Lab 9: Gaussian Processes (GPs)

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Computing the weighted mean and variance

To calculate the mean and variance of the latents, have to use the **weighted** mean and variance of the particles.

$$\mathbb{E}[x] = \int x p(x) dx \approx \sum_{i} X^{(i)} W^{(i)}$$
 (1)

$$\mathbb{V}[x] = \int (x - \mathbb{E}[x])^2 p(x) dx \approx \sum_i (X^{(i)} - \mathbb{E}[x])^2 W^{(i)}$$
 (2)

Updating the last sample

Due to index inside the loop being n-1, have to remember to update the final estimate for the latent mean and variance.

```
### track for last sample:
w = self.compute_w(data[-1, :], z_samp)
self.est_z_mean[-1] = np.sum(w * z_samp)
self.est_z_var[-1] = np.sum(w * (z_samp - self.est_z_mean[-1]) ** 2)
```

Use different instantiation of noise for each particle propagation

Each particle $(Z_t^{(i)})$ is propagated independently of each other.

$$Z_{t+1}^{(i)} = AZ_t^{(i)} + W_t^{(i)}$$
 (3)

where each $W_t^{(i)} \sim \mathcal{N}(0, Q)$. Each of these $W_t^{(i)}$ is sampled **i.i.d**. The following is wrong:

$$Z_{t+1}^{(i)} = AZ_t^{(i)} + W_t (4)$$

where one instance of W_t is used to propagate all particles.

Linear Regression

Given a set of points (x, y), we want to find the points from a modelled signal i.e.

$$y|x \sim \mathcal{N}(f, \sigma^2)$$

 $f(x) = w^{\top} \psi(x).$

- ► Under these assumptions, maximum likelihood is equivalent to linear regression.
- ► With an additional prior on the weights, the maximum a posteriori solution is equivalent to regularized linear regression.
- ► The full Bayesian inference would average over all likely explanations under the posterior distribution.

Bayesian Linear Regression

Assuming:

$$w \sim \mathcal{N}(0, S)$$
$$y|x, w \sim \mathcal{N}(w^{\top}\psi(x), \sigma^2),$$

we can write:

$$\log p(w|\mathcal{D}) = \log p(w) + \log p(\mathcal{D}|w) + K$$

= $-\frac{1}{2}w^{\top}Z^{-1}w - \frac{1}{2\sigma^2}||\psi w - y||^2 + K.$

Bayesian Linear Regression

Developing and completing the square, we can show that this is a multivariate Gaussian distribution :

$$w|\mathcal{D} \sim \mathcal{N}(\mu, \Sigma),$$

where,

$$\mu = \frac{1}{\sigma^2} \Sigma \psi(x)^{\top} y$$
$$\Sigma^{-1} = \frac{1}{\sigma^2} \psi(x)^{\top} \psi(x) + S^{-1}.$$

Bayesian Linear Regression

We want the predictive distribution :

$$p(y_*|x_*,\mathcal{D}) = \int p(t_*|x_*,w)p(w|\mathcal{D}) dw.$$

All quantities are Gaussian, we can compute this integral:

$$\begin{split} & \mu_{\mathsf{pred}} = \mu^\top \psi(x_*) \\ & \sigma_{\mathsf{pred}}^2 = \psi(x_*)^\top \Sigma \psi(x_*) + \sigma^2. \end{split}$$

Note that this quantity could be expressed in terms of :

$$\mu_f = \mu_w^\top \psi(x)$$

$$\Sigma_f = \psi(x^\top \Sigma_w \psi(x)).$$

We don't need to know w anymore! What if we now let those quantities to be arbitrary?

Gaussian Process

Definition

A Gaussian Process (GP) is a collection of random variables, any finite number of which a have a joint Gaussian distribution.

We write $f(x) \sim \mathcal{GP}(m, k)$:

$$[f(x_1), \dots, f(x_N)] \sim \mathcal{N}(\mu, K)$$

 $\mu_i = m(x_i)$
 $K_{ij} = k(x_i, x_j),$

where x_1, \ldots, x_N are inputs. m is the mean function, k is the covariance kernel.

Inference

We are interested in making predictions f_* at points x_* . Hence we are interested in the distribution $p(f_*|x,y,x_*)$. Under the following assumptions :

$$y(x) \sim \mathcal{N}\left(f(x), \sigma^2\right),$$

 $f(x) \sim \mathcal{GP}(0, k_{\theta}),$

we can write:

$$\begin{bmatrix} y \\ f_* \end{bmatrix} \sim \mathcal{N} \left(0, \begin{bmatrix} k_{\theta}(x, x) + \sigma^2 I & k_{\theta}(x, x_*) \\ k_{\theta}(x_*, x) & h_{\theta}(x_*, x_*) \end{bmatrix} \right).$$

Inference

We are interested in making predictions f_* at points x_* . Hence we are interested in the distribution $p(f_*|x,y,x_*)$. Using gaussian identities,

$$\begin{split} f_* | x_*, x, y, \theta &\sim \mathcal{N}(\bar{f}_*, \mathsf{cov}(f_*)) \\ \bar{f}_* &= k_\theta(x_*, x) \left[k_\theta(x, x) + \sigma^2 I \right]^{-1} y, \\ \mathsf{cov}(f_*) &= k_\theta(x_*, x_*) - k_\theta(x_*, x) \left[k_\theta(x, x) + \sigma^2 I \right]^{-1} k_\theta(x, x_*). \end{split}$$

Choosing hyperparameters

- The problem from the previous expression is to determine the hyperparameters θ .
- ► If we wanted to be fully Bayesian, we could specify a prior over those hyperparameters and marginalize them.
- ► In practice, if the prior is flat, we can use a maximum marginal likelihood approach i.e. maximize :

$$p(y|x,\theta) = \int p(y|f,x,\theta)p(f|x,\theta) df.$$

For GP, it is reasonable to assume that $f|X \sim \mathcal{N}(0, K)$. We then have $y \sim \mathcal{N}(0, K + \sigma_n^2 I)$.

Kernel trick

- ► In the Bayesian linear regression example, we used the kernel trick.
- The aim to express a target in function of a dot product between feature vectors i.e. $\langle x, x' \rangle = \psi(x)^{\top} \psi(x')$.
- A kernel implements an inner product between feature vectors, typically implicitly, and often much more efficiently than the explicit dot product.
- Example :

$$\phi(x) = (1, \sqrt{2}x_1, \dots, \sqrt{2}x_d, \sqrt{2}x_1x_2, \sqrt{2}x_1x_3, \dots, \sqrt{2}x_{d-1}x_d, x_1^2, \dots, \sqrt{2}x_d, x$$

but:

$$k(x, x') = \langle \phi(x), \phi(x') \rangle = (1 + \langle x, x' \rangle)^2,$$

which is linear!

Kernel trick

- We often think directly in kernel space, rather than in feature space.
- ► The kernel allows to use very high dimensional feature spaces, but we do have a computational cost.

GP regression:

$$\begin{split} f_*|x_*,x,y,\theta &\sim \mathcal{N}(\bar{f}_*,\mathsf{cov}(f_*)) \\ \bar{f}_* &= k_\theta(x_*,x) \left[k_\theta(x,x) + \sigma^2 I \right]^{-1} y, \\ \mathsf{cov}(f_*) &= k_\theta(x_*,x_*) - k_\theta(x_*,x) \left[k_\theta(x,x) + \sigma^2 I \right]^{-1} k_\theta(x,x_*). \end{split}$$

We need to invert an $N \times N$ matrix!

▶ The $O(N^3)$ cost is typical of kernel methods.

Sensitivity to initial conditions

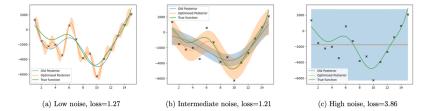
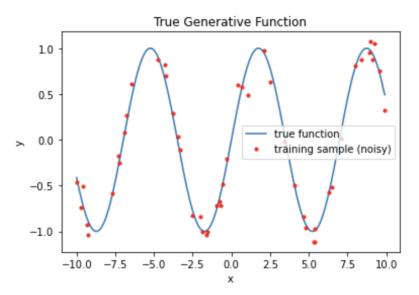


Figure 1: Initial conditions

References

- ► CSC 411 Lecture 20: Gaussian Processes
- ► Pattern Recognition and Machine Learning, Bishop

Part I: Data generation



Part II: Kernels

Part II GP with sklearn

Sklearn has a very handy API for Gaussian Process regression.

http://scikit-learn.org/stable/modules/gaussian_process.html

Kernel functions

Kernels to parametrize covariance structure

Constant Kernel: covariance is defined by a constant value

RBF (squared exponential) Kernel:

$$K(x_m, x_n) = exp\left(-\frac{||x_m - x_n||^2}{2 * l^2}\right)$$

White Kernel: accords for noise-component

$$K(x_m, x_n) = \text{noise}$$
 if $x_m = x_n$ else 0

Part II: Kernels

fitting the GP model

The fit() method automatically selects the hyper-parameters of given kernels.

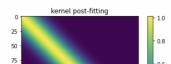
```
gp = gaussian_process.GaussianProcessRegressor(kernel=kernel)
gp.fit(X_train.reshape(-1,1), y_train.reshape(-1,1))
/opt/miniconda3/envs/main/lib/python3.9/site-packages/sklearn/gaussian_process/kernels.py:
e optimal value found for dimension 0 of parameter kl_kl_constant_value is close to the
5. Decreasing the bound and calling fit again may find a better value.
warnings.warn(
```

GaussianProcessRegressor(kernel=1**2 + RBF(length_scale=2) + WhiteKernel(noise_level=1))

```
# print the kernel with fitted parameters
print(gp.kernel_)
plt.figure()
plt.imshow(gp.kernel_(np.array([all_x]).T))
plt.colorbar()
plt.title('kernel post-fitting')
```

0.00316**2 + RBF(length_scale=2.02) + WhiteKernel(noise_level=0.011)

Text(0.5, 1.0, 'kernel post-fitting')



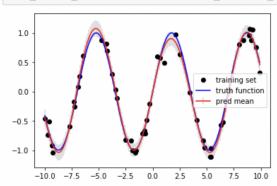
Part II: inference with sklearn

prediction of new values

The predict method returns both mean and std.

```
mus, sigmas = gp.predict(all_x.reshape(-1,1), return_std=True)
```

plot_gp(all_x, mus[:, 0], sigmas, X_train, y_train, true_y=true_y)



Part III: Cholensky predictions

$$p(\mathbf{t}^{pred}|\mathbf{t}^{train}) = N\left(\mu_{\mathbf{t}^{pred}|\mathbf{t}^{train}}, V_{\mathbf{t}^{pred}|\mathbf{t}^{train}}
ight) \ \mu_{\mathbf{t}^{pred}|\mathbf{t}^{train}} = K^T (C^{train})^{-1} \mathbf{t}^{train} \ V_{\mathbf{t}^{pred}|\mathbf{t}^{train}} = C^{pred} - K^T (C^{train})^{-1} K$$

Note that here, we assume zero mean

inference using the Cholesky Decomposition

- faster and more stable way to compute $\mu_{\mathbf{p}^{red}|\mathbf{p}^{train}}$ and $V_{\mathbf{p}^{red}|\mathbf{p}^{train}}$ given that $(C^{train})^{-1}$ is not guaranteed to be non-singular
- The Cholesky decomposition converts a (Hermitian, positive-definite) matrix A into the product of a lower triangular matrix L and its conjugate transpose L*
- ullet We use the Cholesky decomposition to get $C^{\it train} = LL^T$

Because our covariance matrix $(C^{train})^{-1}$ is positive-definite and a real matrix that mirrors itself along the diagonal, it is a Hermitian matrix

L will be a real-value matrix so its conjugate is itself

From this we get:

$$\mu_{t^{prod} \mid t^{prois}} = K^T (C^{train})^{-1} t^{train} = K^T (LL^T)^{-1} t^{train} = K^T (L^T)^{-1} L^{train} = (L^{-1}K)^T (L^{-1} t^{train})$$

$$V_{train total} = C^{prod} - K^T (C^{train})^{-1} K = C^{prod} - (L^{-1}K)^T (L^{-1}K)$$

where L=choleskv(C)

 $L^{-1}K$ and $L^{-1}t^{train}$ can be obtained by solving the linear system Lx=K and $Lx=t^{train}$ using np.linalg.solve

Part III: inference

prediction giving varying number of training data points

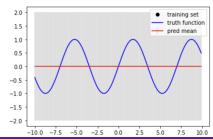
Prior distribution

$$y \sim N(\mu_0, \sigma_0^2)$$

Since we assume a zero mean function, we have $\mu_0 = E[y] = 0$.

```
mu_0 = np.zeros(len(all_x))
sigma_0 = np.sqrt(exponential_cov(0, 0, kernel_parameters))
plot_gp(all_x, mu_0, sigma_0, [], [], true_y)
print("rmse = {0}".format(np.sqrt(mean_squared_error(mu_0, true_y))))
```

rmse = 0.7216677922512522



Part IV: sampling

Part IV: Sampling

For this part, we implement the sample cholesky function.

sampling from multivariate Gaussian

use property of multivariate Gaussian where if $z \sim N(0, I)$ then $x = \mu + Lz$ gives $x \sim N(\mu, LL^T)$ where $L = cholesky(LL^T)$