Function Approximation

1: MLP approximation:

Algorithm: MLP with gradient descent, error function Least square **Activation Function:** 1: Sigmoid activation function in hidden layer

2. No activation in output layer

Loss function: Least square

Parameters to tune:

• Learning rate

• Number of hidden units

Observations and Analysis:

For cross validation, I have used 90% training data as training set and remaining 10% for testing. Analysis is given with the graphs:

Optimal parameters:

| mg85 | |
|-----------------------------|--|
| | |
| hid - 5 | |
| epoch - 600 | |
| learning rate - 1e-4 | |
| | |
| Training Error - 0.22097608 | |
| Testing Error - 0.54073818 | |

| bj |
|----------------------------|
| |
| hid - 4 |
| epoch - 400 |
| learning rate - 1e-2 |
| |
| Training Error - 0.020174 |
| Testing Error - 0.03007362 |

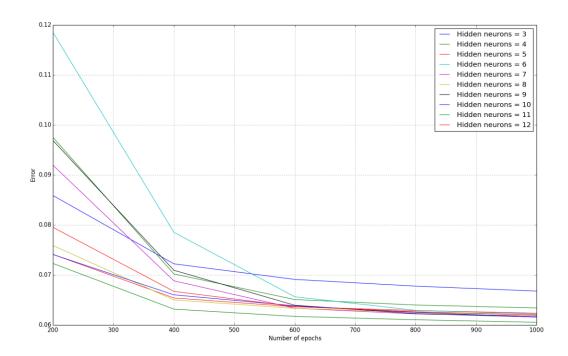
| SI |
|-----------------------------|
| |
| hid - 5 |
| epoch - 600 |
| learning rate - 1e-5 |
| |
| Training Error - 0.06300885 |
| Testing Error - 0.98317534 |

Graphs:

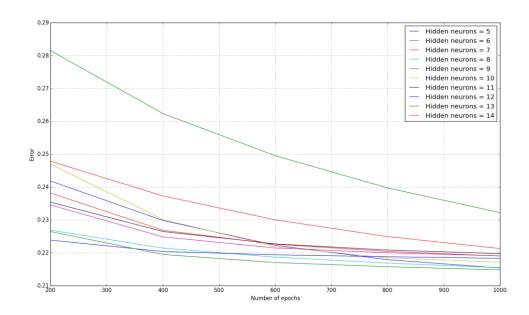
As number of hidden units increases the error is decreasing. But the decrease in error is not significant after a certain number of units (On the test data). After sometime it may be increase on test data.

Number of epochs: The main problem in gradient descent algorithm is that it is susceptible to local optima. It may get stuck on an optimum that may not be even close to the global optima. Here random initialization becomes savior. We have to run the code for 4-5 times and then we saved the best one.

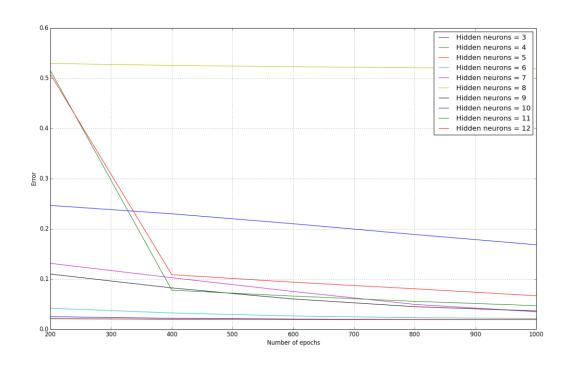
SI:



MG85:



BJ:



2: RBF approximation:

Algorithm: RBF with Pseudo Inverse with clustering and gradient descent different

version.

Activation Function: 1: RBF kernel in the first layer

Loss function: No need Parameters to tune:

Number of Clusters

Algorithm:

Step:1 Normalized the features

Step:2 K-means Clustering, sigma initialized to a random number near 0.5

Step:3 Use pseudo inverse to learn weights in one-shot.

Observations and Analysis:

For cross validation I have used 90% training data as training set and remaining 10% for testing.

- Number of cluster centers
 - Number of cluster centers K was first initialized to (No. features)*2
 - Then I increased the K as long as I got higher accuracy on validation data. I have used a greedy technique in which for more accuracy I have increased K as much as I can. The stopping criteria for learning is (Training error-validation error <= 0.25).
 - For better accuracy we can do more computations. Here main problem is that we are trying to invert a matrix. Taking inverse of a matrix is computationally very expensive algorithm. So I have increased K but not too much.
 - That's why number of clusters are more than the thumb rule we have discussed in class.
- After that for visualization purpose we have plotted error vs epochs (A different version of RBF approx which is based on gradient descent)
- Other technique to plot the True function and the approximated function in same plot and see the actual fitting.

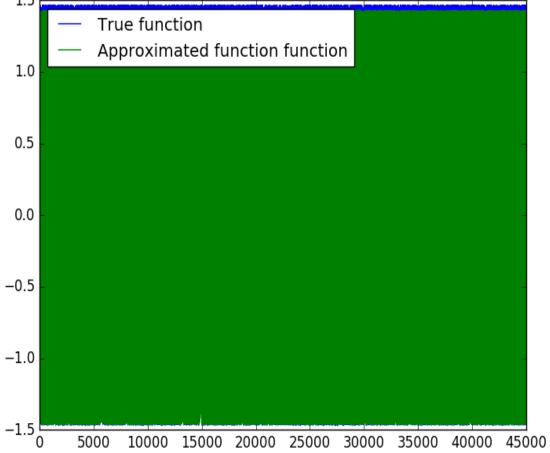
How am I choosing the parameters and Plots:

SI dataset: SI dataset is a very simple function to approximate. The data is given in its normalized form and function is not very steeply changing.

| DATASET | | SI | |
|--------------------|----|----------|----------|
| Number of clusters | | Train | test |
| | 5 | - | - |
| | 10 | 0.022456 | 0.022455 |
| | 20 | 0.015076 | 0.015155 |
| | 40 | 0.01073 | 0.010792 |

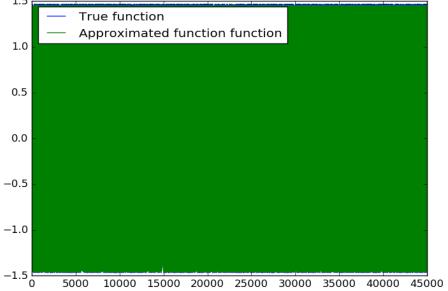
Using 10 clusters: Here you can see that this model so accurately approximates the function. The green (approximated) function is totally overlaps the True function. (On Test data set.)

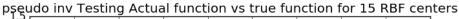


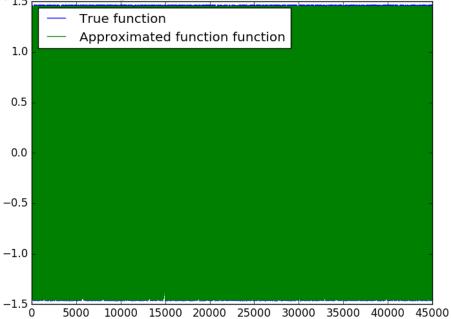


After Increasing number of RBF clusters: The error decreased by a very small amount. But it is accurately approximates the function.

pseudo inv Training Actual function vs true function for 15 RBF centers



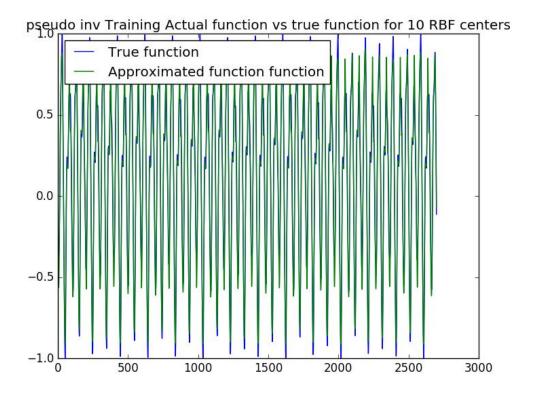


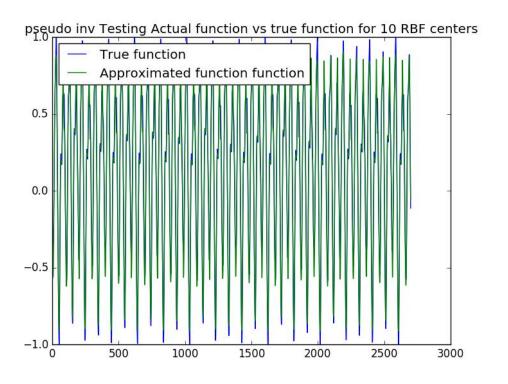


MG85 dataset:

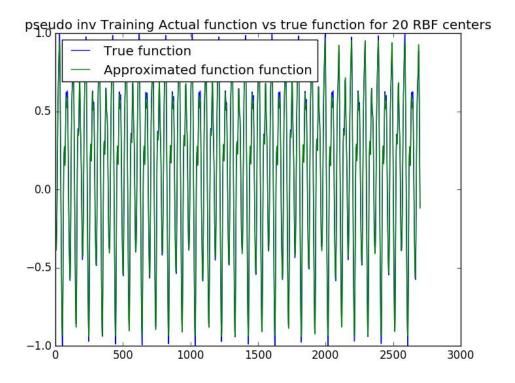
| DATASET | | mg85 | |
|--------------------|----|----------|----------|
| Number of clusters | | train | test |
| | 5 | 1 | - |
| | 10 | 0.092433 | 0.093222 |
| | 20 | 0.042252 | 0.043253 |
| | 40 | 0.027819 | 0.029408 |

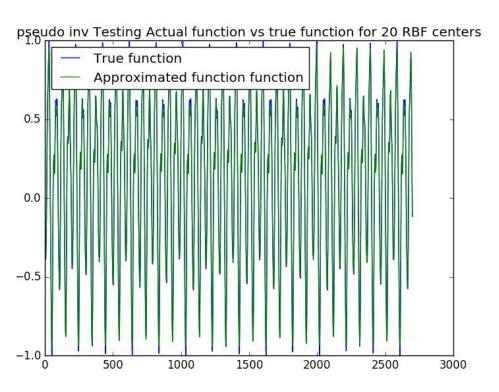
10 neurons:



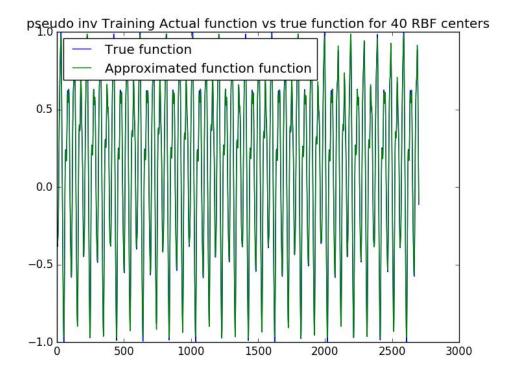


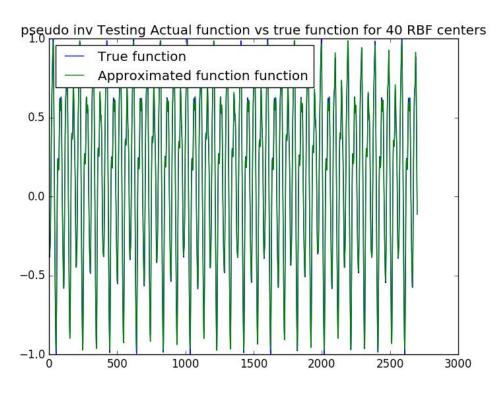
20 neurons:





40 neurons:

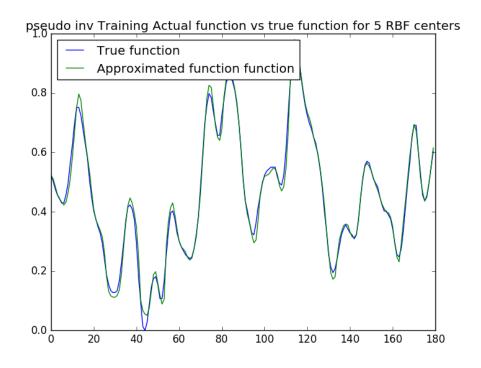


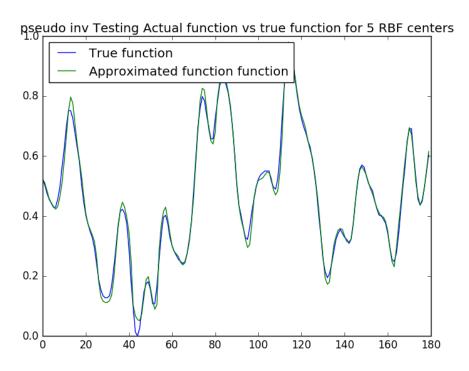


BJ dataset:

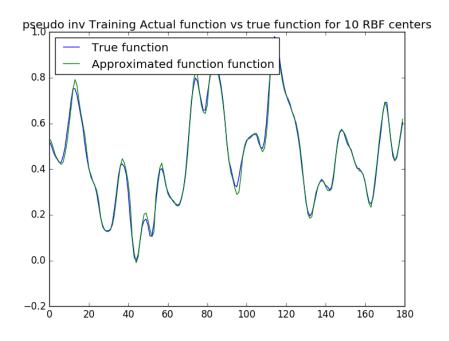
| DATASET | | | bj |
|--------------------|----|----------|----------|
| Number of clusters | | train | test |
| | 5 | 0.021595 | 0.203708 |
| | 10 | 0.01893 | 0.206543 |
| | 20 | 0.016416 | 0.198388 |
| | 40 | 0.015671 | 0.213052 |

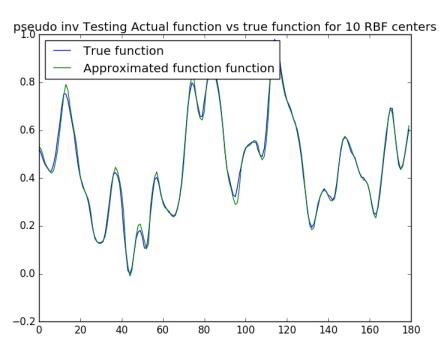
5 Neurons:





10 Neurons:





20 Neurons:

