### **Graduate Admission**

This dataset is created for prediction of Graduate Admissions from an Indian perspective.

The dataset contains several parameters which are considered important during the application for Masters Programs. The parameters included are:

- · GRE Scores (out of 340)
- TOEFL Scores (out of 120)
- University Rating (out of 5)
- Statement of Purpose and Letter of Recommendation Strength (out of 5)
- Undergraduate GPA (out of 10)
- Research Experience (either 0 or 1)
- Chance of Admit (ranging from 0 to 1)

# **Import the Required Libraries**

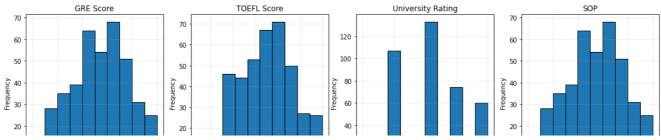
### Extract headers and data

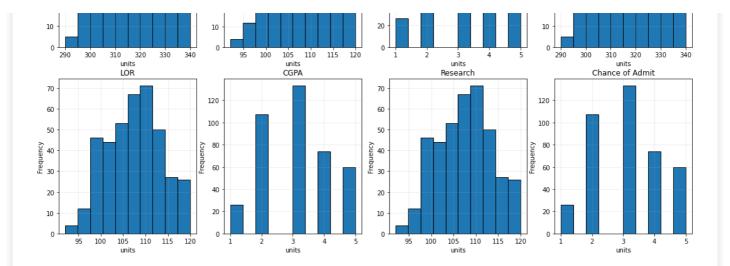
```
In [3]:
headers = df[0,1:]; # TO not take serial no
print(headers)
data = np.array(df[1:,1:], dtype=float); # This will take from the GRE Score
print(data)
['GRE Score' 'TOEFL Score' 'University Rating' 'SOP' 'LOR ' 'CGPA'
 'Research' 'Chance of Admit']
                                   0.92]
[[337. 118. 4. ... 9.65 1.
              4. ... 8.87 1.
[324. 107.
                                     0.761
      104.
[316.
               3.
                   ... 8.
                              1.
                                     0.721
[330. 116.
              4. ... 9.45 1.
                                    0.91]
                                   0.67]
[312. 103.
             3. ... 8.78 0.
 [333. 117.
              4. ... 9.66 1.
                                    0.95]]
```

# Visualise the distribution of independent and dependent variables

In [4]:

```
'''Create subplots in 1 row and 3 columns'''
fig, ax = plt.subplots(2,4)
fig.set figheight(10)
fig.set_figwidth(20)
fig.subplots adjust(left=.2, bottom=None, right=None, top=None, wspace=.2, hspace=.2)
plt1 = plt.subplot(2,4,1)
plt2 = plt.subplot(2,4,2)
plt3 = plt.subplot(2,4,3)
plt4 = plt.subplot(2,4,4)
plt5 = plt.subplot(2,4,5)
plt6 = plt.subplot(2,4,6)
plt7 = plt.subplot(2,4,7)
plt8 = plt.subplot(2,4,8)
plt1.hist(data[:,0], label='GRE Score', edgecolor='black')
plt1.set_title('GRE Score')
plt1.set_xlabel('units')
plt1.set ylabel('Frequency')
plt1.grid(axis='both', alpha=.25)
plt2.hist(data[:,1], label='TOEFL Score', edgecolor='black')
plt2.set_title('TOEFL Score')
plt2.set_xlabel('units')
plt2.set ylabel('Frequency')
plt2.grid(axis='both', alpha=.25)
plt3.hist(data[:,2], label='University Rating', edgecolor='black')
plt3.set_title('University Rating')
plt3.set xlabel('units')
plt3.set_ylabel('Frequency')
plt3.grid(axis='both', alpha=.25)
plt4.hist(data[:,0], label='SOP', edgecolor='black')
plt4.set title('SOP')
plt4.set xlabel('units')
plt4.set ylabel('Frequency')
plt4.grid(axis='both', alpha=.25)
plt5.hist(data[:,1], label='LOR', edgecolor='black')
plt5.set title('LOR')
plt5.set_xlabel('units')
plt5.set ylabel('Frequency')
plt5.grid(axis='both', alpha=.25)
plt6.hist(data[:,2], label='CGPA', edgecolor='black')
plt6.set title('CGPA')
plt6.set_xlabel('units')
plt6.set ylabel('Frequency')
plt6.grid(axis='both', alpha=.25)
plt7.hist(data[:,1], label='Research', edgecolor='black')
plt7.set_title('Research')
plt7.set xlabel('units')
plt7.set ylabel('Frequency')
plt7.grid(axis='both', alpha=.25)
plt8.hist(data[:,2], label='Chance of Admit', edgecolor='black')
plt8.set title('Chance of Admit')
plt8.set xlabel('units')
plt8.set_ylabel('Frequency')
plt8.grid(axis='both', alpha=.25)
```





#### In [5]:

```
data_norm = (data-np.mean(data, axis = 0))/np.std(data, axis = 0)
```

#### In [6]:

```
# Extract y from data

y_label = 'Chance of Admit';
y_index = np.where(headers == y_label)[0][0];
y = data_norm[:,y_index];

# Extract x from data

X = data_norm[:,0:y_index];
```

#### In [7]:

```
# Insert column of 1's for intercept column
X = np.insert(X, 0, 1, axis=1) #Here you added the intercept right
```

#### In [8]:

```
print(X.shape)
```

(400, 8)

#### In [9]:

```
print(X[0]) #Normalized values
```

[1. 1.76210664 1.74697064 0.79882862 1.09386422 1.16732114 1.76481828 0.90911166]

#### In [10]:

```
print(headers) #You dont need serial No. ok. There are total 9. +1 intercept
```

```
['GRE Score' 'TOEFL Score' 'University Rating' 'SOP' 'LOR ' 'CGPA' 'Research' 'Chance of Admit']
```

#### In [11]:

```
y.shape
```

#### Out[11]:

(400,)

```
In [12]:
m = X.shape[0]
n = X.shape[1]
In [13]:
def h(X, theta):
   return X.dot(theta)
In [14]:
def cost(theta, X, y):
   return (h(X, theta) - y).T.dot(h(X, theta) - y)/2
In [15]:
# Gradient of cost function
def gradient(X, y, theta):
   grad = X.T.dot(h(X, theta) - y)
   return grad
In [16]:
def gradient descent(X, y, theta initial, alpha, num iters):
   J per iter = np.zeros(num iters)
   gradient_per_iter = np.zeros((num_iters,len(theta_initial)))
   theta = theta initial
   for iter in np.arange(num iters):
      grad = gradient(X, y, theta)
       theta = theta - alpha * grad
       J per iter[iter] = cost(theta, X, y)
       gradient_per_iter[iter] = grad.T
   return (theta, J_per_iter, gradient_per_iter)
```

# Optimize the parameters using gradient descent

```
theta_initial = np.zeros((X.shape[1],1))
alpha = 0.0005
iterations = 1000
theta, costs, grad = gradient_descent(X,np.array([y]).T,theta_initial,alpha,iterations)
print('Theta values ', theta)
```

```
Theta values [[ 9.07968145e-16] [ 1.39783600e-01] [ 1.24258432e-01] [ 4.58476504e-02] [-2.33355774e-02] [ 1.40830779e-01] [ 4.97342141e-01] [ 8.57053356e-02]]
```

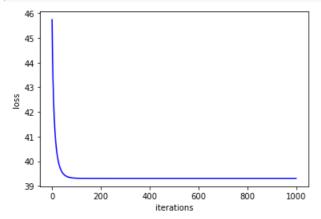
### Visualize the loss

```
In [18]:
```

In [17]:

```
x_loss = np.arange(0,iterations,1)
plt.plot(x_loss, costs, 'b-')
plt.ylabel('iterations')
```

```
htr · vtanet / tretartous /
plt.ylabel('loss')
plt.show()
```



### Goodness of fit

\$R^2\$ is a statistic that will give some information about the goodness of fit of a regression model. The \$R^2\$ coefficient of determination is 1 when the regression predictions perfectly fit the data. We have the theta values from the gradient method. We need to use then to predict target variable(Y predicted) then use it to calculate the R squared value(\$R^2\$).  $\label{eq:left(i/right)} $$ \left( y^{\left(i\right)}-\hat{y}\right) \ R^2 = 1 - \frac{i-1}^{m} \left( y^{\left(i\right)}-\hat{y}\right) \right)^2 {\sum_{i=1}^{m} \left( y^{\left(i\right)}-\hat{y}\right)^2 } (y^{i-1})^2$ \bar{y}^\left(i\right) \right)^2} \end{align}

```
In [19]:
```

```
y \text{ predicted} = h(X, \text{theta})
r_square = 1 - np.square(y - y_predicted.T).sum()/np.square(y - y.mean()).sum()
print(r square)
```

0.8034713719824395

An R2 of 0.80 indicates an extremely good fit to the data.

# Transform standardized data back to original scale

We can transform standardized predicted values, y\_predicted into the orginal data scale using \$\$y\_{\text{norm}} = \sigma\_y y + \mu\_y\$\$

#### In [20]:

```
# Compute mean and standard deviation of data
sigma = np.array(np.std(data,axis=0))
mu = np.array(np.mean(data,axis=0))
# De-normalize y
y predicted = np.round(h(X, theta) * sigma[-1] + mu[-1], 2)
# Print first five values of y_predicted
print(y_predicted[0:5,:])
[[0.95]
[0.81]
 [0.65]
 [0.74]
 [0.64]]
```

# **Normal Equation**

theta = T1.dot(T2)
return(theta,T1,T2)

Let's try to do normal equation and see the performance of the model.

```
In [21]:

def normal_equation(X,y,theta):
    X_transpose = np.transpose(X) #calculate tranpose
    X_transpose_dot_X = X_transpose.dot(X) #Calculate the dot product
    T1 = np.linalg.inv(X_transpose_dot_X) #Calulate inverse
    T2 = X_transpose.dot(y)
```

```
In [22]:
```

```
theta_initial = np.zeros((X.shape[1],1))
iterations = 1000
theta, costs, grad = normal_equation(X,np.array([y]).T,theta_initial)
print('Theta values ', theta)
Theta values [[ 8.70408612e-16]
```

```
[ 1.39783600e-01]
[ 1.24258432e-01]
[ 4.58476504e-02]
[-2.33355774e-02]
[ 1.40830779e-01]
[ 4.97342141e-01]
[ 8.57053356e-02]]
```

#### In [23]:

```
# Goodness of fit
y_predicted = h(X,theta)
r_square = 1 - np.square(y - y_predicted.T).sum()/np.square(y - y.mean()).sum()
print(r_square)
```

0.8034713719824395

An R2 of 0.80 indicates an extremely good fit to the data. In fact ,the accuracy is same as gradient descent.

```
In [ ]:
```