# Strategy (same for both datasets):

Master node, rank==0 is

- loading the datasets
- Normalizing if needed,
- Splitting data into train and test,
- We'll concatenate y\_train and x\_train in one array so it is easier to send to other workers.

A fixed seed number is considered to provide the same random number every time. We make random numbers for weights in a specific range and size depending on the dataset columns, e.g range = 0,5 or size = (1,261).

Learning rate and a fixed number of epochs and lists for the losses are considered.

- Depending on the number of epochs we choose, we loop and check if it is converged or not.
  - o In each epoch, when we are in the master process, the data is splitted by the number of workers we have .
  - Each data part is sent via scatter.
  - But the weights is sent via broadcast as it is the same for all the workers.
  - o In each worker the part of the data received is shuffled.
    - For each instance/row of the data received in each worker.
      - SGD is calculated.
      - Weights are updated.
      - After all the instance weights are updated.
      - Master node gathers all the weights, predicted y and true y.
- Master node finds the mean of the weights, calculates the RMSE and append it to our list for train and it uses new mean weights to predict new values for test data and computes the RMSE of the true test target and predicted ones to append it to out test loss list.
- If the test oss of each instance is less than a specific value then it is converged, it means it can't improve more.
  - If it is then broadcast the converge value so the program stops.
- And finally we save the losses and show the final time.

### Virus dataset.

## The STRATEGY is the same as above.

For this dataset, we define load\_virus() functions which goes into the dataset directory and list all of them into a list called files.

We go through each file and append them in another list, later we concat them to make a bigger dataframe.

The first value of each input is our target y, all the others, there is a key and a value pointing to a feature and its values exist here, we make a sparse matrix and go through each file and split them the first value is the column name and the second one is the value of that column. We make ones as bias and add them to x and we have our final datasets.

```
20 def load virus():
       #freeing up the memory
21
22
       gc.collect()
23
       path = 'dataset/dataset/'
       files = os.listdir(path)
24
25
       li = []
26
       for filename in files:
27
            df = pd.read csv(path+filename, header=None)
28
            li.append(df)
29
30
       dat file = pd.concat(li, axis=0, ignore index=True)
31
       y = np.zeros((107856,1), float)
32
33
       x = np.zeros((107856,479))
       for i in dat_file.index:
            doc = dat_file.iloc[i][0].strip().split()
35
36
            y[i] = float(doc[0])
37
            for element in doc[1:]:
38
39
                k, v = element.split(":")
40
                x[i,int(k)] = int(v)
41
       bias = np.ones((107856,1))
       x = np.append(bias, x, axis=1)
42
43
44
       return x,y
```

The rest of the functions are as below.

- train\_test\_split() makes data into two splits by chosen fraction.
- row normalizer(), normalize each row.
- predic(), predict new y based on x and weights received.
- sgd(), calculates the main part of our program, stochastic gradient descent.
- calculate RMSE(), calculate RMSE between prediction and true values.

```
46 | #splitting dataset
47 def train_test_split(data ,frac=0.7):
48
        data = pd.DataFrame(data)
49
        #train test split
50
        train = data.sample(frac=frac, random state=10)
51
        test = data.drop(train.index)
52
53
        return train.to numpy(), test.to numpy()
54
55 #normalzining based on rows
56 def row_normalizer(x: np.ndarray):
        return x/np.linalg.norm(x, ord=2, axis=1, keepdims=True)
57
58
59 #prediciton
60 def predict(x, betas):
61
        return np.matmul(x, betas.T)
62
63 #SGD function
64 def sgd(x, weight, y, y_hat, lr):
65
        #Computing Derivatives using Chain Rule
66
67
        dv_{loss} = x * (-2*(y - y_{hat}))
68
        #Computing new weights
69
        weight = weight - lr * dv loss
70
        return weight
71
72 #calculate RMSE
73 def calculate RMSE(true y, pred y):
74
        return np.sqrt(mt.mean squared error(true y, pred y))
75
```

## Starting from the master node.

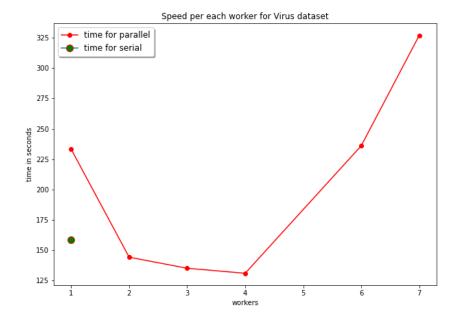
```
79 if rank == 0:
80
        print('Loading data...', flush=True)
81
        x,y = load virus()
        print(f"X shape: {x.shape}")
82
83
        x = normalize rows(x)
        print('Splitting into train and test sets...', flush=True)
84
85
        x_train, x_test = train_test_split(x ,frac=0.7)
        y_train, y_test = train_test_split(y ,frac=0.7)
 86
87
        data = np.concatenate((y train.reshape(-1,1), x train), axis=1)
88
89
90
91 #Initializing weights
92
93 w = np.random.randint(0,2,size=((1,480)))
94
95 #learning rate
96 lr = 0.00001
97
98 #Setting the number of epochs
99 N = 100
100 train_loss = []
101 test \overline{loss} = []
102
103 converged = False #For checking convergence
104 epoch = 0
105
106 if rank == 0:
107
        #Starting time
108
        t0 = MPI.Wtime()
```

```
110 #Training for N Epochs
111 while (not converged) and (epoch < N):</pre>
         if rank == 0:
    #print onlt every 10 steps
114
115
              if epoch%10==0:
                   print("Epoch : ", epoch, flush=True)
116
117
118
              #splitting
119
              data_part = np.array_split(data,size)
120
          else:
121
              data_part = None
          #Sending data using scatter and weights via bcast
123
          recvd dt = comm.scatter(data part,root=0)
124
          weights = comm.bcast(w,root=0)
          row_, col_ = np.shape(recvd_dt)
125
126
          y_hat = np.zeros((row_,1))
127
128
          #Shuffling the received dataset at each worker
129
          np.take(recvd_dt,np.random.permutation(recvd_dt.shape[0]),axis=0,out=recvd_dt)
130
131
          for i in range(0, row_):
              X_part = recvd_dt[i,1:]
133
              y part = recvd dt[i,0]
134
              #Prediction of instance i, forward pass
135
              y hat[i,0] = predict(X part, weights.T)
136
137
              #Computing new weights
138
              weights = sgd(X part, weights, y part, y hat[i,0], lr)
139
140
          #Gathering weights at Worker 0
141
          w r = comm.gather(weights, root=0)
142
          #Gathering predicted output (yy) and original data (true dt) at Worker 0
143
          y hat recvd = comm.gather(y hat, root=0)
          true dt = comm.gather(recvd dt[:,0], root=0)
145
146
        if rank == 0 and converged == False:
147
             #Worker 0 computing mean of the weights
148
             w = np.mean(w_r, axis=0)
149
150
             pred_y = np.vstack(y_hat_recvd)
151
             true_y = np.hstack(true_dt)
            #Worker 0 computing calculate_RMSE calculate_RMSE_train = calculate_RMSE(true_y, pred_y)
154
155
             train loss.append(calculate RMSE train)
156
             #test prediction
            pred_y_test = predict(x_test, w.T)
#Computing RME Testing
calculate_RMSE_test = calculate_RMSE(y_test, pred_y_test)
158
159
160
161
             test loss.append(calculate RMSE test)
162
             #Checking for convergence in RMSE test
if epoch > 0 and abs((test_loss[epoch-1] - test_loss[epoch])) < 10**-6:</pre>
163
164
165
                 print(test_loss[epoch-1] - test_loss[epoch], flush=True)
166
                 converged = True
167
         #the converged flag is sent back for stopping epochs
168
        converged = comm.bcast(converged,root=0)
169
        epoch += 1
170
171 if rank == 0:
        #Ending time
        t1 = MPI.Wtime()
173
        df = pd.DataFrame(list(zip(train_loss, test_loss)),columns =['train_loss', 'test_loss'])
175
        df.to_csv(f"loss_virus{size}.csv", index=False)
#t1 = MPI.Wtime()
176
        time_final = t1 - t0
print("With P = {} Workers, the process took {} seconds" .format(size, time_final), flush=True)
```

Here is the screenshot of running with 4 workers.

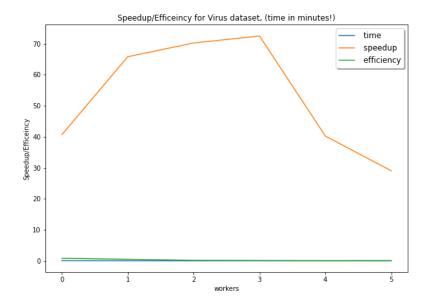
#### 1 !mpirun -n 4 python virus.py Loading data... X shape: (107856, 480) Splitting into train and test sets... Epoch: 0 Epoch: 10 Epoch: 20 30 Epoch: Epoch: 40 Epoch: 50 Epoch: 60 Epoch: 70 Epoch: 80 Epoch: 90 With P = 4 Workers, the process took 130.96906805038452 seconds

Let's look at the time elapsed using different workers.



Looking at the graph we see that the sequential code is faster than parallel code using only one worker. Using 2 workers is definitely faster than one and so until 4th workers. Using 6th and more is not fast anymore as my machine has only 4 cores.

Let's look at the speedup/efficiency graph.

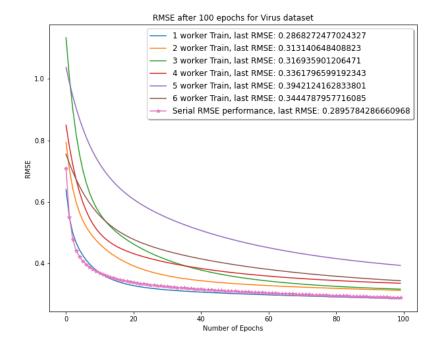


Looking at the above graph we see that we have a huge speed up until using 3 workers and then the speed goes down. The efficiency overall goes down. But more details are observed in this table.

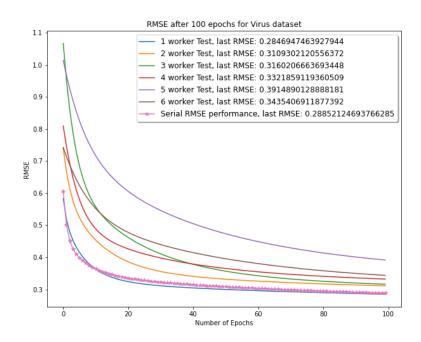
1	time_virus				
	workers	time	speedup	efficiency	
0	1	0.064786	40.718604	0.848725	
1	2	0.040067	65.840266	0.504409	
2	3	0.037536	70.278991	0.186993	
3	4	0.036380	72.511816	0.113861	
4	6	0.065559	40.238802	0.027713	
5	7	0.090825	29.044867	0.019868	

# RMSE Performance.

Looking at the graph the RMSE is going down and we have a good accuracy. But still sequential performance and using one worker is giving us better accuracy than more workers.



## Test RMSE Observation:



Even in the test set we see better performance in serial code and using one core, also 3 workers give us a good RMSE score.

• Using one worker it didn't converge but we have a good RMSE score.

- 2 workers did not converge, RMSE is worse than using one worker.
- 3 workers did not converge, RMSE is worse than using 1/2 workers.
- 4 workers did not converge, RMSE is worse than before.
- 6 workers did not converge, RMSE is worse than before.
- 7 workers did not converge, RMSE is worse than before.

## KDD dataset.

For this data set to save time I first concat the three datasets for learning and validation and validation target in one data frame.

Replaced " " with 0 and applied one hot encoding for features which are categorical plus the columns that have less than 5 unique values, they are converted to categorical features and one hot encoding applied.

Next I checked the Pearson correlation and those which had positive correlation were chosen. Finally we have a data set with shape (191779,261) bias added.

\_\_\_\_

#### First we define some functions:

- load\_data\_kdd() to load csv file we already cleaned and preprocessed, it adds bias to it and create x and y matrix.
- train\_test\_split() makes data into two splits by chosen fraction.
- row\_normalizer(), normalize each row.
- predic(), predict new y based on x and weights received.
- sgd(), calculates the main part of our program, stochastic gradient descent.
- calculate\_RMSE(), calculate RMSE between prediction and true values.

```
20 #reading dataset
21 def load_data_kdd(data = "kdd_final_filtered.csv"):
        #freeing up the memory
23
        gc.collect()
24
        #reading the saved dataset
25
        kdd = pd.read csv(data, low memory=False)
26
        #target
27
       y = kdd["TARGET D"].to numpy()
28
        x = kdd.drop(columns=["TARGET D"], axis=1)
29
30
        bias = pd.DataFrame(np.ones(x.shape[0]).reshape(-1,1), columns = ["bias"])
31
        x = pd.concat([bias, x], axis=1).to_numpy()
32
33
        return x, y
34
35 #splitting dataset
def train_test_split(data ,frac=0.7):
    data = pd.DataFrame(data)
38
        #train_test split
39
        train = data.sample(frac=frac, random_state=10)
40
        test = data.drop(train.index)
41
42
        return train.to_numpy(), test.to_numpy()
43
44 #normalzining based on rows
45 def row normalizer(x: np.ndarray):
        return x/np.linalg.norm(x, ord=2, axis=1, keepdims=True)
46
47
48 #prediciton
49 def predict(x, betas):
        return np.matmul(x, betas.T)
50
51
52 #SGD function
53 def sgd(x, weight, y, y_hat, lr):
55
        #Computing Derivatives using Chain Rule
        dv_loss = x * (-2*(y - y_hat))
#Computing new weights
56
57
58
        weight = weight - lr * dv_loss
59
60
        return weight
61 #calculate RMSE
62 def calculate RMSE(true y, pred y):
63
        return np.sqrt(mt.mean squared error(true y, pred y))
64
```

The rest of the code is the same as explained in the STRATEGY.

```
69 if rank == 0:
70
         print('Loading data...', flush=True)
71
         x,y = load data kdd()
72
         print(f"X shape: {x.shape}")
73
         #print("Normalizing")
74
75
         x = row normalizer(x)
76
77
         print('Splitting into train and test sets...', flush=True)
78
         x_train, x_test = train_test_split(x ,frac=0.7)
79
         y_train, y_test = train_test_split(y ,frac=0.7)
80
81
         data = np.concatenate((y_train.reshape(-1,1), x_train), axis=1)
82
83
84 #fixing seed
85 np.random.seed(2021)
86 #random weights
87 w = np.random.randint(0,5,size=((1,261)))
88
89 #learning rate
90 lr = 0.00001
91
92 #Setting the number of epochs
93 N = 100
94 train_loss = []
95 test_loss = []
96
97 converged = False #For checking convergence
98 epoch = 0
99
100 if rank == 0:
101
         #Starting time
         t0 = MPI.Wtime()
102
105 while (not converged) and (epoch < N):
106
        if rank == 0:
107
108
            #print onlt every 10 steps
109
            if epoch%10==0:
110
                print("Epoch : ", epoch, flush=True)
111
            #splitting
112
113
            data part = np.array split(data, size)
114
        else:
115
            data_part = None
116
        #Sending data using scatter and weights via bcast
        recvd_dt = comm.scatter(data_part,root=0)
117
        weights = comm.bcast(w,root=0)
row_, col_ = np.shape(recvd_dt)
118
119
        y_hat = np.zeros((row_,1))
120
121
122
        #Shuffling the received dataset at each worker
123
        np.take(recvd_dt,np.random.permutation(recvd_dt.shape[0]),axis=0,out=recvd_dt)
124
        #Epoch
125
        for i in range(0, row_):
126
            X part = recvd dt[i,1:]
            y_part = recvd_dt[i,0]
127
            #Prediction of instance i, forward pass
128
129
            y_hat[i,0] = predict(X_part, weights.T)
130
131
            #Computing new weights
            weights = sgd(X_part, weights, y_part, y_hat[i,0], lr)
132
133
134
        #Gathering weights at Worker 0
135
        w_r = comm.gather(weights, root=0)
        #Gathering predicted output (yy) and original data (true_dt) at Worker 0 y_hat_recvd = comm.gather(y_hat, root=0)
136
137
138
        true_dt = comm.gather(recvd_dt[:,0], root=0)
```

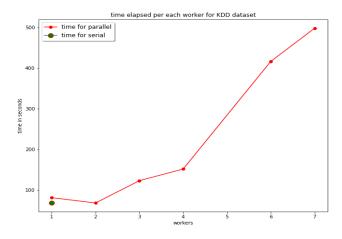
```
40
        if rank == 0 and converged == False:
             #Worker 0 computing mean of the weights
42
             w = np.mean(w_r, axis=0)
43
             pred_y = np.vstack(y_hat_recvd)
44
45
             true_y = np.hstack(true_dt)
46
             #Worker 0 computing calculate_RMSE
48
             calculate_RMSE_train = calculate_RMSE(true_y, pred_y)
49
             train_loss.append(calculate_RMSE_train)
50
51
             #test prediction
             pred_y_test = predict(x_test, w.T)
#Computing RME Testing
calculate_RMSE_test = calculate_RMSE(y_test, pred_y_test)
52
53
             test_loss.append(calculate_RMSE_test)
56
            #Checking for convergence in RMSE test
if epoch > 0 and abs((test_loss[epoch-1] - test_loss[epoch])) < 10**-6:</pre>
57
58
59
                 print(test_loss[epoch-1] - test_loss[epoch], flush=True)
60
                 converged = True
        #the converged flag is sent back for stopping epochs
        converged = comm.bcast(converged,root=0)
63
        epoch += 1
64
65 if rank == 0:
        #End time
66
67
        t1 = MPI.Wtime()
        df = pd.DataFrame(list(zip(train_loss, test_loss)),columns =['train_loss', 'test_loss'])
69
        df.to_csv(f"loss{size}.csv", index=False)
70
        #t1 = MPI.Wtime()
        time_final = t1 - t0
print("With P = {} Workers, the process took {} seconds" .format(size, time_final), flush=True)
```

## Running the code.

```
1 !mpiexec -n 2 python kdd_.py

Loading data...
X shape: (191779, 261)
Splitting into train and test sets...
Epoch : 0
Epoch : 10
Epoch : 20
B.017752826106062e-07
With P = 2 Workers, the process took 68.61729502677917 seconds
```

Looking at the time elapsed for each time with different workers.

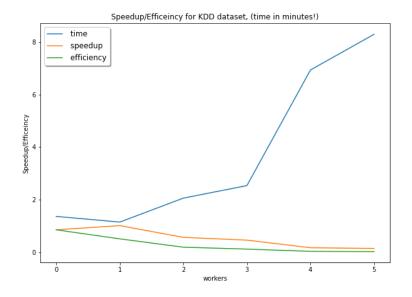


- Looking at this graph we see that using 2 workers gave us the quickest run time, as my machine has only 4 cores it is understandable why it takes longer time for more workers.
- Below is the total time, speedup and efficiency for this program.

	workers	time	speedup	efficiency
0	1	1.359333	0.848725	0.848725
1	2	1.143617	1.008817	0.504409
2	3	2.056583	0.560979	0.186993
3	4	2.533133	0.455444	0.113861
4	6	6.938417	0.166277	0.027713
5	7	8.295667	0.139073	0.019868

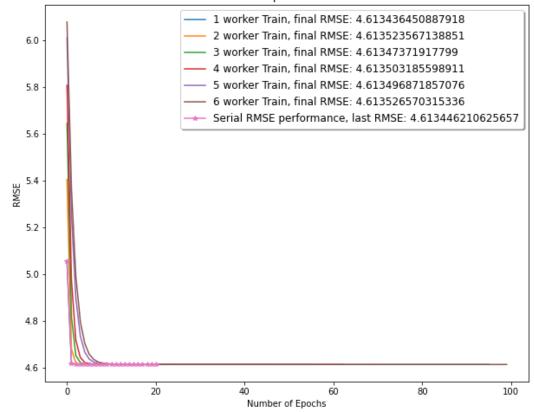
- Below is the graph for speedup and efficiency , time is divided by 60 seconds so we can see more details.
- The efficiency overall goes down with the number of processes increasing! I don't know why this happens, the time for the serial program is 69.222 here how I calculated the speedup and efficiency.
- The speedup increase and later it decreases.

```
time_["speedup"] = (69.222)/time_.time
time_["efficiency"] = time_["speedup"]/time_.workers
```



Looking at RMSE result for train and test sets.





- Using one worker it converged at the 23rd epoch.
- Using 2 workers it converged at the 30th epoch.
- Using 3 workers it converged at the 56th epoch
- Using 4 workers it converged at the 63rd epoch.
- Using 6 workers it converged at the 96th epoch
- Using 7 workers it did not converge.

Overall the RMSE is going lower but i could not get a lower RMSE even training for longer epochs and different learning rates.

Below is the testing RMSE values for KDD dataset. The behavior is as the same as train set.



