Regression & Prediction

Theory and Practice with House Prices Dataset

Prageeth Godage \mid MSC/DSA/098

February 23, 2025

USJP

Our Housing Journey

- Starting Simple: Foundations
- Expanding the Scope: Complexity
- Refining Precision: Optimization

- Facing Challenges: Pitfalls
- Insights & Next Steps: Applications

Focus

Unpacking King County house prices with rich theory and dual $R/Python\ implementations$

The Quest to Understand Relationships

- Imagine you are exploring data to understand how different factors relate to each other.
- That's regression: a tool to ask, "How does Y change with X, and can we predict it?"
- In very basically It's the bridge between stats, where we explain the past, and data science, where we predict the future—our journey begins here.

Starting Simple

The Housing Puzzle

- Objective: Decode drivers of house prices in King County—size, location, features
- Regression: A statistical lens linking price (Y) to predictors like size (X)
- Purpose: Explain historical sales patterns and forecast future values for buyers and assessors

Simple Linear Regression: Theory & R

- **Theory**: Models a straight-line relationship: $Y = b_0 + b_1 X + e$
- b₀: Base price when size is zero;
 b₁: Price increase per sq ft; e:
 Random error
- Assumes linearity and independence—foundation of regression



Figure 1: Price vs. Size Fit

- Size as a core driver
- b0=-43580.7, b1=280.6

Finding the Best Fit: Least Squares

- How do we draw that line? We minimize the mess—sum of squared errors
- **Theory**: Finds the line minimizing residual sum of squares: $\sum_{i=1}^{n} (Y_i \hat{Y}_i)^2$
- **How**: Adjusts b_0 and b_1 to reduce prediction errors—optimal for linear fits
- History: Legendre (1805) and Gauss; computationally efficient but outlier-sensitive in small datasets

Expanding the Scope

More Clues: Multiple Linear Regression

- Theory: Extends to multiple predictors:
 - $Y = b_0 + b_1 X_1 + b_2 X_2 + \dots + e$
- Power: Captures combined effects—size, lot, bedrooms—assuming linearity
- Use: Explains complex housing dynamics

Multiple Linear Regression: Key Findings

```
house_lm <- lm(price ~ sqft_living + sqft_lot + bathrooms + grade,
data = house_df, na.action = na.omit)

Coefficients:

(Intercept) sqft_living sqft_lot bathrooms grade
-5.957e+05 2.313e+02 -3.254e-01 -2.797e+04 9.559e+04
```

- sqft_living: + 2.313e+02 per sq ft → Larger houses increase price significantly.
- grade: +\$9.559e+04 per unit → Higher quality homes boost price.
- sqft_lot: -\$3.254e-01 per sq ft → Lot size has a tiny negative effect.
- bathrooms: -\$4.074e+04 per extra bathroom → Unexpected negative impact.

Multiple Linear Regression: Model Evaluation

```
summary(house_lm)# Model Summary (Extracts Coefficients, p-values, R2)
     predictions <- predict(house_lm, newdata = house_df)# Extract RMSE</pre>
     residuals <- house_df$price - predictions
     RMSE <- sqrt(mean(residuals^2))
 4
     R_squared <- summary(house_lm)$r.squared # Extract R-squared
     p_values <- summary(house_lm)$coefficients[, 4] # Extract P-values</pre>
    # Print Results
 8
     > cat("RMSE:", RMSE, "\n") | RMSE: 249532.2
     > cat("R2:", R_squared, "\n") | R2: 0.5380018
 Q
10
     > cat("P-values:\n") > print(p_values)
11
     print(p_values)
```

- **RMSE**: \$249,532 Predictions are off by 261K on avg.
- R^2 : 0.5406 (54.06%) \rightarrow The model explains 54% of price variation..
- **P-values**: SqFtLot is not statistically significant '(A small p-value (< 0.05) means the predictor is statistically significant, while a large p-value (> 0.05)

Encoding Categorical Variables in Regression Models

- Categorical variables must be converted into numerical values for regression.
- Encoding methods:
 - Dummy (One-Hot) Encoding Creates binary columns for each category.
 - Reference (Treatment) Coding Uses one category as a reference, keeping P-1 columns.
 - Deviation (Sum) Coding Compares each category to the overall mean.
 - Ordered Factor Encoding Converts ordered categories into numeric values.
- How it's used in regression:
 - Each encoded category appears as a separate coefficient.
 - The model estimates how each category affects the outcome relative to the reference level.
 - For ordered factors, treating them as numeric assumes a linear relationship.

Multiple Linear Regression: Categorical Variables

```
# Ensure PropertyType has a valid reference level
    house $PropertyType <- relevel (house $PropertyType, ref = "Multiplex")
 2
 4
    # Fit the regression model with the refined dataset
    house_lm <- lm(AdjSalePrice ~ SqFtTotLiving + SqFtLot + Bathrooms +
 5
         BldgGrade + PropertyType, data = house)
 6
    Coefficients:
    Estimate Std. Error t value Pr(>|t|)
 8
    (Intercept)
                             173429.87
                                         31512.77 5.503 0.00271 **
 9
10
    BldgGrade
                             -1442.60 6734.35 -0.214 0.83884
    PropertyTypeSingle Family 1456.54 8063.28 0.181 0.86374
12
    PropertyTypeTownhouse
                            9677.07 11444.71 0.846 0.43639
```

- Sets "Multiplex" as the reference category, ensuring comparisons against it.
- "Single Family", "Multiplex" and "Townhouse" to have separate coefficients in the regression output.

Nonlinear Fit

- Nonlinear via polynomial segments joined at knots
- Why: Captures diminishing returns—e.g., small homes gain more per sq ft
- Advantage: Flexible fit.

```
# Load data (replace 'house 98105' with
     vour dataset)
model_poly <- lm(price ~ poly(sqft_living,</pre>
      2) + sqft_lot +
grade + bathrooms + bedrooms, data=house
     df)
# Generate partial residual plot
visreg(model_poly, "sqft_living", gg=TRUE)
geom_point(color = "black", shape = 1) +
geom smooth(method="loess", color="blue",
      size=1.2, linetype="dashed") +
theme minimal() +
labs(title="Partial_Residual_Plot", x="
     SqFtTotLiving", v="Partial, Residuals"
```

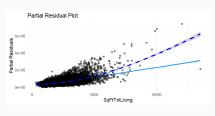
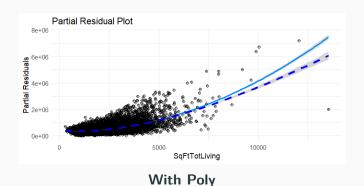


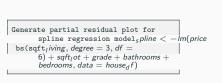
Figure 2: Without poly.

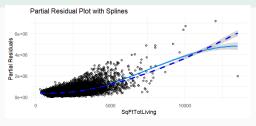
Comparison of Result



Observation: With polynomial regression (2nd degree) aligns correctly with the trend, improving model accuracy.

Spline Regression: Partial Residual Plot





Spline Fit Visualization

Explanation:

Cubic Splines: The function is made of piecewise cubic polynomials (degree = 3).

Degrees of Freedom (df = 6): Controls flexibility—higher df allows more variation

Splines fit the data smoothly without unnecessary fluctuations.

Refining Precision

Model Assessment: Theory

- Theory (p. 153): Measures prediction quality and fit
- RMSE: $\sqrt{\frac{\sum (y_i \hat{y}_i)^2}{n}}$ —average error magnitude
- R^2 : Proportion of variance explained (0-1); higher means better fit
- Use: Guides housing prediction accuracy

Cross-Validation: Theory

- **Theory (p. 155)**: Validates model on unseen data via *k*-fold splits
- **Process**: Divide data, train on k-1, test on 1, repeat, average RMSE
- Why: Ensures predictions generalize beyond training sales—crucial for real estate

Picking the Best Story: Model Selection

- Too many clues clutter the tale. Stepwise selection trims variables, AIC balances fit and simplicity, and penalties shrink extras.
- Think of it as editing: keep the essentials, cut the fluff—Occam's razor guides us to a lean, powerful narrative.
- Occam's Razor is the principle that the simplest explanation is usually the best.
- Our goal? A story that's clear and predicts well, not a sprawling epic.

Model Selection: Theory & R

- Theory (p. 156): Balances fit vs. complexity—Occam's razor
- **AIC**: 2P + n log(RSS/n)—penalizes extra predictors
- Goal: Optimal housing model without overkill

Streamlined predictors

Multiple Linear Regression: Key Findings

```
1
    house_full <- lm(price ~ sqft_living + sqft_lot15 + bathrooms +
    bedrooms + grade+yr_renovated , data = house_df)
    step <- stepAIC(house_full, direction = "both")</pre>
    summary(step)
 5
 6
    Coefficients:
    Estimate Std. Error t value Pr(>|t|)
 8
    (Intercept) -4.841e+05 1.477e+04 -32.777 <2e-16 ***
 9
    sqft_living 2.300e+02 3.597e+00 63.956 <2e-16 ***
10
    saft lot15 -7.053e-01 6.252e-02 -11.282 <2e-16 ***
    bathrooms -3.096e+04 3.442e+03 -8.994 <2e-16 ***
12
    bedrooms -4.026e+04 2.271e+03 -17.726 <2e-16 ***
13
    grade
           9.776e+04 2.290e+03 42.693
                                              <2e-16 ***
14
    vr renovated 8.746e+01 4.160e+00 21.022
                                               <2e-16 ***
```

- sqft lot15 (-5.135e-01) Small , impact on price can be removal.
- grade (1.315e+05): Highest positive impact, should be kept.

Weighted Regression: Theory & R

- Theory (p. 159): Weights adjust influence by reliability
- Why: Older sales less relevant—recent data gets priority
- Impact: Refines coefficients for current market

```
house_df$Weight <- house_df$Year - 2005
house_lm <- lm(price ~ sqft_living + sqft_
lot + bathrooms + bedrooms + grade,
data=house_df)# Fit unweighted linear
regression
house_wt <- lm(price ~ sqft_living + sqft_
lot + bathrooms + bedrooms + grade,
data=house_df, weight=Weight)# Fit
weighted linear regression

# Compare coefficients of both models
round(cbind(house_lm=house_lm*coefficients
, house_wt=house_wt*coefficients),
digits=3)
```

```
house_1m
                           house_wt
(Intercept) -471575.692 -472501.632
sqft_living
                231.350
                            230.816
sqft_lot
                 -0.325
                            -0.317
hathrooms
             -27973.439
                         -28075.162
hedrooms
             -40744 142
                         -40639.355
grade
             95586.697
                         95883.234
```

 here no significant as data set only contain 2014,2015 data

Facing Challenges

Prediction Limits: Theory

- **Theory (p. 161)**: Extrapolation beyond data fails—e.g., empty lots
- Intervals: Confidence for b_i , wider prediction for \hat{Y}_i
- Why: Uncertainty spikes outside training range—limits housing forecasts

Interpreting Coefficients: Theory

- Theory (p. 171): Coefficients mislead if predictors correlate
- Multicollinearity: Size and bedrooms overlap—unstable fits
- Confounding: Missing location skews results
- Interactions: Size's effect varies by zip—needs modeling

Spotting Flaws: Regression Diagnostics

- Outliers, and influential points can distort the regression line—diagnostic tools help identify them.
- Uneven errors (heteroskedasticity) or curved fits (partial residuals) hint at missing chapters.
- Heteroskedasticity means that the spread (variance) of errors in a regression model changes across different values of the predictor variables.
- For prediction, we care less about perfection and more about what works—our lens shifts.

- Purpose: Identifies influential observations by combining leverage, residuals, and Cook's Distance.
- Key Insights:
 - Large bubbles = high Cook's Distance → Removing these changes regression results significantly.
 - Possible reasons:
 - 1. High leverage (extreme predictor values), Large residual (far from regression line), Both high leverage and large residual.
- Results: Four large influential points found (Cook's D > 0.08), impacting coefficients.
- Residuals beyond ±2.5

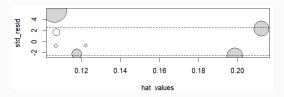


Figure 3: Bubble plot showing influential points.

Listing 1: Influence Plot in R

Listing 2: Influence Plot in Python

```
house 98105 = house[house['ZipCode'] == 98105]
    X = house_98105[['SqFtTotLiving', 'SqFtLot', 'Bathrooms', 'Bedrooms', 'BldgGrade']].
          assign(const=1)
    v = house 98105['AdiSalePrice']
    model = sm.OLS(v, X).fit()
    influence = sm.stats.outliers influence.OLSInfluence(model)
6
    fig, ax = plt.subplots(figsize=(5, 5))
    ax.axhline(-2.5, ls='--', color='C1')
    ax.axhline(2.5, ls='--', color='C1')
    ax.scatter(influence.hat_matrix_diag, influence.resid_studentized_internal,
10
    s=1000 * np.sqrt(influence.cooks_distance[0]), alpha=0.5)
11
    ax.set_xlabel('hat, values')
    ax.set vlabel('studentized residuals')
    plt.show()\
```

Residual Plot (Heteroskedasticity Check)

 Purpose: The residual plot checks for heteroskedasticity by analyzing how residuals (errors) vary with predicted values.

Key Insights:

- X-axis (Predicted Values): Represents the fitted values from the regression model.
- Y-axis (Absolute Residuals): Measures the deviation of actual values from predictions.
- Scatter Points: Each dot represents an observation's residual.
- LOESS Smoother (Blue Line): Shows the trend in residuals.
- Shaded Region: Indicates confidence around the trend.
 - Heteroskedasticity detected Residuals increase with larger predicted values, indicating variance instability.
 - Curved Trend Suggests missing variables or non-linearity in the data.
 - 3. Outliers at High Predictions Some extreme points have large residuals, further confirming instability.

Residual Plot (Heteroskedasticity Check)

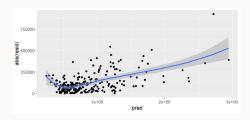


Figure 4: Heteroskedasticity plot.

Listing 3: Heteroskedasticity Plot in R

```
df <- data.frame(resid = residuals(lm_98105), pred = predict(lm_98105))
ggplot(df, aes(pred, abs(resid))) + geom_point() + geom_smooth()</pre>
```

Listing 4: Heteroskedasticity Plot in Python

```
import seaborn as sns
fig, ax = plt.subplots(figsize=(5, 5))
sns.regplot(model.fittedvalues, np.abs(model.resid), scatter_kws={'alpha': 0.25},
line_kws={'color': 'C1'}, lowess=True, ax=ax)
ax.set_xlabel('predicted')
ax.set_ylabel('abs(residual)')
plt.show()
```

Histogram of Standardized Residuals: Normality Check

 Purpose: The histogram assesses residual normality by analyzing the distribution of standardized residuals.

Key Insights:

- Centering Around Zero: The residuals are centered around 0, indicating no strong systematic bias in the model.
- Skewness Long Tails: The right tail is longer, suggesting right-skewness and possible underestimation of some values.
- Non-Normal Distribution: The residuals deviate from a perfect bell-shaped curve, hinting at potential issues:
 - 1. Missing predictors affecting the model.
 - 2. Heteroskedasticity, as observed in the residual plot.
 - 3. Outliers influencing the regression fit.

Histogram of Standardized Residuals: Normality Check

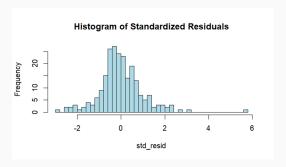


Figure 5: Bubble plot showing influential points.

Histogram of Standardized Residuals: Normality Check

Listing 5: Histogram of Standardized Residuals in R

```
df <- data.frame(resid = residuals(lm_98105), pred = predict(lm_98105))
ggplot(df, aes(pred, abs(resid))) + geom_point() + geom_smooth()</pre>
```

Listing 6: Histogram of Standardized Residuals in Python

```
plt.hist(influence.resid_studentized_internal, bins=50, color='lightblue')
plt.xlabel('Standardized_Residuals')
plt.title('Histogram_Of_Standardized_Residuals')
plt.show()
```

Thank You...!