

Departamento Ingeniería
Eléctrica

Breast Cancer Recurrence Prediction using a Quantitative Evaluation of the Nuclear Pleomorphism

February 25, 2019

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Pathological examination issues

- Due to the heterogeneity between the healthy tissue and the disease, most of the morphological indicators of prognosis are not quantified

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- Due to the heterogeneity between the healthy tissue and the disease, most of the morphological indicators of prognosis are not quantified
- The observations are susceptible to inter and intra observer variations

The Suitability of a Quantitative Estimation in Pathology Workflow

- Digital version of the whole glass slide and image analysis are used to improve the diagnosis

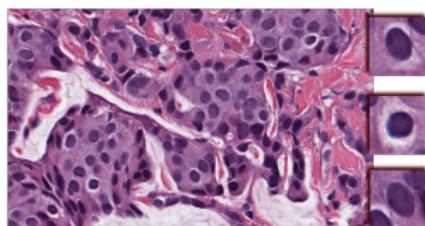
The Suitability of a Quantitative Estimation in Pathology Workflow

- Digital version of the whole glass slide and image analysis are used to improve the diagnosis
- The Quantification of morphological structures to determine the predisposition of a disease and to deliver healthcare

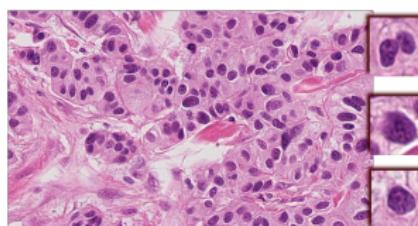
Nuclear Pleomorphism (NP) in Ductal carcinoma in situ (DCIS) - case of Study

Nuclear Pleomorphism

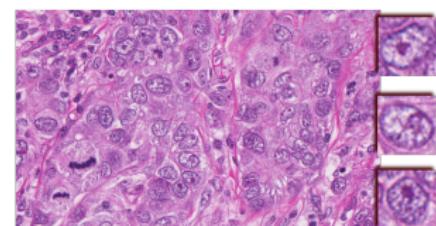
- The current research is on automatic strategies to quantify nuclear pleomorphism in breast cancer(NPBca).



grade 1



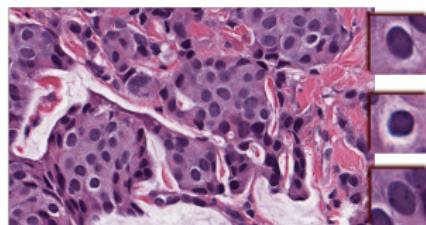
grade 2



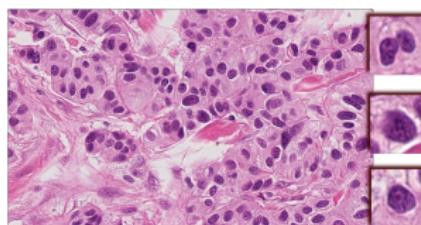
grade 3

Nuclear Pleomorphism

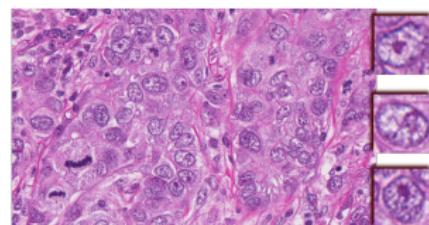
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- NPBca is an indicator of the aggressiveness of the disease



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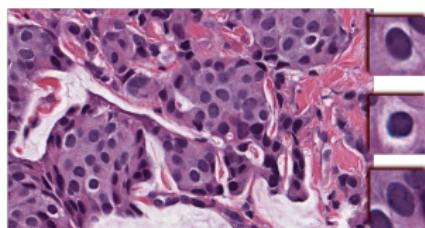
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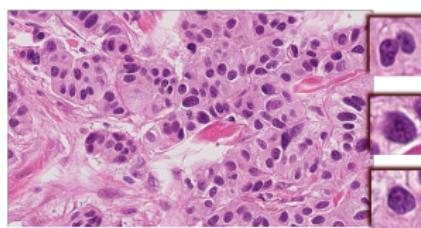
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Nuclear Pleomorphism

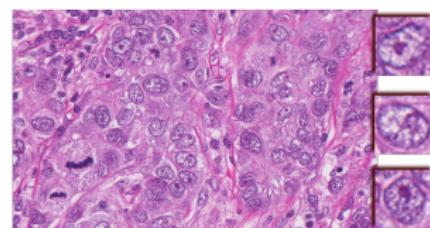
- The current research is on automatic strategies to quantify nuclear pleomorphism in breast cancer(NPBca).
- NPBca is an indicator of the aggressiveness of the disease
- NP has a low inter-observer agreement ($0.3 < \kappa < 0.5$)



grade 1



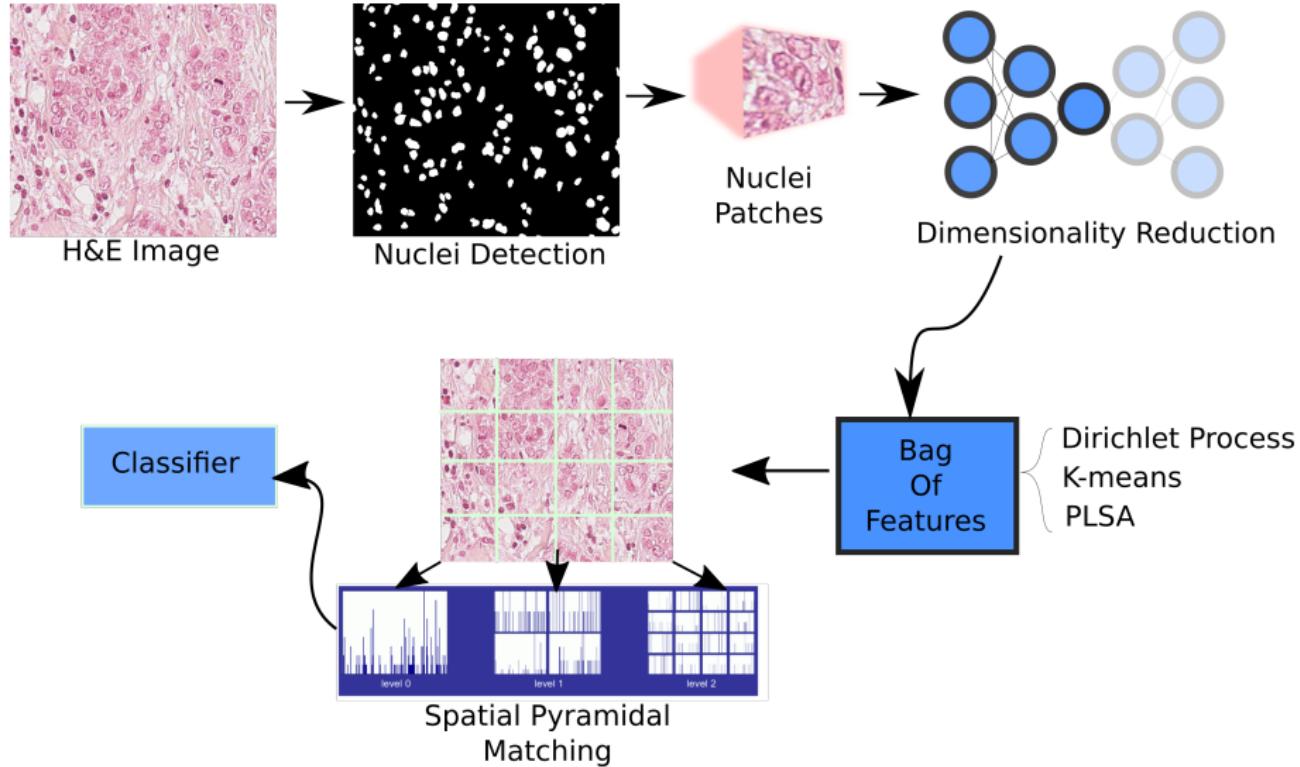
grade 2



grade 3

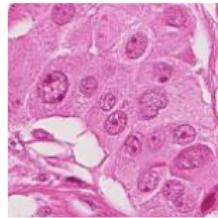
Proposed Framework

Spatial Pyramidal Characterization (SPYC)



Nuclei Detection

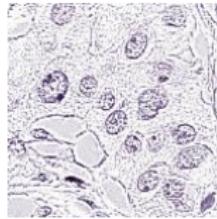
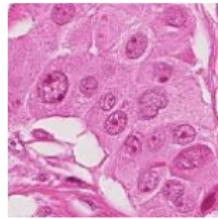
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¹ A method for normalizing histology slides for quantitative analysis, Macenko et al. ISBI 2009

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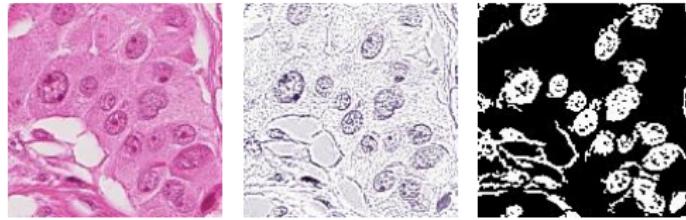
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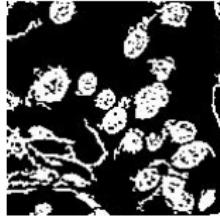
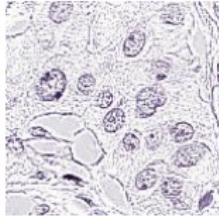
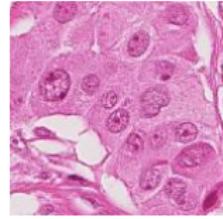
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- Watershed algorithm is used to detect nuclei candidates in H stain



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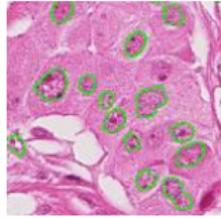
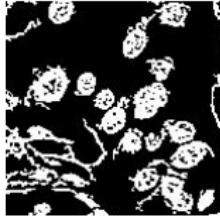
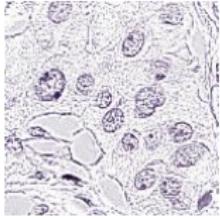
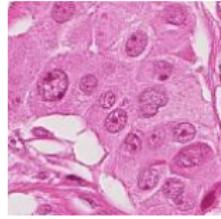
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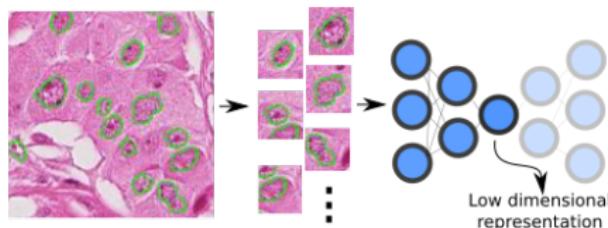
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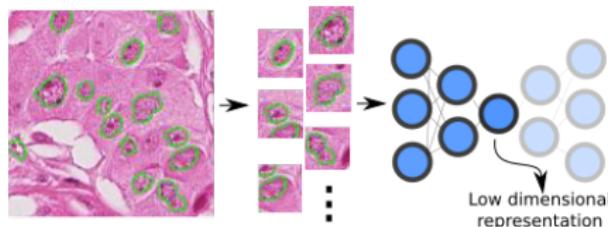
Nuclei Patches And Dimensionality Reduction

- A patch of $n \times n$ pixels around the candidate nucleus is extracted



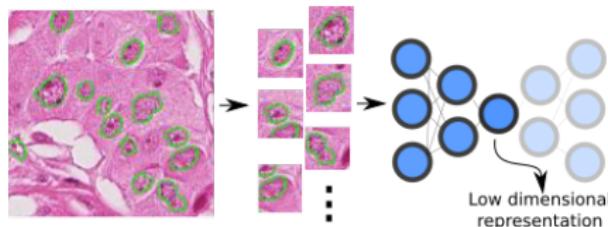
Nuclei Patches And Dimensionality Reduction

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- An autoencoder with three convolutional layers for encoding and 3 for decoding

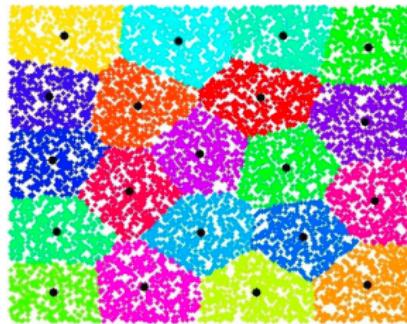


Nuclei Patches And Dimensionality Reduction

- A patch of $n \times n$ pixels around the candidate nucleus is extracted
- An autoencoder with three convolutional layers for encoding and 3 for decoding
- The coded representation for each nucleus is a vector of 2048 dimensions (22.5 compression factor)



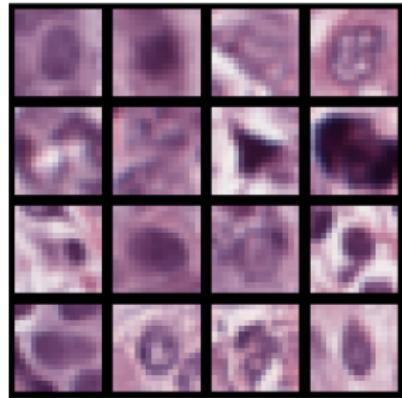
Bag of Features Building



- The multi-scale feature space is partitioned with a k -means algorithm

Figure: *K-means Algorithm*

Bag of Features Building



- The multi-scale feature space is partitioned with a k -means algorithm
- Each centroid correspond to a visual word in the BoF

Figure: *Decoded Dictionary Representation*

Bag of Features Building

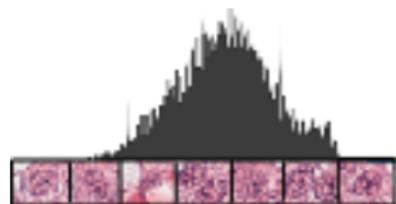


Figure: *Ocurrence representation*

- The multi-scale feature space is partitioned with a k -means algorithm
- Each centroid correspond to a visual word in the BoF
- Each new candidate from a train or test image is represented by some atom in the dictionary, building an histogram.

Spatial Pyramidal Matching

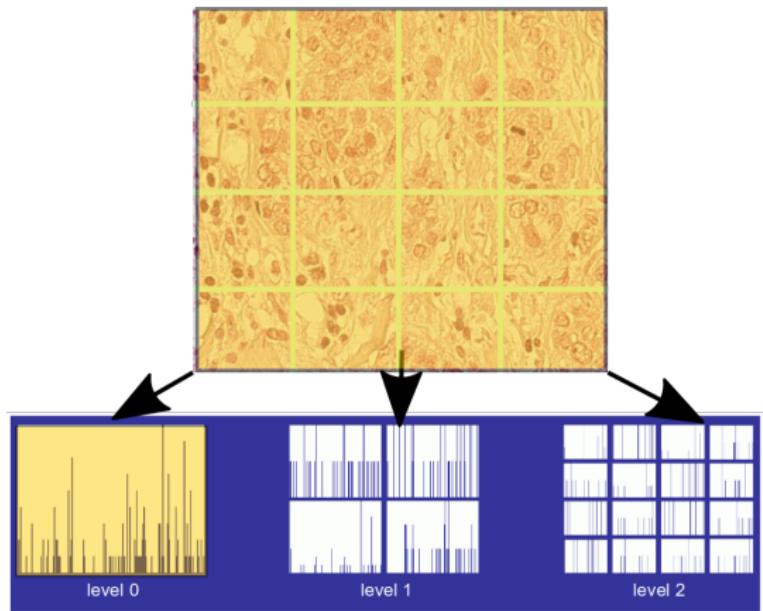
- Locally representation at several levels of resolution

Spatial Pyramidal Matching

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- The construction is as follow:

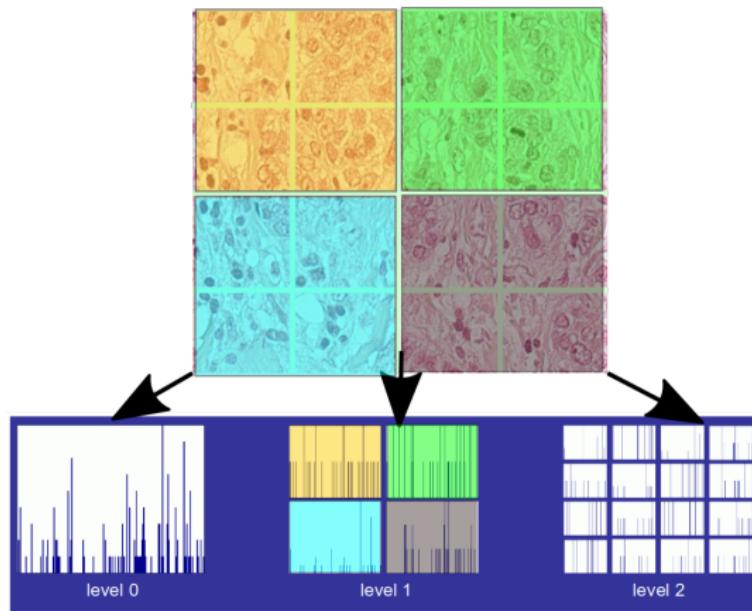
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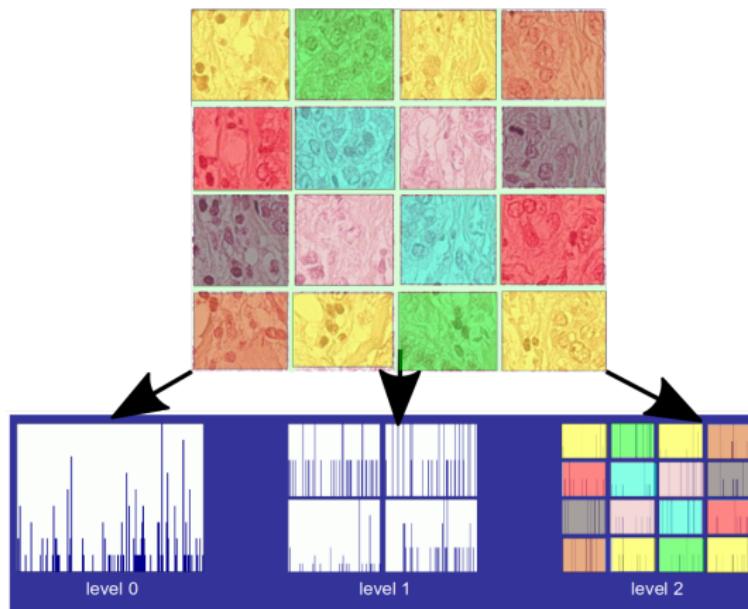
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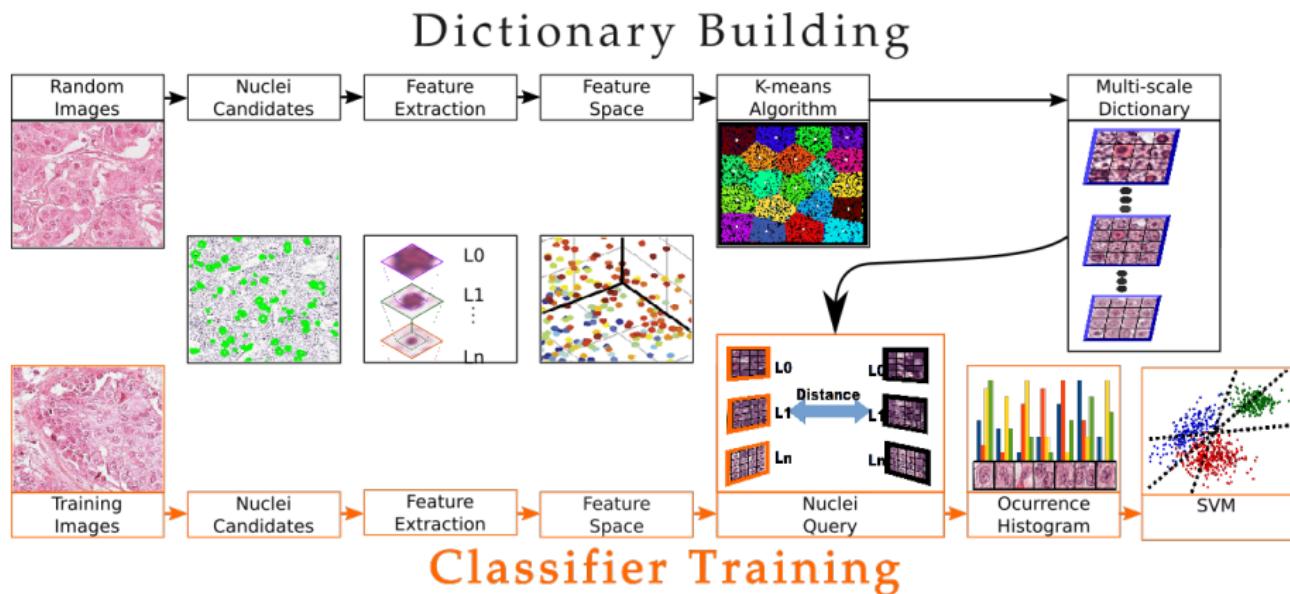


Spatial Pyramidal Matching

- Locally representation at several levels of resolution
- The construction is as follow:
- The histograms are concatenated in a single vector and feed the classifier

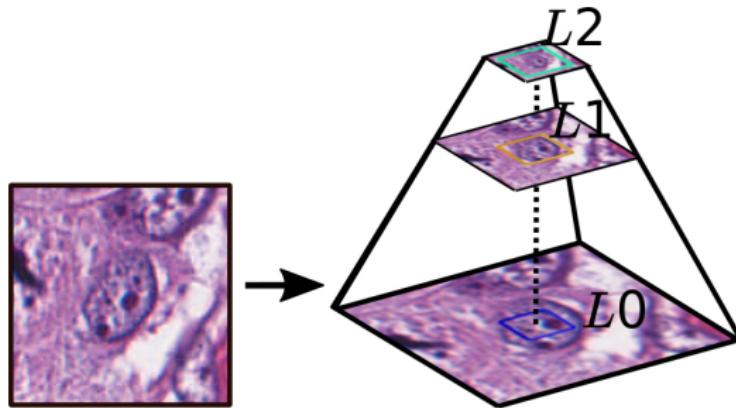
$$f(Y) = [Hist(I0), hist(I1), hist(I2)]$$

Bag of Features with Multiscale Descriptor Baseline Framework(BoF)



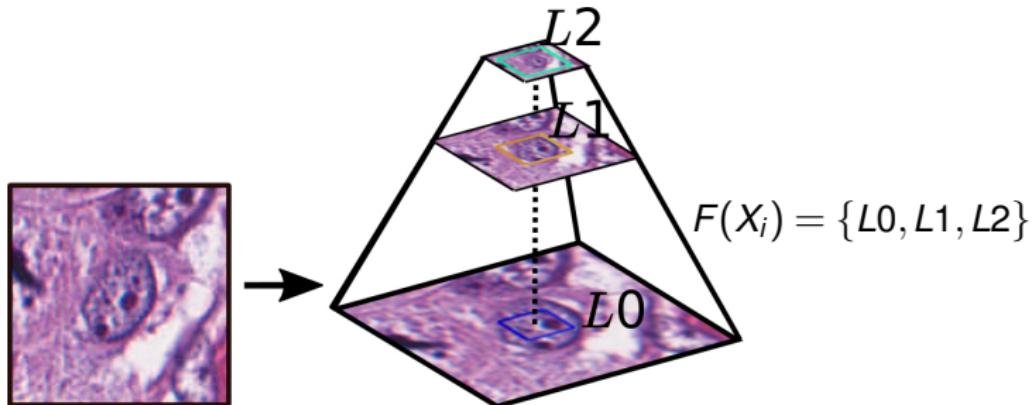
Multi-scale Feature Extractor

- The characterization of each nucleus candidate was performed by analyzing multiple scales

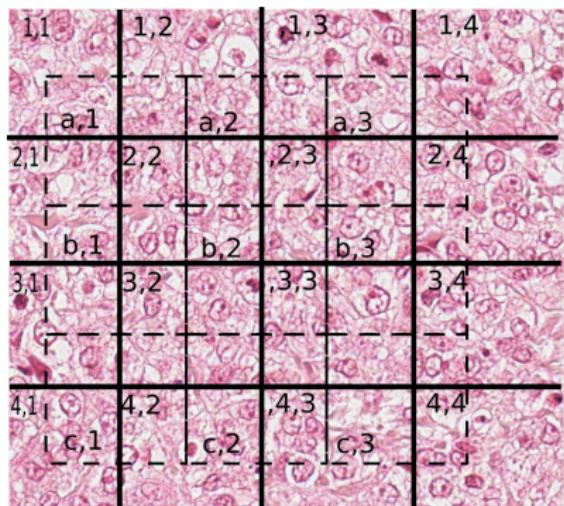


Multi-scale Feature Extractor

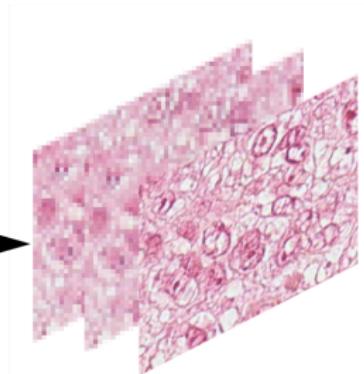
- The characterization of each nucleus candidate was performed by analyzing multiple scales
- The feature vector corresponds to the information from RGB patches which are concatenated along one dimension.



Convolutional Neural Networks Baseline Framework (CNN)



Flip,
Rotations,
color



CNN

Classifier

$[V_{1,1}, V_{1,2}, \dots, V_{4,4}]$

CNN Baseline Framework

- The Image is divided in several patches

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- The feature vector is used to feed a classifier

Experiments and Results

Experiments

- Public database - Mitos Atypia 14
- Patches of 72x72 pixels are extracted using the nuclei centroid as seed
- SPYA framework was evaluated also using each level independently and altogether

Dataset- Mitos Atypia 14

| Original | | After Augmentation | | |
|----------|-------------|---------------------------------|--------|------------|
| Grade | #40X images | Total Patches 344x344 pixels | Train | Validation |
| NP 1 | 92 | 21,344 | 18,144 | 3,200 |
| NP 2 | 900 | 22,500 | 19,096 | 3,404 |
| NP 3 | 208 | 20,800 | 17,764 | 3,036 |

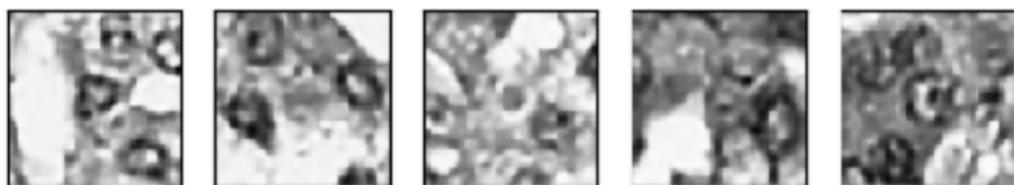
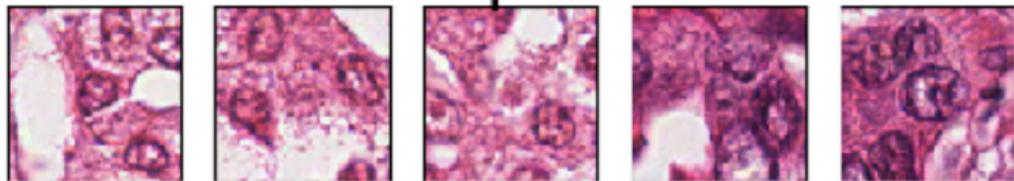
Table: *Training and validation - 11 cases for training*

| Grade | # 40X images |
|-------|--------------|
| NP 1 | 152 |
| NP 2 | 252 |
| NP 3 | 92 |

Table: *Test dataset - 5 cases*

Visual Performance of the Autoencoder

Input



Output

Autoncoder Reconstruction

Results using F1-Score

The F-score is used to evaluate the classification

$$F_{\beta} = (1 + \beta^2) \cdot \frac{\text{precision}.\text{recall}}{(\beta^2 \cdot \text{precision}) + \text{recall}} \quad (1)$$

$\beta = 1$ is used this is the harmonic mean between precision and recall

Results - Proposed Framework

| Task | SPYA | SPYA L0 | SPYA L1 | SPYA L2 |
|---------------|--------------|--------------|---------|--------------|
| 3 Vs. 1 and 2 | 0.578 | 0.630 | 0.565 | 0.530 |
| 1 Vs. 2 and 3 | 0.624 | 0.564 | 0.520 | 0.489 |
| 1 Vs 2 | 0.473 | 0.597 | 0.542 | 0.573 |
| 2 Vs 3 | 0.560 | 0.603 | 0.606 | 0.632 |
| 1 Vs 3 | 0.623 | 0.716 | 0.639 | 0.632 |

Table: *Results using a coded representation of the nuclei*

Results of CNN Baseline

| | CNN (1024dim) | CNN (3dim) | VGG16 | Alexnet | Resnet50 |
|----------------|------------------|---------------|-------|---------|--------------|
| 3Vs.1 and 2 | 0.635 | 0.523 | 0.629 | 0.493 | 0.685 |
| 1Vs.2 and 3 | 0.676 | 0.645 | 0.564 | 0.581 | 0.573 |
| 1 Vs 2 | 0.637 | 0.664 | 0.453 | 0.623 | 0.592 |
| 2 Vs 3 | 0.570 | 0.567 | 0.544 | 0.619 | 0.670 |
| 1 Vs 3 | 0.666 | 0.672 | 0.620 | 0.649 | 0.563 |

Results of BoF multiscale descriptor

| Experiment/S.Patch | 32x32 L2 | 20x20 L2 |
|--------------------|---------------|---------------|
| 1 vs 2 and 3 | 0.593 | 0.434 |
| 3 vs 1 and 2 | 0.642 | 0.47 |
| 1 vs 2 | 0.7128 | 0.47 |
| 1 vs 3 | 0.660 | 0.7029 |
| 2 vs 3 | 0.6759 | 0.5813 |

Table: *Results Bag Of Features*

Results of BoF multiscale descriptor

| | | |
|--------------------|-------------|-------------|
| Experiment/S.Patch | 32x32 L2 | 20x20 L2 |
|--------------------|-------------|-------------|

The BoF with the multiscale descriptor presents a better performance than the SPYA!, a new experiment of this without autoencoder is evaluated...

| | | |
|--------|---------------|---------------|
| 1 vs 3 | 0.660 | 0.7029 |
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Table: *Results Bag Of Features*

Results without Autoencoder

| | BOF PY | BOF PY L0 | BOF PY L1 |
|---------------|--------------|--------------|--------------|
| 3 Vs. 1 and 2 | 0.624 | 0.593 | 0.583 |
| 1 Vs. 2 and 3 | 0.692 | 0.642 | 0.678 |
| 1 Vs 2 | 0.650 | 0.713 | 0.660 |
| 2 Vs 3 | 0.722 | 0.660 | 0.729 |
| 1 Vs 3 | 0.628 | 0.675 | 0.740 |

Results without Autoencoder

| | BOF PY | BOF PY L0 | BOF PY L1 |
|---------------|--------------|-----------|-----------|
| 3 Vs. 1 and 2 | 0.624 | 0.593 | 0.583 |
| 1 Vs. 2 and 3 | 0.692 | 0.642 | 0.678 |
| 1 Vs. 0 | 0.650 | 0.710 | 0.660 |

Then, we obtain a better model of NP using the spatial pyramidal representation approach.

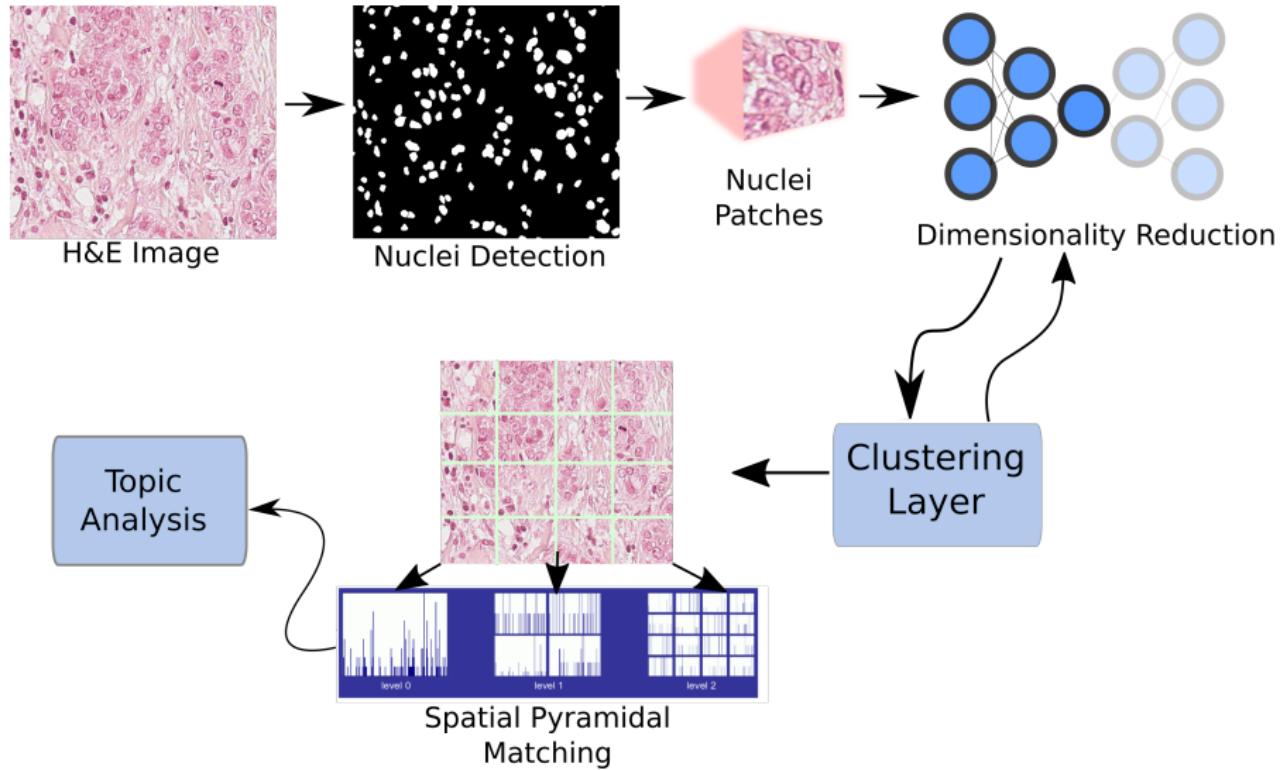
Final Remarks

- The Slightly variation between the NP grades and the high tissue variability makes it challenging to find an adequate model
- The CNN model could be improved if take advantage of the spatial information
- The auto-encoder representation and vocabulary construction maybe are not the most accurate

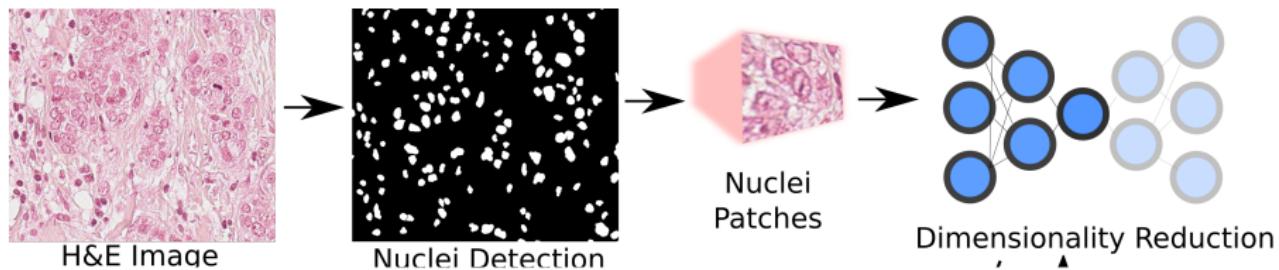
Current Work

- To determine a better cluster representation
- To evaluate the frameworks in a new database
- To determine if there exists a relationship between NP grading and the cancer recurrence (Future work)

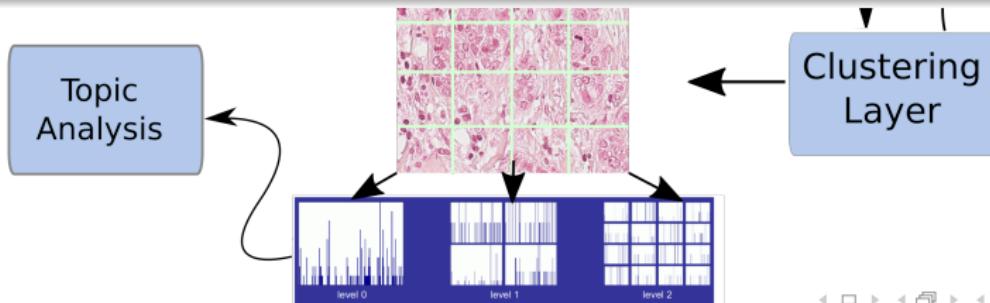
New proposal



New proposal

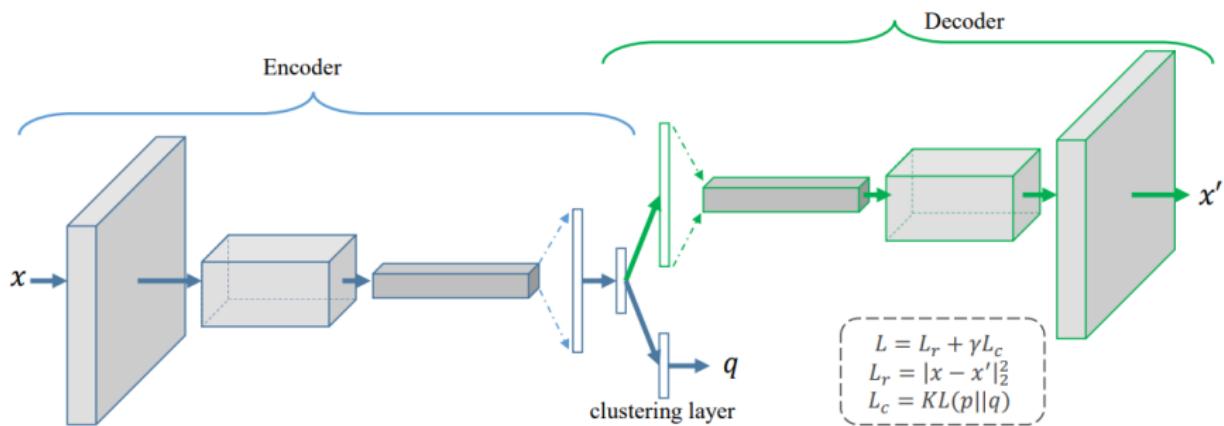


Two main differences of the previous approach are presented: 1) The clustering step, 2) A topic representation



Clustering Step

- The proposed deeplearning architecture² minimize the reconstruction loss of the autoencoder and the clustering loss
 - Intends to preserve the structure of the data



²Guo et al., ICONIP17

New Dataset

- 300 cases of DCIS with recurrence time from Sunnybrook Health Institute
- 306 marked regions from 24 cases grading by an expert pathologist

| Grading | # Marked Areas |
|--------------------|----------------|
| Grade 1 | 135 |
| Grade 2 | 132 |
| Grade 3 | 39 |
| Total Cases | 24 |

Thank you...