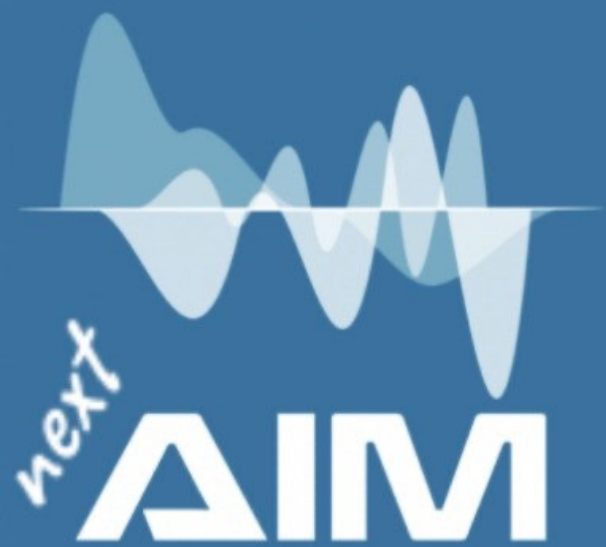


Artificial Intelligence in Medicine



An explainable unified approach for mass detection and classification in DBT

Andrea Berti



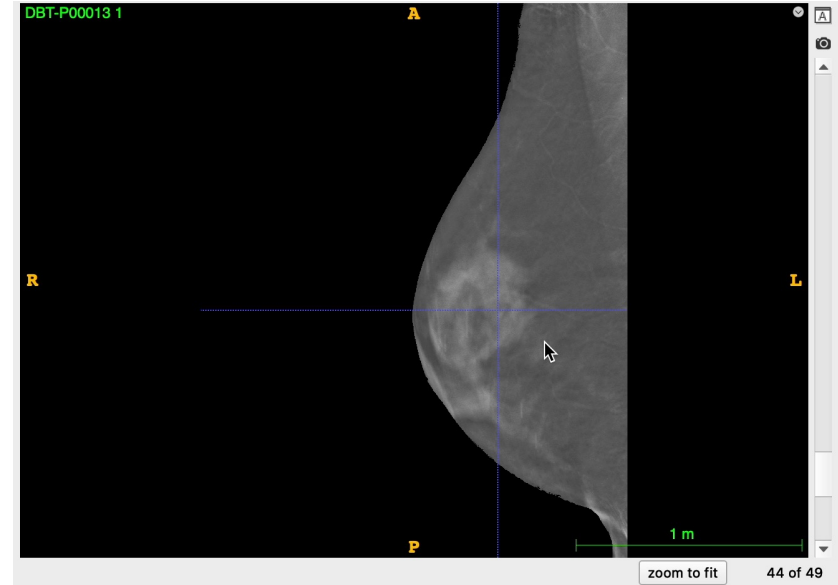
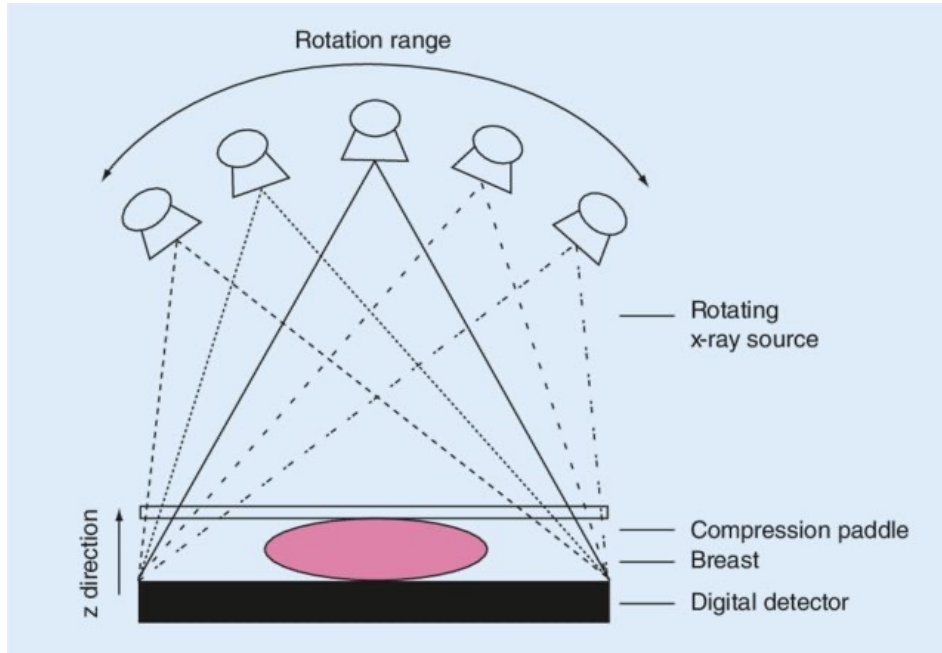
Istituto di Scienza e Tecnologie
dell'Informazione "A. Faedo"
Consiglio Nazionale delle Ricerche



Overview

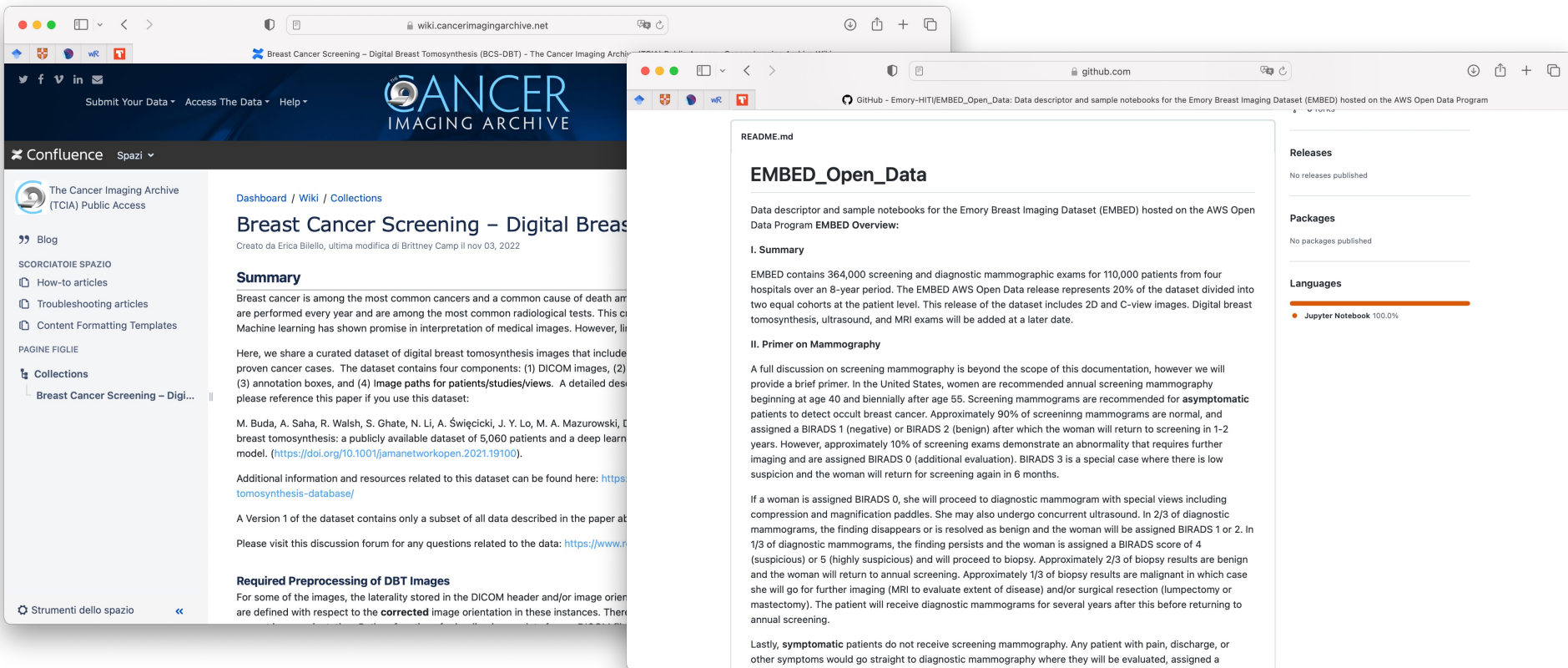
- Digital Breast Tomosynthesis
- The datasets: BCS-DBT & EMBED
- Proposed framework
 - Deep Learning
 - Radiomics
- Explainability in classification

Digital Breast Tomosynthesis



Digital Breast Tomosynthesis

<https://wiki.cancerimagingarchive.net/pages/viewpage.action?pageId=64685580>



The screenshot displays two overlapping browser windows. The background window shows the Cancer Imaging Archive (TCIA) website, specifically the page for 'Breast Cancer Screening – Digital Breast Tomosynthesis (DBT)'. The page includes a sidebar with navigation links like 'Dashboard / Wiki / Collections', 'Summary', and 'Required Preprocessing of DBT Images'. The main content area provides a detailed summary of the dataset, including its components (DICOM images, annotation boxes, image paths, and a deep learning model) and a list of authors (M. Buda, A. Saha, R. Walsh, S. Ghate, N. Li, A. Świąćicki, J. Y. Lo, M. A. Mazurowski, et al.).

The foreground window shows the GitHub repository for the 'EMBED_Open_Data' project. The repository page includes a 'README.md' file, a 'Releases' section, a 'Packages' section, and a 'Languages' section. The 'README.md' file provides a brief overview of the dataset, its purpose, and the recommended screening protocol for patients assigned BIRADS 0, 1, 2, 3, 4, 5, and 6. It also mentions that the dataset is hosted on the AWS Open Data Program.

<https://registry.opendata.aws/emory-breast-imaging-dataset-embed/>

The BCS-DBT Dataset

- Normal:
5129 (91.4%) studies
4609 patients
- Actionable:
280 (5.0%) studies
278 patients
- Biopsy-proven Benign:
112 (2.0%) studies
112 patients
- Biopsy-proven Malignant:
89 (1.6%) studies
89 patients

Normal vs non-Normal

```
### RMLO ###
Number of rmlo = 4781
Number of Normal = 4558
Number of non-Normal = 223
Percentage of Normal = 0.953
```

```
### LMLO ###
Number of lmlo = 4788
Number of Normal = 4558
Number of non Normal = 230
Percentage of Normal = 0.952
```

```
### RCC ###
Number of rcc = 4779
Number of Normal = 4558
Number of non Normal = 221
Percentage of Normal = 0.954
```

```
### LCC ###
Number of lcc = 4791
Number of Normal = 4558
Number of non Normal = 233
Percentage of Normal = 0.951
```

Benign and Malignant

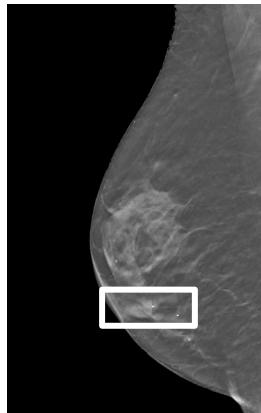
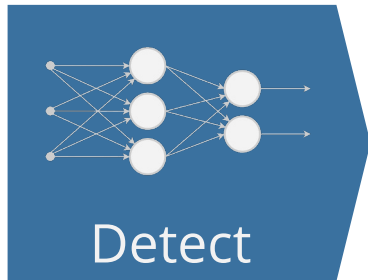
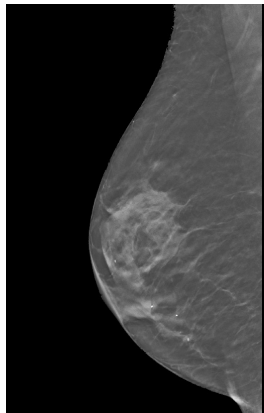
```
### RMLO ###
Number of rmlo = 4781
Number of Benign = 24
Number of Malignant = 20
Number of Benign + Malignant = 44
Percentage of Benign = 0.005
Percentage of Malignant = 0.004
```

```
### LMLO ###
Number of lmlo = 4788
Number of Benign = 36
Number of Malignant = 15
Number of Benign + Malignant = 51
Percentage of Benign = 0.008
Percentage of Malignant = 0.003
```

```
### RCC ###
Number of rcc = 4779
Number of Benign = 23
Number of Malignant = 19
Number of Benign + Malignant = 42
Percentage of Benign = 0.005
Percentage of Malignant = 0.004
```

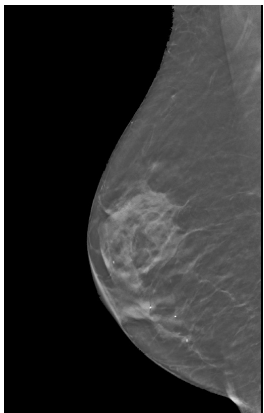
```
### LCC ###
Number of lcc = 4791
Number of Benign = 36
Number of Malignant = 18
Number of Benign + Malignant = 54
Percentage of Benign = 0.008
Percentage of Malignant = 0.004
```

Our approach



Benign

Malignant



Extract
Features

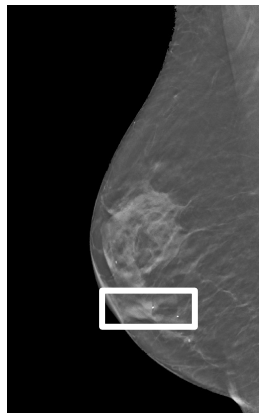
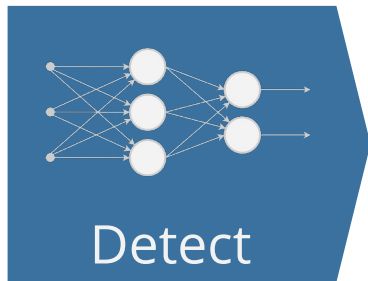
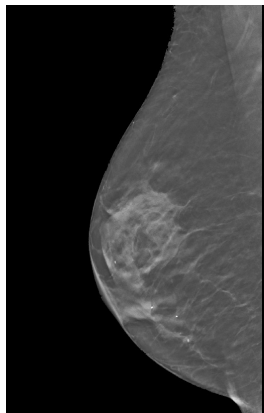
Segment

Classify

Benign

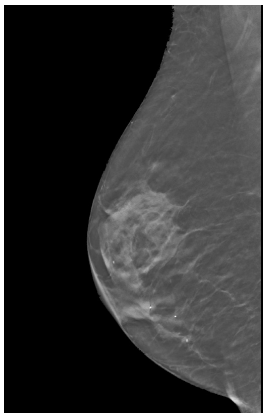
Malignant

Our approach



Benign

Malignant



MatRadiomics

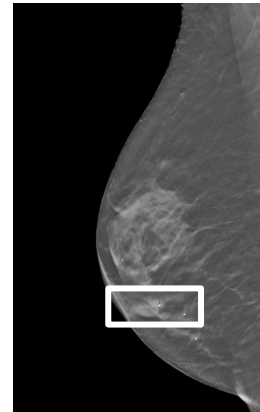
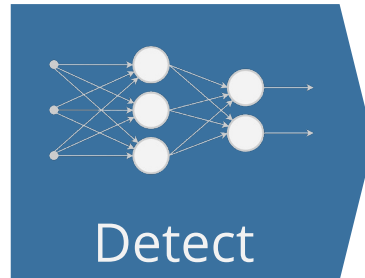
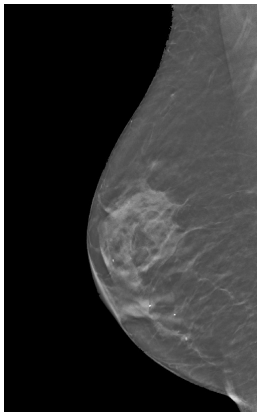
with Alessandro Stefano & Giovanni Pasini

Benign

Malignant

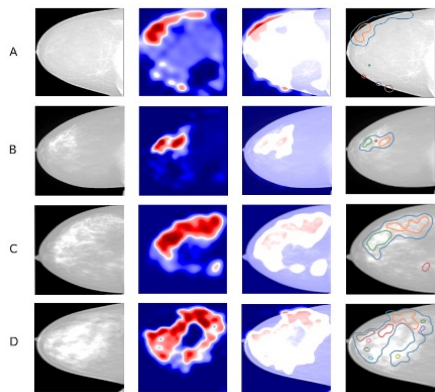
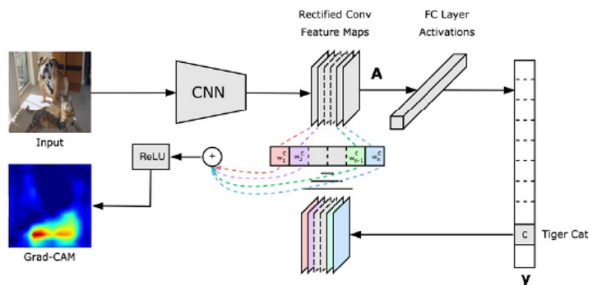
Detection

- Large number of "normal" patients (no mass)
- GAN & UNet
- Anomaly detection approach

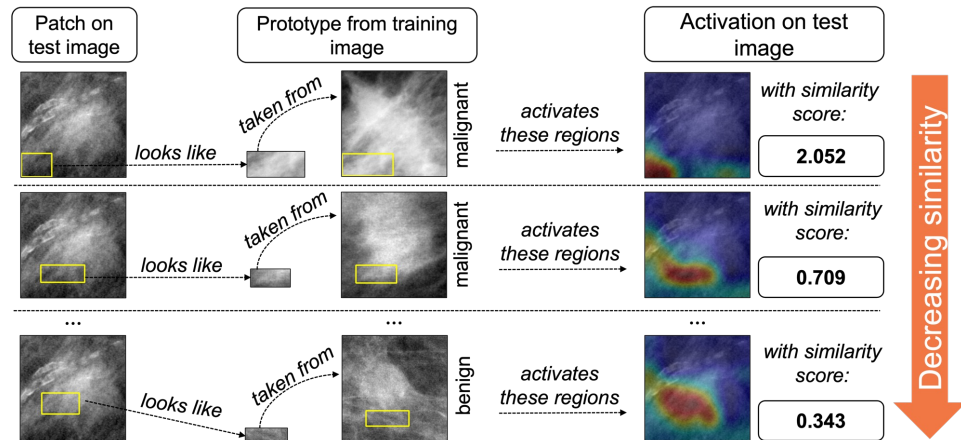


Explainable Classification

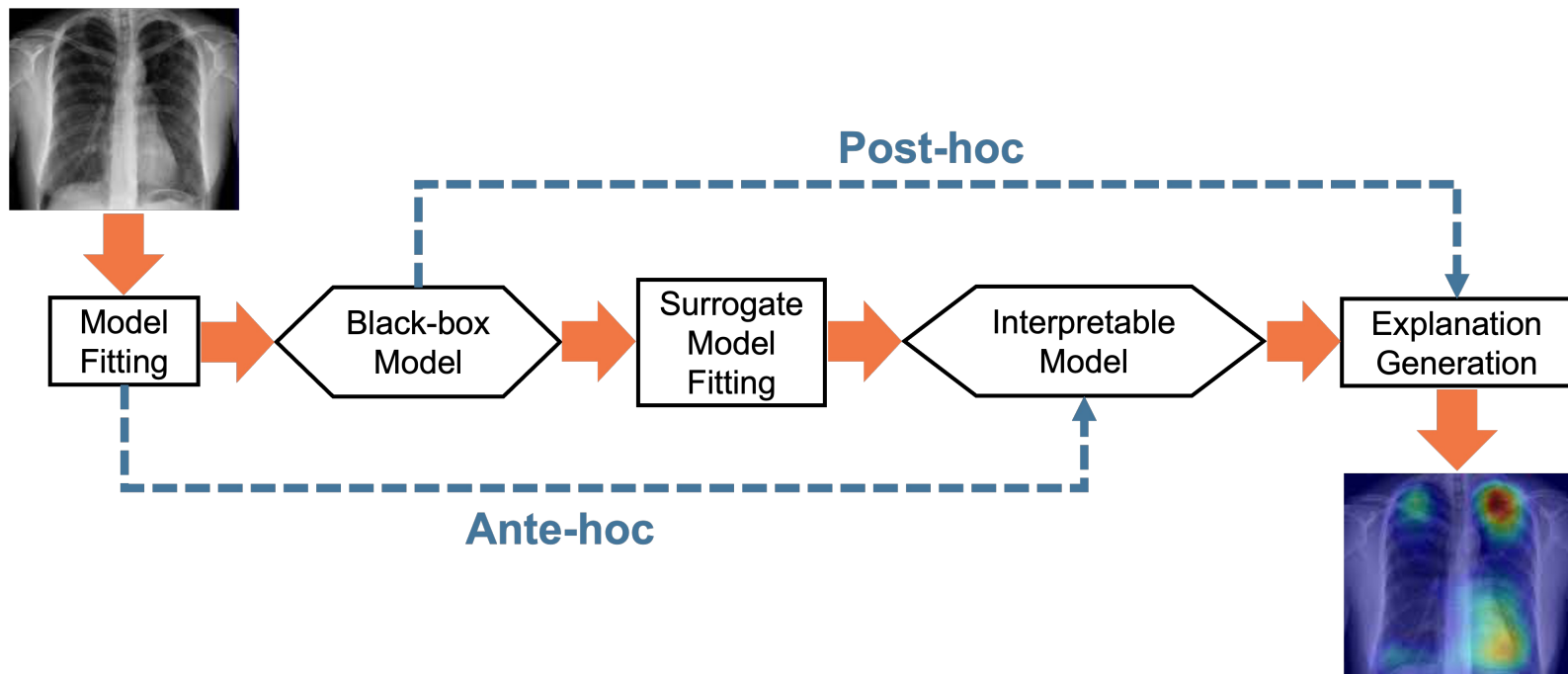
CNN + GradCam



ProtoPNet

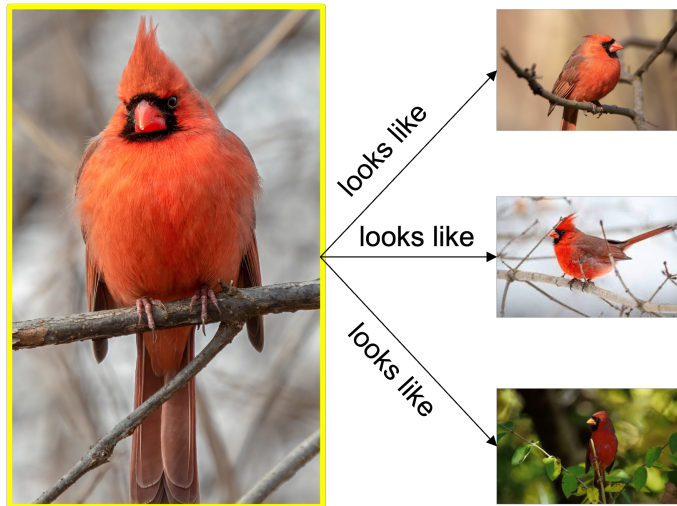


Ante-hoc explainability



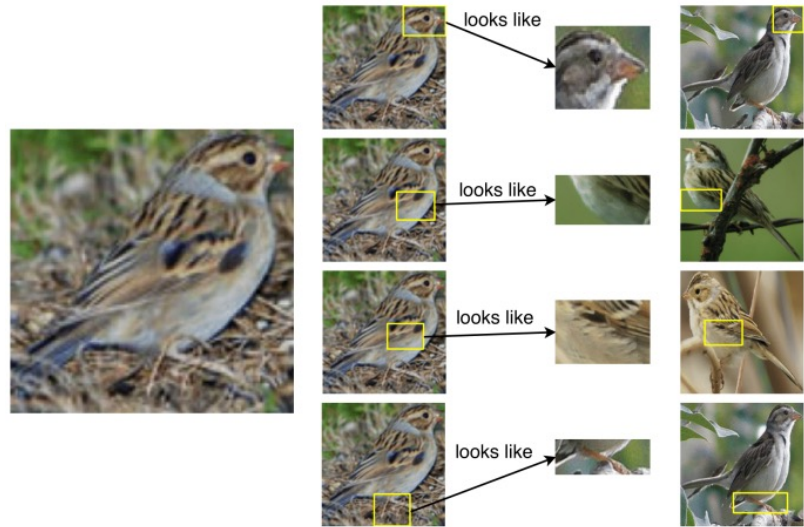
Case-based reasoning

Prototypical Learning



Northern Cardinal

Prototypical Part Learning

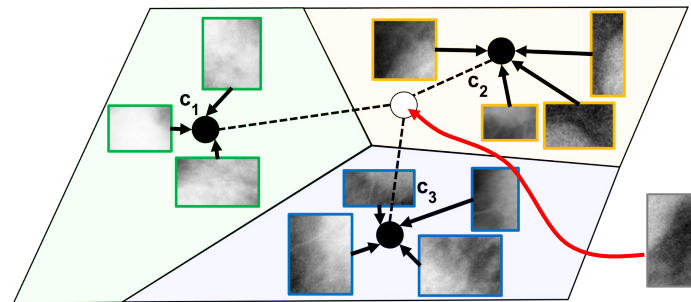


Chen et al. 2019

ProtoPNet on benign vs malignant breast masses

Previously...

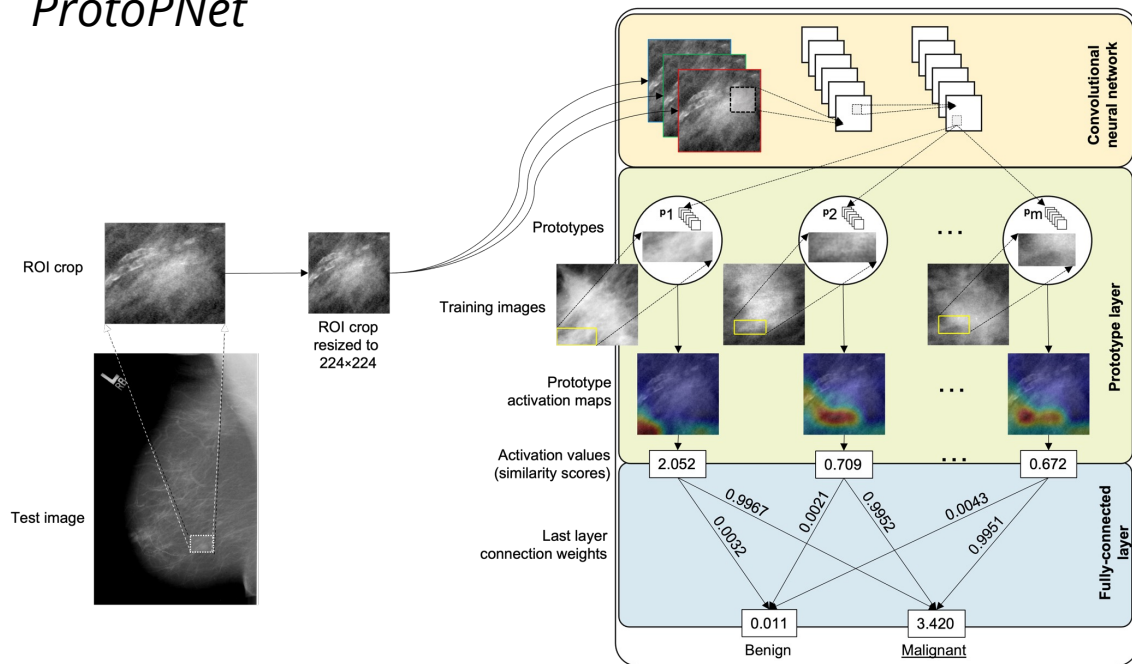
On the Applicability of Prototypical Part Learning in Medical Images: Breast Masses Classification Using ProtoPNet



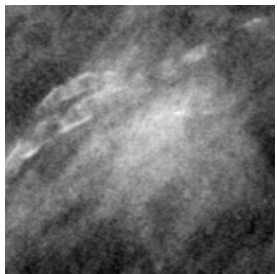
$$\text{Loss} := \text{CrossEntr} + \alpha \text{Clst} + \beta \text{Sep}$$

AIHA Workshop
International Conference on
Pattern Recognition (Canada)

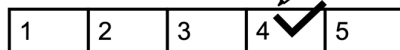
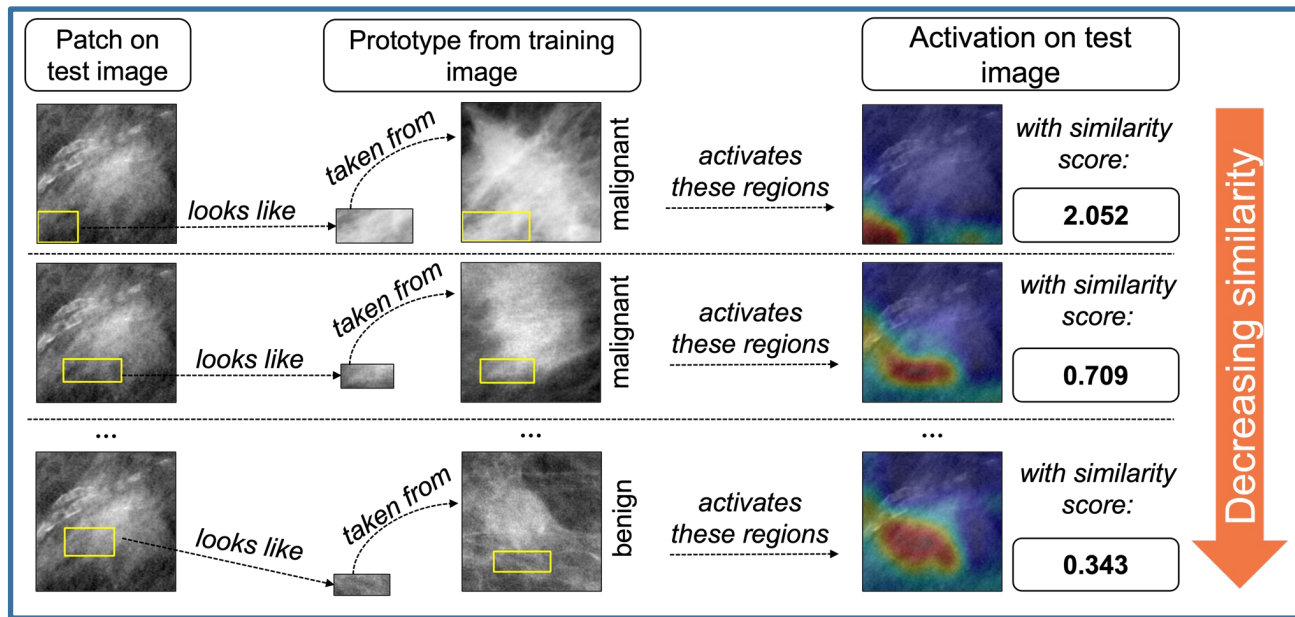
with Gianluca Carloni and Sara
Colantonio (ISTI-CNR)



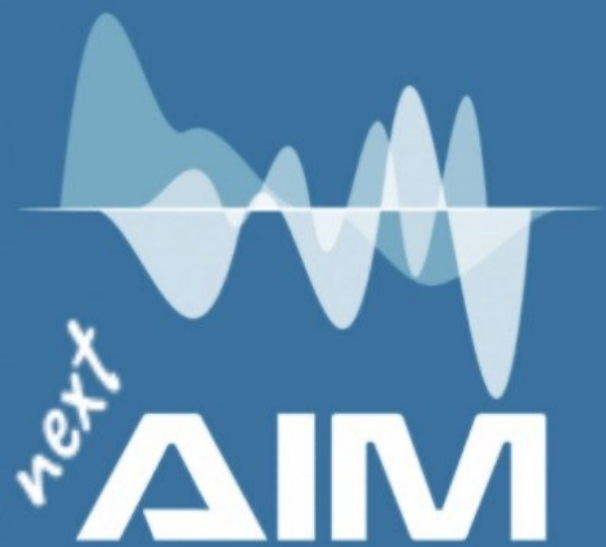
The output of ProtoPNet



Test image: **malignant**
 Predicted as: **malignant**



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I paper su DBT

2D Convolutional Neural Networks for 3D Digital Breast Tomosynthesis Classification (*Zhang et al.*)

- *Classification maligni vs negativi*
- *2D-CNN preallennata su ImageNet per estrarre le feature sulle singole slice*
- *Pooling sulle feature-map delle slice per classificazione 3D*
- *Dataset privato:*
 - *3018 negativi (non maligni)*
 - *272 maligni*
- *AUC = 0.854 (meglio delle 3D-CNN)*

I paper su DBT

A deep learning classifier for digital breast tomosynthesis

(Ricciardi, Mettivier et al.)

- *Classification massa vs non-massa*
- *Custom D-CNN vs AlexNet/VGG*
- *Sembra che non sia patient-stratified*
- *Grad-CAM*
- *Dataset privato 109 pazienti:*
 - *3166 (H1) + 152 (H2) massa*
 - *1526 (H1) + 90 (H2) non-massa*
- *Acc = 0.94*

I paper su BCS-DBT

Intelligent Computer-Aided Model for Efficient Diagnosis of Digital Breast Tomosynthesis 3D Imaging Using Deep Learning (*Adel El-Shazli et al.*)

- *Classification normali vs maligni vs benigni*
- *Mod_AlexNet vs modelli standard*
- *Transfer learning*
- *Dataset:*
 - *499 Normal, 62 Benign, 39 Malignant*
- *Acc = 91.61% sul test*

I paper su BCS-DBT

Applying Graph Convolution Neural Network in Digital Breast Tomosynthesis for Cancer Classification (*Bai et al.*)

- *Classification normal vs cancer*
- *Graph CNN con self-attention pooling layer (risolvere 2D e 3D)*
- *Dataset BCS + privato:*
 - *158 (BCS) + 75 (priv) normal*
 - *75 (BCS) + 94 (priv) cancer (da dove i 75?)*
- *$Acc = 0.84$, $AUC = 0.87$*

I paper su BCS-DBT

Trainable Summarization to Improve Breast Tomosynthesis Classification (*Tardy et al.*)

- *Classification normal vs cancer*
- *Multiple instance learning, slabbing + classification (ResNet)*
- *Dataset privato mammo \rightarrow pretraining, BCS \rightarrow fine-tuning:*
 - *Priv: 1250 benign, 1250 malignant*
 - *BCS: 100 normal, 75 cancer*
- *$AUC = 0.73$*

Altri paper su BCS-DBT: object detection

- **Detection of masses...** (*Buda et al.*): *Focal loss per sbilanciamento*
- **Developing breast leasion detection...** (*Hossain et al.*): *3 slice consecutive per RGB, (falsi positivi)*
- **Lightweight transformer...** (*Zhang et al.*): *VIT per risolvere scarsità casi positivi. ResNet - LightweightVIT - ResNet*

Classificazione

- Possibili classificazioni:
 - Normal vs (Actionable + Benign + Malignant)
 - Normal vs (Benign + Malignant)
 - Benign vs Malignant -> ma sono pochi casi
- Come utilizzare le immagini:
 - 2D vs 3D