# C1\_W1\_Lab\_4\_patient\_overlap\_and\_data\_leakage

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## 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()
```

#### 1.1 1. Data

#### 1.1.1 1.1 Loading the Data

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
                                               0
                                                              1
                                                                            0
        2
                             0
                                       0
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                  0
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                                               0
                                                              0
                                                                    0
                                                                            0
                                       0
                                               0
                                                                    0
                                                                            0
           PatientId Pleural_Thickening Pneumonia Pneumothorax
        0
                8270
                                                  0
                                                                 0
                                       0
        1
               29855
                                       0
                                                   0
                                                                 0
        2
                                       1
                                                   0
                                                                 0
                1297
        3
               12359
                                       0
                                                   0
                                                                 0
        4
               17951
                                       0
                                                   0
                                                                 0
In [3]: # Read csv file containing validation data
       valid_df = pd.read_csv("data/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

There are 109 rows and 16 columns in the validation dataframe

valid\_df.head()

Out[3]:	Image		electasis	Cardiomegaly		Consolidation		Edema	\
0	00027623_007.png		0	0		0		1	
1	00028214_000.png		0	0			0	0	
2	00022764_014.png		0	0		0		0	
3	00020649_001.png		1	0			0	0	
4	00022283_023.png		0	0			0	0	
	Effusion	Emphysema	Fibrosis	Hernia	Infil	tration	Mass	Nodule	\
0	1	0	0	0		0	0	0	
1	0	0	0	0		0	0	0	
2	0	0	0	0		0	0	0	
3	1	0	0	0		0	0	0	
4	0	0	0	0		0	0	0	
	PatientId	Pleural T	ural_Thickening		ia Pn	eumothor	ax		
0	27623	_	0		0		0		
1	28214		0		0		0		
2	22764		0		0		0		
3	20649		0		0		0		
4	22283		0		0		0		

#### 1.1.2 1.2 Extracting Patient IDs

By running the next three cells you will do the following: 1. Extract patient IDs from the train and validation 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
```

#### 1.1.3 1.3 Comparing PatientIDs for Train & 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

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]
```

#### 1.1.4 1.4 Identifying & Removing Overlapping Patients

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.

Note: You could also choose to drop them from train set.

```
In [7]: train_overlap_idxs = []
     valid_overlap_idxs = []
     for idx in range(n_overlap):
         train_overlap_idxs.extend(train_df.index[train_df['PatientId'] == patient_overlap[
```

```
valid_overlap_idxs.extend(valid_df.index[valid_df['PatientId'] == patient_overlap[
    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)
```

#### 1.1.5 1.5 Sanity Check

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!

```
# 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 training set')

There are 86 unique Patient IDs in the training set

In [10]: # Identify patient overlap by looking at the intersection between the sets
patient_overlap = list(ids_train_set.intersection(ids_valid_set))
```

print(f'There are {n\_overlap} Patient IDs in both the training and validation sets')

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

### 1.1.6 Congratulations! You removed overlapping patients from the validation set!

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

n\_overlap = len(patient\_overlap)

In [9]: # Extract patient id's for the validation set
 ids\_valid = valid\_df.PatientId.values

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