We'll generate a synthetic dataset that simulates the complex process

```
pip install seaborn
Collecting seaborn
  Using cached seaborn-0.13.2-py3-none-any.whl.metadata (5.4 kB)
Requirement already satisfied: numpy!=1.24.0,>=1.20 in c:\users\yoga\
appdata\local\programs\python\python39\lib\site-packages (from
seaborn) (1.26.4)
Reguirement already satisfied: pandas>=1.2 in c:\users\yoga\appdata\
local\programs\python\python39\lib\site-packages (from seaborn)
Requirement already satisfied: matplotlib!=3.6.1,>=3.4 in c:\users\
yoga\appdata\local\programs\python\python39\lib\site-packages (from
seaborn) (3.8.4)
Requirement already satisfied: contourpy>=1.0.1 in c:\users\yoga\
appdata\local\programs\python\python39\lib\site-packages (from
matplotlib!=3.6.1,>=3.4->seaborn) (1.2.1)
Requirement already satisfied: cycler>=0.10 in c:\users\yoga\appdata\
local\programs\python\python39\lib\site-packages (from matplotlib!
=3.6.1,>=3.4->seaborn) (0.12.1)
Requirement already satisfied: fonttools>=4.22.0 in c:\users\voga\
appdata\local\programs\python\python39\lib\site-packages (from
matplotlib!=3.6.1,>=3.4->seaborn) (4.51.0)
Requirement already satisfied: kiwisolver>=1.3.1 in c:\users\yoga\
appdata\local\programs\python\python39\lib\site-packages (from
matplotlib!=3.6.1,>=3.4->seaborn) (1.4.5)
Requirement already satisfied: packaging>=20.0 in c:\users\yoga\
appdata\roaming\python\python39\site-packages (from matplotlib!
=3.6.1.>=3.4->seaborn) (23.2)
Requirement already satisfied: pillow>=8 in c:\users\yoga\appdata\
local\programs\python\python39\lib\site-packages (from matplotlib!
=3.6.1,>=3.4->seaborn) (10.3.0)
Requirement already satisfied: pyparsing>=2.3.1 in c:\users\yoga\
appdata\local\programs\python\python39\lib\site-packages (from
matplotlib!=3.6.1,>=3.4->seaborn) (3.1.2)
Requirement already satisfied: python-dateutil>=2.7 in c:\users\yoga\
appdata\roaming\python\python39\site-packages (from matplotlib!
=3.6.1,>=3.4->seaborn) (2.8.2)
Requirement already satisfied: importlib-resources>=3.2.0 in c:\users\
yoga\appdata\local\programs\python\python39\lib\site-packages (from
matplotlib!=3.6.1,>=3.4->seaborn) (6.4.0)
Requirement already satisfied: pytz>=2020.1 in c:\users\yoga\appdata\
local\programs\python\python39\lib\site-packages (from pandas>=1.2-
>seaborn) (2024.2)
Requirement already satisfied: tzdata>=2022.7 in c:\users\yoga\
appdata\local\programs\python\python39\lib\site-packages (from
```

```
pandas >= 1.2 - seaborn) (2024.2)
Requirement already satisfied: zipp>=3.1.0 in c:\users\yoga\appdata\
roaming\python\python39\site-packages (from importlib-
resources>=3.2.0->matplotlib!=3.6.1,>=3.4->seaborn) (3.17.0)
Requirement already satisfied: six>=1.5 in c:\users\yoga\appdata\
roaming\python\python39\site-packages (from python-dateutil>=2.7-
>matplotlib!=3.6.1,>=3.4->seaborn) (1.16.0)
Downloading seaborn-0.13.2-py3-none-any.whl (294 kB)
Installing collected packages: seaborn
Successfully installed seaborn-0.13.2
Note: you may need to restart the kernel to use updated packages.
[notice] A new release of pip is available: 24.2 -> 24.3.1
[notice] To update, run: python.exe -m pip install --upgrade pip
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
```

Set Random Seed To ensure reproducibility.

```
np.random.seed(42)
```

Generate Input Variables Let's assume we have 10 input variables.

```
n_samples = 1000
n_features = 10

# Generate random input features between 0 and 10
X = np.random.uniform(0, 10, size=(n_samples, n_features))
```

Assign Coefficients Randomly assign coefficients β for each term.

```
beta_linear = np.random.uniform(-5, 5, size=n_features)
# Coefficients for interaction terms (n_features choose 2)
n_interactions = int(n_features * (n_features - 1) / 2)
beta_interaction = np.random.uniform(-1, 1, size=n_interactions)
# Linear combination of inputs
linear_terms = np.dot(X, beta_linear)
# Initialize interaction terms array
interaction_terms = np.zeros(n_samples)
# Index to keep track of interaction coefficients
idx = 0
```

```
# Loop over all unique pairs of features
for i in range(n features):
   for j in range(i + 1, n_features):
       # Add interaction term
       interaction terms += beta interaction[idx] * X[:, i] * X[:, j]
       idx += 1
epsilon = np.random.normal(0, 1, n samples)
# Total output
y = beta linear[0] + linear terms + interaction terms + epsilon
# Feature names
columns = [f'x{i+1}' for i in range(n_features)]
# Combine features and target into a DataFrame
data = pd.DataFrame(X, columns=columns)
data['y'] = y
print(data.head(5))
                  x2
                           x3
                                     x4
                                               x5
                                                        x6
        x1
x7 \
0 3.745401 9.507143 7.319939 5.986585 1.560186 1.559945
0.580836
1 0.205845 9.699099 8.324426 2.123391 1.818250
                                                   1.834045
3.042422
2 6.118529 1.394939 2.921446 3.663618 4.560700 7.851760
1.996738
3 6.075449 1.705241 0.650516 9.488855 9.656320
                                                   8.083973
3.046138
4 1.220382 4.951769 0.343885 9.093204 2.587800 6.625223
3.117111
                  x9
                          x10
        x8
0 8.661761 6.011150 7.080726 178.654324
1 5.247564 4.319450 2.912291
                                85.918166
2 5.142344 5.924146 0.464504
                                76.613757
3 0.976721 6.842330 4.401525
                                -94.338920
4 5.200680 5.467103 1.848545
                               -38.892191
```

1. Data Preprocessing Handling Missing Values Our synthetic data has no missing values, but in real scenarios, you'd handle them here.

Feature Scaling Neural networks converge faster with scaled features.

```
from sklearn.preprocessing import StandardScaler
# Initialize scaler
scaler = StandardScaler()
```

```
# Fit and transform features
X_scaled = scaler.fit_transform(data.drop('y', axis=1))
# Update DataFrame
data_scaled = pd.DataFrame(X_scaled, columns=columns)
data_scaled['y'] = data['y']
```

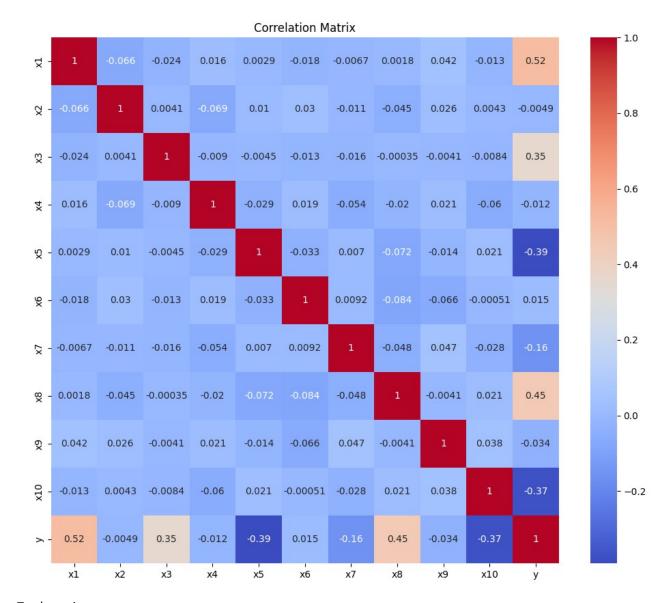
Check Scaled Data

```
print(data_scaled.head())
        x1
                 x2
                           x3
                                    x4
                                              x5
                                                       x6
x7 \
0 -0.363066 1.547478 0.821234 0.361551 -1.177298 -1.196034 -
1.532007
1 -1.603520 1.612962 1.167403 -1.011189 -1.087152 -1.101643 -
0.670775
2 0.468607 -1.219938 -0.694585 -0.463887 -0.129166 0.970644 -
1.036627
3 0.453509 -1.114080 -1.477198 1.606041 1.650825 1.050610 -
0.669475
4 -1.247970 -0.006553 -1.582870 1.465451 -0.818335 0.548268 -
0.644644
        x8 x9
                          x10
0 1.302662 0.337660 0.761895 178.654324
1 0.121575 -0.252714 -0.690874
                                85.918166
2 0.085176 0.307297 -1.543968
                                76.613757
3 -1.355854 0.627727 -0.171851 -94.338920
4 0.105356 0.147797 -1.061607 -38.892191
```

1. Feature Selection Correlation Analysis

```
# Compute correlation matrix
corr_matrix = data_scaled.corr()

# Plot heatmap
plt.figure(figsize=(12, 10))
sns.heatmap(corr_matrix, annot=True, cmap='coolwarm')
plt.title('Correlation Matrix')
plt.show()
```



Explanation:

Purpose: Identify highly correlated features. Action: We can decide to remove features with high multicollinearity. Select Features For simplicity, we'll proceed with all features, but in practice, you might select a subset.

1. Feature Engineering Creating Interaction Features Using DOE principles to include interaction terms.

```
from sklearn.preprocessing import PolynomialFeatures

# Create polynomial features (degree=2 includes interactions)
poly = PolynomialFeatures(degree=2, interaction_only=True,
include_bias=False)

# Fit and transform features
X_poly = poly.fit_transform(data_scaled.drop('y', axis=1))
```

```
# Get feature names
feature_names = poly.get_feature_names_out(columns)
```

Explanation PolynomialFeatures: degree=2: Includes all combinations of features up to degree 2. interaction_only=True: Excludes squared terms, focusing on interactions.

Update DataFrame with Interaction Features

```
# Create DataFrame with interaction features
data_poly = pd.DataFrame(X_poly, columns=feature_names)
data_poly['y'] = data_scaled['y']
```

1. Neural Network Model Implementation Libraries

Split Data into Training and Testing Sets

Define the Neural Network Architecture

```
# Input dimension after adding interaction terms
input_dim = X_train.shape[1]

# Initialize the model
model = Sequential()

# Input layer
model.add(Dense(128, input_dim=input_dim, activation='relu'))

# Hidden layers
model.add(Dense(256, activation='relu'))
model.add(Dropout(0.3)) # Dropout for regularization
model.add(Dense(128, activation='relu'))

# Output layer
model.add(Dense(1, activation='linear'))
```

```
WARNING:tensorflow:From c:\Users\Yoga\AppData\Local\Programs\Python\Python39\lib\site-packages\keras\src\backend.py:873: The name tf.get_default_graph is deprecated. Please use tf.compat.v1.get_default_graph instead.
```

model.summary()

Model: "sequential"

Param #
7168
33024
0
32896
129
:=========

Total params: 73217 (286.00 KB)
Trainable params: 73217 (286.00 KB)
Non-trainable params: 0 (0.00 Byte)

```
model.compile(
    optimizer='adam',
    loss='mean_squared_error',
    metrics=['mean_absolute_error']
)
```

WARNING:tensorflow:From c:\Users\Yoga\AppData\Local\Programs\Python\Python39\lib\site-packages\keras\src\optimizers__init__.py:309: The name tf.train.Optimizer is deprecated. Please use tf.compat.v1.train.Optimizer instead.

Explanation:

Optimizer: Adam optimizer adapts learning rates during training. Loss Function: Mean Squared Error for regression tasks. Metrics: Mean Absolute Error to evaluate model performance.

1. Model Training Set Up Early Stopping

```
from tensorflow.keras.callbacks import EarlyStopping
early_stop = EarlyStopping(
    monitor='val_loss',
```

```
patience=10,
  restore_best_weights=True
)
```

Train the Model

```
history = model.fit(
   X train, y train,
   validation_split=0.2,
   epochs=100,
   batch size=32,
   callbacks=[early stop],
   verbose=1
)
Epoch 1/100
WARNING:tensorflow:From c:\Users\Yoga\AppData\Local\Programs\Python\
Python39\lib\site-packages\keras\src\utils\tf utils.py:492: The name
tf.ragged.RaggedTensorValue is deprecated. Please use
tf.compat.v1.ragged.RaggedTensorValue instead.
WARNING:tensorflow:From c:\Users\Yoga\AppData\Local\Programs\Python\
Python39\lib\site-packages\keras\src\engine\base layer utils.py:384:
The name tf.executing_eagerly_outside_functions is deprecated. Please
use tf.compat.v1.executing eagerly outside functions instead.
- mean absolute error: 73.9305 - val_loss: 6188.2852 -
val mean absolute error: 59.7385
Epoch 2/100
- mean absolute error: 71.0626 - val loss: 5332.3345 -
val mean absolute error: 55.5473
Epoch 3/100
20/20 [============== ] - 0s 3ms/step - loss: 4886.9434
- mean absolute error: 53.6557 - val_loss: 2312.6646 -
val mean absolute error: 36.2888
Epoch 4/100
20/20 [============== ] - 0s 3ms/step - loss: 1039.7753
- mean absolute error: 24.2208 - val loss: 453.5901 -
val mean absolute error: 16.8014
Epoch 5/100
- mean absolute error: 13.1802 - val loss: 213.3192 -
val mean absolute error: 11.3393
Epoch 6/100
- mean absolute error: 10.2701 - val loss: 151.1905 -
val mean absolute error: 9.8167
```

```
Epoch 7/100
20/20 [============ ] - Os 3ms/step - loss: 167.8574
- mean absolute error: 10.1185 - val loss: 151.0729 -
val mean absolute error: 9.5091
Epoch 8/100
20/20 [============== ] - 0s 3ms/step - loss: 136.5610
- mean absolute error: 9.0322 - val loss: 117.6982 -
val mean absolute error: 8.3595
Epoch 9/100
20/20 [============= ] - 0s 3ms/step - loss: 133.2912
- mean absolute error: 8.8254 - val loss: 111.0519 -
val mean absolute error: 7.9624
Epoch 10/100
20/20 [============== ] - 0s 3ms/step - loss: 114.8808
- mean absolute error: 8.2795 - val loss: 115.4543 -
val mean absolute error: 8.3367
Epoch 11/100
20/20 [=============== ] - 0s 3ms/step - loss: 123.8664
- mean absolute error: 8.6454 - val loss: 102.8765 -
val mean absolute error: 7.8222
Epoch 12/100
20/20 [=========== ] - Os 3ms/step - loss: 106.8634
- mean absolute error: 7.9053 - val loss: 94.6192 -
val mean absolute error: 7.4617
Epoch 13/100
20/20 [============== ] - 0s 3ms/step - loss: 100.8406
- mean_absolute_error: 7.5977 - val_loss: 98.9632 -
val mean absolute error: 7.5522
Epoch 14/100
20/20 [============ ] - Os 3ms/step - loss: 108.3899
- mean absolute error: 7.7403 - val loss: 102.0158 -
val mean absolute error: 7.5881
Epoch 15/100
20/20 [============ ] - Os 3ms/step - loss: 106.1804
- mean absolute error: 7.8147 - val loss: 101.9140 -
val mean absolute error: 7.7441
Epoch 16/100
20/20 [=========== ] - Os 3ms/step - loss: 102.3587
- mean absolute error: 7.4590 - val loss: 79.9470 -
val mean absolute error: 6.7715
Epoch 17/100
mean absolute error: 6.8022 - val loss: 81.3400 -
val mean absolute error: 6.8737
Epoch 18/100
mean absolute error: 7.0133 - val loss: 85.9735 -
val mean absolute error: 7.0195
Epoch 19/100
```

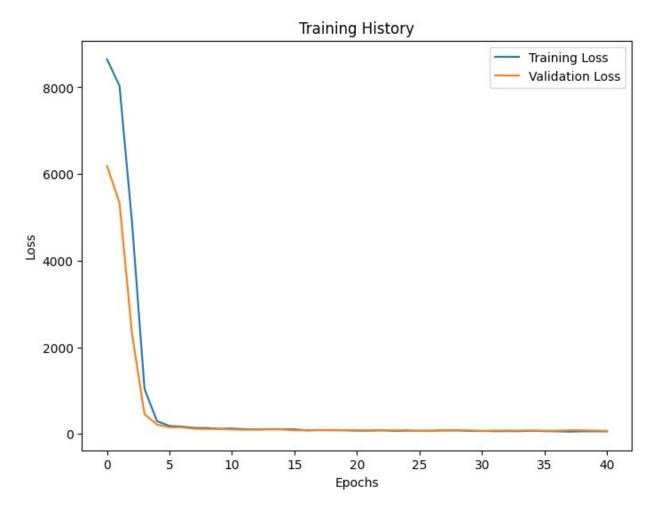
```
mean absolute error: 6.6208 - val loss: 89.1710 -
val mean absolute error: 7.2989
Epoch 20/100
20/20 [============= ] - 0s 3ms/step - loss: 80.7513 -
mean absolute error: 6.7058 - val loss: 84.8360 -
val mean absolute error: 7.0509
Epoch 21/100
20/20 [============== ] - 0s 3ms/step - loss: 74.2774 -
mean absolute error: 6.5526 - val loss: 79.8263 -
val mean absolute error: 6.7771
Epoch 22/100
20/20 [============= ] - Os 3ms/step - loss: 72.4148 -
mean absolute error: 6.4641 - val loss: 79.8499 -
val mean absolute error: 6.8056
Epoch 23/100
mean absolute_error: 6.5561 - val_loss: 77.0856 -
val mean absolute error: 6.5755
Epoch 24/100
20/20 [============= ] - 0s 3ms/step - loss: 67.6780 -
mean absolute error: 6.1833 - val loss: 80.7937 -
val mean absolute error: 6.7611
Epoch 25/100
20/20 [============== ] - 0s 2ms/step - loss: 73.6766 -
mean_absolute_error: 6.2056 - val loss: 80.0893 -
val mean absolute error: 6.8974
Epoch 26/100
mean absolute error: 6.3182 - val loss: 66.9290 -
val mean absolute error: 6.4197
Epoch 27/100
20/20 [============== ] - 0s 3ms/step - loss: 73.5757 -
mean absolute error: 6.4198 - val loss: 69.1794 -
val mean absolute error: 6.3932
Epoch 28/100
20/20 [============== ] - 0s 3ms/step - loss: 79.4177 -
mean absolute error: 6.4063 - val loss: 77.1633 -
val mean absolute error: 6.6143
Epoch 29/100
mean absolute error: 6.6154 - val loss: 76.0644 -
val mean absolute error: 6.7377
Epoch 30/100
mean absolute error: 6.1363 - val_loss: 78.9291 -
val mean absolute error: 6.9183
Epoch 31/100
```

```
mean absolute error: 6.1937 - val_loss: 66.3639 -
val mean absolute error: 6.2720
Epoch 32/100
20/20 [============= ] - 0s 2ms/step - loss: 60.3523 -
mean absolute error: 5.7764 - val loss: 74.7099 -
val mean absolute error: 6.6413
Epoch 33/100
mean absolute error: 6.0050 - val loss: 75.5275 -
val mean absolute error: 6.6400
Epoch 34/100
20/20 [============== ] - 0s 2ms/step - loss: 61.7266 -
mean absolute error: 5.7045 - val_loss: 73.9673 -
val mean absolute error: 6.5526
Epoch 35/100
mean absolute error: 6.1549 - val loss: 77.3238 -
val mean absolute error: 6.5474
Epoch 36/100
20/20 [============== ] - 0s 2ms/step - loss: 62.0706 -
mean absolute error: 5.8968 - val loss: 71.5218 -
val mean absolute error: 6.5054
Epoch 37/100
20/20 [============== ] - 0s 2ms/step - loss: 57.7333 -
mean absolute error: 5.6403 - val loss: 70.3059 -
val mean absolute error: 6.5438
Epoch 38/100
mean absolute error: 5.3124 - val loss: 82.3516 -
val mean absolute error: 7.0180
Epoch 39/100
mean absolute error: 5.6879 - val loss: 81.5721 -
val mean absolute error: 6.9372
Epoch 40/100
20/20 [============= ] - 0s 3ms/step - loss: 57.5124 -
mean absolute error: 5.6813 - val loss: 74.0030 -
val mean absolute error: 6.8042
Epoch 41/100
20/20 [============= ] - 0s 3ms/step - loss: 56.5801 -
mean absolute error: 5.6828 - val loss: 68.1339 -
val mean absolute error: 6.5257
```

Plot Training History

```
# Plot loss over epochs
plt.figure(figsize=(8, 6))
plt.plot(history.history['loss'], label='Training Loss')
plt.plot(history.history['val_loss'], label='Validation Loss')
```

```
plt.legend()
plt.xlabel('Epochs')
plt.ylabel('Loss')
plt.title('Training History')
plt.show()
```



Explanation:

Early Stopping: Prevents overfitting by stopping training when validation loss doesn't improve. Batch Size: Set to 32 for efficient computation. Validation Split: 20% of training data used for validation

1. Model Evaluation Evaluate on Test Data

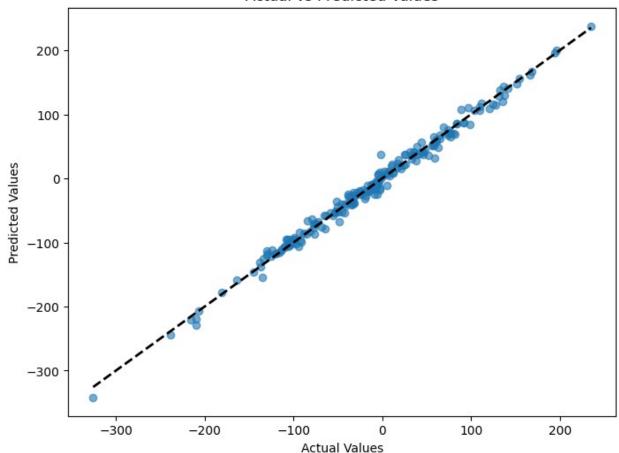
```
test_loss, test_mae = model.evaluate(X_test, y_test, verbose=0)
print(f'Test MSE: {test_loss:.4f}')
print(f'Test MAE: {test_mae:.4f}')
Test MSE: 73.7218
Test MAE: 6.5903
```

Predict on Test Data

Plot Predicted vs Actual Values

```
plt.figure(figsize=(8, 6))
plt.scatter(y_test, y_pred, alpha=0.6)
plt.xlabel('Actual Values')
plt.ylabel('Predicted Values')
plt.title('Actual vs Predicted Values')
plt.plot([y_test.min(), y_test.max()], [y_test.min(), y_test.max()],
'k--', lw=2)
plt.show()
```

Actual vs Predicted Values



Explanation:

R^2 Score: Indicates how well the model explains the variability of the target variable. Mean Squared Error: Average squared difference between predicted and actual values.

1. Optimization and Conclusion Hyperparameter Tuning To improve the model, consider tuning hyperparameters:

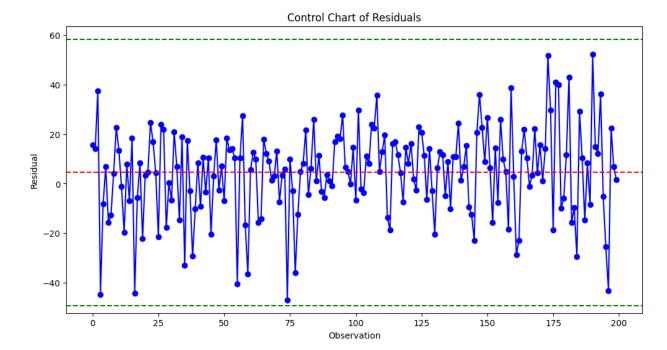
Number of Layers and Neurons: Experiment with different architectures. Learning Rate: Adjust optimizer learning rate. Batch Size and Epochs: Modify based on convergence behavior. Activation Functions: Try different activation functions like tanh or leaky ReLU. Cross-Validation Implement k-fold cross-validation to ensure model robustness.

```
from sklearn.model_selection import KFold

# Define KFold
kfold = KFold(n_splits=5, shuffle=True, random_state=42)

# Placeholder for results
mse_scores = []
```

```
for train index, val index in kfold.split(X):
   # Split data
   X train fold, X val fold = X[train index], X[val index]
   y train fold, y val fold = y[train index], y[val index]
   # Build model (you might want to define a function for this)
   model fold = Sequential()
   model_fold.add(Dense(64, input dim=n features, activation='relu'))
   model fold.add(Dense(128, activation='relu'))
   model fold.add(Dropout(0.2))
   model fold.add(Dense(64, activation='relu'))
   model fold.add(Dense(1, activation='linear'))
   model fold.compile(optimizer='adam', loss='mean squared error')
   # Train model
   model fold.fit(X train fold, y train fold, epochs=50,
batch size=32, verbose=0)
   # Evaluate model
   y val pred = model fold.predict(X val fold)
   mse fold = mean squared error(y val fold, y val pred)
   mse scores.append(mse fold)
print(f'Cross-Validated MSE: {np.mean(mse scores):.4f} ±
{np.std(mse scores):.4f}')
7/7 [=======] - 0s 1ms/step
7/7 [=======] - 0s lms/step
7/7 [======== ] - 0s 1ms/step
7/7 [======= ] - 0s 1ms/step
7/7 [======= ] - 0s 1000us/step
Cross-Validated MSE: 487.9427 ± 56.5823
# Calculate residuals
residuals = y test - y pred.flatten()
# Plot control chart
plt.figure(figsize=(12, 6))
plt.plot(residuals, marker='o', linestyle='-', color='b')
plt.axhline(y=np.mean(residuals), color='r', linestyle='--')
plt.axhline(y=np.mean(residuals) + 3*np.std(residuals), color='g',
linestvle='--')
plt.axhline(y=np.mean(residuals) - 3*np.std(residuals), color='g',
linestyle='--')
plt.title('Control Chart of Residuals')
plt.xlabel('Observation')
plt.ylabel('Residual')
plt.show()
```



Explanation:

Control Limits: Set at ±3 standard deviations from the mean residual. Purpose: Detect any out-of-control signals indicating model issues. Conclusion Model Performance: Our neural network captures the complex relationships in the data. Optimization: Further tuning and feature engineering can enhance performance. SPC Application: Helps in monitoring and maintaining model reliability over time.

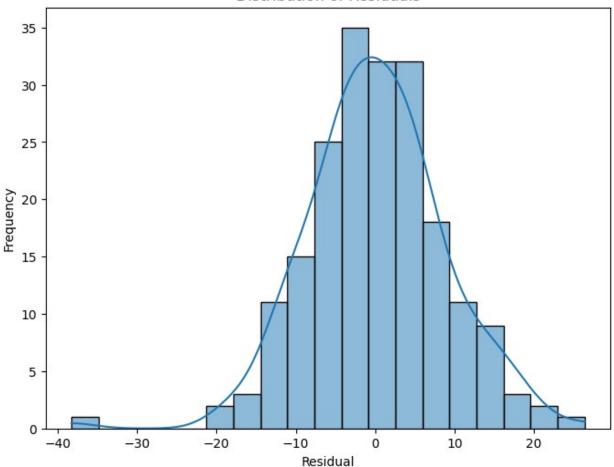
Next Steps Feature Engineering: Incorporate more interaction terms and nonlinear transformations. Advanced Techniques: Use models like XGBoost or Random Forests for comparison. Deployment: Implement the model in a production environment with continuous monitoring.

1. Applying Statistical Process Control Residual Analysis

```
# Calculate residuals
residuals = y_test - y_pred.flatten()

# Plot histogram of residuals
plt.figure(figsize=(8, 6))
sns.histplot(residuals, kde=True)
plt.title('Distribution of Residuals')
plt.xlabel('Residual')
plt.ylabel('Frequency')
plt.show()
```

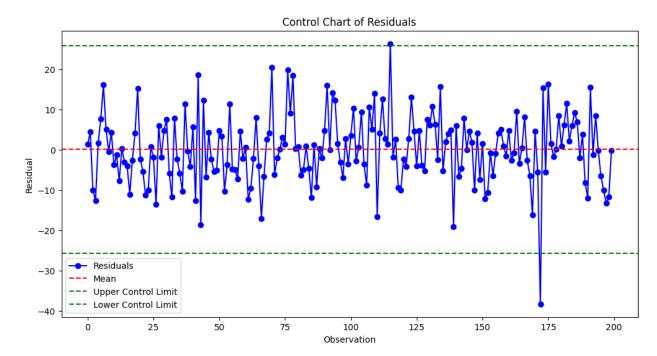
Distribution of Residuals



Control Chart of Residuals

```
# Control chart parameters
mean residual = np.mean(residuals)
std residual = np.std(residuals)
upper control limit = mean residual + 3 * std residual
lower control limit = mean residual - 3 * std residual
# Plot control chart
plt.figure(figsize=(12, 6))
plt.plot(residuals, marker='o', linestyle='-', color='b',
label='Residuals')
plt.axhline(y=mean_residual, color='r', linestyle='--', label='Mean')
plt.axhline(y=upper_control_limit, color='g', linestyle='--',
label='Upper Control Limit')
plt.axhline(y=lower_control_limit, color='g', linestyle='--',
label='Lower Control Limit')
plt.title('Control Chart of Residuals')
plt.xlabel('Observation')
plt.ylabel('Residual')
```

```
plt.legend()
plt.show()
```



Explanation Purpose: Identify any out-of-control points indicating model issues. Control Limits: Set at ±3 standard deviations from the mean residual.

Process Capability Analysis

```
# Process capability indices
process_std = np.std(y_test)
spec_limits = [np.min(y_test), np.max(y_test)] # Assuming
specifications are the min and max of y_test
process_mean = np.mean(y_test)

# Calculate Cp and Cpk
Cp = (spec_limits[1] - spec_limits[0]) / (6 * process_std)
Cpk = min((spec_limits[1] - process_mean), (process_mean -
spec_limits[0])) / (3 * process_std)

print(f'Process Capability Cp: {Cp:.4f}')
print(f'Process Capability Cpk: {Cpk:.4f}')
Process Capability Cp: 1.0575
Process Capability Cpk: 0.9323
```

Explanation Cp and Cpk: Indices to assess the process capability in meeting specifications. Interpretation: Cp > 1: Process has the potential to meet specifications. Cpk > 1: Process is centered between the specification limits.

- 1. Quality Management Considerations Plan-Do-Check-Act (PDCA) Cycle Plan: Defined the problem, objectives, and planned data generation and modeling approach. Do: Executed data generation, preprocessing, and model training. Check: Evaluated model performance and applied SPC tools. Act: Identified areas for improvement, such as model tuning or data quality enhancements. Continuous Improvement Feedback Loop: Use evaluation results to refine the model. Employee Training: Ensure team members understand SPC and DOE principles applied in modeling. Documentation: Maintain detailed records of modeling procedures and findings. Compliance with ISO Standards ISO 9001 Principles: Customer Focus: Model aims to improve process quality, benefiting customers. Process Approach: Systematic modeling process aligns with process approach principle. Improvement: Continuous model evaluation and refinement.
- 2. Conclusion We successfully developed a neural network model to optimize a complex process by incorporating SPC and DOE principles. The model captures linear relationships and interactions between variables, providing valuable insights into the process dynamics.

Key Takeaways Data Generation: Simulated a realistic process with interactions and variability. Modeling Approach: Employed neural networks capable of handling high-dimensional data. Quality Tools: Applied SPC methods to monitor and control model performance. Quality Management: Aligned the project with quality management principles for continuous improvement.