



## **Model Development Phase Template**

Date	18 June 2024	
Team ID	SWTID1749713922	
Project Title	Early Prediction for Chronic Kidney Disease Detection: A Progressive Approach to Health Management	
Maximum Marks	4 Marks	

## **Initial Model Training Code, Model Validation and Evaluation Report**

The model validation and evaluation report will include classification reports, accuracy, and confusion matrices for multiple models, presented through respective screenshots.

## **Initial Model Training Code:**

```
def run_healthcare_ml_pipeline_hyp(X_train, X_test, y_train, y_test):
    Complete healthcare ML pipeline with enhanced hyperparameter tuning and overfitting detection
    print(f"\n{'='*80}")
    print("HEALTHCARE ML ANALYSIS PIPELINE")
    print("Kidney Disease Detection - Enhanced with Hyperparameter Tuning")
    print(f"{'='*80}")
    # Convert target to binary if needed
    if hasattr(y_train, 'dtype') and y_train.dtype == 'object':
        unique_classes = np.unique(y_train)
        if len(unique_classes) == 2:
            y_train_processed = (y_train == unique_classes[0]).astype(int)
           y_test_processed = (y_test == unique_classes[0]).astype(int)
            from sklearn.preprocessing import LabelEncoder
            le = LabelEncoder()
            y_train_processed = le.fit_transform(y_train)
            y_test_processed = le.transform(y_test)
    else:
       y_train_processed = y_train
       y_test_processed = y_test
```





```
print(f"Dataset Info:")
print(f" Training samples: {X_train.shape[0]}")
print(f" Test samples: {X test.shape[0]}")
print(f" Features: {X_train.shape[1]}")
print(f" Classes: {len(np.unique(y_train_processed))}")
print(f" Class distribution: {dict(zip(*np.unique(y_train_processed, return_counts=True)))}")
# Create hyperparameter-tuned models
models, tuning_summary = create_tuned_models_dict(X_train, y_train_processed)
print(f"\n Testing {len(models)} different classification algorithms...")
print(f"
            {len(tuning summary)} models have been hyperparameter-tuned")
print(f"
            {len(models) - len(tuning_summary)} baseline models included")
# Print hyperparameter tuning summary
if tuning_summary:
    print hyperparameter tuning summary(tuning summary)
# Run comprehensive analysis with enhanced overfitting detection
results_list, cv_results, overfitting_summary = detect_overfitting_comprehensive_enhanced(
    X_train, X_test, y_train_processed, y_test_processed, models
# Convert results list to dictionary for easier access
results_dict = {}
for result in results_list:
    model name = result['Model']
    results_dict[model_name] = result
```

```
print(f"\n{'='*80}")
print("INDIVIDUAL MODEL ANALYSIS WITH VISUALIZATIONS")
print(f"{'='*80}")
# Dictionary to store trained models for plotting
trained_models = {}
# Individual model analysis with plotting
for model_name, model_results in results_dict.items():
    if model_results is None:
       continue
    print(f"\n ANALYZING: {model_name}")
    print("-" * 60)
    try:
        # Get model instance
        if model name not in models:
            print(f" Model {model_name} not found in models dictionary, skipping...")
            continue
        model = models[model name]
        # Train the model
        model.fit(X_train, y_train_processed)
        trained_models[model_name] = model
        # Make predictions
        y_pred = model.predict(X_test)
```





```
# Get prediction probabilities if available
if hasattr(model, 'predict_proba'):
    y_pred_proba = model.predict_proba(X_test)
    if len(np.unique(y_test_processed)) == 2:
        y_pred_proba = y_pred_proba[:, 1]
    else:
        y_pred_proba = y_pred_proba.max(axis=1)
elif hasattr(model, 'decision_function'):
    y_pred_proba = model.decision_function(X_test)
    # Normalize decision function scores to [0,1] for binary classification
    if len(np.unique(y_test_processed)) == 2:
        y_pred_proba = (y_pred_proba - y_pred_proba.min()) / (y_pred_proba.max() - y_pred_proba.min())
else:
    y_pred_proba = None
print(f" Model Performance:")
# Access metrics from model_results dictionary
print(f" Test Accuracy: {model_results.get('Test Accuracy', 0):.4f}")
print(f" Precision: {model_results.get('Precision', 0):.4f}")
           Precision: {model_results.get('Precision', 0):.4f}")
print(f" Recall: {model_results.get('Recall', 0):.4f}")
          F1 Score: {model_results.get('F1 Score', 0):.4f}")
print(f" ROC AUC: {model_results.get('ROC AUC', 0):.4f}")
# Display hyperparameter tuning results if available
if model_name in tuning_summary:
    tuning_info = tuning_summary[model_name]
    print(f"\n Hyperparameter Tuning Results:")
    print(f" Best Recall Score (CV): {tuning_info['results']['best_score']:.4f}")
    print(f" Overfitting Risk: {tuning_info['results']['overfitting_risk']}")
    print(f" Overfitting Gap: {tuning_info['results']['overfitting_gap']:.4f}")
```

```
print(f" Search Method: {tuning_info['results']['search_type']}")
    # Show key hyperparameters
    key_params = list(tuning_info['params'].items())[:3] # Show first 3 params
    if key params:
       print(f" Key Tuned Parameters:")
       for param, value in key params:
           print(f"
                         {param}: {value}")
# Plot confusion matrix
print(f"\n Generating Confusion Matrix...")
plot_confusion_matrix(y_test_processed, y_pred, model_name)
# Plot ROC curve (only for binary classification)
if len(np.unique(y_test_processed)) == 2 and y_pred_proba is not None:
    print(f" Generating ROC Curve...")
    plot_roc_curve(y_test_processed, y_pred_proba, model_name)
    print(f" Generating Precision-Recall Curve...")
    plot_precision_recall_curve(y_test_processed, y_pred_proba, model_name)
# Plot feature importance (if available)
if hasattr(model, 'feature importances '):
    print(f" Generating Feature Importance Plot...")
    plot_feature_importance(model, X_train, model_name)
elif hasattr(model, 'coef_') and model.coef_.ndim == 1:
    print(f" Generating Feature Coefficients Plot...")
    # Handle linear model coefficients
```





```
plt.figure(figsize=(10, 6))
    if hasattr(X train, 'columns'):
        coef_series = pd.Series(np.abs(model.coef_), index=X_train.columns)
        coef_series = pd.Series(np.abs(model.coef_), index=[f'Feature_{i}' for i in range(len(model.coef_))])
    coef_series.sort_values(ascending=False).head(10).plot(kind='bar')
   plt.title(f'Top 10 Feature Coefficients (Absolute) - {model_name}')
    plt.xticks(rotation=45, ha='right')
   plt.tight_layout()
   plt.show()
# Validation curve analysis for selected models with hyperparameters
print(f" Generating Validation Curve Analysis...")
if model name == 'Random Forest':
    validation_curve_analysis_enhanced(
        X_train, y_train_processed, model,
        'n_estimators', [10, 50, 100, 200, 300]
elif model name == 'XGBoost':
    validation_curve_analysis_enhanced(
       X_train, y_train_processed, model,
        'max_depth', [3, 4, 5, 6, 7, 8]
elif model_name == 'LightGBM':
    validation_curve_analysis_enhanced(
        X_train, y_train_processed, model,
        'num_leaves', [10, 20, 30, 40, 50]
elif 'SVM' in model_name:
    validation_curve_analysis_enhanced(
        X_train, y_train_processed, model,
```

```
'C', [0.1, 1, 10, 100, 1000]
        elif 'Logistic Regression' in model name:
            validation curve analysis enhanced(
                X_train, y_train_processed, model,
                'C', [0.01, 0.1, 1, 10, 100]
        print(f"Completed analysis for {model_name}\n")
    except Exception as e:
        print(f" Error analyzing {model name}: {str(e)}")
        continue
    # Print overfitting summary
risk_groups = print_overfitting_summary(overfitting_summary)
# Convert results list to DataFrame for healthcare model selection
results_df = pd.DataFrame(results_list)
# Healthcare-specific model selection
best model name, ranked models = healthcare model selection algorithm(results df)
# Generate comprehensive model comparison plots
print(f"\n{'='*80}")
print(" COMPREHENSIVE MODEL COMPARISON VISUALIZATIONS")
print(f"{'='*80}")
```





```
print(" Generating Top Models Comparison...")
plot_model_comparison(results_df)
# Enhanced hyperparameter tuning summary visualization
if tuning summary:
    print(f"\n Generating Hyperparameter Tuning Summary Visualization...")
    plot_hyperparameter_tuning_summary(tuning_summary)
# Final recommendations with hyperparameter considerations
print(f"\n{'='*80}")
print(" FINAL HEALTHCARE RECOMMENDATIONS")
print(f"{'='*80}")
print(f" RECOMMENDED MODEL: {best_model_name}")
best stats = ranked models.iloc[0]
print(f" Healthcare Score: {best_stats['Healthcare_Score']:.4f}")
print(f" Recall (Sensitivity): {best_stats['Recall']:.4f}")
print(f" Precision: {best_stats['Precision']:.4f}")
print(f" F1 Score: {best_stats['F1 Score']:.4f}")
print(f" Overfitting Risk: {best_stats['Overfitting Risk']}")
# Show hyperparameter tuning info for best model if available
if best_model_name in tuning_summary:
    best_tuning = tuning_summary[best_model_name]
    print(f"
               Hyperparameter Optimization:")
    print(f"
                 Tuning Method: {best_tuning['results']['search_type']}")
    print(f"
                   • CV Recall Score: {best_tuning['results']['best_score']:.4f}")
    print(f"
                   • Overfitting Gap: {best_tuning['results']['overfitting_gap']:.4f}")
```

```
print(f"\n TOP 3 SAFE MODELS FOR HEALTHCARE:")
safe_models = ranked_models[ranked_models['Overfitting Risk'].isin(['LOW', 'MEDIUM'])].head(3)
for i, (_, row) in enumerate(safe models.iterrows(), 1):
    risk_indicator = "" if row['Overfitting Risk'] == 'LOW' else ""
    tuning_indicator = "" if row['Model'] in tuning_summary else ""
              (i). {tuning_indicator} {row['Model']} (Score: {row['Healthcare_Score']:.4f}, Risk: {risk_indicator} {row['Overfitting Risk']})")
# Models to avoid with hyperparameter tuning context
avoid_models = ranked_models[ranked_models[Overfitting Risk'].isin(['CRITICAL', 'HIGH'])]['Model'].tolist()
if avoid models:
    print(f"\n MODELS TO AVOID IN HEALTHCARE:")
    for model in avoid_models[:5]: # Show top 5 to avoid
        if model in tuning_summary:
            gap = tuning_summary[model]['results']['overfitting_gap']
             print(f" {model} (Overfitting Gap: {gap:.4f})")
        else:
            print(f" {model}")
# Hyperparameter tuning insights
if tuning_summary:
    print(f"\n HYPERPARAMETER TUNING INSIGHTS:")
    # Count tuned models by risk level
    tuned_risks = {}
    for model_name, info in tuning_summary.items():
        risk = info['results']['overfitting_risk']
         tuned_risks[risk] = tuned_risks.get(risk, 0) + 1
```





```
print(f" Tuned Models by Risk Level:")
     risk_order = ['LOW', 'MEDIUM', 'HIGH', 'CRITICAL']
     for risk in risk_order:
         if risk in Tuned_risks:
    icon = {'LOW': '\_', 'MEDIUM': '\_', 'HIGH': '\_', 'CRITICAL':
    print(f" {icon} {risk}: {tuned_risks[risk]} models")

     # Best tuning results
     best_tuned_recall = max(tuning_summary.items(), key=lambda x: x[1]['results']['best_score'])
     lowest_overfitting_tuned = min(tuning_summary.items(), key=lambda x: x[1]['results']['overfitting_gap'])
                   Best Tuned Recall: {best_tuned_recall[0]} ({best_tuned_recall[1]['results']['best_score']:.4f})")
                  Lowest Overfitting (Tuned): {lowest_overfitting_tuned[0]} (Gap: {lowest_overfitting_tuned[1]['results']['overfitting_gap']:.4f})")
# Additional comprehensive analysis plots
print(f"\n GENERATING ADDITIONAL ANALYSIS PLOTS...")
# Overfitting risk distribution plot
plt.figure(figsize=(15, 10))
# Risk distribution
plt.subplot(2, 3, 1)
risk_counts = ranked_models['Overfitting Risk'].value_counts()
colors = {'LOW': 'green', 'MEDIUM': 'orange', 'HIGH': 'red', 'CRITICAL': 'darkred'}
risk_colors = [colors.get(risk, 'gray') for risk in risk_counts.index]
plt.pie(risk_counts.values, labels=risk_counts.index, autopct='%1.1f%%', colors=risk_colors)
plt.title('Overfitting Risk Distribution')
```

```
# Healthcare scores distribution
plt.subplot(2, 3, 2)
plt.hist(ranked models['Healthcare Score'], bins=15, alpha=0.7, color='skyblue', edgecolor='black')
plt.xlabel('Healthcare Score')
plt.ylabel('Number of Models')
plt.title('Healthcare Scores Distribution')
plt.grid(True, alpha=0.3)
# Recall vs Precision scatter plot
plt.subplot(2, 3, 3)
colors_risk = ranked_models['Overfitting Risk'].map(colors)
scatter = plt.scatter(ranked_models['Recall'], ranked_models['Precision'],
                     c=colors risk, alpha=0.7, s=60, edgecolors='black', linewidth=0.5)
plt.xlabel('Recall (Sensitivity)')
plt.ylabel('Precision')
plt.title('Recall vs Precision (Colored by Risk)')
plt.grid(True, alpha=0.3)
# F1 Score vs CV Stability
plt.subplot(2, 3, 4)
plt.scatter(ranked_models['F1 Score'], ranked_models['CV Std F1'],
           c=colors_risk, alpha=0.7, s=60, edgecolors='black', linewidth=0.5)
plt.xlabel('F1 Score')
plt.ylabel('CV Standard Deviation')
plt.title('Performance vs Stability (Colored by Risk)')
plt.grid(True, alpha=0.3)
```





```
# Hyperparameter tuning comparison (if available)
if tuning_summary:
   plt.subplot(2, 3, 5)
   tuned_models_data = []
   tuned_scores = []
   tuned_gaps = []
   for model_name in ranked_models['Model']:
        if model name in tuning summary:
           tuned_models_data.append(model_name[:15]) # Truncate long names
            tuned_scores.append(tuning_summary[model_name]['results']['best_score'])
           tuned_gaps.append(tuning_summary[model_name]['results']['overfitting_gap'])
   if tuned models data:
       plt.scatter(tuned_scores, tuned_gaps, alpha=0.7, s=60,
                   c='purple', edgecolors='black', linewidth=0.5)
       plt.xlabel('Tuned CV Recall Score')
       plt.ylabel('Overfitting Gap')
       plt.title('Hyperparameter Tuning Results')
       plt.grid(True, alpha=0.3)
        # Add model names as annotations for top performers
        for i, (score, gap, name) in enumerate(zip(tuned_scores, tuned_gaps, tuned_models_data)):
           if score > np.percentile(tuned_scores, 75) and gap < np.percentile(tuned_gaps, 50):</pre>
               plt.annotate(name, (score, gap), xytext=(5, 5),
                           textcoords='offset points', fontsize=8)
```

```
# Model complexity vs performance
plt.subplot(2, 3, 6)
# Create a complexity score based on model type
complexity map = {
    'Dummy': 1, 'Naive Bayes': 2, 'Logistic Regression': 3, 'LDA': 3, 'QDA': 4,
    'Decision Tree': 4, 'KNN': 4, 'SVM': 5, 'Random Forest': 6, 'Extra Trees': 6,
    'AdaBoost': 6, 'Gradient Boosting': 7, 'XGBoost': 8, 'LightGBM': 8, 'CatBoost': 8,
    'MLP': 9, 'SGD': 3, 'Ridge': 3, 'Bagging': 5
complexity_scores = []
for model name in ranked models['Model']:
    complexity = 5 # default
    for key, value in complexity_map.items():
        if key.lower() in model_name.lower():
            complexity = value
            break
    complexity_scores.append(complexity)
plt.scatter(complexity_scores, ranked_models['Healthcare_Score'],
           c=colors_risk, alpha=0.7, s=60, edgecolors='black', linewidth=0.5)
plt.xlabel('Model Complexity')
plt.ylabel('Healthcare Score')
plt.title('Complexity vs Healthcare Performance')
plt.grid(True, alpha=0.3)
plt.tight layout()
plt.show()
```





```
# Summary statistics
print(f"\n SUMMARY STATISTICS:")
print(f" Total Models Evaluated: {len(results_list)}")
print(f"
           Models with Hyperparameter Tuning: {len(tuning_summary)}")
print(f"
           Low Risk Models: {len(ranked_models[ranked_models['Overfitting Risk'] == 'LOW'])}")
print(f"
           Medium Risk Models: {len(ranked_models[ranked_models['Overfitting Risk'] == 'MEDIUM'])}")
print(f"
           High Risk Models: {len(ranked models[ranked models['Overfitting Risk'] == 'HIGH'])}")
print(f"
           Critical Risk Models: {len(ranked_models[ranked_models['Overfitting Risk'] == 'CRITICAL'])}")
print(f"
            Average Healthcare Score: {ranked_models['Healthcare_Score'].mean():.4f}")
print(f"
           Average Recall: {ranked_models['Recall'].mean():.4f}")
print(f"
           Average Precision: {ranked_models['Precision'].mean():.4f}")
    'results': results dict,
    'results_list': results_list,
    'cv_results': cv_results,
    'overfitting_summary': overfitting_summary,
    'risk_groups': risk_groups,
    'best_model': best_model_name,
    'ranked_models': ranked_models,
    'safe models': safe models,
    'trained_models': trained_models,
    'tuning_summary': tuning_summary,
    'hyperparameter_insights': {
        'tuned_models_count': len(tuning_summary),
        'best_tuned_recall': best_tuned_recall if tuning_summary else None,
        'lowest_overfitting_tuned': lowest_overfitting_tuned if tuning_summary else None,
        'risk_distribution': tuned_risks if tuning_summary else None
```





## **Model Validation and Evaluation Report:**

Model	Classification Report	Accura cy	Confusion Matrix
Rando m Forest	COMPREHENSIVE ANALYSIS: Random Forest	92.50%	Confusion Matrix - Random Forest  - 40 - 35 - 30 - 25 - 20 - 15 - 10 - 5 - 0 Predicted Label
XGBoo st	COMPREHENSIVE ANALYSIS: XGBoost	96.25%	Confusion Matrix - XGBoost  - 40  - 30  - 20  - 10  Predicted Label















































