COMP0085 Summative Assignment

Jan 4, 2023

Question 1

(a)

The 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.

The junction tree redrawn as a factor graph:



where the circular nodes are cliques and the square nodes are separators/factors.

(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)

(e)

Question 2

(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:

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

Moreover, for our data $\mathcal{D} = \{X, Y\}$:

$$P(\mathcal{D}|\mathbf{w}) = \mathcal{N}\left(\mathbf{Y} - \mathbf{w}^T \mathbf{X}, \sigma^2 \mathbf{I}\right)$$

where
$$\mathbf{X} = \begin{bmatrix} t_1 & t_2 \cdots t_N \\ 1 & 1 \cdots 1 \end{bmatrix} \in \mathbb{R}^{2 \times N}$$
 and $\mathbf{Y} \in \mathbb{R}^{1 \times N}$.

Knowing:

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

we can substitute the above distributions:

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

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 terms:

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

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

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

Thus we can identify the posterior covariance:

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

and the posterior mean:

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

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

		value
arameters	а	1.828457
	_	
	b	334.203782

Figure 1: The Posterior Mean

Figure 2: t=The Posterior Covariance

(b)

Plotting the residuals:

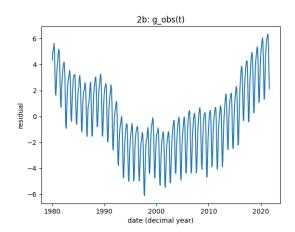


Figure 3: $g_{obs}(t)$

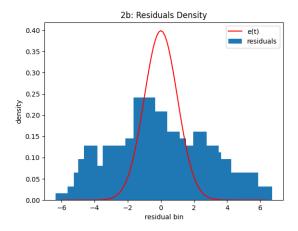


Figure 4: Density Estimation of Residuals vs $e(t) \sim \mathcal{N}(0,1)$

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.

(c & d)

We are considering the kernel:

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

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



Figure 5: Gram 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 sinusoidal 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 our CO_2 data, we would want a class of functions which exhibit both of these behaviours as the data looks sinusoidal (seasonal with respect to each year) and correlations locally.

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.

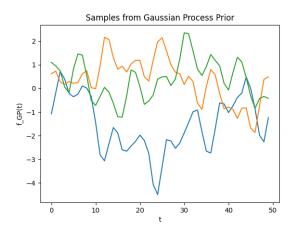


Figure 6: Samples from a zero mean GP with the provided covariance kernel

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

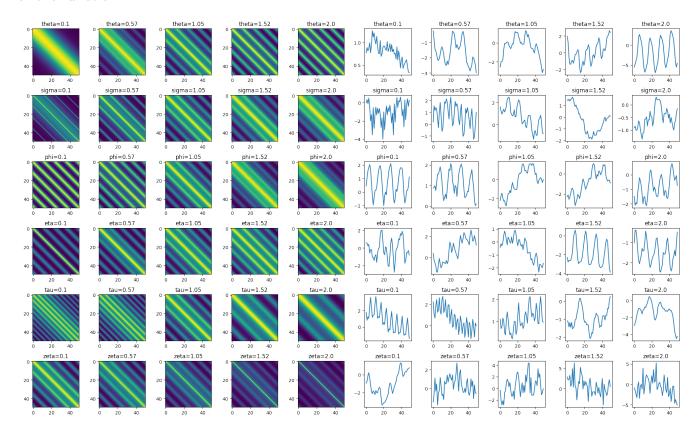


Figure 7: Covariances for different parameters

 η :

 τ :

 ζ :

Figure 8: Samples for different parameters

- θ : As θ increases, we see more pronounced periodic behavior in the sample function. The covariance matrix shows how increasing θ visually reveals the striped periodic component. This is expected because it is the parameter that adjusts the weight of $\exp\left(-\frac{2\sin^2(\pi(s-t)/\tau)}{\sigma^2}\right)$.
- σ : As σ increases, we see smoother periodic behaviour in the sample function. The covariance matrix shows how increasing σ will increase covariance values in the off-diagonals. This is expected because it adjusts the lengthscale of the periodic portion of the kernel.
- ϕ : As ϕ increases, we see less smooth behaviour in the sample function. The covariance matrix shows how increasing σ will increase covariance values in the off-diagonals. This is expected because it adjusts the lengthscale of the periodic portion of the kernel.

(e)

(f)

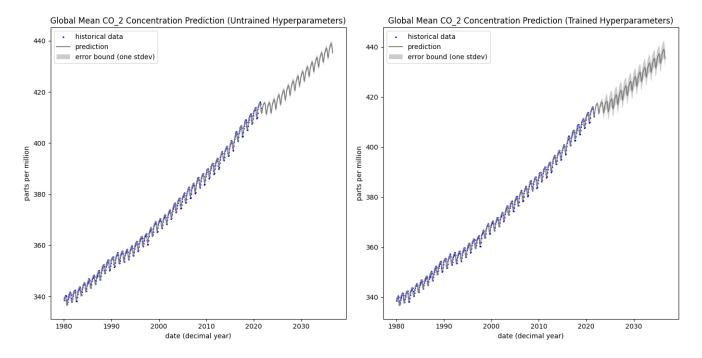


Figure 9: Without hyperparameter tuning

Figure 10: With hyperparameter tuning

(g)

The Python code for Bayesian Linear Regression:

```
from dataclasses import dataclass
 3
     import numpy as np
 5
6
7
     @dataclass
      class LinearRegressionParameters:
          mean: np.ndarray
covariance: np.ndarray
10
          @property
def precision(self):
    return np.linalg.inv(self.covariance)
11
12
13
14
          def predict(self, x: np.ndarray) -> np.ndarray:
16
17
                return self.mean.T @ x
18
19
20
     @dataclass
     class Theta:
21
          linear_regression_parameters: LinearRegressionParameters
          sigma: float
23
24
25
          def variance(self):
return self.sigma**2
26
27
28
          @property
def precision(self):
    return 1 / self.variance
29
30
31
33
34
     def compute_linear_regression_posterior (
          x: np.ndarray,
35
          y: np.ndarray,
           prior_linear_regression_parameters: LinearRegressionParameters,
residuals_precision: float,
36
38
     ) -> LinearRegressionParameters:
40
          Compute the parameters of the posterior distribution on the linear regression weights
41
42
          :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
44
           weights
45
           :param residuals_precision: the precision of the residuals of the linear regression
          return: parameters for the posterior distribution on the linear regression weights
46
47
48
          {\tt posterior\_covariance} \ = \ {\tt np.linalg.inv} \, (
                residuals\_precision \ * \ x \ @ \ x.T + prior\_linear\_regression\_parameters.precision
49
50
51
52
53
           posterior_mean = posterior_covariance @ (
               residuals_precision * x @ y.T + prior_linear_regression_parameters.precision
54
55
56
                @\ prior\_linear\_regression\_parameters.mean\\
           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
                  \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
183
            def sigma(self) -> float:
    return jnp.exp(self.log_sigma)
184
186
187
             @theta.setter
            def theta(self, value: float) -> None:
189
                   self.log_theta = jnp.log(value)
190
```

```
191
                                   @sigma.setter
 192
                                   def sigma(self, value: float) -> None:
                                                self.log_sigma = jnp.log(value)
 193
 194
  195
                                  def phi(self, value: float) -> None:
    self.log-phi = jnp.log(value)
 196
 197
 198
                                   @eta.setter
 199
                                   def eta(self, value: float) -> None:
 200
201
                                                   self.log_eta = jnp.log(value)
202
 203
                                   def tau(self, value: float) -> None:
    self.log_tau = jnp.log(value)
204
 205
 206
207
                                  @zeta.setter
def zeta(self, value: float) -> None:
 208
209
                                                  self.log_zeta = jnp.log(value)
210
 211
212
213
                    {\tt class} \;\; {\tt CombinedKernel(Kernel):} \\
                                  The kernel defined as:  k(x, y) = \frac{1}{2} \left( \frac{2\sin^2(\pi - y)}{\tan^2(x - y)} + \frac{1}{2} \left( \frac{\sin^2(\pi - y)}{\tan^2(x - y)} \right) + \frac{1}{2} \left( \frac{\sin^2(\pi - y)}{2} + \frac{1}{2} \left( \frac{\sin^2(\pi - y)}{2} \right) + \frac{1}{2} \left( \frac{\sin^2(\pi - y)}{2} + \frac{1}{2} \left( \frac{\sin^2(\pi - y)}{2} \right) + \frac{1}{2} \left( \frac{\sin^2(\pi - y)}{2} + \frac{1}{2} \left( \frac{\sin^2(\pi - y)}{2} \right) + \frac{1}{2} \left( \frac{\sin^2(\pi - y)}{2} + \frac{1}{2} \left( \frac{\sin^2(\pi - y)}{2} \right) + \frac{1}{2} \left( \frac{\sin^2(\pi - y)}{2} + \frac{1}{2} \left( \frac{\sin^2(\pi - y)}{2} \right) + \frac{1}{2} \left( \frac{\sin^2(\pi - y)}{2} + \frac{1}{2} \left( \frac{\sin^2(\pi - y)}{2} \right) + \frac{1}{2} \left( \frac{\sin^2(\pi - y)}{2} + \frac{1}{2} \left( \frac{\sin^2(\pi - y)}{2} \right) + \frac{1}{2} \left( \frac{\sin^2(\pi - y)}{2} + \frac{1}{2} \left( \frac{\sin^2(\pi - y)}{2} + \frac{1}{2} \left( \frac{\sin^2(\pi - y)}{2} \right) + \frac{1}{2} \left( \frac{\sin^2(\pi - y)}{2} + \frac{1}{2} \right) \right) + \frac{1}{2} \left( \frac{\sin^2(\pi - y)}{2} + \frac{1}{2} \right) \right) \right) + \frac{1}{2} \left( \frac{\sin^2(\pi - y)}{2} + \frac{1}{2} \right) \right) \right) + \frac{1}{2} \left( \frac{\sin^2(\pi - y)}{2} + \frac{1}{2} \right) \right) \right) \right) \right) \right) + \frac{1}{2} \left( \frac{\sin^2(\pi - y)}{2} + \frac{1}{2} \left( \frac{\sin^2(\pi - y)}{2} + \frac{1}{2} \left( \frac{\sin^2(\pi - y)}{2} + \frac{1}{2} \right) \right) \right) \right) \right) \right) + \frac{1}{2} \left( \frac{\sin^2(\pi - y)}{2} + \frac{1}{2} \right) \right) \right) \right) \right) \right) \right) \right) 
 214
215
216
218
219
                                   Parameters = CombinedKernelParameters
220
221
                                   def _kernel(
 222
223
                                                   parameters: CombinedKernelParameters,
 224
                                                  x: jnp.ndarray,
y: jnp.ndarray,
226
                                   ) -> jnp.ndarray:
 227
                                                            "Kernel evaluation between a single feature x and a single feature y.
229
                                                                 parameters: parameters dataclass for the Gaussian kernel
 230
                                                                 x: ndarray of shape (1,)
y: ndarray of shape (1,)
 231
232
 233
 234
                                                   \begin{array}{cccc} {\rm Returns:} & \\ & {\rm The \ kernel \ evaluation} \ . \end{array} 
 236
 237
                                                  return jnp.dot(
238
                                                                 jnp.ones(1),
                                                                                   (parameters.theta**2)
240
                                                                                ( p a * (
241
                                                                                                                \begin{array}{l} \texttt{jnp.exp(} \\ (-2 * \texttt{jnp.sin(jnp.pi *} (x - y) \ / \ \texttt{parameters.tau)} ** 2) \\ \\ / \ (\texttt{parameters.sigma**2)} \end{array}
243
244
246
248
                                                                                 \begin{array}{l} ' + (parameters.phi**2) \\ * (jnp.exp(-((x - y) ** 2) / (2 * parameters.eta**2))) \\ + parameters.zeta**2 * (x == y) \end{array} 
249
 250
251
252
```

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 GaussianProcess:
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
an.
                 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
      import optax
import dataframe_image as dfi
       import jax
      import jax.numpy as jnp
import matplotlib.pyplot as plt
import numpy as np
      import pandas as pd
import scipy
10
      \begin{array}{c} from \quad src.models.bayesian\_linear\_regression \quad import \quad (\\ LinearRegressionParameters \ , \end{array}
11
13
14
            compute_linear_regression_posterior ,
16
17
      from src.models.gaussian_process_regression import (
             GaussianProcess
            Gaussian Process Parameters,
19
20
      from src.models.kernels import CombinedKernel, CombinedKernelParameters
21
22
      jax.config.update("jax_enable_x64", True)
23
24
25
      def construct_design_matrix(t: np.ndarray):
    return np.stack((t, np.ones(t.shape)), axis=1).T
26
27
28
29
      def a(
30
            t: np.ndarray,
            y: np.ndarray, sigma: float,
             {\tt prior\_linear\_regression\_parameters}: \ {\tt LinearRegressionParameters} \ ,
            save_path: str.
35
      ) -> LinearRegressionParameters:
            x = construct_design_matrix(t)
prior_theta = Theta(
36
37
38
                   linear_regression_parameters=prior_linear_regression_parameters ,
39
                   sigma=sigma,
40
41
             posterior_linear_regression_parameters = compute_linear_regression_posterior(
42
                  prior_linear_regression_parameters , residuals_precision=prior_theta.precision ,
44
45
            df_mean = pd.DataFrame(
    posterior_linear_regression_parameters.mean, columns=["value"]
47
48
49
            df_mean.index = ["a", "b"]
df_mean = pd.concat([df_mean], keys=["parameters"])
dfi.export(df_mean, save_path + "-mean.png")
50
51
             df_covariance = pd.DataFrame(
                   posterior_linear_regression_parameters.covariance, columns=["a", "b"]
56
            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")
58
59
60
            return posterior_linear_regression_parameters
61
62
63
      def b(
65
            t_year ,
66
67
            linear_regression_parameters: LinearRegressionParameters,
            error_mean ,
70
71
72
73
74
75
76
            error_variance,
            save_path ,
            x = construct_design_matrix(t)
            x = construct_design_matrix(t)
residuals = y - linear_regression_parameters.predict(x)
plt.plot(t_year.reshape(-1), residuals.reshape(-1))
plt.xlabel("date (decimal year)")
plt.ylabel("residual")
plt.title("2b: g_obs(t)")
plt.savefig(save_path + "-residuals-timeseries")
plt.savefig(save_path + "-residuals-timeseries")
77
78
79
80
81
            count, bins = np.histogram(residuals, bins=100, density=True)
83
84
            plt.bar(bins[1:], count, label="residuals")
             plt.plot(
                  scipy.stats.norm.pdf(bins [1:], loc=error_mean, scale=error_variance), color="red", label="e(t)",
86
87
89
            plt.xlabel("residual bin")
plt.ylabel("density")
plt.title("2b: Residuals Density")
90
91
92
             plt.legend()
             plt.savefig(save_path + "-residuals-density-estimation")
```

```
plt.close()
 97
 98
       def c(
 aa
             kernel: CombinedKernel,
             kernel\_parameters: \ Combined Kernel Parameters \, ,
             log_theta_range: np.ndarray,
                 np.ndarray,
             number_of_samples: int,
103
             save_path: str,
       ):
             \begin{array}{ll} {\rm gram} \ = \ kernel\,(\,t\,\,,\  \  **asdict\,(\,kernel\,\text{-}parameters\,)\,) \\ {\rm plt\,\,.imshow}\,(\,{\rm gram}\,) \end{array}
106
107
             plt.xlabel("t")
plt.ylabel("t")
plt.title("Gram Matrix (Prior)")
plt.savefig(save_path + "-gram-matrix")
108
109
             plt.close()
113
             for _ in range(number_of_samples):
114
                   plt.plot(
116
                         np.random.multivariate_normal(
117
                              jnp.zeros(gram.shape[0]), gram, size=1
                         ). reshape(-1)
             plt.xlabel("t")
120
             plt.xlabel("f.GP(t)")
plt.ylabel("f.GP(t)")
plt.title("Samples from Gaussian Process Prior")
plt.savefig(save_path + "-samples")
123
124
             plt.close()
             fig_samples, ax_samples = plt.subplots(
126
                   len (fields (kernel.parameters._class__)), len (log_theta_range),
figsize=(len (log_theta_range) * 2, len (fields (kernel.parameters.__class__))*2),
127
128
                   frameon=False,
130
                  __class__)):
134
136
                                     jnp.zeros(gram.shape[0]), gram, size=1
                               ). reshape(-1),
140
             ax_samples[i][j].set_title(f"{field.name.strip('log_')}={np.round(np.exp(log_value), 2)}")
setattr(kernel_parameters, field.name, default_value)
plt.tight_layout()
142
             plt.savefig(save_path + f"-parameter-samples", bbox_inches='tight')
144
145
             plt.close(fig_samples)
             fig_gram , ax_gram = plt.subplots(
    len(fields(kernel_parameters.__class__)), len(log_theta_range),
    figsize=(len(log_theta_range) * 2, len(fields(kernel_parameters.__class__))*2),
147
148
149
                   frameon=False,
151
                       field in enumerate (fields (kernel_parameters
                   default_value = getattr(kernel_parameters, field name)
for j, log_value in enumerate(log_theta_range):
154
                   setattr(kernel-parameters, field.name, log.value)
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)}")
setattr(kernel-parameters, field.name, default_value)
156
158
             plt.tight_layout()
             plt.savefig(save_path + f"-parameter-grams", bbox_inches='tight')
161
             plt.close(fig_gram)
162
164
       def f(
165
                   t_train: np.ndarray,
y_train: np.ndarray,
t_test: np.ndarray,
166
167
168
169
                   min_year: float ,
                   \label{linear_regression_parameters: LinearRegressionParameters, linear\_regression\_sigma: float,
172
                   kernel: CombinedKernel,
                   {\tt gaussian\_process\_parameters}: \ \ {\tt GaussianProcessParameters}, \\ {\tt learning\_rate}: \ {\tt float} \ ,
\frac{175}{176}
                   number_of_iterations: int ,
save_path: str ,
       ):
            # Train Bayesian Linear Regression
x_train = construct_design_matrix(t_train)
prior_theta = Theta(
178
179
180
                   linear\_regression\_parameters = prior\_linear\_regression\_parameters \; ,
181
182
                   sigma=linear_regression_sigma ,
183
             posterior_linear_regression_parameters = compute_linear_regression_posterior(
184
185
                  x_train ,
186
                   v_train
                   prior_linear_regression_parameters .
187
                   residuals_precision=prior_theta.precision,
189
190
```

```
residuals = y\_train - posterior\_linear\_regression\_parameters.predict(x\_train)\\ gaussian\_process = GaussianProcess(kernel, t\_train.reshape(-1, 1), residuals.reshape(-1))
191
 192
193
 194
 195
                            x_test = construct_design_matrix(t_test)
                           \label{linear_prediction} I in ear\_prediction = posterior\_linear\_regression\_parameters.predict(x\_test).reshape(-1) \\ mean\_prediction, covariance\_prediction = gaussian\_process.posterior\_distribution(
196
 197
 198
                                        t_{test.reshape}(-1, 1), **asdict(gaussian_process_parameters)
199
200
201
                           # Plot
                           plt.figure(figsize=(7, 7))
plt.scatter(t_train+min_year, y_train.reshape(-1), s=2, color='blue', label="historical data")
plt.plot(t_test+min_year, linear_prediction + mean_prediction, color="gray", label="prediction")
plt.fill_between(
202
203
204
205
                                        t_test+min_year,
206
207
                                        linear_prediction+mean_prediction-1*jnp.diagonal(covariance_prediction),
linear_prediction+mean_prediction+1*jnp.diagonal(covariance_prediction),
208
                                        facecolor = (0.8, 0.8, 0.8),
label="error bound (one stdev)
209
210
211
                           plt.xlabel("date (decimal year)")
plt.ylabel("parts per million")
plt.title("Global Mean CO_2 Concentration Prediction (Untrained Hyperparameters)")
212
213
214
                            plt.legend()
216
                            plt.tight_layout()
                            plt.savefig(save_path+"-extrapolation-untrained", bbox_inches='tight')
218
                            plt.close()
219
220
                           # Train Gaussian Process Regression (Hyperparameter Tune)
                           optimizer = optax.adam(learning_rate)
gaussian_process_parameters = gaussian_process.train(
222
223
                                       optimizer, number_of_iterations, **asdict(gaussian_process_parameters)
224
226
                           # Prediction
227
                            x_{test} = construct_{design_matrix}(t_{test})
                            \frac{1}{1} = \frac{1}
                           mean_prediction, covariance_prediction = gaussian_process.posterior_distribution( t_{test.reshape}(-1, 1), **asdict(gaussian_process_parameters)
230
231
233
234
                           plt.figure(figsize=(7, 7))
                           plt.scatter(t_train+min_year, y_train.reshape(-1), s=2, color='blue', label="historical data") plt.plot(t_test+min_year, linear_prediction + mean_prediction, color="gray", label="prediction")
236
237
                            plt.fill_between (
238
                                         t_test+min_vear .
                                        linear_prediction+mean_prediction-1*jnp.diagonal(covariance_prediction),
240
                                        linear_prediction+mean_prediction+1*jnp.diagonal(covariance_prediction),
241
                                        facecolor = (0.8, 0.8, 0.8), label="error bound (one stdev)"
243
                           plt.xlabel("date (decimal year)")
plt.ylabel("parts per million")
plt.title("Global Mean CO_2 Concentration Prediction (Trained Hyperparameters)")
244
245
246
                            plt.legend()
248
                            plt.tight_layout()
                            plt.savefig(save_path+"-extrapolation-trained", bbox_inches='tight')
249
```

src/solutions/q2.py

Question 3

(a)

The free energy is can be calculated 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:

$$q(\mathbf{s}) = \prod_{i=1}^{K} q_i(s_i)$$

where $q_i(s_i) = \lambda_i^{s_i} (1 - \lambda_i)^{(1-s_i)}$.

To compute the first term:

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

substituting the appropriate terms:

$$P(\mathbf{x}|\mathbf{s}, \theta) = 2\pi^{-\frac{d}{2}} |\sigma^2 \mathbf{I}|^{-\frac{1}{2}} \exp\left(-\frac{1}{2} \left(\mathbf{x} - \sum_{i=1}^K s_i \mu_i\right)^T \frac{1}{\sigma^2} \mathbf{I} \left(\mathbf{x} - \sum_{i=1}^K s_i \mu_i\right)\right)$$

with d being the number of dimensions.

Taking the logarithm:

$$\log P(\mathbf{x}|\mathbf{s}, \theta) = -\frac{d}{2}\log(2\pi\sigma^2) - \frac{1}{2\sigma^2} \left(\mathbf{x}^T\mathbf{x} - 2\mathbf{x}^T\sum_{i=1}^K s_i\mu_i + \sum_{i=1}^K\sum_{j=1}^K s_is_j\mu_i^T\mu_j\right)$$

The expectation distributed to the relevant terms:

$$\langle \log P(\mathbf{x}|\mathbf{s}, \theta) \rangle_{q(\mathbf{s})} = -\frac{d}{2} \log(2\pi\sigma^2) - \frac{1}{2\sigma^2} \left(\mathbf{x}^T \mathbf{x} - 2\mathbf{x}^T \sum_{i=1}^K \langle s_i \rangle_{q_i(s_i)} \mu_i + \sum_{i=1}^K \sum_{j=1}^K \langle s_i s_j \rangle_{q_i(s_i)q_j(s_j)} \mu_i^T \mu_j \right)$$

Evaluating the expectations:

$$\langle \log P(\mathbf{x}|\mathbf{s}, \theta) \rangle_{q(\mathbf{s})} = -\frac{d}{2} \log(2\pi\sigma^2) - \frac{1}{2\sigma^2} \left(\mathbf{x}^T \mathbf{x} - 2\mathbf{x}^T \sum_{i=1}^K \lambda_i \mu_i + \sum_{i=1}^K \sum_{j=1, j \neq i}^K \lambda_i \lambda_j \mu_i^T \mu_j + \sum_{i=1}^K \lambda_i \mu_i^T \mu_i \right)$$

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

To compute the second term:

$$P(\mathbf{s}|\theta) = \prod_{i=1}^{K} \pi_i^{s_i} (1 - \pi_i)^{(1-s_i)}$$

Taking the logarithm:

$$\log P(\mathbf{s}|\theta) = \sum_{i=1}^{K} s_i \log \pi_i + (1 - s_i) \log(1 - \pi_i)$$

The expectation distributed to the relevant terms:

$$\langle \log P(\mathbf{s}|\theta) \rangle_{q(\mathbf{s})} = \sum_{i=1}^{K} \langle s_i \rangle_{q_i(s_i)} \log \pi_i + (1 - \langle s_i \rangle_{q_i(s_i)}) \log(1 - \pi_i)$$

Evaluating the expectations:

$$\langle \log P(\mathbf{s}|\theta) \rangle_{q(\mathbf{s})} = \sum_{i=1}^{K} \lambda_i \log \pi_i + (1 - \lambda_i) \log(1 - \pi_i)$$

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

$$H\left[q(\mathbf{s})\right] = \sum_{i=1}^{K} H\left[q_i(s_i)\right]$$

Thus,

$$H[q(\mathbf{s})] = -\sum_{i=1}^{K} \sum_{s_i \in \{0,1\}} q_i(s_i) \log q_i(s_i)$$

Substituting the appropriate values:

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

Combining, we have our free energy expression:

$$\begin{aligned} \mathcal{F}(q,\theta) &= \\ &\frac{-d}{2} \log(2\pi\sigma^2) - \frac{1}{2\sigma^2} \left(\mathbf{x}^T \mathbf{x} - 2\mathbf{x}^T \sum_{i=1}^K \lambda_i \mu_i + \sum_{i=1}^K \sum_{j=1, j \neq i}^K \lambda_i \lambda_j \mu_i^T \mu_j + \sum_{i=1}^K \lambda_i \mu_i^T \mu_i \right) \\ &+ \sum_{i=1}^K \lambda_i \log \pi_i + (1 - \lambda_i) \log(1 - \pi_i) \\ &- \sum_{i=1}^K \lambda_i \log \lambda_i + (1 - \lambda_i) \log(1 - \lambda_i) \end{aligned}$$

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

$$\frac{\partial}{\partial q_i} \left(\mathcal{F}(q, \theta) + \lambda^{LG} \int q_i - 1 \right) = \langle \log P(\mathbf{x}, \mathbf{s} | \theta) \rangle_{\prod_{j \neq i} q_j(s_j)} - \log q_i(s_i) - 1 + \lambda^{LG}$$

where λ^{LG} is the Lagrange multiplier.

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

$$\log q_i(s_i) = \langle \log P(\mathbf{x}, \mathbf{s} | \theta) \rangle_{\prod_{i \neq i} q_i(s_i)} - 1 + \lambda^{LG}$$

Similar to our free energy derivation:

$$\langle \log P(\mathbf{x}|\mathbf{s}, \theta) \rangle_{\prod_{j \neq i} q_j(s_j)} \propto -\frac{1}{2\sigma^2} \left(\mathbf{x}^T \mathbf{x} - 2\mathbf{x}^T \sum_{k=1}^K \langle s_k \rangle_{\prod_{j \neq i} q_j(s_j)} \mu_i + \sum_{k=1}^K \sum_{j=1}^K \langle s_k s_j \rangle_{\prod_{j \neq i} q_j(s_j)} \right)$$

and

$$\langle \log P(\mathbf{s}|\theta) \rangle_{\prod_{j \neq i} q_j(s_j)} = \sum_{k=1}^K \langle s_k \rangle_{\prod_{j \neq i} q_j(s_j)} \log \pi_k + (1 - \langle s_k \rangle_{\prod_{j \neq i} q_j(s_j)}) \log(1 - \pi_k)$$

We can write:

$$\log q_i(s_i) \propto \log P(\mathbf{x}|\mathbf{s}, \theta) \rangle_{\prod_{j \neq i} q_j(s_j)} + \langle \log P(\mathbf{s}|\theta) \rangle_{\prod_{j \neq i} q_j(s_j)}$$

Substituting the relevant terms:

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

Knowing $\log q_i(s_i) = s_i \log \lambda_i + (1 - s_i) \log(1 - \lambda_i)$:

$$\log q_i(s_i) \propto s_i \log \frac{\lambda_i}{1 - \lambda_i}$$

Thus,

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

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

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

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

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

Solving for λ_i :

$$\lambda_i = \frac{1}{1 + \exp\left[-\left(\frac{\mu_i^T}{\sigma^2} \left(\mathbf{x} - \frac{\mu_i}{2} - \sum_{j=1, j \neq i}^K \lambda_j \mu_j\right) + \log\frac{\pi_i}{1 - \pi_i}\right)\right]}$$

we have our partial update.

The	P	ython	code:

src/solutions/q3.py

Question 4

Question 5

(a)

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 $s_i^2 = s_i$:

$$\log p(\mathbf{s}, \mathbf{x}) = \sum_{i=1}^{K} \left(\left(\frac{\mathbf{x}^{T} \mu_{i}}{\sigma^{2}} + \log \frac{\pi_{i}}{1 - \pi_{i}} - \frac{\mu_{i}^{T} \mu_{j}}{2\sigma^{2}} \right) s_{i} \right) + \sum_{i=1}^{K} \sum_{j=1}^{i-1} \left(\frac{-\mu_{i}^{T} \mu_{j}}{\sigma^{2}} s_{i} s_{j} \right) + C$$

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:

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

and

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

as required. Note that C can simply be absorbed into any one of these factors. The Boltzmann Machine can be defined:

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

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

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

and

$$\log Z = -C$$

(b)

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

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

Thus,

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

For $g_{ij}(s_i, s_j)$, we will choose a product of Bernoulli's approximation:

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

Thus,

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

To derive the a message passing scheme, we first define:

$$q(\mathbf{s}) = \left(\prod_{i=1}^K \tilde{f}_i(s_i)\right) \left(\prod_{i=1}^K \prod_{j=1}^{i-1} \tilde{g}_{ij}(s_i, s_j)\right)$$

Thus, we can derive cavity distributions:

$$q_{\neg \tilde{f}_{i}(s_{i})}(\mathbf{s}) = \left(\prod_{i'=1, i' \neq i}^{K} \tilde{f}_{i'}(s_{i'})\right) \left(\prod_{i=1}^{K} \prod_{j=1}^{i-1} \tilde{g}_{ij}(s_{i}, s_{j})\right)$$

and

$$q_{\neg \tilde{g}_{ij}(s_i, s_j)}(\mathbf{s}) = \left(\prod_{i'=1}^K \tilde{f}_{i'}(s_{i'})\right) \left(\prod_{i'=1}^K \prod_{\substack{j'=1\\i', j' \neq i, j}}^{i'-1} \tilde{g}_{i'j'}(s_{i'}, s_{j'})\right)$$

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

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

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

$$\log \tilde{f}_i(s_i) = \log f_i(s_i)$$

$$\log\left(\frac{\lambda_i}{1-\lambda_i}\right)s_i = b_i s_i$$

$$\lambda_i = \frac{1}{1 + \exp(-b_i)}$$

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

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

Note that because $\tilde{g}_{ij}(s_i, s_j)$ is the product of two Bernoulli distributions, we only require the natural parameters:

$$\phi_{ij}(\theta) = \begin{bmatrix} \theta_i \\ \theta_j \end{bmatrix}$$

the mean with respect to s_i and s_j respectively.

We can write:

$$\log \tilde{g}_{ij}(s_i, s_j) q_{\neg \tilde{g}_{ij}(s_i, s_j)}(\mathbf{s}) \propto \log \left(\frac{\theta_i}{1 - \theta_i}\right) s_i + \log \left(\frac{\theta_j}{1 - \theta_j}\right) s_j + \sum_{i'=1}^K \log \left(\frac{\lambda_{i'}}{1 - \lambda_{i'}}\right) s_{i'} + \sum_{i'=1}^K \sum_{\substack{j'=1\\i',j' \neq i,j}}^{i'-1} \log \left(\frac{\theta_{i'}}{1 - \theta_{i'}}\right) s_{i'} + \log \left(\frac{\theta_{j'}}{1 - \theta_{j'}}\right) s_{j'}$$

Only including terms with s_i and s_j :

$$\log \tilde{g}_{ij}(s_i, s_j) q_{\neg \tilde{g}_{ij}(s_i, s_j)}(\mathbf{s}) \propto \left((K - 1) \log \left(\frac{\theta_i}{1 - \theta_i} \right) + \log \left(\frac{\lambda_i}{1 - \lambda_i} \right) \right) s_i + \left((K - 1) \log \left(\frac{\theta_j}{1 - \theta_j} \right) + \log \left(\frac{\lambda_j}{1 - \lambda_j} \right) \right) s_j$$

Moreover:

$$\log g_{ij}(s_i, s_j) q_{\neg \tilde{g}_{ij}(s_i, s_j)}(\mathbf{s}) \propto \frac{-\mu_i^T \mu_j}{\sigma^2} s_i s_j + \sum_{i'=1}^K \log \left(\frac{\lambda_{i'}}{1 - \lambda_{i'}}\right) s_{i'} + \sum_{i'=1}^K \sum_{\substack{i'=1 \ i', j' \neq i, j}}^{i'-1} \log \left(\frac{\theta_{i'}}{1 - \theta_{i'}}\right) s_{i'} + \log \left(\frac{\theta_{j'}}{1 - \theta_{j'}}\right) s_{j'}$$

Only including terms with s_i and s_i :

$$\log g_{ij}(s_i, s_j) q_{\neg \tilde{g}_{ij}(s_i, s_j)}(\mathbf{s}) \propto \frac{\neg \mu_i^T \mu_j}{\sigma^2} s_i s_j + \left((K - 2) \log \left(\frac{\theta_i}{1 - \theta_i} \right) + \log \left(\frac{\lambda_i}{1 - \lambda_i} \right) \right) s_i + \left((K - 2) \log \left(\frac{\theta_j}{1 - \theta_j} \right) + \log \left(\frac{\lambda_j}{1 - \lambda_j} \right) \right) s_j$$

Appendix 1: constants.py

src/constants.py

Appendix 2: main.py

```
import os
      import jax
      import jax.numpy as jnp
import numpy as np
import pandas as pd
      from src.constants import CO2_FILE_PATH, IMAGES_FILE_PATH, OUTPUTS_FOLDER
      from src.generate_images import generate_images
from src.models.bayesian_linear_regression import LinearRegressionParameters
from src.models.kernels import CombinedKernel, CombinedKernelParameters
      from src.models.gaussian_process_regression import GaussianProcessParameters from src.solutions import q2, q3, q4, q5, q6 from dataclasses import asdict jax.config.update("jax_enable_x64", True)
13
15
16
      if --name-- == "--main--":
    if not os.path.exists(OUTPUTS_FOLDER):
        os.makedirs(OUTPUTS_FOLDER)
17
18
19
20
21
            if not os.path.exists(Q2_OUTPUT_FOLDER, "q2")
os.makedirs(Q2_OUTPUT_FOLDER):
23
24
             with open (CO2_FILE_PATH) as fil
                  lines = [line.rstrip().split() for line in file]
26
                  np.array([line for line in lines if line[0] != "#"]).astype(float)
29
30
31
            column_names = lines[max([i for i, line in enumerate(lines) if line[0] == "#"])][1:]
            \begin{array}{ll} df\_co2.columns = column\_names \\ t = df\_co2.decimal.values [:] - np.min(df\_co2.decimal.values [:]) \\ y = df\_co2.average.values [:].reshape(1, -1) \end{array}
34
35
37
38
            {\rm mean} \, = \, {\rm np.array} \, ( \, [ \, 0 \, \, , \, \, \, 360 \, ] \, ) \, . \, {\rm reshape} \, ( \, -1 \, , \, \, \, 1 \, )
            covariance = np.array(
39
                         \begin{bmatrix} 1 \, 0 **2 \;,\;\; 0 \, ] \;, \\ [ \, 0 \;,\;\; 1 \, 0 \, 0 **2 \, ] \;, \\ \\ \end{array}
40
41
43
44
             kernel = CombinedKernel()
            kernel-parameters = CombinedKernelParameters(
log_theta=jnp.log(1),
log_sigma=jnp.log(1),
46
47
48
                  \log_{-p} \text{hi} = \text{jnp.log}(1)
                  log_eta=jnp.log(1),
log_tau=jnp.log(1),
49
                  \log_{z} e t a = j n p \cdot \log (1 e - 1),
53
54
             {\tt prior\_linear\_regression\_parameters} \ = \ {\tt LinearRegressionParameters} \ (
                  mean=mean.
56
                   covariance=covariance,
57
58
            posterior_linear_regression_parameters = q2.a(
60
                  prior_linear_regression_parameters ,
save_path=os.path.join(Q2_OUTPUT_FOLDER, "a"),
62
65
            q2.b(
                  t_year=df_co2.decimal.values[:],
66
68
                  v=v.
                  linear_regression_parameters=posterior_linear_regression_parameters,
69
70
71
                  error_mean=0,
                  error_variance=1,
                  save_path=os.path.join(Q2_OUTPUT_FOLDER, "b"),
73
74
75
76
77
78
            q2.c(
                  kernel=kernel,
                  kernel_parameters=kernel_parameters .
                  log\_theta\_range=jnp.log(jnp.linspace(1e-1, 2, 5)),
79
                  t=t [:50]. reshape(-1, 1),
number_of_samples=3,
80
                  {\tt save\_path$=$os.path.join} \; ( {\tt Q2\_OUTPUT\_FOLDER}, \;\;"c") \; ,
82
            {\tt gaussian\_process\_parameters} \ = \ {\tt GaussianProcessParameters} \, (
85
                  kernel=asdict(kernel\_parameters),
                  log_sigma=jnp.log(1),
88
             vears_to_predict = 15
            90
91
            q2.f(
                  t_train=t,
```

```
y_train=y,
t_test=t_test,
min_year=np.min(df_co2.decimal.values[:]),
prior_linear_regression_parameters=prior_linear_regression_parameters,
linear_regression_sig ma=sigma,
kernel=kernel,
gaussian_process_parameters=gaussian_process_parameters,
learning_rate=le-2,
number_of_iterations=100,
save_path=os.path.join(Q2_OUTPUT_FOLDER, "f"),
  93
94
95
96
97
98
   99
 100
 101
 102
 103
 104
                           # Question 3
Q3.OUTPUT.FOLDER = os.path.join(OUTPUTS.FOLDER, "q3")
if not os.path.exists(Q3.OUTPUT.FOLDER):
    os.makedirs(Q3.OUTPUT.FOLDER)
 105
 \frac{106}{107}
 108
 109
110
                          # q3.learn_binary_factors(
# x=generate_images(),
# k=8,
# em_maximum_iterations
# e_maximum_steps=100,
# e_convergence_criteri
# )
111
112
 113
                                              em_maximum_iterations=5,
                                              e_maximum_steps=100,
e_convergence_criterion=0,
114
115
116
```

main.py