

Problem Statement

- The objective of this research is to optimize and predict the milling parameters using a combination of the Taguchi method and machine learning techniques.
- The Taguchi method provides an efficient experimental design approach, while machine learning algorithms offer the ability to learn from data and make accurate predictions.
- By integrating these two approaches, we aim to enhance the efficiency and effectiveness of milling parameter optimization

Importing the necessary Libraries

```
In [1]: ## Importing the required libraries
import pandas as pd
import numpy as np
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
from sklearn.metrics import mean_squared_error, r2_score
import matplotlib.pyplot as plt
import seaborn as sns
%matplotlib inline
from sklearn.metrics import classification_report, accuracy_score
import warnings
warnings.filterwarnings("ignore")
from sklearn.ensemble import RandomForestRegressor
```

Importing the DataSets and Viewing it

```
In [2]: ## DataSet 1 with input values
df = pd.read_csv(r"C:\Users\lenovo\Desktop\Ra Taguchi Parameters.csv")
df.head()
```

Out[2]:

	Sr no	Size	Speed(Rpm)	Depth Of Cut (mm)	Feed Rate(mm/Rev)
0	NaN	NaN	NaN	NaN	NaN
1	1.0 10 mm Thk X 100 mm x 100 mm Lg.		1500.0	0.5	0.12
2	2.0 10 mm Thk X 100 mm x 100 mm Lg.		1500.0	0.5	0.18
3	3.0 10 mm Thk X 100 mm x 100 mm Lg.		1500.0	0.5	0.24
4	4.0 10 mm Thk X 100 mm x 100 mm Lg.		1500.0	1.0	0.12

```
In [3]: ## DataSet with output values
df_1 = pd.read_csv(r"C:\Users\lenovo\Desktop\Ra Taguchi Readings.csv")
df_1.head()
```

Out[3]:

	Sr no	A1(Trnsverse)	A2(Trnsverse)	B1 (Trnsverse)	B2 (Trnsverse)	C1 (Trnsverse)	C2 (Trnsverse)	A (longitudnal)	B (longitudnal)
0	1	0.210	0.105	0.214	0.144	0.229	0.182	0.213	0.510
1	2	0.184	0.095	0.146	0.101	0.208	0.193	0.314	0.435
2	3	0.580	0.574	0.515	0.385	0.327	0.352	0.570	0.660
3	4	0.267	0.208	0.227	0.249	0.233	0.117	0.197	0.370
4	5	0.266	0.123	0.263	0.144	0.215	0.128	0.506	0.439

Merging the DataSets

```
In [4]: ## Merging both the DataSet
df_2 = pd.merge(df, df_1)
df_2.head(2)
```

Out[4]:

	Sr no	Size	Speed(Rpm)	Depth Of Cut (mm)	Feed Rate(mm/Rev)	A1(Trnsverse)	A2(Trnsverse)	B1 (Trnsverse)	B2 (Trnsverse)	C (Trnsvers
0	1.0	10 mm Thk X 100 mm x 100 mm Lg.	1500.0	0.5	0.12	0.210	0.105	0.214	0.144	0.229
1	2.0	10 mm Thk X 100 mm x 100 mm Lg.	1500.0	0.5	0.18	0.184	0.095	0.146	0.101	0.208

Data Preprocssing

```
In [5]: ## Checking the shape of the data i.e. ( Rows and Columns )
df_2.shape
```

Out[5]: (27, 15)

In [6]: *## Checking the Information of DataSet i.e. Data Type of each column*
df_2.info()

```
<class 'pandas.core.frame.DataFrame'>
Int64Index: 27 entries, 0 to 26
Data columns (total 15 columns):
#   Column                                Non-Null Count  Dtype
---  -
0   Sr no                                27 non-null     float64
1   Size                                27 non-null     object
2   Speed(Rpm)                          27 non-null     float64
3   Depth Of Cut (mm)                   27 non-null     float64
4   Feed Rate(mm/Rev)                   27 non-null     float64
5   A1( Trnsverse)                      27 non-null     float64
6   A2( Trnsverse)                      27 non-null     float64
7   B1 (Trnsverse)                      27 non-null     float64
8   B2 (Trnsverse)                      27 non-null     float64
9   C1 (Trnsverse)                      27 non-null     float64
10  C2 (Trnsverse)                      27 non-null     float64
11  A (longitudnal)                     27 non-null     float64
12  B (longitudnal)                     27 non-null     float64
13  Rat                                  27 non-null     float64
14  Ral                                  27 non-null     float64
dtypes: float64(14), object(1)
memory usage: 3.4+ KB
```

In [7]: *## Dropping the unwanted Column as it is of no use to us i.e. (Sr. No.)*
df_2.drop(columns='Sr no', inplace=True)
df_2.head(2)

Out[7]:

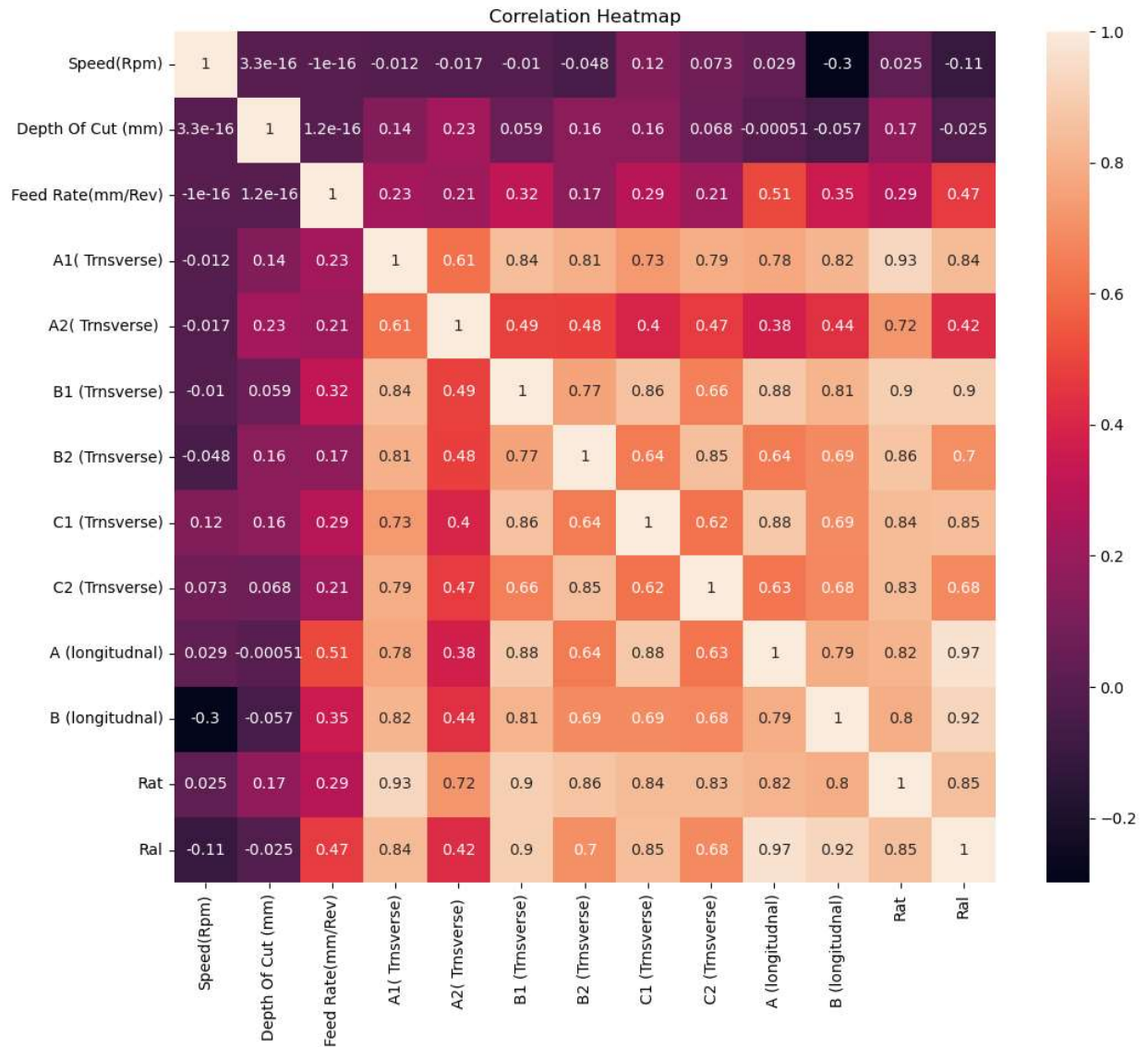
	Size	Speed(Rpm)	Depth Of Cut (mm)	Feed Rate(mm/Rev)	A1(Trnsverse)	A2(Trnsverse)	B1 (Trnsverse)	B2 (Trnsverse)	C1 (Trnsverse)
0	10 mm Thk X 100 mm x 100 mm Lg.	1500.0	0.5	0.12	0.210	0.105	0.214	0.144	0.229
1	10 mm Thk X 100 mm x 100 mm Lg.	1500.0	0.5	0.18	0.184	0.095	0.146	0.101	0.208

```
In [8]: ## Checking for Null Values in merged data  
df_2.isna().sum()
```

```
Out[8]: Size                0  
Speed(Rpm)                0  
Depth Of Cut (mm)         0  
Feed Rate(mm/Rev)         0  
A1( Trnsverse)            0  
A2( Trnsverse)            0  
B1 (Trnsverse)            0  
B2 (Trnsverse)            0  
C1 (Trnsverse)            0  
C2 (Trnsverse)            0  
A (longitudnal)           0  
B (longitudnal)           0  
Rat                        0  
Ral                        0  
dtype: int64
```

In [9]: *## Plotting the HeatMap for checking the Correlation between the Values*

```
correlation_matrix = df_2.corr()
fig, ax = plt.subplots(figsize=(12, 10))
sns.heatmap(correlation_matrix, annot=True, ax=ax)
ax.set_title("Correlation Heatmap")
plt.show()
```



In this HeatMap darker the colour less is the Correlation and Lighter the Colour Heigher is the Correlation with each other.

Conclusion from the HeatMap

Checking for the description of the Data

```
In [10]: ## Checking the descriptive Statistics of the DataSet
df_2.describe().T
```

Out[10]:

	count	mean	std	min	25%	50%	75%	m
Speed(Rpm)	27.0	1766.666667	209.394732	1500.000000	1500.000000	1800.000000	2000.0000	2000.0000
Depth Of Cut (mm)	27.0	0.916667	0.317744	0.500000	0.500000	1.000000	1.2500	1.2500
Feed Rate(mm/Rev)	27.0	0.180000	0.049923	0.120000	0.120000	0.180000	0.2400	0.2400
A1(Trnsverse)	27.0	0.249963	0.135648	0.125000	0.153500	0.215000	0.2665	0.5890
A2(Trnsverse)	27.0	0.225778	0.205467	0.051000	0.100000	0.138000	0.2875	0.9400
B1 (Trnsverse)	27.0	0.259296	0.128337	0.095000	0.185000	0.224000	0.2695	0.5600
B2 (Trnsverse)	27.0	0.191852	0.122062	0.075000	0.097500	0.144000	0.2375	0.5350
C1 (Trnsverse)	27.0	0.268481	0.179552	0.110000	0.164000	0.215000	0.2565	0.8440
C2 (Trnsverse)	27.0	0.191926	0.116344	0.070000	0.115000	0.144000	0.2340	0.4650
A (longitudnal)	27.0	0.320296	0.199535	0.096000	0.198500	0.264000	0.4210	0.8280
B (longitudnal)	27.0	0.375444	0.137178	0.124000	0.291000	0.350000	0.4675	0.6600
Rat	27.0	0.231216	0.123853	0.101167	0.153500	0.180667	0.2605	0.5520
Ral	27.0	0.347870	0.159704	0.111000	0.252250	0.303000	0.4235	0.6720

```
In [11]: ## Checking for Datatypes of Columns
df_2.dtypes
```

```
Out[11]: Size          object
Speed(Rpm)          float64
Depth Of Cut (mm)    float64
Feed Rate(mm/Rev)    float64
A1( Trnsverse)       float64
A2( Trnsverse)       float64
B1 (Trnsverse)       float64
B2 (Trnsverse)       float64
C1 (Trnsverse)       float64
C2 (Trnsverse)       float64
A (longitudnal)      float64
B (longitudnal)      float64
Rat                  float64
Ral                  float64
dtype: object
```

```
In [12]: ## Lets create the dummy variables for all the categorical columns
cat_col = ['Size']
df_3= pd.get_dummies(df_2,columns=cat_col,drop_first=True)
df_3.head()
```

Out[12]:

	Speed(Rpm)	Depth Of Cut (mm)	Feed Rate(mm/Rev)	A1(Trnsverse)	A2(Trnsverse)	B1 (Trnsverse)	B2 (Trnsverse)	C1 (Trnsverse)	(Trnsve
0	1500.0	0.5	0.12	0.210	0.105	0.214	0.144	0.229	0
1	1500.0	0.5	0.18	0.184	0.095	0.146	0.101	0.208	0
2	1500.0	0.5	0.24	0.580	0.574	0.515	0.385	0.327	0
3	1500.0	1.0	0.12	0.267	0.208	0.227	0.249	0.233	0
4	1500.0	1.0	0.18	0.266	0.123	0.263	0.144	0.215	0

```
In [13]: ## Checking the Shape of Dtsaet after creating dummy variables for categorical Columns
df_3.shape
```

Out[13]: (27, 13)

Modelling Part

```
In [14]: ## Assigning the X & Y values
y = df_3[['Ra1']]
X = df_3.drop(columns=['Ra1'])
```

- The StandardScaler is a popular data preprocessing technique used in machine learning and statistics. It is used to standardize features by subtracting the mean and scaling to unit variance. In other words, it transforms the data so that it has a mean of zero and a standard deviation of one.
- The formula for standardization is:

$$z = (x - u) / s$$

Where:

- z is the standardized value
- x is the original value
- u is the mean of the feature
- s is the standard deviation of the feature
- The StandardScaler calculates the mean and standard deviation of each feature in the dataset and then applies the above formula to transform each value. By doing so, it ensures that each feature has the same scale and brings all features to a similar range.
- Standardization is useful when working with algorithms that assume data to be normally distributed and have equal variances. It helps in cases where features have different scales, preventing certain features from dominating others during the learning process. It also helps with gradient-based optimization algorithms that converge faster when features are on a similar scale.

```
In [15]: ## Using scaler for scaling the Data
```

```
sc = StandardScaler()  
X = sc.fit_transform(X)
```

```
In [16]: ## Splitting the data into train & test format
```

```
X_train,X_test,y_train,y_test = train_test_split(X,y,test_size=0.20,random_state=13)
```

```
In [17]: ## Finding the Length of test and train data
```

```
len(X_train),len(X_test),len(y_train),len(y_test)
```

```
Out[17]: (21, 6, 21, 6)
```

Model Selection

Here we are using Regression Model as the Problem we are going to solve is Regression type as we have to predict continuous or numeric values. Here we are going to use Random Forest Regressor because

- **Accuracy:** Random Forest Regressor typically provides high prediction accuracy. It combines multiple decision trees and aggregates their predictions, resulting in more accurate and robust predictions compared to individual trees.
- **Non-linearity:** Random Forest Regressor can capture non-linear relationships between input features and the target variable. It is capable of handling complex interactions and non-linear patterns in the data, making it suitable for a wide range of regression problems.
- **Resistance to overfitting:** Random Forest Regressor reduces the risk of overfitting, which occurs when a model learns too much from the training data and performs poorly on new, unseen data. The randomness introduced during the construction of individual trees helps in decorrelating their predictions, leading to a reduction in overfitting.
- **Robustness to outliers:** Random Forest Regressor is less sensitive to outliers compared to some other regression algorithms. Outliers have a lesser impact on the overall model's performance because they are typically averaged out by the combination of multiple trees.
- **Feature importance:** Random Forest Regressor provides a measure of feature importance. By analyzing the importance scores assigned to different features, you can gain insights into which variables are most influential in the regression task. This information can be valuable for feature selection and understanding the underlying relationships in the data.
- **Handling missing values:** Random Forest Regressor can handle missing values in the dataset. It uses the available features to make predictions without requiring imputation or removal of instances with missing values. This is advantageous when dealing with real-world datasets that often contain missing data.
- **Ease of use:** Random Forest Regressor is relatively easy to use and implement. It does not require extensive data preprocessing or feature scaling and can handle both numerical and categorical features. Additionally, it has fewer hyperparameters to tune compared to some other algorithms.


```

In [18]: ## Selection of Model and Training the Model
rf_regressor = RandomForestRegressor(n_estimators=100, random_state=13)

# Train the model
rf_regressor.fit(X_train, y_train)

# Make predictions on the training set
y_train_pred = rf_regressor.predict(X_train)

# Calculate RMSE and R-squared for training set
mse_train = mean_squared_error(y_train, y_train_pred)
rmse_train = np.sqrt(mse_train)
r2_train = r2_score(y_train, y_train_pred)
print("Training Set:")
print(' ')
print("Mean Squared Error:", mse_train)
print(' ')
print("Root Mean Squared Error:", rmse_train)
print(' ')
print("R-squared:", r2_train)
print("=====")

# Make predictions on the test set
y_test_pred = rf_regressor.predict(X_test)

# Calculate RMSE and R-squared for test set
mse_test = mean_squared_error(y_test, y_test_pred)
rmse_test = np.sqrt(mse_test)
r2_test = r2_score(y_test, y_test_pred)
print("Testing Set:")
print(' ')
print("Mean Squared Error:", mse_test)
print(' ')
print("Root Mean Squared Error:", rmse_test)
print(' ')
print("R-squared:", r2_test)

```

Training Set:

Mean Squared Error: 0.0003762474404761947

Root Mean Squared Error: 0.019397098764407907

R-squared: 0.9862042235498657

=====

Testing Set:

Mean Squared Error: 0.0006994509458333334

Root Mean Squared Error: 0.026447134926742697

R-squared: 0.953410314671729

Conclusion

- On The Basic of above prediction we can see that the R-squared value of Training Set is 98.62% while that of Testing Set is 95.34%. So on the basic of that we can say that Our Model is performing well on the Data We collected and the Accuracy of the Model is 98%.

- So we can say that their is no overfitting condition as the R-squared value of the Training set is Heigher that the Testing Set.

Prediction for the values

```
In [19]: ## Original Y_Test Values
y_test
```

Out[19]:

	RaI
8	0.2780
19	0.2470
1	0.3745
11	0.2575
15	0.6050
7	0.3650

```
In [20]: ## Making prediction on target Column
y_test['Prediction'] = rf_regressor.predict(X_test)
```

```
In [21]: ## Predicted Values of Target Column
y_test
```

Out[21]:

	RaI	Prediction
8	0.2780	0.296445
19	0.2470	0.259530
1	0.3745	0.331220
11	0.2575	0.296785
15	0.6050	0.593385
7	0.3650	0.377170

- - - - - X X X X X X X - - - - -