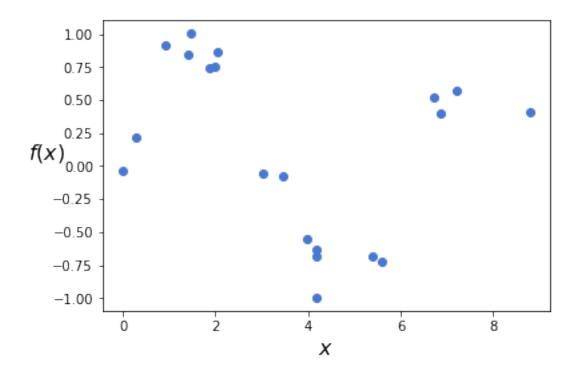
Bayesian Data Analysis Chapter 8

January 20, 2019

```
In [1]: %matplotlib inline
    import pymc3 as pm
    import numpy as np
    import pandas as pd
    import theano.tensor as tt
    import scipy.stats as stats
    import matplotlib.pyplot as plt
    import seaborn as sns;
    palette = 'muted'
    sns.set_palette(palette); sns.set_color_codes(palette)
    np.set_printoptions(precision=2)
```

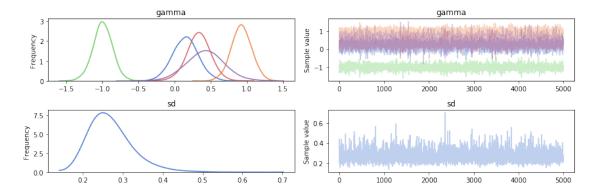
0.1 Kernelized Regression

```
In [2]: np.random.seed(1)
    x = np.random.uniform(0, 10, size=20)
    y = np.random.normal(np.sin(x), 0.2)
    plt.plot(x, y, 'o')
    plt.xlabel('$x$', fontsize=16);
    plt.ylabel('$f(x)$', fontsize=16, rotation=0);
```



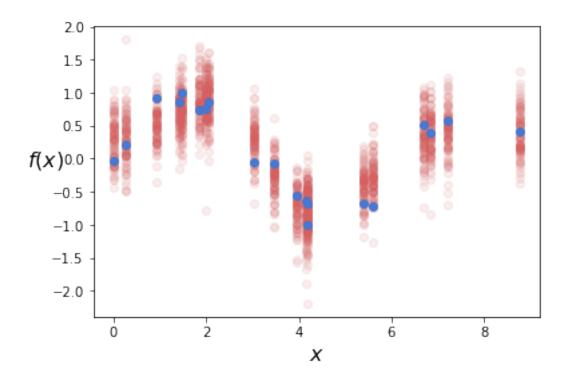
```
In [3]: def gauss_kernel(x, n_knots):
            Simple Gaussian radial kernel
           knots = np.linspace(x.min(), x.max(), n_knots)
            return np.array([np.exp(-(x-k)**2/w) for k in knots])
In [4]: n_knots = 5
In [5]: with pm.Model() as kernel_model:
            gamma = pm.Cauchy('gamma', alpha=0, beta=1, shape=n_knots)
            sd = pm.Uniform('sd',0, 10)
           mu = pm.math.dot(gamma, gauss_kernel(x, n_knots))
            yl = pm.Normal('yl', mu=mu, sd=sd, observed=y)
           kernel_trace = pm.sample(10000, chains=1,njobs=1)
Auto-assigning NUTS sampler...
Initializing NUTS using jitter+adapt_diag...
Sequential sampling (1 chains in 1 job)
NUTS: [sd, gamma]
100%|| 10500/10500 [00:08<00:00, 1168.71it/s]
Only one chain was sampled, this makes it impossible to run some convergence checks
```

In [6]: chain = kernel_trace[5000:] pm.traceplot(chain);

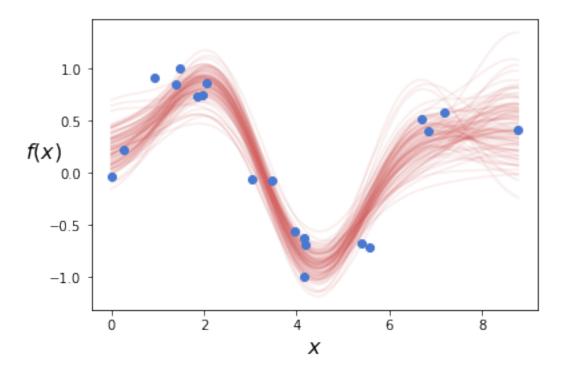


In [7]: pm.summary(chain)

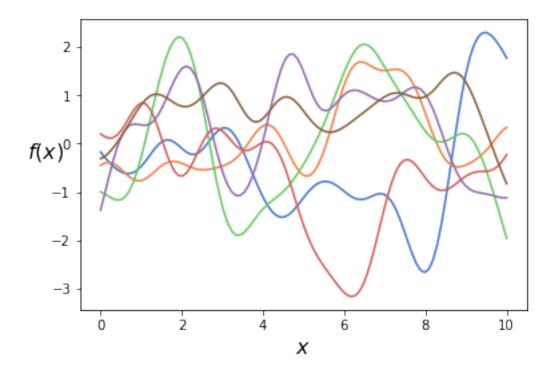
```
Out[7]:
                                  sd mc_error
                                                 hpd_2.5 hpd_97.5
                      mean
                 0.149587
                                      0.002675 -0.188123
                                                          0.513712
        gamma__0
                           0.178287
        gamma__1
                 0.927369
                           0.140474 0.002009 0.640079
                                                          1.194028
                           0.135827
                                      0.002167 -1.271611 -0.731901
        gamma 2 -0.997580
        gamma__3
                  0.335250
                            0.165045 0.002291 -0.006687
                                                          0.649435
                                      0.003824 -0.108575
        gamma__4
                  0.425855
                            0.261586
                                                          0.927639
        sd
                  0.270219
                            0.054199 0.000974 0.185070 0.385145
In [8]: ppc = pm.sample_posterior_predictive(chain, model=kernel_model, samples=100)
       plt.plot(x, ppc['yl'].T, 'ro', alpha=0.1);
       plt.plot(x, y, 'bo');
       plt.xlabel('$x$', fontsize=16);
       plt.ylabel('\frac{f(x)}{f(x)}', fontsize=16, rotation=0);
```



```
In [9]: new_x = np.linspace(x.min(), x.max(), 100)
    k = gauss_kernel(new_x, n_knots)
    gamma_pred = chain['gamma']
    for i in range(100):
        idx = np.random.randint(0, len(gamma_pred))
        # grap a random set of gammas from the MCMC chain
        # e.g. gamma_pred[3642]=[-0.04 0.93 -0.97 0.32 0.05]
        # to get an idea of the uncertainty
        y_pred = np.dot(gamma_pred[idx], k)
        plt.plot(new_x, y_pred, 'r-', alpha=0.1)
    plt.plot(x, y, 'bo')
    plt.xlabel('$x$', fontsize=16)
    plt.ylabel('$x$', fontsize=16, rotation=0);
```



0.2 Gaussian Processes



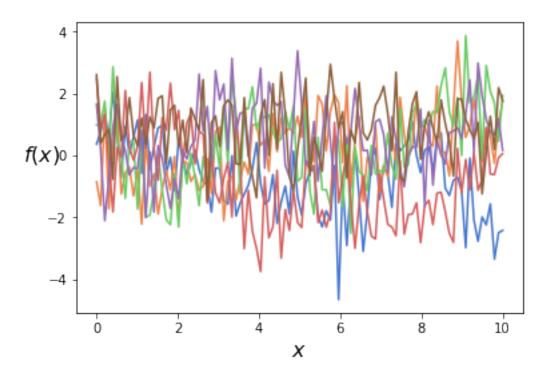
```
In [12]: np.random.seed(1)
    eta = 1
    rho = 0.5
    sigma = 0.03
    D = squared_distance(test_points, test_points)

    cov = eta * np.exp(-rho * D)
    diag = eta * sigma

    np.fill_diagonal(cov, diag)

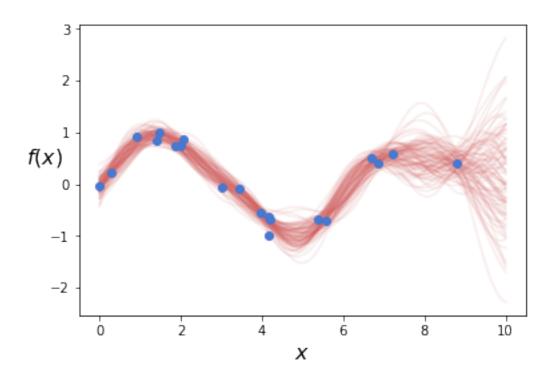
for i in range(6):
        plt.plot(test_points, stats.multivariate_normal.rvs(cov=cov))
    plt.xlabel('$x$', fontsize=16);
    plt.ylabel('$f(x)$', fontsize=16, rotation=0);
```

/home/damianos/miniconda3/envs/pymc3/lib/python3.7/site-packages/scipy/stats/_multivariate.py:out = random_state.multivariate_normal(mean, cov, size)



```
In [13]: np.random.seed(1)
         # K_{**}
         K_{oo} = eta * np.exp(-rho * D)
        D_x = squared_distance(x, x)
         # K
        K = eta * np.exp(-rho * D_x)
        diag_x = eta + sigma
        np.fill_diagonal(K, diag_x)
        D_off_diag = squared_distance(x, test_points)
         # K_{*}
         K_o = eta * np.exp(-rho * D_off_diag)
         # Posterior mean
        mu_post = np.dot(np.dot(K_o, np.linalg.inv(K)), y)
         # Posterior covariance
         SIGMA_post = K_oo - np.dot(np.dot(K_o, np.linalg.inv(K)), K_o.T)
         for i in range(100):
             fx = stats.multivariate_normal.rvs(mean=mu_post, cov=SIGMA_post)
             plt.plot(test_points, fx, 'r-', alpha=0.1)
```

```
plt.plot(x, y, 'o')
plt.xlabel('$x$', fontsize=16);
plt.ylabel('$f(x)$', fontsize=16, rotation=0);
```



Posterior of GP model using Cholesky decomposition

```
# Sample some input points and noisy versions of the function evaluated at
  # these points.
  X = np.random.uniform(0, 10, size=(N,1))
  y = f(X) + sigma * np.random.randn(N)
  K = kernel(X, X)
  L = np.linalg.cholesky(K + sigma * np.eye(N))
  # points we're going to make predictions at.
  Xtest = np.linspace(0, 10, n).reshape(-1,1)
  # compute the mean at our test points.
  Lk = np.linalg.solve(L, kernel(X, Xtest))
  mu = np.dot(Lk.T, np.linalg.solve(L, y))
  # compute the variance at our test points.
  K_ = kernel(Xtest, Xtest)
  sd_pred = (np.diag(K_) - np.sum(Lk**2, axis=0))**0.5
  plt.fill_between(Xtest.flat, mu - 2 * sd_pred, mu + 2 * sd_pred, color="r", alpha=0.2
  plt.plot(Xtest, mu, 'r', lw=2)
  plt.plot(x, y, 'o')
  plt.xlabel('$x$', fontsize=16)
  plt.ylabel('\frac{f(x)}{f(x)}', fontsize=16, rotation=0);
     2.0
     1.5
     1.0
f(x)^{0.5}
```

4

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8

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6

ż

0.0

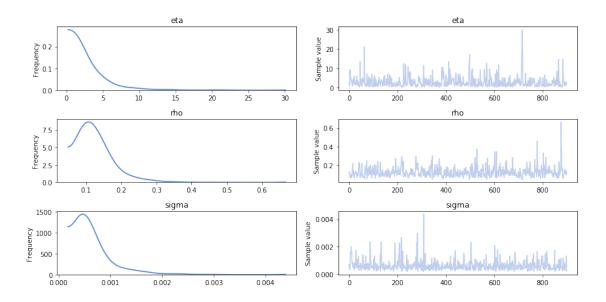
-0.5

-1.0

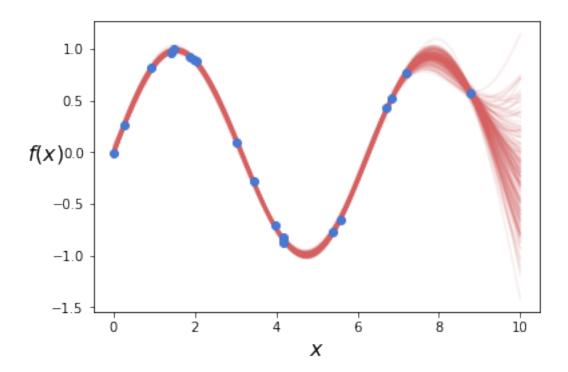
-1.5

Ó

```
In [15]: with pm.Model() as GP:
             mu = np.zeros(N)
             eta = pm.HalfCauchy('eta', 5)
             rho = pm.HalfCauchy('rho', 5)
             sigma = pm.HalfCauchy('sigma', 5)
             D = squared_distance(x, x)
             K = tt.fill_diagonal(eta * pm.math.exp(-rho * D), eta + sigma)
             obs = pm.MvNormal('obs', mu, cov=K, observed=y)
             test_points = np.linspace(0, 10, 100)
             D_pred = squared_distance(test_points, test_points)
             D_off_diag = squared_distance(x, test_points)
             K_{oo} = eta * pm.math.exp(-rho * D_pred)
             K_o = eta * pm.math.exp(-rho * D_off_diag)
             mu_post = pm.Deterministic('mu_post', pm.math.dot(pm.math.dot(K_o, tt.nlinalg.mat.
             SIGMA_post = pm.Deterministic('SIGMA_post', K_oo - pm.math.dot(pm.math.dot(K_o, t
             \#start = pm.find\_MAP()
             trace = pm.sample(1000, chains=1,njobs=1)
Auto-assigning NUTS sampler...
Initializing NUTS using jitter+adapt_diag...
/home/damianos/miniconda3/envs/pymc3/lib/python3.7/site-packages/theano/tensor/basic.py:6611:
  result[diagonal_slice] = x
/home/damianos/miniconda3/envs/pymc3/lib/python3.7/site-packages/theano/tensor/basic.py:6611:
  result[diagonal_slice] = x
/home/damianos/miniconda3/envs/pymc3/lib/python3.7/site-packages/theano/tensor/basic.py:6611:
  result[diagonal_slice] = x
Sequential sampling (1 chains in 1 job)
NUTS: [sigma, rho, eta]
  0%1
               | 0/1500 [00:00<?, ?it/s]/home/damianos/miniconda3/envs/pymc3/lib/python3.7/sit/
  result[diagonal_slice] = x
100%|| 1500/1500 [00:18<00:00, 80.84it/s]
Only one chain was sampled, this makes it impossible to run some convergence checks
In [16]: varnames = ['eta', 'rho', 'sigma']
         chain = trace[100:]
         pm.traceplot(chain, varnames);
```



In [17]: pm.summary(chain, varnames).round(4) Out[17]: mean sd mc_error hpd_2.5 hpd_97.5 2.8176 2.8098 0.1533 0.2049 7.8822 eta 0.1272 0.0537 0.0028 0.0510 0.2265 rho 0.0000 0.0002 0.0006 0.0004 0.0014 sigma In [18]: y_pred = [np.random.multivariate_normal(m, S) for m,S in zip(chain['mu_post'][::5], cd for yp in y_pred: plt.plot(test_points, yp, 'r-', alpha=0.1) plt.plot(x, y, 'bo'); plt.xlabel('\$x\$', fontsize=16); plt.ylabel('\$f(x)\$', fontsize=16, rotation=0);



1 Periodic Kernel

```
In [19]: periodic = lambda x, y: np.array([[np.sin((x[i] - y[j])/2)**2 for i in range(len(x))]
In [20]: with pm.Model() as GP_periodic:
    mu = np.zeros(N)
    eta = pm.HalfCauchy('eta', 5)
    rho = pm.HalfCauchy('rho', 5)
    sigma = pm.HalfCauchy('sigma', 5)

P = periodic(x, x)

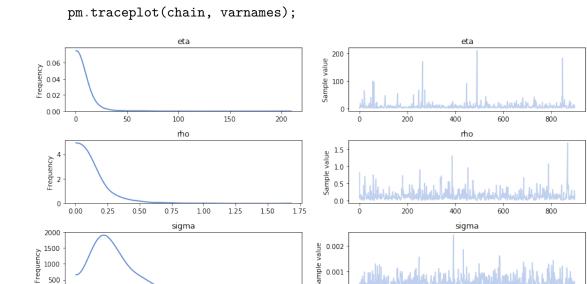
K = tt.fill_diagonal(eta * pm.math.exp(-rho * P), eta + sigma)

obs = pm.MvNormal('obs', mu, cov=K, observed=y)

test_points = np.linspace(0, 10, 100)
    D_pred = periodic(test_points, test_points)
    D_off_diag = periodic(x, test_points)

K_oo = eta * pm.math.exp(-rho * D_pred)
    K_o = eta * pm.math.exp(-rho * D_off_diag)
```

```
mu_post = pm.Deterministic('mu_post', pm.math.dot(pm.math.dot(K_o, tt.nlinalg.mat)
             SIGMA_post = pm.Deterministic('SIGMA_post', K_oo - pm.math.dot(pm.math.dot(K_o, to))
             start = pm.find_MAP()
             trace = pm.sample(1000, start=start,chains=1,njobs=1)
/home/damianos/miniconda3/envs/pymc3/lib/python3.7/site-packages/pymc3/tuning/starting.py:61:
  warnings.warn('find_MAP should not be used to initialize the NUTS sampler, simply call pymc3
/home/damianos/miniconda3/envs/pymc3/lib/python3.7/site-packages/theano/tensor/basic.py:6611:
  result[diagonal_slice] = x
/home/damianos/miniconda3/envs/pymc3/lib/python3.7/site-packages/theano/tensor/basic.py:6611:
  result[diagonal_slice] = x
               | 0/5000 [00:00<?, ?it/s]/home/damianos/miniconda3/envs/pymc3/lib/python3.7/site
  0%1
 result[diagonal_slice] = x
logp = 23.985, ||grad|| = 1.9188: 100%|| 18/18 [00:00<00:00, 839.37it/s]
Auto-assigning NUTS sampler...
Initializing NUTS using jitter+adapt_diag...
Sequential sampling (1 chains in 1 job)
NUTS: [sigma, rho, eta]
100%|| 1500/1500 [00:19<00:00, 75.12it/s]
There was 1 divergence after tuning. Increase `target_accept` or reparameterize.
Only one chain was sampled, this makes it impossible to run some convergence checks
In [21]: varnames = ['eta', 'rho', 'sigma']
         chain = trace[100:]
```



0.0025

0.0005

0.0010

0.0015

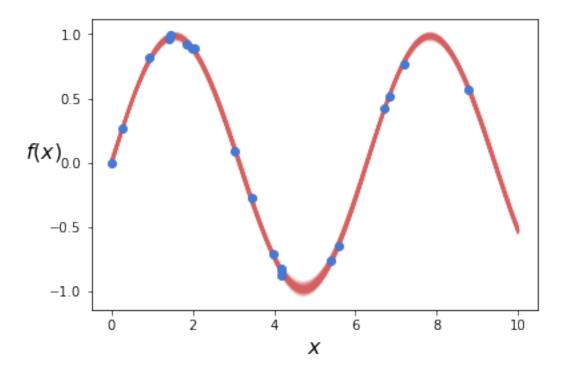
0.0020

In [22]: y_pred = [np.random.multivariate_normal(m, S) for m,S in zip(chain['mu_post'][::5], cd

```
for yp in y_pred:
    plt.plot(test_points, yp, 'r-', alpha=0.1)

plt.plot(x, y, 'bo')
plt.xlabel('$x$', fontsize=16)
plt.ylabel('$f(x)$', fontsize=16, rotation=0);
```

/home/damianos/miniconda3/envs/pymc3/lib/python3.7/site-packages/ipykernel_launcher.py:1: Runt """Entry point for launching an IPython kernel.



In [23]: import sys, IPython, scipy, matplotlib, platform print("This notebook was created on a computer %s running %s and using:\nPython %s\nI

This notebook was created on a computer x86_64 running debian buster/sid and using:

Python 3.7.2

IPython 7.2.0

PyMC3 3.6

NumPy 1.16.0

SciPy 1.2.0

Pandas 0.23.4

Matplotlib 3.0.2

Seaborn 0.9.0

/home/damianos/miniconda3/envs/pymc3/lib/python3.7/site-packages/ipykernel_launcher.py:2: Depre