COMP0085 Summative Assignment

Jan 6, 2023

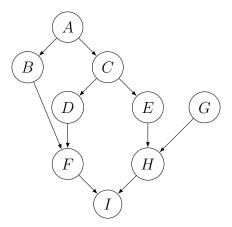
Question 1: A biochemical pathway

(a)

We are given that:

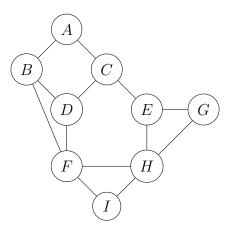
- The concentrations of both species B and C depend on that of A.
- Molecule C seems to redirect the process that produces enzyme D to produce enzyme E instead.
- D catalyses the production of F from B, while E catalyses the production of H from G.
- F and H then combine to form the end product I.

The corresponding directed acyclic graph:

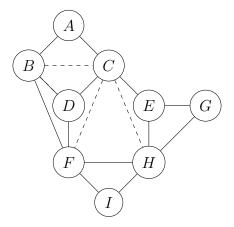


(b)

The moralised graph:

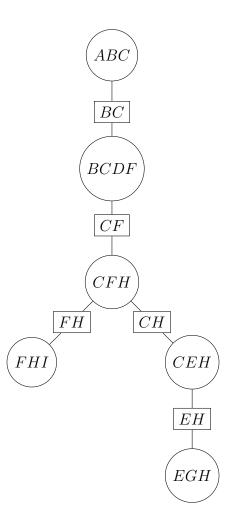


An effective triangulation:

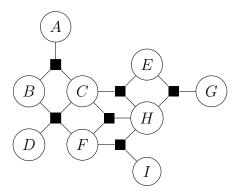


where the dashed lines are edges added to triangulate the moralised graph.

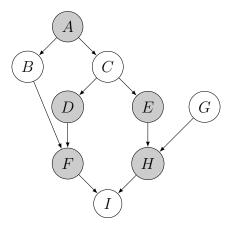
The resulting junction tree:



where the circular nodes are cliques and the square nodes are separators/factors. The junction tree redrawn as a factor graph:



(c)



The set $\{A, D, E, F, H\}$ is a non-unique smallest set of molecules such that if the concentrations of the species within the set are known, the concentrations of the others $\{B, C, G, I\}$ would all be independent (conditioned on the measured ones).

(d)

Using our factor analysis model, we can describe the biochemical pathway as:

$$\delta[\mathbf{x}] = \Lambda \mathbf{z} + \epsilon$$

where $\delta[\mathbf{x}]$ are the concentration perturbations, $\epsilon \sim \mathcal{N}(0, \Psi)$ are the reaction specific Gaussian noise variables, $\Lambda isthefactorloading matrix$, and $\mathbf{z} \sim \mathcal{N}(0, I)$ are the latent factors, the parent concentration perturbations.

From the graph structure, we know that:

Having observations for $\delta[B]$, $\delta[D]$, $\delta[E]$ and $\delta[G]$:

Knowing z are the parent concentration perturbations, we can see that these simplify to a factor analysis set up:

$$\begin{bmatrix} \delta[B] \\ \delta[D] \\ \delta[E] \end{bmatrix} = \begin{bmatrix} \Lambda_{BA} & 0 \\ 0 & \Lambda_{DC} \\ 0 & \Lambda_{EC} \end{bmatrix} \begin{bmatrix} \delta[A] \\ \delta[C] \end{bmatrix} + \begin{bmatrix} \epsilon_B \\ \epsilon_D \\ \epsilon_E \end{bmatrix}$$

Thus, we would expect to recover the factors of A and C, $\delta[A]$ and $\delta[C]$, the two parent nodes of the observations.

(e)

From factor analysis, we would recover $\delta[A]$ and $\delta[C]$. In theory, if these concentration perturbations had unknown parent factors, we could use the results to recover the concentration perturbations of those parent features. However, in our DAG this is not the case, so we would not be able to recover any other species in the cascade because they are all downstream from the known concentration perturbations and those recovered from factor analysis. Similarly, the downstream weights in the DAG from the known measurements would also be unidentifiable. Using factor analysis, we would only be able to identify the upstream weights from the measurements, up to an unknown scale factor. These are the weights Λ_{BA} , Λ_{DC} , and Λ_{EC} . Moreover, knowing $\delta[C]$ from factor analysis, we would also be able to determine Λ_{CA} , up to an unknown scale factor.

Question 2: Bayesian linear and Gaussian process regression

(a)

We want the posterior mean and covariance over a and b. Defining a weight vector \mathbf{w} :

$$\mathbf{w} = \begin{bmatrix} a \\ b \end{bmatrix}$$

Our distribution for w:

$$P(\mathbf{w}) = \mathcal{N}(\mu_{\mathbf{w}}, \Sigma_{\mathbf{w}}) = \mathcal{N}\left(\begin{bmatrix} \mu_a \\ \mu_b \end{bmatrix}, \begin{bmatrix} \sigma_a^2 & 0 \\ 0 & \sigma_b^2 \end{bmatrix}\right)$$

For our data $\mathcal{D} = \{\mathbf{X}, \mathbf{y}\}$:

$$P(\mathcal{D}|\mathbf{w}) = \prod_{t=1}^{T} \mathcal{N} \left(y_t - \mathbf{w}^T \mathbf{x}_t, \sigma^2 \right)$$

where $\mathbf{x}_t = \begin{bmatrix} t_1 \\ 1 \end{bmatrix} \in \mathbb{R}^{2 \times 1}$ and $y_t \in \mathbb{R}^{1 \times 1}$ our response $f_{obs}(t_1)$.

Knowing:

$$P(\mathbf{w}|\mathcal{D}) \propto P(\mathcal{D}|\mathbf{w})P(\mathbf{w})$$

we can substitute the above distributions:

$$P(\mathbf{w}|\mathcal{D}) \propto \exp\left(\frac{-1}{2\sigma^2} \left(\mathbf{y} - \mathbf{w}^T \mathbf{X}\right) \left(\mathbf{y} - \mathbf{w}^T \mathbf{X}\right)^T\right) \exp\left(\frac{-1}{2} \left(\mathbf{w} - \mu_{\mathbf{w}}\right)^T \Sigma_{\mathbf{w}}^{-1} \left(\mathbf{w} - \mu_{\mathbf{w}}\right)\right)$$

where $\mathbf{X} = \begin{bmatrix} t_1 & t_2 \cdots t_T \\ 1 & 1 \cdots 1 \end{bmatrix} \in \mathbb{R}^{2 \times T}$ and $\mathbf{y} \in \mathbb{R}^{1 \times T}$. Expanding:

$$\log P(\mathbf{w}|\mathcal{D}) \propto \frac{-1}{2} \left(\frac{\mathbf{y} \mathbf{y}^T}{\sigma^2} - 2\mathbf{w}^T \frac{\mathbf{X} \mathbf{y}^T}{\sigma^2} + \mathbf{w}^T \frac{\mathbf{X} \mathbf{X}^T}{\sigma^2} \mathbf{w} + \mathbf{w}^T \boldsymbol{\Sigma}_{\mathbf{w}}^{-1} \mathbf{w} - 2\mathbf{w}^T \boldsymbol{\Sigma}_{\mathbf{w}}^{-1} \boldsymbol{\mu}_{\mathbf{w}} + \boldsymbol{\mu}_{\mathbf{w}}^T \boldsymbol{\Sigma}_{\mathbf{w}}^{-1} \boldsymbol{\mu}_{\mathbf{w}} \right)$$

collecting w:

$$\log P(\mathbf{w}|\mathcal{D}) \propto \frac{-1}{2} \left(\mathbf{w}^T \left(\frac{\mathbf{X} \mathbf{X}^T}{\sigma^2} + \Sigma_{\mathbf{w}}^{-1} \right) \mathbf{w} - 2 \mathbf{w}^T \left(\frac{\mathbf{X} \mathbf{y}^T}{\sigma^2} + \Sigma_{\mathbf{w}}^{-1} \mu_{\mathbf{w}} \right) \right)$$

Knowing that the posterior $P(\mathbf{w}|\mathcal{D})$ will be Gaussian with mean $\bar{\mu}_{\mathbf{w}}$ and covariance $\bar{\Sigma}_{\mathbf{w}}$, we can see that expanding the exponent component of a Gaussian would have the form:

$$(\mathbf{w} - \bar{\mu}_{\mathbf{w}})^T \bar{\Sigma}_{\mathbf{w}}^{-1} (\mathbf{w} - \bar{\mu}_{\mathbf{w}}) = \mathbf{w}^T \bar{\Sigma}_{\mathbf{w}}^{-1} \mathbf{w} - 2 \mathbf{w}^T \bar{\Sigma}_{\mathbf{w}}^{-1} \bar{\mu}_{\mathbf{w}} + \bar{\mu}_{\mathbf{w}}^T \bar{\Sigma}_{\mathbf{w}}^{-1} \bar{\mu}_{\mathbf{w}}$$

Thus we can identify the posterior covariance:

$$\bar{\Sigma}_{\mathbf{w}} = \left(\frac{\mathbf{X}\mathbf{X}^T}{\sigma^2} + \Sigma_{\mathbf{w}}^{-1}\right)^{-1}$$

and the posterior mean:

$$\bar{\mu}_{\mathbf{w}} = \bar{\Sigma}_{\mathbf{w}} \left(\frac{\mathbf{X} \mathbf{y}^T}{\sigma^2} + \Sigma_{\mathbf{w}}^{-1} \mu_{\mathbf{w}} \right)$$

Computing the posterior mean and covariance over a and b given by the CO_2 data:

value leters a 1.83E+00 b 3.34E+02		
b 3.34E+02	rs a 1.83E+00	

Figure 1: The Posterior Mean

Figure 2: The Posterior Covariance

(b) Plotting the residuals $g_{obs}(t) = f_{obs}(t) - (a_{MAP}t + b_{MAP})$:

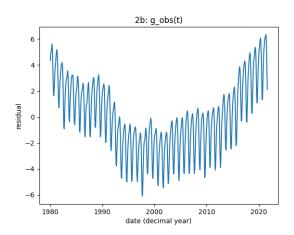


Figure 3: $g_{obs}(t)$

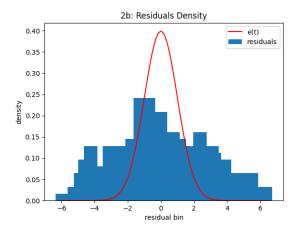


Figure 4: Density Estimation of Residuals vs $e(t) \sim \mathcal{N}(0,1)$

We can see that the residuals do not perfectly conform to our prior over $e(t) \sim \mathcal{N}(0,1)$. The density estimation shows that a mean of zero is a reasonable prior belief however the data does not seem to exhibit unit variance. Also, observing $g_{obs}(t)$ we see that there is periodic structure in the timeseries meaning that the data is not independently and identically distributed as $e(t) \sim \mathcal{N}(0,1)$ would suggest. If this were the case, we would expect g(t) to look like random noise.

(c & d)

We are considering the kernel:

$$k(s,t) = \theta^2 \exp\left(-\frac{2\sin^2(\pi(s-t)/\tau)}{\sigma^2}\right) + \phi^2 \exp\left(-\frac{(s-t)^2}{2\eta^2}\right) + \zeta^2 \delta_{s=t}$$

We can make qualitative observations of this kernel by visualising the covariance (gram) matrix:

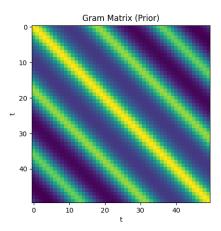


Figure 5: Covariance Matrix

We can observe a striped pattern which indicate higher covariance at regular intervals. This can be attributed to the sinusoidal term in the kernel and encourages periodic structure in functions. Additionally, we can see that covariance values also decay as they are further away from the diagonal. This can be attributed to the exponential term in the kernel, encouraging points closer in time to be more correlated and vice versa. From visualising g(t), we would want to model this with a class of functions which exhibit both of these behaviours as g(t) looks periodic (seasonal with respect to each year) and locally correlated (smooth).

We can also visualise some samples from a Gaussian Process with the same covariance matrix and zero mean. This verifies our observations about the covariance matrix.

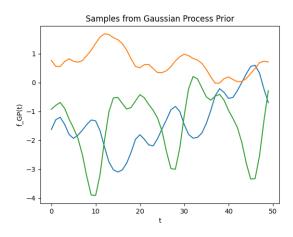


Figure 6: Samples from a zero mean GP with the provided covariance kernel

More specifically, we can see how changing each hyper-parameter will affect the characteristics of the function.

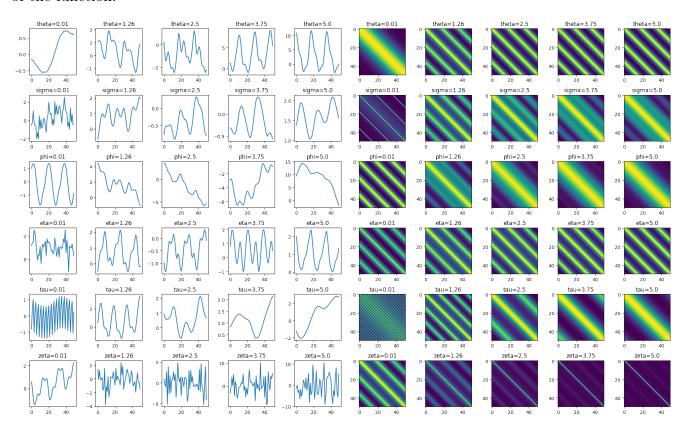


Figure 7: Samples for different parameters

Figure 8: Covariances for different parameters

- θ : As θ increases, we see more pronounced periodic behavior in the sample function by increasing its amplitude. The covariance matrix shows how increasing θ visually reveals the striped periodic component. This is expected because it is the parameter that adjusts the amplitude of the periodic component.
- σ : As σ increases, we see smoother periodic behaviour in the sample function. The covariance matrix shows how increasing σ will increase covariance values around each periodic stripe of the covariance matrix. This is expected because it adjusts the lengthscale of the periodic portion of the kernel.
- ϕ : As ϕ increases, we see the ratio of the amplitude of the periodicity component of the sample function reduces compared to the baseline. The covariance matrix shows how increasing ϕ will start to increase the non-periodic component masking the periodic stripes. This is expected because it adjusts the weight of the squared exponential portion of the kernel. The periodic component remains the same (i.e.same amplitude) but the large baseline shifts from increasing ϕ ends up dominating the function visually.
- η : As η increases we see smoother sample functions. This is expected because the η increases the lengthscale of the squared exponential component, allowing for smoother functions. This causes the off-diagonals of the gram matrix to increase, however the periodic component is still maintained because η doesn't affect the relative weighting between the two components.

- τ : As τ increases, the period of the periodic function increases. We can see this reflected in the stripes in the gram matrix getting further apart and the period of the samples. This makes sense because we are adjusting the period parameter in the sinusoid function of the periodic term.
- ζ : As ζ increases, the function becomes less smooth. This is because the ζ parameter adjusts the weight of the $\delta_{s=t}$ kernel. This places stronger emphasis on the independence of each timestep, which can be seen with the reduction of relative magnitude of off-diagonals in the gram matrix. This masks the periodic and squared-exponential components as we can see with the increased magnitude of the functions as ζ increases.

(e)

Suitable values for hyper-parameters can be chosen through a combination of visual inspection and prior knowledge. For example, it is a reasonable assumption that the CO_2 concentration levels have a strong yearly seasonality behaviour due to the cyclic changes in temperature, humidity, etc. This behaviour cannot be modelled by our Bayesian linear regression and is reflected in our residuals g(t). Thus we can choose $\tau = 1$ to ensure functions with a period of one year to reflect this prior knowledge. It can be difficult to quantitatively choose values for the other parameters as they can relate to the uncertainty exhibited in the data (i.e. the smoothness of the function). One approach is to maximise:

$$\log P(\mathbf{y}|\mathbf{X}) = -\frac{1}{2}\mathbf{y}^T(\mathbf{K} + \sigma^2 \mathbf{I})^{-1}\mathbf{y} - \frac{1}{2}\log|\mathbf{K} + \sigma^2 \mathbf{I}| - \frac{n}{2}\log(2\pi)$$

the log-likelihood of the posterior distribution with respect to the given data where \mathbf{K} is the gram matrix for the kernel (equation 2.30 from http://gaussianprocess.org/gpml/chapters/RW2.pdf). We define a loss function as the negative log-likelihood and employ gradient-based algorithms to find optimal values for these parameters.

Comparing the hyperparameters corresponding to before and after training side by side:

	value
parameter	
eta (kernel)	5.00E+00
phi (kernel)	1.00E+01
sigma	1.00E+00
sigma (kernel)	5.00E+00
tau (kernel)	1.00E+00
theta (kernel)	5.00E+00
zeta (kernel)	2.00E+00

Figure 9: Untrained hyperparameters

Figure 10: Trained Hyperparmaeters

We can analyse some of the changes in these parameters from optimisation to gain some insights. We can see that τ remains the same as we would expect, given the yearly seasonality we knew apriori. On the other hand, the value for ζ is significantly reduced signifying that $\delta_{s=t}$ is not a very good kernel for representing the data as datapoints at different timesteps do exhibit correlations and are not independently and identically distributed.

(f)

Extrapolating the CO_2 concentration levels using both the untrained and trained hyperparameters:

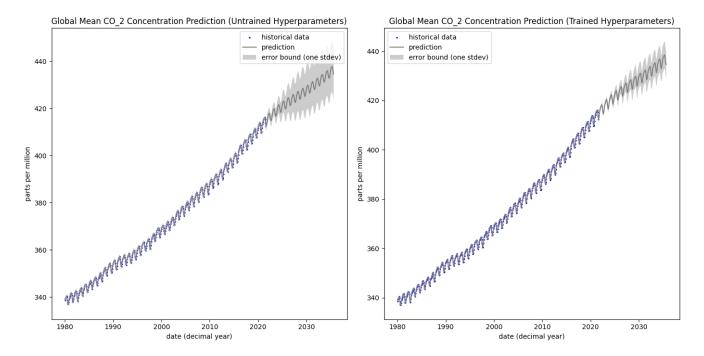


Figure 11: Untrained extrapolation

Figure 12: Trained extrapolation

We can see that the extrapolation shows a continued increase in CO_2 in the future. This follows our expectations given that the levels has been steadily increasing in the past and our Bayesian linear regression model assumption. Moreover, the concentration continues to exhibit yearly seasonality (for the trained extrapolation) as we would expect through the choice of our kernel for modelling the residuals g(t). We can see that the conclusions can be quite sensitive to kernel hyperparameters when comparing the extrapolations from before and after training. Prior to training, the extrapolated prediction is not representative of the given data, with very large uncertainties. After training, we can see that the prediction is much more reasonable, and qualitatively the uncertainty bounds seem to exhibit the historical variability in the data.

(g)

This procedure is not fully Bayesian because despite using a posterior estimate of our linear regression terms, we only use a MAP point estimate when making prediction. For a fully Bayesian approach, we should also incorporate the uncertainty of the linear regression parameters into our extrapolation/uncertainty bounds. For our procedure, we only include the uncertainty of g(t) however it can be observed in the plots that the trend is not perfectly linear so this should be reflected in the uncertainty of our extrapolation. Another approach could be to add a linear kernel to our combined kernel function and model f(t) directly with our kernel, removing the linear regression component in our procedure. Thus our kernel extrapolation would incorporate the uncertainty of all components of our signal.

The Python code for Bayesian Linear Regression:

```
from dataclasses import dataclass
3
     import numpy as np
 6
     @dataclass
      class LinearRegressionParameters:
           Parameters for linear regression
10
11
12
           mean: np.ndarray # weight vector (1, number of features)
13
14
           covariance: np.ndarray # covariance matrix on mean (number of features, number of features)
           @property
16
17
           def precision(self) -> np.ndarray:
    return np.linalg.inv(self.covariance)
18
19
           \begin{array}{lll} \textbf{def} & \texttt{predict} \, (\, \texttt{self} \, \, , \, \, \, \texttt{x: np.ndarray} \, ) \, \, -\!\!\!> \, \texttt{np.ndarray} \, : \end{array}
20
21
22
                Linear regression prediction.
                :param x: design matrix (number of features, number of data points) :return: predicted response matrix (1, number of data points)
23
24
25
26
                return self.mean.T @ x
27
28
29
30
     {\bf class\ Theta:} \\ {\bf linear-regression-parameters:\ LinearRegressionParameters}
31
32
33
34
           @property
35
           def variance(self) -> float:
36
                return self.sigma**2
38
          def precision(self) -> float:
return 1 / self.variance
40
41
42
     def compute_linear_regression_posterior(
44
           x: np.ndarray,
45
           y: np.ndarray,
           prior_linear_regression_parameters: LinearRegressionParameters,
47
            esiduals_precision: float
     ) -> LinearRegressionParameters:
48
49
           Compute the parameters of the posterior distribution on the linear regression weights
51
           : param \ x: \ design \ matrix \ (number \ of \ features \ , \ number \ of \ data \ points)
           :param y: response matrix (1, number of data points)
:param prior_linear_regression_parameters: parameters for the prior distribution on the linear regression
           :param residuals-precision: the precision of the residuals of the linear regression :return: parameters for the posterior distribution on the linear regression weights
57
58
           posterior_covariance = np.linalg.inv(
59
                residuals\_precision * x @ x.T + prior\_linear\_regression\_parameters.precision
           posterior_mean = posterior_covariance @ (
61
                residuals_precision * x @ y.T
+ prior_linear_regression_parameters.precision
@ prior_linear_regression_parameters.mean
62
65
66
           return LinearRegressionParameters (
                mean=posterior_mean, covariance=posterior_covariance
```

src/models/bayesian_linear_regression.py

The Python code for kernels:

```
from abc import ABC, abstractmethod
     from dataclasses import dataclass
3
     import jax.numpy as jnp
     from jax import vmap
 6
     @dataclass
     class KernelParameters (ABC):
10
          An abstract dataclass containing the parameters for a kernel.
11
12
13
14
     class Kernel (ABC):
16
17
          An abstract kernel.
19
20
          Parameters: KernelParameters = None
21
22
          @abstractmethod
23
          def _kernel(
24
                self , parameters: KernelParameters , x: jnp.ndarray , y: jnp.ndarray
          ) -> jnp.ndarray:
25
26
                   Kernel evaluation between a single feature x and a single feature y.
27
28
29
                    parameters: parameters dataclass for the kernel
30
                    x: ndarray of shape (number_of_dimensions,)
y: ndarray of shape (number_of_dimensions,)
               The kernel evaluation. (1, 1)
34
35
36
                raise NotImplementedError
38
          self, parameters: KernelParameters, x: jnp.ndarray, y: jnp.ndarray = None
) -> jnp.ndarray:
          def kernel (
39
40
41
               """ Kernel evaluation for an arbitrary number of x features and y features. Compute k(x, x) if y is None. This method requires the parameters dataclass and is better suited for parameter optimisation.
42
44
45
                     parameters: parameters dataclass for the kernel
                    x: ndarray of shape (number_of_x_features, number_of_dimensions)
y: ndarray of shape (number_of_y_features, number_of_dimensions)
47
48
49
               A gram matrix k(x, y), if y is None then k(x,x). (number_of_x_features, number_of_y_features)
50
51
                # compute k(x, x) if y is None
               if y is None:
y = x
56
               # add dimension when x is 1D, assume the vector is a single feature
               x = jnp.atleast_2d(x)
58
               y = jnp.atleast_2d(y)
59
60
               61
62
63
                return vmap(
lambda x_i: vmap(
65
66
               )(x) (y),
                          lambda y_i: self._kernel(parameters, x_i, y_i),
67
          def __call__(
    self, x: jnp.ndarray, y: jnp.ndarray = None, **parameter_args
) -> jnp.ndarray:
70
71
72
73
74
75
76
               > jnp.ndarray:
""" Kernel evaluation for an arbitrary number of x features and y features.
This method is more user-friendly without the need for a parameter data class.
It wraps the kernel computation with the initial step of constructing the parameter data class from the provided parameter arguments.
77
78
79
                    x: ndarray of shape (number_of_x_features, number_of_dimensions)
y: ndarray of shape (number_of_y_features, number_of_dimensions)
**parameter_args: parameter arguments for the kernel
80
81
83
               A gram matrix k(x, y), if y is None then k(x,x). (number_of_x_features, number_of_y_features).
84
                parameters = self.Parameters(**parameter_args)
86
                return self.kernel(parameters, x, y)
          def diagonal (
89
90
                self,
91
                x: jnp.ndarray,
                y: jnp.ndarray = None,
92
                 **parameter_args ,
          ) -> jnp.ndarray:
```

```
""" Kernel evaluation of only the diagonal terms of the gram matrix.
 96
 97
                        x: ndarray of shape (number_of_x_features, number_of_dimensions)
y: ndarray of shape (number_of_y_features, number_of_dimensions)
**parameter_args: parameter arguments for the kernel
 98
 aa
101
                        A diagonal of gram matrix k(x, y), if y is None then trace(k(x, x)). (number_of_x_features, number_of_y_features)
103
                   \# compute k(x, x) if y is None if y is None:
106
107
108
                        y = x
109
110
                   \# add dimension when x is 1D, assume the vector is a single feature
                  x = jnp.atleast_2d(x)

y = jnp.atleast_2d(y)
113
114
                   assert (
                   x.shape[1] == y.shape[1]
), f"Dimension Mismatch: {x.shape[1]=} != {y.shape[1]=}"
116
117
                   assert (
                  x.shape[0] = y.shape[0]), f"Must have same number of features for diagonal: \{x.shape[0]=\}! = \{y.shape[0]=\}"
120
                   return vmap(
                        lambda x_i, y_i: self._kernel(
parameters=self.Parameters(**parameter_args),
123
124
                               y=y_i ,
126
127
                   )(x, y)
128
             def trace(
130
                   \verb|self|, x: jnp.ndarray|, y: jnp.ndarray| = None, **parameter\_args|
             ) \rightarrow jnp.ndarray: "" Trace of the gram matrix, calculated by summation of the diagonal matrix.
134
                   Args:
                        x: ndarray of shape (number_of_x_features, number_of_dimensions)
y: ndarray of shape (number_of_y_features, number_of_dimensions)
**parameter_args: parameter arguments for the kernel
135
136
                   The trace of the gram matrix k(x, y).
140
141
                   \begin{array}{ll} parameters = self. Parameters (**parameter\_args) \\ \hline return \ jnp.trace (self.kernel (parameters, \ x, \ y)) \end{array}
142
144
145
       class CombinedKernelParameters (KernelParameters):
147
148
149
             Parameters for the Combined Kernel:
151
             log_theta: float
             log_sigma: float log_phi: float
154
             log_eta: float
log_tau: float
156
             log_zeta: float
158
             @property
             def theta(self) -> float:
    return jnp.exp(self.log_theta)
160
161
162
163
             @property
             def sigma(self) -> float:
164
                  return jnp.exp(self.log_sigma)
165
166
             @property
167
            def phi(self) -> float:
    return jnp.exp(self.log_phi)
168
169
             @property
172
173
             def eta(self) -> float:
                  return jnp.exp(self.log_eta)
174
\frac{175}{176}
            @property
def tau(self) -> float:
                  return jnp.exp(self.log_tau)
178
179
             @property
             def zeta(self) -> float:
    return jnp.exp(self.log_zeta)
180
181
182
             @theta.setter
def theta(self, value: float) -> None:
    self.log_theta = jnp.log(value)
183
184
185
186
187
             @sigma.setter
             def sigma(self, value: float) -> None:
189
                   self.log_sigma = jnp.log(value)
190
```

```
191
          @phi.setter
          def phi(self, value: float) -> None:
    self.log_phi = jnp.log(value)
192
193
194
195
          def eta(self , value: float) -> None:
    self.log_eta = jnp.log(value)
196
197
198
          @tau.setter
199
          def tau(self, value: float) -> None:
200
201
               self.log\_tau = jnp.log(value)
202
203
          def zeta(self, value: float) -> None:
    self.log_zeta = jnp.log(value)
204
205
206
207
      class CombinedKernel(Kernel):
208
209
210
          The kernel defined as:  k(x, y) = \frac{2 \sin^2 2 (\exp(-(2 \sin^2 2(\pi - y)/\tan 2)) + \pi^2 * \exp(-(x - y)^2/(2 * \cot^2 2)) }{2 + 2 \cot^2 2 * \det(x - y)} 
211
212
213
          Parameters = CombinedKernelParameters
215
216
          def _kernel(
               self,
218
219
               parameters: CombinedKernelParameters,
          x: jnp.ndarray,
y: jnp.ndarray,
) -> jnp.ndarray:
""" Kernel evaluation between a single feature x and a single feature y.
220
221
223
224
226
                   parameters: parameters dataclass for the Gaussian kernel
227
                   x: ndarray of shape (1,)
y: ndarray of shape (1,)
229
230
               Returns:
               The kernel evaluation.
231
232
233
               return jnp.dot(
234
                   jnp.ones(1),
236
                         (parameters.theta**2)
                         * (
* (
237
238
                                  240
241
243
                             )
244
                        246
249
               )
```

src/models/kernels.py

The Python code for Gaussian Process Regression:

```
from dataclasses import dataclass
from typing import Any, Dict, Tuple
 3
      import jax.numpy as jnp
      import optax
      from jax import grad
from optax import GradientTransformation
10
      from src.models.kernels import Kernel
11
13
14
      @\,d\,a\,t\,a\,c\,l\,a\,s\,s
      class Gaussian Process Parameters:
           Parameters for a Gaussian Process: log_sigma: logarithm of the noise parameter
16
17
                 kernel: parameters for the chosen kernel
19
20
           log_sigma: float
kernel: Dict[str, Any]
21
22
23
24
           def variance(self) -> float:
    return self.sigma**2
25
26
27
28
           @property
29
           def sigma(self) -> float:
30
                 return jnp.exp(self.log_sigma)
           def sigma(self, value: float) -> None:
    self.log_sigma = jnp.log(value)
36
      class Gaussian Process:
38
           A Gaussian measure defined with a kernel, better known as a Gaussian Process.
39
40
41
           Parameters = GaussianProcessParameters
42
           def __init__(self, kernel: Kernel, x: jnp.ndarray, y: jnp.ndarray) -> None:
    """ Initialising requires a kernel and data to condition the distribution.
44
45
47
48
                       kernel: kernel for the Gaussian Process
49
                       x: design matrix (number_of_features, number_of_dimensions)
                 y: response vector (number_of_features, )
50
51
                 self.number_of_train_points = x.shape[0]
                 self.x = x
self.y = y
                  self.kernel = kernel
56
           \textcolor{red}{\textbf{def}} \hspace{0.2cm} \texttt{\_compute\_kxx\_shifted\_cholesky\_decomposition} \hspace{0.1cm} (
           self, parameters
) -> Tuple[jnp.ndarray, bool]:
58
59
60
                 Cholesky decomposition of (kxx + (1/ ^2)*I)
61
62
63
                       parameters: parameters dataclass for the Gaussian Process
                       cholesky_decomposition_kxx_shifted: the cholesky decomposition (number_of_features,
67
             number_of_features)
                 lower\_flag: ^{'}flag \ indicating \ whether \ the \ factor \ is \ in \ the \ lower \ or \ upper \ triangle \ """
68
                 kxx = self.kernel(self.x, **parameters.kernel)
                 kxx_shifted = kxx + parameters.variance * jnp.eye(self.number_of_train_points)
kxx_shifted_cholesky_decomposition , lower_flag = jax.scipy.linalg.cho_factor(
    a=kxx_shifted , lower=True
71
72
73
74
75
                 return kxx_shifted_cholesky_decomposition, lower_flag
76
77
78
           def posterior_distribution (
           self, x: jnp.ndarray, **parameter_args
) -> Tuple[jnp.ndarray, jnp.ndarray]:
    """Compute the posterior distribution for test points x.
    Reference: http://gaussianprocess.org/gpml/chapters/RW2.pdf
79
80
82
83
                       x: test points (number_of_features, number_of_dimensions)
85
                       **parameter\_args:\ parameter\ arguments\ for\ the\ Gaussian\ Process
86
                      mean: the distribution mean (number_of_features, )
covariance: the distribution covariance (number_of_features, number_of_features)
88
89
QΩ
                 parameters = self.Parameters(**parameter.args)
kxy = self.kernel(self.x, x, **parameters.kernel)
kyy = self.kernel(x, **parameters.kernel)
91
```

```
kxx_shifted_cholesky_decomposition,
              lower_flag ,
) = self._compute_kxx_shifted_cholesky_decomposition(parameters)
96
97
98
99
              mean = (
100
                  kxy.T
                  @ jax.scipy.linalg.cho_solve(
                       c_and_lower=(kxx_shifted_cholesky_decomposition, lower_flag), b=self.y
104
              ).reshape(
105
                   -1.
              107
108
109
              return mean, covariance
         112
113
114
                   **parameter_args: parameter arguments for the Gaussian Process
118
              Returns:
119
              The negative log likelihood.
122
              parameters = self.Parameters(**parameter_args)
124
                   kxx_shifted_cholesky_decomposition,
125
                   lower_flag
126
              ) = self._compute_kxx_shifted_cholesky_decomposition(parameters)
              negative_log_likelihood = -(
                  -0.5
* (
130
                        self.y.T
                       @ jax.scipy.linalg.cho_solve(
    c_and_lower=(kxx_shifted_cholesky_decomposition, lower_flag),
132
                            b=self.y,
136
                  / jnp.trace(jnp.log(kxx_shifted_cholesky_decomposition))
- (self.number_of_train_points / 2) * jnp.log(2 * jnp.pi)
138
139
140
              return negative_log_likelihood
141
         def -compute-gradient(self, **parameter-args) -> Dict[str, Any]:
    """ Calculate the gradient of the posterior negative log likelihood with respect to the parameters.
143
144
              Args:
146
                   **parameter_args: parameter arguments for the Gaussian Process
147
              A dictionary of the gradients for each parameter argument.
149
150
              gradients = grad(
              lambda params: self.posterior_negative_log_likelihood(**params))(parameter_args)
153
154
              return gradients
157
              optimizer: GradientTransformation,
              number_of_training_iterations: int,
160
              **parameter_args
         ) -> GaussianProcessParameters:
161
               ""Train the parameters for a Gaussian Process by optimising the negative log likelihood.
163
164
                  optimizer: jax optimizer object number_of_training_iterations: number_of_training_iterations: number of iterations to perform the optimizer
165
                   **parameter_args: parameter arguments for the Gaussian Process
167
168
              Returns:
              A parameters dataclass containing the optimised parameters. ""
171
172
              opt_state = optimizer.init(parameter_args)
for _ in range(number_of_training_iterations):
\frac{174}{175}
                   gradients = self._compute_gradient(**parameter_args)
updates, opt_state = optimizer.update(gradients, opt_state)
                   parameter_args = optax.apply_updates(parameter_args, updates)
              return self.Parameters(**parameter_args)
```

src/models/gaussian_process_regression.py

The rest of the Python code for question 2:

```
from dataclasses import asdict, fields from decimal import Decimal
 3
       import dataframe_image as dfi
       import jax
import jax.numpy as jnp
import matplotlib.pyplot as plt
       import numpy as np
import optax
10
       import pandas as pd
11
       import scipy
13
14
       from src.models.bayesian_linear_regression import (
              {\tt Linear Regression Parameters} \ ,
16
17
              compute_linear_regression_posterior,
                src.models.gaussian_process_regression import (
              GaussianProcess,
GaussianProcessParameters,
19
20
21
22
       from src.models.kernels import CombinedKernel, CombinedKernelParameters
23
24
25
       jax.config.update("jax_enable_x64", True)
26
27
28
       \begin{array}{lll} \textbf{def} & \texttt{construct\_design\_matrix} \, (\, t : \, \, \texttt{np.ndarray} \, ) : \end{array}
              \begin{array}{lll} \textbf{return} & \texttt{np.stack} \left( \left( \begin{smallmatrix} t \end{smallmatrix}, & \texttt{np.ones} \left( \begin{smallmatrix} t \end{smallmatrix}. \texttt{shape} \right) \right) \end{smallmatrix}, & \texttt{axis} \! = \! 1 \right) . T \end{array}
29
30
       def a(
    t: np.ndarray,
31
32
              y: np.ndarray,
34
              sigma:
                          float.
              prior_linear_regression_parameters: LinearRegressionParameters,
35
       save_path: str;
) -> LinearRegressionParameters:
36
             x = construct_design_matrix(t)
prior_theta = Theta(
38
39
                     linear_regression_parameters=prior_linear_regression_parameters ,
41
                     sigma=sigma,
42
              posterior_linear_regression_parameters = compute_linear_regression_posterior(
44
45
                     v .
                     prior_linear_regression_parameters,
                     residuals_precision=prior_theta.precision,
47
48
49
              df_mean = pd.DataFrame(
             posterior-linear_regression_parameters.mean, columns=["value"]).apply(lambda col: ["%.2E" % Decimal(val) for val in col]) df.mean.index = ["a", "b"] df.mean = pd.concat([df.mean], keys=["parameters"]) dfi.export(df.mean, save_path + "-mean.png")
50
51
56
              df-covariance = pd.DataFrame(
              df_covariance = pd.DataFrame(
    posterior_linear_regression_parameters.covariance, columns=["a", "b"]
).apply(lambda col: ["%.2E" % Decimal(val) for val in col])
df_covariance.index = ["a", "b"]
df_covariance = pd.concat([df_covariance], keys=["parameters"])
df_covariance = pd.concat([df_covariance.T], keys=["parameters"])
dfi.export(df_covariance, save_path + "-covariance.png")
return posterior_linear_regression_parameters
58
59
60
61
62
63
66
       def b(
              t_year: np.ndarray,
67
              t: np.ndarray,
              y: np.ndarray
              linear_regression_parameters: LinearRegressionParameters, error_mean: float,
70
71
72
73
74
75
76
              error_variance: float ,
              save_path,
             plt.xlabel("date (decimal year)")
plt.ylabel("residual")
plt.title("2b: g-obs(t)")
plt.savefig(save_path + "-residuals-timeseries")
80
81
83
84
              count, bins = np.histogram(residuals, bins=100, density=True)
plt.bar(bins[1:], count, label="residuals")
              plt.plot(
bins[1:],
86
87
                     scipy.stats.norm.pdf(bins [1:], loc=error_mean, scale=error_variance), color="red", label="e(t)",
89
90
91
              plt.xlabel("residual bin")
plt.ylabel("density")
plt.title("2b: Residuals Density")
92
```

```
plt.legend()
             plt.savefig(save_path + "-residuals-density-estimation")
97
             plt.close()
 98
 aa
       def c(
101
             kernel: CombinedKernel,
             {\tt kernel\_parameters}: \ \ Combined Kernel Parameters \ ,
            log_theta_range: np.ndarray,
t: np.ndarray,
103
             number_of_samples: int ,
            save_path: str,
106
       ) -> None:
107
108
            \begin{array}{ll} gram \ = \ kernel\,(\,t\,,\ **asdict\,(\,kernel\_parameters\,)\,) \\ plt\,.imshow\,(\,gram\,) \end{array}
109
            plt.:msnow(gram)
plt.:xlabel("t")
plt.:ylabel("t")
plt.:title("Gram Matrix (Prior)")
plt.:savefig(save_path + "-gram-matrix")
113
114
            plt.close()
116
                    in range (number_of_samples):
                  plt.plot(
117
                       np.random.multivariate_normal(
                             jnp.zeros(gram.shape[0]), gram, size=1
120
                        ). reshape(-1)
            plt.xlabel("t")
plt.ylabel("f-GP(t)")
plt.title("Samples from Gaussian Process Prior")
plt.savefig(save_path + "-samples")
123
124
126
            plt.close()
127
            fig_samples , ax_samples = plt.subplots(
    len(fields(kernel_parameters.__class__)),
128
130
                  len (log_theta_range),
                   figsize=(
                        len(log_theta_range) * 2,
                        len (fields (kernel_parameters.__class__)) * 2,
134
                  frameon=False,
136
            for i, field in enumerate (fields (kernel-parameters.
                                                                                       __class__)):
                  138
140
142
144
                                   jnp.zeros(gram.shape[0]), gram, size=1
                              ). reshape(-1),
145
                        /ax_samples[i][j].set_title(
    f"{field.name.strip('log_')}={np.round(np.exp(log_value), 2)}"
147
148
149
                  {\color{red} \textbf{setattr}} \, (\, \texttt{kernel\_parameters} \, , \, \, \, \texttt{field.name} \, , \, \, \, \texttt{default\_value} \, )
151
             plt.tight_layout()
             plt.savefig(save_path + f"-parameter-samples", bbox_inches="tight")
             plt.close(fig_samples)
154
            fig_gram , ax_gram = plt.subplots(
    len(fields(kernel_parameters.__class__)),
    len(log_theta_range),
155
156
                   figsize=(
    len(log_theta_range) * 2,
    len(fields(kernel_parameters.__class__)) * 2,
158
161
                  frameon=False,
162
163
            for i, field in enumerate (fields (kernel_parameters.
164
                                                                                       __class__)):
                  default_value = getattr(kernel_parameters, field name)
for j, log_value in enumerate(log_theta_range):
165
166
                        gram = kernel(t, **asdict(kernel-parameters))
ax-gram[i][j].imshow(gram)
ax-gram[i][j].set_title(
    f"{field.name.strip('log_')}={np.round(np.exp(log_value), 2)}"
167
168
169
172
                  setattr(kernel_parameters, field.name, default_value)
174
             plt.tight_layout()
            plt.savefig(save_path + f"-parameter-grams", bbox_inches="tight")
plt.close(fig_gram)
\frac{175}{176}
178
179
       def f(
180
             t_train: np.ndarray,
            y_train: np.ndarray,
t_test: np.ndarray,
min_year: float,
181
182
183
             prior_linear_regression_parameters: LinearRegressionParameters, linear_regression_sigma: float,
184
185
186
             kernel: CombinedKernel,
             gaussian_process_parameters: GaussianProcessParameters,
187
             learning_rate: float,
189
            number_of_iterations: int ,
save_path: str ,
190
```

```
191
         ) -> None:
                # Train Bayesian Linear Regression
192
                x_train = construct_design_matrix(t_train)
prior_theta = Theta(
193
194
195
                        {\tt linear\_regression\_parameters=prior\_linear\_regression\_parameters}\;,
196
                        sigma=linear_regression_sigma ,
197
198
                 posterior_linear_regression_parameters = compute_linear_regression_posterior(
                        x_train ,
199
200
                        y_train ,
201
                        prior_linear_regression_parameters
                        residuals_precision=prior_theta.precision,
202
203
204
205
                 residuals = y_train - posterior_linear_regression_parameters.predict(x_train)
206
                 gaussian_process = GaussianProcess(
207
                      kernel, t-train.reshape(-1, 1), residuals.reshape(-1)
208
209
                # Prediction
210
211
                 x_test = construct_design_matrix(t_test)
212
                 linear\_prediction = posterior\_linear\_regression\_parameters.predict(x\_test).reshape(test) = testing the properties of t
213
214
                 mean_prediction, covariance_prediction = gaussian_process.posterior_distribution( t_test.reshape(-1, 1), **asdict(gaussian_process_parameters)
216
218
219
                # Plot
220
                plt.figure(figsize=(7, 7))
                 plt.scatter(
t_train + min_year
222
223
                        y_train.reshape(-1),
224
                        s=2.
                        color="blue",
226
                        label="historical data",
                 plt.plot(
                        t_test + min_year,
linear_prediction + mean_prediction,
230
                        color="gray",
label="prediction",
231
233
234
                 plt.fill_between(
                         t_test + min_year,
                        linear_prediction
236
237
                        + mean_prediction
                        - 1 * jnp.sqrt(jnp.diagonal(covariance_prediction)), linear_prediction
238
                       + mean_prediction

+ 1 * jnp.sqrt(jnp.diagonal(covariance_prediction)),

facecolor=(0.8, 0.8, 0.8),

label="error bound (one stdev)",
240
241
243
244
                 plt.xlabel("date (decimal year)")
plt.ylabel("parts per million")
plt.title("Global Mean CO_2 Concentration Prediction (Untrained Hyperparameters)")
245
246
247
248
                 plt.legend()
249
                 plt.tight_layout()
250
                 plt.savefig(save_path + "-extrapolation-untrained", bbox_inches="tight")
251
                 plt.close()
252
                 df_parameters = pd.DataFrame(
254
255
                                       256
257
258
259
                                for x in gaussian_process_parameters.kernel.keys()
260
                        261
262
263
264
                 265
266
                # Train Gaussian Process Regression (Hyperparameter Tune)
268
                 optimizer = optax.adam(learning_rate)
                269
270
271
272
                 df_parameters = pd.DataFrame(
273
                        [
274
                                      276
277
                                for x in gaussian_process_parameters.kernel.keys()
279
                        280
281
282
                df-parameters = df-parameters.set_index("parameter").sort_values(by=["parameter"])
dfi.export(df-parameters, save_path + "-trained-parameters.png")
283
285
                # Prediction
286
```

```
287
                  x_test = construct_design_matrix(t_test)
288
                  linear\_prediction = posterior\_linear\_regression\_parameters.predict(x\_test).reshape(
289
290
                 \label{eq:continuous} \begin{array}{lll} \text{mean\_prediction} \;\;,\;\; & \text{covariance\_prediction} \;\; = \;\; \text{gaussian\_process.posterior\_distribution} \, (\\ & \text{t\_test.reshape} (-1, \;\; 1) \;\;, \;\; ** \text{asdict} \big( \text{gaussian\_process\_parameters} \big) \end{array}
291
292
293
294
                 # Plot
295
296
                 plt.figure(figsize=(7, 7))
                 plt.scatter(
t_train + min_year,
297
298
299
                          y_train.reshape(-1),
300
                         s=2,
color="blue",
301
                          label="historical data",
302
303
304
                         .piot(
   t_test + min_year,
   linear_prediction + mean_prediction,
   color="gray",
   label="prediction",
305
306
307
308
309
310
                 plt.fill_between (
                         t_test + min_year,
linear_prediction
311
312
                         + mean_prediction
313
                         \begin{array}{ll} + \mbox{ mean\_prediction} \\ - 1 * jnp.sqrt(jnp.diagonal(covariance\_prediction)), \\ linear\_prediction \\ + \mbox{ mean\_prediction} \\ + 1 * jnp.sqrt(jnp.diagonal(covariance\_prediction)), \\ facecolor = (0.8, \ 0.8, \ 0.8), \\ label="error bound (one stdev)", \\ \end{array}
314
315
316
317
318
319
320
                 plt.xlabel("date (decimal year)")
plt.ylabel("parts per million")
plt.title("Global Mean CO_2 Concentration Prediction (Trained Hyperparameters)")
plt.legend()
321
322
323
                  plt.tight_layout()
plt.savefig(save_path + "-extrapolation-trained", bbox_inches="tight")
325
326
```

src/solutions/q2.py

Question 3: Mean-field learning

(a)

The free energy is can be expressed as:

$$\mathcal{F}(q, \theta) = \langle \log P(\mathbf{x}, \mathbf{s} | \theta) \rangle_{q(\mathbf{s})} + H[Q(\mathbf{s})]$$

Knowing,

$$\log P(\mathbf{x}, \mathbf{s}|\theta) = \log P(\mathbf{x}|\mathbf{s}, \theta) + \log P(\mathbf{s}|\theta)$$

we can write:

$$\mathcal{F}(Q, \theta) = \langle \log P(\mathbf{x}|\mathbf{s}, \theta) \rangle_{q(\mathbf{s})} + \langle \log P(\mathbf{s}|\theta) \rangle_{q(\mathbf{s})} + H[q(\mathbf{s})]$$

Moreover, our mean field approximation is:

$$q(\mathbf{s}) = \prod_{k=1}^{K} q_k(s_k)$$

where $q_k(s_k) = (\lambda_k)^{s_k} (1 - \lambda_k)^{(1-s_k)}$. Given that:

$$P(\mathbf{x}|\mathbf{s}, \theta) = \mathcal{N}\left(\sum_{k=1}^{K} s_k \mu_k, \sigma^2 \mathbf{I}\right)$$

we can substitute the appropriate terms:

$$P(\mathbf{x}|\mathbf{s},\theta) = 2\pi^{-\frac{D}{2}}|\sigma^2 \mathbf{I}|^{-\frac{1}{2}} \exp\left(-\frac{1}{2}\left(\mathbf{x} - \sum_{k=1}^K s_k \mu_k\right)^T \frac{1}{\sigma^2} \mathbf{I}\left(\mathbf{x} - \sum_{k=1}^K s_k \mu_k\right)\right)$$

with d being the number of dimensions.

Taking the logarithm:

$$\log P(\mathbf{x}|\mathbf{s}, \theta) = -\frac{D}{2}\log(2\pi\sigma^2) - \frac{1}{2\sigma^2}\left(\mathbf{x}^T\mathbf{x} - 2\mathbf{x}^T\sum_{k=1}^K s_k \mu_k + \sum_{k=1}^K \sum_{k=1}^K s_k s_{k'} \mu_k^T \mu_{k'}\right)$$

The expectation distributed to the relevant terms:

$$\langle \log P(\mathbf{x}|\mathbf{s},\theta) \rangle_{q(\mathbf{s})} = -\frac{D}{2} \log(2\pi\sigma^2) - \frac{1}{2\sigma^2} \left(\mathbf{x}^T \mathbf{x} - 2\mathbf{x}^T \sum_{k=1}^K \langle s_k \rangle_{q_k(s_k)} \mu_k + \sum_{k=1}^K \sum_{k'=1}^K \langle s_k s_{k'} \rangle_{q_k(s_k)q_{k'}(s_k')} \mu_k^T \mu_{k'} \right)$$

Evaluating the expectations:

$$\langle \log P(\mathbf{x}|\mathbf{s}, \theta) \rangle_{q(\mathbf{s})} = -\frac{D}{2} \log(2\pi\sigma^2) - \frac{1}{2\sigma^2} \left(\mathbf{x}^T \mathbf{x} - 2\mathbf{x}^T \sum_{k=1}^K \lambda_k \mu_k + \sum_{k=1}^K \sum_{k'=1, k' \neq k}^K \lambda_k \lambda_{k'} \mu_k^T \mu_{k'} + \sum_{k=1}^K \lambda_k \mu_k^T \mu_k \right)$$

where $\langle s_k s_k \rangle_{q_k(s_k)} = \langle s_k \rangle_{q_k(s_k)}$ because $s_k \in \{0, 1\}$.

Given that:

$$P(\mathbf{s}|\theta) = \prod_{k=1}^{K} \pi_k^{s_k} (1 - \pi_k)^{(1 - s_k)}$$

Taking the logarithm:

$$\log P(\mathbf{s}|\theta) = \sum_{k=1}^{K} s_k \log \pi_k + (1 - s_k) \log(1 - \pi_k)$$

The expectation distributed to the relevant terms:

$$\langle \log P(\mathbf{s}|\theta) \rangle_{q(\mathbf{s})} = \sum_{k=1}^{K} \langle s_k \rangle_{q_k(s_k)} \log \pi_k + (1 - \langle s_k \rangle_{q_k(s_k)}) \log(1 - \pi_k)$$

Evaluating the expectations with respect to $q(\mathbf{s})$:

$$\langle \log P(\mathbf{s}|\theta) \rangle_{q(\mathbf{s})} = \sum_{k=1}^{K} \lambda_k \log \pi_k + (1 - \lambda_k) \log(1 - \pi_k)$$

To compute the third term, we use the mean field factorisation:

$$H\left[q(\mathbf{s})\right] = \sum_{k=1}^{K} H\left[q_k(s_k)\right]$$

Thus,

$$H[q(\mathbf{s})] = -\sum_{k=1}^{K} \sum_{s_k \in \{0,1\}} q_k(s_k) \log q_k(s_k)$$

Substituting the appropriate values:

$$H[q(\mathbf{s})] = -\sum_{k=1}^{K} \lambda_k \log \lambda_k + (1 - \lambda_k) \log(1 - \lambda_k)$$

Combining, we have our free energy expression:

$$\mathcal{F}(q,\theta) = \frac{-D}{2} \log(2\pi\sigma^{2}) - \frac{1}{2\sigma^{2}} \left(\mathbf{x}^{T} \mathbf{x} - 2\mathbf{x}^{T} \sum_{k=1}^{K} \lambda_{k} \mu_{k} + \sum_{k=1}^{K} \sum_{k'=1, k' \neq k}^{K} \lambda_{k} \lambda_{k'} \mu_{k}^{T} \mu_{k'} + \sum_{k=1}^{K} \lambda_{k} \mu_{k}^{T} \mu_{k} \right) + \sum_{k=1}^{K} \lambda_{k} \log \pi_{k} + (1 - \lambda_{k}) \log(1 - \pi_{k}) - \sum_{k=1}^{K} \lambda_{k} \log \lambda_{k} + (1 - \lambda_{k}) \log(1 - \lambda_{k})$$

To derive the partial update for $q_k(s_k)$ we take the variational derivative of the Lagrangian, enforcing the normalisation of q_k :

$$\frac{\partial}{\partial q_k} \left(\mathcal{F}(q, \theta) + \lambda^{LG} \int q_k - 1 \right) = \left\langle \log P(\mathbf{x}, \mathbf{s} | \theta) \right\rangle_{\prod_{k' \neq k} q_{k'}(s_{k'})} - \log q_k(s_k) - 1 + \lambda^{LG}$$

where λ^{LG} is the Lagrange multiplier.

Setting this to zero we can solve for the λ_k that maximises the free energy:

$$\log q_k(s_k) = \langle \log P(\mathbf{x}, \mathbf{s} | \theta) \rangle_{\prod_{k' \neq k} q_{k'}(s_{k'})} - 1 + \lambda^{LG}$$

Similar to our free energy derivation:

$$\langle \log P(\mathbf{x}|\mathbf{s},\theta) \rangle_{\prod_{k' \neq k} q_{k'}(s_{k'})} \propto -\frac{1}{2\sigma^2} \left(\mathbf{x}^T \mathbf{x} - 2\mathbf{x}^T \sum_{k=1}^K \langle s_k \rangle_{\prod_{k' \neq k} q_{k'}(s_{k'})} \mu_k + \sum_{k=1}^K \sum_{k'=1}^K \langle s_k s_{k'} \rangle_{\prod_{k' \neq k} q_{k'}(s_{k'})} \right)$$

and

$$\langle \log P(\mathbf{s}|\theta) \rangle_{\prod_{k' \neq k} q_{k'}(s_{k'})} = \sum_{k=1}^{K} \langle s_k \rangle_{\prod_{k' \neq k} q_{k'}(s_{k'})} \log \pi_k + (1 - \langle s_k \rangle_{\prod_{k' \neq k} q_{k'}(s_{k'})}) \log(1 - \pi_k)$$

We can write:

$$\log q_k(s_k) \propto \log P(\mathbf{x}|\mathbf{s}, \theta)_{\prod_{k' \neq k} q_{k'}(s_{k'})} + \langle \log P(\mathbf{s}|\theta) \rangle_{\prod_{k' \neq k} q_{k'}(s_{k'})}$$

Substituting the relevant terms:

$$\log q_k(s_k) \propto -\frac{1}{2\sigma^2} \left(-2s_k \mathbf{x}^T \mu_k + s_k s_k \mu_k^T \mu_k + 2 \sum_{k'=1, k' \neq k}^K s_k \lambda_{k'} \mu_k^T \mu_{k'} \right) + s_k \log \pi_k + (1-s_k) \log(1-\pi_k)$$

Knowing $\log q_k(s_k) = s_k \log \lambda_k + (1 - s_k) \log(1 - \lambda_k)$:

$$\log q_k(s_k) \propto s_k \log \frac{\lambda_k}{1 - \lambda_k}$$

Thus,

$$s_k \log \frac{\lambda_k}{1 - \lambda_k} \propto -\frac{1}{2\sigma^2} \left(-2s_k \mathbf{x}^T \mu_k + s_k s_k \mu_k^T \mu_k + 2 \sum_{k'=1}^K s_k \lambda_{k'} \mu_k^T \mu_{k'} \right) + s_k \log \frac{\pi_k}{1 - \pi_k}$$

Also, because $s_k \in \{0, 1\}$ we know that $s_k^2 = s_k$:

$$s_k \log \frac{\lambda_k}{1 - \lambda_k} \propto -\frac{1}{2\sigma^2} \left(-2s_k \mathbf{x}^T \mu_k + s_k \mu_k^T \mu_k + 2 \sum_{k'=1, k' \neq k}^K s_k \lambda_{k'} \mu_k^T \mu_{k'} \right) + s_k \log \frac{\pi_k}{1 - \pi_k}$$

Because we have only kept terms with s_k , this is an equality:

$$s_k \log \frac{\lambda_k}{1 - \lambda_k} = \frac{s_k \mu_k^T}{2\sigma^2} \left(2\mathbf{x} - \mu_k - 2\sum_{k'=1, k' \neq k}^K \lambda_{k'} \mu_{k'} \right) + s_k \log \frac{\pi_k}{1 - \pi_k}$$

Solving for λ_k :

$$\lambda_k = \frac{1}{1 + \exp\left[-\left(\frac{\mu_k^T}{\sigma^2} \left(\mathbf{x} - \frac{\mu_k}{2} - \sum_{k'=1, k' \neq k}^K \lambda_{k'} \mu_{k'}\right) + \log\frac{\pi_k}{1 - \pi_k}\right)\right]}$$

we have our partial update for $q_k(s_k)$

(b)

The provided derivations for the M step of the mean parameter μ :

$$\mu = \left(\left\langle \mathbf{s} \mathbf{s}^T \right\rangle_{q(\mathbf{s})} \right)^{-1} \left\langle \mathbf{s} \right\rangle_{q(\mathbf{s})} \mathbf{x}$$

where $\mu \in \mathbb{R}^{K \times D}$, $\mathbf{s} \in \mathbb{R}^{K \times N}$, and $\mathbf{x} \in \mathbb{R}^{N \times D}$.

This mimics the least squares solution:

$$\hat{\beta} = (\mathbf{X}\mathbf{X}^T)^{-1}\mathbf{X}\mathbf{Y}$$

for the linear regression problem $\mathbf{Y} = \mathbf{X}^T \boldsymbol{\beta}$ where $\boldsymbol{\beta}$ corresponds to our mean parameters $\boldsymbol{\mu}$, the design matrix \mathbf{X} corresponds to our input \mathbf{s} and the response Y corresponds to the image pixels we denoted as \mathbf{x} . This makes sense because our resulting images \mathbf{x} are modeled as linear combinations of features $\boldsymbol{\mu}$, weighted by \mathbf{s} , $\mathbf{x} = \boldsymbol{\mu}\mathbf{s}$ where $\mathbf{x} \in \mathbb{R}^{D \times 1}$, $\boldsymbol{\mu} \in \mathbb{R}^{D \times K}$, and $\mathbf{s} \in \mathbb{R}^{K \times 1}$.

(c)

The computational complexity of the implemented M step function can be broken down for each parameter:

 μ : - The inversion ESS⁻¹ where ESS $\in \mathbb{R}^{K \times K}$ is $\mathcal{O}(K^3)$

- The dot product $ESS^{-1}ES^T$ where $ESS^{-1} \in \mathbb{R}^{K \times K}$ and $ES \in \mathbb{R}^{N \times K}$ is $\mathcal{O}(K^2N)$

- The dot product $(ESS^{-1}ES^T)\mathbf{x}$ where $(ESS^{-1}ES^T) \in \mathbb{R}^{K \times N}$ and $\mathbf{x} \in \mathbb{R}^{N \times D}$ is $\mathcal{O}(KND)$

 σ : - The dot product $(\mathbf{x}^T\mathbf{x})$ where $\mathbf{x} \in \mathbb{R}^{N \times D}$ is $\mathcal{O}(D^2N)$

- The dot product $\mu^T \mu$ where $\mu \in \mathbb{R}^{D \times K}$ is $\mathcal{O}(K^2 D)$

– The dot product $(\mu^T \mu)$ ESS where $\mu^T \mu \in \mathbb{R}^{K \times K}$ and ESS $\in \mathbb{R}^{K \times K}$ is $\mathcal{O}(K^3)$

- The dot product $ES^T \mathbf{x}$ where $ES \in \mathbb{R}^{N \times K}$ and $\mathbf{x} \in \mathbb{R}^{N \times D}$ is $\mathcal{O}(KND)$

- The dot product $(ES^T\mathbf{x})\mu$ where $ES^T\mathbf{x} \in \mathbb{R}^{K \times D}$ and $\mu \in \mathbb{R}^{D \times K}$ is $\mathcal{O}(K^2D)$

 π : - The mean operation for ES $\in \mathbb{R}^{N \times K}$ along the first dimension is $\mathcal{O}(NK)$

Thus, the computational complexity of the M step is $\mathcal{O}(K^3 + K^2N + KND + D^2N + K^2D)$ where we do not assume that any of N, K, or D is large compared to the others.

(d)

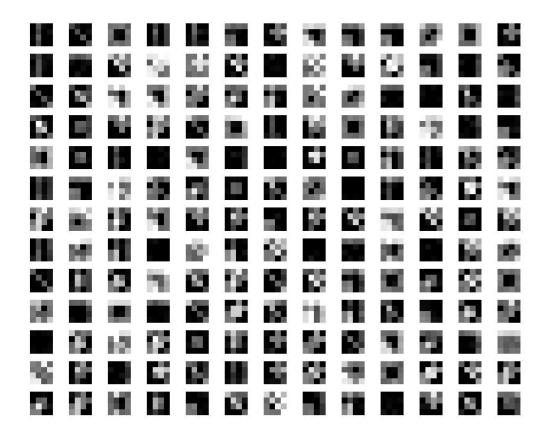


Figure 13: Images generated by randomly combined features with noise Examining the generated images, we can see eight features:

- (1) a cross
- (2) a border
- (3) a two by two square in the middle
- (4) a two by two square in the bottom left corner
- (5) a diagonal from top left to bottom right
- (6) a vertical line in the second column
- (7) a vertical line in the fourth column
- (8) a horizontal line in the first row

Factor analysis assumes a model:

$$\mathbf{x} = \mathbf{W}\mathbf{s} + \epsilon$$

where $\epsilon \sim \mathcal{N}(0, \Psi)$ and $\mathbf{s} \sim \mathcal{N}(0, \mathbf{I})$. Factor analysis would be inappropriate for this data because the our latent variables are binary (i.e. whether or not a feature is present) and not Gaussians.

A mixture of Gaussians assumes as model:

$$\mathbf{x} = \sum_{k=1}^{K} s_k \mu_k + \epsilon \text{ and } \sum_{k=1}^{K} s_k = 1$$

where $\epsilon \sim \mathcal{N}(0, \Sigma_{\epsilon})$. This also wouldn't be appropriate because it assumes a dirichlet distribution on the vector **s**. Our data is constructed by having each s_k independently sampled from a Bernoulli distribution indicating the presence of feature μ_k .

Independent component analysis assumes a model:

$$\mathbf{x} = \mathbf{W}\mathbf{s} + \epsilon$$

where $\epsilon \sim \mathcal{N}(0, \sigma^2 \mathbf{I})$ and $P(\mathbf{s}) = \prod_{k=1}^K P(s_k)$. This is appropriate for our data because $P(s_k)$ are our independent Bernoulli distributions and we are linearly combining different features with $\mathbf{W} \in \mathbb{R}^{D \times K}$ and then adding noise ϵ .

Thus, it would be expected that ICA does a good job modelling this data while factor analysis and mixture of Gaussians models would not.

(e)

We can plot the free energy at each EM step to make sure it increases each iteration:

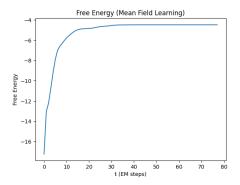


Figure 14: Free Energy

(f)

The initialised features:



Figure 15: Initial Latent Factors

The features learned by the algorithm:

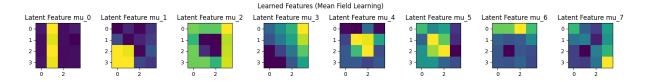


Figure 16: Learned Latent Factors

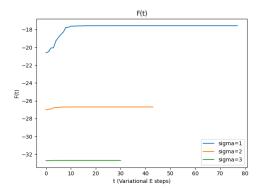
We can see that this approach has learned some of the previously identified features, such as the vertical line in the second column, the two by two square in the bottom left corner, the border, the horizontal line in the first row, and the a two by two square in the middle. The other features seem to be some linear combination of two or more features, such as μ_4 which looks like a combination of the cross and two by two square in the middle.

A possible way to improve our algorithm is reinitialising our algorithm a few times to find better potential convergence results (i.e. choose the model with the highest free energy convergence). We can also increase the number of data samples, with the hopes of learning better features. Finally, we can perform Variational Bayes by setting priors on π , σ^2 , and μ for better estimation of these parameters.

When implementing the algorithm, the mean field parameters were initialised randomly, each independently from a uniform distribution on [0,1]. π , σ , and μ were initialised by running a maximisation step using these randomly initialised mean field parameters. K was set to eight, after visually identifying eight features in part d. Moreover, for computational stability, when calculating $\log(\lambda_i)$ and $\log(1-\lambda_i)$, we manually shifted λ_i by $1e^{-10}$ to prevent $\lambda_i = 0$ or $\lambda_i = 1$.

(g)

Plotting the free energy at each partial expectation step of the variational approximation for different σ 's:



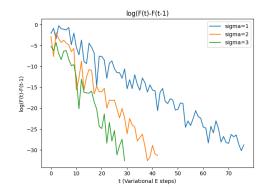


Figure 17: Free energy vs σ

Figure 18: Free energy convergence vs σ

We know that our free energy is a upper bounded on the log likelihood:

$$\log P(\mathcal{X}|\theta) \ge \mathcal{F}(q,\theta)$$

In the variational expectation step, $\log P(\mathcal{X}|\theta)$ is fixed by the parameters π , σ , and μ and we adjust our approximation q with parameter λ to try to reach this upper bound by increasing $\mathcal{F}(q,\theta)$. We know that σ quantifies the noise of \mathbf{x} , thus a higher σ means a wider spread in our distribution $\log P(\mathcal{X}|\theta)$, meaning we are reducing our upper bound for $\mathcal{F}(q,\theta)$. As such, we can see in the plot for free energy above that when σ is increased, our free energy converges to a lower value, due to being bounded above by a lower log-likelihood. Moreover, by reducing the upper bound, we see in the plot of $\log(F(t) - F(t-1))$ that our free energy is able to converge faster. Because we have reduced the upper bound by increasing σ , our free energy reaches the upper bound faster.

The Python code for the binary latent factor model:

```
from typing import Tuple
     import numpy as np
     from demo_code.m_step import m_step
           src.models.binary_latent_factor_approximations.abstract_binary_latent_factor_approximation import (
         AbstractBinaryLatentFactorApproximation\;,\\
            rc.models.binary_latent_factor_models.abstract_binary_latent_factor_model import (
         AbstractBinaryLatentFactorModel\;,\\
12
13
14
     {\bf class}\ Binary Latent Factor Model (\,Abstract Binary Latent Factor Model):
15
16
         def __init__(
self,
17
              mu: np.ndarray,
18
19
              sigma: float
              pi: np.ndarray,
              21
22
24
          @property
         def mu(self):
26
27
              return self._mu
29
         @mu.setter
30
         def mu(self, value):
              self._mu = value
32
33
         @property
34
         def sigma(self):
35
              return self._sigma
36
         @sigma.setter
def sigma(self, value):
    self._sigma = value
38
39
40
41
         @property
         def pi(self):
43
               return self._pi
44
         @pi.setter
         def pi(self, value):
    self._pi = value
46
47
49
         @property
         def variance(self) -> float:
return self.sigma**2
50
51
54
         {\tt def} \ \ {\tt calculate\_maximisation\_parameters} \, (
              x: np.ndarray
              binary_latent_factor_approximation: AbstractBinaryLatentFactorApproximation,
57
58
         ) \rightarrow Tuple [np.ndarray, float, np.ndarray]:
              return m_step(
59
60
                   es{=}binary\_latent\_factor\_approximation.expectation\_s\ ,
61
                   ess=binary_latent_factor_approximation.expectation_ss,
64
         def maximisation_step (
              self,
              x: np.ndarray,
66
67
              binary_latent_factor_approximation: AbstractBinaryLatentFactorApproximation,
68
         ) -> None:
              mu, sigma, pi = self.calculate_maximisation_parameters(
70
71
72
                   x, binary_latent_factor_approximation
              self.mu = mu
73
74
              self.sigma = sigma
              self.pi = pi
     def init_binary_latent_factor_model(
          \begin{array}{lll} \textbf{x:} & \textbf{np.ndarray}\,, \\ \textbf{binary\_latent\_factor\_approximation:} & \textbf{AbstractBinaryLatentFactorApproximation}\,, \end{array} 
79
80
     ) -> BinaryLatentFactorModel:
82
         Initialise by running a maximisation step with the parameters of the binary latent factor approximation
83
         :param binary_latent_factor_approximation: a binary_latent_factor_approximation :return: an initialised binary latent factor model """
          :param x: data matrix (number_of_points, number_of_dimensions)
85
86
         \begin{array}{lll} mu, & sigma\,, & pi &= BinaryLatentFactorModel.\, calculate\_maximisation\_parameters\,(\\ & x\,, & binary\_latent\_factor\_approximation \end{array}
88
89
         return BinaryLatentFactorModel(mu, sigma, pi)
```

src/models/binary_latent_factor_models/binary_latent_factor_model.py

The Python code for mean field learning:

```
from typing import List
 3
     import numpy as np
     from src.models.binary_latent_factor_approximations.abstract_binary_latent_factor_approximation import (
          AbstractBinaryLatentFactorApproximation,
     from src.models.binary_latent_factor_models.binary_latent_factor_model import (
          AbstractBinaryLatentFactorModel,
10
11
13
14
     {\bf class} \quad {\bf Mean Field Approximation (Abstract Binary Latent Factor Approximation):}
          def __init__(
               self, lambda_matrix: np.ndarray, max_steps: int, convergence_criterion: float
16
17
               self._lambda_matrix = lambda_matrix
               self.max_steps = max_steps
self.convergence_criterion = convergence_criterion
19
20
21
22
          def lambda_matrix(self) -> np.ndarray:
23
24
               lambda_matrix: parameters variational approximation (number_of_points, number_of_latent_variables)
25
26
               return self._lambda_matrix
27
28
          @lambda_matrix.setter
29
          def lambda_matrix(self, value):
30
               self._lambda_matrix = value
          {\tt def\ lambda\_matrix\_exclude(self\ ,\ exclude\_latent\_index:\ int)\ ->\ np.\,ndarray:}
               # (number_of_points, number_of_latent_variables-1)
return np.concatenate(
34
35
                         self.lambda_matrix[:, :exclude_latent_index],
self.lambda_matrix[:, exclude_latent_index + 1 :],
36
38
39
                     axis=1.
41
          def _partial_expectation_step(
42
               self,
44
               x: np.ndarray
               binary_latent_factor_model: AbstractBinaryLatentFactorModel,
45
               latent_factor: int,
          ) -> np.ndarray:
""" Partial Variational E step for factor i for all data points
47
48
49
               :param x: data matrix (number_of_points, number_of_dimensions)
:param binary_latent_factor_model: a binary_latent_factor_model
:param latent_factor: latent factor to compute partial update
:return: lambda_vector: new lambda parameters for the latent factor (number_of_points, 1)
"""
50
51
               lambda_matrix_excluded = self.lambda_matrix_exclude(latent_factor)
56
               mu\_excluded \ = \ binary\_latent\_factor\_model. \, mu\_exclude \, (\, latent\_factor \, )
58
               \verb|mu_latent| = \verb|binary_latent_factor_model.mu[:, latent_factor]|
59
               # (number_of_points, 1)
partial_expectation_log_p_x_given_s_theta_proportion = (
61
                    \verb|binary_latent_factor_model.precision|
62
                         63
65
66
67
                    @ mu_latent # (number_of_dimensions, 1)
               )
70
71
72
73
74
75
76
               # (1, 1)
partial_expectation_log_p_s_given_theta_proportion = np.log(
    binary_latent_factor_model.pi[0, latent_factor]
    / (1 - binary_latent_factor_model.pi[0, latent_factor])
               partial_expectation_log_p_x_s_given_theta_proportion = (
    partial_expectation_log_p_x_given_s_theta_proportion
80
                    + partial_expectation_log_p_s_given_theta_proportion
81
83
84
               # (number_of_points, 1)
               lambda_vector = 1 / (
    1 + np.exp(-partial_expectation_log_p_x_s_given_theta_proportion)
86
               [lambda\_vector[lambda\_vector == 0] = 1e-10
87
               lambda\_vector[lambda\_vector == 1] = 1 - 1e-10
89
               return lambda_vector
90
91
          def variational_expectation_step (
          self , x: np.ndarray , binary_latent_factor_model: AbstractBinaryLatentFactorModel
) -> List[float]:
92
                   'Variational E step
```

```
96
97
                   :param binary_latent_factor_model: a binary_latent_factor_model
                   :param x: data matrix (number_of_points, number_of_dimensions)
 98
                  99
101
102
103
105
                                    self.compute_free_energy(x, binary_latent_factor_model)
106
107
108
                                \begin{array}{lll} \textbf{if} & \texttt{free\_energy} \, [\, -1] \, - \, \, \texttt{free\_energy} \, [\, -2] \, <= \, \, \texttt{self.convergence\_criterion} \, : \end{array} 
109
110
                          \text{if } \text{free\_energy} \left[-1\right] \ - \ \text{free\_energy} \left[-2\right] \ <= \ \text{self.convergence\_criterion} : 
\begin{array}{c} 111 \\ 112 \end{array}
                  break
return free_energy
113
114
       def init_mean_field_approximation (
116
117
       k: int, n: int, max_steps, convergence_criterion ) -> MeanFieldApproximation:
             return MeanFieldApproximation (
119
                  \label{eq:lambda_matrix} \begin{array}{l} \texttt{lambda_matrix} = \texttt{np.random.random} \left( \, \, \texttt{size} = (\texttt{n} \,, \  \, \texttt{k} \,) \, \right) \,, \end{array}
120
                   max_steps=max_steps,
                   convergence_criterion=convergence_criterion,
```

src/models/binary_latent_factor_approximations/mean_field_approximation.py

The Python code for expectation maximisation:

```
from __future__ import annotations
 3
     from typing import List, Tuple
     import numpy as np
     from src.models.binary_latent_factor_approximations.abstract_binary_latent_factor_approximation import (
          AbstractBinaryLatentFactorApproximation\;,\\
            src.models.binary_latent_factor_models.binary_latent_factor_model import (
11
          AbstractBinaryLatentFactorModel,
12
13
14
     def is_converge(
          free_energies: List[float],
current_lambda_matrix: np.ndarray,
previous_lambda_matrix: np.ndarray,
16
17
19
     ) -> bool:
20
21
          Check for convergence of free energy and lambda matrix
22
23
          :param free_energies: list of free energies
          :param current_lambda_matrix: current lambda matrix
:param previous_lambda_matrix: previous lambda matrix
24
25
          :return: boolean indicating convergence
26
27
           \begin{array}{lll} \textbf{return} & (abs(free\_energies[-1] - free\_energies[-2]) == 0) \ and \ np. \\ linalg.norm(current\_lambda\_matrix - previous\_lambda\_matrix \end{array} 
28
29
30
          ) == 0
     def learn_binary_factors(
34
          x: np.ndarray,
em_iterations: int,
35
          binary_latent_factor_model: AbstractBinaryLatentFactorModel, binary_latent_factor_approximation: AbstractBinaryLatentFactorApproximation,
36
     ) -> Tuple [
AbstractBinaryLatentFactorApproximation,
38
39
40
          AbstractBinaryLatentFactorModel,
41
          List [float],
42
     1:
44
          Expectation maximisation algorithm to learn binary factors.
45
          :param x: data matrix (number_of_points, number_of_dimensions)
          :param em_iterations: number of iterations to run EM
:param binary_latent_factor_model: a binary_latent_factor_model
:param binary_latent_factor_approximation: a binary_latent_factor_approximation
47
48
49
          :return: a Tuple containing the updated binary_latent_factor_model, updated binary_latent_factor_approximation,
50
                      and free energies during each step of EM
          free\_energies: List[float] = [
54
               binary\_latent\_factor\_approximation.compute\_free\_energy (
                    x, binary_latent_factor_model
57
58
          for _ in range (em_iterations):
59
               previous_lambda_matrix = np.copy(
                    \verb|binary_latent_factor_approximation.lambda_matrix|
               )
61
               # E step
               binary_latent_factor_approximation.variational_expectation_step(
65
                    binary_latent_factor_model=binary_latent_factor_model,
66
               )
68
               # M step
69
                binary_latent_factor_model.maximisation_step(
71
72
73
74
75
                     binary_latent_factor_approximation,
               free_energies.append(
76
77
78
                    binary_latent_factor_approximation.compute_free_energy(
                        x, binary_latent_factor_model
79
80
               if is_converge(
                     free_energies ,
82
                     binary_latent_factor_approximation.lambda_matrix,
                    previous_lambda_matrix ,
83
85
          {\bf return \ binary\_latent\_factor\_approximation \ , \ binary\_latent\_factor\_model \ , \ free\_energies \ }
```

src/expectation_maximisation.py

The rest of the Python code for question 3:

```
from typing import List
     import matplotlib.pyplot as plt
     import numpy as np
     from src.expectation_maximisation import is_converge, learn_binary_factors
     from src.models.binary_latent_factor_approximations.mean_field_approximation import (
          MeanFieldApproximation, init_mean_field_approximation,
10
     from src.models.binarv_latent_factor_models.binarv_latent_factor_model import (
12
          AbstractBinaryLatentFactorModel,
13
14
          init_binary_latent_factor_model,
15
16
17
     def e_and_f(
          x: np.ndarray,
          k: int,
19
20
          em_iterations: int,
          e_maximum_steps:
          e_convergence_criterion: float.
22
23
          save_path: str,
24
     ) \ -\!\!\!> \ [\,AbstractBinaryLatentFactorModel\,\,,\,\,\,MeanFieldApproximation\,] \, :
25
          n = x.shape[0]
26
          mean_field_approximation = init_mean_field_approximation (
27
               k, n, max_steps=e_maximum_steps, convergence_criterion=e_convergence_criterion
28
29
          binary_latent_factor_model = init_binary_latent_factor_model(
30
               x, mean_field_approximation
          fig, ax = plt.subplots(1, k, figsize=(k * 2, 2))
          fig , ax = pit.subplots(i, k, figstze=(k * 2, 2))
for i in range(k):
    ax[i].imshow(binary_latent_factor_model.mu[:, i].reshape(4, 4))
    ax[i].set_title(f"Latent Feature mu-{i}")
fig.suptitle("Initial Features (Mean Field Learning)")
plt.tight_layout()
34
36
38
          plt.savefig(save_path + "-init-latent-factors", bbox_inches="tight")
39
          plt.close()
40
41
                mean_field_approximation
42
                binary_latent_factor_model .
                free_energy ,
44
          ) = learn_binary_factors (
45
               х.
                em_iterations,
               binary_latent_factor_model,
binary_latent_factor_approximation=mean_field_approximation,
47
48
49
          fig , ax = plt.subplots(1, k, figsize=(k * 2, 2))
for i in range(k):
    ax[i].imshow(binary_latent_factor_model.mu[:, i].reshape(4, 4))
50
51
          ax[i].set_title(f"Latent Feature mu_{i}")
fig.suptitle("Learned Features (Mean Field Learning)")
          plt.tight_layout()
          plt.savefig(save_path + "-latent-factors", bbox_inches="tight")
56
58
59
          plt.title("Free Energy (Mean Field Learning)")
          plt.xlabel("t (EM steps)")
plt.ylabel("Free Energy")
60
61
          plt.plot(free_energy)
62
          plt.savefig(save_path + "-free-energy", bbox_inches="tight")
          plt.close()
65
          return binary_latent_factor_model, mean_field_approximation
66
67
    def g(
    x: np.ndarray,
    binary_latent_factor_model: AbstractBinaryLatentFactorModel,
    mean_field_approximation: MeanFieldApproximation,
    -:rmas: List[float],
68
70
73
74
75
76
          save_path: str,
     ) -> None:
free_energies = []
77
78
79
          for sigma in sigmas:
               binary_latent_factor_model.sigma = sigma
mean_field_approximation_single_point = MeanFieldApproximation(
80
                     lambda\_matrix = mean\_field\_approximation. \\ lambda\_matrix \\ [:1\ , \ :]\ ,
                     max_steps=mean_field_approximation.max_steps
81
                     convergence_criterion=mean_field_approximation.convergence_criterion,
83
84
               free_energy: List[float] = [
                     mean_field_approximation_single_point.compute_free_energy(
86
                         x, binary_latent_factor_model
89
               for _ in range(em_iterations):
                     free_energy.pop(-1)
previous_lambda_matrix = np.copy(
90
91
                          mean_field_approximation_single_point.lambda_matrix
92
                     new_free_energy = (
```

```
mean\_field\_approximation\_single\_point.\ variational\_expectation\_step\ (
                                     binary_latent_factor_model=binary_latent_factor_model,
 97
                                     x=x,
 98
 99
                         free_energy.extend(new_free_energy)
                        if (
    free_energy[-1] - free_energy[-2]
    <= mean_field_approximation_single_point.convergence_criterion</pre>
101
103
                              \operatorname{free\_energy}.\operatorname{pop}(-1)
106
                              break
107
                         if is_converge (
108
                              free_energy ,
mean_field_approximation_single_point .lambda_matrix ,
109
110
                              previous_lambda_matrix,
\begin{array}{c} 111 \\ 112 \end{array}
                        ):
113
                   free_energies.append(free_energy)
114
115
             for i, free_energy in {\tt enumerate}({\tt free\_energies}):
                   plt.plot(
free_energy,
label=f"sigma={sigmas[i]}",
116
117
119
             plt.title(f"F(t)")
plt.xlabel("t (Variational E steps)")
plt.ylabel("F(t)")
plt.tight_layout()
120
123
124
             plt.legend()
             plt.savefig(save_path + f"-free-energy-sigma.png", bbox_inches="tight") plt.close()
126
127
             for i, free_energy in enumerate(free_energies):
    diffs = np.log(np.diff(free_energy))
128
130
                   plt.plot(
131
                         diffs
                         label=f"sigma={sigmas[i]}",
133
             )    plt.title(f"\log(F(t)-F(t-1)"))    plt.xlabel("t (Variational E steps)")    plt.ylabel("\log(F(t)-F(t-1)"))    plt.tight_layout()
134
135
136
137
138
             plt.legend()
             plt.savefig(save_path + f"-free-energy-diff-sigma.png", bbox_inches="tight")
             plt.close()
140
```

src/solutions/q3.py

Question 4: Variational Bayes for binary factors

(a)

To derive a Variational Bayesian hyperparameter optimisation algorithm to automatically determine K, the number of hidden binary variables in this model, we begin by writing the expression for x_d :

$$P(x_d|\mathbf{s}, \mathbf{w}_d, \sigma^2) = \mathcal{N}\left(\mathbf{s}^T \mathbf{w}_d, \sigma^2\right)$$

where we know from the diagonal covariance of $P(\mathbf{x}|\mathbf{s}, \mu, \sigma^2)$ that each dimension is independent. Moreover, $\mathbf{w}_d \in \mathbb{R}^{K \times 1}$, which is the d^{th} row of $\mu \in \mathbb{R}^{D \times K}$

Thus, we can express the posterior as:

$$\log P(\mathbf{x}, \mathbf{s}, \mu | \pi, \sigma^2, \alpha) = \log P(\mathbf{s} | \pi) + \sum_{d=1}^{D} \log P(x_d | s, \mathbf{w}_d, \sigma^2) + \log P(\mathbf{w}_d | \alpha)$$

where we introduce priors on each \mathbf{w}_d with $\alpha \in \mathbb{R}^{K \times 1}$. We choose each prior to be:

$$P(\mathbf{w}_d|\alpha) = \mathcal{N}(0, \mathbf{A}^{-1})$$

where $\mathbf{A} = diag(\alpha)$, the precision matrix. Combining, we have our expression:

$$\log P(\mathbf{x}, \mathbf{s}, \mu | \pi, \sigma^2, \alpha) = \sum_{d=1}^{D} \frac{-1}{2} \log(2\pi\sigma^2) - \frac{1}{2\sigma^2} \left(x_d^2 - 2x_d \mathbf{s}^T \mathbf{w}_d + \mathbf{w}_d^T \mathbf{s} \mathbf{s}^T \mathbf{w}_d \right) \\ + \sum_{k=1}^{K} s_k \log \pi_k + (1 - s_k) \log(1 - \pi_k) \\ + \sum_{d=1}^{D} \left(\frac{-K}{2} \log(2\pi) + \frac{1}{2} \sum_{k=1}^{K} (\log \alpha_k) - \frac{1}{2} \mathbf{w}_d^T \mathbf{A} \mathbf{w}_d \right)$$

For the Variational Bayes expectation step, we minimise $\mathbf{KL}[q_s(\mathbf{s}|\text{everything else})||P(\mathbf{s}|\text{everything else})]$ by setting:

$$q_s(\mathbf{s}) \propto \exp \left\langle \log P(\mathbf{x}, \mathbf{s}, \mu | \pi, \sigma^2, \alpha) \right\rangle_{q(\mu)}$$

Substituting the relevant terms:

$$q_s(\mathbf{s}) \propto \exp\left\{-\frac{1}{2\sigma^2}\left(-2\mathbf{x}^T\sum_{k=1}^K s_k \mu_k + \sum_{k=1}^K \sum_{k'=1, k' \neq k}^K s_k s_{k'} \mu_k^T \mu_{k'} + \sum_{k=1}^K s_k \mu_k^T \mu_k\right) + \sum_{k=1}^K s_k \log\frac{\pi_k}{1 - \pi_k}\right\}_{q(\mu)}$$

Given our factored approximation $q(\mathbf{s}) = \prod_{k=1}^{K} q_k(s_k)$, we can see that we can derive a similar partial update for $q_k(s_k)$ as in Question 3, by taking the variation derivative of the Lagrangian to enforce the normalisation of q_k :

$$\frac{\partial}{\partial q_k} \left(\exp \left\langle \log P(\mathbf{x}, \mathbf{s}, \mu | \pi, \sigma^2, \alpha) \right\rangle_{q(\mu)} + \lambda^{LG} \int q_k - 1) \right) \propto \exp \left\langle \log P(\mathbf{x}, \mathbf{s}, \mu | \pi, \sigma^2, \alpha) \right\rangle_{q(\mu) \prod_{k' \neq k} q_{k'}(s_k)} - \log q_k(s_k)$$

Setting this to zero we can solve for λ_i where $q_k(s_k) = \lambda_k^{s_k} (1 - \lambda_k)^{(1-s_k)}$:

$$\lambda_k = \frac{1}{1 + \exp\left[-\left(\frac{\langle \mu_k \rangle_{q_{\mu_k}}^T}{\sigma^2} \left(\mathbf{x} - \frac{\langle \mu_k \rangle_{q_{\mu_k}}}{2} - \sum_{k'=1, k' \neq k}^K \lambda_{k'} \langle \mu_k \rangle_{q_{\mu_k}}\right) + \log\frac{\pi_k}{1 - \pi_k}\right)\right]}$$

we have our partial E step update.

For the maximisation step, we perform maximisation steps for the parameters σ and π in the same way as question 3. However, having defined a prior on μ (through **w**) we will have to derive our expression for $\langle \mu_k \rangle_{q_{\mu_k}}$ the expectation of the posterior on μ_k . This involves deriving the posterior distribution of \mathbf{w}_d

$$q_{\mathbf{w}_d}(\mathbf{w}_d) \propto P(\mathbf{w}_d) \exp \left\langle \log P(\mathbf{x}, \mathbf{s}, \mu | \pi, \sigma^2, \alpha) \right\rangle_{q_{\mathbf{s}}(\mathbf{s})q_{\neg \mathbf{w}}}$$

Substituting the appropriate terms:

$$q_{\mathbf{w}_d}(\mathbf{w}_d) \propto \exp\left(-\frac{1}{2}\mathbf{w}_d^T \mathbf{A} \mathbf{w}_d\right) \exp\left\langle-\frac{1}{2\sigma^2} \left(-2x_d \mathbf{s}^T \mathbf{w}_d + \mathbf{w}_d^T \mathbf{s} \mathbf{s}^T \mathbf{w}_d\right)\right\rangle_{q_{\mathbf{s}}(\mathbf{s})q_{\neg \mathbf{w}_d}}$$

Simplifying:

$$q_{\mathbf{w}_d}(\mathbf{w}_d) \propto \exp\left(-\frac{1}{2}\left(\mathbf{w}_d^T\left(\mathbf{A} + \frac{\langle \mathbf{s}\mathbf{s}^T\rangle_{q_{\mathbf{s}}(\mathbf{s})}}{\sigma^2}\right)\mathbf{w}_d - 2\left(\frac{x_d\langle \mathbf{s}^T\rangle_{q_{\mathbf{s}}(\mathbf{s})}}{\sigma^2}\right)\mathbf{w}_d\right)\right)$$

We see that the posterior:

$$q_{\mathbf{w}_d}(\mathbf{w}_d) = \mathcal{N}\left(\mu_{\mathbf{w}_d}, \Sigma_{\mathbf{w}_d}\right)$$

where:

$$\Sigma_{\mathbf{w}_d} = \left(\frac{\langle \mathbf{s}\mathbf{s}^T \rangle_{q_{\mathbf{s}}(\mathbf{s})}}{\sigma^2} + \mathbf{A}\right)^{-1}$$

and

$$\mu_{\mathbf{w}_d} = \Sigma_{\mathbf{w}_d} \left(\frac{x_d \langle \mathbf{s}^T \rangle_{q_{\mathbf{s}}(\mathbf{s})}}{\sigma^2} \right)$$

Thus, $\langle \mu_k \rangle_{q_{\mu_k}} \in \mathbb{R}^{D \times 1}$ is the concatenation of the k^{th} elements of $\mu_{\mathbf{w}_d} \in \mathbb{R}^{K \times 1}$ for $d \in \{1, ..., D\}$ For ARD, we must also optimise α with a hyper-M step. We start by choosing $Ga(\alpha_k|a,b)$, a Gamma prior on α_k , with a and b being hyperparameters. Thus, to optimise α we want to maximise the penalised objective:

$$\alpha = \arg \max_{\alpha} \left\langle \log P(\mathbf{x}, \mathbf{s}, \mu | \pi, \sigma^2, \alpha) \right\rangle_{q(\mathbf{w})} + \sum_{k=1}^{K} \log P(\alpha_k | a, b)$$

Substituting the appropriate terms, we have our penalised objective Q:

$$Q = \left\langle \sum_{d=1}^{D} \frac{1}{2} \sum_{k=1}^{K} (\log \alpha_k) - \frac{1}{2} \mathbf{w}_d^T \mathbf{A} \mathbf{w}_d \right\rangle_{q(\mathbf{w})} + \sum_{k=1}^{K} (a-1) \log \alpha_k - b\alpha_k$$

Simplifying:

$$Q = \frac{D}{2} \sum_{k=1}^{K} (\log \alpha_k) - \frac{1}{2} \sum_{d=1}^{D} \left(tr \left[\mathbf{A} \left\langle \mathbf{w}_d \mathbf{w}_d^T \right\rangle_{q(\mathbf{w}_d)} \right] \right) + \sum_{k=1}^{K} (a-1) \log \alpha_k - b\alpha_k$$

Setting $\frac{dQ}{d\alpha_k} = 0$ we get:

$$\frac{D}{2\alpha_k} - \frac{1}{2} \sum_{d=1}^{D} \left\langle (w_{d,k})^2 \right\rangle_{q(\mathbf{w}_d)} + \frac{a-1}{\alpha_k} - b = 0$$

where $w_{d,k}$ is the k^{th} element of \mathbf{w}_d . Knowing $\langle (w_{d,k})^2 \rangle_{q(\mathbf{w}_d)} = (\mu_{\mathbf{w}_{d,k}})^2 + \Sigma_{\mathbf{w}_{d,(k,k)}}$, the k^{th} element of $\mu_{\mathbf{w}_d}$ and element (k,k) of $\Sigma_{\mathbf{w}_d}$ respectively, we can solve for α_k :

$$\alpha_k = \frac{2a + D - 2}{2b + \sum_{d=1}^{D} \left((\mu_{\mathbf{w}_{d,k}})^2 + \sum_{\mathbf{w}_{d,(k,k)}} \right)}$$

we have our hyper-M steps for optimising α .

(b)

Running variational Bayes for different values of k, we can visualise the learned features μ_k and corresponding α_k^{-1} :

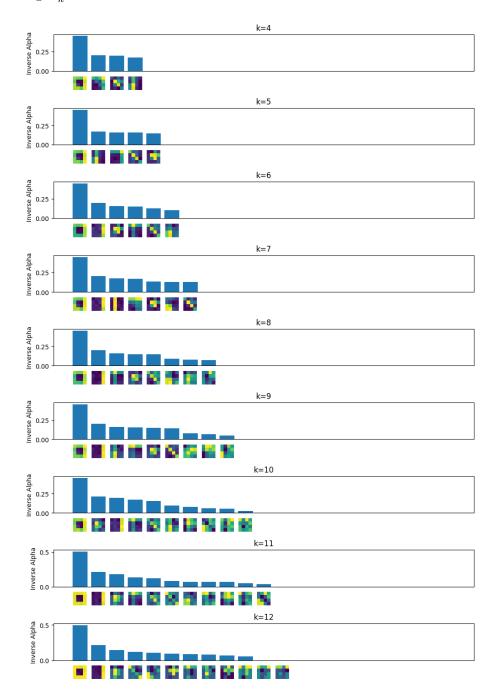


Figure 19: Learned Latent Factors vs Inverse Alpha

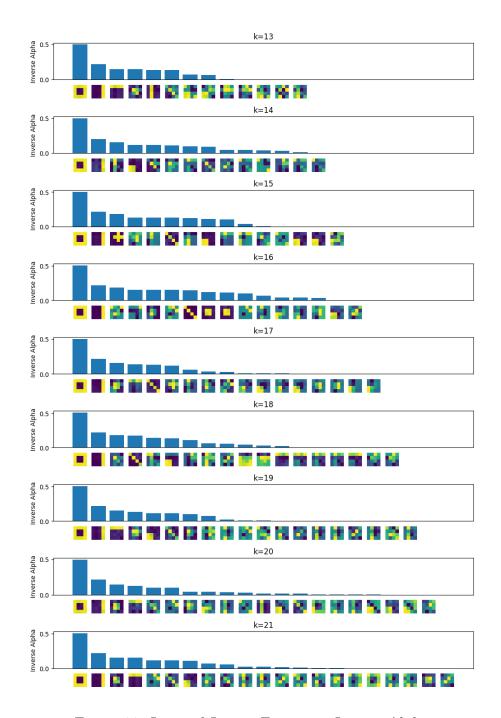


Figure 20: Learned Latent Factors vs Inverse Alpha

As we expect, when running the algorithm for higher K values, many of the features have $\alpha_k \to \infty$, depicted as $\alpha_k^{-1} \to 0$ for visual convenience. Moreover, visualising the learned features, we can see the clearest features often have the highest α_k^{-1} while the features deemed irrelevant are often noisy or duplicates.

Comparing the free energy plots of models trained on different K values:

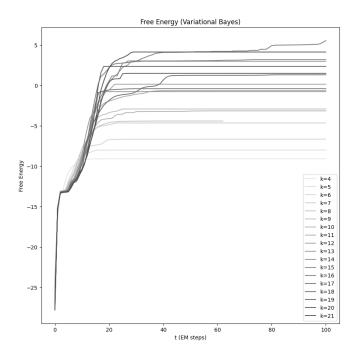


Figure 21: Free Energy for different values of k

We can see that initially for K=4 to K=8, increasing K significantly increases the convergence value of the free energy. However, beyond K=8, the trend of K versus the free energy convergence value is not as clear. We can see that this corresponds to the visualisation of α^{-1} where beyond K=11, the number of relevant features remains more or less the same. We know that there are only eight latent features, thus models with K>8 should be learning duplicate or irrelevant features. As such, we wouldn't expect a model to be able to increase it's free energy significantly when provided with additional degrees of freedom by increasing the value of K beyond eight. We see that for models with K>>8, there are typically ten or eleven features that might be deemed relevant (depending on how you threshold) and this is likely from slight overfitting, noise in the data, or duplicate features. Thus, the relationship between the free energy and the effective number of latent features for each model is as we would expect with ARD.

The Python code for Variational Bayes:

```
import numpy as np
 3
     from src.models.binary_latent_factor_approximations.abstract_binary_latent_factor_approximation import (
          AbstractBinaryLatentFactorApproximation,
          src.models.binary_latent_factor_models.binary_latent_factor_model import (BinaryLatentFactorModel,
     from src.models.binary_latent_factor_models.boltzmann_machine import BoltzmannMachine
10
11
12
     class Gaussian Prior:
13
14
         def __init__(self , a: float , b: float , d: int , k: int):
               Gaussian prior on mu matrix
16
17
               :param a: alpha parameter of Gamma Prior
               :param b: beta parameter of Gamma Prior
               :param d: number of dimensions
:param k: number of latent variables
19
20
21
22
               self.a = a
23
               self.b = b
               24
25
26
27
28
          \begin{array}{lll} \textbf{def} & \textbf{mu\_k} \big(\, \textbf{self} \,\,, \,\,\, \textbf{k} \colon \,\, \textbf{int} \,\big) \,\, -\!\!\!> \,\, \textbf{np.ndarray} \,\colon \\ \end{array}
29
30
               Column vector of mu matrix, the latent feature vector
               : \mathtt{param} \ k \colon \ \mathtt{latent} \ \mathtt{factor} \ \mathtt{index}
               :return: column vector (number_of_dimensions, 1)
34
35
               \textcolor{return}{\texttt{return}} \hspace{0.2cm} \texttt{self.mu[:, k : k + 1]}
36
          def w_d(self , d: int) -> np.ndarray:
38
               Row vector of mu matrix, the weight vector for a particular dimension (pixel) of the data
39
40
               :param d: data dimension index
:return: row vector (1, number_of_latent_variables)
"""
41
42
44
               return self.mu[d : d + 1, :]
45
          def a_matrix(self) -> np.ndarray:
47
48
49
               Precision matrix for a weight vector w_d
               : return: \ matrix \ of \ shape \ ( \ number\_of\_latent\_variables \ , \ number\_of\_latent\_variables )
50
51
               return np.diag(self.alpha)
55
     class VariationalBayes (BoltzmannMachine):
56
          Variational Bayes implementation with prior on mu. Note that we are inheriting from BoltzmannMachine for Question 5d only.
58
59
60
               :param mu: Gaussian prior on latent features
:param variance: Gaussian noise parameter
:param pi: vector of priors (1, number_of_latent_variables)
61
62
65
               super().__init__(mu=mu.mu, sigma=np.sqrt(variance), pi=pi)
               self.gaussian_prior = mu
66
67
               self._variance = variance
self._pi = pi
70
71
72
          @property
          def variance (self) -> float:
              return self._variance
73
74
75
76
          def pi(self) -> np.ndarray:
    return self._pi
77
78
79
          @property
def mu(self) -> np.ndarray:
80
               return self.gaussian_prior.mu
81
          def _update_w_d_covariance(
83
84
               binary_latent_factor_approximation: AbstractBinaryLatentFactorApproximation,
                   (number_of_latent_variables, number_of_latent_variables)
86
87
               self.gaussian_prior.w_covariance = np.linalg.inv(
                   + self.precision * binary_latent_factor_approximation.expectation_ss
89
90
91
          def _update_w_d_mean (
92
94
               x: np.ndarray, # (number_of_points, number_of_dimensions)
```

```
binary_latent_factor_approximation: AbstractBinaryLatentFactorApproximation,
            ) \rightarrow None:
 97
 98
 99
                  Update mean vector for w_d.
101
                  :param x: data matrix (number_of_points, number_of_dimensions)
                  :param binary_latent_factor_approximation: a binary_latent_factor_approximation :param d: index of data dimension to update
103
106
                  # (number_of_latent_variables, 1)
self.gaussian_prior.mu[d : d + 1, :] = (
    self.gaussian_prior.w_covariance
108
109
                        @ (
110
                              self.\ precision \\ *\ binary_latent_factor_approximation.\ expectation\_s.T \\ @\ x[:,\ d\ :\ d\ +\ 1]
113
114
115
116
            \begin{array}{ll} \textbf{def} & \texttt{-hyper\_maximisation\_step} \, (\, \texttt{self} \, ) \, \, -\!\!\!> \, \, \texttt{None} \colon \\ \end{array}
117
119
                  Hyper M step updating alpha, which parameterize the covariance matrix of the Gaussian prior on mu
120
                  for k in range(self.k):
                        123
124
126
127
128
            def maximisation_step (
                  self,
130
                  x: np.ndarray
                  binary\_latent\_factor\_approximation: \ AbstractBinaryLatentFactorApproximation\;,
                  distribution of mu as well as a hyper M-step followed by posterior updates to the distribution of mu as well as a hyper M-step updating the prior parameters on mu, the alpha vector param x: data matrix (number_of_points, number_of_dimensions) :param binary_latent_factor_approximation: a binary_latent_factor_approximation
                  Maximisation step which runs the usual M-step followed by posterior updates to the
134
135
136
                  \verb|--, sigma|, pi = Binary Latent Factor Model. calculate \verb|-maximisation \>--parameters| (
                        x, binary_latent_factor_approximation
140
                  self._variance = sigma**2
142
                  self._pi = pi
self._update_w_d_covariance(binary_latent_factor_approximation)
144
145
                   \begin{array}{lll} & for \ d \ in \ range(self.d): \\ & self.\_update\_w\_d\_mean(x, \ binary\_latent\_factor\_approximation \,, \ d) \end{array} 
                  self._hyper_maximisation_step()
147
```

src/models/binary_latent_factor_models/variational_bayes.py

The Python code for Automatic Relevance Determination:

```
from typing import List, Tuple
3
4
     import numpy as np
     from src.expectation_maximisation import learn_binary_factors
    from \ src.models.binary\_latent\_factor\_approximations.abstract\_binary\_latent\_factor\_approximation \ import \ (AbstractBinaryLatentFactorApproximation\ ,
          src.models.binary_latent_factor_models.binary_latent_factor_model import (
10
         {\tt BinaryLatentFactorModel}\ ,
11
12
     from src.models.binary_latent_factor_models.variational_bayes import (
13
14
         Gaussian Prior,
         VariationalBayes,
15
16
17
18
19
     def run_automatic_relevance_determination (
         x: np.ndarray, binary_latent_factor_approximation: AbstractBinaryLatentFactorApproximation,
20
21
22
         b_parameter: float,
23
         k: int,
24
25
         em_iterations: int
     ) -> Tuple [VariationalBayes, List [float]]:
26
27
28
         Run automatic relevance determination with variational Bayes.
29
         :param x: data matrix (number_of_points, number_of_dimensions)
30
         :param binary_latent_factor_approximation: a binary_latent_factor_approximation
31
         :param a_parameter: alpha parameter for gamma prior :param b_parameter: beta parameter for gamma prior
33
34
         :param k: number of latent variables
:param em_iterations: number of iterations to run EM
         return: a Tuple containing the optimised VB model and a list of free energies during each EM step
35
36
         (_, sigma, pi,) = BinaryLatentFactorModel.calculate_maximisation_parameters(
38
              x, binary_latent_factor_approximation
39
         mu = Gaussian Prior (
41
42
              a=a_parameter ,
              b=b_parameter,
              k=k ,
              d=x . shape [1] ,
44
45
         variational_bayes_model: VariationalBayes = VariationalBayes(
              mu=mu,
47
48
              variance=sigma * * 2,
49
              _{\rm p\,i=p\,i}\;,
51
         , variational_bayes_model, free_energy = learn_binary_factors(
52
53
54
              em_iterations=em_iterations
              binary_latent_factor_model=variational_bayes_model,
55
              binary_latent_factor_approximation=binary_latent_factor_approximation,
56
         return variational_bayes_model, free_energy
```

src/automatic_relevance_determination.py

The rest of the Python code for question 4:

```
from typing import List
     import matplotlib.pyplot as plt
     import numpy as np from matplotlib.offsetbox import AnnotationBbox, OffsetImage
     from src.automatic_relevance_determination import run_automatic_relevance_determination
     from src.models.binary_latent_factor_approximations.mean_field_approximation import (
          init_mean_field_approximation,
     from src.models.binary_latent_factor_models.variational_bayes import VariationalBayes
13
14
     def offset_image(coord: int, path: str, ax: plt.axis):
16
17
          Add image to matplotlib axis.
19
          : \verb"param" coord": coordinate on axis"
20
          :param path: path to image
          :param ax: plot axis
21
22
          img = plt.imread(path)
im = OffsetImage(img, zoom=0.72)
23
24
          im.image.axes = ax
25
26
27
28
          ab = AnnotationBbox(
               im, (coord, 0), xybox = (0.0, -19.0),
29
30
                frameon=False,
xycoords="data"
                boxcoords="offset points",
                pad=0,
36
          ax.add_artist(ab)
39
     def plot_factors (
40
           variational_bayes_binary_latent_factor_models: List[VariationalBayes],
41
          \mathtt{ks} : \ \mathtt{List} \ [\ \mathbf{int}\ ] \ ,
42
          max_k: int,
          save_path: str ,
44
45
          # store each feature as an image for later use
          for i, k in enumerate(ks):
sort_indices = np.argsort(
variational_bayes_binary_latent_factor_models[i].gaussian_prior.alpha
47
48
49
                for j, idx in enumerate(sort_indices):
fig = plt.figure(figsize=(0.3, 0.3))
ax = plt.Axes(fig, [0.0, 0.0, 1.0, 1.0])
50
51
                     ax.set_axis_off()
fig.add_axes(ax)
                     ax.imshow(
56
                          variational_bayes_binary_latent_factor_models[i]
58
                          .reshape (4, 4)
59
                     fig.savefig(save\_path + f"-latent-factor-{i}-{j}", bbox\_inches="tight")
61
                     plt.close()
62
63
          The property of alphas fig, ax = plt. subplots (len(ks), 1, figsize = (12, 2 * len(ks))) plt.subplots_adjust(hspace=1)
65
          for i, k in enumerate(ks):
    sort_indices = np.argsort(
        variational_bayes_binary_latent_factor_models[i].gaussian_prior.alpha
66
67
                y = list(
70
71
72
73
74
75
76
                     / variational_bayes_binary_latent_factor_models[i].gaussian_prior.alpha[
                          sort_indices
               ) + [0] * (max_k - k)
ax[i].set_title(f"{k=}")
ax[i].bar(range(max_k), y)
               ax[i].set_xticks([])
ax[i].set_ylabel("Inverse Alpha")
          # add feature image ticks
for i, k in enumerate(ks):
    sort_indices = np.argsort(
80
81
83
                     variational_bayes_binary_latent_factor_models[i].gaussian_prior.alpha
84
                for j in range(len(sort_indices)):
86
                     path = save_path + f"-latent
                                                           factor -{ i }-{ j }.png"
                     offset_image(j, path, ax[i])
os.remove(path)
           fig.savefig(save_path + f"-latent-factors-comparison", bbox_inches="tight")
89
90
          plt.close()
91
92
     def b(
        x: np.ndarray,
```

```
a_parameter: int ,
b_parameter: int ,
 97
            ks: List[int],
            max_k: int,
 98
 99
             em_iterations: int
100
            e_maximum_steps: int ,
101
             e_convergence_criterion: float,
       save_path: str,
) -> List[List[float]]:
103
             variational_bayes_models: List[VariationalBayes] = []
106
             free_energies = []
for i, k in enumerate(ks):
108
                  n = x.shape[0]
mean_field_approximation = init_mean_field_approximation(
109
110
\begin{array}{c} 111 \\ 112 \end{array}
                        n,
                        max_steps=e_maximum_steps,
113
                        convergence_criterion=e_convergence_criterion,
114
115
                  (variational_bayes_model, free_energy) = run_automatic_relevance_determination(
116
                        x \!\!=\! x\,, binary_latent_factor_approximation=mean_field_approximation ,
117
                        a_parameter=a_parameter,
119
                        b\_parameter = b\_parameter \;,
120
                        k=k.
                        {\tt em\_iterations} {=} {\tt em\_iterations} \ ,
                  /
variational_bayes_models.append(variational_bayes_model)
free_energies.append(free_energy)
123
124
             plot_factors (
126
                  variational_bayes_models ,
127
                  ks,
                  max_k.
128
                  save_path ,
130
             return free_energies
131
133
134
       def free_energy_plot(
135
            ks: List[int], free_energies: List[List[float]], model_name: str, save_path: str
       ):
136
             fig = plt.figure()
137
            fig = plt.figure()
fig.set_figwidth(10)
fig.set_figheight(10)
shades = np.flip(np.linspace(0.3, 0.9, len(ks)))
for i, k in enumerate(ks):
    plt.plot(free_energies[i], label=f"{k=}", color=np.ones(3) * shades[i])
plt.title(f"Free Energy ({model_name})")
plt.xlabel("t (EM steps)")
plt.ylabel("Free Energy")
plt.ylabel("Free Energy")
140
142
144
145
             plt.legend()
             plt.savefig(save_path + "-free-energy", bbox_inches="tight")
147
148
             plt.close()
```

src/solutions/q4.py

Question 5: EP for the binary factor model

(a)

To derive an EP algorithm to infer marginals on the source variables in the binary latent factor model from Question 3, we first write the log-joint probability for a single observation-source pair:

$$\log p(\mathbf{s}, \mathbf{x}) = \log p(\mathbf{s}) + (\mathbf{x}|\mathbf{s})$$

Knowing $p(\mathbf{s}) = \prod_{i=1}^K p(s_i|\pi_i)$ and $p(\mathbf{x}|\mathbf{s}) = \mathcal{N}(\sum_{i=1}^K s_i \mu_i, \sigma^2 \mathbf{I})$:

$$\log p(\mathbf{s}, \mathbf{x}) \propto \frac{-1}{2} \left(\mathbf{x} - \sum_{i=1}^{K} s_i \mu_i \right)^T \frac{1}{\sigma^2} \mathbf{I} \left(\mathbf{x} - \sum_{i=1}^{K} s_i \mu_i \right) + \sum_{i=1}^{K} \left(s_i \log \pi_i + (1 - s_i) \log(1 - \pi_i) \right)$$

Expanding:

$$\log p(\mathbf{s}, \mathbf{x}) \propto \frac{-1}{2\sigma^2} \left(\mathbf{x}^T \mathbf{x} - 2\mathbf{x}^T \sum_{i=1}^K s_i \mu_i + \sum_{i=1}^K \sum_{j=1}^K s_i s_j \mu_i^T \mu_j \right) + \sum_{i=1}^K \left(s_i \log \pi_i + (1 - s_i) \log(1 - \pi_i) \right)$$

Collecting terms pertaining to s_i :

$$\log p(\mathbf{s}, \mathbf{x}) = \sum_{i=1}^{K} \left(\left(\frac{\mathbf{x}^T \mu_i}{\sigma^2} + \log \frac{\pi_i}{1 - \pi_i} \right) s_i \right) + \sum_{i=1}^{K} \sum_{j=1}^{K} \left(\frac{-\mu_i^T \mu_j}{2\sigma^2} s_i s_j \right) + C$$

where C are all other terms without s_i .

Knowing that $s_i^2 = s_i$:

$$\log p(\mathbf{s}, \mathbf{x}) = \sum_{i=1}^{K} \left(\left(\frac{\mathbf{x}^T \mu_i}{\sigma^2} + \log \frac{\pi_i}{1 - \pi_i} - \frac{\mu_i^T \mu_i}{2\sigma^2} \right) s_i \right) + \sum_{i=1}^{K} \sum_{j=1}^{i-1} \left(\frac{-\mu_i^T \mu_j}{\sigma^2} s_i s_j \right) + C$$

Thus:

$$\log p(\mathbf{s}, \mathbf{x}) = \sum_{i=1}^{K} \log f_i(s_i) + \sum_{i=1}^{K} \sum_{j=1}^{i-1} \log g_{ij}(s_i, s_j)$$

where the factors are defined:

$$\log f_i(s_i) = \left(\frac{\mathbf{x}^T \mu_i}{\sigma^2} + \log \frac{\pi_i}{1 - \pi_i} - \frac{\mu_i^T \mu_i}{2\sigma^2}\right) s_i$$

and

$$\log g_{ij}(s_i, s_j) = \frac{-\mu_i^T \mu_j}{\sigma^2} s_i s_j$$

as required.

The Boltzmann Machine can be defined as:

$$P(\mathbf{s}|\mathbf{W}, \mathbf{b}) = \frac{1}{Z} \exp\left(\sum_{i=1}^{K} \sum_{j=1}^{i-1} W_{ij} s_i s_j - \sum_{i=1}^{K} b_i s_i\right)$$

where $s_i \in \{0, 1\}$, the same as our source variables.

From our factorisation, we can see that $p(\mathbf{s}, \mathbf{x})$ is a Boltzmann Machine with:

$$W_{ij} = \frac{-\mu_i^T \mu_j}{\sigma^2}$$

and

$$b_i = -\left(\frac{\mathbf{x}^T \mu_i}{\sigma^2} + \log \frac{\pi_i}{1 - \pi_i} - \frac{\mu_i^T \mu_i}{2\sigma^2}\right)$$

and

$$\log Z = -C$$

(b)

For $f_i(s_i)$, we will choose a Bernoulli approximation:

$$\tilde{f}_i(s_i) = (\theta_{ii})^{s_i} + (1 - \theta_{ii})^{1 - s_i}$$

Thus,

$$\log \tilde{f}_i(s_i) \propto \log \left(\frac{\theta_{ii}}{1 - \theta_{ii}}\right) s_i$$

we can define $\eta_{ii} = \log \left(\frac{\theta_{ii}}{1 - \theta_{ii}} \right)$:

$$\log \tilde{f}_i(s_i) \propto \eta_{ii} s_i$$

For $g_{ij}(s_i, s_j)$, we can choose a product of Bernoulli distributions for our approximation:

$$\tilde{g}_{ij}(s_i, s_j) = \tilde{g}_{ij, \neg s_j}(s_i) \tilde{g}_{ij, \neg s_i}(s_j)$$

where

$$\tilde{g}_{ij,\neg s_j}(s_i) = (\theta_{ji})^{s_i} + (1 - \theta_{ji})^{1 - s_i}$$

and

$$\tilde{g}_{ij,\neg s_i}(s_j) = (\theta_{ij})^{s_j} + (1 - \theta_{ij})^{1-s_j}$$

Thus.

$$\log \tilde{g}_{ij}(s_i, s_j) \propto \log \left(\frac{\theta_{ji}}{1 - \theta_{ji}}\right) s_i + \log \left(\frac{\theta_{ij}}{1 - \theta_{ij}}\right) s_j$$

we can define $\eta_{ji} = \log\left(\frac{\theta_{ji}}{1-\theta_{ji}}\right)$ and $\eta_{ij} = \log\left(\frac{\theta_{ij}}{1-\theta_{ij}}\right)$:

$$\log \tilde{g}_{ij}(s_i, s_j) \propto \eta_{ji} s_i + \eta_{ij} s_j$$

To derive a message passing scheme, we define the incoming message to node i from the singleton factor:

$$\mathcal{M}_{i \to i}(s_i) = \tilde{f}_i(s_i)$$

and the incoming message to node i from node j:

$$\mathcal{M}_{j\to i}(s_i) = \sum_{s_1 \in \{0,1\}} \cdots \sum_{s_{i-1} \in \{0,1\}} \sum_{s_{i+1} \in \{0,1\}} \cdots \sum_{s_1 \in \{0,1\}} \tilde{g}_{ji}(s_j, s_i) \mathcal{M}_{j\to j}(s_j) \prod_{k \in ne(j), k \neq i}^K \mathcal{M}_{k\to j}(s_j)$$

where ne(j) are indices of neighbouring nodes of node j. This is the product of the incoming messages to node j from its neighbours, the singleton message from node j, and $\tilde{g}_{ji}(s_j, s_i)$ with all nodes except s_i marginalised out.

Because $\tilde{g}_{ji}(s_j, s_i)$ is a product:

$$\mathcal{M}_{j \to i}(s_i) = \tilde{g}_{ji, \neg s_j}(s_i) \sum_{s_1 \in \{0,1\}} \cdots \sum_{s_{i-1} \in \{0,1\}} \sum_{s_{i+1} \in \{0,1\}} \cdots \sum_{s_1 \in \{0,1\}} \tilde{f}_j(s_j) \tilde{g}_{ji, \neg s_i}(s_j) \prod_{k \in ne(j), k \neq i}^K \mathcal{M}_{k \to j}(s_j)$$

We are left with:

$$\mathcal{M}_{j\to i}(s_i) = \tilde{g}_{ji,\neg s_j}(s_i)$$

and

$$\mathcal{M}_{j\to i}(s_i) \propto \exp\left(\eta_{ji}s_i\right)$$

The cavity distributions are:

$$q_{\neg \tilde{f}_i(s_i)}(s_i) = \prod_{j \in ne(i)}^K \mathcal{M}_{j \to i}(s_i)$$

and

$$q_{\neg \tilde{g}_{ij}(s_i, s_j)}(s_i, s_j) = \left(\mathcal{M}_{i \to i}(s_i) \prod_{k \in ne(i), k \neq j}^K \mathcal{M}_{k \to i}(s_i)\right) \left(\mathcal{M}_{j \to j}(s_j) \prod_{k \in ne(j), k \neq i}^K \mathcal{M}_{k \to j}(s_j)\right)$$

For $\tilde{f}_i(s_i)$, we do not need to make an approximation step. This is because we are minimising:

$$\tilde{f}_i(s_i) = \arg\min_{\tilde{f}_i(s_i)} \mathbf{KL} \left[f_i(s_i) q_{\neg \tilde{f}_i(s_i)}(s_i) \| \tilde{f}_i(s_i) q_{\neg \tilde{f}_i(s_i)}(s_i) \right]$$

We know that the factor $\log f_i(s_i)$ is a Bernoulli of the form $b_i s_i$. Because our approximation for this site is also Bernoulli, we can simply solve for θ_{ii} in $\log \tilde{f}_i(s_i)$:

$$\log \tilde{f}_i(s_i) = \log f_i(s_i)$$
$$\log \left(\frac{\theta_{ii}}{1 - \theta_{ii}}\right) s_i = b_i s_i$$
$$\theta_{ii} = \frac{1}{1 + \exp(-b_i)}$$

On the other hand, for $\tilde{g}_{ij}(s_i, s_j)$, we will approximate with:

$$\tilde{g}_{ij}(s_i, s_j) = \arg\min_{\tilde{g}_{ij}(s_i, s_j)} \mathbf{KL} \left[g_{ij}(s_i, s_j) q_{\neg \tilde{g}_{ij}(s_i, s_j)}(s_i, s_j) \left\| \tilde{g}_{ij}(s_i, s_j) q_{\neg \tilde{g}_{ij}(s_i, s_j)}(s_i, s_j) \right\| \right]$$

We can define natural parameters $\eta_{i,\neg s_{j}}$ and $\eta_{j,\neg s_{i}}$ for $q_{\neg \tilde{g}_{ij}(s_{i},s_{j})}(s_{i},s_{j})$ such that:

$$\mathcal{M}_{i \to i}(s_i) \prod_{k \in ne(i), k \neq j}^{K} \mathcal{M}_{k \to i}(s_i) \propto \exp(\eta_{i, \neg s_j} s_i)$$

$$\mathcal{M}_{j o j}(s_j) \prod_{k \in ne(j), k \neq j}^K \mathcal{M}_{k o j}(s_j) \propto \exp(\eta_{j, \neg s_i} s_j)$$

Note that $\tilde{g}_{ij}(s_i, s_j)$ was chosen as the product of two Bernoulli distributions, so updates to this site approximation involves updating the parameters η_{ji} and η_{ij} , for s_i and s_j respectively.

We can write:

$$\log\left(\tilde{g}_{ij}(s_i,s_j)q_{\neg\tilde{q}_{ij}(s_i,s_j)}(s_i,s_j)\right) \propto \eta_{ji}s_i + \eta_{ij}s_j + \eta_{i,\neg s_i}s_i + \eta_{j,\neg s_i}s_j$$

Simplifying:

$$\log\left(\tilde{g}_{ij}(s_i, s_j) q_{\neg \tilde{g}_{ij}(s_i, s_j)}(s_i, s_j)\right) \propto \left(\eta_{ji} + \eta_{i, \neg s_j}\right) s_i + \left(\eta_{ij} + \eta_{j, \neg s_i}\right) s_j$$

Thus, the first moments:

$$\mathbb{E}_{s_i} \left[\sum_{s_j \in \{0,1\}} \tilde{g}_{ij}(s_i, s_j) q_{-\tilde{g}_{ij}(s_i, s_j)}(s_i, s_j) \right] = \frac{1}{1 + \exp\left(-\left(\eta_{ji} + \eta_{i, \neg s_j}\right)\right)}$$

and

$$\mathbb{E}_{s_j} \left[\sum_{s_i \in \{0,1\}} \tilde{g}_{ij}(s_i, s_j) q_{\neg \tilde{g}_{ij}(s_i, s_j)}(s_i, s_j) \right] = \frac{1}{1 + \exp\left(-\left(\eta_{ij} + \eta_{j, \neg s_i}\right)\right)}$$

Moreover:

$$\log\left(g_{ij}(s_i,s_j)q_{\neg\tilde{q}_{ij}(s_i,s_j)}(s_i,s_j)\right) \propto W_{ij}s_is_j + \eta_{i,\neg s_i}s_i + \eta_{j,\neg s_i}s_j$$

To derive the first moment for $g_{ij}(s_i, s_j)q_{\neg \tilde{g}_{ij}(s_i, s_j)}(s_i, s_j)$ with respect to s_i , we first marginalise out s_j :

$$\sum_{s_j \in \{0,1\}} g_{ij}(s_i, s_j) q_{\neg \tilde{g}_{ij}(s_i, s_j)}(\mathbf{s}) \propto \exp\left(W_{ij} s_i + \eta_{i, \neg s_j} s_i + \eta_{j, \neg s_i}\right) + \exp\left(\eta_{i, \neg s_j} s_i\right)$$

Thus, the first moment:

$$\mathbb{E}_{s_i} \left[\sum_{s_j \in \{0,1\}} g_{ij}(s_i, s_j) q_{\neg \tilde{g}_{ij}(s_i, s_j)}(s_i, s_j) \right] = \frac{\exp\left(W_{ij} + \eta_{i, \neg s_j} + \eta_{j, \neg s_i}\right) + \exp\left(\eta_{i, \neg s_j}\right)}{\left[\exp\left(W_{ij} + \eta_{i, \neg s_j} + \eta_{j, \neg s_i}\right) + \exp\left(\eta_{i, \neg s_j}\right)\right] + \left[\exp\left(\eta_{j, \neg s_i}\right) + 1\right]}$$

Simplifying:

$$\mathbb{E}_{s_i} \left[\sum_{s_j \in \{0,1\}} g_{ij}(s_i, s_j) q_{\neg \tilde{g}_{ij}(s_i, s_j)}(s_i, s_j) \right] = \frac{\exp\left(\eta_{i, \neg s_j}\right) \left(\exp\left(W_{ij} + \eta_{j, \neg s_i}\right) + 1\right)}{\left[\exp\left(\eta_{i, \neg s_j}\right) \left(\exp\left(W_{ij} + \eta_{j, \neg s_i}\right) + 1\right)\right] + \left[\exp\left(\eta_{j, \neg s_i}\right) + 1\right]}$$

Similarly:

$$\mathbb{E}_{s_{j}} \left[\sum_{s_{i} \in \{0,1\}} g_{ij}(s_{i}, s_{j}) q_{\neg \tilde{g}_{ij}(s_{i}, s_{j})}(s_{i}, s_{j}) \right] = \frac{\exp\left(\eta_{j, \neg s_{i}}\right) \left(\exp\left(W_{ij} + \eta_{i, \neg s_{j}}\right) + 1\right)}{\left[\exp\left(\eta_{j, \neg s_{i}}\right) \left(\exp\left(W_{ij} + \eta_{i, \neg s_{j}}\right) + 1\right)\right] + \left[\exp\left(\eta_{i, \neg s_{j}}\right) + 1\right]}$$

By setting:

$$\mathbb{E}_{s_i} \left[\sum_{s_j \in \{0,1\}} \tilde{g}_{ij}(s_i, s_j) q_{\neg \tilde{g}_{ij}(s_i, s_j)}(s_i, s_j) \right] = \mathbb{E}_{s_i} \left[\sum_{s_j \in \{0,1\}} g_{ij}(s_i, s_j) q_{\neg \tilde{g}_{ij}(s_i, s_j)}(s_i, s_j) \right]$$

and

$$\mathbb{E}_{s_j} \left[\sum_{s_i \in \{0,1\}} \tilde{g}_{ij}(s_i, s_j) q_{\neg \tilde{g}_{ij}(s_i, s_j)}(s_i, s_j) \right] = \mathbb{E}_{s_j} \left[\sum_{s_i \in \{0,1\}} g_{ij}(s_i, s_j) q_{\neg \tilde{g}_{ij}(s_i, s_j)}(s_i, s_j) \right]$$

we can solve for the parameters of $\tilde{g}_{ij}(s_i, s_j)$ with moment matching:

$$\frac{1}{1+\exp\left(-\left(\eta_{ji}+\eta_{i,\neg s_{j}}\right)\right)}=\frac{\exp\left(\eta_{i,\neg s_{j}}\right)\left(\exp\left(W_{ij}+\eta_{j,\neg s_{i}}\right)+1\right)}{\left[\exp\left(\eta_{i,\neg s_{j}}\right)\left(\exp\left(W_{ij}+\eta_{j,\neg s_{i}}\right)+1\right)\right]+\left[\exp\left(\eta_{j,\neg s_{i}}\right)+1\right]}$$

Simplifying:

$$\exp(\eta_{j,\neg s_{i}}) + 1 = \exp\left(-\left(\eta_{ji} + \eta_{i,\neg s_{j}}\right)\right) \exp\left(\eta_{i,\neg s_{j}}\right) \left(\exp\left(W_{ij} + \eta_{j,\neg s_{i}}\right) + 1\right)$$

$$\frac{\exp(\eta_{j,\neg s_{i}}) + 1}{\exp(W_{ij} + \eta_{i,\neg s_{i}}) + 1} = \exp\left(-\eta_{ji}\right)$$

Our parameter update:

$$\eta_{ji} = \log \left(\frac{1 + \exp\left(W_{ij} + \eta_{j, \neg s_i}\right)}{1 + \exp\left(\eta_{i, \neg s_i}\right)} \right)$$

Similarly:

$$\eta_{ij} = \log \left(\frac{1 + \exp\left(W_{ij} + \eta_{i, \neg s_j}\right)}{1 + \exp\left(\eta_{i, \neg s_j}\right)} \right)$$

(c)

Using factored approximate messages, we see that:

$$\eta_{i,\neg s_j} = \log\left(\frac{\theta_{ii}}{1 - \theta_{ii}}\right) + \sum_{k \in ne(i), k \neq j}^{K} \log\left(\frac{\theta_{ki}}{1 - \theta_{ki}}\right)$$

Knowing $\eta_{ii} = \log \left(\frac{\theta_{ii}}{1 - \theta_{ii}} \right)$ and $\eta_{ki} = \log \left(\frac{\theta_{ki}}{1 - \theta_{ki}} \right)$:

$$\eta_{i,\neg s_j} = \eta_{ii} + \sum_{k \in ne(i), k \neq j}^K \eta_{ki}$$

and

$$\eta_{j,\neg s_i} = \eta_{jj} + \sum_{k \in ne(j), k \neq i}^K \eta_{kj}$$

The summation of the natural parameters of the singleton factor for node i with the natural parameters of messages from all the neighbouring nodes that aren't j.

This leads to a loopy BP algorithm because the nodes are fully connected (i.e. every node is the neighbour of all other nodes). Thus, we cannot simply move from one end of the graph to the other like BP for tree structured graphs.

Moreover, our factored approximations:

$$q(s_i) \propto \sum_{j=1}^K \eta_{ij} s_i$$

and so:

$$\lambda_i = \frac{1}{1 + \exp(-\sum_{j=1}^K \eta_{ij})}$$

our parameter for $q(s_i)$.

(d)

We can use automatic relevance determination (ARD) as a hyperparameter method to select relevant features by placing a prior on μ_k in the same way as Question 4. With a hyper-M step, certain features will have diverging precision, leaving us with the relevant features, from which we get our K. We can implement this by running the message passing algorithm (loopy BP) which was implemented for Question 6 for the expectation step and the VB algorithm for the maximisation step (along with its hyper-M Step) which was implemented for Question 4:

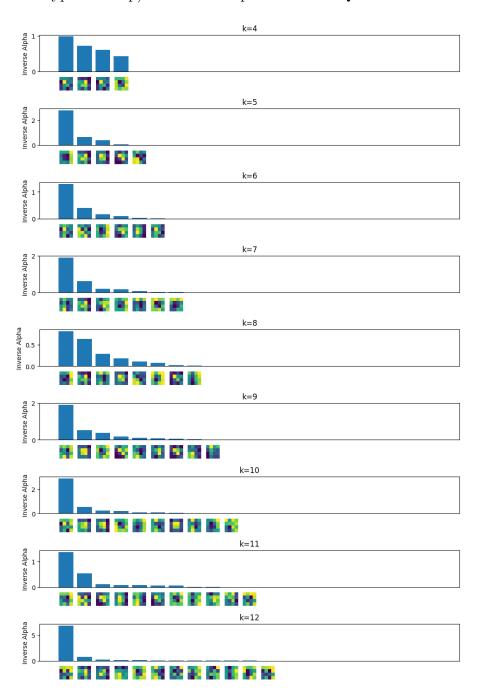


Figure 22: Learned Latent Factors vs Inverse Alpha

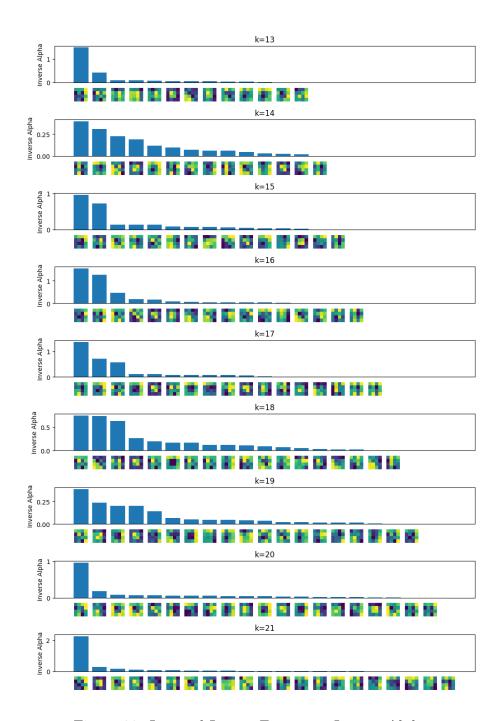


Figure 23: Learned Latent Factors vs Inverse Alpha

We can see that ARD has more difficulty finding relevant features when using loop BP. However, looking at the features themselves, we can see that the features are not very clear compared those from the mean field approximation or Variational Bayes in Question 4. This is probably why we do not consistently have the same number of α_k values that don't diverge as compared with Question 4, loopy BP seems to be unable to identify too many clear features that we would want to clearly deem relevant.

Comparing the free energy plots of models trained on different K values:

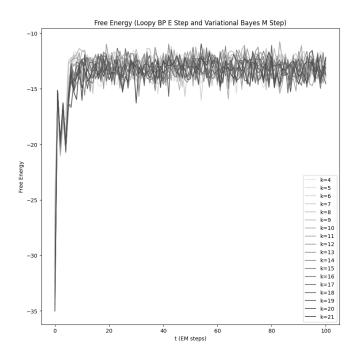


Figure 24: Free Energy for different values of k

We can see the difficulties of this approach due to the inability for loopy BP to converge for this particular problem. This likely contributes to the difficulty for ARD to find relevant features when the message passing algorithm is unable to converge on generating meaningful features. To improve this, we could also put priors on the other parameters σ^2 and π in the hopes that this will help stabilise the algorithm. Moreover, because loopy BP is unable to converge for this problem, it is quite computationally intensive as EM will always run the maximum number of iterations without improvement in the results. Knowing that it won't converge, employing early stopping or fewer iterations can also be helpful. Moreover, Boltzmann machines do not scale well with respect to the number of nodes, K, introducing computational difficulties. This was experienced when generating the above results, where for larger K values, loopy BP E steps took much longer to perform. To tackle this, a restricted Boltzmann Machine could be implemented to reduce the number of connections in the graph, alleviating these computational difficulties.

The Python code for question 5:

```
from typing import List
                  import numpy as np
from tqdm import tqdm
                  \begin{array}{lll} from & src.automatic\_relevance\_determination & import & run\_automatic\_relevance\_determination \\ from & src.models.binary\_latent\_factor\_approximations.message\_passing\_approximation & import & (instance of the context of the
                                   \verb|init_message_passing|,
                  from src.models.binary_latent_factor_models.variational_bayes import VariationalBayes from src.solutions.q4 import plot_factors
10
11
12
13
14
15
                  def d(
    x: np.ndarray,
                                 x: np.ndarray,
a_parameter: int,
b_parameter: int,
ks: List[int],
max_k: int,
em_iterations: int,
16
17
18
19
20
                 save_path: str,
) -> List[List[float]]:
21
23
24
25
                                    variational_bayes_models: List[VariationalBayes] = []
                                   free_energies = []
for i, k in tqdm(enumerate(ks)):
    n = x.shape[0]
    message_passing_approximation = init_message_passing(k, n)
    (variational_bayes_model, free_energy) = run_automatic_relevance_determination(
26
27
28
 29
30
31
                                                                     x = x \,, \\ binary\_latent\_factor\_approximation = message\_passing\_approximation \,, \\ a\_parameter = a\_parameter \,, \\ \\
33
34
                                                                      b\_parameter {=} b\_parameter \;,
                                                                     k=k,
                                                                      {\tt em\_iterations} {=} {\tt em\_iterations} \ ,
 36
                                                    variational_bayes_models.append(variational_bayes_model)
38
                                                    free_energies.append(free_energy)
                                    plot_factors (
                                                    variational_bayes_models,
41
42
                                                   ks,
max_k,
                                                    save_path,
 44
                                   return free_energies
```

src/solutions/q5.py

Question 6: EP/Loopy BP Implementation

Implementing the EP/loopy-BP algorithm, we can compare the learned latent factors with those of the variational mean-field algorithm:

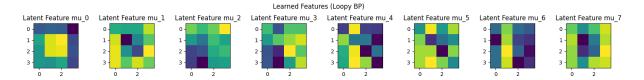


Figure 25: Learned Latent factors learned with EP/Loopy-BP

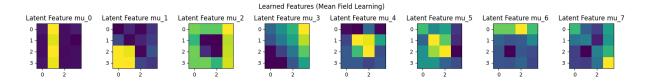
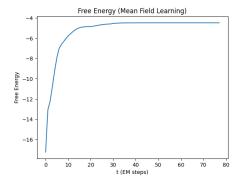


Figure 26: Learned Latent Factors with Mean Field Approximation

We can see that the mean field algorithm seems to learn better latent features. In particular loopy BP that has a few duplicates and some of the learned features are quite noisy. For example μ_0 for the mean field algorithm looks almost like a binary image whereas μ_3 and μ_5 from loopy BP are almost completely noise. We can understand the reason for this by comparing the free energies of the two algorithms:



Free Energy (Loopy BP)

-12

-14

-14

-18

-20

0 20 40 60 80 100

t (EM steps)

Figure 27: Mean Field Approximation

Figure 28: Loopy BP

We can observe that the free energy of the mean field algorithm converges while our loopy belief propagation is unable to converge to a free energy. Because loopy BP does not have convergence guarantees, one of the limitations of this approach, we can see that in this case loopy BP is unable to identify many latent factors.

The Python code for the Boltzmann machine:

```
import numpy as np
3
     from src.models.binary_latent_factor_approximations.abstract_binary_latent_factor_approximation import (
          AbstractBinaryLatentFactorApproximation,
           src.models.binary_latent_factor_models.binary_latent_factor_model import (
          BinaryLatentFactorModel,
10
11
     class BoltzmannMachine (BinaryLatentFactorModel):
          def __init__(
self,
12
13
14
              mu: np.ndarray,
sigma: float,
16
17
              pi: np.ndarray,
19
               Binary latent factor model with Boltzmann Machine terms
20
               super().__init__(mu, sigma, pi)
22
23
          @property
          def w_matrix(self) -> np.ndarray:
24
25
26
               Weight matrix of the Boltzmann machine
27
               : return: \ matrix \ of \ weights \ (number\_of\_latent\_variables \, , \ number\_of\_latent\_variables \, )
28
29
30
               return -self.precision * (self.mu.T @ self.mu)
          def w_matrix_index(self, i, j) \rightarrow float:
34
               Weight matrix at a specific index
36
               :param i: row index
               :param j: column index
               :return: weight value
39
40
               return -self.precision * (self.mu[:, i] @ self.mu[:, j])
41
          def b(self, x) \rightarrow np.ndarray:
42
44
              b term in the Boltzmann machine for all data points
45
               :param x: design matrix (number_of_points, number_of_dimensions)
              :return: matrix of shape (number_of_points, number_of_latent_variables)
47
48
                   irn -(
   self.precision * x @ self.mu
+ self.log_pi_ratio
- 0.5 * self.precision * np.multiply(self.mu, self.mu).sum(axis=0)
49
50
51
55
          def b_index(self, x, node_index) -> float:
56
               b term for a specific node in the Boltzmann machine for all data points
58
59
               :param x: design matrix (number_of_points, number_of_dimensions)
               :param node_index: node index
              :return: vector of shape (number_of_points, 1)
61
62
                   self.precision * x @ self.mu[:, node_index] + (self.log_pi[0, node_index] - self.log_one_minus_pi[0, node_index]) - 0.5 * self.precision * self.mu[:, node_index] @ self.mu[:, node_index]
67
               ).reshape(
70
71
72
73
74
75
76
          def log_pi_ratio(self) -> np.ndarray:
    return self.log_pi - self.log_one_minus_pi
     def init_boltzmann_machine (
     x: np.ndarray,
binary_latent_factor_approximation: AbstractBinaryLatentFactorApproximation,
) -> BinaryLatentFactorModel:
80
81
          Initialise by running a maximisation step with the parameters of the binary latent factor approximation
          : param \ x: \ data \ matrix \ (number\_of\_points \ , \ number\_of\_dimensions) \\ : param \ binary\_latent\_factor\_approximation : \ a \ binary\_latent\_factor\_approximation
83
84
          return: an initialised Boltzmann machine model
86
          mu, sigma, pi = BinaryLatentFactorModel.calculate_maximisation_parameters(
               x, binary_latent_factor_approximation
89
          return BoltzmannMachine(mu=mu, sigma=sigma, pi=pi)
```

src/models/binary_latent_factor_models/boltzmann_machine.py

The Python code for message passing:

```
from typing import List
 3
      import numpy as np
      from \  \  src.models.binary\_latent\_factor\_approximations.abstract\_binary\_latent\_factor\_approximation \  \  import \  \  (abstract\_binary\_latent\_factor\_approximation \  \  import \  \  )
            AbstractBinaryLatentFactorApproximation,
      from src.models.binary_latent_factor_models.boltzmann_machine import BoltzmannMachine
10
11
      {\bf class} \ \ {\bf MessagePassingApproximation} \ ({\bf \, AbstractBinaryLatentFactorApproximation}): \\
12
           bernoulli_parameter_matrix (theta): matrix of parameters bernoulli_parameter_matrix [n, i, j] off diagonals corresponds to \tilda\{g\}_{\{i\}}, \neq s_i\}(s_j) for data point n diagonals correspond to \tilda\{f\}_{\{i\}}(s_i) (number_of_points, number_of_latent_variables, number_of_latent_variables)
13
14
16
17
           def __init__(self, bernoulli_parameter_matrix: np.ndarray):
    self.bernoulli_parameter_matrix = bernoulli_parameter_matrix
19
20
21
22
23
            def \ lambda\_matrix(self) \rightarrow np.ndarray:
24
                  Aggregate messages and compute parameter for Bernoulli distribution
25
26
                  :return:
27
                   \begin{array}{ll} lambda\_matrix = 1 \ / \ (1 \ + \ np. \exp(- \, self. \, natural\_parameter\_matrix. \, sum(\, axis = 1))) \\ lambda\_matrix \, [\, lambda\_matrix \, = \, 0] \ = \ 1e - 10 \end{array} 
28
29
30
                  lambda_matrix[lambda_matrix == 1] = 1 - 1e-10
                  return lambda_matrix
            @property
34
            def natural_parameter_matrix(self) -> np.ndarray:
35
                 The matrix containing natural parameters (eta) of each factor off diagonals corresponds to \tilda\{g\}_{\{i\}}, \tilda\{g\}_{\{i\}} for data point n diagonals correspond to \tilda\{f\}_{\{i\}}
36
38
                              (number_of_points, number_of_latent_variables, number_of_latent_variables)
39
40
                  :return:
41
42
                  return np.log(
                       np. divide
44
                              self.bernoulli_parameter_matrix , 1 - self.bernoulli_parameter_matrix
45
47
48
           def aggregate_incoming_binary_factor_messages(
    self , node_index: int , excluded_node_index: int
49
            ) -> np.ndarray:
# (number_of_points,
50
51
                      exclude message from excluded_node_index -> node_index
                  return (
np.sum(
                              self.natural_parameter_matrix[:, :excluded_node_index, node_index],
56
                              axis=1,

/ np.sum(
self.natural_parameter_matrix[:, excluded_node_index + 1 :, node_index],

58
59
60
61
62
                  ).reshape(
63
65
66
            @staticmethod
67
            def calculate_bernoulli_parameter(
                  natural_parameter_matrix: np.ndarray,
            ) -> np.ndarray:
                  bernoulli_parameter = 1 / (1 + np.exp(-natural_parameter_matrix))
bernoulli_parameter[bernoulli_parameter == 0] = 1e-10
bernoulli_parameter[bernoulli_parameter == 1] = 1 - 1e-10
70
71
72
73
74
75
76
                  return bernoulli-parameter
           def variational_expectation_step(
    self, x: np.ndarray, binary_latent_factor_model: BoltzmannMachine
) -> List[float]:
77
78
79
                  Iteratively update singleton and binary factors
80
                  :param x: data matrix (number_of_points, number_of_dimensions)
:param binary_latent_factor_model: a binary_latent_factor_model
81
                  return: free energies after each update
83
84
                  free_energy = [self.compute_free_energy(x, binary_latent_factor_model)]
for i in range(self.k):
    # singleton factor update
86
                        natural_parameter_ii = self.calculate_singleton_message_update(
87
                              boltzmann_machine=binary_latent_factor_model,
89
                              x=x,
90
                              i=i,
91
                        self.bernoulli_parameter_matrix[
92
                        ] = self.calculate_bernoulli_parameter(natural_parameter_ii)
```

```
free_energy.append(self.compute_free_energy(x, binary_latent_factor_model))
97
                   for j in range(i):
98
                       # binary factor update
99
                       natural_parameter_ij = self.calculate_binary_message_update(
                            boltzmann_machine=binary_latent_factor_model,
                            x=x,
                            j=j ,
                       self.bernoulli_parameter_matrix[
                       :, i, j
] = self.calculate_bernoulli_parameter(natural_parameter_ij)
106
108
                       natural_parameter_ji = self.calculate_binary_message_update(
    boltzmann_machine=binary_latent_factor_model,
109
                            j=i,
113
                       self.bernoulli_parameter_matrix[
114
                         :, j, i
= self.calculate_bernoulli_parameter(natural_parameter_ji)
116
                       free_energy.append(
                            self.compute_free_energy(x, binary_latent_factor_model)
              return free_energy
120
         def calculate_binary_message_update(
123
              self.
124
              x: np.ndarray
              boltzmann\_machine:\ BoltzmannMachine\,,
126
              i: int,
j: int,
127
         ) -> float:
128
130
              Calculate new parameters for a binary factored message.
              :param x: data matrix (number_of_points, number_of_dimensions)
              :param boltzmann_machine: Boltzmann machine model
:param i: starting node for the message
:param j: ending node for the message
134
135
              return: new parameter from aggregating incoming messages
136
              natural\_parameter\_i\_not\_j \ = \ boltzmann\_machine.b\_index (
              x=x, node_index=i
) + self.aggregate_incoming_binary_factor_messages(
140
                  node_index=i, excluded_node_index=j
142
              w_i_j = boltzmann_machine.w_matrix_index(i, j)
              144
145
147
148
          @staticmethod
149
          def calculate_singleton_message_update(
              x: np.ndarray,
boltzmann_machine: BoltzmannMachine,
151
         ) -> float:
              Calculate the parameter update for the singleton message.
155
156
              Note that this does not require any approximation.
              :param x: data matrix (number_of_points, number_ot:param boltzmann_machine: Boltzmann machine model
158
                                                            number_of_dimensions)
              :param i: node to update
              :return: new parameter
161
162
              return boltzmann_machine.b_index(x=x, node_index=i)
164
165
166
     def init_message_passing(k: int, n: int) -> MessagePassingApproximation:
167
168
         Message passing initialisation
169
          :param k: number of latent variables
          :param n: number of data points
          :return: message passing
172
173
          bernoulli_parameter_matrix = np.random.random(size=(n, k, k))
         return MessagePassingApproximation(bernoulli_parameter_matrix)
```

src/models/binary_latent_factor_approximations/message_passing_approximation.py

The rest of the Python code for question 6:

```
import matplotlib.pyplot as plt import numpy as np
 3
4
       from src.expectation_maximisation import learn_binary_factors
       {\bf from } \  \  {\bf src.models.binary\_latent\_factor\_approximations.message\_passing\_approximation \ \ \underline{\bf import} \ \ (
             \verb"init_message_passing",
      from src.models.binary_latent_factor_models.boltzmann_machine import ( init_boltzmann_machine ,
10
11
12
13
14
       def run(x: np.ndarray, k: int, em_iterations: int, save_path: str) -> None:
             n = x.shape[0]
message\_passing = init\_message\_passing(k, n)
15
16
17
             boltzmann\_machine = init\_boltzmann\_machine(x, message\_passing)
18
19
             # pre-training features plot
fig , ax = plt.subplots(1, k, figsize=(k * 2, 2))
for i in range(k):
20
             ax[i].imshow(boltzmann_machine.mu[:, i].reshape(4, 4))
ax[i].set_title(f"Latent Feature mu_{i}")
fig.suptitle("Initial Features (Loopy BP)")
plt_tight_layout()
21
22
23
24
25
             plt.savefig(save_path + "-init-latent-factors", bbox_inches="tight")
26
             plt.close()
27
28
29
             message_passing, boltzmann_machine, free_energy = learn_binary_factors(
30
31
                    em_iterations=em_iterations
                    binary_latent_factor_model=boltzmann_machine,
32
33
34
                    \verb|binary_latent_factor_approximation=message\_passing|,
35
            # post training features plot
fig , ax = plt.subplots(1, k, figsize=(k * 2, 2))
for i in range(k):
    ax[i].imshow(boltzmann_machine.mu[:, i].reshape(4, 4))
    ax[i].set_title(f"Latent Feature mu_{i})
fig.suptitle("Learned Features (Loopy BP)")
plt.tight_layout()
plt.savefig(save_path + "-latent-factors", bbox_inches="tight")
plt_clase()
36
38
41
42
44
             plt.close()
45
            # free energy plot
plt.title("Free Energy (Loopy BP)")
plt.xlabel("t (EM steps)")
plt.ylabel("Free Energy")
47
48
49
             plt.plot(free_energy)
plt.savefig(save_path + "-free-energy", bbox_inches="tight")
50
51
             plt.close()
```

src/solutions/q6.py

Appendix 1: abstract_binary_latent_factor_model.py

```
from __future__ import annotations
      from abc import ABC, abstractmethod from typing import TYPE_CHECKING
       import numpy as np
       if TYPE_CHECKING:
             from \  \  src.models.binary\_latent\_factor\_approximations.abstract\_binary\_latent\_factor\_approximation \  \  import \  \  (abstract\_binary\_latent\_factor\_approximation \  \  import \  \  )
                   AbstractBinaryLatentFactorApproximation,
11
12
14
15
16
17
18
       class AbstractBinaryLatentFactorModel(ABC):
             @property
             \begin{array}{ll} \textbf{def} & \text{mu(self)} & -\!\!\!> & \text{np.ndarray:} \end{array}
19
                    matrix \ of \ means \ (number\_of\_dimensions \, , \ number\_of\_latent\_variables \, )
20
22
23
             @property
             @abstractmethod
25
26
             def variance(self) -> float:
                   gaussian noise parameter
28
29
                    pass
             @property
31
32
             @abstractmethod
33
             \begin{array}{ll} \textbf{def} & \text{pi(self)} & -\!\!\!> & \text{np.ndarray:} \end{array}
34
                   (1, number_of_latent_variables)
36
37
                    pass
39
             @abstractmethod
40
             def maximisation_step(
                   x: np.ndarray, binary_latent_factor_approximation: AbstractBinaryLatentFactorApproximation,
42
43
45
46
             def mu_exclude(self, exclude_latent_index: int) -> np.ndarray:
    return np.concatenate( # (number_of_dimensions, number_of_latent_variables -1)
        (self.mu[:, :exclude_latent_index], self.mu[:, exclude_latent_index + 1 :]),
47
48
50
51
                          axis=1,
53
54
             def log_pi(self) -> np.ndarray:
    return np.log(self.pi)
56
             @property
             def log_one_minus_pi(self) -> np.ndarray:
    return np.log(1 - self.pi)
60
61
             def precision(self) -> float:
    return 1 / self.variance
62
64
             @property
def d(self) -> int:
    return self.mu.shape[0]
65
67
68
             def k(self) -> int:
    return self.mu.shape[1]
70
71
```

src/models/binary_latent_factor_models/abstract_binary_latent_factor_model.py

Appendix 2: abstract_binary_latent_factor_approximation.py

```
from __future__ import annotations
     from abc import ABC, abstractmethod
     from typing import TYPE_CHECKING, List
     if TYPE CHECKING:
         from src.models.binary_latent_factor_models.binary_latent_factor_model import (
              AbstractBinaryLatentFactorModel\;,\\
11
12
13
     import numpy as np
14
15
     class AbstractBinaryLatentFactorApproximation(ABC):
         @property
16
17
18
         \begin{array}{lll} d\,ef & lambda\_matrix\,(\,\,s\,elf\,) \,\,\longrightarrow\,\, np\,.\,ndarray: \end{array}
19
              lambda_matrix: parameters variational approximation (number_of_points, number_of_latent_variables)
20
22
23
         @abstractmethod
24
         def variational_expectation_step (
25
26
              self,
              x: np.ndarray.
         binary_latent_factor_model: AbstractBinaryLatentFactorModel, ) -> List[float]:
28
29
              pass
30
31
         @property
32
         def expectation_s (self):
33
              return self.lambda_matrix
34
         def expectation_ss(self):
36
              ess = self.lambda_matrix.T @ self.lambda_matrix
37
              np.fill_diagonal(ess, self.lambda_matrix.sum(axis=0))
39
40
         def log.lambda_matrix(self) -> np.ndarray:
    return np.log(self.lambda_matrix)
42
43
44
45
46
         def log_one_minus_lambda_matrix(self) -> np.ndarray:
47
48
              return np.log(1 - self.lambda_matrix)
49
         @property
50
51
         def n(self) \rightarrow int:
              Number of data points
54
              return self.lambda_matrix.shape[0]
56
         def k(self) \rightarrow int:
57
58
              Number of latent variables
60
61
              return self.lambda_matrix.shape[1]
62
         def compute_free_energy(
64
              self,
65
              x: np.ndarrav
              binary_latent_factor_model: AbstractBinaryLatentFactorModel,
67
68
69
              free energy associated with current E\!M parameters and data x
70 \\ 71
              :param x: data matrix (number_of_points, number_of_dimensions)
              :param binary_latent_factor_model: a binary_latent_factor_model :return: average free energy per data point """
75
76
              expectation_log_p_x_s_given_theta = (
                   self._compute_expectation_log_p_x_s_given_theta(
    x, binary_latent_factor_model
77
78
79
              approximation_model_entropy = self._compute_approximation_model_entropy()
81
82
                   \verb|expectation_log_p_x_s_given_theta| + approximation_model_entropy|
83
84
         def _compute_expectation_log_p_x_s_given_theta(
86
              x: np.ndarray
87
              binary_latent_factor_model: AbstractBinaryLatentFactorModel,
89
90
              The first term of the free energy, the expectation of log P(X,S | theta)
```

```
:param x: data matrix (number_of_points, number_of_dimensions) :param binary_latent_factor_model: a binary_latent_factor_model :return: the expectation of log P(X,S \mid theta) """
 95
 96
                 # (number_of_points, number_of_dimensions)
                 mu_lambda = self.lambda_matrix @ binary_latent_factor_model.mu.T
98
 99
100
                 # (number_of_latent_variables, number_of_latent_variables)
                 expectation_s_i_s_j_mu_i_mu_j = np.multiply(
self.lambda_matrix.T @ self.lambda_matrix
                       \verb|binary_latent_factor_model.mu.T @ binary_latent_factor_model.mu|,
104
                 106
107
108
                      0.5 \ * \ binary\_latent\_factor\_model.precision
110
                      np.sum(np.multiply(x, x))
- 2 * np.sum(np.multiply(x, mu_lambda))
+ np.sum(expectation_s_i_s_j_mu_i_mu_j)
113
114
                         np.trace(
                      expectation_s_i_s_j_mu_i_mu_j
) # remove incorrect E[s_i s_i] = lambda_i * lambda_i
+ np.sum( # add correct E[s_i s_i] = lambda_i
self.lambda_matrix
116
117
118
                            @ np.multiply(
120
                                  binary\_latent\_factor\_model.mu, \ binary\_latent\_factor\_model.mu
121
                 expectation_log_p_s_given_theta = np.sum(
124
125
                      np.multiply(
                            self.lambda_matrix,
binary_latent_factor_model.log_pi,
126
128
                      + np.multiply(
1 - self.lambda_matrix,
                            binary_latent_factor_model.log_one_minus_pi,
133
134
                 return expectation_log_p_x_given_s_theta + expectation_log_p_s_given_theta
135
           {\tt def\_compute\_approximation\_model\_entropy(self)} \; -\!\!\!> \; {\tt float}:
138
                 Compute the model entropy
139
                 :return: model entropy
140
142
                 {\tt return -np.sum}\,(
143
                      np.multiply(
                            self.lambda_matrix,
145
                            self.log\_lambda\_matrix,
146
                      + np. multiply (
                            1 - self.lambda_matrix,
self.log_one_minus_lambda_matrix,
148
149
```

src/models/binary_latent_factor_approximations/abstract_binary_latent_factor_approximation.py

Appendix 3: main.py

```
import os
             from dataclasses import asdict
             import jax
            import jax.numpy as jnp
import numpy as np
import pandas as pd
            from src.constants import CO2_FILE_PATH, DEFAULT_SEED, OUTPUTS_FOLDER from src.generate_images import generate_images from src.models.bayesian_linear_regression import LinearRegressionParameters
11
12
            \begin{array}{lll} from & src.models.gaussian\_process\_regression & import & GaussianProcessParameters \\ from & src.models.kernels & import & CombinedKernel, & CombinedKernelParameters \\ \end{array}
14
            from src.solutions import q2, q3, q4, q5, q6
             jax.config.update("jax_enable_x64", True)
17
18
             if __name__ == "__main__":
19
                         np.random.seed(DEFAULT\_SEED)
20
                         if not os.path.exists(OUTPUTS_FOLDER):
22
23
                                     os.makedirs(OUTPUTS_FOLDER)
                        # Question 2
Q2_OUTPUT_FOLDER = os.path.join(OUTPUTS_FOLDER, "q2")
if not os.path.exists(Q2_OUTPUT_FOLDER):
    os.makedirs(Q2_OUTPUT_FOLDER)
with open(CO2_FILE_PATH) as file:
25
26
28
29
                                     lines = [line.rstrip().split() for line in file]
                         df_co2 = pd.DataFrame(
32
                                    np.array([line for line in lines if line[0] != "#"]).astype(float)
33
                         column_names = lines [max([i for i, line in enumerate(lines) if line[0] == "#"])][1:]
34
                        df_co2.columns = column_names

t = df_co2.decimal.values[:] - np.min(df_co2.decimal.values[:])

y = df_co2.average.values[:].reshape(1, -1)
36
37
39
                         sigma = 1
                        \begin{array}{ll} \operatorname{Signat} & -1 \\ \operatorname{mean} & = \operatorname{np.array} \left( [0\,,\ 360] \right).\operatorname{reshape} \left( -1,\ 1 \right) \\ \operatorname{covariance} & = \operatorname{np.array} \left( \end{array} \right) \end{array}
40
42
43
                                                   [10**2, 0]
44
                                                  [0, 100**2],
45
46
47
48
                          kernel = CombinedKernel()
                         kernel_parameters = CombinedKernelParameters( log_theta=jnp.log(1),
49
                                     log_sigma=jnp.log(1),
log_phi=jnp.log(5e-1),
log_eta=jnp.log(1),
50
                                     \log_{-} tau = jnp.log(1.5)
54
                                     \log zeta=jnp.\log (1e-2),
56
57
                          prior_linear_regression_parameters = LinearRegressionParameters (
58
                                     covariance=covariance,
60
61
                          posterior_linear_regression_parameters = q2.a(
                                    t,
62
64
                                     prior_linear_regression_parameters
65
                                     save_path=os.path.join(Q2_OUTPUT_FOLDER, "a"),
67
68
69
                                     t_year=df_co2.decimal.values[:],
70 \\ 71
                                     t=t ,
                                     y=y,
                                     linear\_regression\_parameters = posterior\_linear\_regression\_parameters \; ,
                                     error_mean=0,
                                     error_variance=1,
75
76
                                     \verb|save-path| = \verb|os.path|.join| (Q2-OUTPUT-FOLDER, "b") ,
77
78
79
                         q2.c(
                                     kernel=kernel,
                                     kernel_parameters=kernel_parameters ,
                                     \begin{array}{l} \log_{-1} + \log_{-1
81
82
83
                                     \verb|save_path| = \verb|os.path.join| (Q2\_OUTPUT\_FOLDER, "c") \;,
84
86
                          init\_kernel\_parameters = Combined Kernel Parameters (
87
                                     log_theta=jnp.log(5),
                                     log_sigma=jnp.log(5),
log_phi=jnp.log(10),
log_eta=jnp.log(5),
89
90
                                     log_tau=jnp.log(1)
```

```
\log_z z e t a = j n p \cdot \log(2),
 94
           gaussian_process_parameters = GaussianProcessParameters(
 95
 96
                kernel=asdict(init_kernel_parameters),
                log_sigma=jnp.log(1),
 97
 98
 99
          100
           t_{test} = np.concatenate((t, t_{new}))
                \dot{t}_{-}train=t,
                y_train=y,
t_test=t_test,
104
106
                \verb|min_year=np.min| (|df_co2.decimal.values[:])|
                prior_linear_regression_parameters=prior_linear_regression_parameters,
107
108
                linear_regression_sigma=sigma,
                kernel=kernel,
110
                gaussian_process_parameters=gaussian_process_parameters,
                learning_rate=1e-2,
112
                number_of_iterations=100.
                save_path=os.path.join(Q2_OUTPUT_FOLDER, "f"),
113
114
115
116
          # Q3_OUTPUT_FOLDER = os.path.join(OUTPUTS_FOLDER, "q3")
if not os.path.exists(Q3_OUTPUT_FOLDER):
118
                os.makedirs(Q3_OUTPUT_FOLDER)
120
          number_of_images = 2000
          x = generate_images (n=number_of_images)

k = 8
121
           {\tt em\_iterations} \ = \ 100
          e-maximum-steps = 50
124
           e_convergence_criterion = 0
126
           binary\_latent\_factor\_model \;, \; mean\_field\_approximation \; = \; q3 \,.\, e\_and\_f \, (
128
                k=k
                em_iterations=em_iterations,
                e_maximum_steps=e_maximum_steps,
e_convergence_criterion=e_convergence_criterion,
                save_path=os.path.join(Q3_OUTPUT_FOLDER, "f"),
134
135
          q3.g(
               x=x[:1, :],
binary_latent_factor_model=binary_latent_factor_model,
mean_field_approximation=mean_field_approximation,
136
138
139
                sigmas = [1, 2, 3],
                em iterations=em iterations
140
                save_path=os.path.join(Q3_OUTPUT_FOLDER, "g"),
141
142
          )
143
          os.makedirs(Q4_OUTPUT_FOLDER):

os.makedirs(Q4_OUTPUT_FOLDER):
145
146
           \max k = 21
148
149
           free\_energies\_1 = q4.b(
150
               x=x,
                a-parameter=1,
152
                b_parameter=0,
153
                ks=np.arange(4, 13),
                max_k=max_k,
                em_iterations=em_iterations,
156
                e\_maximum\_steps = e\_maximum\_steps \;,
                e_convergence_criterion=e_convergence_criterion,
                save_path=os.path.join(Q4_OUTPUT_FOLDER, "b-1"),
160
           free_energies_2 = q4.b(
162
                a_parameter=1,
163
                b_parameter=0,
164
                ks=np.arange(13, 22),
                \max_{k=\max_{k}}
                em_iterations=em_iterations
166
167
                e\_maximum\_steps = e\_maximum\_steps \;,
                e_convergence_criterion=e_convergence_criterion, save_path=os.path.join(Q4_OUTPUT_FOLDER, "b-2"),
168
170
171
          ) q4.free_energy_plot(
                free_energies=free_energies_1 + free_energies_2 ,
model_name="Variational Bayes",
\frac{173}{174}
175
                \verb|save_path| = \verb|os.path.join(Q4\_OUTPUT\_FOLDER, "b")|,
\frac{176}{177}
178
          # Question 5
Q5_OUTPUT_FOLDER = os.path.join(OUTPUTS_FOLDER, "q5")
if not os.path.exists(Q5_OUTPUT_FOLDER):
179
180
181
                os.makedirs(Q5_OUTPUT_FOLDER)
          max_k = 21
182
183
           free\_energies\_1 = q5.d(
184
185
                a_parameter=1.
                b-parameter=0,
187
               ks=np.arange(4, 13),
max_k=max_k,
188
```

```
189
190
                               em_iterations=em_iterations , save_path=os.path.join (Q5_OUTPUT_FOLDER, "d-1") ,
191
192
                      free_energies_2 = q5.d(
                              x=x,
a_parameter=1,
b_parameter=0,
ks=np.arange(13, 22),
max_k=max_k,
em_iterations=em_iterations,
 193
194
195
196
197
198
199
200
201
                               \verb|save_path| = \verb|os.path.join| (Q5\_OUTPUT\_FOLDER, "d-2") \;,
                     )
q4.free_energy_plot(
    ks=np.arange(4, 22),
    free_energies=free_energies_1 + free_energies_2,
    model_name="Loopy BP E Step and Variational Bayes M Step",
    save_path=os.path.join(Q5_OUTPUT_FOLDER, "d"),
202
203
204
205
206
207
208
                    # Question 6
Q6_OUTPUT_FOLDER = os.path.join(OUTPUTS_FOLDER, "q6")
if not os.path.exists(Q6_OUTPUT_FOLDER):
   os.makedirs(Q6_OUTPUT_FOLDER)
q6.run(x, k, em_iterations, save_path=os.path.join(Q6_OUTPUT_FOLDER, "all"))
209
210
211
```

main.py

Appendix 4: constants.py

```
import os

DATAFOLDER = "data"

CO2.FILE.PATH = os.path.join(DATAFOLDER, "co2.txt")
IMAGES.FILE.PATH = os.path.join(DATAFOLDER, "images.jpg")

OUTPUTS.FOLDER = "outputs"

DEFAULT.SEED = 0

II

M1 = [0, 0, 1, 0, 0, 1, 1, 1, 0, 0, 1, 0, 0, 0, 0]

M2 = [0, 1, 0, 0, 0, 1, 0, 0, 0, 1, 0, 0, 0, 0, 0]

M3 = [1, 1, 1, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0]

M4 = [1, 0, 0, 0, 0, 1, 0, 0, 0, 1, 1, 0, 0, 0, 1]

M5 = [0, 0, 0, 0, 0, 0, 0, 0, 1, 1, 0, 0, 1, 1, 1, 1]

M6 = [1, 1, 1, 1, 0, 0, 0, 1, 1, 0, 0, 1, 1, 1, 1]

M7 = [0, 0, 0, 0, 0, 1, 0, 0, 0, 1, 1, 0, 0, 0, 0, 0]

M8 = [0, 0, 0, 1, 0, 0, 0, 1, 0, 0, 0, 1, 1, 0, 0, 0, 0]

M8 = [0, 0, 0, 1, 0, 0, 0, 1, 0, 0, 0, 1, 0, 0, 0, 0]

M8 = [0, 0, 0, 1, 0, 0, 0, 1, 0, 0, 0, 1, 0, 0, 0, 0, 0]

M8 = [0, 0, 0, 1, 0, 0, 0, 1, 0, 0, 0, 1, 0, 0, 0, 0, 1]
```

src/constants.py

Appendix 5: generate_images.py

```
import numpy as np
     from src.constants import DEFAULT_SEED, M1, M2, M3, M4, M5, M6, M7, M8
         n: int = 400, seed: int = DEFAULT_SEED, sigma: float = 0.1
     ) -> np.ndarray:
         Image generation, adapted from provided demo code
11
12
13
         :param n: number of data points
         :param seed: random seed
:param sigma: Gaussian noise
14
15
16
         :return: images as a data matrix (number_of_points, number_of_dimensions)
17
18
19
         d=16 # dimensionality of the data
         {\tt np.random.seed (seed)}
20
21
         # Define the basic shapes of the features
         number_of_features = 8 # number of features
23
24
             0.5 + np.random.rand(number_of_features , 1) * 0.5 \# weight of each feature between 0.5 and 1
25
26
         mut = np.array(
27
                   rr[0] * M1,
                   rr [1]
rr [2]
rr [3]
rr [4]
rr [5]
rr [6]
28
29
                          * M3,
* M4,
* M5,
30
31
32
                          * M6,
* M7,
34
35
                          * M8
37
38
              np.random.rand(n, number_of_features) < 0.3
39
             # each feature occurs with prob 0.3 independently
40
         # Generate Data - The Data is stored in Y
43
              np.dot(s, mut) + np.random.randn(n, d) * sigma
             # some Gaussian noise is added
```

src/generate_images.py

Appendix 6: m_step.py

```
import numpy as np
           from typing import Tuple
  5
6
7
8
9
            def m_step(x: np.ndarray, es: np.ndarray, ess: np.ndarray) -> Tuple[np.ndarray, float, np.ndarray]:
                      mu, sigma, pie = m\_step(x, es, ess)
                      Inputs:
11
12
13
                                   x: shape (n, d) data matrix es: shape (n, k) E_{-q}[s] ess: shape (k, k) sum over data points of E_{-q}[ss'] (n, k, k) if E_{-q}[ss'] is provided, the sum over n is done for you.
14
15
16
17
18
                      Outputs:
                             mu: shape (d, k) matrix of means in p(y|\{s_i\},mu,sigma) sigma: shape (,) standard deviation in same pie: shape (1, k) vector of parameters specifying generative distribution for s
19
\frac{20}{21}
22
23
24
                     n, d = x.shape
if es.shape[0] != n:
    raise TypeError('es must have the same number of rows as x')
k = es.shape[1]
if ess.shape == (n, k, k):
    ess = np.sum(ess, axis=0)
if ess.shape != (k, k):
    raise TypeError('ess must be square and have the same number of columns as es')
25
26
27
28
29
31
                     \begin{array}{lll} mu = np. \det(np. \det(np. \operatorname{linalg.inv}(\operatorname{ess}), \operatorname{es.T}), x).T \\ sigma = np. \operatorname{sqrt}\left((np. \operatorname{trace}(np. \det(x.T, x)) + np. \operatorname{trace}(np. \det(\operatorname{np. dot}(\operatorname{mu.T}, \operatorname{mu}), \operatorname{ess})) \\ & - 2 * np. \operatorname{trace}(\operatorname{np. dot}(\operatorname{np. dot}(\operatorname{es.T}, x), \operatorname{mu}))) \ / \ (n * d)) \\ pie = np. \operatorname{mean}(\operatorname{es}, \operatorname{axis} = 0, \operatorname{keepdims} = \operatorname{True}) \end{array}
33
36
                      return mu, sigma, pie
```

demo_code/m_step.py