Gradient Boosted Decision Trees (GBDT)

Tutorial on Implementation and Interpretability in MoDeVa

July 5, 2025

Tutorial Overview:

- GBDT fundamentals and mathematical foundations
- Implementation with XGBoost, LightGBM, and CatBoost
- Interpretability through Functional ANOVA
- Special cases: GAM and GAMI structures
- Monotonicity constraints and domain knowledge
- Global and local interpretation techniques

Outline

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What is Gradient Boosted Decision Trees?

Definition: GBDT is a powerful ensemble learning algorithm that builds a sequence of decision trees, where each subsequent tree is trained to correct the errors of its predecessors.

Key Characteristics:

- Sequential Learning: Trees built one after another
- Error Correction: Each tree learns from previous mistakes
- Gradient-Based: Uses gradients of loss function for optimization
- Versatile: Handles both regression and classification tasks

vs. Random Forests:

- GBDT: Sequential, dependent trees
- Random Forests: Parallel, independent trees

Why GBDT Matters

Advantages:

- **High Accuracy**: Often achieves state-of-the-art performance
- Flexibility: Works well with various data types
- Feature Handling: Robust to missing values and outliers
- Interpretability: Can be made interpretable with constraints

Applications:

- Credit scoring and risk assessment
- Sales forecasting and demand prediction
- Medical diagnosis and treatment planning
- Marketing response modeling

Popular Implementations:

- XGBoost: Performance and speed optimization
- LightGBM: Memory efficiency and fast training
- CatBoost: Categorical feature handling

GBDT Mathematical Framework

Overall Ensemble Model:

$$F_M(x) = F_0(x) + \sum_{m=1}^{M} \gamma_m T_m(x)$$
 (1)

Where:

- $F_0(x)$: Initial model (often constant, e.g., mean for regression)
- $T_m(x)$: Decision tree added at iteration m
- γ_m : Learning rate controlling tree contribution
- M: Total number of trees

Key Insight: Each tree adds a small correction to the ensemble, gradually improving predictions through iterative refinement.

Pseudo-Residual Calculation

Gradient-Based Learning:

At each iteration m, compute pseudo-residuals:

$$r_{im} = -\frac{\partial L(y_i, F_{m-1}(x_i))}{\partial F_{m-1}(x_i)}$$
 (2)

Components:

- y_i : True target value for sample i
- $F_{m-1}(x_i)$: Current model prediction for sample i
- r_{im}: Pseudo-residual (negative gradient) for sample i
- L: Loss function (e.g., squared loss, log-loss)

Intuition: Pseudo-residuals point in the direction of steepest descent, indicating how to adjust predictions to reduce loss.

Model Update Rule

Sequential Update Process:

After fitting tree $h_m(x)$ to pseudo-residuals:

$$F_m(x) = F_{m-1}(x) + \gamma_m h_m(x) \tag{3}$$

Complete GBDT Algorithm:

- **1** Initialize: $F_0(x) = \arg\min_{\gamma} \sum_{i=1}^n L(y_i, \gamma)$
- ② For m = 1 to M:
 - Compute pseudo-residuals: $r_{im} = -\frac{\partial L(y_i, F_{m-1}(x_i))}{\partial F_{m-1}(x_i)}$
 - Fit tree $h_m(x)$ to residuals
 - Find optimal step size: $\gamma_m = \arg\min_{\gamma} \sum_{i=1}^n L(y_i, F_{m-1}(x_i) + \gamma h_m(x_i))$
 - Update model: $F_m(x) = F_{m-1}(x) + \gamma_m h_m(x)$
- **3** Return final model: $F_M(x)$

MoDeVa GBDT Ecosystem

Unified Wrapper Architecture:

MoDeVa provides comprehensive wrappers around leading GBDT implementations:

XGBoost

- Speed optimization
- Advanced regularization
- Cross-validation built-in
- Monotonicity constraints

LightGBM

- Memory efficient
- Fast training speed
- Categorical features
- Large-scale datasets

CatBoost

- Categorical handling
- Overfitting resistant
- No preprocessing
- Symmetric trees

Benefits of Unified Interface:

- Consistent API across implementations
- Easy model comparison and switching
- Integrated interpretability tools
- Unified hyperparameter management

Data Setup and Preprocessing

Preparing Data for GBDT:

```
1 from modeva import DataSet
3 # Create dataset object
4 ds = DataSet()
5 # Load built-in dataset
6 ds.load(name="BikeSharing")
8 # Preprocessing pipeline
9 ds.scale_numerical(features=("cnt",), method="log1p")
ds.set_feature_type(feature="hr",
                     feature_type="categorical")
ds.set_feature_type(feature="mnth".
                     feature_type="categorical")
14 ds.scale_numerical(features=ds.feature_names_numerical,
                    method="standardize")
16 ds.set_inactive_features(features=("yr", "season", "temp"))
ds.preprocess()
19 # Split data
ds.set_random_split(test_ratio=0.2, random_state=42)
```

Model Configuration

Setting Up GBDT Models:

```
Regression Models:
```

```
from modeva.models import
                             (MoLGBMRegressor,
                             MoXGBRegressor,
                             MoCatBoostRegressor)
    LightGBM
6 model_lgbm = MoLGBMRegressor(name="LGBM_model",
                               max_depth=2,
                               n_{estimators} = 100
10 # XGBoost
11 model_xgb = MoXGBRegressor(name="XGB_model",
                             max_depth=2,
                             n_estimators=100)
14
15 # CatBoost
16 model_cat = MoCatBoostRegressor(name="CBoost_model".
                                  max_depth=2,
                                  n_estimators=100)
```

Model Configuration

Classification Models:

Model Training and Evaluation

Training Process:

```
# Train the model
model_gbdt.fit(ds.train_x, ds.train_y)

# Make predictions
train_pred = model_gbdt.predict(ds.train_x)
test_pred = model_gbdt.predict(ds.test_x)
```

Performance Assessment:

```
from modeva import TestSuite
from modeva
```

Model Training and Evaluation

Performance Metrics Include:

- Regression: RMSE, MAE, R², MAPE
- Classification: Accuracy, Precision, Recall, F1, AUC
- Cross-validation scores
- Training vs. validation performance

Making GBDT Interpretable

The Challenge: GBDT models are typically considered "black boxes"

The Solution: Functional ANOVA Decomposition

Mathematical Framework:

$$f(x) = \mu + \sum_{j} f_j(x_j) + \sum_{j < k} f_{jk}(x_j, x_k) + \dots$$
 (4)

Components:

- μ : Global intercept (average prediction)
- $f_i(x_i)$: Main effects (individual feature contributions)
- $f_{jk}(x_j, x_k)$: Pairwise interaction effects
- Higher-order terms: Complex interactions (limited by tree depth)

Key Insight: Deep trees create complex interactions; shallow trees enable clearer interpretation.

Special Cases: GAM and GAMI Structures

Depth-1 Trees: GAM Structure

$$f(x) = \mu + \sum_{j} f_{j}(x_{j}) \tag{5}$$

- Each tree makes single split: Results in Generalized Additive Model
- Only main effects, no interactions
- Highly interpretable but limited flexibility

Depth-2 Trees: GAMI Structure

$$f(x) = \mu + \sum_{j} f_j(x_j) + \sum_{j < k} f_{jk}(x_j, x_k)$$

- Each tree makes up to two splits: GAM with Interactions (GAMI)
- Balances interpretability with interaction modeling
- Most practical choice for interpretable GBDT

(6)

ANOVA Decomposition Process

Two-Stage Process:

1. Aggregation Stage:

- Start with tree ensemble: $f(x) = \sum_k \eta_k T_k(x)$
- Decompose into leaf nodes:

$$f(x) = \sum_{m} v_{m} \prod_{j \in S_{m}} I(s_{mj}^{l} \le x_{j} < s_{mj}^{u})$$

- Assign effects based on split variables:
 - Main effects: 1 split variable
 - Pairwise interactions: 2 split variables
 - Higher-order: More split variables

2. Purification Stage:

- Ensure orthogonality: $\int f_{i_1...i_t} dx_k = 0$
- Remove identifiability issues
- Center effects to have zero mean
- Cascade from high-order to low-order interactions

Feature Importance Analysis

Understanding Overall Feature Impact:

```
# Global feature importance
result = ts.interpret_fi()
result.plot()
```

Importance Metrics:

- Based on variance of marginal effects
- Normalized to sum to 1
- Accounts for feature scale differences
- Shows relative importance across features

Interpretation:

- Higher values indicate stronger influence on predictions
- Helps identify key drivers in the model
- Supports feature selection decisions
- Guides business focus areas

Effect Importance Analysis

ANOVA Component Analysis:

```
# Global effect importance
result = ts.interpret_ei()
result.plot()
```

Effect Types:

- Main effects: Individual feature contributions
- Interaction effects: Pairwise feature interactions
- **Higher-order**: Complex multi-feature interactions

Applications:

- Understand model complexity
- Identify significant interactions
- Validate domain knowledge
- Guide feature engineering

Effect Visualization

Main Effect Plots:

```
# Visualize main effect for specific feature
result = ts.interpret_effects(features="hr")
result.plot()
```

Plot Interpretation:

- Shows how feature affects predictions
- Reveals non-linear relationships
- Indicates feature importance regions
- Supports business understanding

Interaction Effect Plots:

```
# Visualize pairwise interaction
result = ts.interpret_effects(features=("temp", "hum"))
result.plot()
```

Individual Prediction Analysis

Local Feature Importance:

Local Effect Importance:

```
# ANOVA-based local explanation
result = ts.interpret_local_ei(sample_index=10,
centered=True)
result.plot()
```

Local Explanation Components:

- Feature/effect contributions to specific prediction sample
- Comparison to average behavior
- Direction and magnitude of effects

Introduction to Monotonicity Constraints

What are Monotonicity Constraints?

Enforce that certain feature-response relationships follow domain knowledge:

Common Examples:

- Credit Scoring: Score should increase with income
- Risk Assessment: Risk should decrease with credit rating
- Real Estate: Property value should increase with square footage
- Marketing: Response rate should increase with previous purchases

Mathematical Constraint: For monotonic increasing feature x_j :

$$\frac{\partial f(x)}{\partial x_j} \ge 0 \quad \forall x \tag{7}$$

Implementation: Available in XGBoost through MoDeVa's MoXGBRegressor and MoXGBClassifier

Benefits of Monotonicity Constraints

Interpretability Enhancement:

- Smoother Functions: Reduces noise and fluctuations
- Clearer Patterns: More interpretable global trends
- Consistent SHAP: More stable local explanations
- Business Logic: Aligned with domain knowledge

Model Quality Improvements:

- Reduced Overfitting: Constraints act as regularization
- Better Generalization: More robust to new data
- Stable Performance: Often maintains or improves accuracy

Interaction with ANOVA:

- Guarantees monotonic shape for constrained features
- Preserves interpretability in decomposition
- Maintains partial monotonicity in interactions
- Simplifies effect visualization

Model Selection Guidelines

Choosing the Right GBDT Implementation:

Use XGBoost when:

- Need monotonicity constraints
- Want extensive hyperparameter control
- Performance is critical
- Working with structured data

Use LightGBM when:

- Large datasets (100K+ samples)
- Memory is limited
- Training speed matters
- Categorical features present

Use CatBoost when:

- Many categorical features
- Minimal preprocessing desired
- Overfitting is a concern
- New to GBDT

Tree Depth Selection:

- **Depth 1**: Maximum interpretability (GAM)
- **Depth 2**: Good balance (GAMI) recommended
- **Depth 3+**: Higher performance, reduced interpretability

Hyperparameter Tuning Strategy

Key Parameters to Optimize:

1. Tree Structure:

- max_depth: Start with 2, increase if needed
- n_estimators: Start with 100, use early stopping
- min_child_weight: Increase to prevent overfitting

2. Learning Control:

- learning_rate: Start with 0.1, decrease for more trees
- subsample: Try 0.8-1.0 for regularization
- colsample_bytree: Use 0.8-1.0 for feature sampling

3. Tuning Process:

- Fix depth at 2 for interpretability
- 2 Tune number of estimators with early stopping
- Optimize learning rate and regularization
- Apply monotonicity constraints if needed

Key Takeaways

MoDeVa Integration:

- Unified interface across XGBoost, LightGBM, and CatBoost
- Integrated interpretability through Functional ANOVA
- Easy model comparison and deployment

Interpretability:

- Depth constraints enable GAM/GAMI structures
- Global and local explanation methods
- Monotonicity constraints for domain knowledge integration

Best Practices:

- Start with depth-2 trees for interpretability
- Use early stopping and cross-validation
- Apply monotonicity constraints when appropriate
- Validate interpretations with domain experts