                             statistics, encrypted_message, decrypter, proposal_decrypter
331
332
                  return decrypter, logged_decryption_message
333
334
335
       def _construct_decryptions_table(
336
            decryption_messages: List[str], decryption_interval: int, columns: List[str]
       ) -> pd.DataFrame:
337
             decrypted_message_iterations_table = pd.DataFrame(
339
                         np.arange(0, len(decryption_messages)) * decryption_interval,
340
                         decryption_messages ,
342
343
             ).transpose()
             decrypted_message_iterations_table.columns = columns
344
345
             return decrypted_message_iterations_table.set_index(columns[0])
347
348
       def a(
             symbols: List[str],
             training_text: str,
transition_matrix_path: str,
350
351
             invariant_distribution_path: str,
353
       ) -> None:
354
             Produces answers for question 5a
            :param symbols: symbols in the training text
:param training.text: training text for calculating transition and invariant probability
:param transition_matrix_path: path to store transition matrix
:param invariant_distribution_path: path to store invariant distribution
356
357
358
359
             :return:
360
361
             statistics = Statistics (
362
                  training_text ,
364
                  symbols,
365
366
             statistics.transition_table.to_csv(transition_matrix_path)
367
             statistics.invariant_distribution_table.to_csv(invariant_distribution_path, sep="|")
368
369
370
       def d(
371
             encrypted_message: str,
             symbols: List[str],
             training_text: str ,
number_trials: int ,
374
             number_of_mh_loops: int ,
             number_start_attempts: int
             log_decryption_interval: int,
378
             log_decryption_size: int,
             trial_decryptions_table_path: str,
379
             decryptor_table_path: str
381
             decrypted_message_iterations_table_path: str,
382
      ) -> None:
```

```
383
384
              Produces answers for question 5d
             Produces answers for question 5d
:param encrypted_message: the encrypted message
:param symbols: symbols in the training text
:param training_text: training text for calculating transition and invariant probability
:param number_trials: number of times to restart and run the sampler
:param number_of_mh_loops: number of loops to run the metropolis hastings sampler
:param number_start_attempts: number of possible starting decrypters to check
:param log_decryption_interval: number of samples between logging the decrypted message
:param log_decryption_size: number of symbols to decrypt when logging the decrypted message
:param trial_decryptions_table_path: path to store decryption messages for each trial
:param decrypted_message_iterations_table_path: path to store logged decryption messages
:return:
385
386
387
388
389
390
391
303
394
395
              :return:
396
397
398
              statistics = Statistics (
399
                    training_text ,
400
                    symbols,
401
              np.random.seed (DEFAULT_SEED)
402
403
              metropolis_hastings_decryption = MetropolisHastingsDecryption(symbols)
              decrypters: List[Decrypter] = []
log_likelihoods: List[float] = []
logged_decryption_messages: List[List[str]] = []
404
405
406
              407
408
409
410
411
412
                           number_of_mh_loops,
413
                           number_start_attempts ,
log_decryption_interval ,
414
415
                           log_decryption_size,
416
                     decrypters.append(decrypter)
                     log_likelihoods.append(
418
                           statistics.compute_log_probability(decrypter.decrypt(encrypted_message))
419
420
                     logged_decryption_messages.append(logged_decryption_message) decryption_messages.append(
421
422
                           decrypter.decrypt(encrypted_message)[:log_decryption_size]
424
              df_trial_decryptions = _construct_decryptions_table(
425
                    decryption.messages=[x[:log_decryption_size] for x in decryption_messages], decryption_interval=1, columns=["Trial", "Decryption"],
426
427
428
429
              df_trial_decryptions.to_csv(trial_decryptions_table_path, sep="|")
430
431
432
              # sort trials by log likelihood
              433
435
437
                     decryption_interval=log_decryption_interval
438
                    columns=["MH Iteration",
                                                            "Current Decryption"],
439
              df_logged_decryptions.to_csv(decrypted_message_iterations_table_path, sep="|")
440
```

src/solutions/q5.py

- (e) When some values of $\Psi(\alpha, \beta) = 0$, this affects the ergodicity of the chain. An ergodic chain is one that is irreducible (i.e. all possible transitions between symbols, including to itself, have probability greater than zero). If $\Psi(\alpha, \beta) = 0$, this means that there is zero probability that β will transition to α , breaking our definition. To restore ergodicity, we can add a small transition probability between all symbols of the chain. This essentially acts as a prior, stating that the probability of a symbol to transition to any other symbol (including itself) should never be zero.
- (f) If we were to use symbol probabilities alone for decoding, the joint probability would be:

$$P(e_1, e_2, ..., e_n | \sigma) = \prod_{i=1}^n P(\sigma^{-1}(e_i))$$

the product of the likelihoods of the decoded letters. In this case, the optimal decoding would simply replace the most frequent symbols in the encrypted message with the most frequent symbols in the training text. This decoding approach is much more difficult because each letter is assumed to be independent of its neighbours. For a first order Markov chain, we exploit the structure of language by considering pairs of letters. Assuming that as the training text size approaches infinity and the size of the encrypted message also approaches infinity, that the two will have the same symbol frequency and that the probability of each symbol is unique, (i.e. two different symbols can't have the same frequency), then using symbol probabilities alone should theoretically work by matching symbol probabilities. However, in practise it would be unlikely to be able to make these assumptions about symbol frequencies, especially with the finite size of our training set and encrypted message. Therefore in practise, symbol probabilities alone would not be sufficient.

A second-order chain should also work in theory. However, with this approach it is probably practically more difficult for finding a suitable decoding. This is because our transition tensor would contain N^3 elements, where N is the number of symbols, to account for all possible second order transitions. Our training text would need to increase quadratically to maintain the same ratio of possible transitions to example transitions (number of first order transitions in a text of length N is N-1 and second order its N-2). This can also introduce sparsity (in this case, small non-zero probabilities because ergodicity is maintained) in our transition tensor. Thus, the log-likelihood of many areas of σ space might be very small or the same as their neighbours, when the transition probabilities are mostly just the offset probability added to maintain ergodicity. Navigating this space will be much more difficult for the sampler.

For an encryption scheme where two symbols map to the same encrypted value:

$$\exists \alpha, \beta, \sigma(\alpha) = \sigma(\beta), \alpha \neq \beta$$

this approach can become much more complicated. Our σ is no longer as easily inverted and therefore for each duplicate mapping, we would have to integrate out the probability for the two possible decrypted symbols when computing the log-likelihood. Moreover, generating proposal encodings is not as simple as swapping the encryption for two symbols. This is because we do not know which two symbols map to the same encrypted symbol and simply swapping would preserve the same collision mapping of the current encoding. Moreover, the number of proposal σ' 's will depend on how many duplicates exist in the current σ .

Thus $S(\sigma \to \sigma')$ would no longer be symmetric, complicating the acceptance probability calculation as it would be dependent on the σ and σ' . Overall, this approach could work but would require many changes to accommodate for these complications. Integrating out collision mappings in the log-likelihood, non-symmetric proposal probabilities, and a much larger σ space because duplicates are allowed, means that it will take much longer for the sampler to find a reasonable σ .

If we used this approach for Chinese with ≥ 10000 symbols, we would be attempting to solve the same problem but with $N \geq 10000$ instead of N=53. Similar to the second order Markov chain, although this is theoretically possible, it would require a transition matrix of size $\geq 10000^2$ which is quite impractical and we'd run into similar problems as for second order Markov Chains. An alternative set up could be with using Chinese phonetics, for which there are much fewer than 10000, however this would require a mapping from a phonetic to an encrypted phonetic.

Question 7

(a) To find the local extrema of the function f(x,y) = x+2y subject to the constraint $y^2+xy=1$, first we define g(x,y):

$$g(x,y) = y^2 + xy - 1$$

where g(x,y) = 0 is an equivalent representation of the given constraint.

We can therefore construct the optimisation problem:

$$\min_{\mathbf{x}} f(\mathbf{x})$$

such that $g(\mathbf{x}) = 0$ and $\mathbf{x} := [x, y]^T$.

We can calculate $\nabla f(\mathbf{x})$:

$$\nabla f(\mathbf{x}) = \left[\frac{\partial}{\partial x}(x+2y), \frac{\partial}{\partial y}(x+2y)\right]^T$$

$$\nabla f(\mathbf{x}) = [1, 2]^T$$

and calculating $\nabla g(\mathbf{x})$:

$$\nabla g(\mathbf{x}) = \left[\frac{\partial}{\partial x}(y^2 + xy - 1), \frac{\partial}{\partial y}(y^2 + xy - 1)\right]^T$$

$$\nabla g(\mathbf{x}) = [y, 2y + x]^T$$

Solving the constraint optimisation problem with Lagrange multipliers, we set up the equations:

$$\nabla f(\mathbf{x}) + \lambda \nabla g(\mathbf{x}) = \mathbf{0}$$

and

$$g(\mathbf{x}) = 0$$

Giving us the three equations:

$$1 + \lambda y = 0$$

$$2 + \lambda(2y + x) = 0$$

$$y^2 + xy - 1 = 0$$

Substituting $y = \frac{-1}{\lambda}$ from the first equation into the second equation:

$$2 + \lambda(2(\frac{-1}{\lambda}) + x) = 0$$

$$x = 0$$

Solving for y in our third equation with x = 0:

$$y^2 - 1 = 0$$

We see that $y = \pm 1$ and from the first equation $\lambda \mp 1$.

The local extrema are (x=0,y=1) when $\lambda=-1$ and (x=0,y=-1) when $\lambda=1$.

(b)

(i) Given that $g(a) = \ln(a)$, we want to transform this to the form f(x, a) = 0 where x = g(a):

$$x = \ln(a)$$
$$\exp(x) - a = 0$$

Thus,

$$f(x,a) = \exp(x) - a$$

(ii) We know that for Newton's method's

$$x_{n+1} = x_n - \frac{f(x_n)}{f'(x_n)}$$

where $f(x_n) = \exp(x_n) - a$

We can calculate:

$$f'(x) = \frac{\partial f(x, a)}{\partial x} = \exp(x)$$

Assuming we can evaluate $\exp(x)$, our update equation is:

$$x_{n+1} = x_n - \frac{\exp(x_n) - a}{\exp(x_n)}$$

Simplifying:

$$x_{n+1} = x_n + \frac{a}{\exp(x_n)} - 1$$

we have our update equation in Newton's algorithm for this problem.

Question 8

(a) For:

$$\sup_{\{\mathbf{X}\in\mathbb{R}^n\}}R_A(\mathbf{x})$$

where $R_A(\mathbf{x}) = \frac{\mathbf{x}^T \mathbf{A} \mathbf{x}}{\|\mathbf{x}\|^2}$, we want to show that a maximum is attained.

To do this, we will first show that the above optimisation can be equivalently formulated as:

$$\sup_{\left\{\mathbf{x}\in\mathbb{R}^n\big|\,\|\mathbf{x}\|=1\right\}}R_A(\mathbf{x})$$

We begin by considering any $\mathbf{w} \in \mathbb{R}^n$ and let $\mathbf{x} = \frac{\mathbf{w}}{\|\mathbf{w}\|}$. Because $\|\mathbf{x}\| = 1$ we can substitute:

$$\sup_{\left\{\mathbf{X}\in\mathbb{R}^{n}\big|\|\mathbf{X}\|=1\right\}}R_{A}(\mathbf{x}) = \sup_{\left\{\frac{\mathbf{W}}{\|\mathbf{W}\|}\in\mathbb{R}^{n}\bigg|\|\frac{\mathbf{W}}{\|\mathbf{W}\|}\|=1\right\}} \frac{\mathbf{w}^{T}\mathbf{A}\mathbf{w}\|\mathbf{w}\|^{2}}{\|\mathbf{w}\|^{2}\mathbf{w}^{T}\mathbf{w}}$$

where $\|\mathbf{x}\|^2 = \mathbf{x}^T \mathbf{x}$.

The set $\left\{\frac{\mathbf{w}}{\|\mathbf{w}\|} \in \mathbb{R}^n \mid \|\frac{\mathbf{w}}{\|\mathbf{w}\|}\| = 1\right\}$ contains all $\mathbf{w} \in \mathbb{R}^n$ so we can rewrite:

$$\sup_{\left\{\mathbf{x} \in \mathbb{R}^n \mid \|\mathbf{x}\|=1\right\}} R_A(\mathbf{x}) = \sup_{\left\{\mathbf{w} \in \mathbb{R}^n\right\}} \frac{\mathbf{w}^T \mathbf{A} \mathbf{w} \|\mathbf{w}\|^2}{\|\mathbf{w}\|^2 \mathbf{w}^T \mathbf{w}}$$

We can simplify the expression:

$$\sup_{\left\{\mathbf{x} \in \mathbb{R}^n \middle| \|\mathbf{x}\| = 1\right\}} R_A(\mathbf{x}) = \sup_{\left\{\mathbf{w} \in \mathbb{R}^n\right\}} \frac{\mathbf{w}^T \mathbf{A} \mathbf{w}}{\mathbf{w}^T \mathbf{w}}$$

$$\sup_{\left\{\mathbf{X}\in\mathbb{R}^n\big|\|\mathbf{X}\|=1\right\}}R_A(\mathbf{x})=\sup_{\left\{\mathbf{W}\in\mathbb{R}^n\right\}}R_A(\mathbf{w})$$

and recover our original optimisation problem by letting $\mathbf{x} = \mathbf{w}$, showing that it is equivalent to the supremum over the unit sphere. Assuming the set containing the unit sphere is compact, the extreme value theory of calculus states that $\sup_{\{\mathbf{x} \in \mathbb{R}^n | ||\mathbf{x}|| = 1\}} R_A(\mathbf{x})$ is attained so equivalently $\sup_{\{\mathbf{x} \in \mathbb{R}^n\}} R_A(\mathbf{x})$ is attained as required.

(b) We can now reformulate the optimisation as:

$$\sup_{\left\{\mathbf{x} \in \mathbb{R}^n \middle| \|\mathbf{x}\| = 1\right\}} \frac{\mathbf{x}^T \mathbf{A} \mathbf{x}}{\|\mathbf{x}\|^2}$$

Because $\|\mathbf{x}\| = 1$ (i.e. choosing $\mathbf{w} \in \mathbb{R}^n$ and $\mathbf{x} = \frac{\mathbf{w}}{\|\mathbf{w}\|}$ to reformulate the problem over the unit sphere), we can equivalently write:

$$\sup_{\left\{\mathbf{X} \in \mathbb{R}^n \middle| \|\mathbf{X}\| = 1\right\}} \mathbf{x}^T \mathbf{A} \mathbf{x}$$

Thus, showing $R_A(\mathbf{x}) \leq \lambda_1$ will be equivalent to showing $\mathbf{x}^T \mathbf{A} \mathbf{x} \leq \lambda_1$ for $||\mathbf{x}|| = 1$. We know that for all $\mathbf{x} \in \mathbb{R}^n$:

$$\mathbf{x} = \sum_{i=1}^{n} (\xi_i^T \mathbf{x}) \xi_i$$

so we can write:

$$\mathbf{x}^T A x = \left(\sum_{i=1}^n (\xi_i^T \mathbf{x}) \xi_i^T\right) \mathbf{A} \left(\sum_{i=1}^n (\xi_i^T \mathbf{x}) \xi_i\right)$$

Given that ξ_i R are eigenvectors of **A** corresponding to eigenvalues λ_i :

$$\mathbf{x}^T \mathbf{A} \mathbf{x} = \left(\sum_{i=1}^n (\xi_i^T \mathbf{x}) \xi_i^T \right) \left(\sum_{i=1}^n \lambda_i (\xi_i^T \mathbf{x}) \xi_i \right)$$

Given that the eigenvectors ξ_i form an orthonormal basis, we know that $\xi_i^T \xi_j = 0$ when $i \neq j$ and $\xi_i^T \xi_j = 1$ when i = j, so:

$$\mathbf{x}^T \mathbf{A} \mathbf{x} = \sum_{i=1}^n \lambda_i (\xi_i^T \mathbf{x})^2$$

From our above reformulation with the unit sphere, we know that $\|\mathbf{x}\|^2 = 1$ so $\|\mathbf{x}\|^2 = \sum_{i=1}^n x_i^2 = \sum_{j=i}^n (\mathbf{x})^2 = 1$. Thus the quantity $\sum_{i=1}^n \lambda_i (\xi_i^T \mathbf{x})^2$ is a weighted average of λ_i 's with weights $(\xi_i^T \mathbf{x})^2$, which is always less than or equal to the largest λ_i value so:

$$\mathbf{x}^T \mathbf{A} \mathbf{x} = \sum_{i=1}^n \lambda_i (\xi_i^T \mathbf{x})^2 \le \lambda_1$$

where λ_1 is the largest eigenvalue of eigenvalues λ_i . Therefore, $R_A(\mathbf{x}) \leq \lambda_1$ as required.

(c) Given that $\mathbf{x} \in span\{\xi_{k+1},...,\xi_n\}$, we can rewrite \mathbf{x} :

$$\mathbf{x} = \sum_{i=k+1}^{n} (\xi_i^T \mathbf{x}) \xi_i$$

Using the same line of argument as in part (b) we can bound $\mathbf{x}^T \mathbf{A} \mathbf{x}$:

$$\mathbf{x}^T \mathbf{A} \mathbf{x} = \sum_{i=k+1}^n \lambda_i (\xi_i^T \mathbf{x})^2 \le \max \{\lambda_{k+1}, ..., \lambda_n\}$$

But given that the maximum eigenvalue λ_1 is not contained in $\{\lambda_{k+1},...,\lambda_n\}$:

$$\max\{\lambda_{k+1},...,\lambda_n\} < \lambda_1$$

and therefore $R_A(\mathbf{x}) < \lambda_1$ as required.

Appendix 1: constants.py

```
import os

DATA_FOLDER = "data"

BINARY_DIGITS_FILE_PATH = os.path.join(DATA_FOLDER, "binarydigits.txt")

MESSAGE_FILE_PATH = os.path.join(DATA_FOLDER, "message.txt")

SYMBOLS_FILE_PATH = os.path.join(DATA_FOLDER, "symbols.txt")

TRAINING_TEXT_FILE_PATH = os.path.join(DATA_FOLDER, "war_and_peace.txt")

OUTPUTS_FOLDER = "outputs"

DEFAULT_SEED = 0
```

src/constants.py

Appendix 2: main.py

```
import os
 2 3
       import numpy as np
              src.constants import (
BINARY_DIGITS_FILE_PATH,
 6
7
              MESSAGE_FILE_PATH,
              OUTPUTS FOLDER.
              SYMBOLS_FILE_PATH
              TRAINING_TEXT_FILE_PATH,
11
12
       \stackrel{'}{	ext{from}} src.solutions import q1, q2, q3, q5
13
       if --name__ == "--main_-":
    if not os.path.exists(OUTPUTS_FOLDER):
        os.makedirs(OUTPUTS_FOLDER)
14
15
16
17
18
              x \; = \; \mathrm{np.loadtxt} \; (\, \mathrm{BINARY\_DIGITS\_FILE\_PATH})
             # Question 1
Q1_OUTPUT_FOLDER = os.path.join(OUTPUTS_FOLDER, "q1")
if not os.path.exists(Q1_OUTPUT_FOLDER):
    os.makedirs(Q1_OUTPUT_FOLDER)
19
20
23
                     figure_path=os.path.join(Q1_OUTPUT_FOLDER, "qld.png"),
figure_title="Qld: Maximum Likelihood Estimate",
26
28
              q1.e(
29
30
                     alpha=3,
31
                     beta=3,
32
                      \label{eq:figure_path}  figure\_path=os.path.join\left(Q1\_OUTPUT\_FOLDER,\ "q1e"\right),
33
                     figure_title="Q1e: Maximum A Prior",
34
36
              Q2_OUTPUT_FOLDER = os.path.join(OUTPUTS_FOLDER, "q2")
37
              if not os.path.exists(Q2_OUTPUT_FOLDER):
              \begin{array}{ll} os.makedirs(Q2\_OUTPUT\_FOLDER) \\ q2.c(x, table\_path=os.path.join(Q2\_OUTPUT\_FOLDER, "q2c.csv")) \end{array}
39
40
42
43
              Q3_OUTPUT_FOLDER = os.path.join(OUTPUTS_FOLDER, "q3")
              if not os.path.exists(Q3_OUTPUT_FOLDER):
    os.makedirs(Q3_OUTPUT_FOLDER)
45
47
                     \begin{array}{l} {\tt alpha\_parameter=1} \,+\, 1\,{\tt e}-5\,,\\ {\tt beta\_parameter=1} \,+\, 1\,{\tt e}-5\,, \end{array}
48
                     \begin{array}{l} \text{number_of_trials=4,} \\ \text{ks} = [2, \ 3, \ 4, \ 7, \ 10], \\ \text{epsilon=1e-5,} \end{array}
50
                     max_number_of_steps=int(1e2),
figure_path=os.path.join(Q3_OUTPUT_FOLDER, "q3e"),
figure_title="Q3e",
54
                     \verb|compression_csv_path| = \verb|os.path.join| (Q3\_OUTPUT\_FOLDER, "q3e-compression") \;,
56
57
58
              Q5_OUTPUT_FOLDER = os.path.join(OUTPUTS_FOLDER, "q5")
              if not os.path.exists(Q5_OUTPUT_FOLDER):
    os.makedirs(Q5_OUTPUT_FOLDER)
61
               with open (TRAINING_TEXT_FILE_PATH) as fp:
              training_text = fp.read().replace("\n", "").lower()
with open(SYMBOLS_FILE_PATH) as fp:
    symbols = fp.read().split("\n")
with open(MESSAGE_FILE_PATH) as fp:
64
65
67
                     encrypted_message = fp.read()
68
70
                     symbols,
71
72
73
74
75
76
77
78
79
                     training_text
                     transition_matrix_path=os.path.join(Q5_OUTPUT_FOLDER, "q5a-transition.csv"), invariant_distribution_path=os.path.join(Q5_OUTPUT_FOLDER, "q5a-invariant.csv"),
                     encrypted_message,
                     training_text
                     number_trials=10.
                     number_of_mh_loops=int(1e4)
81
                     number_start_attempts=int(1e4), log_decryption_interval=100,
82
83
                     log_decryption_size=60,
                     trial_decryptions_table_path=os.path.join(Q5_OUTPUT_FOLDER, "q5d-trials.csv"),
decryptor_table_path=os.path.join(Q5_OUTPUT_FOLDER, "q5d-decrypter.csv"),
decrypted_message_iterations_table_path=os.path.join(
Q5_OUTPUT_FOLDER, "q5d-iterations.csv")
84
86
                     ),
```

main.py