```
import pandas as pd
import numpy as np
from sklearn.model_selection import train_test_split, learning_curve
from sklearn.preprocessing import StandardScaler, OneHotEncoder
from sklearn.ensemble import RandomForestRegressor, GradientBoostingRegressor
from sklearn.linear_model import LinearRegression
from sklearn.metrics import mean_absolute_error, mean_squared_error, r2_score
from sklearn.pipeline import Pipeline
from sklearn.compose import ColumnTransformer
from sklearn.impute import KNNImputer
import matplotlib.pyplot as plt
import seaborn as sns
import plotly.express as px
import plotly.graph_objects as go
from scipy import stats
try:
  from statsmodels.stats.outliers_influence import variance_inflation_factor
  statsmodels_available = True
except ImportError:
  statsmodels available = False
import warnings
warnings.filterwarnings('ignore')
# Set random seed for reproducibility
np.random.seed(42)
# 1. Load the dataset
def load_data(url):
  try:
     df = pd.read csv (url)
     print("Data loaded successfully.")
     print("\nDataset Column Types:")
     print(df.dtypes)
     return df
  except Exception as e:
     print(f"Error loading data: {e}")
     return None
url =
"https://raw.githubusercontent.com/ageron/handson-ml2/master/datasets/housing/housing.csv"
df = load data(url)
if df is None:
  raise SystemExit("Exiting due to data loading failure.")
```

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# Subsample dataset for faster training (comment out to use full dataset)
df = df.sample(frac=0.3, random_state=42)
print(f"\nSubsampled dataset to {len(df)} rows for faster training.")
# 2. Data Preprocessing
def preprocess data(df):
  print("\n=== Data Preprocessing ===")
  # Display missing values before imputation
  print("Missing Values Before Imputation:")
  print(df.isnull().sum())
  # Separate categorical and numerical columns
  categorical_cols = ['ocean_proximity'] if 'ocean_proximity' in df.columns else []
  numerical cols = [col for col in df.columns if col not in categorical cols]
  # Debug: Confirm columns
  print("\nNumerical Columns for Imputation:", numerical cols)
  print("Categorical Columns:", categorical_cols)
  # Handle missing values with KNN imputation for numerical columns
  try:
    if numerical_cols:
       imputer = KNNImputer(n_neighbors=5)
       df[numerical_cols] = pd.DataFrame(
          imputer.fit transform(df[numerical cols]),
          columns=numerical_cols,
          index=df.index
       )
  except Exception as e:
     print(f"Error during KNN imputation: {e}")
     print("Falling back to median imputation.")
     for col in numerical cols:
       df[col].fillna(df[col].median(), inplace=True)
  # Display missing values after imputation
  print("\nMissing Values After Imputation:")
  print(df.isnull().sum())
  # Remove duplicates
  initial rows = len(df)
  df.drop_duplicates(inplace=True)
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print(f"\nDuplicates Removed: {initial rows - len(df)}")
  # Cap outliers for numerical columns and track counts
  outlier counts = {}
  def cap outliers(series, col name):
     Q1 = series.quantile(0.25)
     Q3 = series.quantile(0.75)
     IQR = Q3 - Q1
    lower bound = Q1 - 1.5 * IQR
     upper_bound = Q3 + 1.5 * IQR
     outliers = ((series < lower_bound) | (series > upper_bound)).sum()
     outlier counts[col name] = outliers
     return series.clip(lower_bound, upper_bound)
  for col in numerical_cols:
     df[col] = cap_outliers(df[col], col)
  print("\nOutliers Detected and Capped:")
  for col, count in outlier_counts.items():
     print(f"{col}: {count} outliers")
  # Display skewness before log-transformation
  print("\nSkewness Before Log-Transformation:")
  print(df[numerical_cols].skew())
  # Log-transform skewed features
  skewed_cols = ['total_rooms', 'population', 'median_house_value']
  for col in skewed_cols:
     if col in df.columns:
       df[col] = np.log1p(df[col])
  # Display skewness after log-transformation
  print("\nSkewness After Log-Transformation:")
  print(df[numerical_cols].skew())
  # Verify median_house_value presence
  print("\nColumns After Preprocessing:", df.columns.tolist())
  return df
df = preprocess_data(df)
#3. Feature Engineering
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def engineer features(df):
  print("\n=== Feature Engineering ===")
  # Add core features (reduced set)
  df['rooms per household'] = df['total rooms'] / df['households']
  df['bedrooms per room'] = df['total bedrooms'] / df['total rooms']
  df['population_per_household'] = df['population'] / df['households']
  # Removed 'distance_to_coast' and 'median_income_poly1' to reduce feature count
  # Summarize new features
  print("New Features Created:")
  print(df[['rooms_per_household', 'bedrooms_per_room', 'population_per_household']].head())
  return df
df = engineer features(df)
# 4. Statistical Analysis
def statistical analysis(df):
  print("\n=== Statistical Analysis ===")
  stat, p value = stats.shapiro(df['median house value'])
  normality_result = f"Shapiro-Wilk Test for median_house_value: p-value = {p_value:.4f}"
  # Filter numerical columns for descriptive stats, skewness, and kurtosis
  numerical_cols = df.select_dtypes(include=['float64', 'int64']).columns
  print("\nNumerical Columns for Statistical Analysis:", numerical_cols.tolist())
  desc stats = df[numerical cols].describe().T
  desc_stats['skewness'] = df[numerical_cols].skew()
  desc stats['kurtosis'] = df[numerical cols].kurtosis()
  # Multcollinearity analysis (VIF)
  if statsmodels available:
     numerical_df = df[numerical_cols]
     vif data = pd.DataFrame()
     vif_data['Feature'] = numerical_df.columns
     vif_data['VIF'] = [variance_inflation_factor(numerical_df.values, i) for i in
range(numerical df.shape[1])]
     print("\nVariance Inflation Factor (VIF) Analysis:")
     print(vif_data)
  return normality_result, desc_stats
normality_result, desc_stats = statistical_analysis(df)
```

```
print("\nNormality Test:")
print(normality_result)
print("\nDescriptive Statistics with Skewness and Kurtosis:")
print(desc stats)
# 5. Exploratory Data Analysis (EDA)
def perform eda(df):
  print("\n=== Exploratory Data Analysis ===")
  # Correlation matrix
  corr_matrix = df.select_dtypes(include=['float64', 'int64']).corr()
  fig corr = go.Figure(data=go.Heatmap(
     z=corr matrix.values,
    x=corr_matrix.columns,
    y=corr_matrix.columns,
    colorscale='RdBu',
     zmin=-1, zmax=1,
     text=corr_matrix.values.round(2),
    texttemplate="%{text}",
    textfont={"size": 10}
  ))
  fig_corr.update_layout(title='Interactive Correlation Matrix', width=800, height=800)
  fig_corr.show()
  # Scatter plot (fixed syntax error)
  fig_scatter = px.scatter(df, x='median_income', y='median_house_value', title='Median_
Income vs Median House Value (USD)',
                 hover data=['longitude', 'latitude'],
                 trendline='ols')
  fig_scatter.update_yaxes(title_text='Median House Value (USD)')
  fig scatter.update traces(hovertemplate='Income: %{x}House Value: $%{y:,.2f} USD')
  fig scatter.show()
  # Geographical plot (requires Mapbox token)
  px.set mapbox access token('your mapbox token')
  fig_geo = px.scatter_mapbox(df, lat='latitude', lon='longitude', color='median_house_value',
                    size='population', zoom=5, mapbox_style='open-street-map',
                    title='House Prices by Location (USD)',
                    color continuous scale=px.colors.sequential.Plasma)
  fig_geo.update_coloraxes(colorbar_title='Median House Value (USD)')
  fig_geo.show()
  # Distribution plot
  plt.figure(figsize=(10, 6))
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sns.histplot(df['median house value'], kde=True)
  plt.title('Distribution of Median House Value (USD)')
  plt.xlabel('Median House Value (USD)')
  plt.show()
perform_eda(df)
#6. Prepare Data for Modeling
def prepare data(df):
  X = df.drop('median_house_value', axis=1)
  y = df['median house value']
  X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)
  return X_train, X_test, y_train, y_test
X_train, X_test, y_train, y_test = prepare_data(df)
#7. Modeling and Impact Analysis
def build pipeline(model):
  numerical_cols = X_train.select_dtypes(include=['float64', 'int64']).columns
  categorical cols = ['ocean proximity'] if 'ocean proximity' in df.columns else []
  preprocessor = ColumnTransformer([
     ('num', StandardScaler(), numerical cols),
     ('cat', OneHotEncoder(drop='first', handle_unknown='ignore'), categorical_cols)
  ])
  pipeline = Pipeline([
     ('preprocessor', preprocessor),
     ('regressor', model)
  ])
  return pipeline
def plot feature importance(model, feature names):
  importances = model.feature_importances_
  indices = np.argsort(importances)[::-1]
  plt.figure(figsize=(10, 6))
  plt.bar(range(len(importances)), importances[indices], align='center')
  plt.xticks(range(len(importances)), [feature_names[i] for i in indices], rotation=90)
  plt.title('Feature Importance')
  plt.tight_layout()
  plt.show()
```

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def plot learning curves(model, X train, y train):
  train_sizes, train_scores, val_scores = learning_curve(
     model, X_train, y_train, cv=3, scoring='r2', n_jobs=-1, train_sizes=np.linspace(0.1, 1.0, 5))
  train mean = np.mean(train scores, axis=1)
  train std = np.std(train scores, axis=1)
  val_mean = np.mean(val_scores, axis=1)
  val_std = np.std(val_scores, axis=1)
  plt.figure(figsize=(10, 6))
  plt.plot(train_sizes, train_mean, label='Training R2')
  plt.plot(train sizes, val mean, label='Validation R2')
  plt.fill between(train sizes, train mean - train std, train mean + train std, alpha=0.1)
  plt.fill_between(train_sizes, val_mean - val_std, val_mean + val_std, alpha=0.1)
  plt.xlabel('Training Examples')
  plt.ylabel('R2 Score')
  plt.title('Learning Curves')
  plt.legend(loc='best')
  plt.grid(True)
  plt.show()
def train_and_evaluate(X_train, X_test, y_train, y_test):
  print("\n=== Model Training and Evaluation ===")
  # Define models
  models = {
     'Linear Regression': LinearRegression(),
     'Random Forest': RandomForestRegressor(n estimators=50, max depth=10,
random_state=42, n_jobs=-1),
     'Gradient Boosting': GradientBoostingRegressor(n_estimators=50, random_state=42)
  }
  results = []
  best model = None
  best r2 = -np.inf
  numerical_cols = X_train.select_dtypes(include=['float64', 'int64']).columns
  categorical_cols = ['ocean_proximity'] if 'ocean_proximity' in df.columns else []
  feature_names = numerical_cols.tolist() + [f"ocean_proximity_{cat}" for cat in
OneHotEncoder(drop='first').fit(X_train[categorical_cols]).get_feature_names_out()] if
categorical_cols else numerical_cols.tolist()
  for name, model in models.items():
     print(f"\nTraining {name}...")
```

```
pipeline = build pipeline(model)
pipeline.fit(X_train, y_train)
# Predictions
y_pred = pipeline.predict(X_test)
# Evaluation metrics (convert back to USD)
y_test_usd = np.expm1(y_test)
y_pred_usd = np.expm1(y_pred)
mae = mean_absolute_error(y_test_usd, y_pred_usd)
rmse = np.sqrt(mean_squared_error(y_test_usd, y_pred_usd))
r2 = r2_score(y_test_usd, y_pred_usd)
results.append({
  'Model': name,
  'MAE (USD)': mae,
  'RMSE (USD)': rmse,
  'R2': r2
})
print(f"\n{name} Performance (in USD):")
print(f"Mean Absolute Error: ${mae:,.2f}")
print(f"Root Mean Squared Error: ${rmse:,.2f}")
print(f"R2: {r2:.4f}")
# Feature importance (for Random Forest and Gradient Boosting)
if name in ['Random Forest', 'Gradient Boosting']:
  plot_feature_importance(pipeline.named_steps['regressor'], feature_names)
# Learning curves
plot_learning_curves(pipeline, X_train, y_train)
# Residual analysis
residuals = y_test_usd - y_pred_usd
plt.figure(figsize=(10, 6))
stats.probplot(residuals, dist="norm", plot=plt)
plt.title(f'Q-Q Plot of Residuals ({name})')
plt.show()
# Update best model
if r2 > best_r2:
  best r2 = r2
  best_model = pipeline
```

```
# Display model comparison
  print("\n=== Model Comparison ===")
  results df = pd.DataFrame(results)
  print(results df)
  return best model
best_model = train_and_evaluate(X_train, X_test, y_train, y_test)
# 8. Gradio Interface for Hugging Face Spaces
try:
  import gradio as gr
  def predict_house_value(longitude, latitude, housing_median_age, total_rooms,
total bedrooms,
                 population, households, median_income, ocean_proximity):
     input data = pd.DataFrame({
       'longitude': [longitude],
       'latitude': [latitude],
       'housing_median_age': [housing_median_age],
       'total_rooms': [np.log1p(total_rooms)],
       'total bedrooms': [total bedrooms],
       'population': [np.log1p(population)],
       'households': [households],
       'median income': [median income],
       'ocean proximity': [ocean proximity],
       'rooms_per_household': [np.log1p(total_rooms) / households],
       'bedrooms_per_room': [total_bedrooms / np.log1p(total_rooms)],
       'population_per_household': [np.log1p(population) / households]
    })
     prediction = best_model.predict(input_data)
     usd value = np.expm1(prediction[0]) # Reverse log transformation
     return f"Predicted House Value: ${usd value:,.2f} USD"
  iface = gr.Interface(
     fn=predict house value,
     inputs=[
       gr.Slider(-124, -114, step=0.1, label="Longitude"),
       gr.Slider(32, 42, step=0.1, label="Latitude"),
       gr.Slider(0, 52, step=1, label="Housing Median Age"),
       gr.Slider(0, 40000, step=100, label="Total Rooms"),
       gr.Slider(0, 7000, step=10, label="Total Bedrooms"),
```

```
gr.Slider(0, 50000, step=100, label="Population"),
gr.Slider(0, 7000, step=10, label="Households"),
gr.Slider(0, 15, step=0.1, label="Median Income"),
gr.Dropdown(choices=['<1H OCEAN', 'INLAND', 'NEAR OCEAN', 'NEAR BAY',
'ISLAND'], label="Ocean Proximity")
],
outputs="text",
title="California House Price Predictor",
description="Enter features to predict the median house value (in USD)."
)

print("\nGradio interface is ready. Run iface.launch() in a Hugging Face Space to use it.")
except ImportError:
print("\nGradio not installed. Skipping Gradio interface. Install gradio to enable.")
```