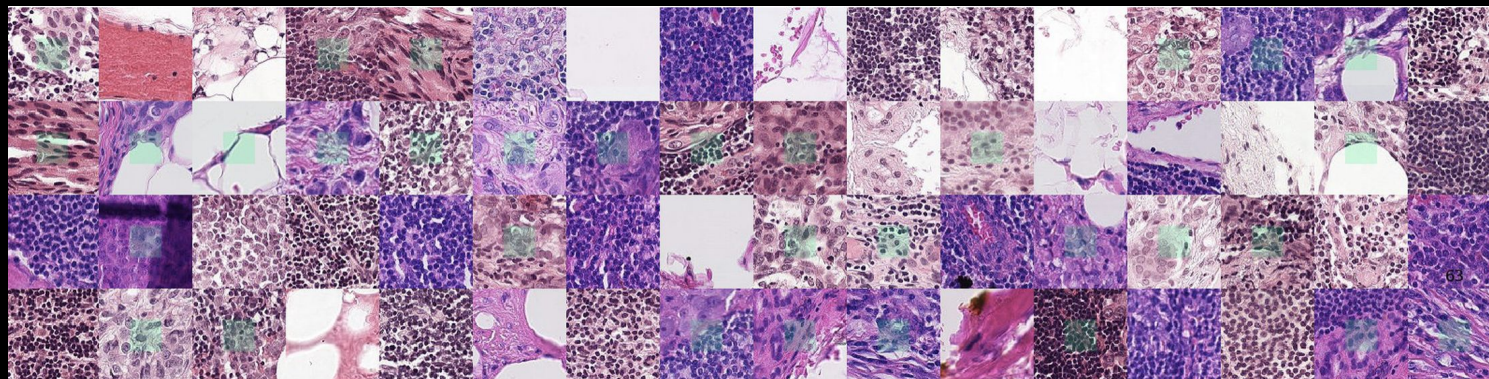


Classification of images from histopathologic scans of lymph node sections

The PatchCamelyon dataset

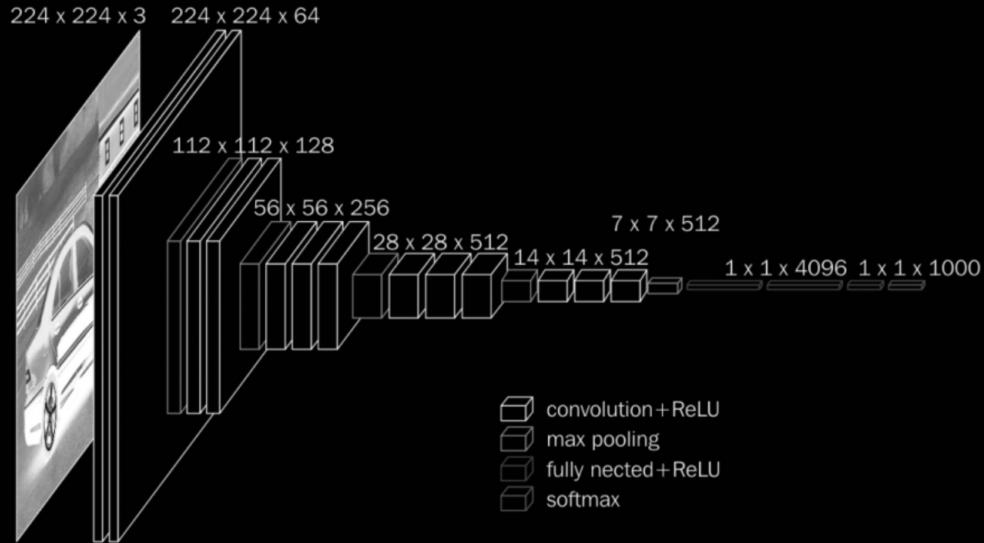


- 327,680 color images ($96 * 96 * 3$)
- Division into training, validation and test datasets
- All sets are class balanced
- Each image is annotated with a binary label
- Positive label indicates that center $32 * 32$ region of the patch contains metastatic tissue

Comparative study: Analysis of different network architectures

VGG 16
ResNet50
DenseNet121
Ensemble learner

VGG 16

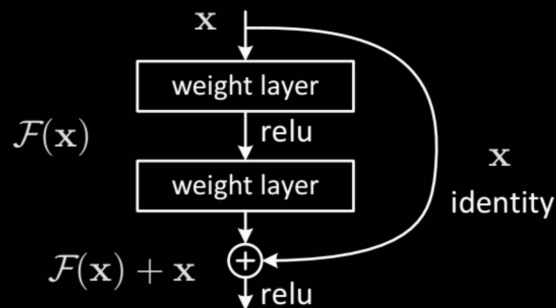


- base: 16
- MLP : {256, 256, 1}
- ReLu
- sigmoid
- batchnorm: CNN part
- dropout for MLP part

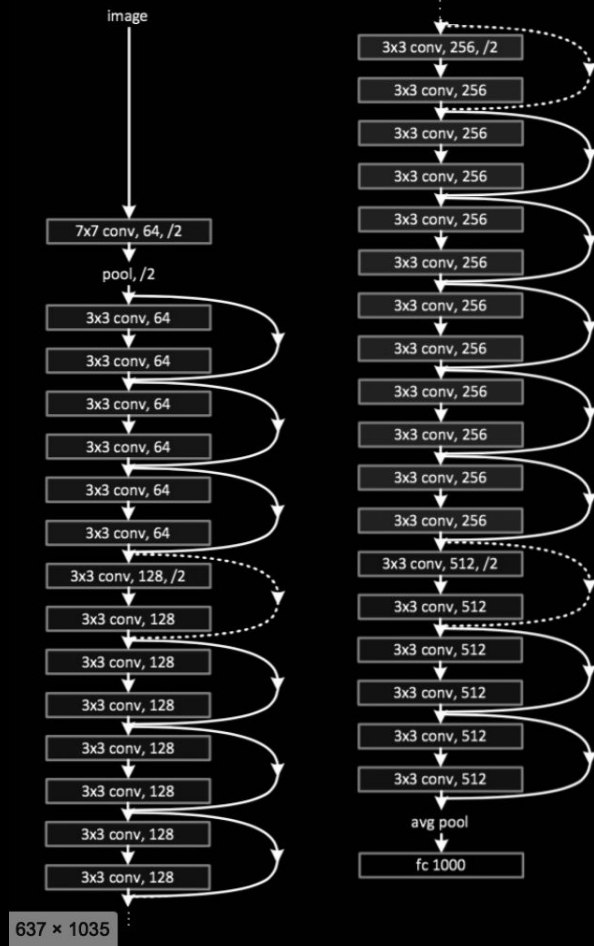
Resnet 50

- base: 16
- Blocks: {3, 4, 6, 3}
- ReLu
- sigmoid
- batchnorm: CNN part
- 5*5 conv + 3*3 stride initially
- Global average pooling
- No MLP!

Residual block:



34-layer residual

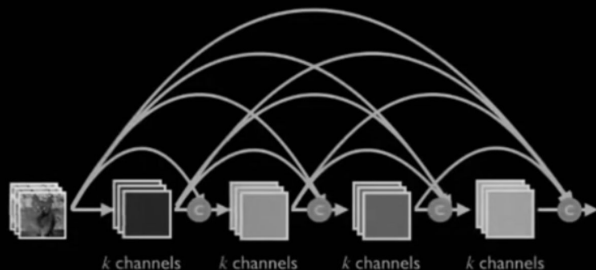


DenseNet 121

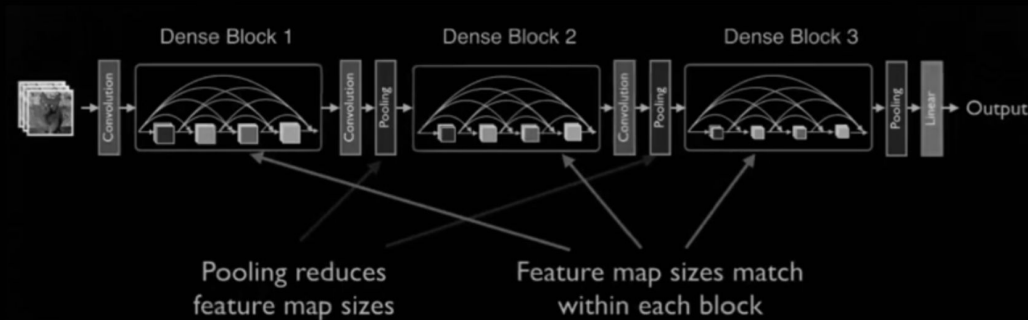
- base: 16
- growth_rate : 8
- Blocks: {6, 12, 24, 16}
- Transition layer
- Bottleneck layer + conv layer
- 5*5 conv + 3*3 stride initially

- Global average pooling.
- No MLP!
- ReLu
- sigmoid
- batchnorm: CNN part

Block:



Network:



Training regimen

Data augmentation

- Horizontal flip
- Vertical flip
- Translation +/- 4 pixels in both x- and y-axis

Metrics

- binary cross entropy
- binary_accuracy
- precision
- sensitivity

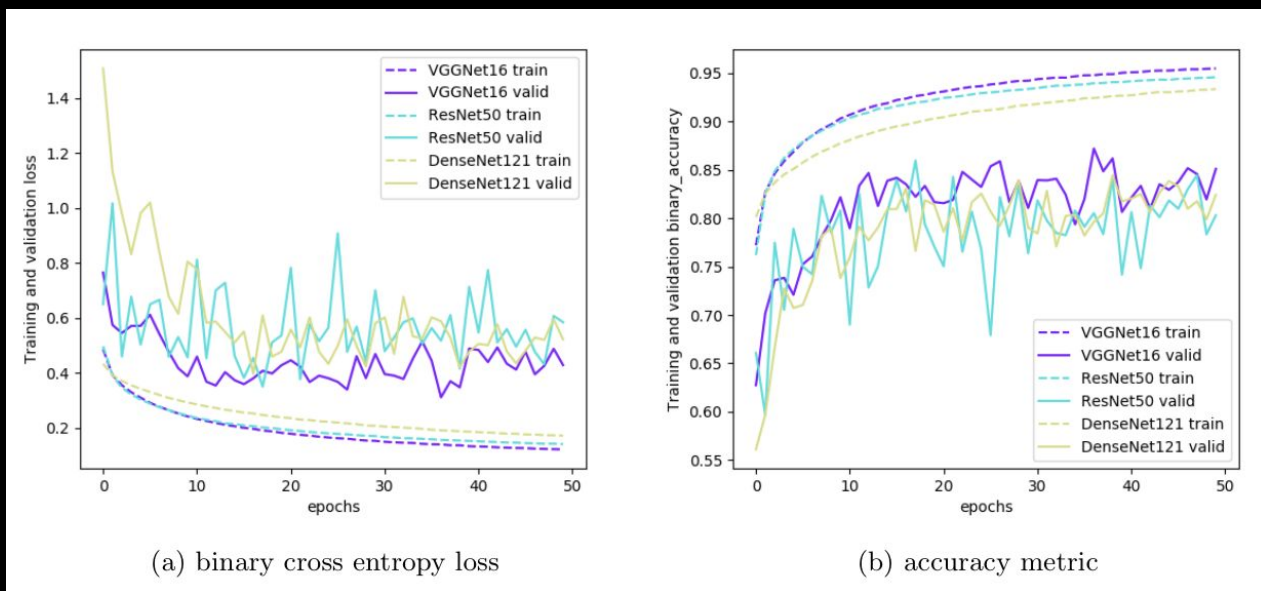
Data handling

- Saved model with lowest validation loss
- History to csv

Learning parameters

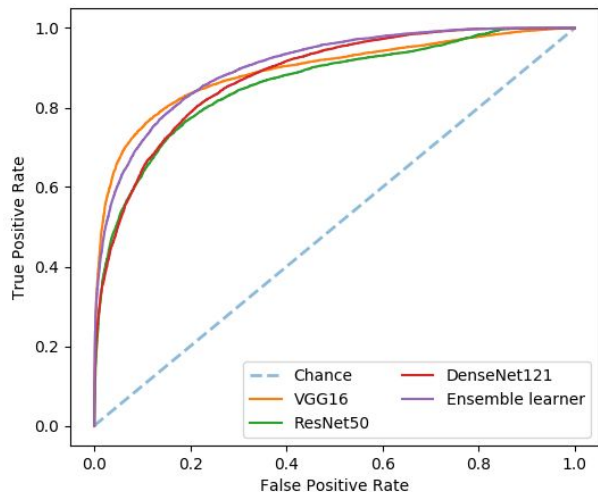
- Different learning rates
- 50 epoch of training/model

Findings: Learning curves



Findings: evaluation on test data

ROC curve:



Test scores:

	accuracy	AUC	NLL	precision	sensitivity
VGG	81.5 (87.2)	89.2 (94.8)	.562 (.312)	61.9 (62.4)	45.7 (55.1)
ResNet	76.8 (86.0)	85.7 (93.4)	.706 (.350)	60.9 (61.8)	43.8 (56.9)
DenseNet	74.8 (83.1)	87.9 (93.4)	.680 (.397)	61.7 (62.7)	39.1 (50.6)
Ensemble	78.8 (88.0)	90.4 (95.4)	.495 (.289)	84.5 (85.6)	61.7 (76.8)

Table 1: Loss and metric scores for the networks on the PatchCamelyon test and validation datasets. The test scores are shown regularly and the validation scores are within parentheses. AUC is the area under curve measure and NLL is the negative log loss, i.e. the binary cross entropy loss.

Implementation - structure

*Tensorflow + Keras
Images stored in
Hierarchical Data Format*

- *File system*
- *For teachers: If you want to run the code, the Results folder and its subfolders are created automatically*

```
InputFiles
├── densenet.json
├── prediction.json
├── resnet.json
└── vgg.json

Models
├── __init__.py
├── densenet.py
├── resnet.py
└── vgg.py

Results
├── History
│   ├── DenseNet121.csv
│   ├── ResNet50.csv
│   └── VGG16.csv
├── Models
│   ├── DenseNet121.h5
│   ├── ResNet50.h5
│   └── VGG16.h5
└── Plots
    ├── ROC_curve_test.png
    ├── accuracy.png
    ├── loss.png
    ├── precision.png
    └── sensitivity.png

Utils
├── __init__.py
├── dataloader.py
├── generator.py
├── metrics.py
├── plotter.py
├── show_dataset.py
└── utilities.py

dl_data
├── Camelyon
│   ├── camelyonpatch_level_2_split_test_meta.csv
│   ├── camelyonpatch_level_2_split_test_x.h5
│   ├── camelyonpatch_level_2_split_test_y.h5
│   ├── camelyonpatch_level_2_split_train_meta.csv
│   ├── camelyonpatch_level_2_split_train_x.h5
│   ├── camelyonpatch_level_2_split_train_y.h5
│   ├── camelyonpatch_level_2_split_valid_meta.csv
│   ├── camelyonpatch_level_2_split_valid_x.h5
│   └── camelyonpatch_level_2_split_valid_y.h5
├── plot_history.py
├── predict.py
├── show_dataset.py
└── train.py
```

Implementation - execution

- *Generic network definitions*
- *Config. files (json)*
- *Argparsers*

```
{
  "model" : "DenseNet",
  "modelparams" : {
    "base" : 16,
    "dense_blocks" : 4,
    "layers_per_block" : [6, 12 , 24, 16],
    "growth_rate" : 8,
    "dense_depth" : 5,
    "out_fnc" : "sigmoid",
    "classes" : 1,
    "bottleneck" : true,
    "dropout_rate" : 0.5
  },
  "regimen" : {
    "lr" : 0.00001,
    "batch_size" : 16,
    "epochs" : 50,
    "loss" : "binary_crossentropy",
    "optimizer" : "Adam",
    "data_augmentation" : true,
    "metrics" : ["binary_accuracy", "precision", "sensitivity"]
  },
  "augmentation" : {
    "rotation_range" : 10,
    "width_shift_range" : 4,
    "height_shift_range" : 4,
    "horizontal_flip" : true,
    "vertical_flip" : true
  },
  "result" : {
    "history_file_name" : "DenseNet121.csv",
    "model_file_name" : "DenseNet121.h5"
  }
}
```

The end