Strategies for Large-Scale Renewable Integration in Power Systems: A Focus on Sensitive Bus Analysis

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Appendix

Draft Version for Review.

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holder at danish@mdanish.me.
# This script is designed to be executed within the Jupyter Lab environment,
a web-based interactive development interface used for Python programming.
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
import statsmodels.api as sm
import warnings
# The script loads the dataset corresponding to the first feeder's 15 load
bus data from the Iowa 240 bus test system.
# The feeder 1 dataset can be found from this url:
https://github.com/mirsayedshah/230704.git
df = pd.read_csv('feeder1_load_data.csv')
# Step 0: Sensitivity analysis
# Calculate mean and standard deviation for each bus.
mean loads = df.mean()
std devs = df.std()
# Calculate normalized standard deviation for each bus.
norm std devs = std devs / mean loads
# Print out the buses with the highest normalized standard deviations. These
are the buses that are most sensitive to load changes over time.
most_sensitive_buses = norm_std_devs.sort_values(ascending=False)
print(most sensitive buses)
# Step 1: Data Preparation
# No specific data preparation steps are considered for this case study.
# Step 2: Correlation Analysis
# Select numeric columns from the dataframe.
numeric columns = df.select dtypes(include=[float, int]).columns
# Calculate correlation matrix.
correlation matrix = df[numeric columns].corr()
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# Visualize correlation matrix using a heatmap.
plt.figure(figsize=(10, 8))
sns.heatmap(correlation matrix, annot=True, cmap='coolwarm')
plt.title('Correlation Matrix')
plt.savefig('correlation matrix.png')
plt.show()
plt.close()
# Export correlation matrix to CSV file.
correlation matrix.to csv('correlation matrix.csv', index=True)
# Step 3: Regression Analysis
# Prepare the target variable and explanatory variables.
X = df[['B01', 'B02', 'B04', 'B05', 'B06', 'B07', 'B08', 'B09', 'B10', 'B11',
'B12', 'B13', 'B14', 'B15']]
y = df['B03']
# Add a constant term to the explanatory variables.
X = sm.add constant(X)
# Fit the regression model.
model = sm.OLS(y, X)
results = model.fit()
# Print regression results summary.
print(results.summary())
# Export regression results to CSV file.
results summary = results.summary().tables[1].as csv()
with open ('regression results.csv', 'w') as f:
    f.write(results summary)
# Visualization of regression results.
plt.figure(figsize=(10, 6))
plt.scatter(y, results.fittedvalues)
plt.xlabel('Actual B03')
plt.ylabel('Predicted B03')
plt.title('Regression Results - Actual vs Predicted B03')
plt.savefig('regression results.png')
plt.show()
plt.close()
# Create a DataFrame with actual and predicted values.
regression results df = pd.DataFrame({
    'Actual B03': y,
    'Predicted B03': results.fittedvalues
})
# Export regression results (actual vs predicted B03) to CSV.
regression results df.to csv('regression results actual vs predicted.csv',
index=False)
# Step 4: Granger Causality Analysis
# Perform the Granger causality test.
with warnings.catch warnings():
    warnings.simplefilter("ignore", category=FutureWarning)
    granger results = sm.tsa.stattools.grangercausalitytests(df[['B03',
'B01']], maxlag=3, verbose=False)
# Export Granger causality results to CSV file.
granger_results_summary = """
for lag in granger results.keys():
    result = granger results[lag][0]['ssr chi2test']
    result summary = f"Lag: {lag}\n"
    result_summary += f"F-statistic: {result[0]}\n"
    result summary += f"p-value: {result[1]}\n\n"
    granger results summary += result summary
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with open('granger causality results.csv', 'w') as f:
    f.write(granger results summary)
# Print Granger causality results.
for lag in granger results.keys():
    print(f'Lag: {lag}')
   print(granger results[lag])
# Visualization of Granger causality results.
lag values = []
p values = []
for lag in granger results.keys():
    lag values.append(lag)
    p values.append(granger results[lag][0]['ssr chi2test'][1])
plt.figure(figsize=(10, 6))
plt.plot(lag_values, p_values, marker='o')
plt.xlabel('Lag')
plt.ylabel('p-value')
plt.title('Granger Causality Results')
plt.xticks(lag values)
plt.axhline(y=0.05, color='red', linestyle='--', label='Significance
Threshold (p=0.05)')
plt.legend()
plt.savefig('granger_causality_results.png')
plt.show()
plt.close()
# This script is designed to be flexible and adaptable. Additional
visualizations, tables, calculations, and summaries can be incorporated as
needed to present the results. These elements can be tailored and modified
based on specific requirements, ensuring a comprehensive and customized data
analysis outcome.
# ...
# End of the script
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