# Wine Quality Prediction Using Support Vector Machines and Logistic Regression

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October 2, 2025

# **Declaration**

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# 1 Introduction

This project implements binary classification algorithms from scratch to predict wine quality based on physicochemical properties. Wines are classified as "good" (quality  $\geq 6$ ) or "bad" (quality < 6). We implement **Support Vector Machines (SVM)** and **Logistic Regression (LR)** with their kernelized variants (polynomial and RBF kernels) to capture non-linear decision boundaries.

**Objectives:** (1) Implement SVM and LR from scratch in Python; (2) Extend both with kernel methods; (3) Apply rigorous ML methodology with cross-validation; (4) Compare model performance comprehensively.

# 2 Theoretical Background

#### 2.1 Logistic Regression

Logistic Regression models the probability of binary outcomes using the sigmoid function:

$$\sigma(z) = \frac{1}{1 + e^{-z}}, \quad P(y = 1|\mathbf{x}) = \sigma(\mathbf{w}^T \cdot \mathbf{x} + b)$$
(1)

The model minimizes binary cross-entropy loss with L2 regularization:

$$\mathcal{L} = -\frac{1}{m} \sum_{i=1}^{m} \left[ y_i \log(\hat{y}_i) + (1 - y_i) \log(1 - \hat{y}_i) \right] + \frac{\lambda}{2m} \|\mathbf{w}\|^2$$
 (2)

Gradients for parameter updates:

$$\frac{\partial \mathcal{L}}{\partial \mathbf{w}} = \frac{1}{m} \mathbf{X}^{T} (\hat{\mathbf{y}} - \mathbf{y}) + \frac{\lambda}{m} \mathbf{w}, \quad \frac{\partial \mathcal{L}}{\partial b} = \frac{1}{m} \sum_{i=1}^{m} (\hat{y}_{i} - y_{i})$$
(3)

# 2.2 Support Vector Machine

SVM finds a maximum-margin hyperplane using hinge loss. For labels  $y \in \{-1, +1\}$ :

$$\min_{\mathbf{w},b} \frac{1}{2C} \|\mathbf{w}\|^2 + \frac{1}{m} \sum_{i=1}^m \max(0, 1 - y_i(\mathbf{w}^T \mathbf{x}_i + b))$$
 (4)

Hinge loss penalizes margin violations: samples with  $y \cdot (\mathbf{w}^T \mathbf{x} + b) < 1$  contribute to the gradient, while correctly classified samples beyond the margin do not.

# 2.3 Kernel Methods

Kernels map data to higher-dimensional spaces without explicit transformation via the kernel trick:

$$K(\mathbf{x}_i, \mathbf{x}_j) = \phi(\mathbf{x}_i)^T \cdot \phi(\mathbf{x}_j) \tag{5}$$

Decision function in dual space:

$$f(\mathbf{x}) = \sum_{i=1}^{m} \alpha_i y_i K(\mathbf{x}_i, \mathbf{x})$$
 (6)

Implemented kernels:

1. Linear:  $K(\mathbf{x}, \mathbf{y}) = \mathbf{x}^T \cdot \mathbf{y}$ 

2. Polynomial:  $K(\mathbf{x}, \mathbf{y}) = (\mathbf{x}^T \cdot \mathbf{y} + c)^d$ 

3. **RBF:**  $K(\mathbf{x}, \mathbf{y}) = \exp(-\gamma ||\mathbf{x} - \mathbf{y}||^2)$ 

# 3 Dataset and Preprocessing

# 3.1 Dataset Description

The UCI Wine Quality dataset contains 6,497 wines (1,599 red, 4,898 white) with 11 physicochemical features plus wine type. Quality scores range from 3 to 9. We converted this to binary classification: bad (quality < 6, 36.7%) vs good (quality  $\ge 6, 63.3\%$ ), resulting in a 1.73:1 class imbalance.

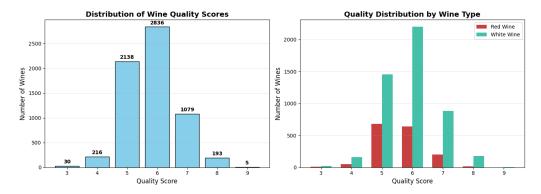


Figure 1: Wine quality distribution showing most wines rated 5 or 6.

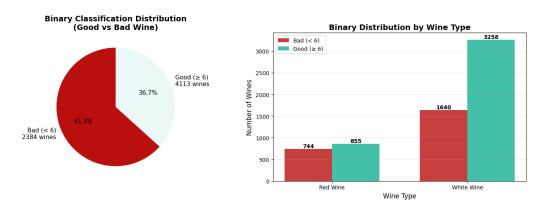


Figure 2: Binary classification distribution (63.3% good vs 36.7% bad).

# 3.2 Feature Analysis

The dataset contains 12 features describing physicochemical properties and wine type: Features:

- 1. Fixed Acidity (g/L): Non-volatile acids (tartaric acid)
- 2. Volatile Acidity (g/L): Acetic acid, can cause vinegar taste
- 3. Citric Acid (g/L): Adds freshness and flavor
- 4. Residual Sugar (g/L): Remaining sugar after fermentation
- 5. Chlorides (g/L): Salt content
- 6. Free Sulfur Dioxide (mg/L): Prevents microbial growth
- 7. **Total Sulfur Dioxide** (mg/L): Total  $SO_2$  (free + bound)
- 8. **Density** (g/cm<sup>3</sup>): Related to alcohol and sugar content

- 9. **pH**: Acidity level (3-4 for most wines)
- 10. Sulphates (g/L): Wine additive, affects  $SO_2$  levels
- 11. **Alcohol** (% vol): Alcohol percentage
- 12. Wine Type: Binary (0=red, 1=white)

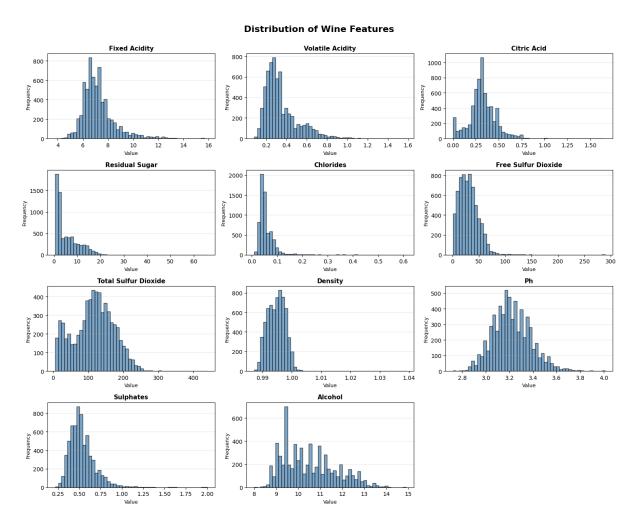


Figure 3: Distribution histograms for all features showing varying skewness patterns.

# Distribution patterns observed:

- **Highly right-skewed:** Residual sugar (most wines dry, few sweet), chlorides, free and total sulfur dioxide
- Moderately right-skewed: Sulphates, citric acid
- Approximately normal: Fixed acidity, volatile acidity, density, pH
- Multimodal: Alcohol (distinct peaks at different concentrations)

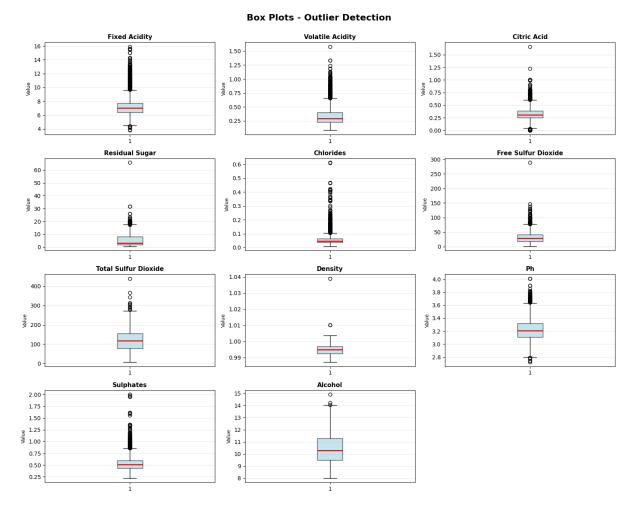


Figure 4: Box plots revealing outliers across features. Residual sugar, chlorides, and sulfur compounds show numerous extreme values.

Outlier analysis: Box plots reveal significant outliers in multiple features. Residual sugar shows the most extreme outliers (some wines with 60+ g/L vs median 2-3 g/L), indicating a few dessert wines. Chlorides and sulfur dioxide compounds also exhibit many outliers. Fixed acidity shows some high outliers (15+ g/L). These outliers likely represent legitimate wines with unusual characteristics rather than measurement errors. Our z-score normalization mitigates outlier impact while preserving relative relationships.

**Key observations:** Most wines clustered around quality 5-6; very few extreme ratings (30 wines at quality 3, only 5 at quality 9). The right-skewed distributions suggest most wines have moderate levels of sugar, chlorides, and sulfur compounds, with occasional wines having much higher concentrations.

# 3.3 Data Preprocessing

Stratified train-test split: 80% training (5,199 samples), 20% test (1,298 samples). Stratification maintained class balance in both sets.

**Z-score normalization:** Features standardized using training set statistics only to prevent data leakage:

$$\mathbf{X}_{norm} = \frac{\mathbf{X} - \mu_{train}}{\sigma_{train}} \tag{7}$$

# 3.4 Hyperparameter Tuning

**5-fold stratified cross-validation** used for hyperparameter selection. For Logistic Regression regularization  $\lambda \in \{0.001, 0.01, 0.1, 1.0\}$ , we found  $\lambda = 1.0$  optimal (CV accuracy: 73.84%  $\pm 1.46\%$ ).

Other hyperparameters: LR learning rate 0.1, SVM learning rate 0.001, 1,000 iterations for standard models, 500 for kernel models, SVM C=1.0. Kernel parameters tested: polynomial degrees 2 and 3, RBF gamma values 0.1 and 0.5.

# 4 Results

#### 4.1 Model Performance

Table 1 summarizes test set performance for all models.

Model	Accuracy	Precision	Recall	F1-Score
Logistic Regression	75.65%	78.24%	85.28%	81.61%
SVM	69.34%	74.04%	79.44%	76.64%
LR - Poly $(d=2)$	63.87%	83.16%	53.89%	65.39%
LR - Poly $(d=3)$	65.79%	72.88%	72.88%	72.96%
LR - RBF ( $\gamma = 0.1$ )	$\pmb{76.19\%}$	77.71%	87.47%	82.31%
LR - RBF ( $\gamma = 0.5$ )	74.19%	74.72%	89.54%	81.46%
SVM - Poly $(d=2)$	57.47%	81.40%	42.58%	55.91%
SVM - Poly $(d=3)$	68.80%	75.79%	75.79%	75.47%
SVM - RBF ( $\gamma = 0.1$ )	68.03%	68.78%	90.63%	78.22%
SVM - RBF ( $\gamma = 0.5$ )	68.80%	68.57%	92.82%	79.03%

Table 1: Test set performance comparison

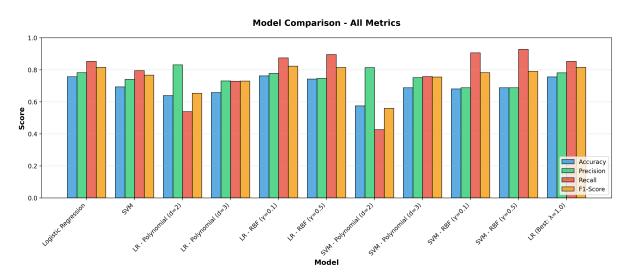


Figure 5: Performance comparison across all models and metrics.

Best model: Kernel LR with RBF ( $\gamma=0.1$ ) achieved 82.31% F1-score and 76.19% accuracy.

**Key findings:** (1) LR consistently outperformed SVM (6.3% accuracy advantage); (2) RBF kernels significantly better than polynomial kernels; (3) Lower gamma (0.1) generalized better than higher gamma (0.5); (4) Polynomial degree 2 severely underfit, degree 3 showed

improvement but remained suboptimal; (5) Standard LR competitive with kernel variants, suggesting linear relationships dominate.

# 4.2 Best Model Analysis

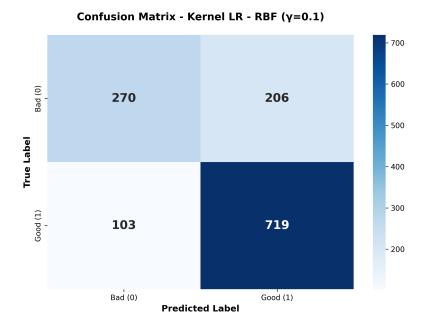


Figure 6: Confusion matrix for best model (Kernel LR with RBF,  $\gamma = 0.1$ ).

The best model correctly classified 989/1,298 samples (76.19

#### Error breakdown:

- True Positives: 719 (correctly identified good wines)
- True Negatives: 270 (correctly identified bad wines)
- False Positives: 206 (bad wines misclassified as good)
- False Negatives: 103 (good wines misclassified as bad)

The 2:1 FP:FN ratio indicates bias toward predicting "good," reflecting the 63.3% good-wine prevalence in the dataset.

#### 4.2.1 Misclassification Analysis

We examined misclassified samples to understand model limitations. Of 309 errors:

- False Positives (206): Bad wines predicted as good
- False Negatives (103): Good wines predicted as bad

### Sample misclassified examples:

Example 1 - False Positive (Bad predicted as Good):

- Actual: Bad (quality < 6)
- Features: High volatile acidity (0.558), high pH (1.270), high sulphates (1.230), red wine
- Issue: High sulphates and certain acid levels may correlate with good wines, but high volatile acidity (vinegar taste) makes it bad

Example 2 - False Positive (Bad predicted as Good):

- Actual: Bad (quality < 6)
- Features: Very high residual sugar (3.596), high chlorides (1.831), high alcohol (1.931), white wine
- **Issue:** Model likely associates high alcohol with quality, but excessive sugar and salt create imbalance

Example 3 - False Negative (Good predicted as Bad):

- Actual: Good (quality  $\geq 6$ )
- Features: High volatile acidity (0.925), low citric acid (-0.965), low alcohol (-1.079), red wine
- **Issue:** Low alcohol strongly associated with bad wines, but other factors compensate in reality

#### Patterns identified:

False Positives characteristics:

- 1. **Conflicting features:** Wines with some good indicators (e.g., high alcohol, proper pH) but critical flaws (e.g., high volatile acidity, excessive chlorides)
- 2. **Imbalanced chemistry:** High sugar with high salt, or high alcohol with poor acidity balance
- 3. Single strong features: Model overweights individual positive features, ignoring negative combinations

False Negatives characteristics:

- 1. Low alcohol bias: Wines with alcohol < 9.5% often predicted as bad despite other good qualities
- 2. Threshold proximity: Many are borderline quality=6 wines where small errors matter
- 3. **Atypical profiles:** Good wines with unusual feature combinations not well-represented in training data

#### Model limitations revealed:

- 1. **Feature interaction blindness:** Linear (even kernelized) models struggle with complex feature interactions. A wine might have high alcohol (good) and high volatile acidity (bad), but the model doesn't capture that this specific combination is unacceptable.
- 2. Threshold artifacts: Binary classification at quality=6 creates a sharp boundary where wines rated 5.5 and 6.5 (if continuous) would be very similar but are forced into different classes.
- 3. **Majority class bias:** With 63.3% good wines, the model defaults to "good" when uncertain, explaining the 2:1 FP:FN ratio.
- 4. Rare profile handling: Wines with unusual but valid feature combinations (e.g., low alcohol but excellent acid balance) are often misclassified due to insufficient training examples.

5. **Missing domain knowledge:** The model doesn't understand that certain feature combinations are chemically incompatible with quality, treating all features independently.

These limitations suggest that future improvements should focus on: (1) explicit feature interaction terms; (2) ensemble methods to capture different decision patterns; (3) class-weighted loss to reduce majority bias; (4) multi-class formulation to preserve gradual quality transitions.

# 4.3 Training Curves

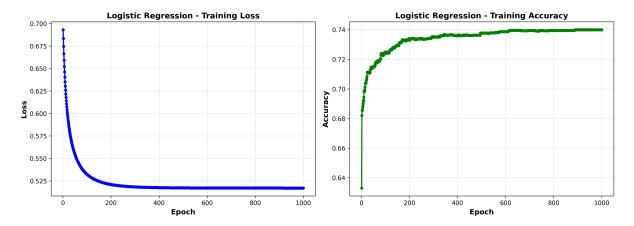


Figure 7: LR training curves showing smooth convergence without overfitting.

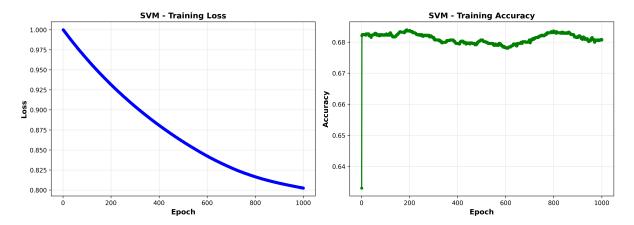


Figure 8: SVM training curves showing slower convergence with more oscillation.

LR converged rapidly (loss  $0.70 \rightarrow 0.52$  by epoch 300) with stable accuracy plateau at 74%. SVM converged more slowly (loss  $1.00 \rightarrow 0.80$ ) with greater oscillation, plateauing at 68%. Both models showed no overfitting signs.

# 5 Discussion

# 5.1 Why Logistic Regression Outperformed SVM

LR's superior performance (75.65% vs 69.34% accuracy) stems from several factors:

**Theoretical:** (1) LR's smooth cross-entropy loss enables more stable gradient descent than SVM's sharp hinge loss; (2) LR provides calibrated probabilities suitable for overlapping class distributions, while SVM's hard margins may be too rigid; (3) LR's uniform weight penalization differs from SVM's support-vector-focused approach.

**Practical:** LR converged  $3-5 \times$  faster (0.16s vs 4.91s) and showed less sensitivity to hyperparameters. SVM required careful learning rate tuning and exhibited training instability.

# 5.2 Kernel Analysis

**RBF** success: The 82.31% F1-score suggests non-linear relationships exist between features and quality. RBF's local similarity emphasis and smooth boundaries effectively captured wine quality patterns.

**Polynomial failure:** Degree 2 insufficient for complex interactions (63.87% accuracy); degree 3 overfitted (65.79%). Polynomial kernels' global transformation nature ill-suited this dataset's localized patterns.

**Gamma selection:** Lower  $\gamma = 0.1$  (broader influence, smoother boundaries) generalized better than  $\gamma = 0.5$  (tighter influence, overfitting risk).

# 5.3 Model Complexity Assessment

No overfitting: Test accuracy (76.19%) close to CV accuracy (73.84%), training curves plateaued, regularization effective.

No underfitting: 74% training accuracy, RBF kernels captured non-linearity, multiple models converged similarly.

The  $\sim 76\%$  accuracy ceiling likely reflects: (1) inherent noise in subjective quality ratings; (2) missing quality-relevant features; (3) information loss from continuous-to-binary conversion.

# 6 Conclusion

# 6.1 Summary

We successfully implemented SVM and Logistic Regression from scratch with kernel extensions, achieving 82.31% F1-score on wine quality prediction. Rigorous methodology (stratified splitting, proper normalization, cross-validation) ensured reliable results.

### **Key findings:**

- LR superior to SVM (6.3% accuracy gain, faster training)
- RBF kernel most effective for non-linearity
- Lower gamma optimal for generalization
- Polynomial kernels underperformed

### 6.2 Limitations and Future Work

**Limitations:** (1) Kernel SVM computationally expensive (34-145s training); (2) class imbalance addressed only via stratification; (3) binary classification discards fine-grained quality information; (4) limited hyperparameter search space.

## **Future directions:**

- Class-weighted loss functions or SMOTE for imbalance
- Multi-class classification preserving full quality scale
- Feature engineering (domain-specific interactions, PCA)
- Ensemble methods and deep learning
- Bayesian hyperparameter optimization