

# stable\_tree\_bertsimas

April 17, 2025

## 0.1 Notebook implementing the algorithm in Bertsimas et al. (<https://arxiv.org/abs/2305.17299>)

```
[1]: import numpy as np
import itertools
from sklearn.model_selection import train_test_split
from sklearn.tree import DecisionTreeClassifier
from sklearn.datasets import load_breast_cancer
from sklearn.metrics import accuracy_score
```

### 0.1.1 Import dataset and split

```
[2]: data_breast_cancer = load_breast_cancer(as_frame=True)
X_full = data_breast_cancer["data"]
y_full = data_breast_cancer["target"]
```

```
[3]: print("X_full shape: ", X_full.shape)
X_full.head()
```

X\_full shape: (569, 30)

```
[3]:
```

	mean radius	mean texture	mean perimeter	mean area	mean smoothness	\
0	17.99	10.38	122.80	1001.0	0.11840	
1	20.57	17.77	132.90	1326.0	0.08474	
2	19.69	21.25	130.00	1203.0	0.10960	
3	11.42	20.38	77.58	386.1	0.14250	
4	20.29	14.34	135.10	1297.0	0.10030	

  

	mean compactness	mean concavity	mean concave points	mean symmetry	\
0	0.27760	0.3001	0.14710	0.2419	
1	0.07864	0.0869	0.07017	0.1812	
2	0.15990	0.1974	0.12790	0.2069	
3	0.28390	0.2414	0.10520	0.2597	
4	0.13280	0.1980	0.10430	0.1809	

  

	mean fractal dimension	...	worst radius	worst texture	worst perimeter	\
0	0.07871	...	25.38	17.33	184.60	
1	0.05667	...	24.99	23.41	158.80	

2	0.05999	...	23.57	25.53	152.50
3	0.09744	...	14.91	26.50	98.87
4	0.05883	...	22.54	16.67	152.20

	worst area	worst smoothness	worst compactness	worst concavity	\
0	2019.0	0.1622	0.6656	0.7119	
1	1956.0	0.1238	0.1866	0.2416	
2	1709.0	0.1444	0.4245	0.4504	
3	567.7	0.2098	0.8663	0.6869	
4	1575.0	0.1374	0.2050	0.4000	

	worst concave points	worst symmetry	worst fractal dimension
0	0.2654	0.4601	0.11890
1	0.1860	0.2750	0.08902
2	0.2430	0.3613	0.08758
3	0.2575	0.6638	0.17300
4	0.1625	0.2364	0.07678

[5 rows x 30 columns]

```
[4]: print("y_full shape: ", y_full.shape)
      y_full.head()
```

y\_full shape: (569,)

```
[4]: 0    0
      1    0
      2    0
      3    0
      4    0
      Name: target, dtype: int64
```

```
[5]: X_train, X_test, y_train, y_test = train_test_split(X_full, y_full, test_size=0.
      ↪2, random_state=42)

      print("X_train shape: {}, X_test shape: {}".format(X_train.shape, X_test.shape))
      print("y_train shape: {}, y_test shape: {}".format(y_train.shape, y_test.shape))
```

X\_train shape: (455, 30), X\_test shape: (114, 30)  
y\_train shape: (455,), y\_test shape: (114,)

### 0.1.2 Generate first collection (T0) of trees (trained on X0)

```
[6]: def train_trees(X, y, depths=[3,5,7], min_samples=[5,10]):
      """Train multiple trees for different hyperparams & possibly bootstrap."""
      trees = []
      for depth, min_leaf in itertools.product(depths, min_samples):
          # need to bootstrap later
```

```

        clf = DecisionTreeClassifier(
            max_depth=depth,
            min_samples_leaf=min_leaf,
            random_state=42
        )
        clf.fit(X, y)
        trees.append(clf)
    return trees

T0 = train_trees(X_train, y_train)
print("Generated {} trees for T0".format(len(T0)))

```

Generated 6 trees for T0

### 0.1.3 Generate second collection of trees (T) (trained on full data)

```
[7]: T = train_trees(X_full, y_full)
```

### 0.1.4 Get global ranges of numerical features and their names

(todo: categorical features)

```
[8]: feature_names = X_full.columns
    global_lower = X_full.min().values
    global_upper = X_full.max().values

```

### 0.1.5 Compute average distance of each tree in T to the T0 collection

```
[9]: from bertsimas_stable.Paths import tree_distance
```

Distances to T0

```
[10]: distances = []
    max_depth = 7 # largest in our hyperparameter search
    for i, tree_b in enumerate(T):
        # average distance to all trees in T0
        d_b = 0.0
        for tree_beta in T0:
            d_b += tree_distance(
                tree_beta, tree_b,
                global_lower=global_lower,
                global_upper=global_upper,
                lambda_val=2*max_depth
            )
        d_b /= len(T0)
        distances.append(d_b)
    distances

```

```
[10]: [np.float64(9.941638853041566),
      np.float64(9.726568660248738),
      np.float64(9.34090448178685),
      np.float64(9.872533253986653),
      np.float64(9.716180344437472),
      np.float64(9.824395104992929)]
```

### 0.1.6 Distances to T0 (using given method)

```
[11]: from dt_distance_repo.dt_distance.distance_calculator import DistanceCalculator
      X_train = X_train.values
      X_test = X_test.values
      distances_method2 = []
      for i, tree_b in enumerate(T):
          # average distance to all trees in T0
          d_b = 0.0
          for tree_beta in T0:
              distance_calculator = DistanceCalculator(tree_beta, tree_b, X=X_train,
          ↪ y=y_train)
              d_b += distance_calculator.compute_tree_distance()
          d_b /= len(T0)
          distances_method2.append(d_b)
      distances_method2
```

```
[11]: [np.float64(0.05715),
      np.float64(0.05145),
      np.float64(0.038016666666666664),
      np.float64(0.026416666666666668),
      np.float64(0.020283333333333334),
      np.float64(0.014333333333333335)]
```

### 0.1.7 Compute predictive performance for each tree

```
[12]: performances = []
      for i, tree_b in enumerate(T):
          y_pred = tree_b.predict(X_test)
          acc = accuracy_score(y_test, y_pred)
          performances.append(acc)

      print("Accuracies on test set: {%s}" % ", ".join("%.3f" % a for a in
          ↪ performances))
```

Accuracies on test set: {0.974, 0.965, 0.991, 0.965, 0.991, 0.956}

```
/Users/adb/stuff/gitclones/Suicide_Project/.venv/lib/python3.11/site-
packages/sklearn/utils/validation.py:2739: UserWarning: X does not have valid
feature names, but DecisionTreeClassifier was fitted with feature names
  warnings.warn(
```

```

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  warnings.warn(

```

### 0.1.8 Identify Pareto frontier

```

[13]: pairs = list(zip(distances, performances)) # (distance, performance)

def is_dominated(i, pairs):
    di, pi = pairs[i]
    for j, (dj, pj) in enumerate(pairs):
        if j != i:
            # Condition for i is dominated by j: dj <= di and pj >= pi
            # with at least one strict inequality
            if (dj <= di and pj >= pi) and (dj < di or pj > pi):
                return True
    return False

pareto_indices = [i for i in range(len(pairs)) if not is_dominated(i, pairs)]
pareto_trees = [T[i] for i in pareto_indices]

print("Pareto indices:", pareto_indices)
print("Number of Pareto-optimal trees:", len(pareto_trees))

```

```

Pareto indices: [2]
Number of Pareto-optimal trees: 1

```

## 0.2 Choose the “best” stable tree from the Pareto set

```
[14]: distance_threshold = min(distances) + 0.2 * (max(distances) - min(distances))
candidate_indices = [i for i in pareto_indices if distances[i] <=
    ↪ distance_threshold]
if candidate_indices:
    best_idx = max(candidate_indices, key=lambda i: performances[i])
    stable_tree = T[best_idx]
    print(f"Chosen stable tree index = {best_idx}, dist={distances[best_idx]},
    ↪ perf={performances[best_idx]}")
else:
    print("No tree satisfies the distance threshold; picking best accuracy from
    ↪ all T.")
    best_idx = np.argmax(performances)
    stable_tree = T[best_idx]
    print(f"Chosen best accuracy tree index = {best_idx},
    ↪ dist={distances[best_idx]}, perf={performances[best_idx]}")
```

Chosen stable tree index = 2, dist=9.34090448178685, perf=0.9912280701754386