

Duke Classification: with “2D-picture”

Yuhui Cao

20220816 Tue

Roadmap

- Part 1. Convert Threshold to “2D picture”
- Part 2. Shallow Neural Network (SNN)
 - Train on Duke (400)
 - Validate on Duke (100)
 - Test on UAB (301)
- Part 3: Transfer study of **VGG16**
 - Train on Duke (400)
 - Validate on Duke (100)
 - Test on UAB (301)1a

Method 1 - pad with 0

Method 2 – Up-sample smaller img

Part 1. Convert Threshold to “2D picture”

Test Pattern (indexed)										---- index of VF columns				
Right Eye										----- index of 10X10 array				
		1	2	3	4	5	6	7	8	9	10			
S1		0	1	2	1	2	3	4	7	8	9		27	
S2		10		5	6	7	8	9	10		19		21	
S3		20	11	12	13	14	15	16	17	18	29		15	
S4		19	20	21	22	23	24	25	26	27	28		9	
S5		29	30	31	32	33	34	35	36	37	38		3	
S6		39	40	41	42	43	44	45	46	47	48		-3	
S7		49	50	51	52	53	54	55	56	57	58		-9	
S8		70	59	60	61	62	63	64	65	66	79		-15	
S9		80		67	68	69	70	71	72		89		-21	
S10		90	91	92	73	74	75	76	97	98	99		-27	
		-27	-21	-15	-9	-3	3	9	15	21	27			

```
array([[ 0.,  0.,  0., nan, nan, nan, nan,  0.,  0.,  0.],
       [ 0.,  0., nan, 28., 28., 26., 29., nan,  0.,  0.],
       [ 0., nan, 25., 29., 31., 29., 29., 30., nan,  0.],
       [nan, 27., 30., 32., 31., 32., 31., 30., 27., nan],
       [24., 30., 31., 32., 33., 33., 34., 27., 27., nan],
       [22., 27., 30., 32., 34., 32., 32., 16., 27., nan],
       [nan, 21., 23., 24., 30., 33., 31., 27., 25., nan],
       [ 0., nan, 16., 23., 29., 31., 27., 26., nan,  0.],
       [ 0.,  0., nan, 11., 24., 27., 29., nan,  0.,  0.],
       [ 0.,  0.,  0., nan, nan, nan, nan,  0.,  0.,  0.]])
```

Yuhui:

- Fill the rest of (100-76 =) 24 locations with 0?
- Fill NaN of 24-2 with 0?

Part 1. Convert Threshold to “2D picture”

An example of 24-2

```
array([[ 0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.],
       [ 0.,  0.,  0., 28., 28., 26., 29.,  0.,  0.,  0.],
       [ 0.,  0., 25., 29., 31., 29., 29., 30.,  0.,  0.],
       [ 0., 27., 30., 32., 31., 32., 31., 30., 27.,  0.],
       [24., 30., 31., 32., 33., 33., 34., 27., 27.,  0.],
       [22., 27., 30., 32., 34., 32., 32., 16., 27.,  0.],
       [ 0., 21., 23., 24., 30., 33., 31., 27., 25.,  0.],
       [ 0.,  0., 16., 23., 29., 31., 27., 26.,  0.,  0.],
       [ 0.,  0.,  0., 11., 24., 27., 29.,  0.,  0.,  0.],
       [ 0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.,  0.]])
```

An example of 30-2

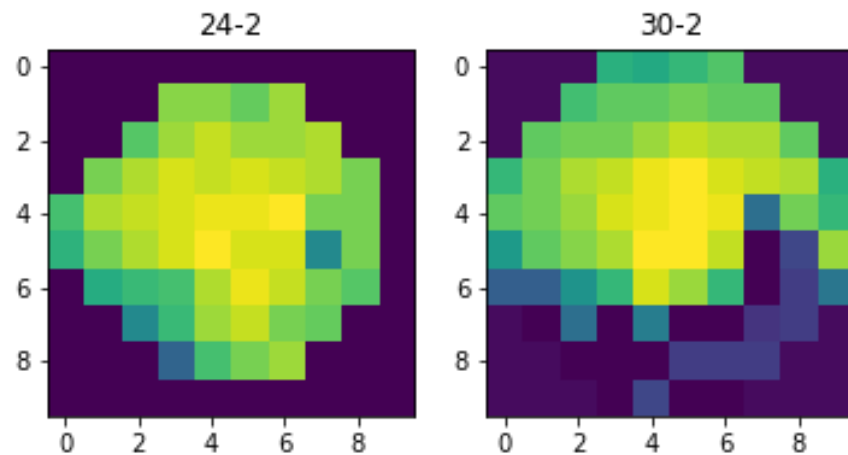
```
array([[ 0.,  0.,  0., 20., 19., 21., 23.,  0.,  0.,  0.],
       [ 0.,  0., 22., 24., 24., 25., 24., 24.,  0.,  0.],
       [ 0., 24., 25., 25., 27., 29., 28., 28., 24.,  0.],
       [21., 25., 28., 29., 31., 32., 30., 29., 28., 20.],
       [24., 25., 27., 30., 31., 32., 31., 11., 25., 21.],
       [17., 24., 26., 28., 32., 32., 29., -1.,  6., 27.],
       [ 9.,  9., 16., 21., 30., 27., 21., -1.,  5., 12.],
       [ 0., -1., 11., -1., 13., -1., -1.,  4.,  5.,  0.],
       [ 0.,  0., -1., -1., -1.,  5.,  5.,  5.,  0.,  0.],
       [ 0.,  0.,  0., -1.,  6., -1., -1.,  0.,  0.,  0.]])
```

500 cases:

- The number of 30-2 pattern is: 14
- The number of 24-2 pattern is: 486

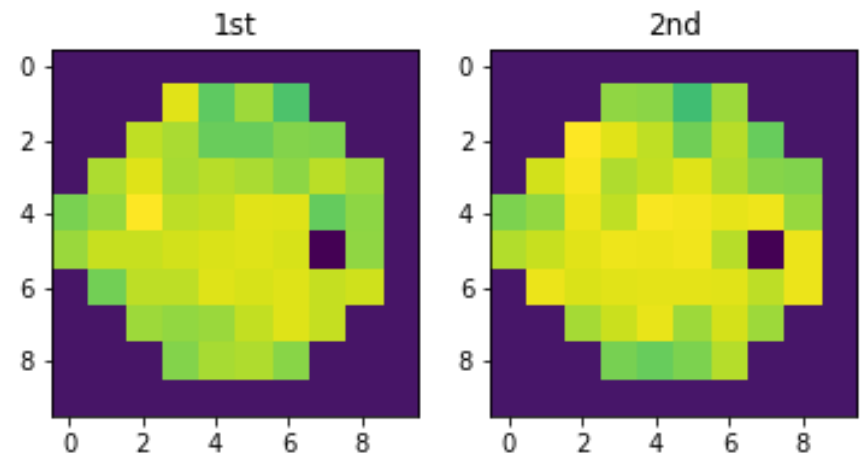
Part 1. Convert Threshold to “2D picture”

Visualization of examples of 24-2 vs. 30-2 (Duke)



500 Duke

Visualization of fist 2 examples (UAB)



301 UAB

Part 2. Shallow Neural Network (SNN)

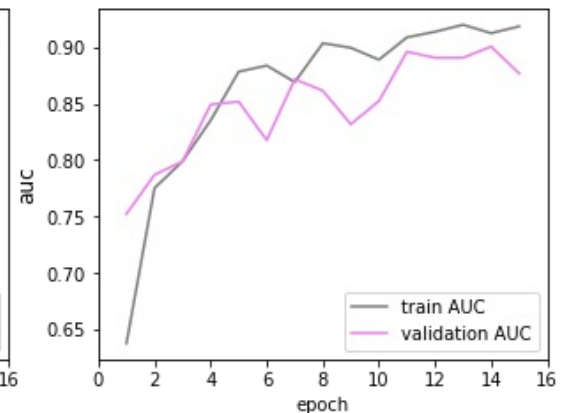
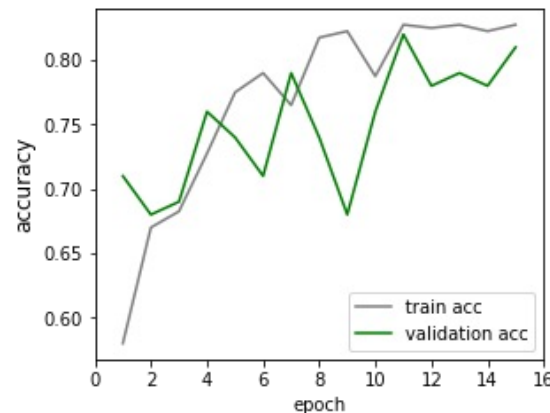
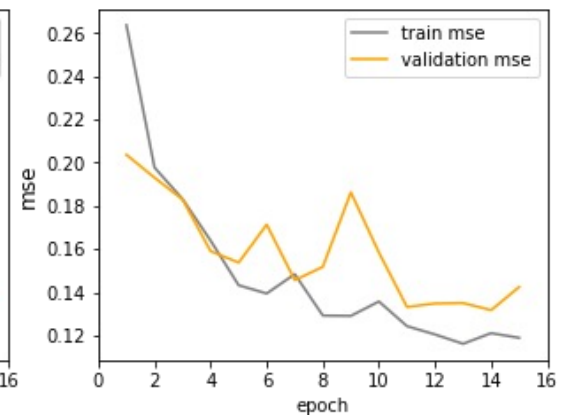
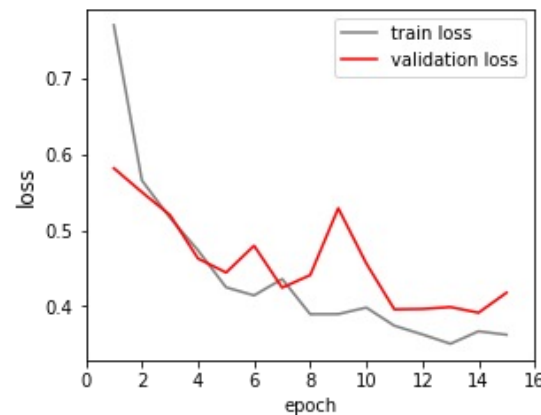
Classification on normal vs. abnormal

SNN 1

```
# create model
model = Sequential()
model.add(Conv2D(16, (2, 2), strides=(1, 1), activation='relu',
                input_shape=(10, 10, 1)))
model.add(MaxPooling2D(pool_size=(2, 2), strides=(2, 2)))

model.add(Flatten())
model.add(Dense(20, activation='relu'))
model.add(Dense(num_classes, activation='softmax'))
```

epochs = 15
batch_size=30



Part 2. Shallow Neural Network (SNN)

Classification on normal vs. abnormal

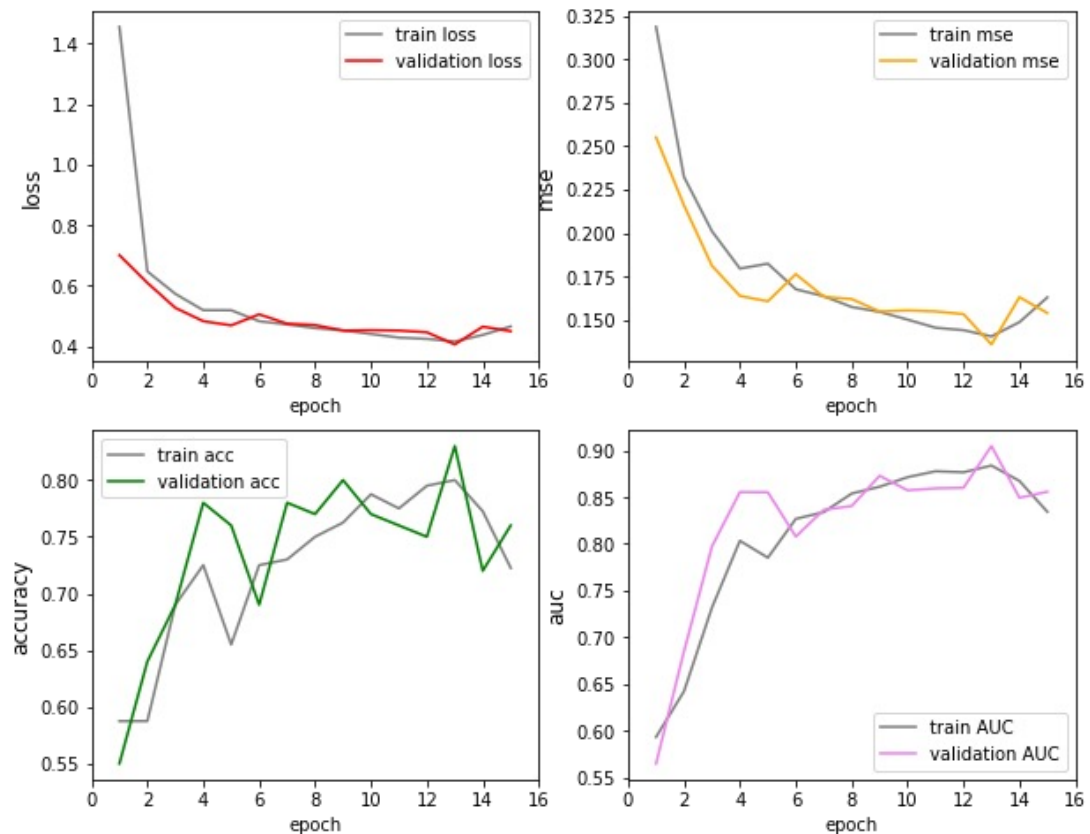
SNN 2

```
# create model
model = Sequential()
model.add(Conv2D(16, (2, 2), strides=(1, 1), activation='relu',
                input_shape=(10, 10, 1)))
model.add(MaxPooling2D(pool_size=(2, 2), strides=(2, 2)))

model.add(Conv2D(8, (2, 2), activation='relu'))
#model.add(MaxPooling2D(pool_size=(2, 2), strides=(2, 2)))

model.add(Flatten())
model.add(Dense(20, activation='relu'))
model.add(Dense(num_classes, activation='softmax'))
```

epochs = 15
batch_size=30



Part 2. Shallow Neural Network (SNN)

Classification on normal vs. abnormal

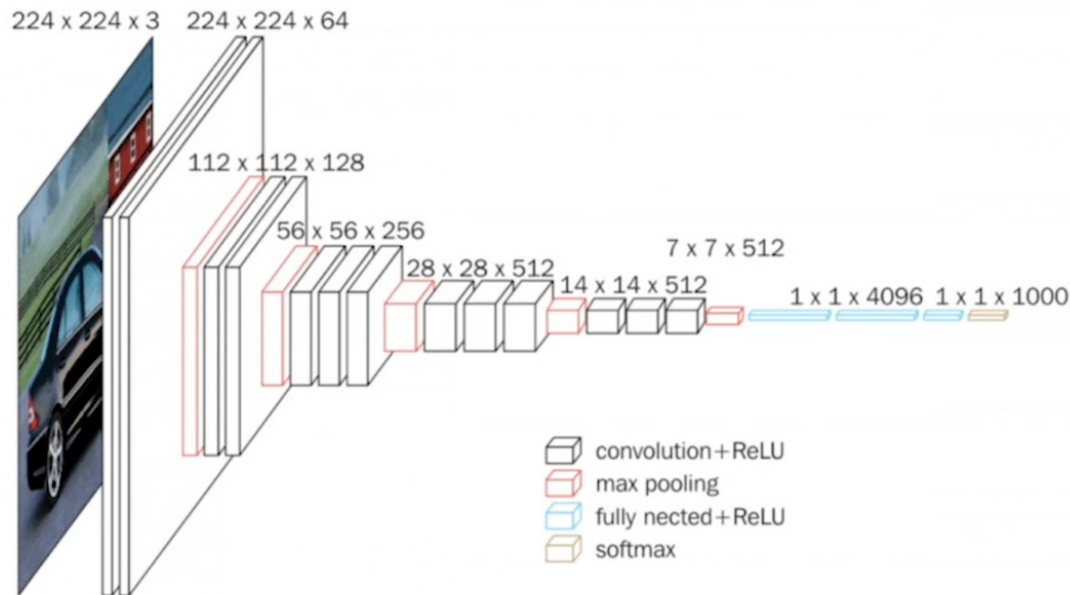
	validation (M1)	test (M1)	validation (M2)	test (M2)
loss	0.417800	0.582213	0.450455	0.563872
mse	0.142439	0.212142	0.153874	0.201238
accuracy	0.810000	0.637874	0.760000	0.657807
auc	0.876650	0.756371	0.855800	0.742376

Validation on Duke (100)
Test on UAB (301)

```
model1.save("SNN_1.h5")  
model2.save("SNN_2.h5")
```


Part 3: Transfer study of **VGG16**

CNNs



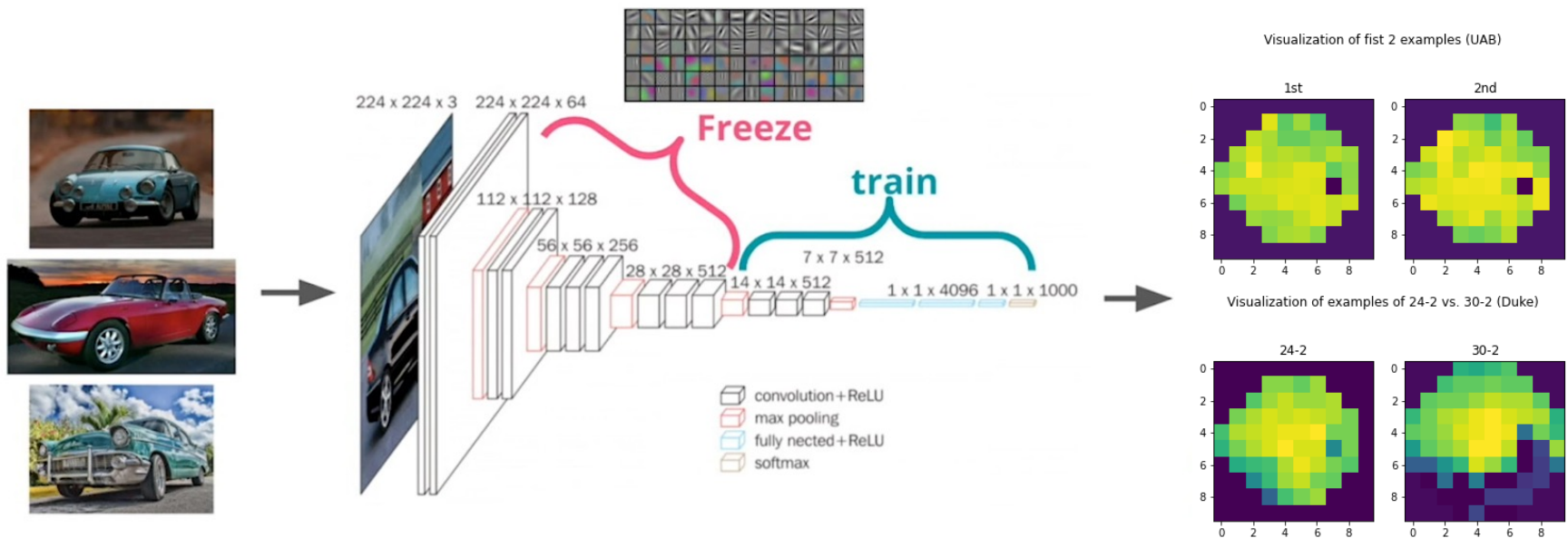
Yuhui strategies:

1. Should I start from 14X14X512 layer? (can very easily pad 10X10 to 14X14)
2. Resize 10X0 to 224X224 so that can use earlier layer? (**preferred**)

Part 3: Transfer study of **VGG16**

Fine-tuning

Repurposing pre-trained architectures



Part 3: Transfer study of **VGG16**

Dimension construction: resize to 224 X 224

1. 10 X 10

2. 224 X 224

3. 224 X 224 X 3

- repeat thr 3 times

- future: use thr, TD, PD

4. 500 X 224 X 224

Method 1 - pad with 0

Method 2 – Up-sample smaller img

Method 3 – Interpolating smaller img

Result 0. Pad with NaN is not better ~

Epoch 1/5

16/16 [=====] - 174s 11s/step - loss: nan - binary_accuracy: 0.630
0 - val_loss: nan - val_binary_accuracy: 0.8472

Epoch 2/5

16/16 [=====] - 188s 12s/step - loss: nan - binary_accuracy: 0.630
0 - val_loss: nan - val_binary_accuracy: 0.8472

Epoch 3/5

16/16 [=====] - 193s 12s/step - loss: nan - binary_accuracy: 0.630
0 - val_loss: nan - val_binary_accuracy: 0.8472

Epoch 4/5

16/16 [=====] - 177s 11s/step - loss: nan - binary_accuracy: 0.630
0 - val_loss: nan - val_binary_accuracy: 0.8472

Epoch 5/5

16/16 [=====] - 184s 12s/step - loss: nan - binary_accuracy: 0.630
0 - val_loss: nan - val_binary_accuracy: 0.8472

Result 1. Method 1 - pad with 0

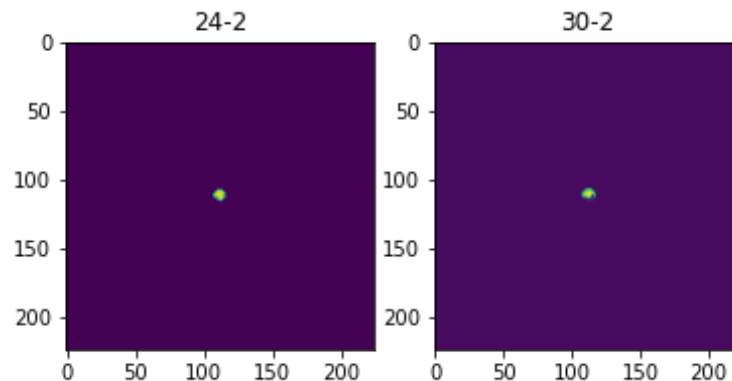
Model 1: a simple sequential model using only the VGG16 architecture

Model 2: add a few more dense layers

Model 3: add dropout and another fully connected layer

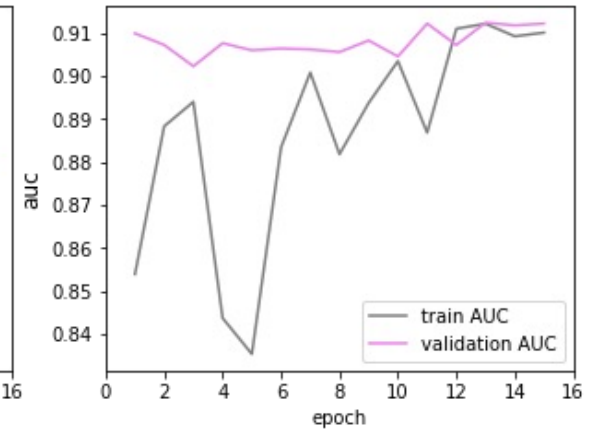
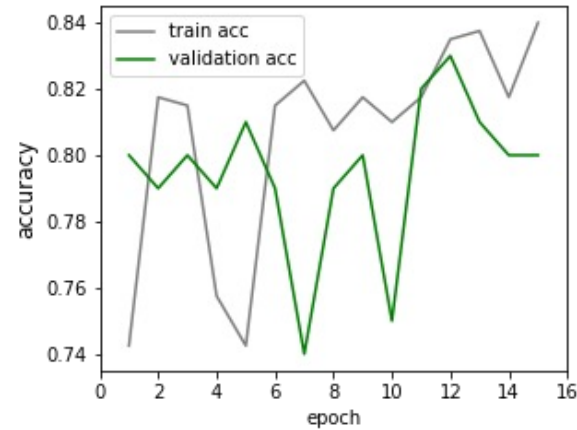
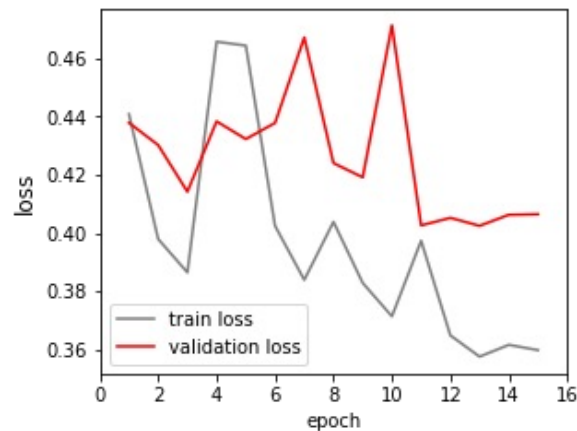
Visualization of examples of 24-2 vs. 30-2 (Duke)

Image Visualization



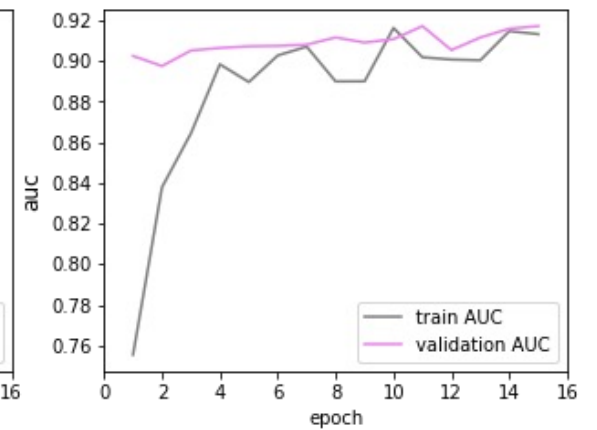
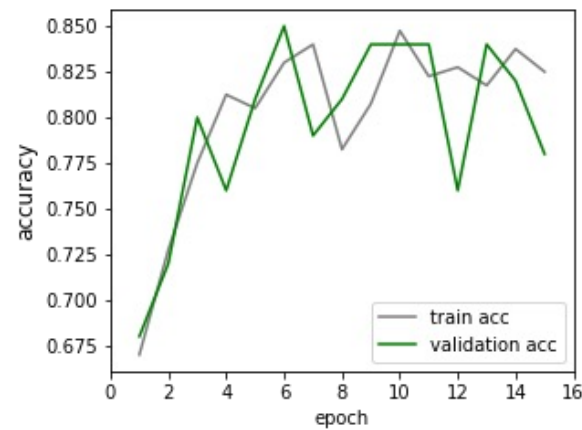
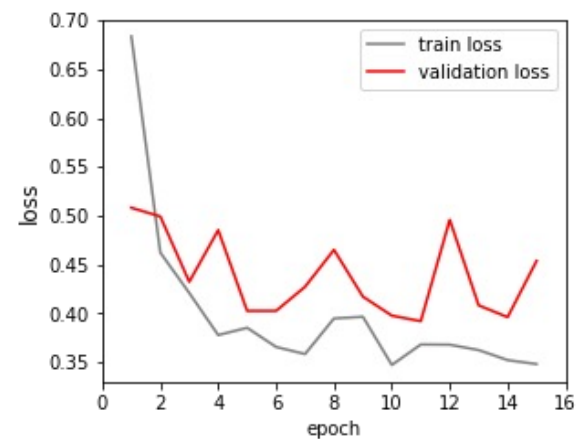
Part 3: Transfer study of VGG16 — model 1

epochs = 15
batch_size = 32 (default)



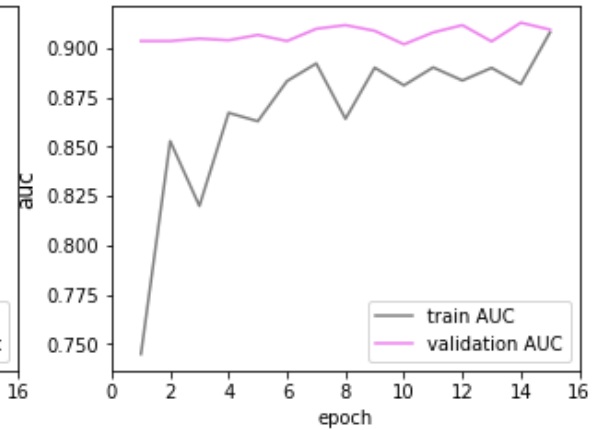
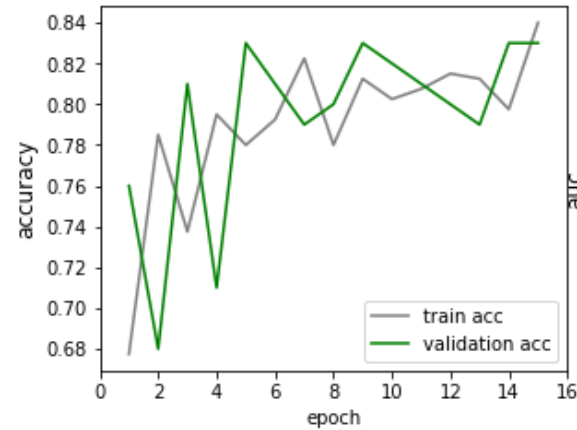
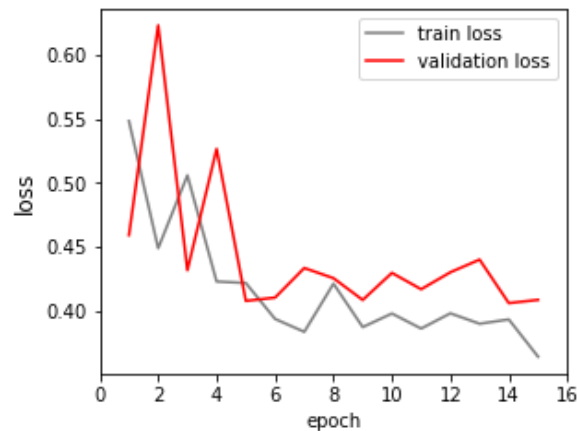
Part 3: Transfer study of VGG16 — model 2

epochs = 15
batch_size = 32 (default)



Part 3: Transfer study of VGG16 — model 3

epochs = 15
batch_size = 32 (default)



Test: Method 1 - pad with 0 (on UAB)

	validation (M1)	test (M1)	validation (M2)	test (M2)	validation (M3)	test (M3)
loss	0.406474	0.590327	0.453722	0.301917	0.408702	0.412213
accuracy	0.800000	0.651163	0.780000	0.857143	0.830000	0.777409
auc	0.912154	0.902899	0.917114	0.911978	0.909260	0.900384

Validation on Duke (100)
Test on UAB (301)

```
model1.save("VGG1.h5")  
model2.save("VGG2.h5")  
model3.save("VGG3.h5")
```

Result 2. Method 2 – Up-sample smaller img

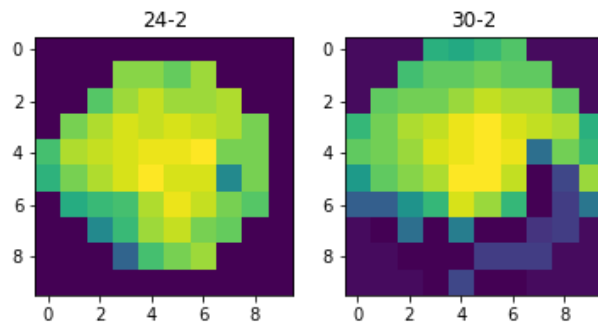
Model 1: a simple sequential model using only the VGG16 architecture

Model 2: add a few more dense layers

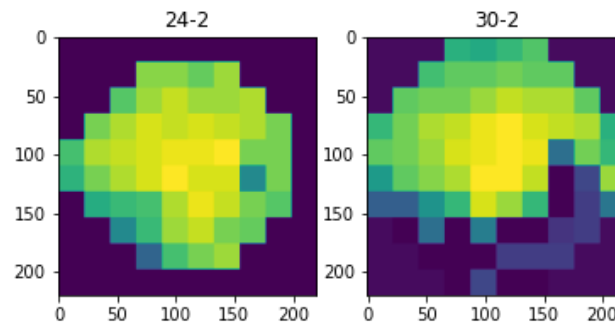
Model 3: add dropout and another fully connected layer

Image Visualization

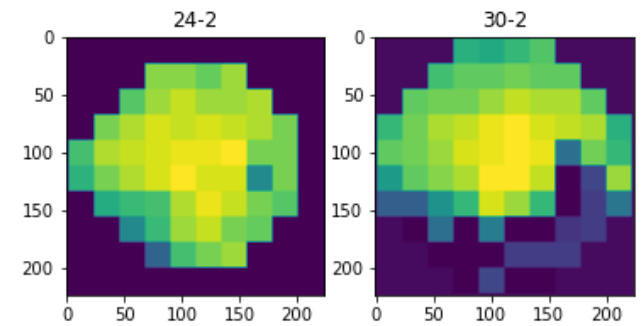
1. Visualization of examples of 24-2 vs. 30-2 (Duke) [10X10]



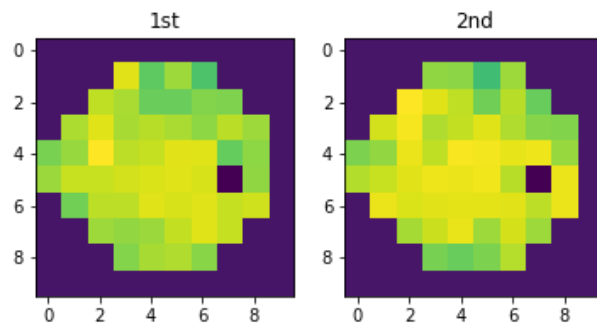
2. Visualization of examples of 24-2 vs. 30-2 (Duke) [220X220]



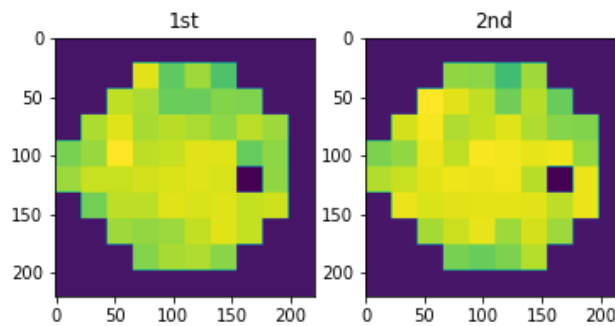
3. Visualization of examples of 24-2 vs. 30-2 (Duke) [224X224]



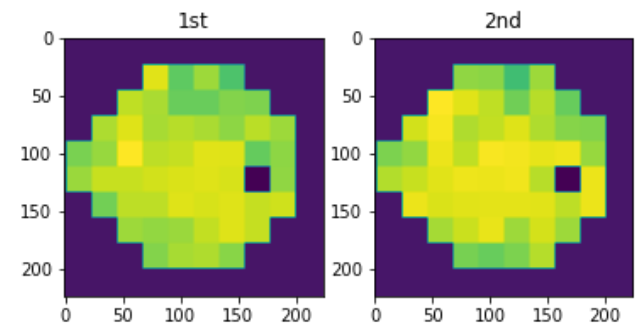
1. Visualization of fist 2 examples (UAB) [10X10]



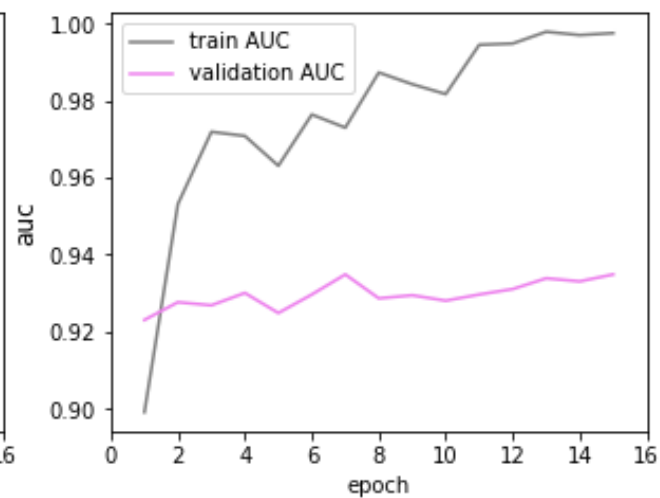
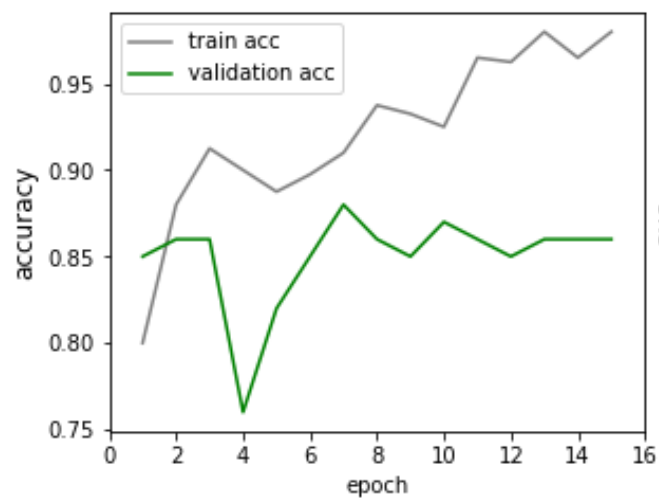
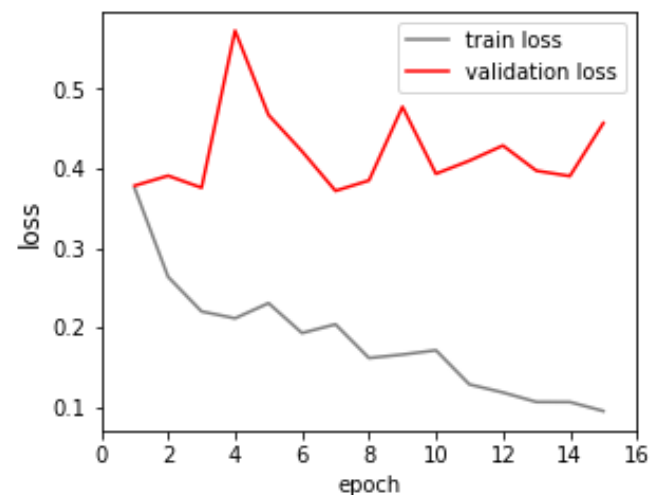
2. Visualization of fist 2 examples (UAB) [220X220]

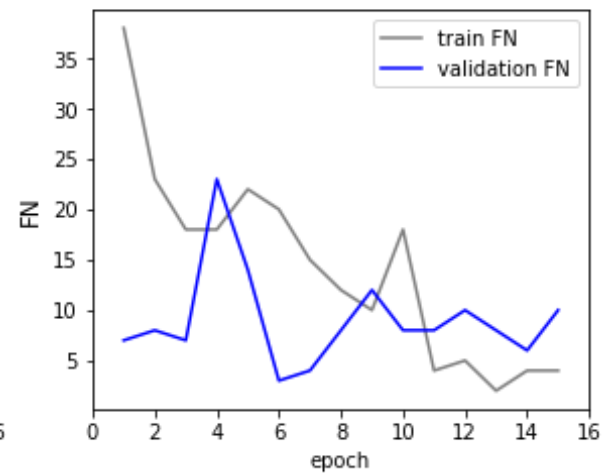
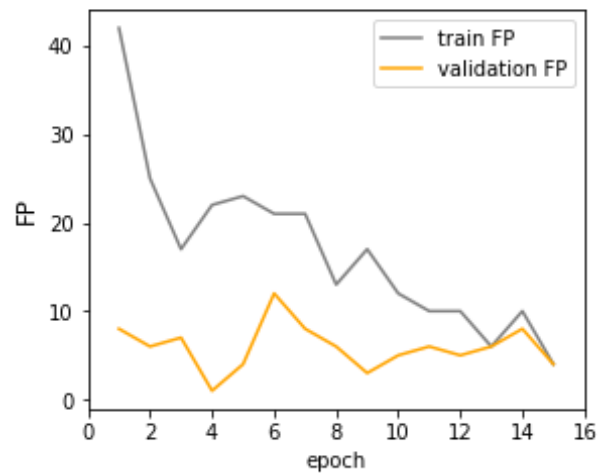
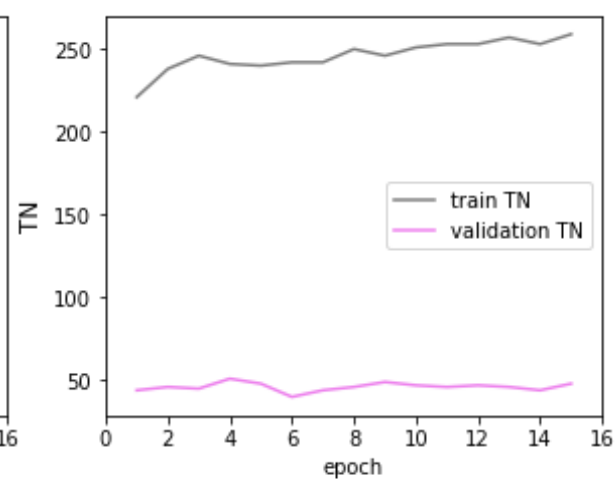
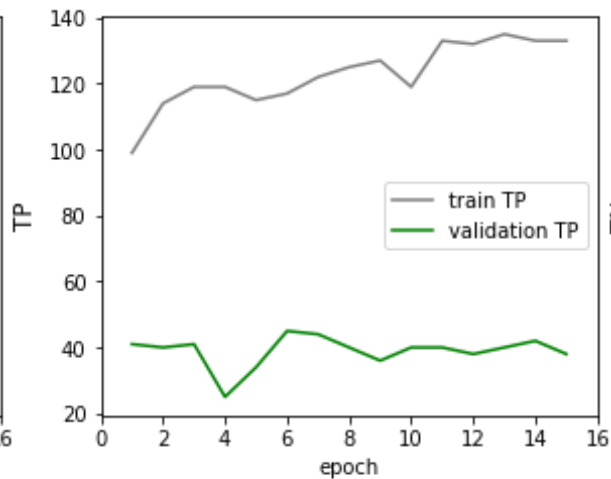
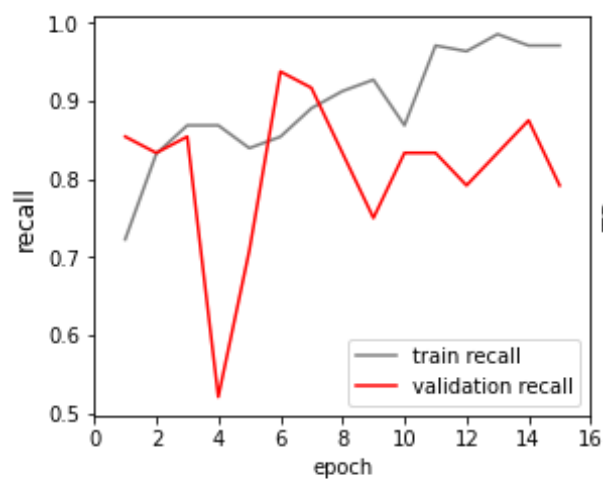


3. Visualization of fist 2 examples (UAB) [224X224]



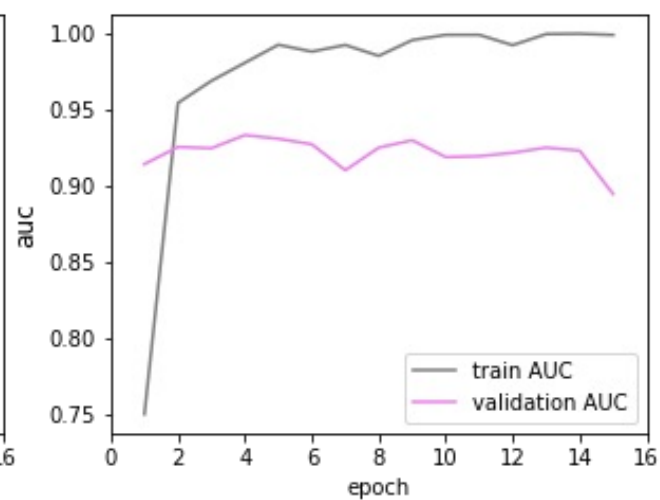
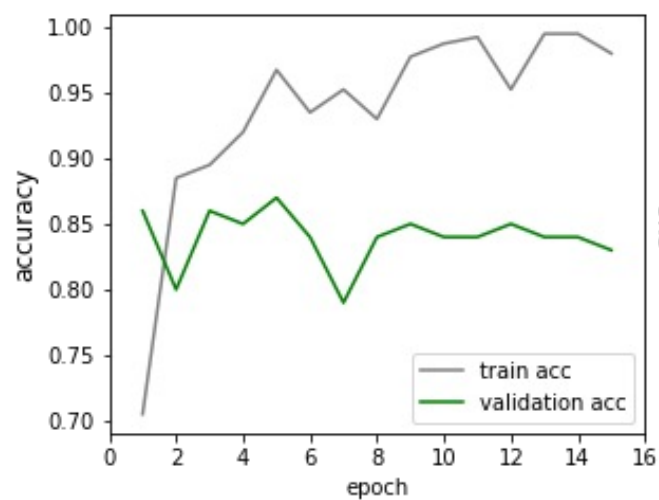
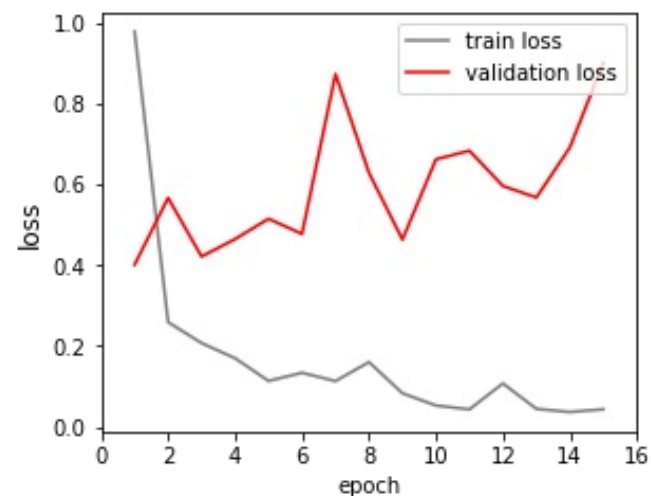
Model 1

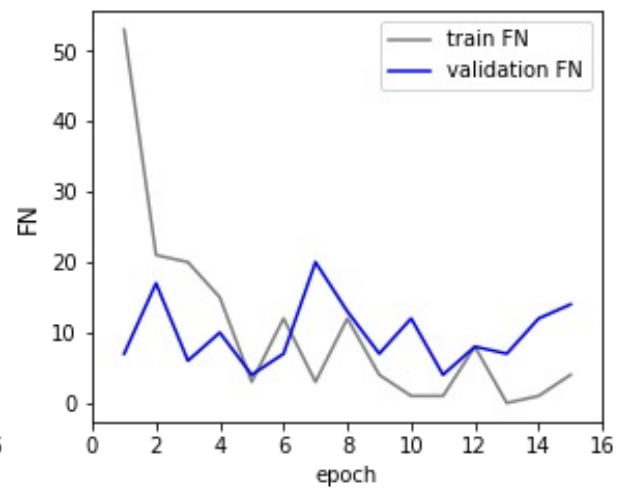
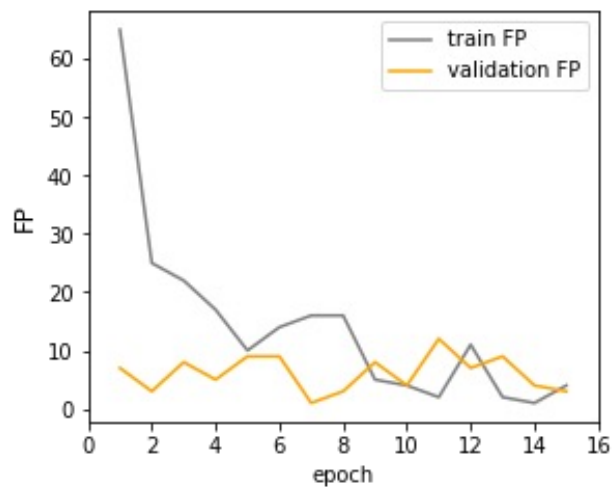
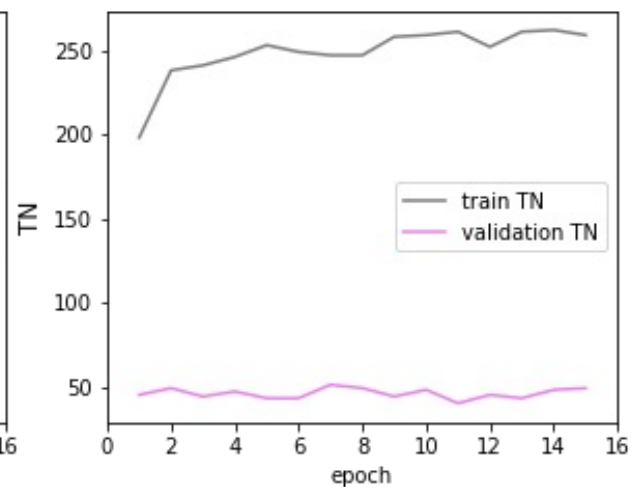
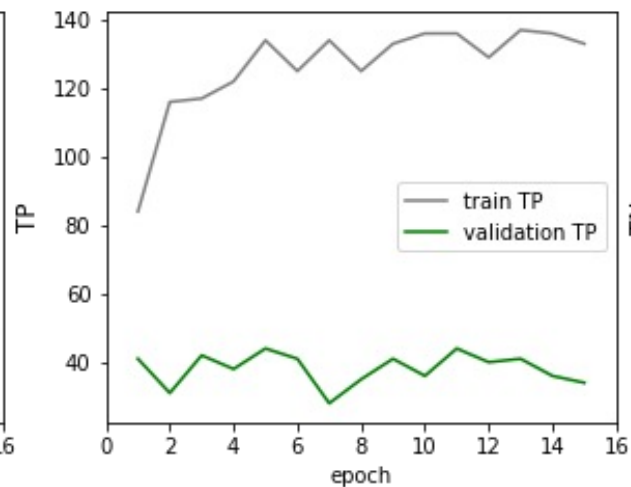
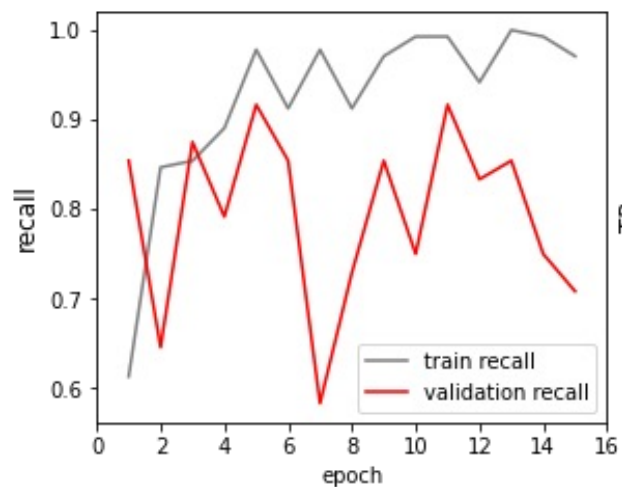




Model 1

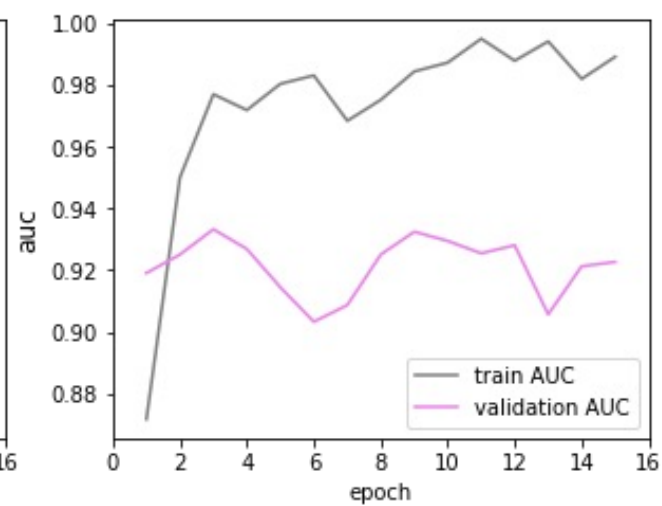
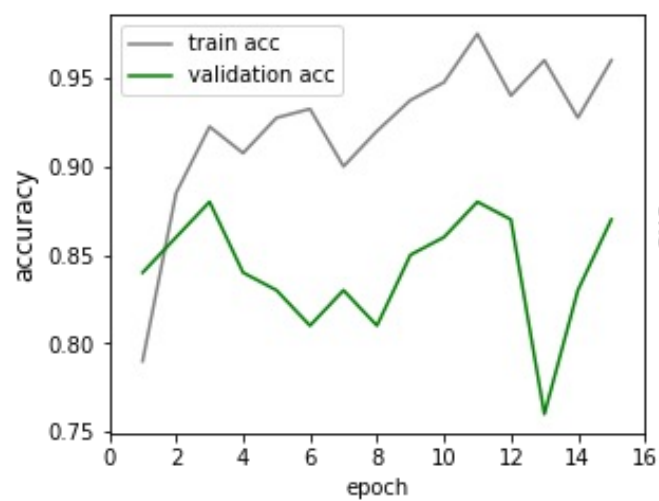
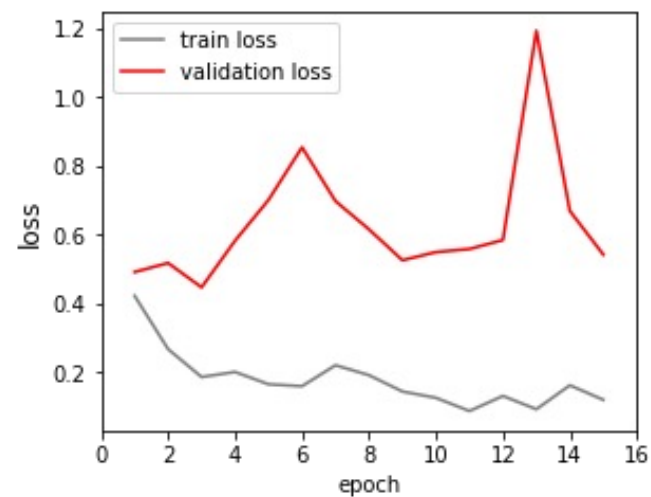
Model 2

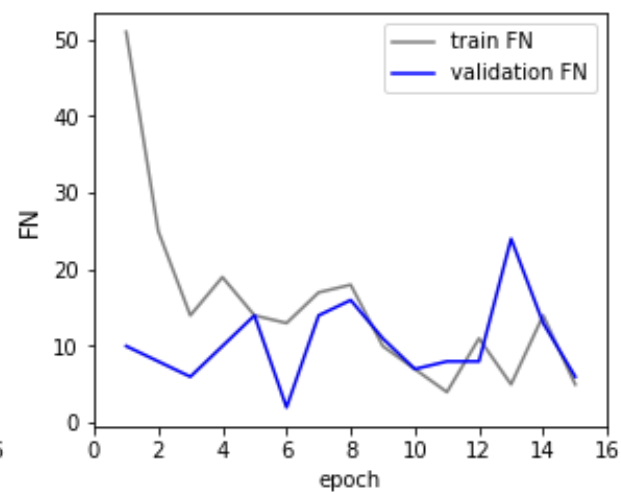
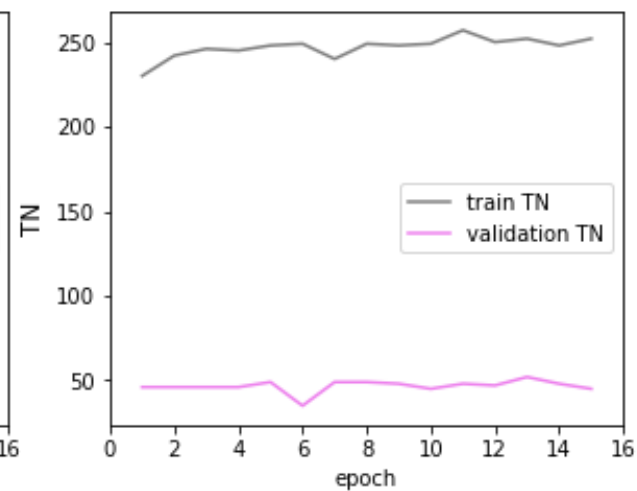
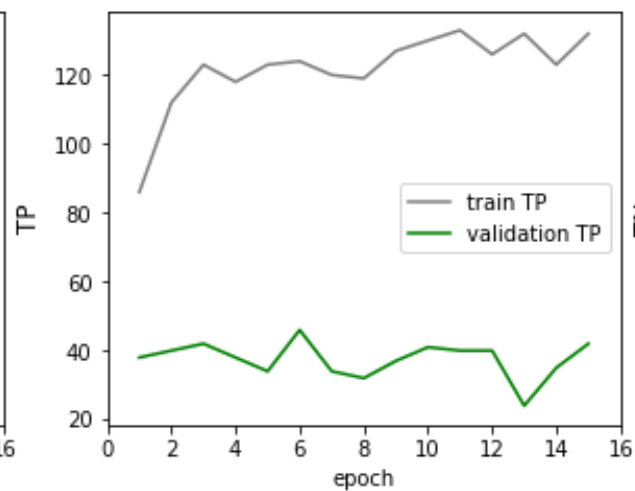
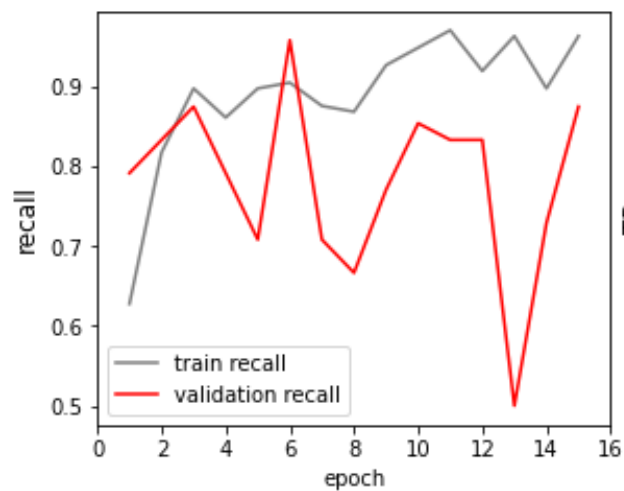




Model 2

Model 3





Model 3

Model 1

**Train
(Duke 400)**

	TP	TN	FP	FN	sens	spec
0	99.0	221.0	42.0	38.0	0.722628	0.840304
1	114.0	238.0	25.0	23.0	0.832117	0.904943
2	119.0	246.0	17.0	18.0	0.868613	0.935361
3	119.0	241.0	22.0	18.0	0.868613	0.916350
4	115.0	240.0	23.0	22.0	0.839416	0.912548
5	117.0	242.0	21.0	20.0	0.854015	0.920152
6	122.0	242.0	21.0	15.0	0.890511	0.920152
7	125.0	250.0	13.0	12.0	0.912409	0.950570
8	127.0	246.0	17.0	10.0	0.927007	0.935361
9	119.0	251.0	12.0	18.0	0.868613	0.954373
10	133.0	253.0	10.0	4.0	0.970803	0.961977
11	132.0	253.0	10.0	5.0	0.963504	0.961977
12	135.0	257.0	6.0	2.0	0.985401	0.977186
13	133.0	253.0	10.0	4.0	0.970803	0.961977
14	133.0	259.0	4.0	4.0	0.970803	0.984791

**Validation
(Duke 100)**

	TP_val	TN_val	FP_val	FN_val	sens_val	spec_val
0	41.0	44.0	8.0	7.0	0.854167	0.846154
1	40.0	46.0	6.0	8.0	0.833333	0.884615
2	41.0	45.0	7.0	7.0	0.854167	0.865385
3	25.0	51.0	1.0	23.0	0.520833	0.980769
4	34.0	48.0	4.0	14.0	0.708333	0.923077
5	45.0	40.0	12.0	3.0	0.937500	0.769231
6	44.0	44.0	8.0	4.0	0.916667	0.846154
7	40.0	46.0	6.0	8.0	0.833333	0.884615
8	36.0	49.0	3.0	12.0	0.750000	0.942308
9	40.0	47.0	5.0	8.0	0.833333	0.903846
10	40.0	46.0	6.0	8.0	0.833333	0.884615
11	38.0	47.0	5.0	10.0	0.791667	0.903846
12	40.0	46.0	6.0	8.0	0.833333	0.884615
13	42.0	44.0	8.0	6.0	0.875000	0.846154
14	38.0	48.0	4.0	10.0	0.791667	0.923077

sensitivity, recall, hit rate, or true positive rate (TPR)

$$\text{TPR} = \frac{\text{TP}}{\text{P}} = \frac{\text{TP}}{\text{TP} + \text{FN}} = 1 - \text{FNR}$$

specificity, selectivity or true negative rate (TNR)

$$\text{TNR} = \frac{\text{TN}}{\text{N}} = \frac{\text{TN}}{\text{TN} + \text{FP}} = 1 - \text{FPR}$$

Validation:

- Specificity was OK.
- Sensitivity was bad

Solution:

- increase power
- Increase sample size

Test: Method 2 – Up-sample (on UAB)

	validation (M1)	test (M1)	validation (M2)	test (M2)	validation (M3)	test (M3)
loss	0.296763	0.378864	0.199720	0.395243	0.244206	0.774924
accuracy	0.940000	0.787375	0.970000	0.797342	0.930000	0.730897
auc	0.980035	0.917008	0.979167	0.902856	0.978733	0.909548
recall	0.916667	0.913043	0.944444	0.869565	0.972222	0.934783
TP	33.000000	42.000000	34.000000	40.000000	35.000000	43.000000
TN	61.000000	195.000000	63.000000	200.000000	58.000000	177.000000
FP	3.000000	60.000000	1.000000	55.000000	6.000000	78.000000
FN	3.000000	4.000000	2.000000	6.000000	1.000000	3.000000

Validation on Duke (100)
Test on UAB (301)

```
model1.save("VGG1_upsampling.h5")  
model2.save("VGG2_upsampling.h5")  
model3.save("VGG3_upsampling.h5")
```

Result 3. Method 3 – Interpolate an image

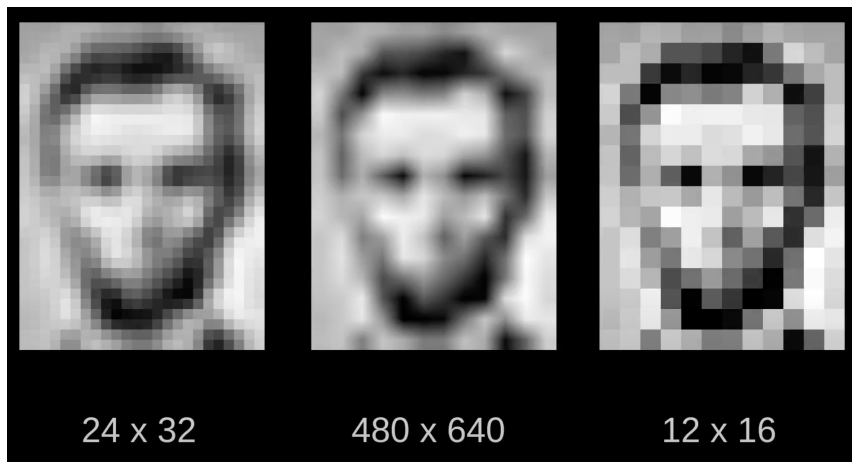
Model 1: a simple sequential model using only the VGG16 architecture

Model 2: add a few more dense layers

Model 3: add dropout and another fully connected layer

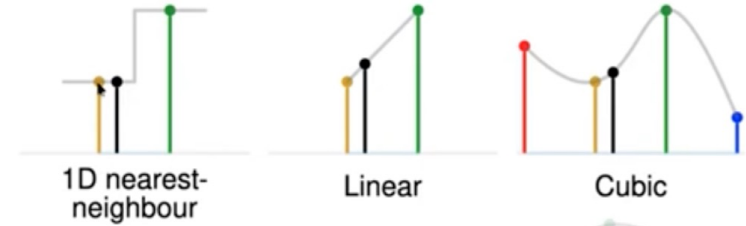
Image Processing

Bilinear Interpolation & Resampling



Types of interpolation

1-D:



2-D:

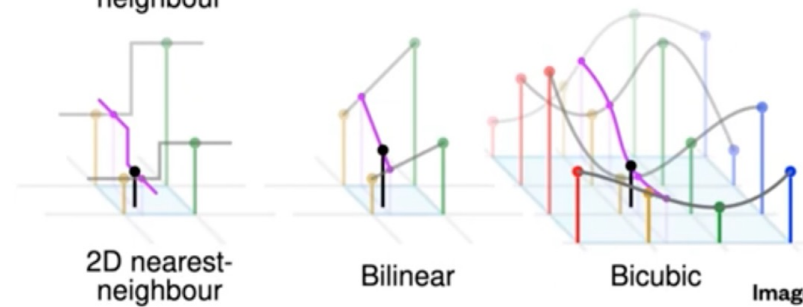
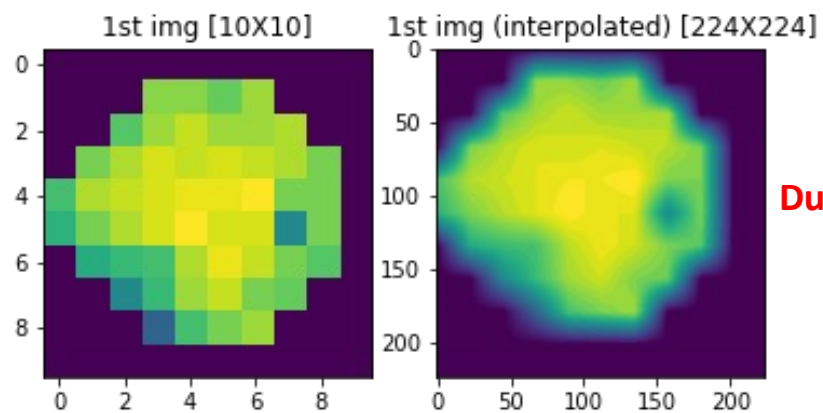
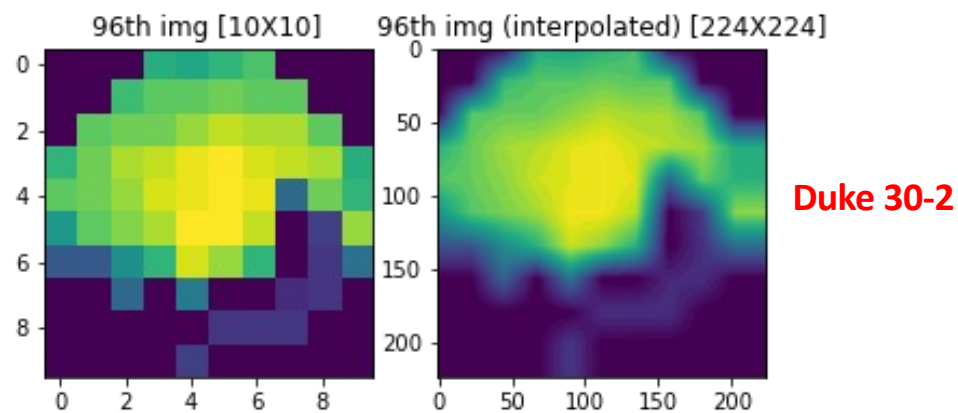


Image credit: C.M.G. Lee

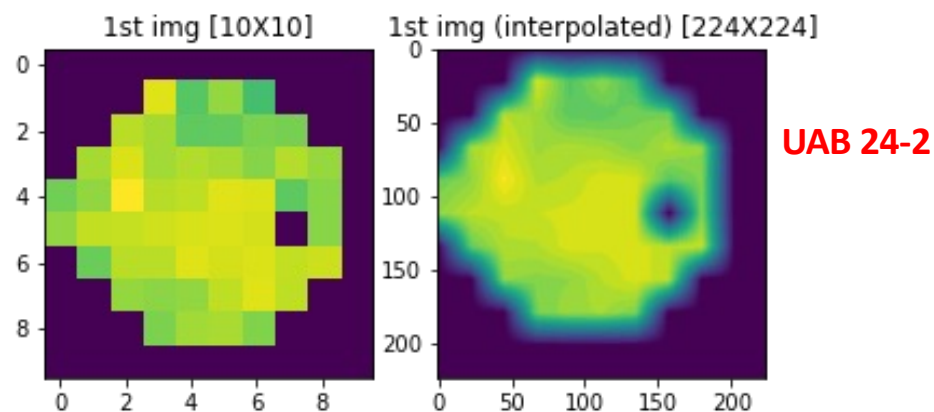
1. Visualization of Duke



1. Visualization of Duke



1. Visualization of UAB

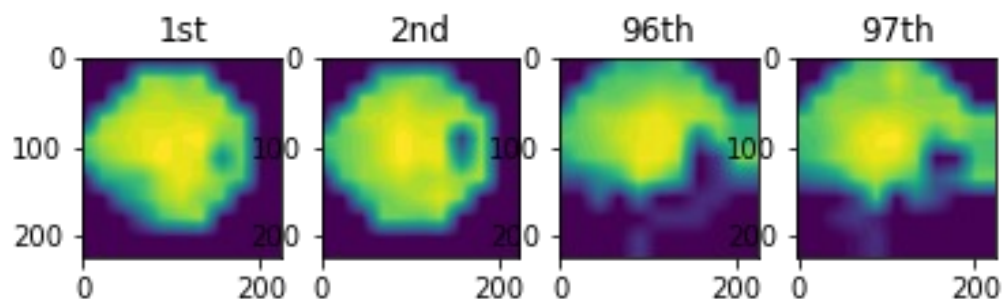


Visualization:

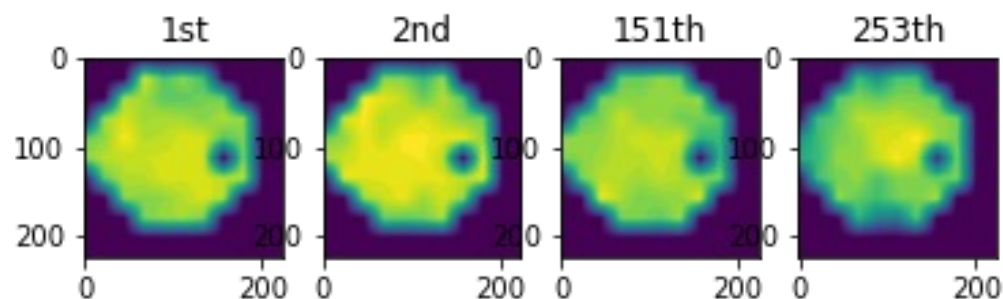
- Original vs. Interpolated

Visualization: Interpolated

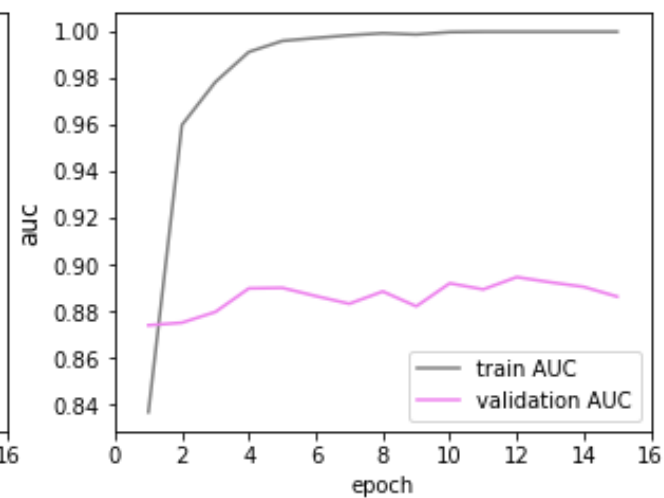
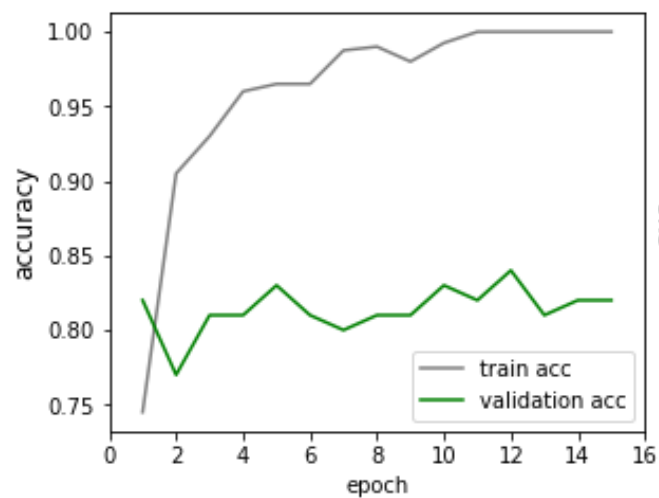
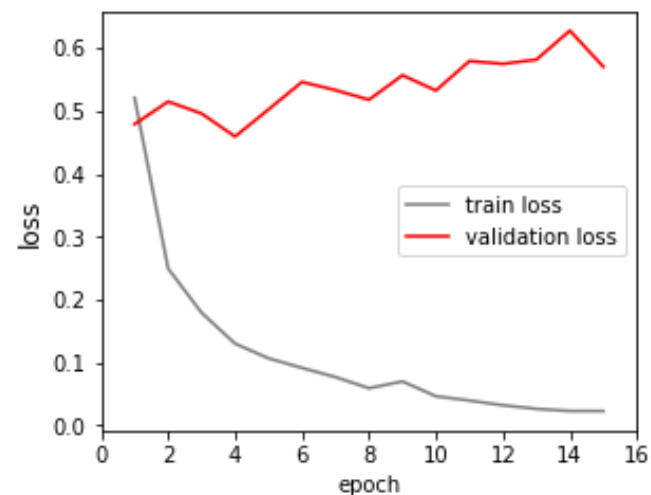
1. Visualization of DUKE [224X24]

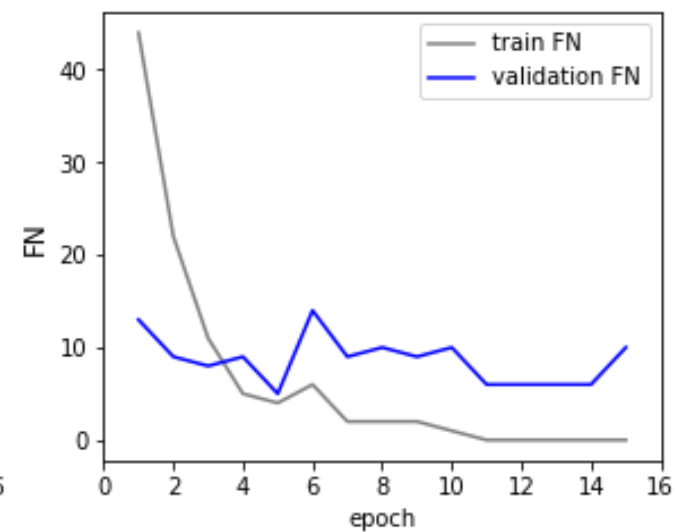
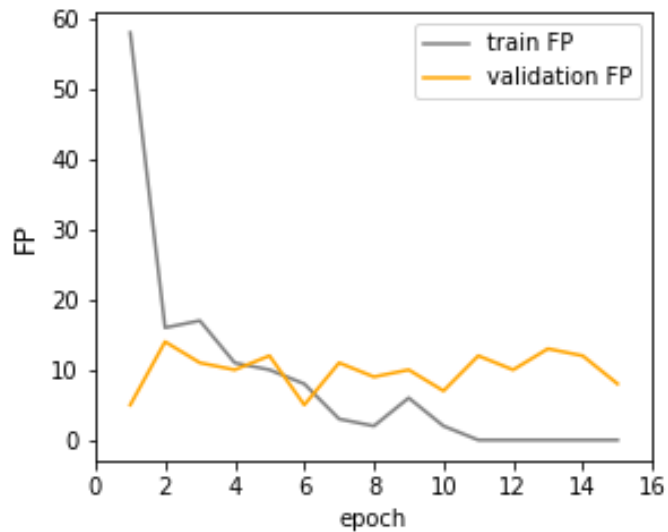
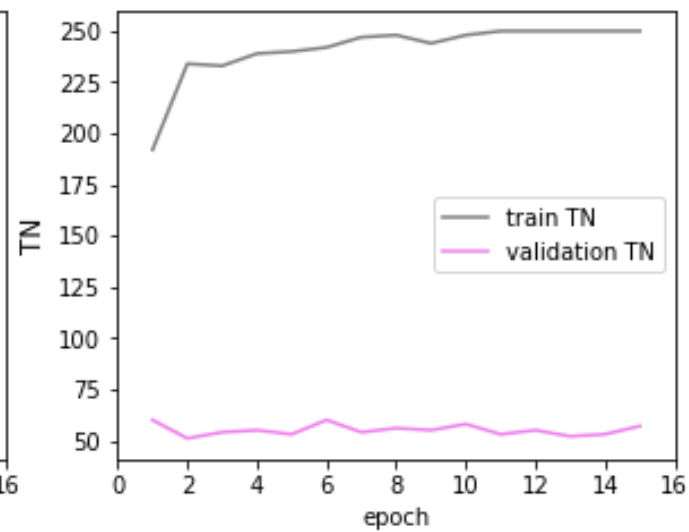
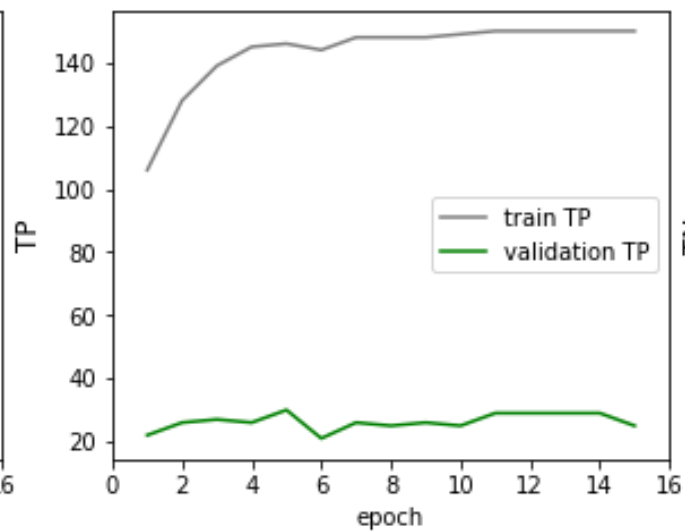
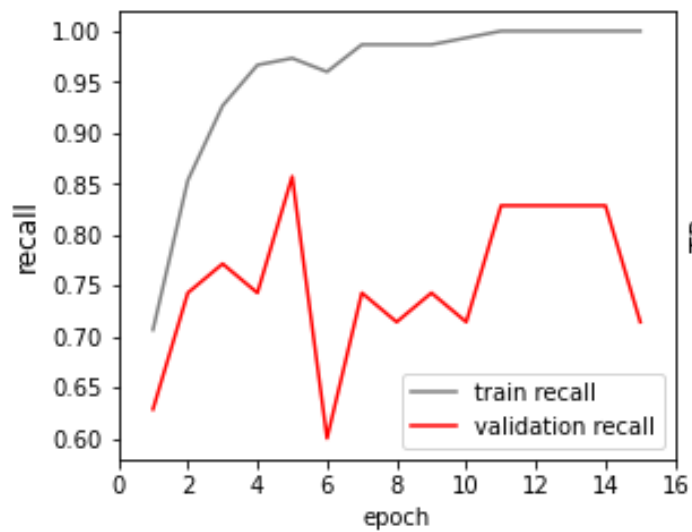


1. Visualization of UAB [224X24]



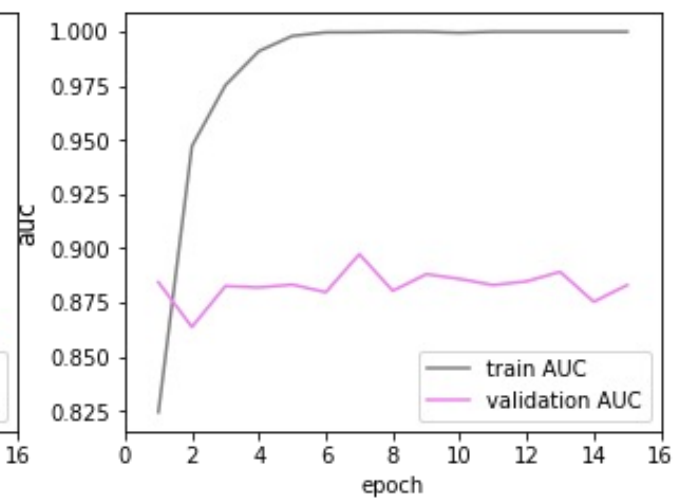
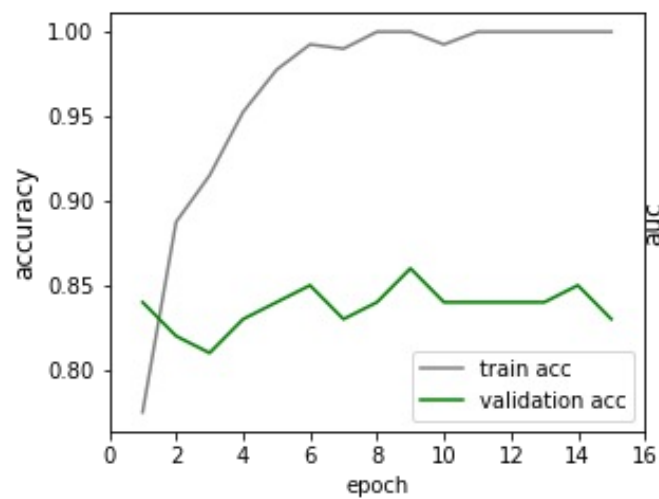
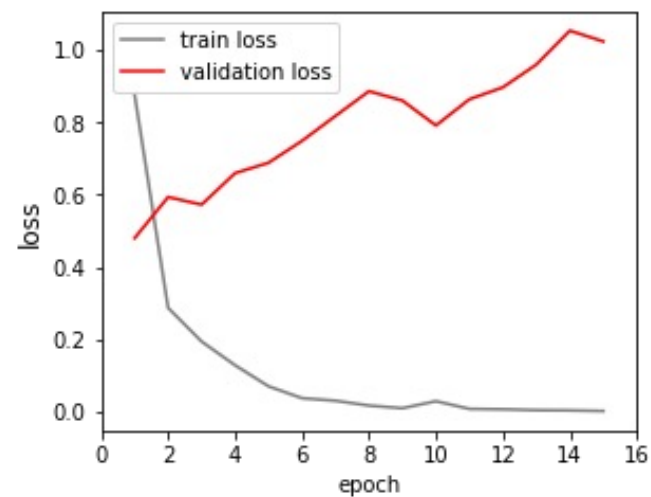
Model 1

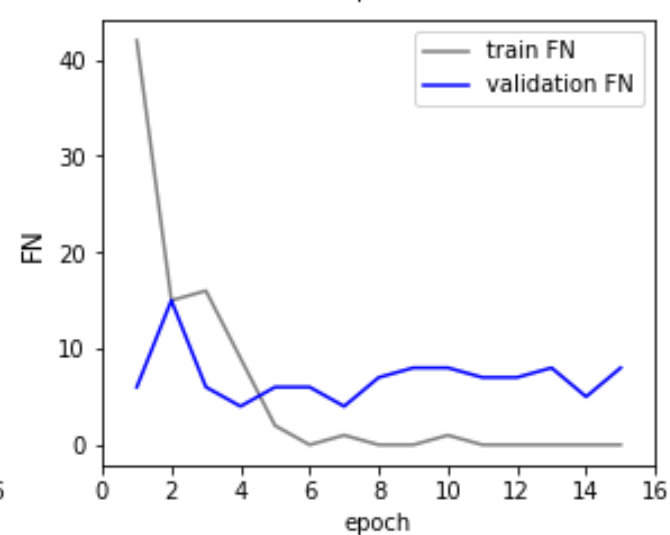
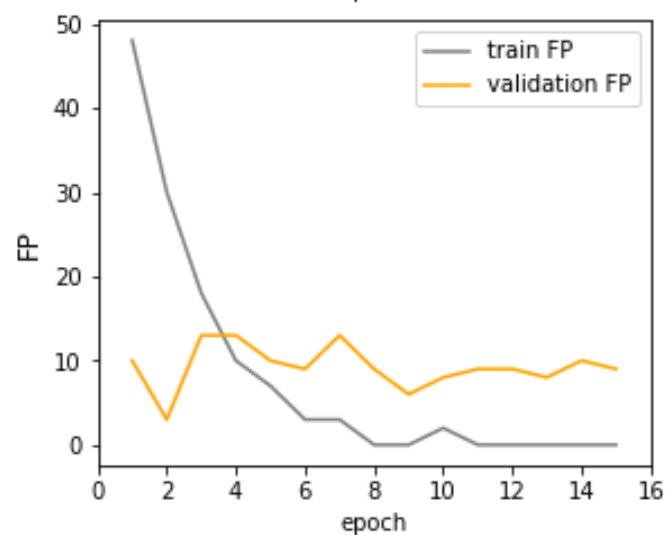
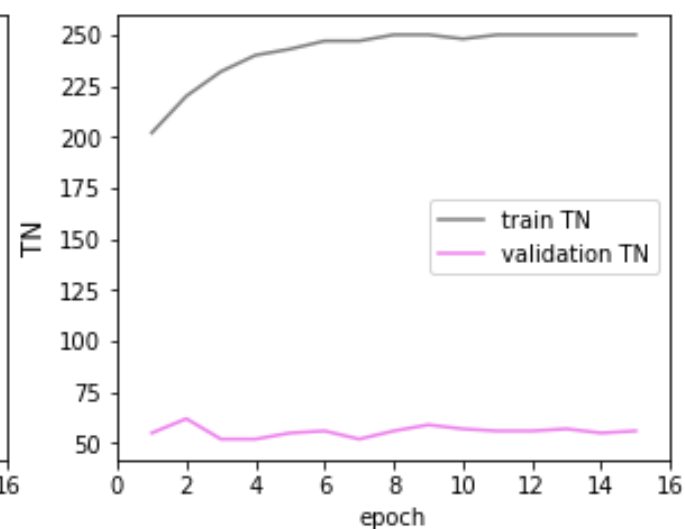
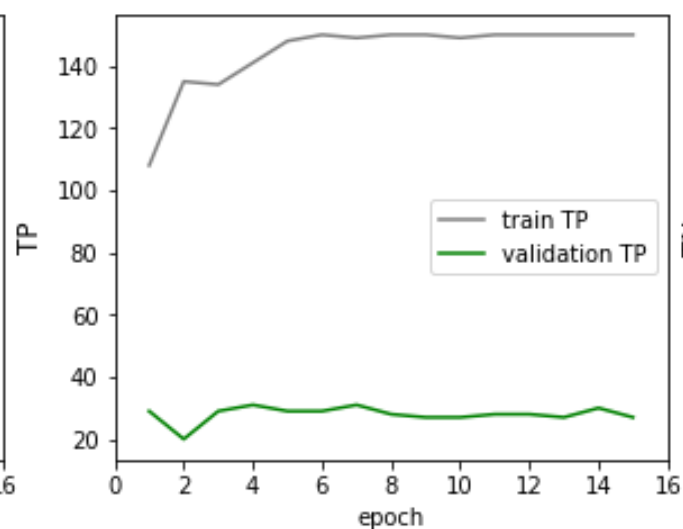
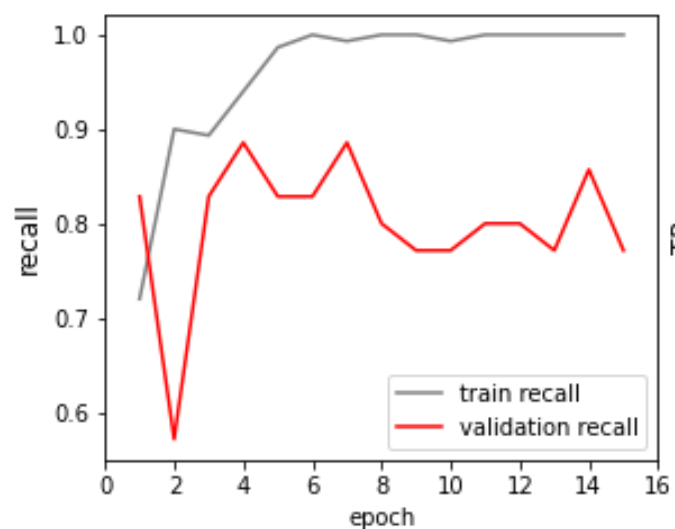




Model 1

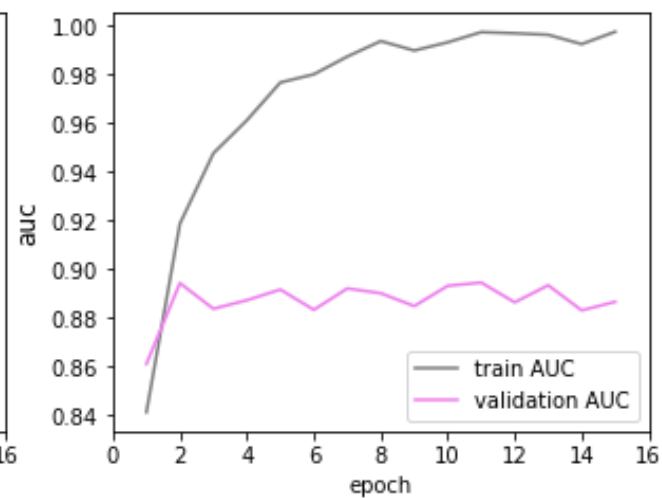
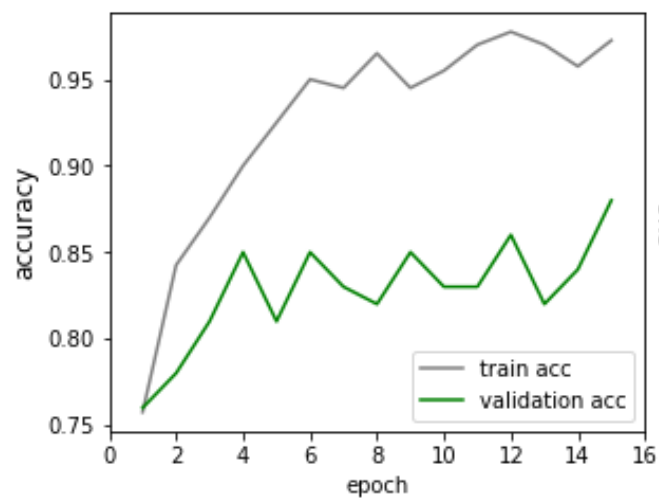
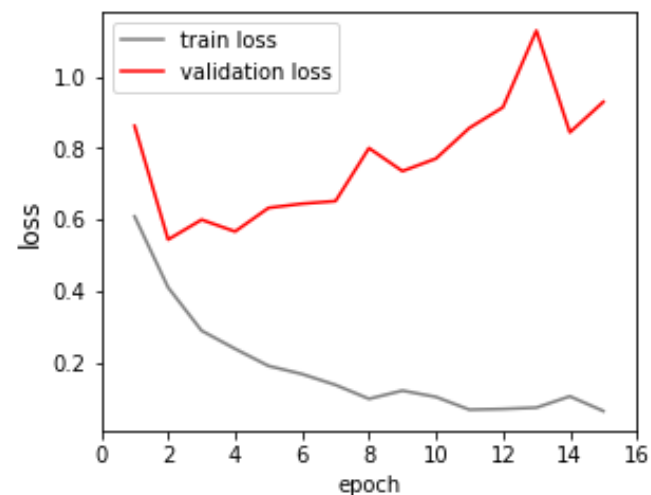
Model 2

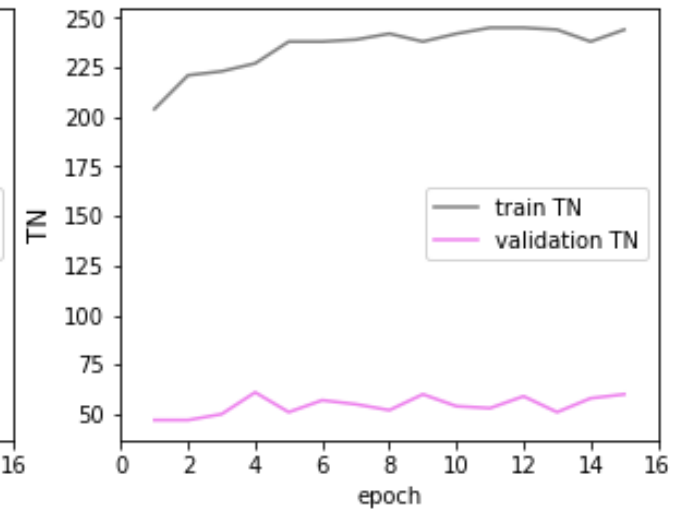
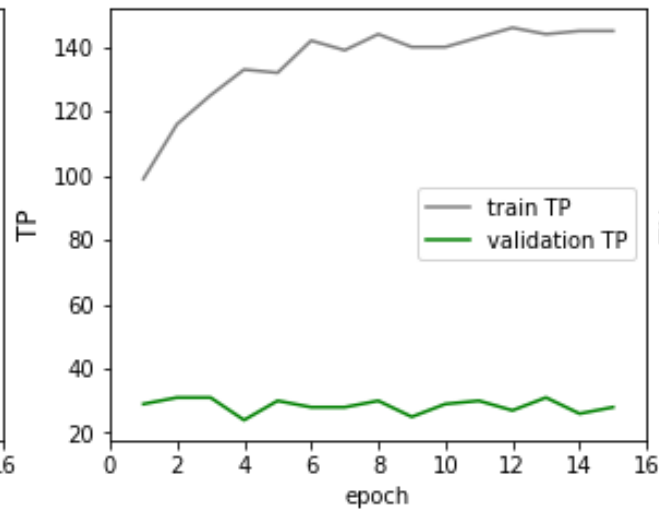
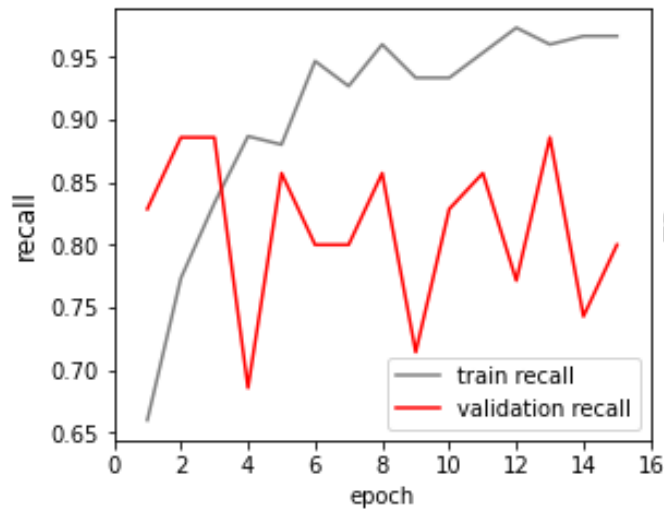




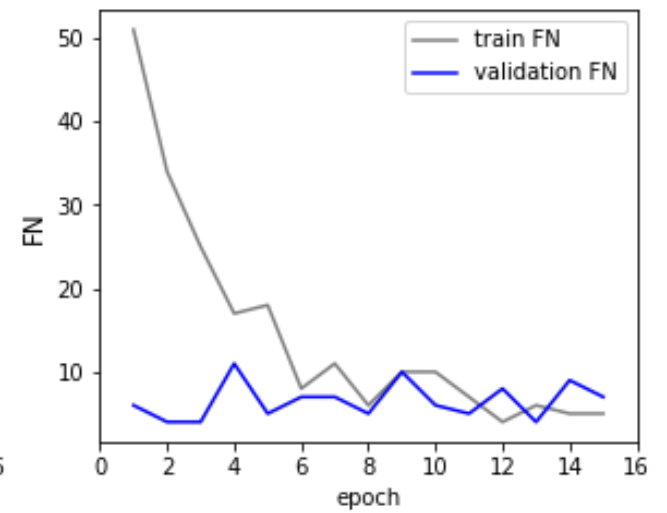
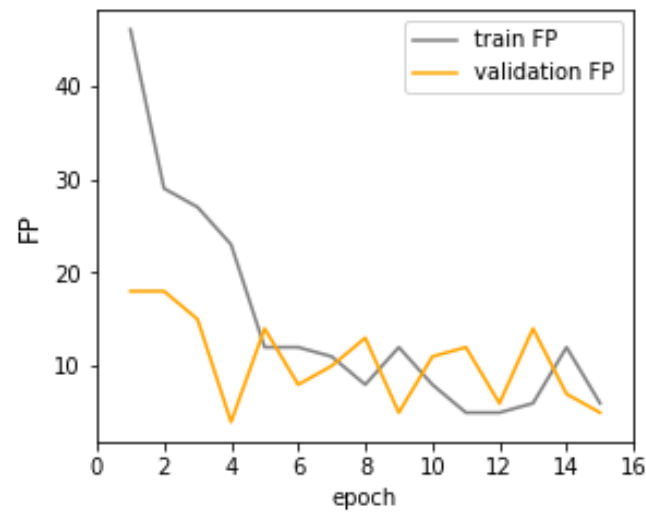
Model 2

Model 3





Model 3



Model 1

**Train
(Duke 400)**

	TP	TN	FP	FN	sens	spec
0	106.0	192.0	58.0	44.0	0.706667	0.768
1	128.0	234.0	16.0	22.0	0.853333	0.936
2	139.0	233.0	17.0	11.0	0.926667	0.932
3	145.0	239.0	11.0	5.0	0.966667	0.956
4	146.0	240.0	10.0	4.0	0.973333	0.960
5	144.0	242.0	8.0	6.0	0.960000	0.968
6	148.0	247.0	3.0	2.0	0.986667	0.988
7	148.0	248.0	2.0	2.0	0.986667	0.992
8	148.0	244.0	6.0	2.0	0.986667	0.976
9	149.0	248.0	2.0	1.0	0.993333	0.992
10	150.0	250.0	0.0	0.0	1.000000	1.000
11	150.0	250.0	0.0	0.0	1.000000	1.000
12	150.0	250.0	0.0	0.0	1.000000	1.000
13	150.0	250.0	0.0	0.0	1.000000	1.000
14	150.0	250.0	0.0	0.0	1.000000	1.000

**Validation
(Duke 100)**

	TP_val	TN_val	FP_val	FN_val	sens_val	spec_val
0	22.0	60.0	5.0	13.0	0.628571	0.923077
1	26.0	51.0	14.0	9.0	0.742857	0.784615
2	27.0	54.0	11.0	8.0	0.771429	0.830769
3	26.0	55.0	10.0	9.0	0.742857	0.846154
4	30.0	53.0	12.0	5.0	0.857143	0.815385
5	21.0	60.0	5.0	14.0	0.600000	0.923077
6	26.0	54.0	11.0	9.0	0.742857	0.830769
7	25.0	56.0	9.0	10.0	0.714286	0.861538
8	26.0	55.0	10.0	9.0	0.742857	0.846154
9	25.0	58.0	7.0	10.0	0.714286	0.892308
10	29.0	53.0	12.0	6.0	0.828571	0.815385
11	29.0	55.0	10.0	6.0	0.828571	0.846154
12	29.0	52.0	13.0	6.0	0.828571	0.800000
13	29.0	53.0	12.0	6.0	0.828571	0.815385
14	25.0	57.0	8.0	10.0	0.714286	0.876923

sensitivity, recall, hit rate, or true positive rate (TPR)

$$\text{TPR} = \frac{\text{TP}}{\text{P}} = \frac{\text{TP}}{\text{TP} + \text{FN}} = 1 - \text{FNR}$$

specificity, selectivity or true negative rate (TNR)

$$\text{TNR} = \frac{\text{TN}}{\text{N}} = \frac{\text{TN}}{\text{TN} + \text{FP}} = 1 - \text{FPR}$$

Validation:

- Specificity was OK.
- Sensitivity was bad

Solution:

- increase power
- Increase sample size

Test: Method 3 – Interpolating (on UAB)

	validation (M1)	test (M1)	validation (M2)	test (M2)	validation (M3)	test (M3)
loss	0.446491	0.327347	0.695937	0.362315	0.928706	0.707755
accuracy	0.830000	0.847176	0.860000	0.860465	0.880000	0.807309
auc	0.896703	0.914109	0.885055	0.912788	0.886593	0.908269
recall	0.657143	0.869565	0.685714	0.891304	0.800000	0.934783
TP	23.000000	40.000000	24.000000	41.000000	28.000000	43.000000
TN	60.000000	215.000000	62.000000	218.000000	60.000000	200.000000
FP	5.000000	40.000000	3.000000	37.000000	5.000000	55.000000
FN	12.000000	6.000000	11.000000	5.000000	7.000000	3.000000

Validation on Duke (100)
Test on UAB (301)

```
model1.save("VGG1_interpolating.h5")  
model2.save("VGG2_interpolating.h5")  
model3.save("VGG3_interpolating.h5")
```

Data backup

4 Folders:

- Competitor Company_Yuhui 20220819
- Duke project_python_Yuhui 20220819
- UAB project_python_Yuhui 20220819
- Zeiss PPT - work report_Yuhui 20220819

3 locations:

- Network
- OneDrive
- Remote computer (desktop)

Future

1. Upsampling:

- Pad out 10X10 to 224X224 (with 0 or NaN) (**done (1)**)
- Upsample smaller image (Keras) (**done (3)**)
- standard interpolator (scipy.interpolate.griddata — SciPy v1.9.0 Manual) (**done (4)**)

2. Fine-tuning the transfer study of VGG16 (**done (2)**)

- Hyperparameter tuning: AutoML