```
In [399... import numpy as np
         import pandas as pd
         import matplotlib.pyplot as plt
         from tqdm import tqdm
In [400... def train_test_split(X, Y, train_size):
             N = X.shape[0] #assumes data of shape (N,1)
             #shuffle data
             indexes = np.arange(N)
             np.random.seed(42)
             np.random.shuffle(indexes)
             #get indexes
             N_train = int(N*train_size) #train_size is in the range [0,1]
             train_idxs = indexes[:N_train]
             test_idxs = indexes[N_train:]
             #partition data
             return X[train_idxs], X[test_idxs], Y[train_idxs], Y[test_idxs]
         def vandermonde(X, k):
             N = X.shape[0] \# X is assumed to be (N, d). In this case d=1
             PhiX = np.zeros((N, k))
             for i in range(k):
                 PhiX[:, i] = (X**i).flatten() \#PhiX[:, i] has shape (N,) while X**i has shape (N,1) => transform X**i to a row
             return PhiX
         def loss(PhiX, Y, theta, **kwargs):
             return 0.5 * np.linalg.norm(Y - PhiX@theta, 2)**2
         def grad_loss(PhiX, Y, theta, **kwargs):
             return -PhiX.T @ (Y - PhiX@theta) #shape (K, 1) => column vector
         def compute_error(Y, pred):
             return np.mean((Y - pred)**2) # MSE
In [401... | def GD(loss, grad_loss, theta0, data, alpha=0.1, epochs=200, early_stopping=False, tolf=1e-4, tolx=1e-4, **kwargs):
             X, Y = data # X.shape (N, d) --- Y.shape (N, 1)
             f_vals = np.zeros((epochs+1,))
             grad_norms = np.zeros((epochs+1,))
             errors = np.zeros((epochs+1,))
             f_vals[0] = loss(X, Y, theta0, **kwargs)
             grad_norms[0] = np.linalg.norm(grad_loss(X, Y, theta0, **kwargs))
             errors[0] = compute_error(Y, X@theta0)
             theta = theta0
             for epoch in tqdm(range(epochs)):
                 gradient = grad_loss(X, Y, theta, **kwargs) # shape (d, 1)
                 new_theta = theta - alpha * gradient # column vector
                 # check stopping conditions if we are using early stopping
                 if early_stopping==True and ((np.linalg.norm(grad_loss(X, Y, new_theta, **kwargs))<(tolf*np.linalg.norm(theta))) or (np.linalg.norm(new_theta-theta)<tolx)):
                 #update current theta
                 theta = new_theta
                 # append values for plots
                 f_vals[epoch+1] = loss(X, Y,theta, **kwargs)
                 grad norms[epoch+1] = np.linalq.norm(gradient)
                 errors[epoch+1] = compute_error(Y, X@theta)
             return theta, f_vals, grad_norms,errors, epoch
```

```
In [402... | def SGD(loss, grad_loss, theta0, data, alpha=0.1, epochs=50, batch_size=10, early_stopping=False, tolf=1e-4, tolx=1e-4, **kwargs):
             X, Y = data # X.shape (N, d) --- Y.shape (N, 1)
             N = X.shape[0]
             idxs = np.arange(0, N)
             loss_vals = np.zeros((epochs+1,))
             grad_norms = np.zeros((epochs+1,))
             errors = np.zeros((epochs+1,))
             loss_vals[0] = loss(X, Y, theta0, **kwargs)
             grad_norms[0] = np.linalg.norm(grad_loss(X, Y, theta0, **kwargs))
             errors[0] = compute_error(Y, X@theta0)
             theta = theta0
             stop = False # stop is a boolean function to break the outer loop
             for epoch in tqdm(range(epochs)): #iter over epochs
                 if stop: #end the training process if stopping condition has occurred
                     break
                 np.random.shuffle(idxs)
                 grad_loss_vec = []
                 for batch_start in range(0, N, batch_size): #iter over batches within an epoch
                     batch_end = min(batch_start + batch_size, N)
                     batch_idxs = idxs[batch_start:batch_end]
                     minibatchX = X[batch_idxs]
                     minibatchY = Y[batch_idxs]
                     gradient = grad_loss(minibatchX, minibatchY, theta, **kwargs)
                     new_theta = theta - alpha * gradient
                     if early_stopping==True and ((np.linalg.norm(grad_loss(X, Y, new_theta, **kwargs))<(tolf*np.linalg.norm(theta))) or (np.linalg.norm(new_theta-theta)<tolx)):</pre>
                         stop = True #stop the iteration over thee epoches at the beginning of the next epoch
                         break
                     theta = new_theta
                     grad_loss_vec.append(np.linalg.norm(gradient, 2))
                 loss_vals[epoch+1] = loss(X, Y, theta, **kwargs)
                 grad_norms[epoch+1] = np.mean(grad_loss_vec)
                 errors[epoch+1] = compute_error(Y, X@theta)
             return theta, loss_vals, grad_norms, errors, epoch
In [403... def cholesky(PhiX, Y):
             L = np.linalg.cholesky(PhiX.T @ PhiX)
             z = np.linalg.solve(L, PhiX.T@Y)
             theta = np.linalg.solve(L.T, z)
             return theta
```

• For a given value of K, write three Python functions computing θ_{MLE} , i.e. the optimal parameters obtained by optimizing the MLE-related loss function with Gaussian assumption on the likelihood $p_{\theta}(y|x)$, by Gradient Descent, Stochastic Gradient Descent (with a batch_size = 5), and Normal Equations method with Cholesky Decomposition.

```
In [445... df = pd.read_csv("./poly_regression_large.csv")
         X, Y = df["x"].to_numpy().reshape(-1,1), df["y"].to_numpy().reshape(-1,1) #shape (1000,1) --> columns
         Xtrain, Xtest, Ytrain, Ytest = train_test_split(X, Y, train_size=0.8) #shapes (N,1)
         K = 6
         PhiX = vandermonde(Xtrain, K) #shape (N, k)
         #PhiX = np.vander(Xtrain.ravel(), N=K, increasing=True)
         theta0 = np.zeros((K, 1)) #column vector
         theta_gd, loss_gd, grad_gd, error_gd, epoch_gd = GD(
             loss,
             grad_loss,
             data=(PhiX,Ytrain),
             epochs=5000,
              theta0=theta0,
             alpha=0.001,
             early_stopping=True,
             tolf=1e-6,
             tolx=1e-6
         theta_sgd, loss_sgd, grad_sgd, error_sgd, epoch_sgd = SGD(
             loss,
             grad_loss,
             data=(PhiX,Ytrain),
             batch_size=5,
             epochs=5000,
             alpha=0.001,
             theta0=theta0,
              early_stopping=True,
             tolf=1e-6,
             tolx=1e-6
         theta_cholesky = cholesky(PhiX, Ytrain)
                        [| 5000/5000 [00:00<00:00, 15266.53it/s]
                      | | 3885/5000 [00:16<00:04, 235.92it/s]
```

• Compare the performance of the three regression model computed above. In particular, if (X_{test}, Y_{test}) is the test set from the poly_regression_large.csv dataset, for each of the model, compute:

$$Err = rac{1}{N_{test}} \sum_{i=1}^{N_{test}} (f_{ heta}(x^i) - y^i)^2,$$

where N_{test} is the number of elements in the test set, (x^i, y^i) are the input and output elements in the test set. Comment the performance of the three models.

```
In [446... PhiX_test = vandermonde(Xtest, K)

test_err_gd = compute_error(Ytest, PhiX_test@theta_gd)

test_err_sgd = compute_error(Ytest, PhiX_test@theta_sgd)

test_err_cho = compute_error(Ytest, PhiX_test@theta_cholesky)

print(f"Test MSE GD: {test_err_gd}\nTest MSE SGD: {test_err_sgd}\nTest MSE Cholesky: {test_err_cho}")

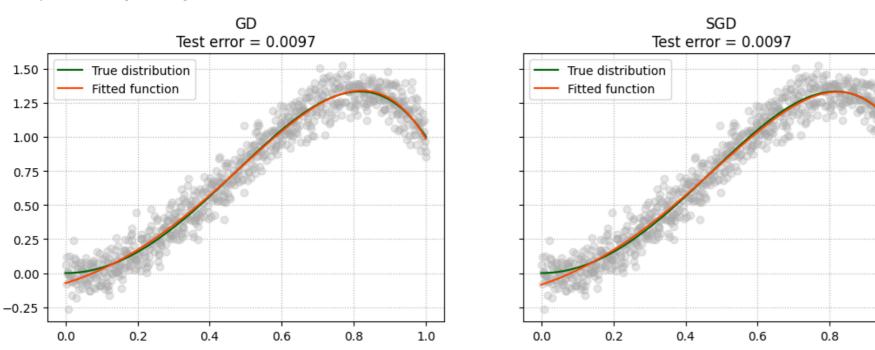
Test MSE GD: 0.00971010047760351

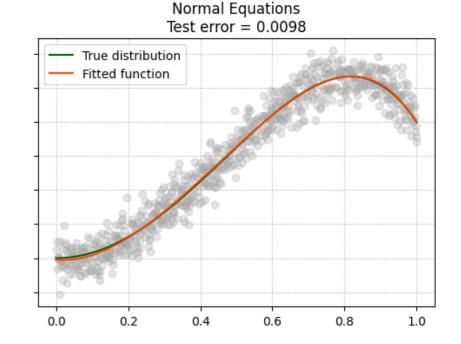
Test MSE SGD: 0.009703817960597794

Test MSE Cholesky: 0.009844990396114582
```

```
In [447... fig, axs = plt.subplots(figsize=(20,4) ,ncols=3, sharey=True)
         PhiX_test = vandermonde(Xtest, K)
         ax = axs[0]
         ax.set_title(f'GD\nTest error = {test_err_gd:.4f}')
         ax.scatter(Xtrain, Ytrain, color='darkgrey', alpha=0.3)
         ax.grid(linestyle=':')
         ax.plot(X, 4*X**2 - 3*X**4, color='darkgreen', label='True distribution')
         ax.plot(np.linspace(0,1, 200), vandermonde(np.linspace(0,1,200), K)@theta_gd, color='orangered', label='Fitted function')
         ax.legend()
         ax = axs[1]
         ax.set_title(f'SGD\nTest error = {test_err_sgd:.4f}')
         ax.scatter(Xtrain, Ytrain, color='darkgrey', alpha=0.3)
         ax.grid(linestyle=':')
         ax.plot(X, 4*X**2 - 3*X**4, color='darkgreen', label='True distribution')
         ax.plot(np.linspace(0,1, 200), vandermonde(np.linspace(0,1,200), K)@theta_sgd, color='orangered', label='Fitted function')
         ax.legend()
         ax = axs[2]
         ax.set_title(f'Normal Equations\nTest error = {test_err_cho:.4f}')
         ax.scatter(Xtrain, Ytrain, color='darkgrey', alpha=0.3)
         ax.grid(linestyle=':')
         ax.plot(X, 4*X**2 - 3*X**4, color='darkgreen', label='True distribution')
         ax.plot(np.linspace(0,1, 200), vandermonde(np.linspace(0,1,200), K)@theta_cholesky, color='orangered', label='Fitted function')
         ax.legend()
```

Out[447]: <matplotlib.legend.Legend at 0x2838b39d0>





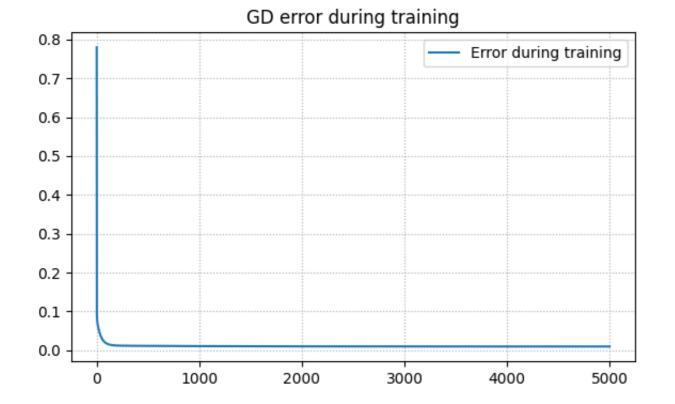
```
In [451... fig, axs = plt.subplots(figsize=(15,4) ,ncols=2, sharey=True)

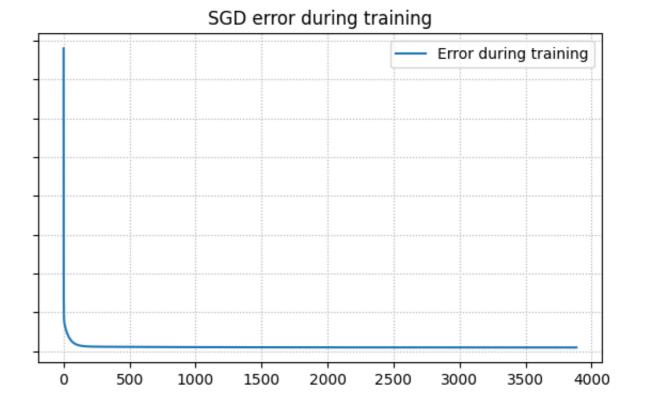
ax = axs[0]
ax.set_title(f'GD error during training')
ax.plot(error_gd[error_gd!=0], label='Error during training')
ax.grid(linestyle=':')
ax.legend()

ax = axs[1]
ax.set_title(f'SGD error during training')
ax.plot(error_sgd[error_sgd!=0], label='Error during training')
ax.grid(linestyle=':')
ax.legend()
```

1.0

Out[451]: <matplotlib.legend.Legend at 0x2839da9e0>



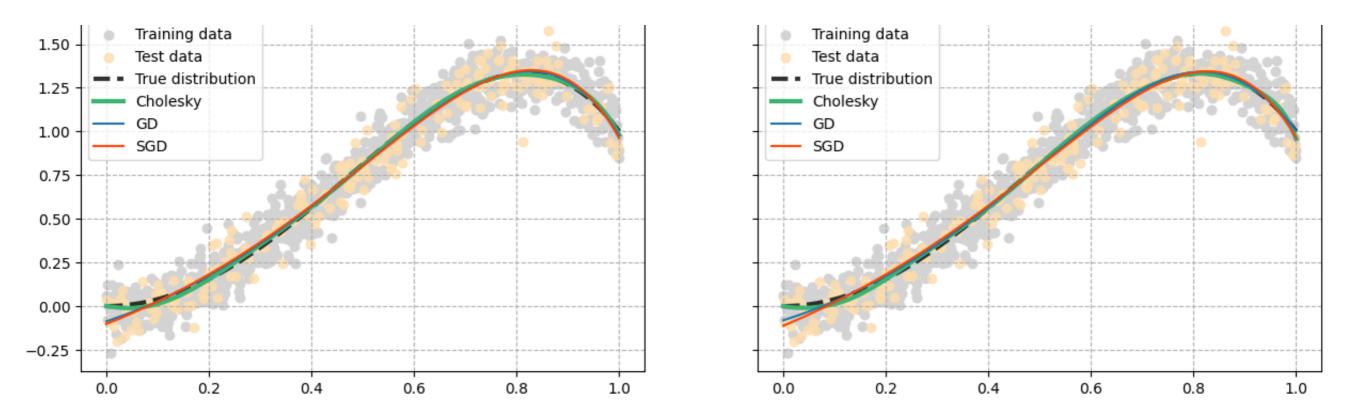


As we can see from the experiments above, all three methods work very well since they show very low test error. However, even though the test error is almost the same for all three methods, the function estimated by the normal equations better resembles the true function from which the data were generated. Furthermore, tolf and tolx were carefully selected to avoid underfitting. Indeed, too high values of tolf and tolf lead to good functions (in the sense that the test error is very low) but they do not fit the data well. Apparently, we can stop training after very few epochs but this would result in a very poor approximation of the true function.

• For different values of K, plot the training datapoints and the test datapoints with different colors and visualize (as a continuous line) the three learned regression model $f_{\theta}(x)$. Comment the results.

```
In [455... | k_vals = np.array([3,4,5,6,7,8,10,12])
         nrows = 4
         ncols = 2
         theta_vals = []
         fig, axs = plt.subplots(figsize=(15,20), ncols=ncols, nrows=nrows, sharex=True, sharey=True)
         iter = 0
         for i in range(nrows):
             for j in range(ncols):
                 k = k_vals[iter]
                 print(f'\nTraining GD and SGD for k = \{k\}')
                 PhiX = vandermonde(Xtrain, k) \#shape (N, k)
                 theta0 = np.zeros((k, 1)) #column vector
                 theta_gd, loss_gd, grad_gd,error_gd, epoch_gd = GD(
                     loss,
                    grad_loss,
                    data=(PhiX,Ytrain),
                    epochs=7000,
                    theta0=theta0,
                    alpha=0.001,
                    early_stopping=True,
                    tolf=1e-6,
                    tolx=1e-6
                 theta_sgd, loss_sgd, grad_sgd, error_sgd, epoch_sgd = SGD(
                    loss,
                    grad_loss,
                    data=(PhiX,Ytrain),
                    batch_size=5,
                    epochs=5000,
                    alpha=0.001,
                    theta0=theta0,
                    early_stopping=True,
                    tolf=1e-6,
                    tolx=1e-6
                 theta_cholesky = cholesky(PhiX, Ytrain)
                 theta_vals.append([theta_gd, theta_sgd, theta_cholesky])
                 ax = axs[i][j]
                 ax.scatter(Xtrain, Ytrain, color='lightgray', label='Training data', alpha=1)
                 ax.scatter(Xtest, Ytest, color='navajowhite', label='Test data', alpha=0.8)
                 ax.plot(X, 4*X**2 - 3*X**4, color='black', label='True distribution', linewidth=3, alpha=0.8, linestyle='--')
                 ax.plot(np.linspace(0,1, 100), vandermonde(np.linspace(0,1,100), k)@theta_cholesky, color='mediumseagreen', label='Cholesky', linewidth=3)
                 ax.plot(np.linspace(0,1, 100), vandermonde(np.linspace(0,1,100), k)@theta_gd, color='tab:blue', label='GD')
                 ax.plot(np.linspace(0,1, 100), vandermonde(np.linspace(0,1,100), k)@theta_sgd, color='orangered', label='SGD')
                 ax.set_title(f"$K = {k}$")
                 ax.grid(linestyle='--')
                 ax.legend()
                 iter += 1
         Training GD and SGD for k = 3
                        | 4057/7000 [00:00<00:00, 17554.04it/s]
                  | 5000/5000 [00:22<00:00, 218.27it/s]
         Training GD and SGD for k = 4
                      7000/7000 [00:00<00:00, 22248.07it/s]
                      ■ | 4540/5000 [00:17<00:01, 253.39it/s]
         Training GD and SGD for k = 5
         100% | 7000/7000 [00:00<00:00, 21558.31it/s]
          56% | 2782/5000 [00:11<00:09, 239.34it/s]
         Training GD and SGD for k = 6
         100% | 7000/7000 [00:00<00:00, 20956.28it/s]
         100% | 5000/5000 [00:21<00:00, 236.77it/s]
         Training GD and SGD for k = 7
         100% | 7000/7000 [00:00<00:00, 20168.86it/s]
           5%|| | 266/5000 [00:01<00:20, 230.39it/s]
         Training GD and SGD for k = 8
         100% | 7000/7000 [00:00<00:00, 21693.93it/s]
          87%| 4346/5000 [00:17<00:02, 243.78it/s]
```

Training GD and SGD for k = 10

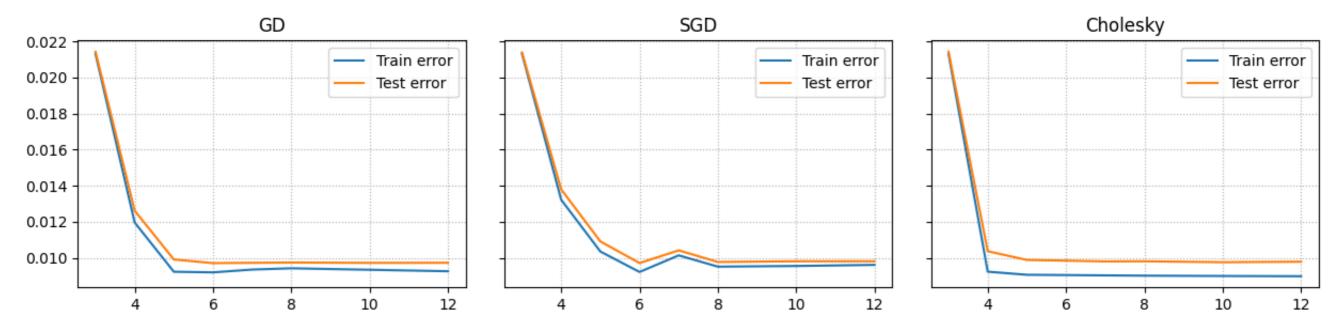


For k=3 all the three methods provide the same curve. The curve predicted underfits the data, meaning that it is not flexible enough to capture the complexity of the data. With k=4, GD and SGD still underfitting the data while the curve predicted by Normal Equations resambles pretty well the data. The best configuration probably is for k=6: all the three methods provide approximatively the same curve (maybe Normal equations method is slightly better). For higher values of k (for $k\geq 8$ for example), the three methods seem to "overfit" a little bit especially Normal equations method

 \bullet For increasing values of K, compute the training and test error as discussed above. Plot the two errors with respect to K. Comment the results.

```
In [456... test err qd = []
         test_err_sgd = []
         test_err_cho = []
         train_err_gd = []
         train err sqd = []
         train_err_cho = []
         iter = 0
         k_{vals} = np.array([3,4,5,6,7,8,10,12])
         ncols = 3
         nrows = 1
         for theta_gd, theta_sgd, theta_cho in theta_vals:
             k = k_vals[iter]
             test err gd.append(compute error(Ytest, vandermonde(Xtest, k)@theta gd))
             test_err_sgd.append(compute_error(Ytest, vandermonde(Xtest, k)@theta_sgd))
             test_err_cho.append(compute_error(Ytest, vandermonde(Xtest, k)@theta_cho))
             train_err_gd.append(compute_error(Ytrain, vandermonde(Xtrain, k)@theta_gd))
             train_err_sgd.append(compute_error(Ytrain, vandermonde(Xtrain, k)@theta_sgd))
             train_err_cho.append(compute_error(Ytrain, vandermonde(Xtrain, k)@theta_cho))
             iter += 1
         fig, axs = plt.subplots(figsize=(15,3), ncols=ncols, nrows=nrows, sharey=True)
         fig.subplots_adjust(wspace=0.1)
         ax = axs[0]
         ax.plot(k_vals, train_err_gd, label='Train error')
         ax.plot(k_vals, test_err_gd, label='Test error')
         ax.set_title('GD')
         ax.grid(linestyle=':')
         ax.legend()
         ax = axs[1]
         ax.plot(k_vals, train_err_sgd, label='Train error')
         ax.plot(k_vals, test_err_sgd, label='Test error')
         ax.set title('SGD')
         ax.grid(linestyle=':')
         ax.legend()
         ax = axs[2]
         ax.plot(k_vals, train_err_cho, label='Train error')
         ax.plot(k_vals, test_err_cho, label='Test error')
         ax.set_title('Cholesky')
         ax.grid(linestyle=':')
         ax.legend()
```

Out[456]: <matplotlib.legend.Legend at 0x2863243d0>



All the three methods have a similar behavior: they have high error valus for smaller values of k. Normal equations is the method with the fastest convergenge w.r.t. k, in the sense that, even for k=4, it has an acceptable test error value. For all the methods, strating from k=6 both the test error and the train error reaches a plateau, meaning that a degree of the polynomial function greater than 6 is not necessary. The error of SGD reaches the minimum for k=6 and then slightly increases suggesting a little overfitting. Moreover, it oscillates and this is due to the stochastic nature of SGD.

```
def loss_map(PhiX, Y, theta, **kwargs):
    lamb = kwargs['lamb']
    return 0.5 * np.linalg.norm(Y - PhiX@theta, 2)**2 + 0.5 * lamb * np.linalg.norm(theta)**2

def grad_loss_map(PhiX, Y, theta, **kwargs):
    lamb = kwargs['lamb']
    return -PhiX.T @ (Y - PhiX@theta) + lamb * theta #shape (K, 1) => column vector
```

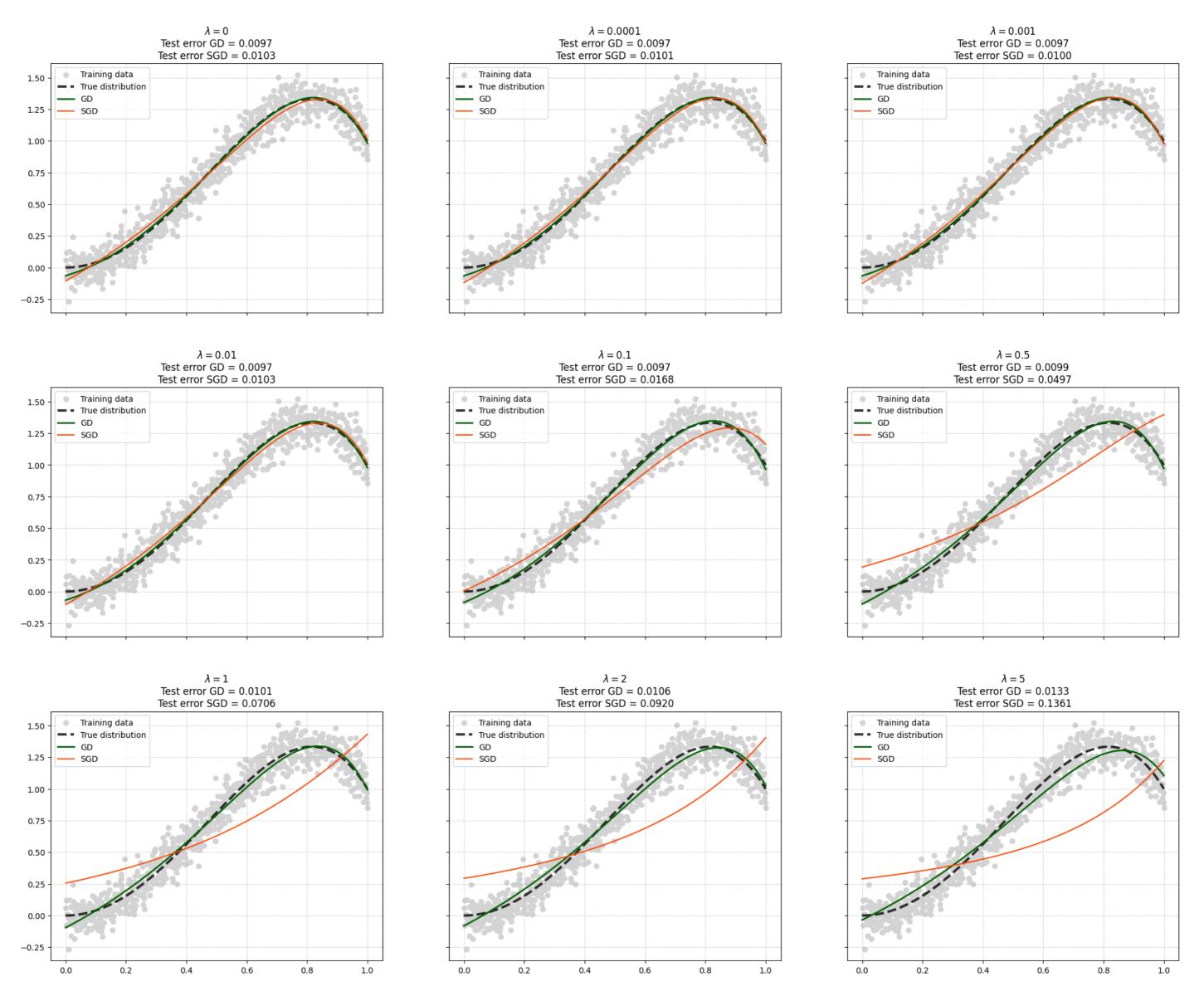
```
Repeat the same experiments by considering the MAP formulation with Gaussian assumption on the prior term p(\theta). Set K=8 and test different values of \lambda>0 in the experiments.
In [463... df = pd.read_csv("./poly_regression_large.csv")
         X, Y = df["x"].to_numpy().reshape(-1,1), df["y"].to_numpy().reshape(-1,1) #shape (1000,1) --> columns
         Xtrain, Xtest, Ytrain, Ytest = train_test_split(X, Y, train_size=0.8) #shapes (N,1)
         K = 8
         PhiX = vandermonde(Xtrain, K) #shape (N, k)
         #PhiX = np.vander(Xtrain, N=K, increasing=True)
         theta0 = np.zeros((K, 1)) #column vector
         theta_gd, loss_gd, grad_gd,error_gd, epoch_gd = GD(
             loss_map,
             grad_loss_map,
             data=(PhiX,Ytrain),
             epochs=20000,
             theta0=theta0,
             alpha=0.001,
             lamb=0.01,
             early_stopping=True,
             tolf=1e-5,
             tolx=1e-5
         theta_sgd, loss_sgd, grad_sgd, error_sgd, epoch_sgd = SGD(
             loss_map,
             grad_loss_map,
             data=(PhiX,Ytrain),
             batch_size=10,
             epochs=20000,
             alpha=0.001,
             lamb=0.01,
             theta0=theta0,
             early_stopping=True,
             tolf=1e-5,
             tolx=1e-5
         theta_cholesky = cholesky(PhiX, Ytrain)
                     20000/20000 [00:00<00:00, 20054.35it/s]
         2%|| | 375/20000 [00:00<00:44, 445.20it/s]
In [464... PhiX_test = vandermonde(Xtest, K)
         test_err_gd = compute_error(Ytest, PhiX_test@theta_gd)
         test_err_sgd = compute_error(Ytest, PhiX_test@theta_sgd)
         test_err_cho = compute_error(Ytest, PhiX_test@theta_cholesky)
         print(f"Test MSE GD: {test_err_gd}\nTest MSE SGD: {test_err_sgd}\nTest MSE Cholesky: {test_err_cho}")
         Test MSE GD: 0.009702565942922842
         Test MSE SGD: 0.01006650267760104
         Test MSE Cholesky: 0.009810708635243658
```

• Compare the different values of $\lambda > 0$ tested,

```
In [465... lambda_vals = [0, 1e-4, 1e-3, 1e-2, 1e-1, 0.5, 1, 2, 5]
         nrows = 3
         ncols = 3
         k = 8
         PhiX = vandermonde(Xtrain, k) #shape (N,k)
         theta0 = np.zeros((k, 1)) #column vector
         theta_vals_gd = []
         theta_vals_sgd = []
         test_errs_gd = []
         test_errs_sgd = []
         train_errs_gd = []
         train_errs_sgd = []
         fig, axs = plt.subplots(figsize=(25,20), ncols=ncols, nrows=nrows, sharex=True, sharey=True)
         fig.subplots_adjust(hspace=0.3)
         iter = 0
         for i in range(nrows):
             for j in range(ncols):
                 print(f'\nTraining GD and SGD for lambda = {lambda_vals[iter]}')
                 theta_gd, loss_gd, grad_gd,error_gd, epoch_gd = GD(
                     loss_map,
                     grad_loss_map,
                     data=(PhiX,Ytrain),
                     epochs=20000,
                     theta0=theta0,
                     alpha=0.001,
                     lamb=lambda_vals[iter],
                     early_stopping=True,
                     tolf=1e-5,
                     tolx=1e-5
                 theta_sgd, loss_sgd, grad_sgd, error_sgd, epoch_sgd = SGD(
                     loss_map,
                     grad_loss_map,
                     data=(PhiX,Ytrain),
                     batch_size=10,
                     epochs=20000,
                     alpha=0.001,
                     lamb=lambda_vals[iter],
                     theta0=theta0,
                     early_stopping=True,
                     tolf=1e-5,
                     tolx=1e-5
                 theta_vals_gd.append(theta_gd)
                 theta_vals_sgd.append(theta_sgd)
                 train_errs_gd.append(compute_error(Ytrain, PhiX@theta_gd))
                 train_errs_sgd.append(compute_error(Ytrain, PhiX@theta_sgd))
                 PhiX_test = vandermonde(Xtest, k)
                 test_err_gd = compute_error(Ytest, PhiX_test@theta_gd)
                 test_errs_gd.append(test_err_gd)
                 test_err_sgd = compute_error(Ytest, PhiX_test@theta_sgd)
                 test_errs_sgd.append(test_err_sgd)
                 ax = axs[i][j]
                 ax.set_title(f'$\lambda = {\lambda_vals[iter]}$\nTest error GD = {\test_err_gd:.4f}\nTest error SGD = {\test_err_sgd:.4f}\)
                 ax.scatter(Xtrain, Ytrain, color='lightgray', label='Training data', alpha=1)
                 ax.plot(X, 4*X**2 - 3*X**4, color='black', label='True distribution', linewidth=3, alpha=0.8, linestyle='--')
                 ax.plot(np.linspace(0,1, 100), vandermonde(np.linspace(0,1,100), k)@theta_gd, color='darkgreen', label='GD', linewidth=2)
                 ax.plot(np.linspace(0,1, 100), vandermonde(np.linspace(0,1,100), k)@theta_sgd, color='orangered', label='SGD')
                 ax.legend()
                 ax.grid(linestyle=':')
                 iter += 1
```

Training GD and SGD for lambda = 0

Training GD and SGD for lambda = 0.0001 100%| 20000/20000 [00:01<00:00, 19560.71it/s] 1%| | 178/20000 [00:00<00:44, 449.35it/s] Training GD and SGD for lambda = 0.001 100%| 20000/20000 [00:01<00:00, 19232.16it/s] 1%| | 248/20000 [00:00<00:43, 450.88it/s] Training GD and SGD for lambda = 0.01 100%| 20000/20000 [00:00<00:00, 20502.23it/s] 1%| | 182/20000 [00:00<00:43, 453.65it/s] Training GD and SGD for lambda = 0.1 94%| | 18822/20000 [00:00<00:00, 20459.32it/s] 10%|■ | 2077/20000 [00:04<00:39, 451.66it/s] Training GD and SGD for lambda = 0.525%| | 4978/20000 [00:00<00:00, 20245.09it/s] 48% | 9629/20000 [00:21<00:23, 450.09it/s] Training GD and SGD for lambda = 1 13%| | 2605/20000 [00:00<00:00, 19513.41it/s] 20000/20000 [00:45<00:00, 440.94it/s] 100% Training GD and SGD for lambda = 27%|▮ | 1399/20000 [00:00<00:00, 19019.47it/s] | 4517/20000 [00:10<00:36, 425.42it/s] 23%| Training GD and SGD for lambda = 53%|| | 643/20000 [00:00<00:01, 19224.84it/s] 40% | 7956/20000 [00:18<00:28, 423.36it/s]

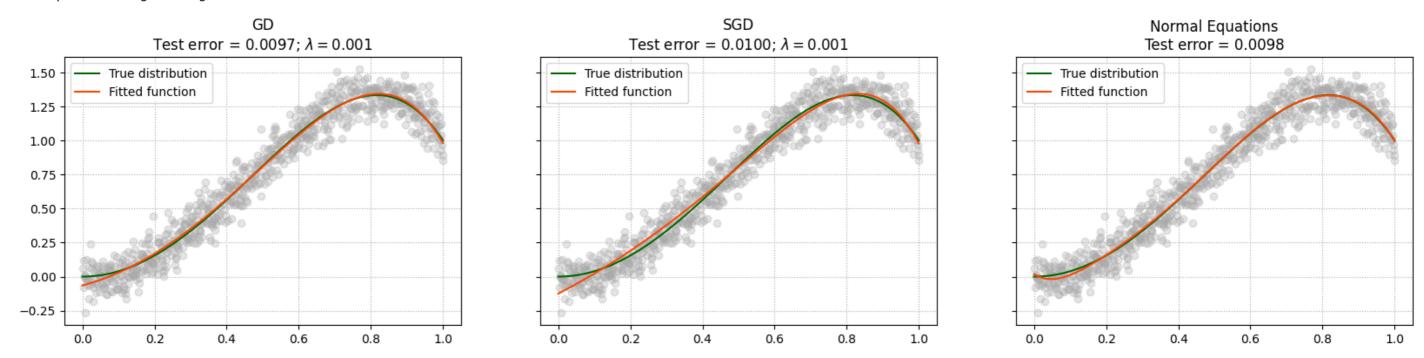


MAP can be considered as a form of regularization for MLE. Regularization is a set of methods for reducing overfitting of a model. Since MLE (MAP with $\lambda=0$) doesn't suffer of overfitting, MAP with high value of λ leads to underfitted models. As we can see from the graph above, for small values of λ (that is $\lambda\leq0.1$) the fitted curve approximate pretty well the data. This is not true for highr values of λ . This phenomenon is more accentuated on SGD because SGD uses a small batch, so each time the loss is computed the term $\lambda||\theta||_1$ dominates over the term $\frac{1}{2}||\Phi(X)\theta-Y||_2^2$

ullet Compare the three optimization method used to obtain $heta_{MAP}$ (i.e. GD, SGD and Normal Equations),

```
In [466... fig, axs = plt.subplots(figsize=(20,4) ,ncols=3, sharey=True)
         ax = axs[0]
         ax.set_title(f'GD\nTest error = {compute_error(Ytest, PhiX_test@theta_vals_gd[2]):.4f}; $\lambda={lambda_vals[2]}$')
         ax.scatter(Xtrain, Ytrain, color='darkgrey', alpha=0.3)
         ax.grid(linestyle=':')
         ax.plot(X, 4*X**2 - 3*X**4, color='darkgreen', label='True distribution')
         ax.plot(np.linspace(0,1, 200), vandermonde(np.linspace(0,1,200), K)@theta_vals_gd[2], color='orangered', label='Fitted function')
         ax.legend()
         ax = axs[1]
         ax.set_title(f'SGD\nTest error = {compute_error(Ytest, PhiX_test@theta_vals_sgd[2]):.4f}; $\lambda={lambda_vals[2]}$')
         ax.scatter(Xtrain, Ytrain, color='darkgrey', alpha=0.3)
         ax.grid(linestyle=':')
         ax.plot(X, 4*X**2 - 3*X**4, color='darkgreen', label='True distribution')
         ax.plot(np.linspace(0,1, 200), vandermonde(np.linspace(0,1,200), K)@theta_vals_sgd[2], color='orangered', label='Fitted function')
         ax.legend()
         ax = axs[2]
         ax.set_title(f'Normal Equations\nTest error = {test_err_cho:.4f}')
         ax.scatter(Xtrain, Ytrain, color='darkgrey', alpha=0.3)
         ax.grid(linestyle=':')
         ax.plot(X, 4*X**2 - 3*X**4, color='darkgreen', label='True distribution')
         ax.plot(np.linspace(0,1, 200), vandermonde(np.linspace(0,1,200), K)@theta_cholesky, color='orangered', label='Fitted function')
         ax.legend()
```

Out[466]: <matplotlib.legend.Legend at 0x28b7ed3f0>



In general GD+MAP behaves better than SGD+MAP because GD is less prone to overfitting. Normal equations method does not depend on λ but behaves preety good. In this case normal equations method resambles true data almost perfectly. It is the same for GD that shows the lowest test error.

- Compare the results obtained by $heta_{MLE}$ vs $heta_{MAP}$.

Since MLE with K=6 does not overfit the data, adding regularization doesn't help the model to reach a better fitting. In general MLE works pretty good with all the three methods, while MAP doesn't work with high values of λ . With small values of λ it achieves the same results of MLE. For all these reasons, with this dataset and with the polynomial model the best choiche is MLE.