AI4M_C1_W1_lecture_ex_04

June 22, 2020

0.1 AI for Medicine Course 1 Week 1 lecture exercises

1 Patient Overlap and Data Leakage

Patient overlap in medical data is a part of a more general problem in machine learning called **data leakage**. To identify patient overlap in this week's graded assignment, you'll check to see if a patient's ID appears in both the training set and the test set. You should also verify that you don't have patient overlap in the training and validation sets, which is what you'll do here.

Below is a simple example showing how you can check for and remove patient overlap in your training and validations sets.

```
In [1]: # Import necessary packages
    import pandas as pd
    import numpy as np
    import matplotlib.pyplot as plt
    %matplotlib inline
    import os
    import seaborn as sns
    sns.set()
```

1.0.1 Read in the data from a csv file

First, you'll read in your training and validation datasets from csv files. Run the next two cells to read these csvs into pandas dataframes.

There are 1000 rows and 16 columns in the training dataframe

```
3 00012359_002.png
                                                      0
                                                                             0
        4 00017951_001.png
                                                      0
                                                                             0
           Effusion Emphysema Fibrosis Hernia
                                                   Infiltration Mass Nodule \
        0
                                                0
                                                                     0
                  0
                             0
                                        0
                                                                             0
        1
                  1
                             0
                                        0
                                                0
                                                               1
                                                                     0
                                                                             0
        2
                                        0
                                                              0
                                                                     0
                  0
                             0
                                                0
                                                                             0
                  0
                             0
                                                              0
                                                                     0
                                        0
                                                0
                                                                             0
        4
                                        0
                                                0
                                                                     0
           PatientId Pleural_Thickening Pneumonia Pneumothorax
        0
                8270
                                                   0
                                                                  0
                                        0
        1
               29855
                                                   0
                                                                  0
                                        0
        2
                                                   0
                                                                  0
                1297
                                        1
        3
               12359
                                        0
                                                   0
                                                                  0
        4
                                                                  0
               17951
                                        0
                                                   0
In [3]: # Read csv file containing validation data
        valid_df = pd.read_csv("nih/valid-small.csv")
        # Print first 5 rows
        print(f'There are {valid_df.shape[0]} rows and {valid_df.shape[1]} columns in the valid
        valid_df.head()
There are 109 rows and 16 columns in the validation dataframe
Out[3]:
                      Image Atelectasis Cardiomegaly Consolidation
        0 00027623_007.png
                                        0
                                                      0
        1 00028214_000.png
                                                      0
                                                                      0
                                                                             0
                                        0
        2 00022764_014.png
                                        0
                                                      0
                                                                      0
                                                                             0
        3 00020649_001.png
                                                      0
                                                                      0
                                                                             0
                                        1
        4 00022283_023.png
                                        0
                                                      0
                                                                      0
                                                                             0
           Effusion Emphysema Fibrosis Hernia
                                                   Infiltration Mass
                                                                        Nodule \
        0
                                                                     0
                  1
                             0
                                        0
                                                0
                                                              0
                                                                             0
                  0
                                        0
                                                              0
                                                                     0
        1
                             0
                                                0
                                                                             0
        2
                  0
                              0
                                        0
                                                0
                                                              0
                                                                     0
                                                                             0
        3
                  1
                             0
                                        0
                                                0
                                                              0
                                                                     0
                                                                             0
```

PatientId Pleural_Thickening Pneumonia Pneumothorax

1.0.2 Extract and compare the PatientId columns from the train and validation sets

By running the next four cells you will do the following: 1. Extract patient IDs from the train and validation sets 2. Convert these arrays of numbers into set() datatypes for easy comparison 3. Identify patient overlap in the intersection of the two sets

```
In [4]: # Extract patient id's for the training set
        ids_train = train_df.PatientId.values
        # Extract patient id's for the validation set
        ids_valid = valid_df.PatientId.values
In [5]: # Create a "set" datastructure of the training set id's to identify unique id's
        ids_train_set = set(ids_train)
        print(f'There are {len(ids_train_set)} unique Patient IDs in the training set')
        # Create a "set" datastructure of the validation set id's to identify unique id's
        ids_valid_set = set(ids_valid)
        print(f'There are {len(ids_valid_set)} unique Patient IDs in the validation set')
There are 928 unique Patient IDs in the training set
There are 97 unique Patient IDs in the validation set
In [6]: # Identify patient overlap by looking at the intersection between the sets
        patient_overlap = list(ids_train_set.intersection(ids_valid_set))
       n overlap = len(patient overlap)
        print(f'There are {n_overlap} Patient IDs in both the training and validation sets')
        print(f'These patients are in both the training and validation datasets:')
       print(f'{patient overlap}')
There are 11 Patient IDs in both the training and validation sets
These patients are in both the training and validation datasets:
[20290, 27618, 9925, 10888, 22764, 19981, 18253, 4461, 28208, 8760, 7482]
```

1.0.3 Identify rows (indices) of overlapping patients and remove from either the train or validation set

Run the next two cells to do the following: 1. Create lists of the overlapping row numbers in both the training and validation sets. 2. Drop the overlapping patient records from the validation set (could also choose to drop from train set)

```
print(f'These are the indices of overlapping patients in the training set: ')
    print(f'{train_overlap_idxs}')
    print(f'These are the indices of overlapping patients in the validation set: ')
    print(f'{valid_overlap_idxs}')

These are the indices of overlapping patients in the training set:
[306, 186, 797, 98, 408, 917, 327, 913, 10, 51, 276]
These are the indices of overlapping patients in the validation set:
[104, 88, 65, 13, 2, 41, 56, 70, 26, 75, 20, 52, 55]

In [8]: # Drop the overlapping rows from the validation set
    valid_df.drop(valid_overlap_idxs, inplace=True)
```

1.0.4 Check that everything worked as planned by rerunning the patient ID comparison between train and validation sets.

When you run the next two cells you should see that there are now fewer records in the validation set and that the overlap problem has been removed!

```
In [9]: # Extract patient id's for the validation set
    ids_valid = valid_df.PatientId.values
    # Create a "set" datastructure of the validation set id's to identify unique id's
    ids_valid_set = set(ids_valid)
    print(f'There are {len(ids_valid_set)} unique Patient IDs in the validation set')
There are 86 unique Patient IDs in the validation set
```

There are O Patient IDs in both the training and validation sets

1.0.5 Congratulations! You removed overlapping patients from the validation set!

You could have just as well removed them from the training set.

Always be sure to check for patient overlap in your train, validation and test sets.