

- CLASSIFYING
BIOLOGICAL IMAGES
USING PRE-TRAINED
CNNs



TEAM

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Problem

● Object Classification in Biological Domain

- Difficult to use a CNN on a small dataset
- Large number of samples required for greater accuracy
- Annotating biological images needs expertise
- Large labelled dataset not available



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Dataset Description

Breast Cancer Histopathological Database

- Benign:

- Adenosis
- Fibroadenoma
- Phyllodes tumor
- Tubular Adenoma

- Malignant:

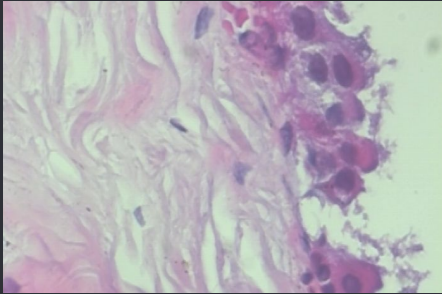
- Carcinoma
- Lobular Carcinoma
- Mucinous Carcinoma
- Papillary Carcinoma

Breast Cancer Histopathological Database

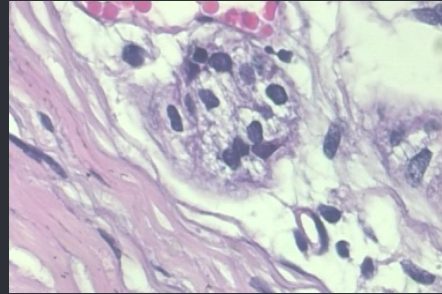
- Data from 82 patients
- RGB images, size 700x460

Magnification	Benign	Malignant	Total
40X	652	1,370	1,995
100X	644	1,437	2,081
200X	623	1,390	2,013
400X	588	1,232	1,820
Total Number of Images	2,480	5,429	7,909

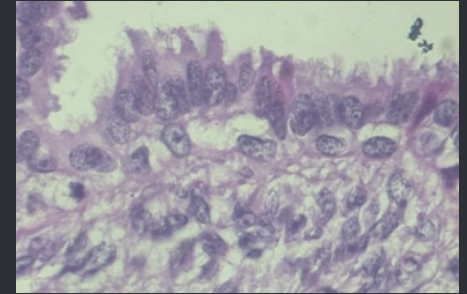
● Samples at 400X magnification



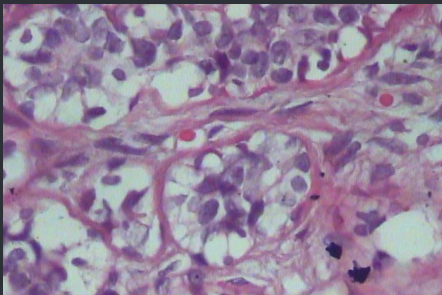
Adenosis



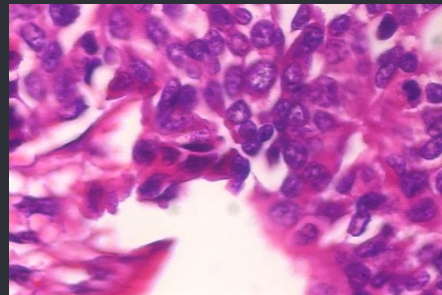
Fibroadenoma



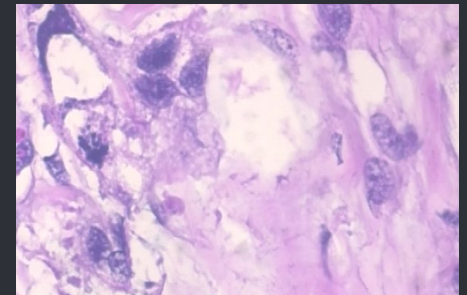
Phyllodes Tumor



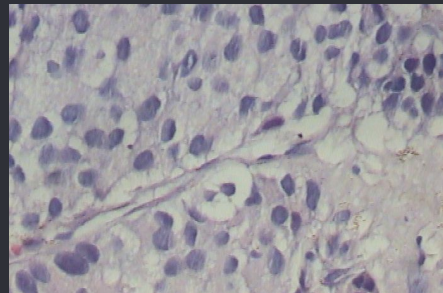
Tubular Adenoma



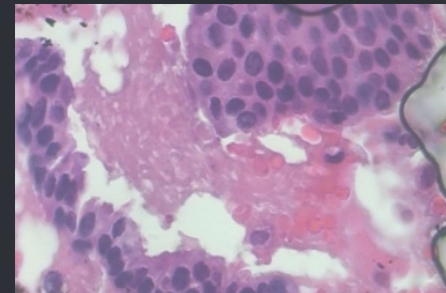
Carcinoma



Lobular Carcinoma



Mucinous Carcinoma



Papillary Carcinoma

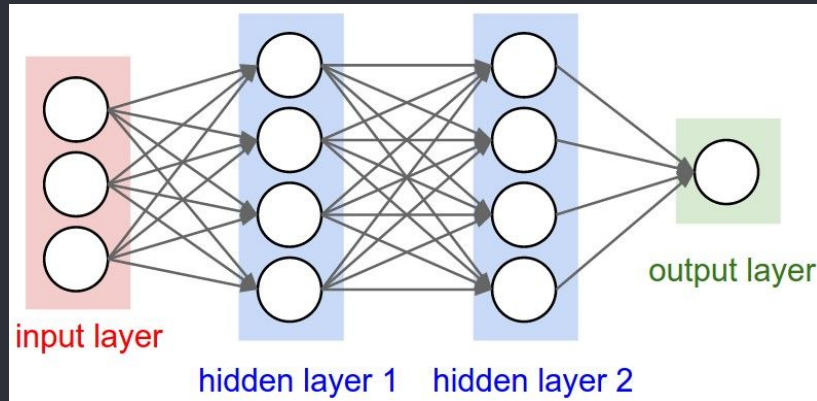
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Convolutional Neural Networks



Convolutional Neural Network

- Similar to Neural Networks



- Encode images into meaningful features

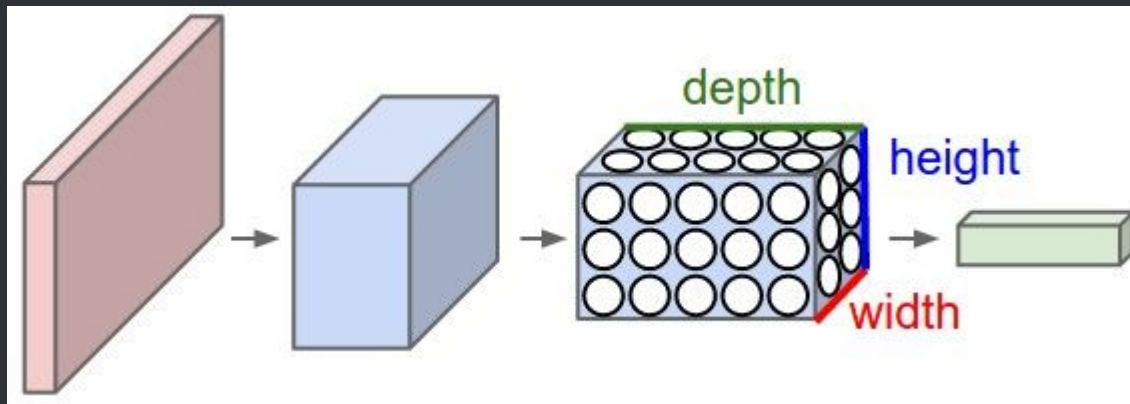
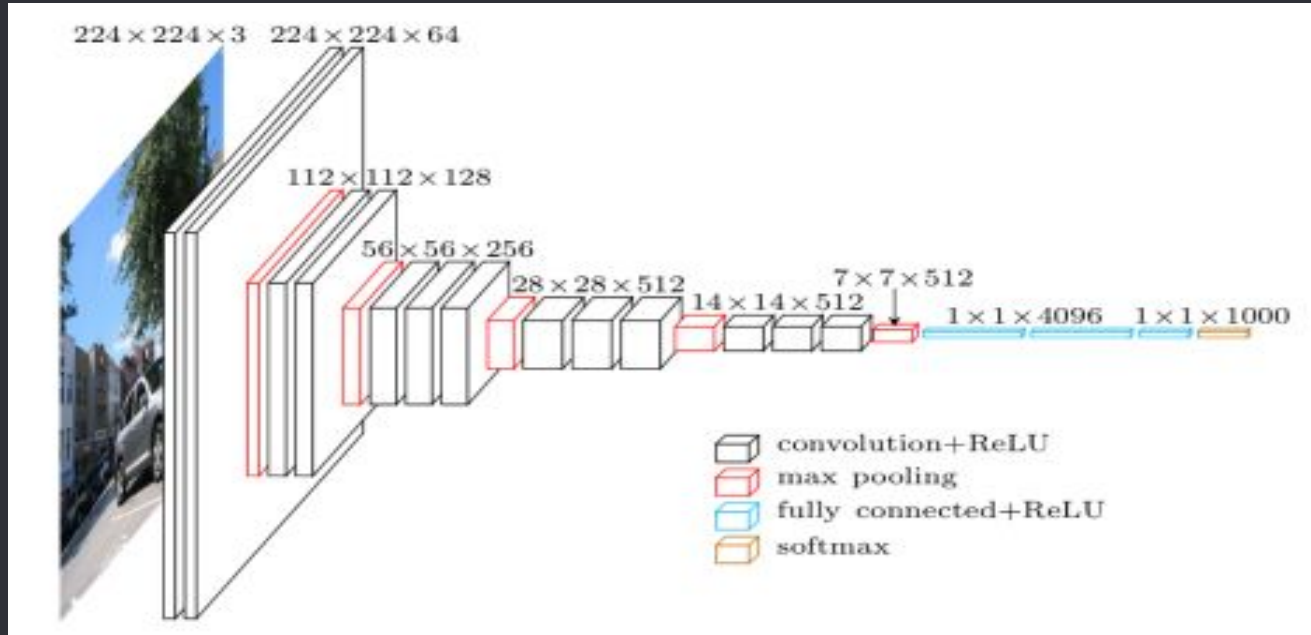


Image references: Convolutional Neural Networks for Visual Recognition

VGG-16 Network



- Depth : critical component in performance
- Components: Conv layers, FC layers, ReLU, Pooling
- 140M parameters

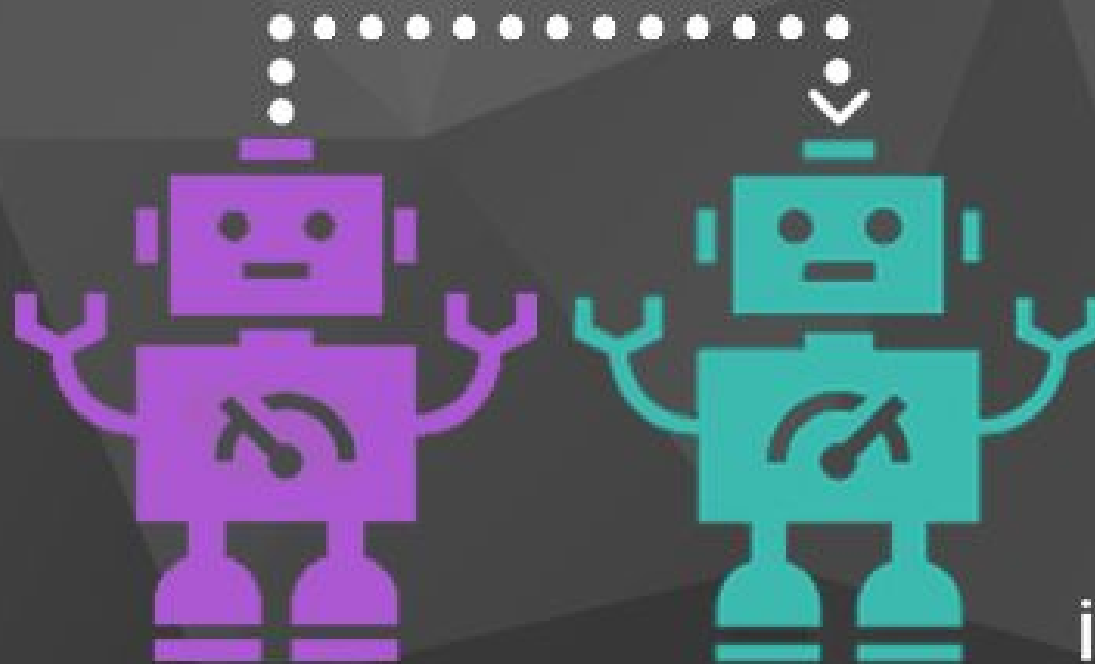
Image reference: Heuritech Blog



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Transfer Learning

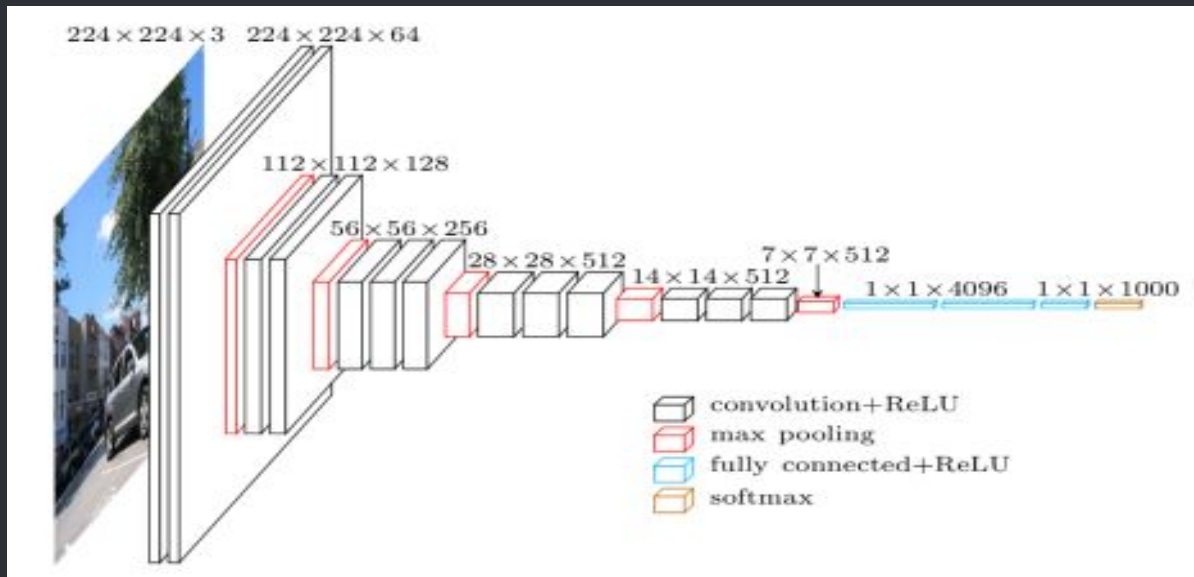
transfer learning



indico

ConvNet as Feature Extractor

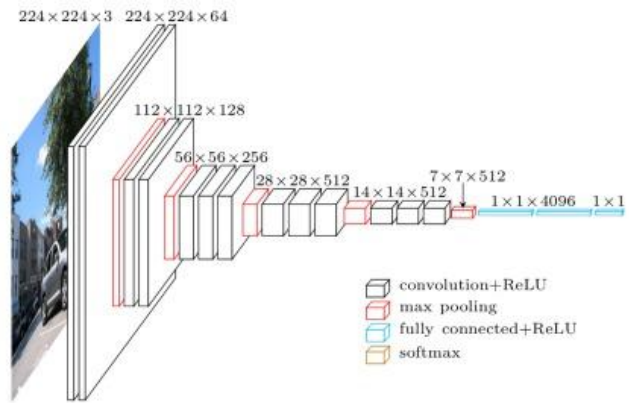
- Use VGG-16 trained on Imagenet (1.2 M images, 1000 classes)
- Remove the last fully-connected layer
- Use as feature extractor, 4096 D vector for every image



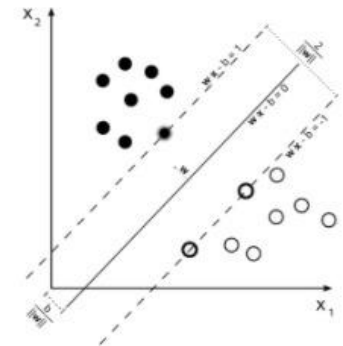


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Approach



Step 2: Getting the features from VGG-16



The diagram shows a root node x at the top. It branches down to a row of trees labeled $tree_1$, $tree_2$, ..., $tree_n$. Each tree is enclosed in a box. Below each tree box is a key: k_1 under $tree_1$, k_2 under $tree_2$, and k_n under $tree_n$. Arrows from k_1 , k_2 , and k_n point to a central node labeled 'voting'. An arrow from 'voting' points down to a final node labeled k .

Step 4: SVM and Random Forest



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Results

● BreakHis Classification

- Dataset is divided according to the magnifications i.e. 40X, 100X, 200X, 400X
- 2 datasets:
 - 2 classes: Benign and Malignant
 - 8 classes: Subtypes of Benign and Malignant
- Classification is done using C-SVC linear, polynomial SVM and Random Forest

BreakHis Classification

2 classes (average accuracy)

	40X	100X	200X	400X
C-SVC Linear SVM	89%	89%	88%	88%
C-SVC Polynomial SVM	88%	90%	89%	85%
Random Forest	89.18%	88%	87.74%	80%

BreakHis Classification

8 classes (average accuracy)

	40X	100X	200X	400X
C-SVC Linear SVM	55.97%	51.59%	55.93%	53.06%
C-SVC Polynomial SVM	53.70%	55.12%	54.78%	53.9%
Random Forest	48.50%	49%	49.81%	48.16%



Conclusion

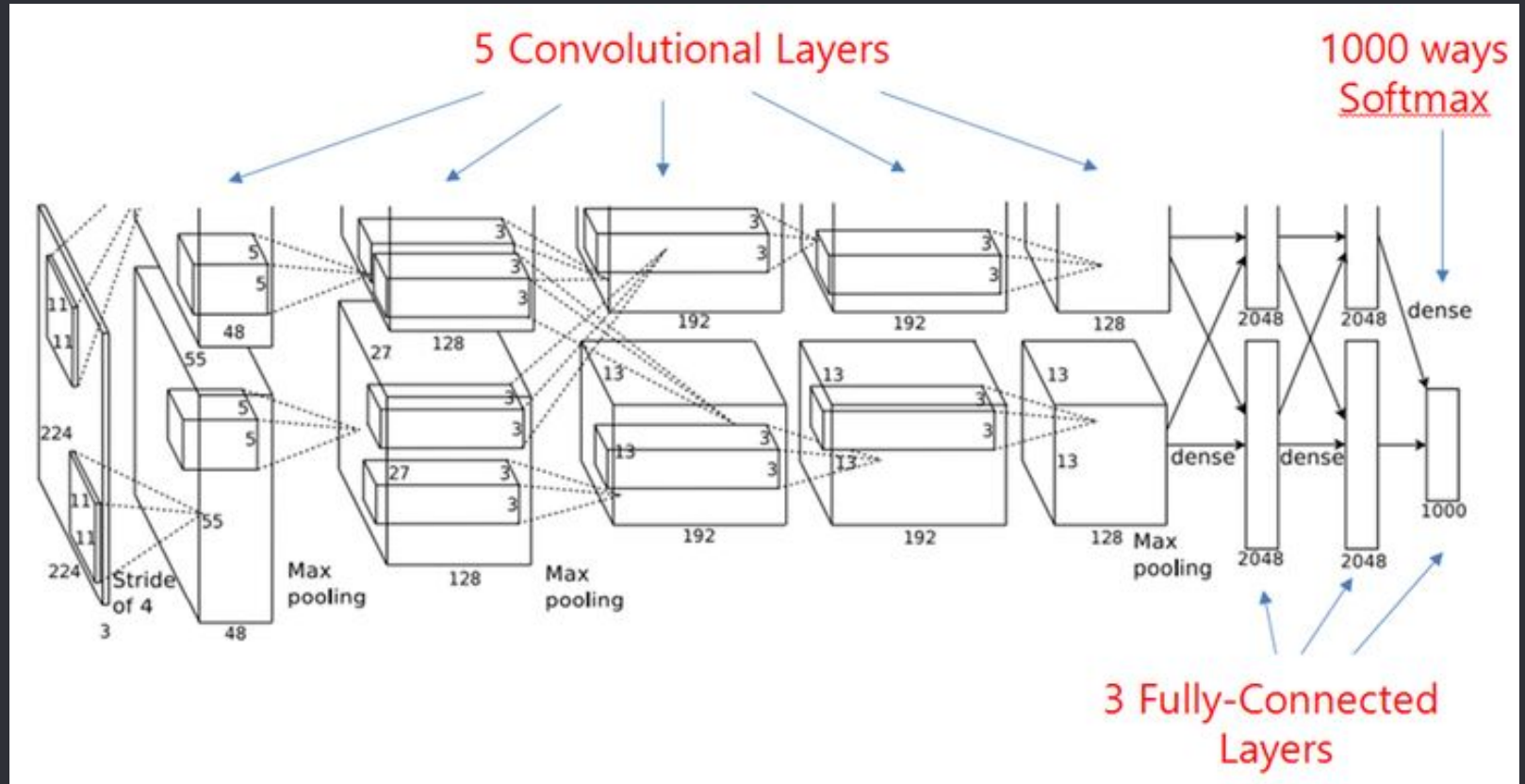
- Pre-training can be used in case of small dataset
- Pre-trained network on natural images, can give significant result in images from an entire different domain
- Performance also depends on complexity of classification task



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Ongoing Work

AlexNet



AlexNet: ~60M parameters

REFERENCES

- Breast Cancer Histopathological Database
<http://web.inf.ufpr.br/vri/breast-cancer-database>
- F. A. Spanhol, L. S. Oliveira, C. Petitjean and L. Heutte, "Breast cancer histopathological image classification using Convolutional Neural Networks," 2016 International Joint Conference on Neural Networks (IJCNN), Vancouver, BC, 2016, pp. 2560-2567.
- Aravindh Mahendran, Andrea Vedaldi, "Visualizing Deep Convolutional Neural Networks Using Natural Pre-Images"
- Nima Tajbakhsh et. al., "Convolutional Neural Networks for Medical Image Analysis: Full Training or Fine Tuning?"
- <http://www.deeplearningmodel.net>
- <https://github.com/tensorflow/models/tree/master/slim>

Thanks!

ANY QUESTIONS?