```
In [1]: !mkdir ~/.keras
!mkdir ~/.keras/models
!cp ../input/keras-pretrained-models/*notop* ~/.keras/models/
!cp ../input/keras-pretrained-models/imagenet_class_index.json ~/.keras/model
s/
!cp ../input/keras-pretrained-models/resnet50* ~/.keras/models/

The syntax of the command is incorrect.
The syntax of the command is incorrect.
'cp' is not recognized as an internal or external command,
```

operable program or batch file.

'cp' is not recognized as an internal or external command, operable program or batch file.

'cp' is not recognized as an internal or external command,

'cp' is not recognized as an internal or external command, operable program or batch file.

```
In [2]: %matplotlib inline
    import numpy as np # linear algebra
    import pandas as pd # data processing, CSV file I/O (e.g. pd.read_csv)
    from glob import glob
    import matplotlib.pyplot as plt
    import os
    import pandas as pd
    import seaborn as sns
    from skimage.util.montage import montage2d
    from skimage.io import imread
    base_dir = os.path.join('D:/Neural Network/','input', 'pulmonary-chest-xray-ab normalities')
```

In []:

```
mont paths = glob(os.path.join(base dir, 'Montgomery', 'MontgomerySet', '*',
In [3]:
        '*.*'))
        shen paths = glob(os.path.join(base dir, 'ChinaSet AllFiles', 'ChinaSet AllFil
        es', '*', '*.*'))
        print('Montgomery Files', len(mont_paths))
        print('Shenzhen Files', len(shen_paths))
        all paths df = pd.DataFrame(dict(path = mont paths + shen paths))
        all paths df['source'] = all paths df['path'].map(lambda x: x.split('/')[2])
        all paths df['file id'] = all paths df['path'].map(lambda x: os.path.splitext(
        os.path.basename(x))[0])
        all paths df['patient group'] = all paths df['file id'].map(lambda x: x.split
        ('_')[0])
        all paths df['file ext'] = all paths df['path'].map(lambda x: os.path.splitext
        (x)[1][1:])
        all_paths_df = all_paths_df[all_paths_df.file_ext.isin(['png', 'txt'])]
        all paths df['pulm state'] = all paths df['file id'].map(lambda x: int(x.spli
        t('_')[-1]))
        all paths df.sample(5)
```

Montgomery Files 277 Shenzhen Files 1325

Out[3]:

	path	source	file_id	patient_group	file_ext	pulr
1467	D:/Neural Network/input\pulmonary- chest-xray-a	input\pulmonary- chest-xray- abnormalities\China	CHNCXR_0529_1	CHNCXR	png	
41	D:/Neural Network/input\pulmonary- chest-xray-a	input\pulmonary- chest-xray- abnormalities\Montg	MCUCXR_0056_0	MCUCXR	txt	
895	D:/Neural Network/input\pulmonary- chest-xray-a	input\pulmonary- chest-xray- abnormalities\China	CHNCXR_0619_1	CHNCXR	txt	
706	D:/Neural Network/input\pulmonary- chest-xray-a	input\pulmonary- chest-xray- abnormalities\China	CHNCXR_0430_1	CHNCXR	txt	
455	D:/Neural Network/input\pulmonary- chest-xray-a	input\pulmonary- chest-xray- abnormalities\China	CHNCXR_0179_0	CHNCXR	txt	

```
In [4]: clean patients df = all paths df.pivot table(index = ['patient group', 'pulm s
        tate', 'file_id'],
                                                      columns=['file ext'],
                                                      values = 'path', aggfunc='first')
         .reset index()
        clean patients df.sample(5)
        from warnings import warn
        def report to dict(in path):
            with open(in_path, 'r') as f:
                all_lines = [x.strip() for x in f.read().split('\n')]
            info dict = {}
            try:
                if "Patient's Sex" in all_lines[0]:
                     info dict['age'] = all lines[1].split(':')[-1].strip().replace('Y'
                     info dict['sex'] = all lines[0].split(':')[-1].strip()
                     info dict['report'] = ' '.join(all lines[2:]).strip()
                     info_dict['age'] = all_lines[0].split(' ')[-1].replace('yrs', '').
        replace('vr', '')
                     info dict['sex'] = all lines[0].split(' ')[0].strip()
                     info_dict['report'] = ' '.join(all_lines[1:]).strip()
                 info_dict['sex'] = info_dict['sex'].upper().replace('FEMALE', 'F').rep
        lace('MALE', 'M').replace('FEMAL', 'F')[0:1]
                 if 'month' in info dict.get('age', ''):
                     info dict.pop('age') # invalid
                if 'day' in info_dict.get('age', ''):
                     info dict.pop('age') # invalid
                elif len(info dict.get('age',''))>0:
                     info_dict['age'] = float(info_dict['age'])
                     info dict.pop('age')
                 return info dict
            except Exception as e:
                 print(all lines)
                warn(str(e), RuntimeWarning)
                 return {}
         report df = pd.DataFrame([dict(**report to dict(c row.pop('txt')), **c row)
                           _, c_row in clean_patients_df.iterrows()])
        report df.sample(5)
```

```
['male 16month', 'normal']
['male 16month', 'normal']
['female24yrs', 'normal']
```

C:\Users\mkahs\Anaconda3\lib\site-packages\ipykernel__main__.py:32: RuntimeW
arning: 'age'

C:\Users\mkahs\Anaconda3\lib\site-packages\ipykernel__main__.py:32: RuntimeW
arning: could not convert string to float: 'female24'

['male35yrs', 'PTB in the left upper field']
['male35yrs', 'Bilateral secondary PTB']

['Male , 38yrs,', 'secondary PTB in the bilateral upper and middle fields, mainly fibrous lesions'] $\[\]$

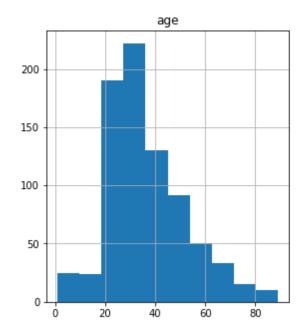
C:\Users\mkahs\Anaconda3\lib\site-packages\ipykernel__main__.py:32: RuntimeW
arning: could not convert string to float: 'male35'

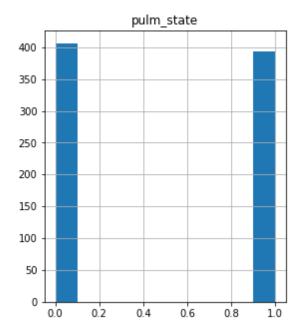
C:\Users\mkahs\Anaconda3\lib\site-packages\ipykernel__main__.py:32: RuntimeW
arning: could not convert string to float: '38,'

Out[4]:

	age	file_id	patient_group	png	pulm_state	report	sex
345	26.0	CHNCXR_0346_1	CHNCXR	D:/Neural Network/input\pulmonary- chest-xray-a	1	STB tuberculosis pleuritis	F
169	23.0	CHNCXR_0170_0	CHNCXR	D:/Neural Network/input\pulmonary- chest-xray-a	0	normal	M
520	34.0	CHNCXR_0521_1	CHNCXR	D:/Neural Network/input\pulmonary- chest-xray-a	1	PTB in the right upper field	M
654	33.0	CHNCXR_0655_1	CHNCXR	D:/Neural Network/input\pulmonary- chest-xray-a	1	secondary PTB in the right upper field,mainly 	M
496	45.0	CHNCXR_0497_1	CHNCXR	D:/Neural Network/input\pulmonary- chest-xray-a	1	bilateral PTB	М

```
In [5]: report_df[['age', 'patient_group', 'pulm_state', 'sex']].hist(figsize = (10, 5
))
```



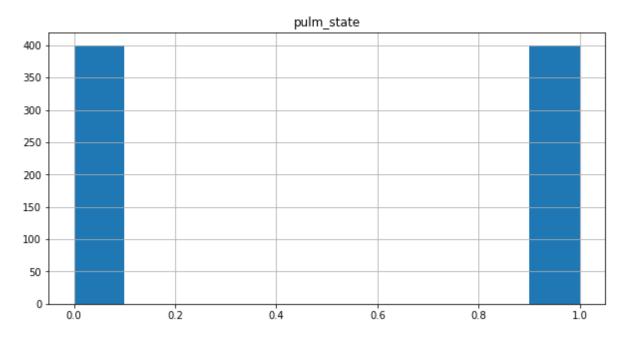


train 600 validation 200

Out[6]:

	age	file_id	patient_group	png	pulm_state	report	sex
628	62.0	CHNCXR_0629_1	CHNCXR	D:/Neural Network/input\pulmonary- chest-xray-a	1	Bilateral secondary PTB	М

New Data Size: 800 Old Size: 600



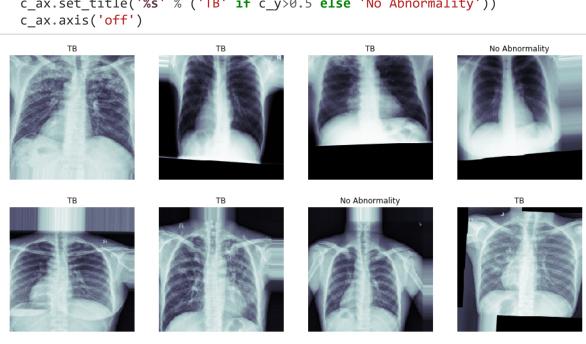
```
from keras.preprocessing.image import ImageDataGenerator
from keras.applications.imagenet utils import preprocess input
from PIL import Image
ppi = lambda x: Image.fromarray(preprocess input(np.array(x).astype(np.float32
)))
IMG SIZE = (224, 224) # slightly smaller than vgq16 normally expects
core_idg = ImageDataGenerator(samplewise_center=False,
                               samplewise std normalization=False,
                               horizontal flip = True,
                               vertical flip = False,
                               height shift range = 0.15,
                               width_shift_range = 0.15,
                               rotation range = 5,
                               shear range = 0.01,
                               fill mode = 'nearest',
                               zoom range=0.2)
```

C:\Users\mkahs\Anaconda3\lib\site-packages\h5py__init__.py:36: FutureWarnin
g: Conversion of the second argument of issubdtype from `float` to `np.floati
ng` is deprecated. In future, it will be treated as `np.float64 == np.dtype(float).type`.

from ._conv import register_converters as _register_converters
Using TensorFlow backend.

```
In [11]: train gen = flow from dataframe(core idg, train df,
                                       path_col = 'png',
                                      y col = 'pulm state',
                                      target size = IMG SIZE,
                                       color mode = 'rgb',
                                      batch size = 32)
         valid gen = flow from dataframe(core idg, valid df,
                                       path_col = 'png',
                                      y_col = 'pulm_state',
                                      target size = IMG SIZE,
                                       color_mode = 'rgb',
                                      batch_size = 256) # we can use much larger batches
         for evaluation
         # used a fixed dataset for evaluating the algorithm
         test_X, test_Y = next(flow_from_dataframe(core_idg,
                                         valid df,
                                       path_col = 'png',
                                      y_col = 'pulm_state',
                                      target size = IMG SIZE,
                                       color_mode = 'rgb',
                                      batch_size = 1024)) # one big batch
```

Ignore next message from keras, values are replaced anyways
Found 0 images belonging to 0 classes.
Reinserting dataframe: 800 images
Ignore next message from keras, values are replaced anyways
Found 0 images belonging to 0 classes.
Reinserting dataframe: 200 images
Ignore next message from keras, values are replaced anyways
Found 0 images belonging to 0 classes.
Reinserting dataframe: 200 images



```
In [33]: from keras.applications.vgg16 import VGG16
         from keras.layers import GlobalAveragePooling2D, Dense, Dropout, Flatten, Inpu
         t, Conv2D, multiply, LocallyConnected2D, Lambda
         from keras.models import Model
         in lay = Input(t x.shape[1:])
         base_pretrained_model = VGG16(input_shape = t_x.shape[1:],
                                        include top = False, weights = 'imagenet')
         base pretrained model.trainable = False
         pt depth = base pretrained model.get output shape at(0)[-1]
         pt features = base pretrained model(in lay)
         from keras.layers import BatchNormalization
         bn features = BatchNormalization()(pt features)
         # here we do an attention mechanism to turn pixels in the GAP on an off
         attn_layer = Conv2D(64, kernel_size = (1,1), padding = 'same', activation = 'r
         elu')(bn features)
         attn layer = Conv2D(16, kernel size = (1,1), padding = 'same', activation = 'r
         elu')(attn layer)
         attn layer = Conv2D(1,
                                          kernel size = (1,1),
                                          padding = 'valid',
                                          activation = 'sigmoid')(attn layer)
         # fan it out to all of the channels
         up_c2_w = np.ones((1, 1, 1, pt_depth))
         up c2 = Conv2D(pt depth, kernel size = (1,1), padding = 'same',
                        activation = 'linear', use bias = False, weights = [up c2 w])
         up_c2.trainable = False
         attn layer = up c2(attn layer)
         mask features = multiply([attn layer, bn features])
         gap features = GlobalAveragePooling2D()(mask features)
         gap mask = GlobalAveragePooling2D()(attn layer)
         # to account for missing values from the attention model
         gap = Lambda(lambda x: x[0]/x[1], name = 'RescaleGAP')([gap_features, gap_mask])
         ])
         gap dr = Dropout(0.5)(gap)
         dr_steps = Dropout(0.25)(Dense(128, activation = 'elu')(gap_dr))
         out layer = Dense(1, activation = 'sigmoid')(dr steps)
         tb model = Model(inputs = [in lay], outputs = [out layer])
         tb model.compile(optimizer = 'adam', loss = 'binary crossentropy',
                                     metrics = ['binary accuracy'])
         # tb model.summary()
         base pretrained model.summary()
```

Layer (type)	Output	Shape	Param #
<pre>input_6 (InputLayer)</pre>	(None,	224, 224, 3)	0
block1_conv1 (Conv2D)	(None,	224, 224, 64)	1792
block1_conv2 (Conv2D)	(None,	224, 224, 64)	36928
block1_pool (MaxPooling2D)	(None,	112, 112, 64)	0
block2_conv1 (Conv2D)	(None,	112, 112, 128)	73856
block2_conv2 (Conv2D)	(None,	112, 112, 128)	147584
block2_pool (MaxPooling2D)	(None,	56, 56, 128)	0
block3_conv1 (Conv2D)	(None,	56, 56, 256)	295168
block3_conv2 (Conv2D)	(None,	56, 56, 256)	590080
block3_conv3 (Conv2D)	(None,	56, 56, 256)	590080
block3_pool (MaxPooling2D)	(None,	28, 28, 256)	0
block4_conv1 (Conv2D)	(None,	28, 28, 512)	1180160
block4_conv2 (Conv2D)	(None,	28, 28, 512)	2359808
block4_conv3 (Conv2D)	(None,	28, 28, 512)	2359808
block4_pool (MaxPooling2D)	(None,	14, 14, 512)	0
block5_conv1 (Conv2D)	(None,	14, 14, 512)	2359808
block5_conv2 (Conv2D)	(None,	14, 14, 512)	2359808
block5_conv3 (Conv2D)	(None,	14, 14, 512)	2359808
block5_pool (MaxPooling2D)	(None,	7, 7, 512)	0

Total params: 14,714,688

Trainable params: 0

Non-trainable params: 14,714,688

In [15]: !rm -rf ~/.keras # clean up the model / make space for other things

'rm' is not recognized as an internal or external command, operable program or batch file.

C:\Users\mkahs\Anaconda3\lib\site-packages\keras\callbacks.py:1065: UserWarni
ng: `epsilon` argument is deprecated and will be removed, use `min_delta` ins
tead.

warnings.warn('`epsilon` argument is deprecated and '

```
In [17]: | tb model.fit generator(train gen,
                                       steps per epoch = 35,
                                       validation_data = (test_X, test_Y),
                                       epochs = 8,
                                       callbacks = callbacks list)
        Epoch 1/8
        35/35 [==========================] - 659s 19s/step - loss: 0.6049 - binar
        y accuracy: 0.6571 - val loss: 0.4442 - val binary accuracy: 0.8000
        Epoch 00001: val_loss improved from inf to 0.44423, saving model to tb_detect
        or weights.best.hdf5
        Epoch 2/8
        35/35 [================ ] - 635s 18s/step - loss: 0.5101 - binar
        y accuracy: 0.7688 - val loss: 0.4420 - val binary accuracy: 0.7800
        Epoch 00002: val loss improved from 0.44423 to 0.44196, saving model to tb de
        tector_weights.best.hdf5
        Epoch 3/8
        y_accuracy: 0.7991 - val_loss: 0.4272 - val_binary_accuracy: 0.7900
        Epoch 00003: val loss improved from 0.44196 to 0.42721, saving model to tb de
        tector_weights.best.hdf5
        Epoch 4/8
        35/35 [================ ] - 636s 18s/step - loss: 0.4316 - binar
        y_accuracy: 0.8036 - val_loss: 0.4249 - val_binary_accuracy: 0.7950
        Epoch 00004: val loss improved from 0.42721 to 0.42488, saving model to tb de
        tector weights.best.hdf5
        Epoch 5/8
        y_accuracy: 0.8446 - val_loss: 0.4316 - val_binary_accuracy: 0.7900
        Epoch 00005: val loss did not improve from 0.42488
        Epoch 6/8
        35/35 [================ ] - 635s 18s/step - loss: 0.3687 - binar
        y accuracy: 0.8446 - val loss: 0.4593 - val binary accuracy: 0.7850
        Epoch 00006: val_loss did not improve from 0.42488
        Epoch 7/8
        35/35 [================ ] - 634s 18s/step - loss: 0.3528 - binar
        y_accuracy: 0.8473 - val_loss: 0.4411 - val_binary_accuracy: 0.8050
        Epoch 00007: val loss did not improve from 0.42488
        Epoch 8/8
        35/35 [================ ] - 634s 18s/step - loss: 0.3321 - binar
        y_accuracy: 0.8589 - val_loss: 0.4731 - val_binary_accuracy: 0.8000
        Epoch 00008: val loss did not improve from 0.42488
Out[17]: <keras.callbacks.History at 0x1b5c88a3b70>
In [45]: | tb_model.load_weights(weight_path)
        tb_model.save('full_tb_model.h5')
```

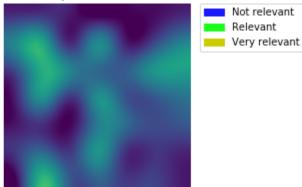
```
In [46]: # get the attention layer since it is the only one with a single output dim
for attn_layer in tb_model.layers:
    c_shape = attn_layer.get_output_shape_at(0)
    if len(c_shape)==4:
        if c_shape[-1]==1:
            print(attn_layer)
            break
```

<keras.layers.convolutional.Conv2D object at 0x000001B584F35198>

In [55]: import keras.backend as K import matplotlib.patches as mpatches rand idx = np.random.choice(range(len(test X)), size = 6) attn func = K.function(inputs = [tb model.get input at(0), K.learning phase ()],outputs = [attn layer.get output at(0)] fig, m axs = plt.subplots(len(rand idx), 2, figsize = (8, 4*len(rand idx))) [c ax.axis('off') for c ax in m axs.flatten()] for c_idx, (img_ax, attn_ax) in zip(rand_idx, m_axs): cur img = test X[c idx:(c idx+1)] attn_img = attn_func([cur_img, 0])[0] img_ax.imshow(cur_img[0,:,:,0], cmap = 'bone') $attn_ax.imshow(attn_img[0, :, :, 0], cmap = 'viridis',$ vmin = 0, vmax = 1, interpolation = 'lanczos') real label = test Y[c idx] img ax.set title('Tb-Class:%s' % (real label)) pred_confidence = tb_model.predict(cur_img)[0] cmap = $\{1:[0.1,0.1,1.0,t],2:[0.1,1.0,0.1,t],3:[0.8,0.8,0.0,t]\}$ labels = {1:'Not relevant',2:'Relevant',3:'Very relevant'} attn ax.set title('Attention Map-Prediction:%s' % (pred confidence[0])) patches =[mpatches.Patch(color=cmap[i],label=labels[i]) for i in cmap] attn ax.legend(handles=patches, bbox to anchor=(1.05, 1), loc=2, borderaxe spad=0.) fig.savefig('attention map.png', dpi = 300)

Tb-Class:1

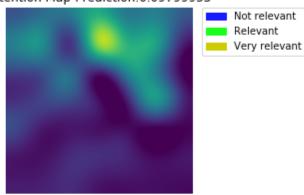
Attention Map-Prediction:0.10857909



Tb-Class:0



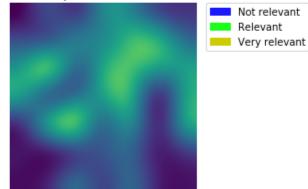
Attention Map-Prediction: 0.09799953



Tb-Class:1



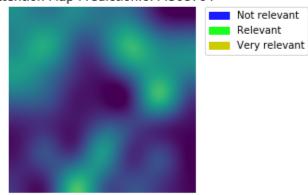
Attention Map-Prediction: 0.116258286



Tb-Class:1



Attention Map-Prediction: 0.44508764



Tb-Class:0



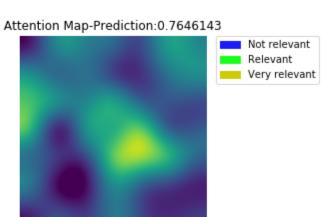
Attention Map-Prediction:0.82040644





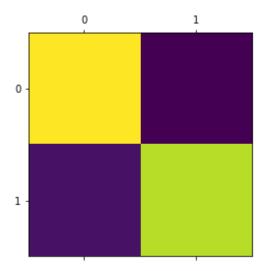
Tb-Class:1





```
In [22]: from sklearn.metrics import classification_report, confusion_matrix
    plt.matshow(confusion_matrix(test_Y, pred_Y>0.5))
    print(classification_report(test_Y, pred_Y>0.5, target_names = ['Healthy', 'T B']))
```

	precision	recall	f1-score	support
Healthy	0.79	0.81	0.80	102
TB	0.80	0.78	0.79	98
micro avg	0.80	0.80	0.80	200
macro avg	0.80	0.79	0.79	200
weighted avg	0.80	0.80	0.79	200



```
In [31]: from sklearn.metrics import roc_curve, roc_auc_score
    fpr, tpr, _ = roc_curve(test_Y, pred_Y)
        fig, ax1 = plt.subplots(1,1, figsize = (5, 5), dpi = 250)
        ax1.plot(fpr, tpr, 'b.-', label = 'Model (AUC:%2.2f)' % roc_auc_score(test_Y, pred_Y))
        ax1.plot(fpr, fpr, 'k-', label = 'Random Guessing')
        ax1.legend(loc = 4)
        ax1.set_xlabel('False Positive Rate')
        ax1.set_ylabel('True Positive Rate');
        fig.savefig('roc.pdf')
```

