Muestra para cada dataset todo lo que se puede hacer con el none_box

In [1]:

```
from demo_utils.demo10 import Demo10
from demo_utils.general import SUPPORTED_DATASETS
from IPython.display import Markdown as md
from demo_utils.get_hyper_params import get_hyper_params
```

In [2]:

```
%%javascript
IPython.OutputArea.prototype._should_scroll = function(lines) {
    return false;
}
```

In [3]:

```
import warnings
warnings.filterwarnings('ignore')
```

In [4]:

```
# d10 data = {
         'dts_name': testing_dataset,
#
#
       'dts size': 1000,
#
       'features_range': (500, 501),
#
       'rbfsampler gamma': 'UNUSED',
       'nystroem gamma': 'UNUSED',
#
       'hparams': {'dt': {'max_depth': None,
#
#
                            'min samples split': 2,
                            'min_samples_leaf': 1,
#
#
                            'min weight fraction leaf': 0.0,
#
                            'max leaf nodes': None,
#
                            'min impurity decrease': 0.0},
                    'logit': \{'\overline{C}': 1000.\overline{0}\},
#
#
                    'linear svc': {'C': 5}}
#
  }
```

In [5]:

```
d10_data = {
#    'dts_name': testing_dataset,
    'dts_size': 1000,
    'features_range': (500, 501),
}
```

In [6]:

```
def get a model(model name, sampler name, dts name):
    box_type = 'black_bag'
    n = 50
    # más adelante habrá que soportar distintas box
          {'model name': model name,
    #
           'sampler name': 'identity',
    #
           'sampler_gamma': None,
    #
           'model_params': {},
    #
           # 'box type': 'none',
    #
           'box type': box type,
    #
           'n estim': None,
    #
           'pca': False,
           'pca first': False}
    ret dic = {'model name': model name,
#
                 'sampler name': 'identity',
               'sampler name': sampler name,
               'sampler gamma': None,
               'model_params': {},
               'box type': box type,
               'n estim': n estim,
               'pca': False,
               'pca first': False}
    hyper params = get hyper params(dts name=dts name, box name=box type,
                                     model name=model name, sampler name=sampler
name)
    gamma = hyper params.pop('qamma', None)
#
      ret dic['sampler gamma'] = gamma
    ret_dic['gamma'] = gamma
#
      ret dic['model params'] = hyper params
    ret dic['base model params'] = hyper params
    if sampler name == 'rff':
        ret dic['sampler name'] = 'rbf'
      elif sampler name == 'nystroem':
#
          ret dic['sampler name'] = 'nystroem'
#
    return ret dic
```

In [7]:

```
def test dt(d10 data):
    d10 = Demo10()
    new data = dict(d10 data)
    dts name = new data['dts name']
    model name = 'dt'
    # dt solo, dt con rff y dt con nystroem
    m1 = get a model(model name=model name, sampler name='identity', dts name=dt
s name)
    m2 = get a model(model name=model name, sampler name='rff', dts name=dts nam
e)
    m3 = get a model(model name=model name, sampler name='nystroem', dts name=dt
s name)
    models = [m1, m2, m3,]
    new data['models'] = models
    d10.non interactive(**new data)
def test_logit(d10_data):
    d10 = Demo10()
    new data = dict(d10 data)
    dts name = new_data['dts_name']
    model name = 'logit'
    # logit solo, logit con rff y logit con nystroem
    m1 = get a model(model name=model name, sampler name='identity', dts name=dt
s name)
    m2 = get a model(model name=model name, sampler name='rff', dts name=dts nam
e)
    m3 = get a model(model name=model name, sampler name='nystroem', dts name=dt
s name)
    models = [m1, m2, m3,]
    new data['models'] = models
    d10.non interactive(**new data)
def test_linear_svc(d10_data):
    d10 = Demo10()
    new data = dict(d10 data)
    dts_name = new_data['dts_name']
    model name = 'linear svc'
    # linear_svc solo, linear_svc con rff y linear_svc con nystroem
    m1 = get a model(model name=model name, sampler name='identity', dts name=dt
s name)
    m2 = get a model(model name=model name, sampler name='rff', dts name=dts nam
e)
    m3 = get a model(model name=model name, sampler name='nystroem', dts name=dt
s_name)
    models = [m1, m2, m3,]
    new data['models'] = models
    d10.non interactive(**new data)
```

In [8]:

```
def test_dataset(d10_data, dts_name):
    new_data = dict(d10_data)
    new_data['dts_name'] = dts_name
    display(md(f'# {dts_name}'))
    test_dt(new_data)
    test_logit(new_data)
    test_linear_svc(new_data)
```

In [9]:

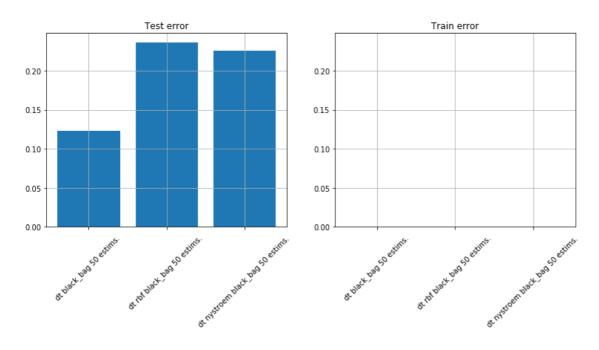
```
def test_everything():
    for dts_name in SUPPORTED_DATASETS:
    for dts_name in ['mnist', 'segment']:
        test_dataset(d10_data, dts_name=dts_name)
```

In [10]:

test_everything()

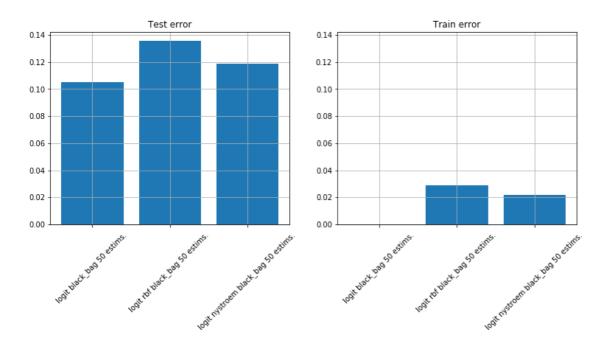
mnist

Demo genérica v10

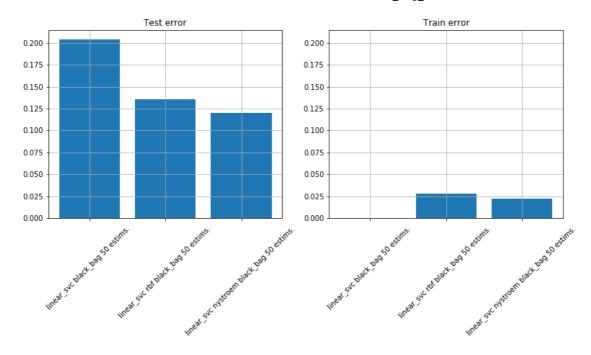


black_bag_box

Demo genérica v10

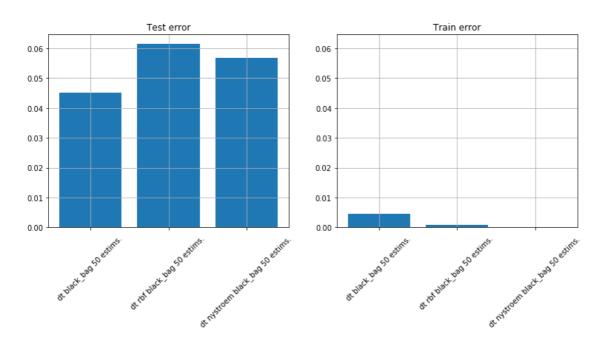


Demo genérica v10

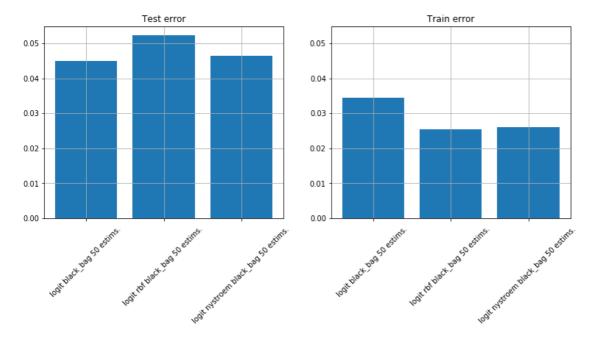


segment

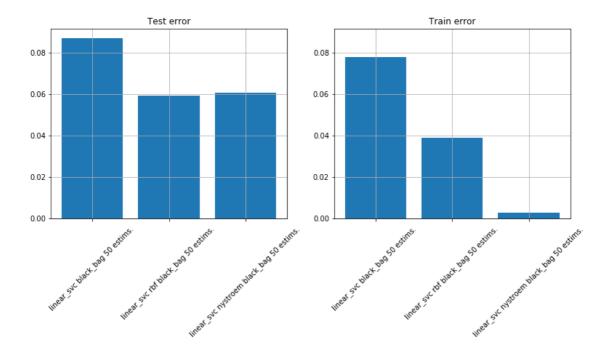
Demo genérica v10



Demo genérica v10



Demo genérica v10



In []: