Étude sur le propagation du Covid 19 avec machine learning

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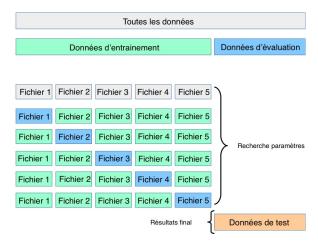
Sommaire

- 1 Première approche : simple regression
 - Validation croisée et hyperparamètres
 - Résultats avec SVR

- 2 Approche multivariées
 - Multiregresseur : 'RegressorChain'
 - Réseau neuronal

Recherche du meilleur paramètre

Principe de la validation croisée:



SVR, premier résultat

Approche à l'aide du modèle SVR. Noyau « rbf » \rightarrow ajustement du paramètre C

— Modèle SVR (prédit les données entre le 02/12/20 et 16/12/20)

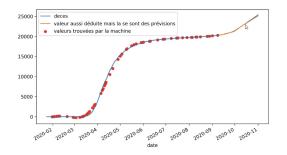


Figure: Premier résultat avec SVR et découpage inadapté, C = 50k

Découpage adapté pour la validation croisée

Remise en question de la méthode de découpage

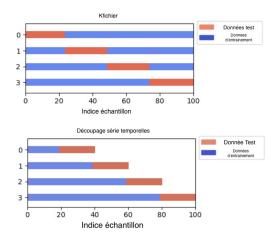


Figure: Comparaison des découpages pour la validation croisée

Avec
$$C = 10^5$$

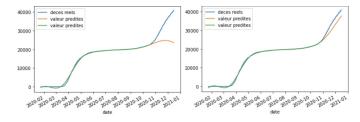


Figure: À gauche: prediction avant point d'inflexion; à droite: après.

⇒ Prédiction inefficace du point d'inflexion.

RegressorChain SVR

Multiregresseur RegressorChain

Corrélation : Cas confirmé \rightarrow réanimation \rightarrow décès

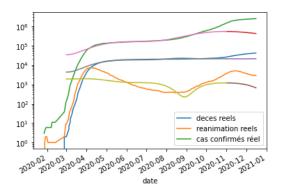


Figure: Résultat avec SVR insatisfaisant

RegressorChain TheilSenRegressor

Changement de régresseur : meilleurs résultats

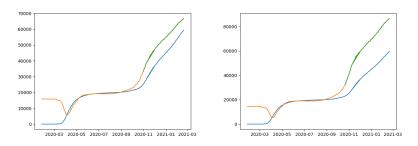
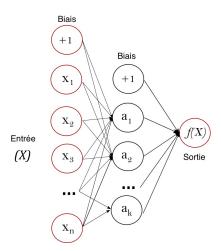


Figure: Prédiction avec et sans point d'inflexion

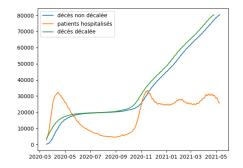
Principe du réseau neuronal



Décalage de courbes

Représentation avant et aprés décalage

date		date	
2020-03-17	175	2020-03-17	1100
2020-03-18	264	2020-03-18	1331
2020-03-19	372	2020-03-19	1696
2020-03-20	450	2020-03-20	1995
2020-03-21	562	2020-03-21	2314



Corrélation et Premier résultat

Corrélation cas confirmés et décès : 0.978

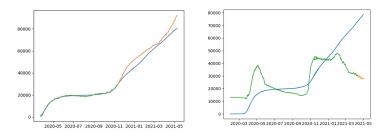
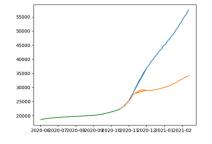


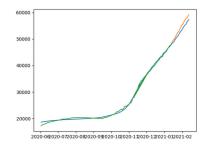
Figure: Tests avec et sans cas confirmés

Échec du modèle sans les cas confirmés \rightarrow cohérent avec corrélation

Résultats

Réseaux neuronaux : Meilleurs paramètres





Neural network avec 7 jour de décalage; max_iter=90k

Augmentation du décalage

Résultats insatisfaisant

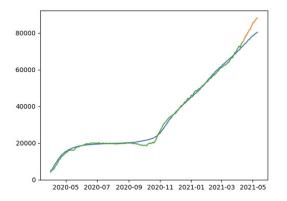


Figure: Tests avec 30 jours de décalage

Conclusion

Modèle peu fiable hors situation stabilisée ou début d'évolution

```
import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
from sklearn pipeline import make pipeline
from sklearn.neural network import MLPRegressor
from pandas. plotting import
   register matplotlib converters
register matplotlib converters()
from sklearn metrics import make scorer
from sklearn.metrics import mean squared error
scorer = make scorer(mean squared error,
   greater is better=False)
from sklearn.preprocessing import
   StandardScaler
from sklearn.model selection import
   TimeSeriesSplit
from sklearn.model selection import
   GridSearchCV
```

```
def timeshift day(data,n):
    data.index = data.index.shift(periods = n,
   freq = 'D'
tscv = TimeSeriesSplit(n splits=5)
covid = pd.read csv('synthese-fra.csv',
   index col='date', parse dates=True)#,
   index col='date', parse dates=True)
for i in covid:
    if not (covid[i].name in ["
   total deces hopital", 'patients reanimation',
```

```
patients hospitalises', 'total cas confirmes'
   1):
        covid.drop([i], axis=1, inplace = True)
jour decale=7
covid = covid.dropna(axis=0)
timeshift day(covid["total deces hopital"], -
   jour decale)
full size=covid.count()[0]
subject = ['patients reanimation',
         'patients hospitalises','
   total cas confirmes'
result = ["total deces hopital"]
\#start:2020-03-17
predit jour='2021-04-01'
```

```
X = covid[:predit jour][subject]
y = covid[result][: predit jour]
y = y.values.reshape(381,)
X result=covid[predit jour:][subject]
y result=covid[result][predit jour:]
y result = y result.values.reshape(40,)
params={
    'mlpregressor max iter': [10000],
    'mlpregressor tol': [0.0001],
    'mlpregressor n iter no change': [2],
model=make pipeline(StandardScaler(),
   MLPRegressor())
grid=GridSearchCV (model, param grid=params, cv=
```

```
tscv, scoring=scorer)
grid.fit(X,y)
timeshift day(covid["total deces hopital"],
  jour decale)
covid=covid[jour decale:]
plt.figure()
plt.plot(covid.index,covid["total deces hopital
plt.plot(covid.index[374:], grid.predict(covid[
   2021-03-25': [subject]))
plt.plot(covid.index[:374], grid.predict(covid[:
   '2021-03-25'][subject]))
```

```
plt.show()
```

NN deces datagouv copie.py

```
import matplotlib.pyplot as plt
import pandas as pd
from sklearn model selection import
   train test split
from sklearn.pipeline import make pipeline
from sklearn.preprocessing import
   StandardScaler
from sklearn.svm import SVR
from sklearn.model selection import
   GridSearchCV
from sklearn.linear model import Lasso
covid = pd.read excel('covid1.xlsx',index col=
   date', parse dates=True)
```

```
covid = covid[covid['granularite']=='pays']
covid.drop(['cas ehpad', 'cas confirmes ehpad',
    'cas possibles ehpad', 'source url',
            'deces ehpad', 'reanimation',
   gueris', 'source nom'.
             'nouvelles hospitalisations', '
   nouvelles reanimations', 'depistes',
              'maille code', 'maille nom','
   source archive', 'source type'], inplace=True,
   axis=1
covid = covid . dropna(axis=0)
subject = ['deces']
jour prevu=75
```

```
size=covid.count()[0]-jour prevu
X = covid.index[:size]
y = covid [: size][subject]
y = y.values.reshape(size)
X = X. values . reshape (size , 1)
X train, y train=X, y
model=make pipeline(StandardScaler(), SVR())
params={
    'svr degree': [0],
    'svr C': [600000],
grid = GridSearchCV (model, param grid=params, cv
   =4)
```

import matplotlib.pyplot as plt
import numpy as np
import pandas as pd
from matplotlib import pyplot as plt
from sklearn.pipeline import make_pipeline
from sklearn.preprocessing import
 StandardScaler
from sklearn.model_selection import
 TimeSeriesSplit

```
from sklearn.svm import SVR
from sklearn.multioutput import RegressorChain
from sklearn.model selection import
   GridSearchCV
from pandas. plotting import
   register matplotlib converters
register matplotlib converters()
from sklearn.metrics import make scorer
from sklearn.metrics import mean squared error
from sklearn.linear model import
   RANSACRegressor
scorer = make scorer (mean squared error,
   greater is better=False)
from sklearn.linear model import
   TheilSenRegressor
tscv = TimeSeriesSplit(n splits=10)
```

```
covid = pd.read excel('chiffre covid 14032021.
   xlsx',index col='date',parse dates=True)#,
   index col='date', parse dates=True)
covid = covid[covid['granularite']=='pays']
for i in covid:
    if not (covid[i].name in ["deces",'
   cas confirmes', 'reanimation', 'hospitalises'
   1):
        covid.drop([i], axis=1, inplace = True)
covid=covid [5:]
covid = covid . dropna(axis=0)
subject = ['hospitalises','cas confirmes','
   reanimation'
result = [ "deces "]
```

```
size = 255
X = covid[: size][subject]
y = covid [: size] [result]
X result=covid[size:][subject]
y result=covid[size:][result]
y = y.values.reshape(size,)
y result = y result.values.reshape(348-size,)
params={
    'theilsenregressor tol': [0.001],
    'theilsenregressor random state': [i for i
    in range (45)],
    'theilsenregressor max iter': [90000],
```

```
model=make pipeline(StandardScaler(),
   TheilSenRegressor())
grid=GridSearchCV (model, param grid=params, cv=
   tscv)
grid. fit (X, y)
p=covid[subject]
plt.figure()
plt . plot ( covid . index , covid [ 'deces '])
plt.plot(covid.index[size:], grid.predict(
   X result))
```

plt.show()

multi regressor tipe copie.py