

piecewise_linear_regression_criterion

February 28, 2023

1 Custom DecisionTreeRegressor adapted to a linear regression

A `DecisionTreeRegressor` can be trained with a couple of possible criterions but it is possible to implement a custom one (see [hellinger_distance_criterion](#)). See also tutorial [Cython example of exposing C-computed arrays in Python without data copies](#) which describes a way to implement fast `cython` extensions.

```
[1]: from jyquickhelper import add_notebook_menu
      add_notebook_menu()
```

```
[1]: <IPython.core.display.HTML object>
```

```
[2]: %matplotlib inline
```

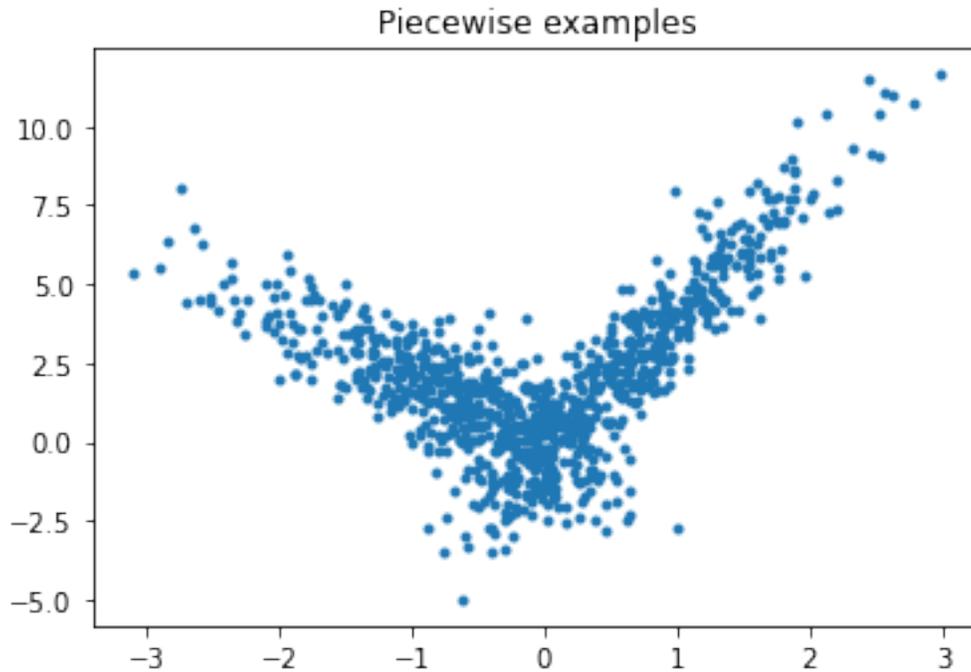
```
[3]: import warnings
      warnings.simplefilter("ignore")
```

1.1 Piecewise data

Let's build a toy problem based on two linear models.

```
[4]: import numpy
      import numpy.random as npr
      X = npr.normal(size=(1000,4))
      alpha = [4, -2]
      t = (X[:, 0] + X[:, 3] * 0.5) > 0
      switch = numpy.zeros(X.shape[0])
      switch[t] = 1
      y = alpha[0] * X[:, 0] * t + alpha[1] * X[:, 0] * (1-t) + X[:, 2]
```

```
[5]: import matplotlib.pyplot as plt
      fig, ax = plt.subplots(1, 1)
      ax.plot(X[:, 0], y, ".")
      ax.set_title("Piecewise examples");
```



1.2 DecisionTreeRegressor

```
[6]: from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(X[:, :1], y)
```

```
[7]: from sklearn.tree import DecisionTreeRegressor

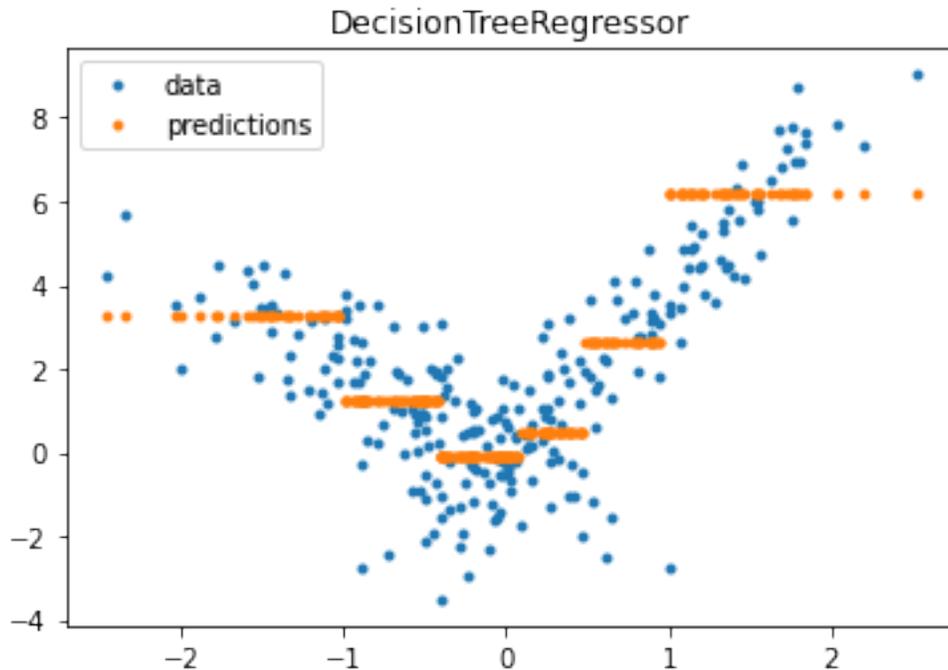
model = DecisionTreeRegressor(min_samples_leaf=100)
model.fit(X_train, y_train)
```

```
[7]: DecisionTreeRegressor(min_samples_leaf=100)
```

```
[8]: pred = model.predict(X_test)
pred[:5]
```

```
[8]: array([ 3.29436256,  0.50924806, -0.07129149,  0.50924806,  2.64957806])
```

```
[9]: fig, ax = plt.subplots(1, 1)
ax.plot(X_test[:, 0], y_test, ".", label='data')
ax.plot(X_test[:, 0], pred, ".", label="predictions")
ax.set_title("DecisionTreeRegressor")
ax.legend();
```



1.3 DecisionTreeRegressor with custom implementation

```
[10]: import sklearn
from pyquickhelper.texthelper import compare_module_version
if compare_module_version(sklearn.__version__, '0.21') < 0:
    print("Next step requires scikit-learn >= 0.21")
else:
    print("sklearn.__version__ =", sklearn.__version__)
```

```
sklearn.__version__ = 1.1.dev0
```

```
[11]: from mlinsights.mlmodel.piecewise_tree_regression_criterion import SimpleRegressorCriterion
```

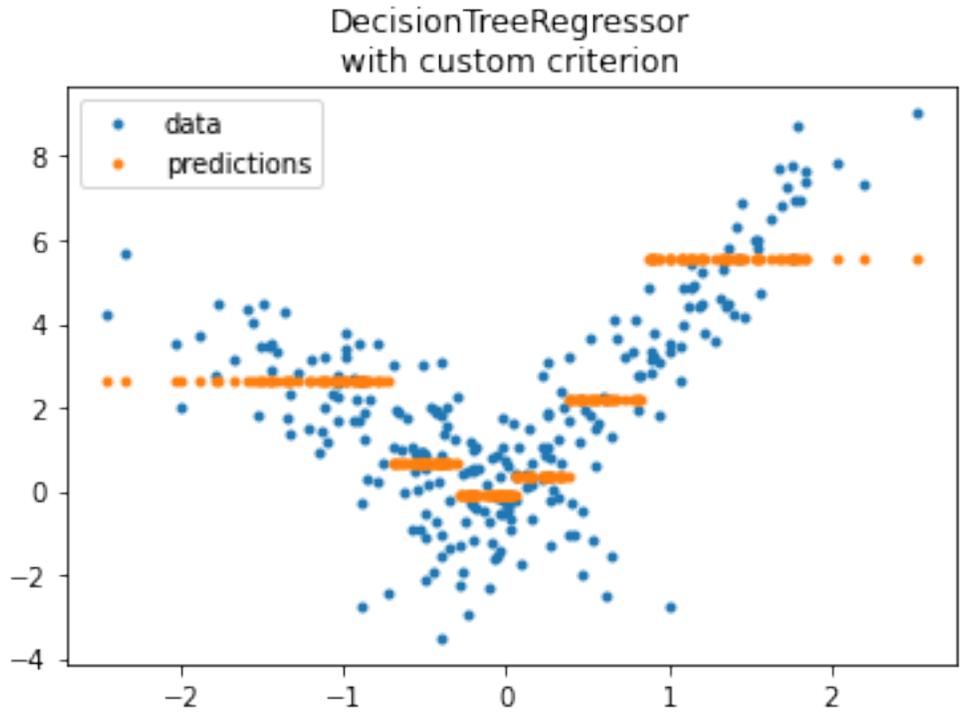
```
[12]: model2 = DecisionTreeRegressor(min_samples_leaf=100,
                                     criterion=SimpleRegressorCriterion(X_train))
model2.fit(X_train, y_train)
```

```
[12]: DecisionTreeRegressor(criterion=<mlinsights.mlmodel.piecewise_tree_regression_criterion.SimpleRegressorCriterion object at 0x000001FC680A0BF0>,
                             min_samples_leaf=100)
```

```
[13]: pred = model2.predict(X_test)
pred[:5]
```

```
[13]: array([ 2.65757699,  0.37665413, -0.07967816,  0.37665413,  5.57229226])
```

```
[14]: fig, ax = plt.subplots(1, 1)
ax.plot(X_test[:, 0], y_test, ".", label='data')
ax.plot(X_test[:, 0], pred, ".", label="predictions")
ax.set_title("DecisionTreeRegressor\nwith custom criterion")
ax.legend();
```



1.4 Computation time

The custom criterion is not really efficient but it was meant that way. The code can be found in [piecewise_tree_regression_criterion](#). Bascially, it is slow because each time the algorithm optimizing the tree needs the class Criterion to evaluate the impurity reduction for a split, the computation happens on the whole data under the node being split. The implementation in [_criterion.pyx](#) does it once.

```
[15]: %timeit model.fit(X_train, y_train)
```

551 μ s \pm 93.7 μ s per loop (mean \pm std. dev. of 7 runs, 1000 loops each)

```
[16]: %timeit model2.fit(X_train, y_train)
```

43.9 ms \pm 2.21 ms per loop (mean \pm std. dev. of 7 runs, 10 loops each)

A loop is involved every time the criterion of the node is involved which raises a the computation cost of lot. The method `_mse` is called each time the algorithm training the decision tree needs to evaluate a cut, one cut involves elements between, position `[start, end]`.

```

cdef void _mean(self, SIZE_t start, SIZE_t end, DOUBLE_t *mean, DOUBLE_t *weight) nogil:
    if start == end:
        mean[0] = 0.
        return
    cdef DOUBLE_t m = 0.
    cdef DOUBLE_t w = 0.
    cdef int k
    for k in range(start, end):
        m += self.sample_wy[k]
        w += self.sample_w[k]
    weight[0] = w
    mean[0] = 0. if w == 0. else m / w

cdef double _mse(self, SIZE_t start, SIZE_t end, DOUBLE_t mean, DOUBLE_t weight) nogil:
    if start == end:
        return 0.
    cdef DOUBLE_t squ = 0.
    cdef int k
    for k in range(start, end):
        squ += (self.y[self.sample_i[k], 0] - mean) ** 2 * self.sample_w[k]
    return 0. if weight == 0. else squ / weight

```

1.5 Better implementation

I rewrote my first implementation to be closer to what *scikit-learn* is doing. The criterion is computed once for all possible cut and then retrieved on demand. The code is below, arrays `sample_wy_left` is the cumulated sum of $weight * Y$ starting from the left side (lower Y). The loop disappeared.

```

cdef void _mean(self, SIZE_t start, SIZE_t end, DOUBLE_t *mean, DOUBLE_t *weight) nogil:
    if start == end:
        mean[0] = 0.
        return
    cdef DOUBLE_t m = self.sample_wy_left[end-1] - (self.sample_wy_left[start-1] if start > 0 else 0)
    cdef DOUBLE_t w = self.sample_w_left[end-1] - (self.sample_w_left[start-1] if start > 0 else 0)
    weight[0] = w
    mean[0] = 0. if w == 0. else m / w

cdef double _mse(self, SIZE_t start, SIZE_t end, DOUBLE_t mean, DOUBLE_t weight) nogil:
    if start == end:
        return 0.
    cdef DOUBLE_t squ = self.sample_wy2_left[end-1] - (self.sample_wy2_left[start-1] if start > 0 else 0)
    # This formula only holds if mean is computed on the same interval.
    # Otherwise, it is squ / weight - true_mean ** 2 + (mean - true_mean) ** 2.
    return 0. if weight == 0. else squ / weight - mean ** 2

```

```

[17]: from mlinsights.mlmodel.piecewise_tree_regression_criterion_fast import SimpleRegressorCriterionFast
      SimpleRegressorCriterionFast
      model3 = DecisionTreeRegressor(min_samples_leaf=100,
      criterion=SimpleRegressorCriterionFast(X_train))
      model3.fit(X_train, y_train)
      pred = model3.predict(X_test)
      pred[:5]

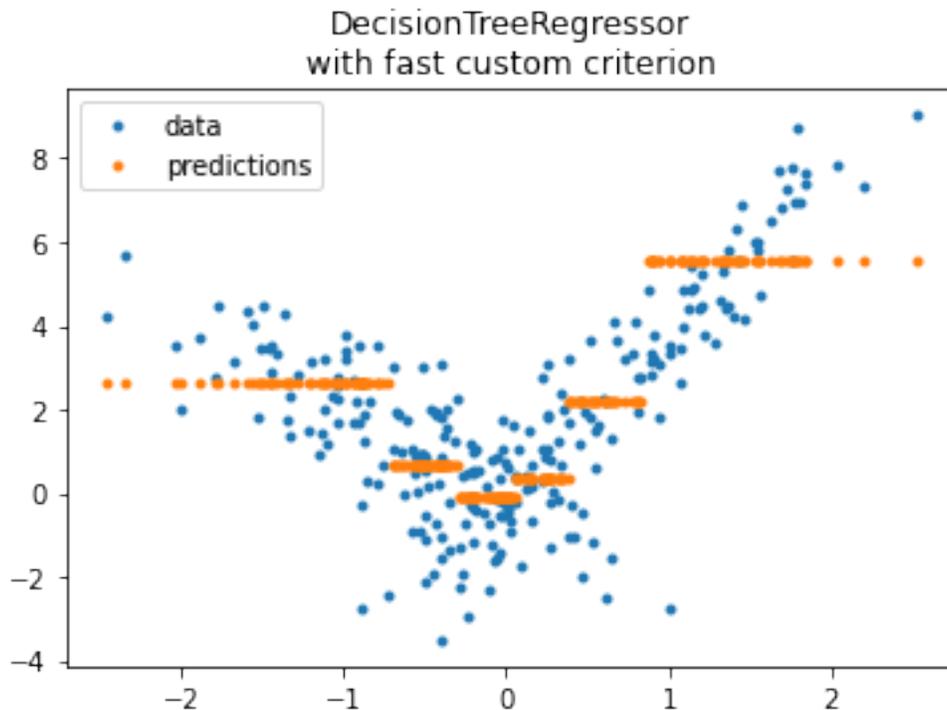
```

```

[17]: array([ 2.65757699,  0.37665413, -0.07967816,  0.37665413,  5.57229226])

```

```
[18]: fig, ax = plt.subplots(1, 1)
ax.plot(X_test[:, 0], y_test, ".", label='data')
ax.plot(X_test[:, 0], pred, ".", label="predictions")
ax.set_title("DecisionTreeRegressor\nwith fast custom criterion")
ax.legend();
```



```
[19]: %timeit model3.fit(X_train, y_train)
```

676 μ s \pm 48.8 μ s per loop (mean \pm std. dev. of 7 runs, 1000 loops each)

Much better even though this implementation is currently 3, 4 times slower than scikit-learn's. Let's check with a datasets three times bigger to see if it is a fix cost or a cost.

```
[20]: import numpy
X_train3 = numpy.vstack([X_train, X_train, X_train])
y_train3 = numpy.hstack([y_train, y_train, y_train])
```

```
[21]: X_train.shape, X_train3.shape, y_train3.shape
```

```
[21]: ((750, 1), (2250, 1), (2250,))
```

```
[22]: %timeit model.fit(X_train3, y_train3)
```

1.36 ms \pm 57 μ s per loop (mean \pm std. dev. of 7 runs, 1000 loops each)

The criterion needs to be reinstanciated since it depends on the features X . The computation does not but the design does. This was introduced to compare the current output with a decision tree optimizing for a piecewise linear regression and not a stepwise regression.

```
[23]: try:
      model3.fit(X_train3, y_train3)
      except Exception as e:
          print(e)
```

X.shape=[750, 1, 0, 0, 0, 0, 0, 0] -- y.shape=[2250, 1, 0, 0, 0, 0, 0, 0]

```
[24]: model3 = DecisionTreeRegressor(min_samples_leaf=100,
                                   criterion=SimpleRegressorCriterionFast(X_train3))
      %timeit model3.fit(X_train3, y_train3)
```

2.03 ms ± 159 µs per loop (mean ± std. dev. of 7 runs, 100 loops each)

Still almost 2 times slower but of the same order of magnitude. We could go further and investigate why or continue and introduce a criterion which optimizes a piecewise linear regression instead of a stepwise regression.

1.6 Criterion adapted for a linear regression

The previous examples are all about decision trees which approximates a function by a stepwise function. On every interval $[r_1, r_2]$, the model optimizes $\sum_i (y_i - C)^2 \mathbf{1}_{r_1 \leq x_i \leq r_2}$ and finds the best constant (= the average) approximating the function on this interval. We would like to approximate the function by a regression line and not a constant anymore. It means minimizing $\sum_i (y_i - X_i \beta)^2 \mathbf{1}_{r_1 \leq x_i \leq r_2}$. Doing this require to change the criterion used to split the space of feature into buckets and the prediction function of the decision tree which now needs to return a dot product.

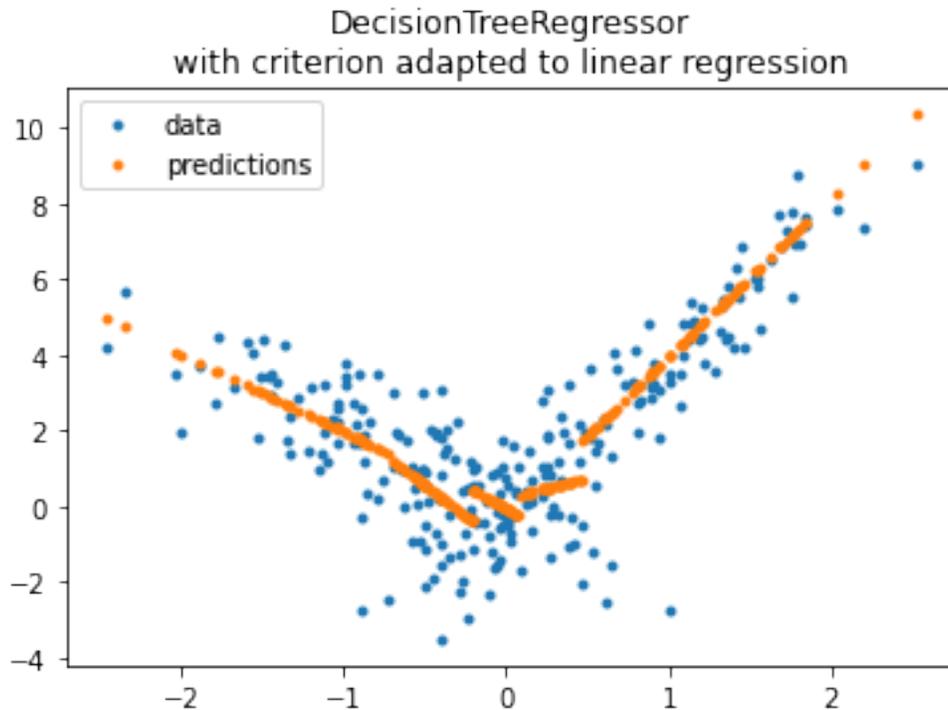
```
[25]: from mlinsights.mlmodel.piecewise_tree_regression import PiecewiseTreeRegressor
      piece = PiecewiseTreeRegressor(criterion='mselin', min_samples_leaf=100)
      piece.fit(X_train, y_train)
```

```
[25]: PiecewiseTreeRegressor(min_samples_leaf=100)
```

```
[26]: pred = piece.predict(X_test)
      pred[:5]
```

```
[26]: array([3.80559618, 0.45086204, 0.42563158, 0.44940565, 3.57423704])
```

```
[27]: fig, ax = plt.subplots(1, 1)
      ax.plot(X_test[:, 0], y_test, ".", label='data')
      ax.plot(X_test[:, 0], pred, ".", label="predictions")
      ax.set_title("DecisionTreeRegressor\nwith criterion adapted to linear regression")
      ax.legend();
```



The coefficients for the linear regressions are kept into the following attribute:

```
[28]: piece.betas_
```

```
[28]: array([[ -2.05163528, -0.07590713],
            [-3.28097357, -1.07747849],
            [-2.37373495, -0.05126022],
            [ 1.10248486,  0.20358196],
            [ 4.21766561, -0.2568136 ]])
```

Mapped to the following leaves:

```
[29]: piece.leaves_index_, piece.leaves_mapping_
```

```
[29]: ([1, 4, 5, 7, 8], {1: 745, 4: 746, 3: 748, 0: 749, 2: 739})
```

We can get the leaf each observation falls into:

```
[30]: piece.predict_leaves(X_test)[:5]
```

```
[30]: array([0, 3, 2, 3, 4])
```

The training is quite slow as it is training many linear regression each time a split is evaluated.

```
[31]: %timeit piece.fit(X_train, y_train)
```

26.9 ms ± 1.98 ms per loop (mean ± std. dev. of 7 runs, 10 loops each)

```
[32]: %timeit piece.fit(X_train3, y_train3)
```

105 ms ± 5.26 ms per loop (mean ± std. dev. of 7 runs, 10 loops each)

It works but it is slow, slower than the slow implementation of the standard criterion for decision tree regression.

1.7 Next

PR [Model trees \(M5P and co\)](#) and issue [Model trees \(M5P\)](#) propose an implementation a piecewise regression with any kind of regression model. It is based on [Building Model Trees](#). It fits many models to find the best splits and should be slower than this implementation in the case of a decision tree regressor associated with linear regressions.

[33] :