Data Science Lab: process and methods

Lab 8 solution

```
[1]: import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
%matplotlib inline
```

```
[2]: plt.rcParams["figure.figsize"] = (10, 6)
```

0.1 Exercise 1

In this exercise, we will try different regression models on points drawn from three different analytical functions. We will see that the overall quality of the models heavily depends on the shape of the functions.

That said, let's define our three analytical functions.

```
[3]: def f1(x):
    return x * np.sin(x) + 2*x
def f2(x):
    return 10 * np.sin(x) + x**2
def f3(x):
    return np.sign(x) * (300 + x**2) + 20 * np.sin(x)
```

0.1.1 Exercises 1.1 and 1.2

As usual, we can define convenient functions to generate and plot our data.

```
[4]: def generate_X_y(f):
    tr = 20
    n_samples = 100
    X = np.linspace(-tr, tr, n_samples)
    y = f(X)
    return X, y
def plot_f(X, y, title):
```

```
LW = 4

fig, ax = plt.subplots()

ax.plot(X, y, color='cornflowerblue', linewidth=.5*LW, label="ground truth")

fig.suptitle(title)

#ax.scatter(X_train, y_train, color='navy', s=30, marker='o', 

⇔label="training points")
```

[5]: for f in [f1, f2, f3]:
 X, y = generate_X_y(f)
 plot_f(X, y, f)



<function f1 at 0x7ffe54e2a158>

<function f2 at 0x7ffe54e2a0d0>



<function f3 at 0x7ffe54e2a2f0>



f1 has two main components: a sine wave with an increasing magnitude (due to the factor x), plus an additive factor that gives a non zero slope to the curve.

f2 has a sine wave with fixed amplitude, modulated by a parabolic function.

f3 presents a sine wave modulated by a parabolic function. However, it is different from f2. The quadratic expression changes concativity (i.e. there is an inflection point) at zero, due to the *sign* component. Also, the factor sign(x) 300 produces a discontinuity point of type one.

The shape of the functions tells us that f2 and f3 have an infinite asymptotic value, for both positive and negative values. Hence, they can be approximated with polynomial regressors. However, f3 has a discontinuity point at 0, which can harden the approximation for classifiers. Even if the ordinary least squares regressor would fit the linear trend present in f1, we can see that all the three functions cannot be approximated with a linear regressor, with sufficient results, for large values of x.

Back to programming, you can notice how the functions behave as simple objects in Python. It is not unsual to assign them to variables or, like in this case, create an array of them. Here, the fvariable points, at each loop cycle, to a different function in memory, and gets invoked as a callable object. Additionally, you can note the printed version of f contains the name of the assigned function and the memory address where its code is stored.

Exercises 1.3, 1.4, and 1.5 Let's now fit our regression models. To do so, we define a function to create the training and test points given a function and a scikit-learn Pipeline to apply to them. For the seek of readability, we inspect one analytic function at a time.

Note that here the cardinality of training and test sets are reversed with respect to common cases. Typically, the 70% of the dataset is kept as training set and the remaining 30% is used to test the model.

In the following cells, we test different regression algorithm and evaluate the regression error through two metrics: the MSE and the R2 score. Thus, we use a convenient Python function that, given an analytical function, a dataset, and a model, produces the value of the two metrics. To better understand the quality of each regressor, the function provides also a graphical representation of the predicted values against the real values drawn from the original curve.

```
[7]: from sklearn.metrics import mean_squared_error
from sklearn.metrics import r2_score
def evaluate_model(f, X, y, model, model_name):
```

```
X_train, X_test, y_train, y_test = generate_train_test(f, X, y)
# plot the real function and the training points
LW = 2
fig, ax = plt.subplots()
ax.plot(X, y, color='cornflowerblue', linewidth=.5*LW, label="ground truth")
ax.scatter(X_train, y_train, color='navy', s=30, marker='o',_u
=label="training points")
# predict the test points and plot them onto the chart
model.fit(X_train.reshape(-1, 1), y_train)
y_hat = model.predict(X_test.reshape(-1, 1))
ax.plot(X_test, y_hat, linewidth=LW, label=name, color='r')
fig.suptitle(f"{f} approximated by {model_name}")
fig.legend()
return mean_squared_error(y_test, y_hat), r2_score(y_test, y_hat)
[8]: from sklearn.linear_model import LinearRegression
from sklearn neural network import MIPBegressor
```

```
from sklearn.neural_network import MLPRegressor
from sklearn.svm import SVR
from sklearn.ensemble import RandomForestRegressor
from sklearn.linear_model import Ridge
from sklearn.preprocessing import FunctionTransformer, PolynomialFeatures
from sklearn.compose import make_column_transformer
from sklearn.pipeline import make_pipeline
```

The performance of each regressor are collected and displayed with the library PrettyTable.

```
[9]: from prettytable import PrettyTable
     degree = 5
     models = [
         LinearRegression(),
         Ridge(random_state=42),
         MLPRegressor(hidden_layer_sizes=(10,), random_state=42, max_iter=10000),
         MLPRegressor(hidden_layer_sizes=(10,10), activation='tanh', solver='lbfgs',
                      alpha=0.000, batch_size='auto', learning_rate='constant',
                      learning_rate_init=0.01, power_t=0.5, max_iter=10000,
      \rightarrow shuffle=True,
                      random_state=42, tol=0.0001, verbose=True, warm_start=False,
                      momentum=0.0, nesterovs_momentum=False, early_stopping=False,
                      validation_fraction=0.0, beta_1=0.9, beta_2=0.999,
      \rightarrowepsilon=1e-08),
         SVR(gamma='scale'),
         RandomForestRegressor(n_estimators=300),
```

```
make_pipeline(
        make column transformer(
            (FunctionTransformer(np.sin), [0]),
            (PolynomialFeatures(degree), [0])
        ),
        LinearRegression()
    ),
    make_pipeline(
        make column transformer(
            (FunctionTransformer(np.sin), [0]),
            (PolynomialFeatures(degree), [0])
        ),
        Ridge(alpha=1)
    )
]
names = [
    'linreg',
    'ridge',
    'mlp_standard',
    'mlp_tuned',
    'svr',
    'rf',
    f'sin+poly{degree}+linreg',
    f'sin+poly{degree}+ridge'
]
```

Before the actual simulation, let's spend a few comments on the code above. We generated a list with a few models to be tested along with a respective textual representation that is used in the title of the each figure. The single-stage models are: - a simple Linear Regression model; - a MultiLayer Perceptron (a.k.a. feed forward neural network) with a single hidden layer with a reasonable number of hidden nodes; - a deeper MultiLayer Perceptron with tuned parameters. We do not spend too much time on them since it goes beyond the scope of the exercise. However, the provided parameters should let you grasp how wide are the tuning possibilities of this estimator; - a Support Vector Regressor (gamma = 'scale' is recommended for the scikit-learn implementation); - a Random Forest Regressor (the higher the number of estimator, the better).

We also adopted two composite pipelines. Each of them has two steps: 1. in the first step we build a ColumnTransformer using an utility method. The Transform objects in scikit-learn's preprocessing module are typically used to perfom some sort of transformation of the input columns (e.g. scaling and normalization, application of a function). The ColumnTransform lets you specify a series of single Transformers to apply to columns of your choice. In our case, we apply two types of column transformation to the first (and only) column (see the parameter [0] of each tuple). One uses a FunctionTransformer to generate one additional feature in the form sin(x). The other creates new polynomial features using the pattern provided by the class PolynomialFeatures. 2. In the second test we apply the regressor model as usual. Specifically, we test LinearRegression a Ridge again to measure the impact of the previous preprocessing on the performance.

R	tesults for $f = f1$				
: t	<pre>t = PrettyTable()</pre>				
t	.field_names = ['m	odel', 'MSE', 'R2']			
Х	K, y = generate_X_y	(f1)			
f	for model, name in	<pre>zip(models, names):</pre>			
	mse, r2 = evalu	ate_model(f1, X, y, n	no	del, name)	
	t.add_row([name	, mse, r2])			
P	orint(t)				
+-	+		+-		-+
I	model	MSE		R2	I
+-			+-		-+
	linreg	65.72227592689515	1	0.8950661957810067	1
1	ridge	65.74765554283465	1	0.8950256740612001	1
I	mlp_standard	75.46463406818695	I	0.879511306857702	Ι
Ι	mlp_tuned	104.46617412806707	I	0.8332067338073432	Ι
Ι	svr	399.9552743551347	L	0.36142155968215894	
Ι	rf	57.024236887693924	L	0.9089536991085534	
Ι	sin+poly5+linreg	95.50251981840506	L	0.8475183250167385	Ι
		02 41070200557275	I.	0 8508581378836467	Т
	sin+poly5+riage	93.410/239055/3/5		0.0000010100000101	- I
	sin+poly5+ridge	93.41072390557375	' +-		י +-

















We see that LinearRegression and Ridge have comparable results. SVR fails at modeling the correct shape of the curve, the standard MLP converged to an approximation of the linear behavior, while the tuned MLP shows a more complex pattern, although an abnormal spike near zero worsen probably affects negatively the performance. The Random Forest regressor achieved the best performance in both the MSE and R2 scores. Finally, it is worth noting that the addition of sin(x) and the polynomial features up to the fifth degree slightly worsened the performance of Linear-Regression and Ridge. Having the real function shape, we can graphically find a motivation for that. For both the regressors, the predictions (the red curve) are able follow a sinusoidal pattern (which we might have expected by the introduction of sin(x)). Nonetheless, for negative values of x, this curve is in couterphase with the real one, increasing the overall error.

0.1.2 Exercise 1.6

In many real tasks, we typically suppose that the measurements of the predictive variables carry some sort of noise. To reflect this aspect to our synthetic data, we can inject it manually.

```
[11]: def inject_noise(x):
    """Add a random noise drawn from a normal distribution."""
    return x + np.random.normal(0, 50, size=x.size)
[12]: X, y = generate_X_y(f1)
    y = inject_noise(y)
```

```
t = PrettyTable()
t.field_names = ['model', 'MSE', 'R2']
```

```
for model, name in zip(models, names):
    mse, r2 = evaluate_model(f1, X, y, model, name)
    t.add_row([name, mse, r2])
```

print(t)

```
/Users/giuseppe/miniconda3/lib/python3.6/site-
packages/sklearn/neural_network/_multilayer_perceptron.py:571:
ConvergenceWarning: Stochastic Optimizer: Maximum iterations (10000) reached and
the optimization hasn't converged yet.
```

<pre>Interim 1 2957.236994778774 0.11280320127342303 Interim 2 2957.0994988641382 0.11280320127342303 Interim 2 2957.0994988641382 0.11284445124274178 Interim 2 2957.0994988641382 0.11284445124274178 Interim 3 296.3441666075573 0.011068102665194646 Interim 3 296.3441666075573 0.011068102665194646 Interim 3 288.7875228403523 -0.10666831479001027 Interim 3 324.973510100103 0.002479050810008321 Interim 4 556.727445167173 -0.36705783444464113 Interim 3 288.756662982372 0.013344419693040566 Interim 4 3288.756662982372 0.013344419693040566 Interim 4 3288.5728746771806 0.01309954909119293 </pre>	+ model +	MSE	R2
	<pre> linreg ridge mlp_standard mlp_tuned svr rf sin+poly5+linreg sin+poly5+ridge</pre>	 2957.236994778774 2957.0994988641382 3296.3441666075573 3688.7875228403523 3324.973510100103 4556.727445167173 3288.756662982372 3289.5728746771806 	<pre> 0.11280320127342303 0.11284445124274178 0.011068102665194646 -0.10666831479001027 0.002479050810008321 -0.36705783444464113 0.013344419693040566 0.01309954909119293 </pre>

















This kind of noise (normal, standard deviation=50) makes the problem extremely more complex. The initial shape of the function is lost and the performance are worse for any classifier.

Note: the reported warning concerns MLP. Even with 10000 iterations is is not able to converge to stable values for the weights in the network.

```
Results for f = f2
```

```
[13]: t = PrettyTable()
t.field_names = ['model', 'MSE', 'R2']
X, y = generate_X_y(f2)
for model, name in zip(models, names):
    mse, r2 = evaluate_model(f2, X, y, model, name)
    t.add_row([name, mse, r2])
print(t)
```

```
/Users/giuseppe/miniconda3/lib/python3.6/site-
packages/sklearn/neural_network/_multilayer_perceptron.py:571:
ConvergenceWarning: Stochastic Optimizer: Maximum iterations (10000) reached and
the optimization hasn't converged yet.
  % self.max_iter, ConvergenceWarning)
/Users/giuseppe/miniconda3/lib/python3.6/site-
packages/sklearn/neural_network/_multilayer_perceptron.py:470:
```

```
ConvergenceWarning: lbfgs failed to converge (status=1):
```

STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.

Increase the number of iterations (max_iter) or scale the data as shown in: https://scikit-learn.org/stable/modules/preprocessing.html self.n_iter_ = _check_optimize_result("lbfgs", opt_res, self.max_iter)

1		+	L
	model	I MSE	R2
Ī	linreg	15300.007189428956	
I	ridge	15299.785419953309	-0.037405519380177665
I	mlp_standard	383.373251167696	0.9740052872751136
I	mlp_tuned	194.42309666497553	0.9868170965775644
۱	svr	14341.447317768201	0.027574819189968847
I	rf	229.9094464786911	0.9844109363505463
I	sin+poly5+linreg	5.419623147322589e-23	1.0
I	sin+poly5+ridge	0.23623830289128636	0.9999839818067652















ò

5

10

15

20

100

0

-20

-15

-10

-5



The crucial result here concerns the LinearRegression and Ridge models. Given the highly nonlinear behavior, without a proper preprocessing, they performed the worst. However, the higher capability achieved including the sinusoidal component, along with polynomial features, drastically improved the performance. The best performing model here is the Ridge regularitazion in sin+poly5+ridge which copied almost identically the true values. This is a clear example of how an accurate preprocessing becomes crucial, especially for linear models.

As a further exercise, we suggest to test the performance of the other regressors with the same preprocessing applied to improve the linear models. Let's test now what happens adding the Gaussian noise.

[14]: $X, y = generate_X_y(f2)$

```
y = inject_noise(y)
t = PrettyTable()
t.field_names = ['model', 'MSE', 'R2']
for model, name in zip(models, names):
    mse, r2 = evaluate_model(f2, X, y, model, name)
    t.add_row([name, mse, r2])
```

```
print(t)
```

/Users/giuseppe/miniconda3/lib/python3.6/sitepackages/sklearn/neural_network/_multilayer_perceptron.py:571: ConvergenceWarning: Stochastic Optimizer: Maximum iterations (10000) reached and the optimization hasn't converged yet.

model	+ MSE	R2
linreg ridge	13844.561167759284 13844.460508413049	−0.015583193249441907 −0.015575809271066499
mlp_standard	2991.1860609583377	0.7805782173532807
svr	12891.137648265858	0.05435625016904311
rf sin+poly5+linreg	<pre>4113.044606937682 3314.773489446183</pre>	0.6982830351012688 0.7568411013886113
sin+poly5+ridge	3298.91855356633	0.7580041578563832

















The overall shape of the function is retained by the standard MLP, and sin+poly5+[linreg/ridge]. This lead to the lowest overall error.

```
Results for f = f3
```

```
[15]: t = PrettyTable()
t.field_names = ['model', 'MSE', 'R2']
X, y = generate_X_y(f3)
for model, name in zip(models, names):
    mse, r2 = evaluate_model(f3, X, y, model, name)
    t.add_row([name, mse, r2])
print(t)
```

/Users/giuseppe/miniconda3/lib/python3.6/sitepackages/sklearn/neural_network/_multilayer_perceptron.py:571: ConvergenceWarning: Stochastic Optimizer: Maximum iterations (10000) reached and the optimization hasn't converged yet.

			L
	model	I MSE	R2
	linreg	15760.86079984127	0.9222471586304892
1	ridge	15763.614946346856	0.9222335716367915
I	mlp_standard	11102.721825876006	0.9452270926150279
	mlp_tuned	13752.433310925047	0.9321553067913714

















The top performing algorithm is Random Forest again. We can see that it is the only model able to handle the discontinuity at 0: values approaching 0 from the left (small negative values) are similar to the real values of the left-most part of the function, the same applies for small positive values.

```
[16]: X, y = generate_X_y(f3)
y = inject_noise(y)
t = PrettyTable()
t.field_names = ['model', 'MSE', 'R2']
for model, name in zip(models, names):
    mse, r2 = evaluate_model(f3, X, y, model, name)
    t.add_row([name, mse, r2])
print(t)
```

```
/Users/giuseppe/miniconda3/lib/python3.6/site-
```

packages/sklearn/neural_network/_multilayer_perceptron.py:571: ConvergenceWarning: Stochastic Optimizer: Maximum iterations (10000) reached and the optimization hasn't converged yet.

		
model	MSE	R2
<pre>+</pre>	17515.338908756312 17517.105255078095 12352.939018232815 22997.03282444203 305167.54329760035 6353.001176701917 eg 11793.78421643006 11650.803186975163	<pre> 0.9121094984003594 0.9121006350226635 0.9380139880707941 0.8846028180914527 -0.5313051372901239 0.9681211729334968 0.9408197799688728 0.941537246782589 </pre>
+	+	-++















The most robust to noise is Random Forest in this case.

Exercise 2

In this exercise, you will carry out a multivariate regression analysis. Technically speaking, the

preprocessing step added in the pipeline in Exercise 1 also lead to a multivariate analyis, considering also the newly generated features. Now, we generate a synthetic, multi-dimensional dataset using scikit-learn. The nature and the importance of each of the features can be fine-tuned using the make_regression function.

```
[17]: from sklearn.datasets import make_regression
```

Exercise 2.1

We can use the make_regression function to generate a synthetic dataset with 2000 points. You should spend enough time inspecting the function parameters. For now, we recall that, by default: - 100 features are generated, 10 of which are informative - the target variable has a single dimension - no noise is applied. You can set a normal noise with the parameter noise

```
[18]: X, y = make_regression(n_samples=2000, random_state=42)
X.shape, y.shape
```

[18]: ((2000, 100), (2000,))

Exercise 2.2

We can now run again the regression simulation developed in Exercise 1.

Note: here we get back to the normal conditions where we adopt 70% of the dataset as training set and the remaining 30% as test set. This can be achieved by simply changing the value to the train_size parameter.

/Users/giuseppe/miniconda3/lib/python3.6/sitepackages/sklearn/neural_network/_multilayer_perceptron.py:470: ConvergenceWarning: lbfgs failed to converge (status=1): STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.

Increase the number of iterations (max_iter) or scale the data as shown in:

https://scikit-learn.org/stable/modules/preprocessing.html
self.n_iter_ = _check_optimize_result("lbfgs", opt_res, self.max_iter)

+		+		+-		-+
1	model		MSE		R2	I
 	linreg ridge	+ 	1.491861127842467e-25 0.026589221217723277	 	1.0 0.9999993361938854	
I	mlp_standard	I	59.67097245363076	I	0.998510300243291	I
I	mlp_tuned	I	9081.738642192302	I	0.773272274785179	Ι
I	svr	I	38892.567627033684	I	0.029037970232504207	Ι
I	rf	I	10655.011193604842	I	0.7339951582793712	I
I	sin+poly5+linreg	I	40959.66218569419	I	-0.022567527960504208	Ι
I	sin+poly5+ridge	I	40620.5489939929	I	-0.01410148796816535	I
+		+		+		-+

The tested model behave differently. Since the target label is generated by make_regression from a random linear model, we find that linear models and the model which can closely approximate a linear model (e.g. the standard MLP with one hidden layer) perform better. The R2 for the unregularized linear model is 1, meaning that it is able to capture all the information in the 10 informative features (the MSE is approximately 0).

Even though these results suggest that the so-defined problem is simple, we can notice that more complex models fail at grasping this simplicity, leading to high errors. Over-parametrized models strongly suffer the redundancy of information carried by the non-informative 90 features.

Exercise 2.3

Let's now inspect how the performance of our models change when: - some noise is introduced; - the number of informative features increases.

/Users/giuseppe/miniconda3/lib/python3.6/sitepackages/sklearn/neural_network/_multilayer_perceptron.py:470: ConvergenceWarning: lbfgs failed to converge (status=1): STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.

```
Increase the number of iterations (max_iter) or scale the data as shown in:
    https://scikit-learn.org/stable/modules/preprocessing.html
    self.n_iter_ = _check_optimize_result("lbfgs", opt_res, self.max_iter)
```

_		<u>.</u>		–		
ר נ	model		MSE		R2	
	linreg ridge mlp_standard mlp_tuned svr rf	+- 	111.1154487123277 111.32455388067977 397.79932377985136 8652.930271788262 39028.249235400996 10049.773799052658	+. 	0.9972285317141203 0.9972233161626446 0.9900779934493669 0.7841765289516479 0.026548006935037116 0.7493361212404551	+-
	sin+poly5+linreg		40425.61438644836	I	-0.008305411242192973	I
	sin+poly5+ridge	I	40291.90586225675	I	-0.004970421025771943	I
-		+-		+.		-+

As we might have expected, the introduction of a random gaussian noise with standard deviation of 10 decreases the performance of every model.

Let's introduce more informative features. However, keep in mind that those will be used to generate a target value with a linear combination: we can expect that models close to linear will be again the top performer.

/Users/giuseppe/miniconda3/lib/python3.6/sitepackages/sklearn/neural_network/_multilayer_perceptron.py:470: ConvergenceWarning: lbfgs failed to converge (status=1): STOP: TOTAL NO. of ITERATIONS REACHED LIMIT. Increase the number of iterations (max_iter) or scale the data as shown in: https://scikit-learn.org/stable/modules/preprocessing.html

```
self.n_iter_ = _check_optimize_result("lbfgs", opt_res, self.max_iter)
```

ᆂ				ㅗ.		
	model		MSE		R2	
+	linreg ridge mlp_standard mlp_tuned svr rf sin+poly5+linreg sin+poly5+ridge	-+- 	99.92193369275083 100.17526177647642 151.2964519724636 70093.10550001937 288970.03486527514 193262.1739056526 287344.53105641843 287575.8006801805	+ · 	0.9996581789295267 0.9996573123242324 0.9994824328026811 0.7602198089111757 0.011467822909904779 0.3388730509404215 0.01702847842834465 0.016237332506315427	+-
+		+-		+.		-+

Comparing the latter tables with the previous one (with 10 informative features) we can notice two different changes. While Linear Regression, Ridge and the standard MLP have improved their performance (in terms of both MSE and R2), the other, more complex classifiers performed significanly worse. We can conclude that increasing the number of informative features not only brings more information (and lowers the redundancy), but also makes the problem harder, and the latter factor has the strongest impact on the tested models.

0.2 Exercise 2

In the second part of the laboratory, we will cover the topic of time series forecasting, working on the dataset of temperatures collected during the Second World War. Specifically, we will focus on the forecasting of the temperature value one - or more - day ahead of the current time, based on the history of available temperatures.

Exercise 2.1

To quickly load and inspect the dataset, we can use pandas.

```
[22]: import pandas as pd
```

```
[23]: df = pd.read_csv('weatherww2/SummaryofWeather.csv')
```

```
/Users/giuseppe/miniconda3/lib/python3.6/site-
packages/IPython/core/interactiveshell.py:3063: DtypeWarning: Columns
(7,8,18,25) have mixed types.Specify dtype option on import or set
low_memory=False.
```

```
interactivity=interactivity, compiler=compiler, result=result)
```

0.2.1 Exercise 2.2

We can inspect the content of the dataset with the method info.

[24]: df.info()

<classian <<="" th=""><th>ss 'pandas.co</th><th>re.frame.DataFrame</th><th>e'></th></classian>	ss 'pandas.co	re.frame.DataFrame	e'>
Rang	eIndex: 11904	0 entries, 0 to 1	19039
Data	columns (tot	al 31 columns):	
#	Column	Non-Null Count	Dtype
0	STA	119040 non-null	int64
1	Date	119040 non-null	object
2	Precip	119040 non-null	object
3	WindGustSpd	532 non-null	float64
4	MaxTemp	119040 non-null	float64
5	MinTemp	119040 non-null	float64
6	MeanTemp	119040 non-null	float64
7	Snowfall	117877 non-null	object
8	PoorWeather	34237 non-null	object
9	YR	119040 non-null	int64
10	MO	119040 non-null	int64
11	DA	119040 non-null	int64
12	PRCP	117108 non-null	object
13	DR	533 non-null	float64
14	SPD	532 non-null	float64
15	MAX	118566 non-null	float64
16	MIN	118572 non-null	float64
17	MEA	118542 non-null	float64
18	SNF	117877 non-null	object
19	SND	5563 non-null	float64
20	FT	0 non-null	float64
21	FB	0 non-null	float64
22	FTI	0 non-null	float64
23	ITH	0 non-null	float64
24	PGT	525 non-null	float64
25	TSHDSBRSGF	34237 non-null	object
26	SD3	0 non-null	float64
27	RHX	0 non-null	float64
28	RHN	0 non-null	float64
29	RVG	0 non-null	float64
30	WTE	0 non-null	float64
dtyp	es: float64(2	0), int64(4), obj	ect(7)
memo	ry usage: 28.	2+ MB	

Please refer to the laboratory text to discover more on the attributes.

The dataset is composed of 119040 rows. From the output of **info**, it is clear that most of the 31 attributes have missing values. 9 of them are completely null (e.g. FT, FB, etc.). This is commonplace in real-life tasks, where different sampling procedures are used during the time, the measurements can fail, etc.

The information relative to each sensor are reported in a second file. Let's read it.

[25]: sensors = pd.read_csv("weatherww2/WeatherStationLocations.csv")
 sensors.info()

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 161 entries, 0 to 160
Data columns (total 8 columns):
 #
     Column
                       Non-Null Count
                                        Dtype
     _____
                       _____
                                        ____
___
 0
     WBAN
                       161 non-null
                                        int64
 1
     NAME
                       161 non-null
                                        object
 2
     STATE/COUNTRY ID
                       161 non-null
                                        object
 3
     LAT
                       161 non-null
                                        object
 4
     LON
                       161 non-null
                                        object
 5
     ELEV
                       161 non-null
                                        int64
 6
     Latitude
                       161 non-null
                                        float64
 7
     Longitude
                       161 non-null
                                        float64
dtypes: float64(2), int64(2), object(4)
memory usage: 10.2+ KB
```

The latter file contains useful information to characterize our sensors. Specifically, the Latitude and Longitude attribute can be used to group sensors near on the map. As we can see, many attributes in both the files are numerical, floating point. The normalization of these attributes is a matter specific to the considered task. Let's skip it for now, we will discuss about it later.

To discover the sensors with most of the temperature readings, we can use pandas. Note: this is not mandatory, every step in the following cells can be implemented with plain Python.

[26]: df.groupby("STA").size().sort_values(0, ascending=False).head(10)

[26]: STA

22508	2192
10701	2185
22502	2154
22504	2118
10803	1750
11610	1631
16405	1622
11601	1604
10502	1527
11604	1514
dtype:	int64

Grouping the rows by sensor id (STA), counting the size of each group, and, finally, sorting them gives us the list of STAs that show up the most.

Exercise 2.3

Transorming features with dates into Datetime objects is usually a good idea. Meanwhile we can set a new index for our dataset.

```
[27]: df["Date"] = pd.to_datetime(df["Date"])
df = df.set_index("Date")
```

We can now keep only the readings of Sensor 22508 and plot the series of MeanTemps.

```
[28]: mtemps = df[df["STA"] == 22508]["MeanTemp"]
    mtemps.head(5)
```

[28]: Date

1940-01-0120.0000001940-01-0219.4444441940-01-0320.0000001940-01-0421.1111111940-01-0518.333333Name: MeanTemp, dtype: float64

```
Exercise 2.4
```

```
[29]: fig, ax = plt.subplots()
_ = ax.plot(mtemps.index.values, mtemps.values)
# or equivalently
# mtemps.plot(ax=ax)
```



This plot tells us that: - the series has a clear seasonality. We may have expected that: the temperatures raise during the summer and fall down in winter; - while the highest temperature of the year remain constant (except for the year 1944) in the period 1940-1946, the lowest one

increases in the late years; - given the observed range, we can suppose that the temperature values are expressed in Celsius degrees.

Exercise 2.5 There are many strategies to address the forecasting task of a time series. The transformation of the series itself into a structured representation (i.e. a set of records sharing some predictive features) enables the use of machine learning models. We also know these models are trained to learn the function mapping predicting features to a desired value. Thus, to complete the design of a forecasting framework, we need to carefully select a target variable.

ML practitioners often use the following strategy: - predictive features: the features to be used in the structured representation are computed from the values already observed from the series. They can be the values themselves (which makes the approach more similar to classical, statistical autoregressive models like ARIMA), or a more complex combination of them. In this exercise, we will adopt the simpler way and use the values in a window of fixed-length W. Specifically, this window will be rolling along the series, to obtain one record for each time step (for a graphical representation, please refer to the laboratory text);

- target variable: to model the task of forecasting, the target variable should encode a future event. Thus, the algorithm will be able to model the relationship between some already-seen feature values and an upcoming behavior of the series. In this exercise, we will use as target variable the value of the series right after the considered window.

[30]: W = 3 X = list() w = list()

```
y = list()
```

```
[31]: for i in range(mtemps.size - W): # range: [0, mtemps.size - W - 1]
X.append(mtemps.iloc[i:i + W].values.T) # transpose to create a row array
y.append(mtemps.iloc[i + W])
# transform the structured representation into numpy arrays
X = np.array(X)
y = np.array(y)
X.shape, y.shape
```

[31]: ((2189, 3), (2189,))

We can check whether the rolling window has worked properly inspecting a few samples.

[32]: mtemps[:5]

[32]: Date

1940-01-01 20.00000 1940-01-02 19.44444 1940-01-03 20.00000 1940-01-04 21.11111 1940-01-05 18.33333 Name: MeanTemp, dtype: float64

```
[33]: X[:3,:]
[33]: array([20. , 19.4444444, 20. ],
       [19.44444444, 20. , 21.1111111],
       [20. , 21.1111111],
       [20. , 21.1111111],
       [34]: y[:3]
[34]: array([21.1111111, 18.3333333, 20. ])
```

Given a record associated with the day t, y[t] is the value taken by the series at the day t+W+1

(i.e. mtemps[t+W]), which seems correct.

Exercise 2.6 Let's use now the values from 1940 to 1944 as training data and test the model on the remaining year. We can leverage pandas to filter data based on time strings. To do so, we can convert our arrays into pandas DataFrame and Series using the DatetimeIndex from mtemps.

```
[35]: X_df = pd.DataFrame(X, index=mtemps.index[:mtemps.size - W],
                          columns=["t0", "t1", "t2"])
      X_df.head()
[35]:
                         t0
                                    t1
                                               t2
      Date
      1940-01-01
                 20.000000 19.44444
                                        20.000000
      1940-01-02 19.444444 20.000000 21.111111
      1940-01-03 20.000000
                            21.111111
                                        18.333333
      1940-01-04 21.111111 18.333333
                                        20.000000
      1940-01-05 18.333333 20.000000 20.555556
[36]: y_s = pd.Series(y, index=mtemps.index[:mtemps.size - W])
      y_s.head()
[36]: Date
      1940-01-01
                    21.111111
      1940-01-02
                    18.333333
      1940-01-03
                    20.000000
      1940-01-04
                    20.555556
      1940-01-05
                    18.888889
      dtype: float64
[37]: X_train, y_train = X_df.loc["1940":"1944"], y_s.loc["1940":"1944"]
      X_train.index
[37]: DatetimeIndex(['1940-01-01', '1940-01-02', '1940-01-03', '1940-01-04',
                     '1940-01-05', '1940-01-06', '1940-01-07', '1940-01-08',
                     '1940-01-09', '1940-01-10',
```

```
'1944-12-22', '1944-12-23', '1944-12-24', '1944-12-25',
'1944-12-26', '1944-12-27', '1944-12-28', '1944-12-29',
'1944-12-30', '1944-12-31'],
dtype='datetime64[ns]', name='Date', length=1827, freq=None)
```

Keep in mind that we used a forward-looking notation for our window, i.e. for the record t we have the values of the series between t and t+W. Under this conditions, at the end of the year, we are going to have W data points that include information from the next year. This is not a problem up if we slightly shift the beginning of our test set. Specifically, we might want to start our test set for the Wth day or the year.

Including some information from what it is consider the test set into the training procedure is a wrong habit known as **data leakage**. You should always stop and take a moment to consider if any data leakage is happening within your pipeline.

```
[38]: from datetime import date
initial_day = date(1944, 12, 31) + pd.Timedelta(f"{W} days")
initial_day
```

```
[38]: datetime.date(1945, 1, 3)
```

```
[39]: X_test, y_test = X_df.loc[initial_day:], y_s.loc[initial_day:]
X_test.shape
```

```
[39]: (360, 3)
```

Note how neat is the indexing on pandas object with Datetime indices. You are allowed to specify a numpy-like interval with dates as datetime.date or strings. Of course, more complex and useful indexings are possibile, but they go beyond the scope of the laboratory.

```
\#\#\#\# Exercise 2.7
```

Since we still have in memory the definitions of models and names (this is one of the beauties and curses of Jupyter notebooks) we can recycle and them for our current task.

```
[40]: t = PrettyTable()
t.field_names = ['model', 'MSE', 'R2']
for model, name in zip(models, names):
    model.fit(X_train, y_train)
    y_hat = model.predict(X_test)
    mse = mean_squared_error(y_test, y_hat)
    r2 = r2_score(y_test, y_hat)
    t.add_row([name, mse, r2])
print(t)
+-----+
    model | MSE | R2 |
```

I	linreg	Ι	0.7708838668711151	Ι	0.6684666582498298	Ι
I	ridge	Ι	0.7708190312576426	Ι	0.6684945420446823	Ι
I	mlp_standard	Ι	0.7756832982117083	Ι	0.6664025710646799	Ι
I	mlp_tuned	Ι	2.407530887487392	Ι	-0.03540467610914	Ι
I	svr	Ι	0.7606632382515228	Ι	0.6728622349464781	Ι
I	rf	Ι	0.8362497037896549	Ι	0.6403548306695499	Ι
I	<pre>sin+poly5+linreg</pre>	Ι	1.1396245117527566	Ι	0.5098826957485592	Ι
I	sin+poly5+ridge	Ι	1.1393130367062887	Ι	0.5100166515458784	Ι
+-		-+-		-+-		-+

Even in this case, the models with behavior more close to linear perform the better. This could imply that some sort of linear relationship exists between the past W values of the series and the target. Additional comments can be made: - the choice of custom tuning of MLP is wrong in this case, since it lead to the worst performance. Keep in mind that we do not have carried out any validation. The validation step is used for hyperparameter tuning, iterating the training process to identify the best configurations. However, validating a machine learning approach based on time series requires further attention. Cross-validation is not allowed, for example, since values have an intrinsc order and cannot be shuffled. Scikit-learn offers a validation strategy for time series thourgh the TimeSeriesSplit class. You can find complete examples on how to use it on the official documentation. The hyperparameter tuning can be mixed with TimeSeriesSplit the already seen GridSearchCV. - our approach is intentionally non incremental, i.e. we train the model once and we use it to forecast an entire future year. Although this is correct for research purposes, in practical applications we might find useful to fed fresh information to the model (e.g. by running the training again) as it comes, while the time passes by and new records are collected. The profitability of this approach, however, depends on the application (e.g. there are cases in which you are not able to train the model iteratively in time) and an improvement in performance should not be taken for granted.

Exercise 2.8 Thanks to matplotlib, we can plot our predictions onto the real series for the test year. For simplicity, we can inspect the best performing model, the Ridge regularizer.

[41]: model = Ridge(random_state=42)



[41]: 'The average |error| is: 0.65 +- 0.59 degrees Celsius'

The figure shows that Ridge provides a good approximation but its predictions are somewhat lagged. The model takes at least one day to react to the current trend and so induces a non-zero error. On the bottom-end of the figure, the plot of residuals shows that the larger is the change in temperature w.r.t. the previous day (i.e. a high volatility among consecutive days is present), the larger is the error due to the lagged limitations of the estimator.

Even though these results suggest that the only values of the series are not sufficient to build a robust forecasting model (e.g. we may think that other, more expressive features are needed), the average absolute error 0.65 and the standard deviation is 0.59, which may be acceptable values for our domain after all.

Predicting a value further than one day ahead is not possibile at the moment, but many strategies exist to expand the forecast horizon. For example, one can train different models on different target variables: one encoding the value one day ahead (i.e. what we have done here), one with the value two days ahead as target, and so on. In this experimental setting, you would end up having one model per horizon step, which could add some computational cost.

Exercise 2.9

As a further exercise, we leave to you the tweaking of this pipeline to test and visually assess the

performance of the remaining regressors.