

# Automated Mitosis Detection in Color and Multi-spectral High-Content Images in Histopathology: Application to Breast Cancer Grading in Digital Pathology

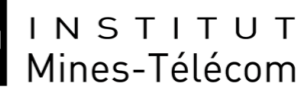
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Co-Supervisor: Dr. Ludovic ROUX

20 Jan, 2014

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## ▶ Context

- Digital Pathology
- Breast Cancer Grading

## ▶ Research Contributions

- Mitosis Detection Framework for Color Images
- Mitosis Detection Framework for Multispectral Images
- Dynamic Sampling Framework for WSI analysis

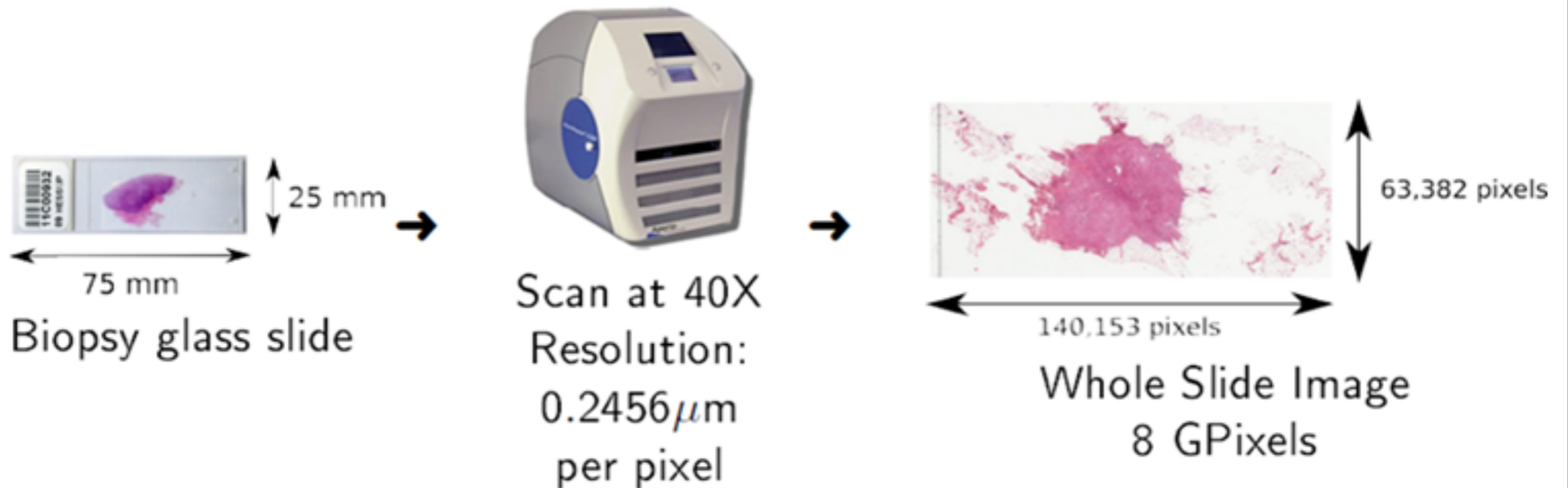
## ▶ Conclusion & Future works

## Slide Preparation

- Dissection / extraction
- Chemical processing
- Sectioning
- Staining - Hematoxylin & Eosin (H&E),  
Immunohistochemistry (IHC)

## Imaging & Visualization

- Microscope
- Slide Scanner

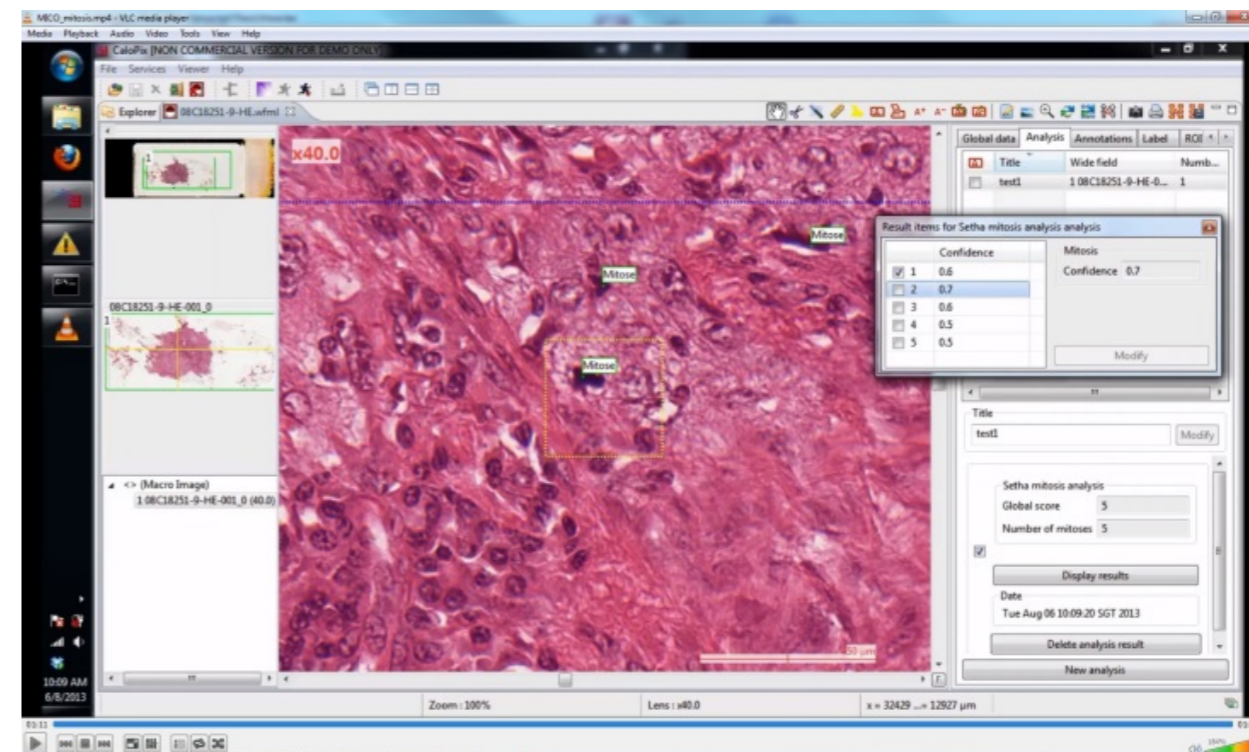


## Pathology

- Manual analysis is labor intensive work
- Inter and intra-reader variations

## Digital Pathology

- Location independence
- Virtual slide sharing with clinicians
- Enables automated analysis techniques



## Breast Cancer

- **Origin** – Ductal, lobule or stromal tissues
- **Proliferation** – Carcinoma in Situ, Invasive
- Worldwide, Breast Cancer accounts for 22.9% of all cancer in women [1]
- 1 in 8 US women estimated to develop Invasive Breast Cancer over the course of her lifetime [2]

### ➤ Nottingham Grading System for Invasive Breast Cancer

- International grading system recommended by the World Health Organization
- 3 Criteria [3]:
  1. Gland Formation
  2. Nuclear Atypia / Pleomorphism
  3. Mitosis Count

1) Peter Boyle, et al. World cancer report 2008. IARC Press, International Agency for Research on Cancer, 2008.

2) US Breast Cancer Statistics, 2013 ([http://www.breastcancer.org/symptoms/understand\\_bc/statistics](http://www.breastcancer.org/symptoms/understand_bc/statistics)).

3) C W Elston & I O Ellis, Pathological prognostic factors in breast cancer. Histopathology, 19(5):403-410, 1991.

## Mitosis Count

- Scan sections to find area with most mitotic activity (often at tumor edge)
- Count mitosis in 10 consecutive high power fields (HPFs) of selected area
- Skip fields with few carcinoma cells or obvious necrosis
  
- Score 1 ( < 10 Mitosis )
- Score 2 ( 10 ~ 19 Mitosis )
- Score 3 ( > 19 Mitosis )
  
- ❖ HPF is an area of microscope field diameter of 58mm (or a digitized square image 512 \* 512  $\mu\text{m}^2$ ).

1. State-of-the-art in nuclei detection, segmentation and classification

## 2. Color Framework

- Selection of color channels for different tissue component
- Intensity (1st order statistical) and texture (2nd order statistical) features
- Region vs patch based texture features analysis for mitosis discrimination
- An inspection of over-sampling method for balancing the training set

## 3. Multispectral Framework

- Spectral absorption responses of different tissue components
- Multispectral-statistical features in selected Spectral Bands (SBs)

## 4. Whole Slide Image Analysis Framework

- Robust strategy to explore WSI using a dynamic sampling framework
- Extension of ITK QuadEdgeMesh data structure to handle duality of meshes

## 5. Proof of concept in MICO platform

## MITOS Benchmark (ICPR 2012)

- 5 breast cancer biopsy slides (H&E stained) provided by [4]
- In each biopsy slide, 10 HPFs at 40X magnification are selected
- 35 Training HPFs 226 mitotic nuclei (69.3%)
- 15 Evaluation HPFs 100 mitotic nuclei (30.7%)

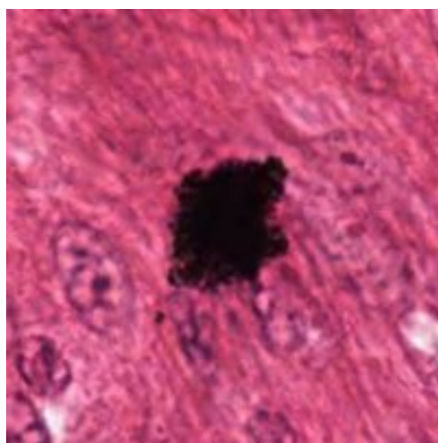
Scanners	Resolution per pixel	HPF dimension to cover area of $512 \times 512 \mu\text{m}^2$
Aperio Scanner	0.2456 $\mu\text{m}$	2084 $\times$ 2084 pixels
Hamamatsu Scanner	0.2273 $\times$ 0.22753 $\mu\text{m}$	2252 $\times$ 2250 pixels
Multispectral Microscope	0.185 $\mu\text{m}$	2767 $\times$ 2767 pixels

4) Team of Prof. Frederique Capron, head of Pathology department, Pitie-Salpetriere Hospital Paris.

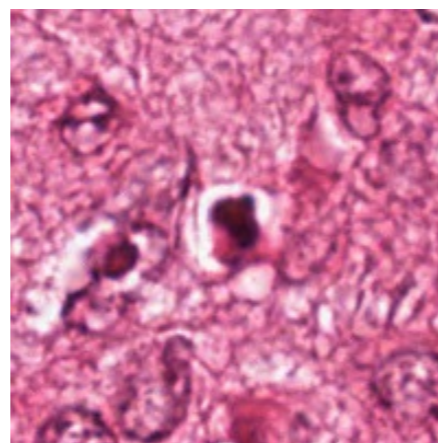


## Mitosis Detection – A Challenge

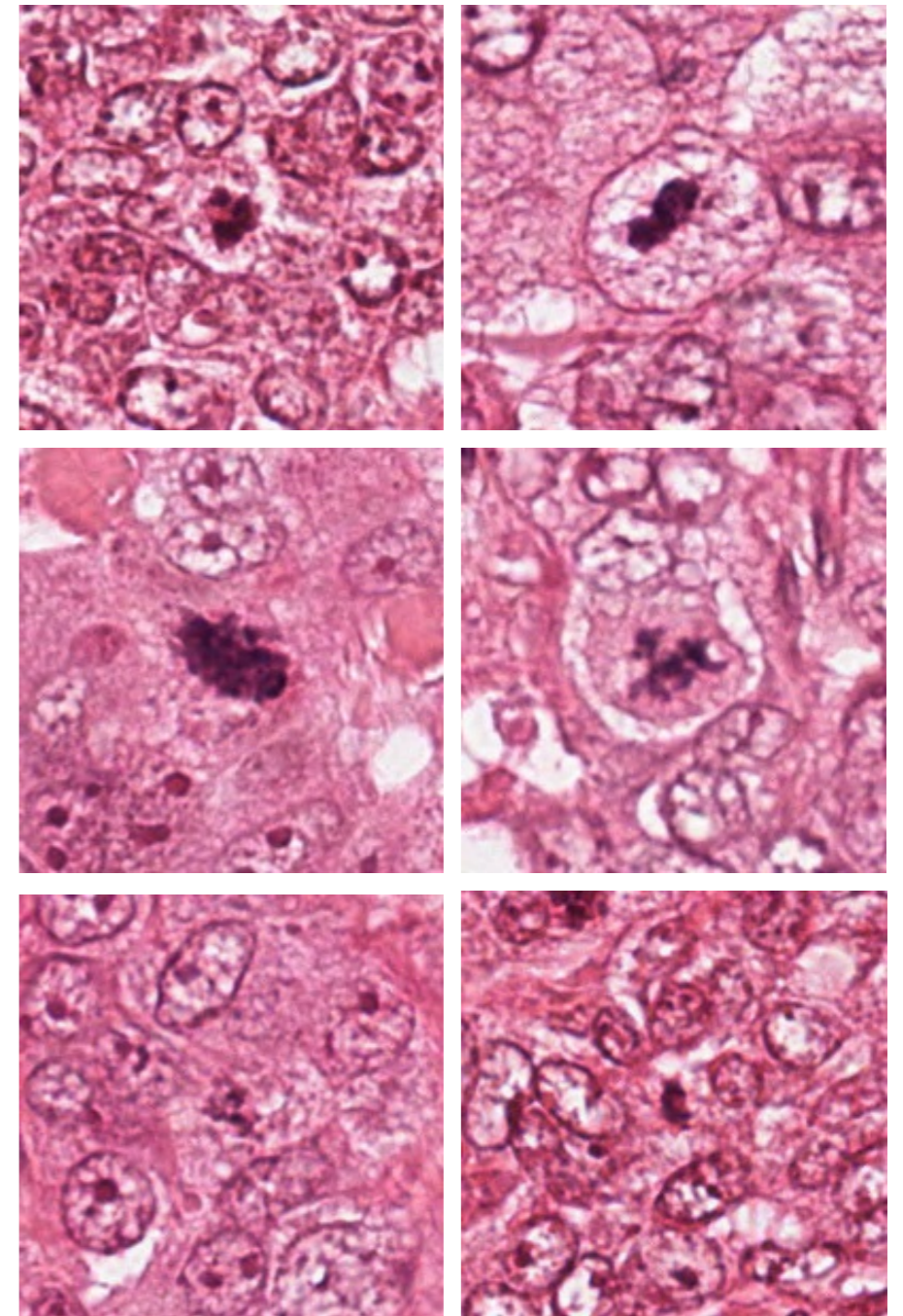
- Mitotic nuclei have
  - large variety of shape configuration
  - Texture variation
  - Low frequency of appearance in HPF
  - Similarity with other types of objects  
( e.g., apoptosis, necrosis, dust particles etc. )



Dust particles



Apoptosis

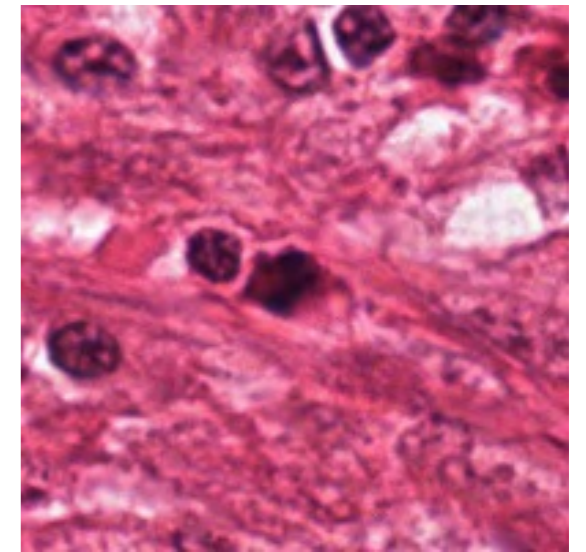
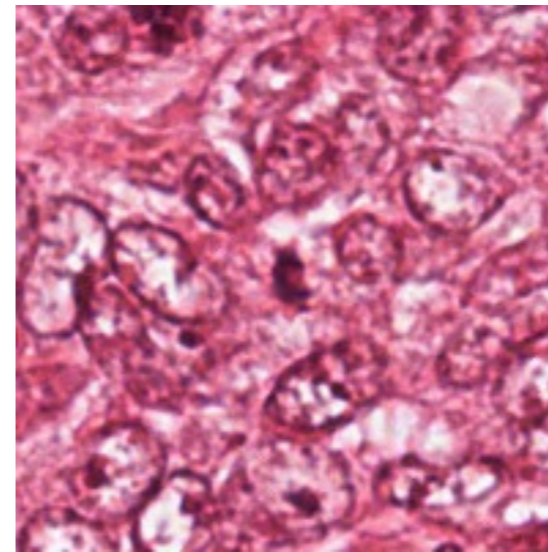
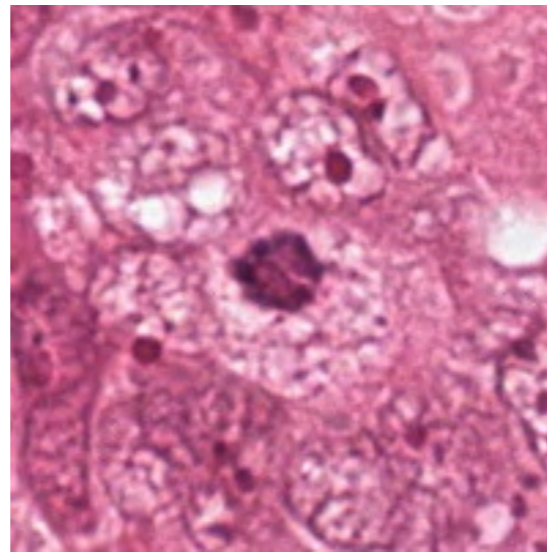
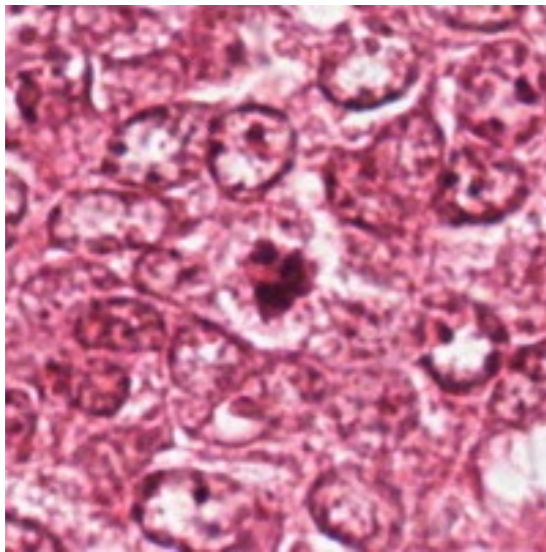


Mitosis

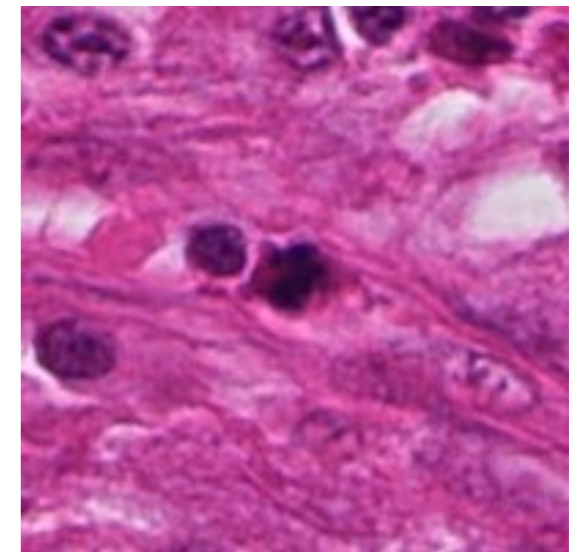
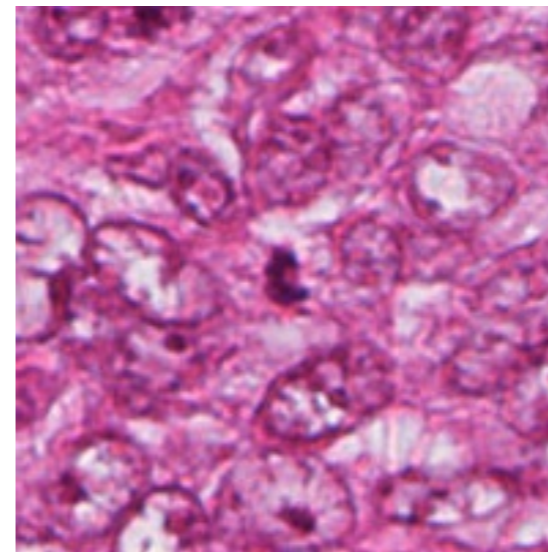
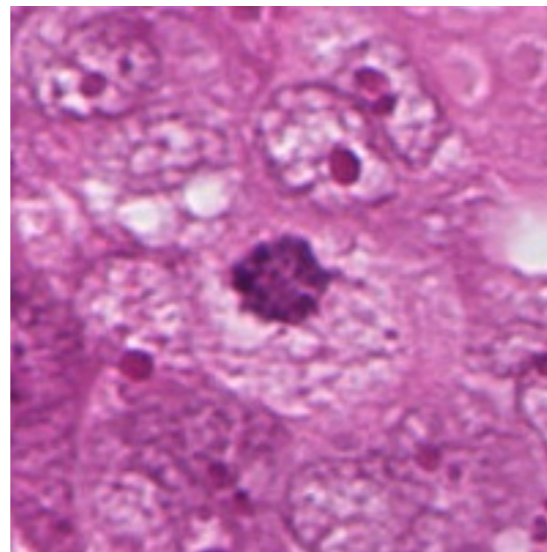
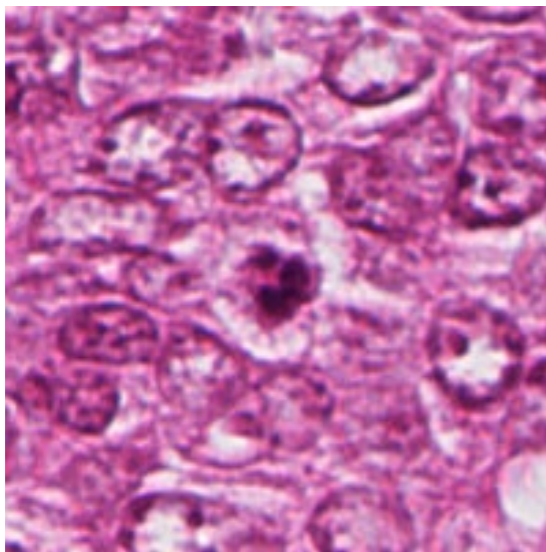
## Mitosis Detection – A Challenge

➤ Which one is mitotic nuclei and which is not?

Aperio



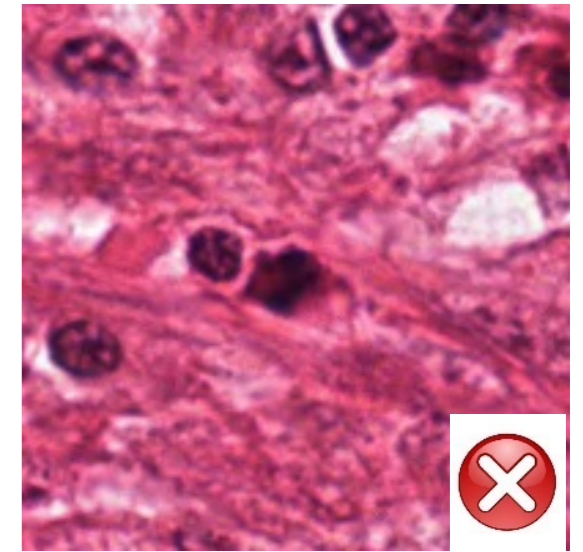
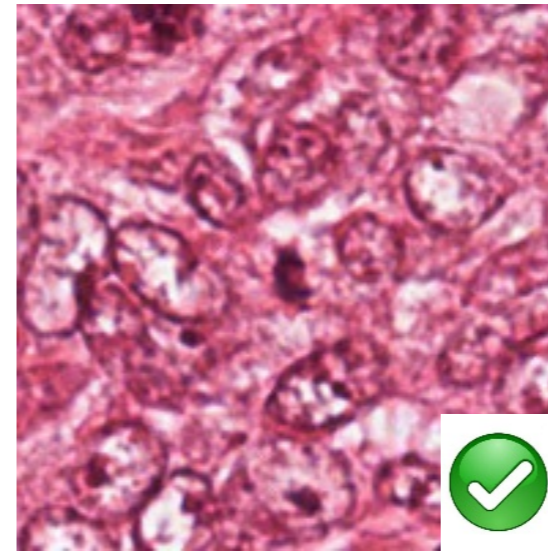
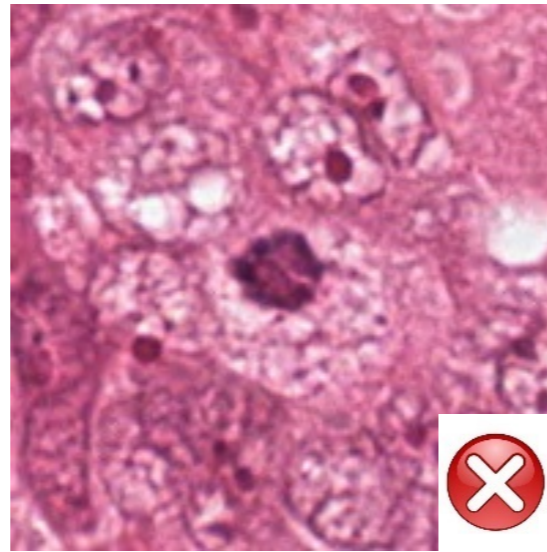
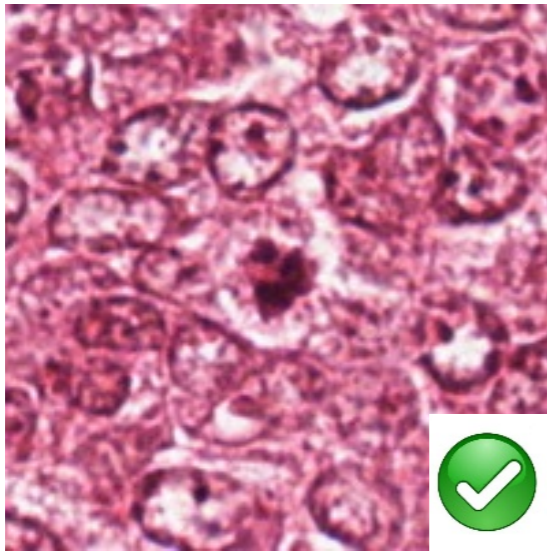
Hamamatsu



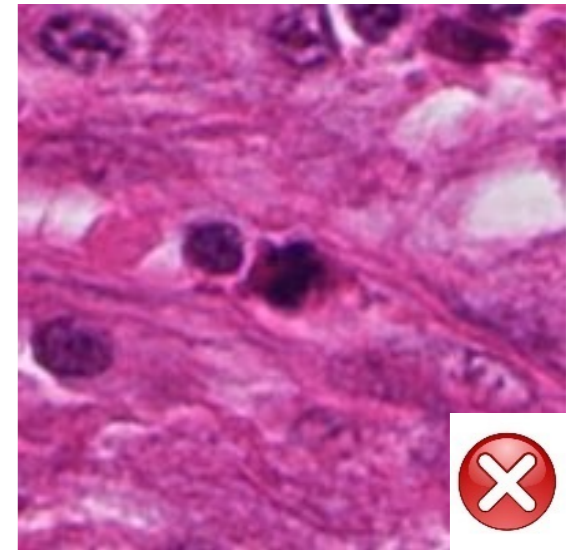
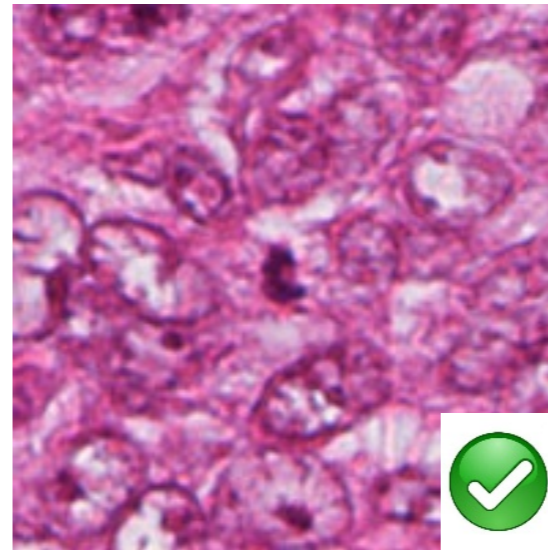
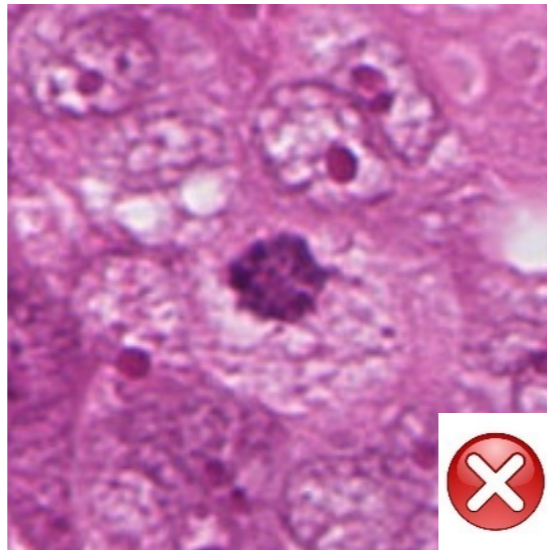
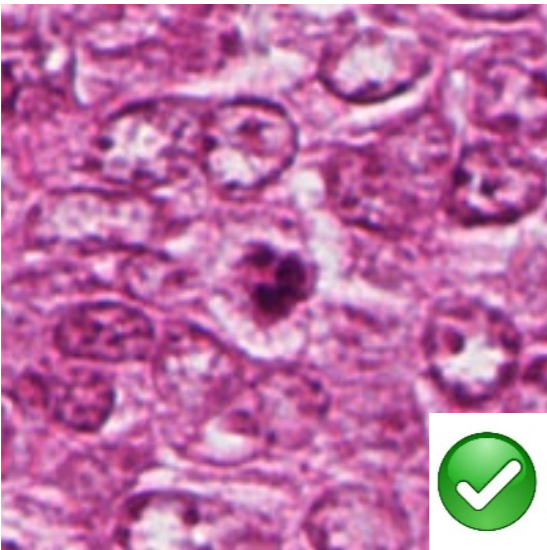
## Mitosis Detection – A Challenge

➤ Which one is mitotic nuclei and which is not?

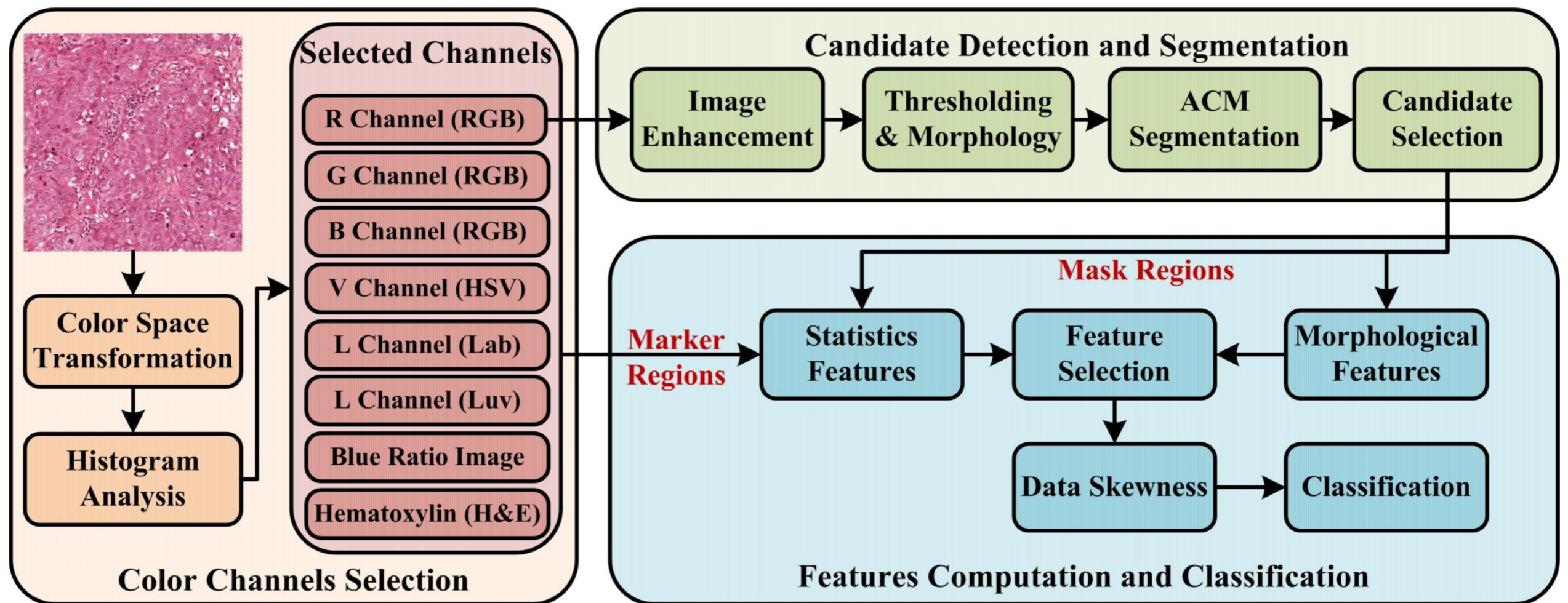
Aperio



Hamamatsu



## Intensity, Texture & Morphology based Mitosis detection in Color images (ITM<sup>2</sup>C) Framework

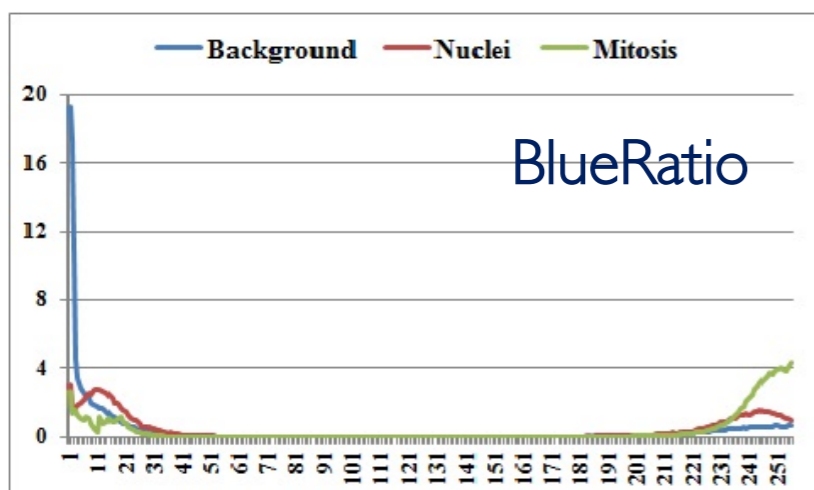
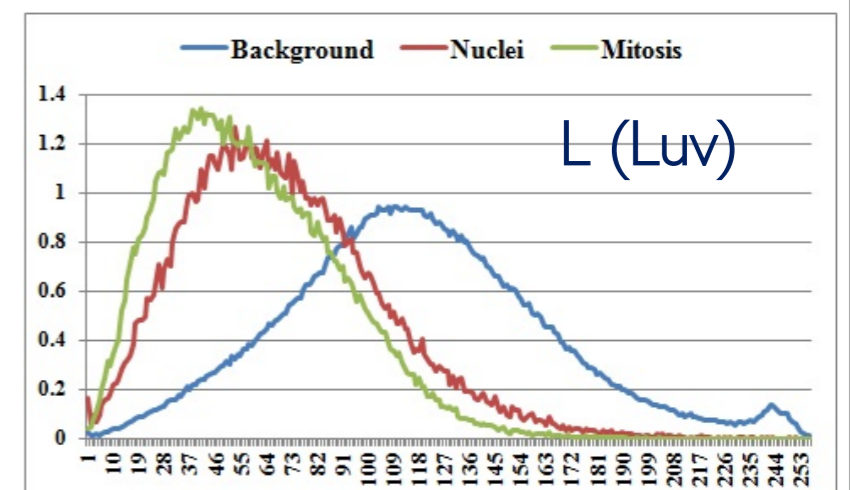
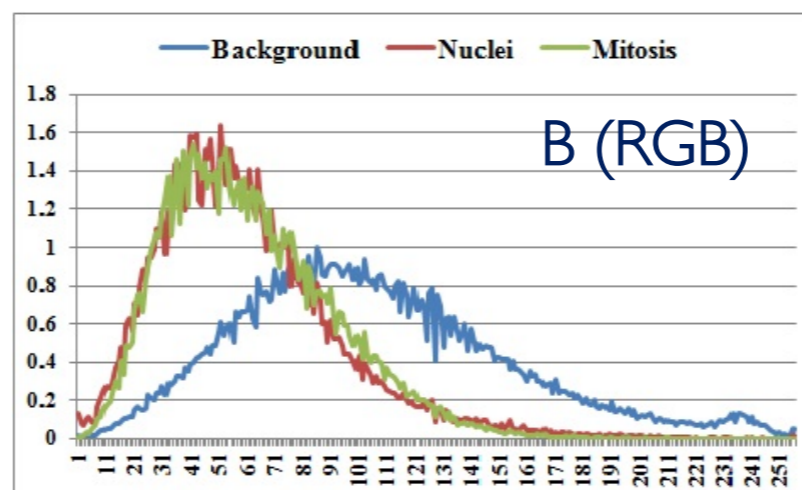
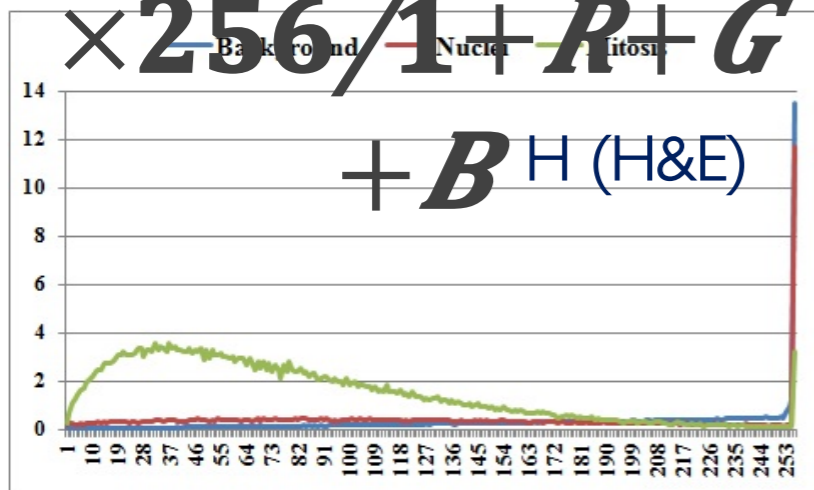
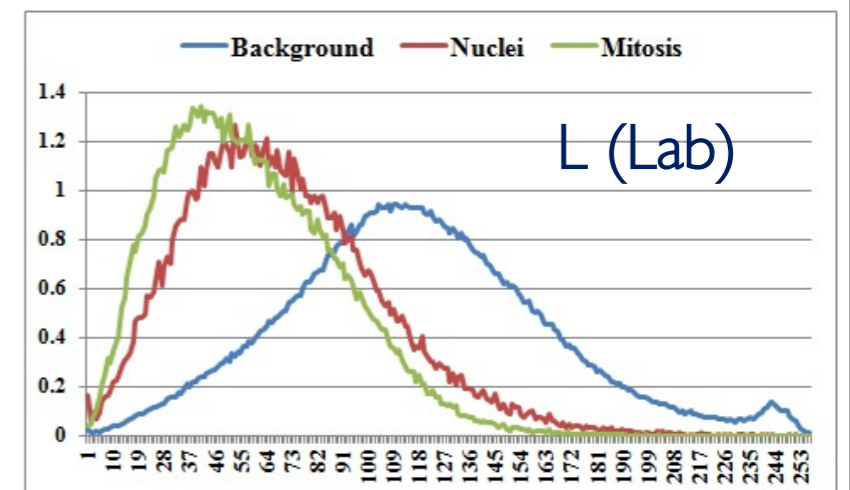
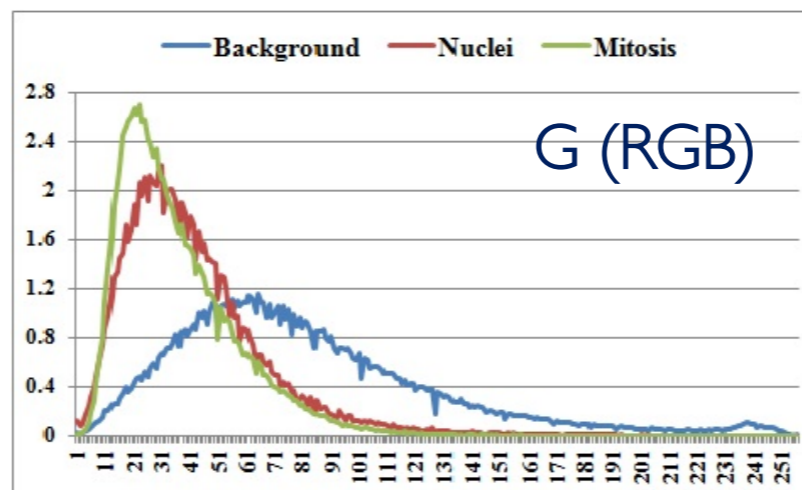
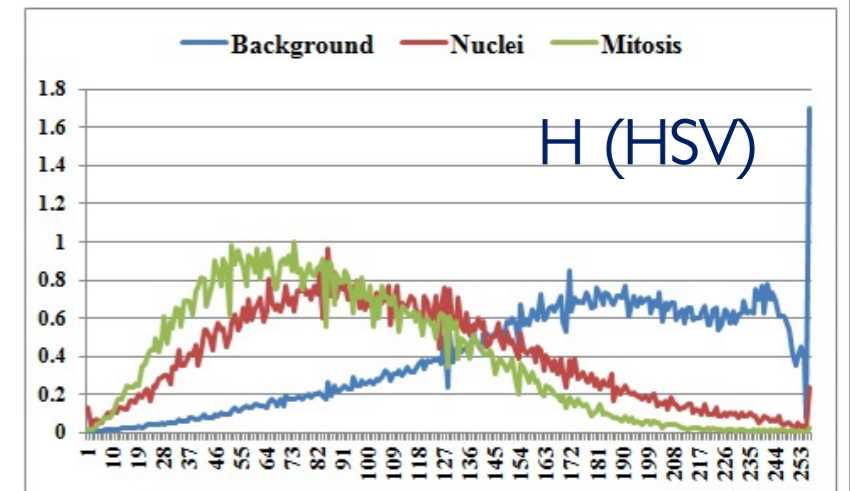
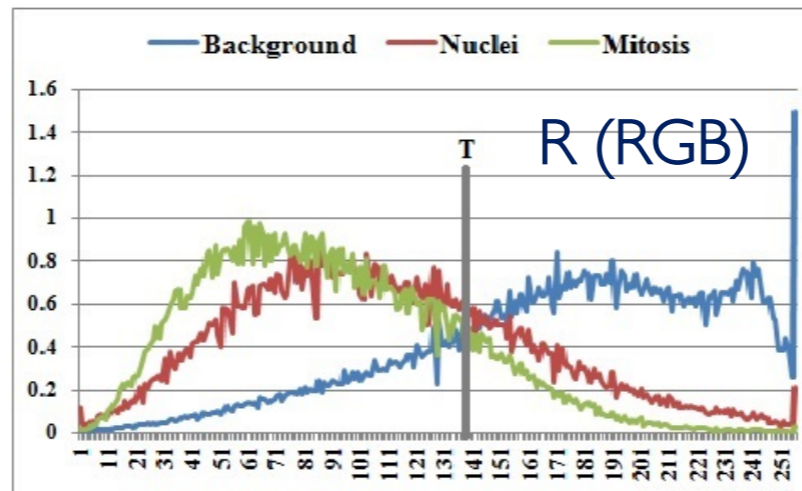


ITM<sup>2</sup>C Framework

## Histogram of Selected Channels on Aperio Dataset

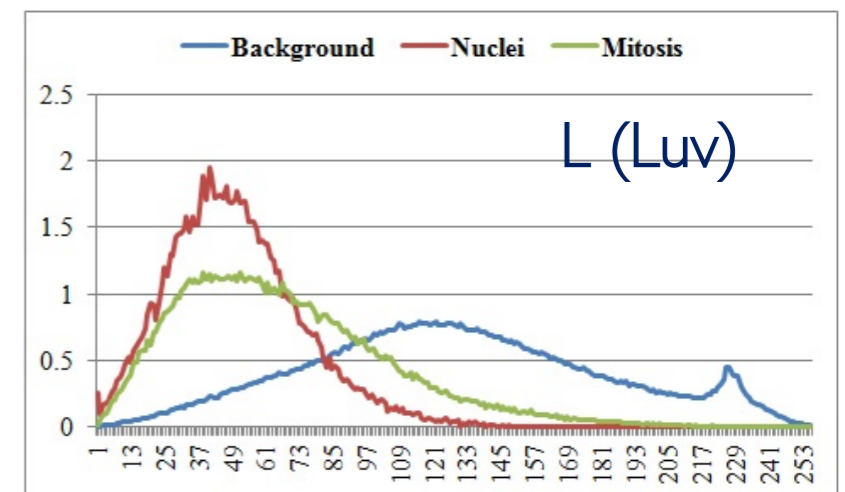
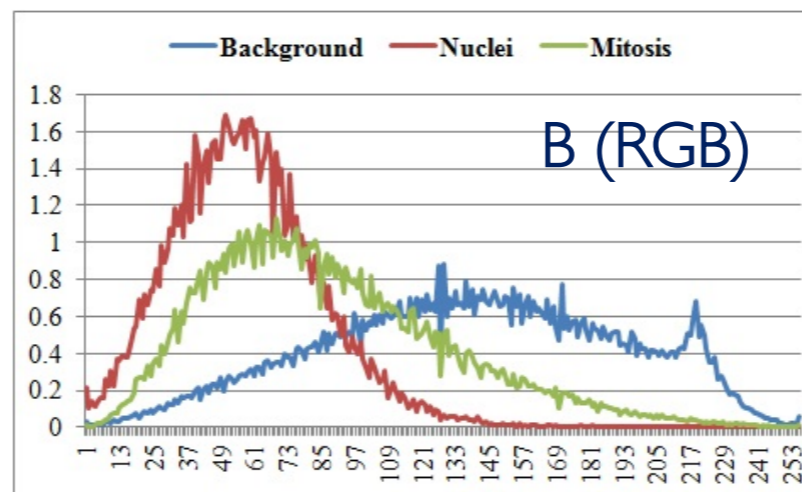
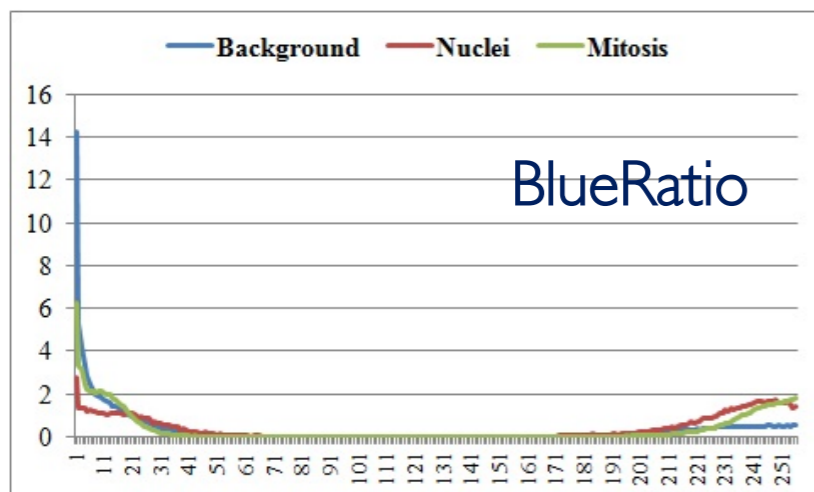
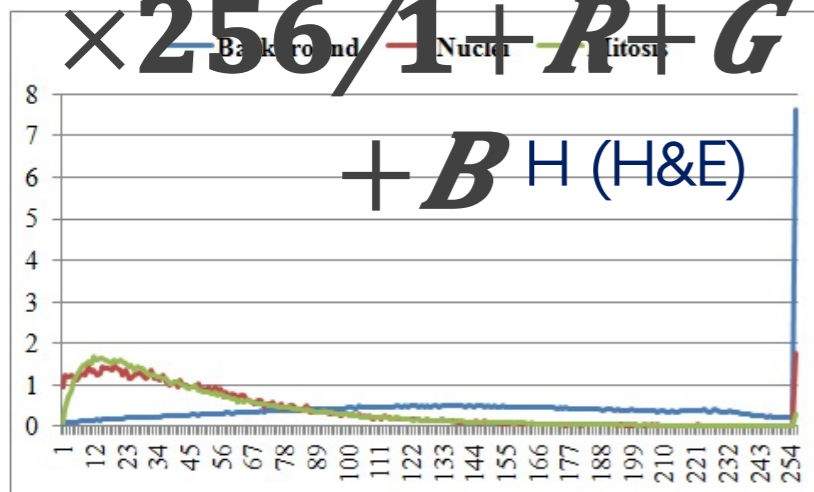
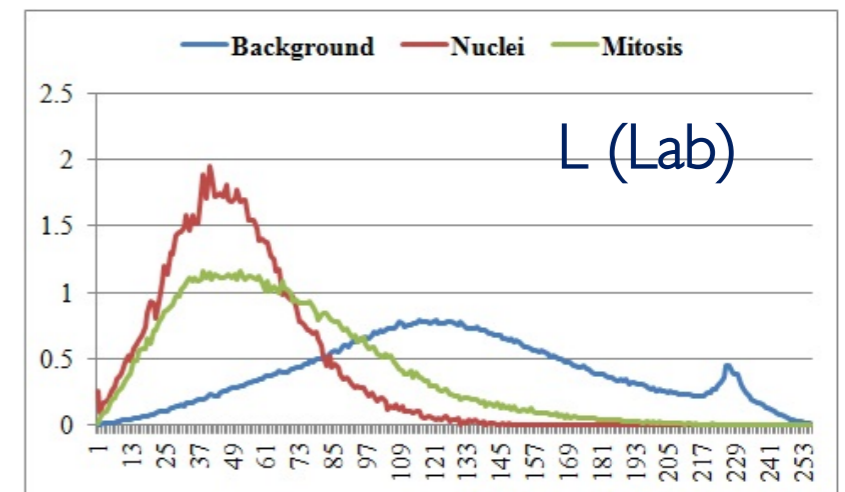
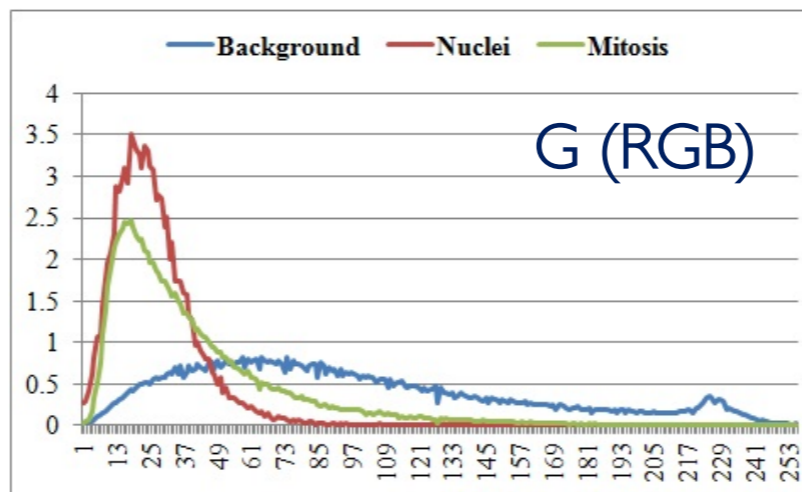
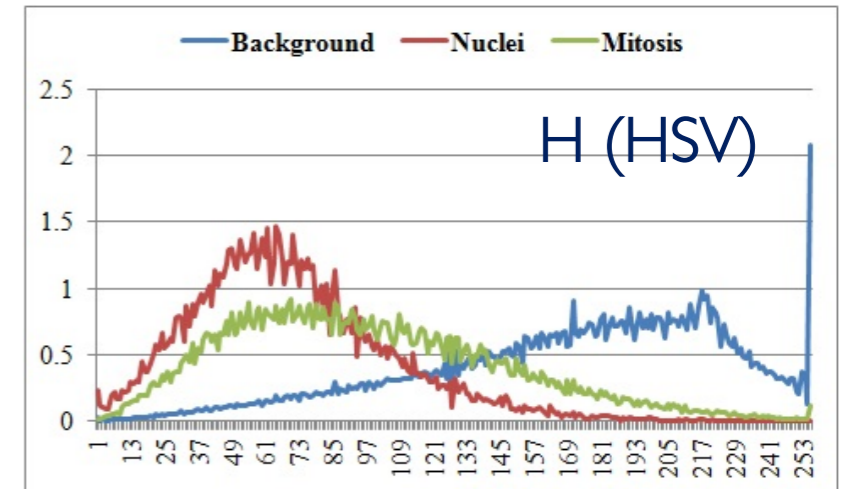
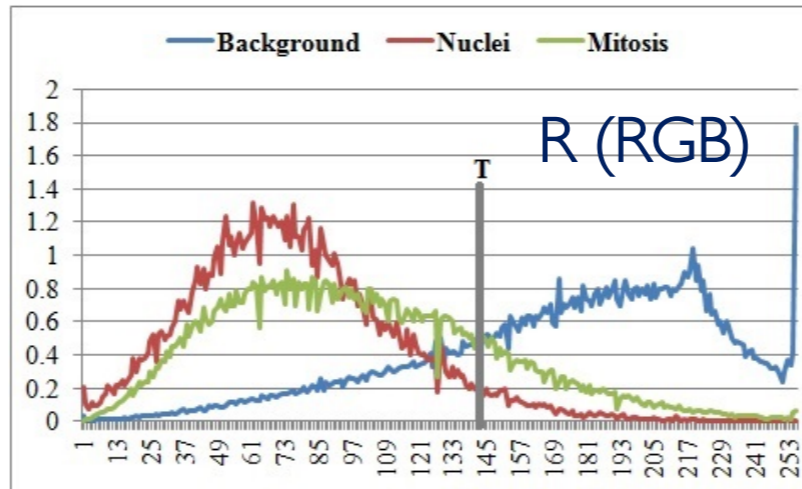
$$BlueRatio = \frac{100 \times B}{1 + R + G} \times \frac{256}{1 + R + G} + B$$

H (H&E)

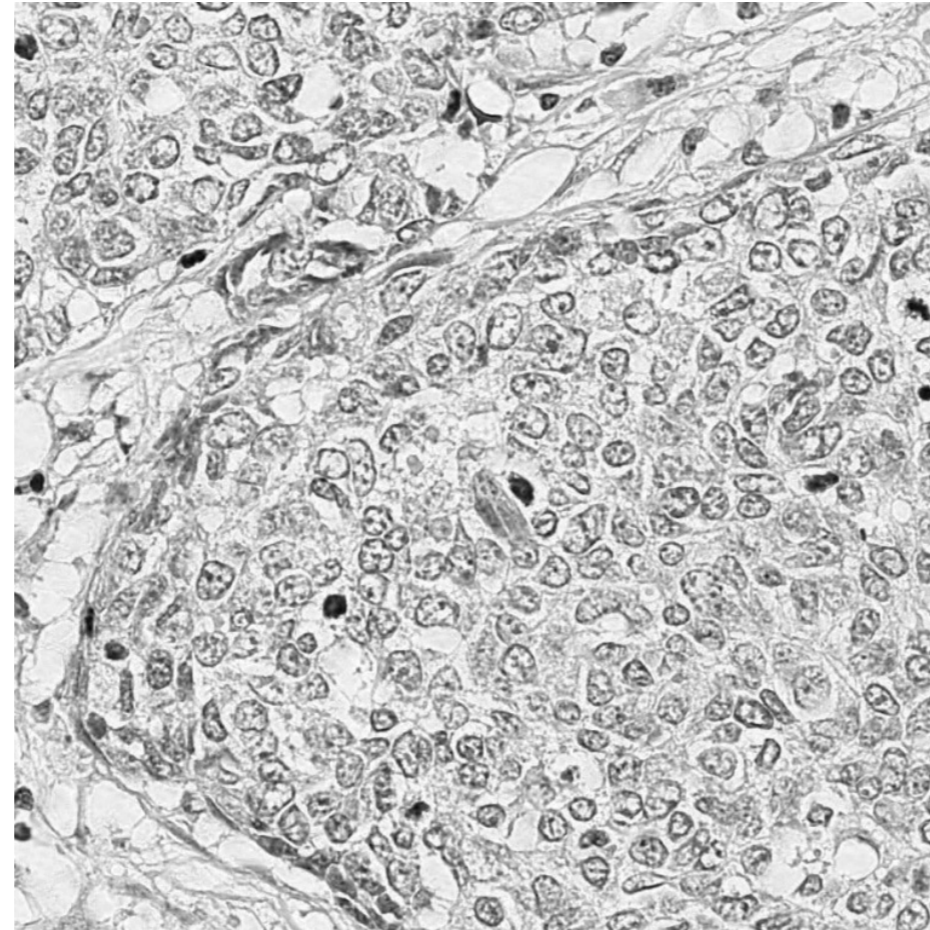
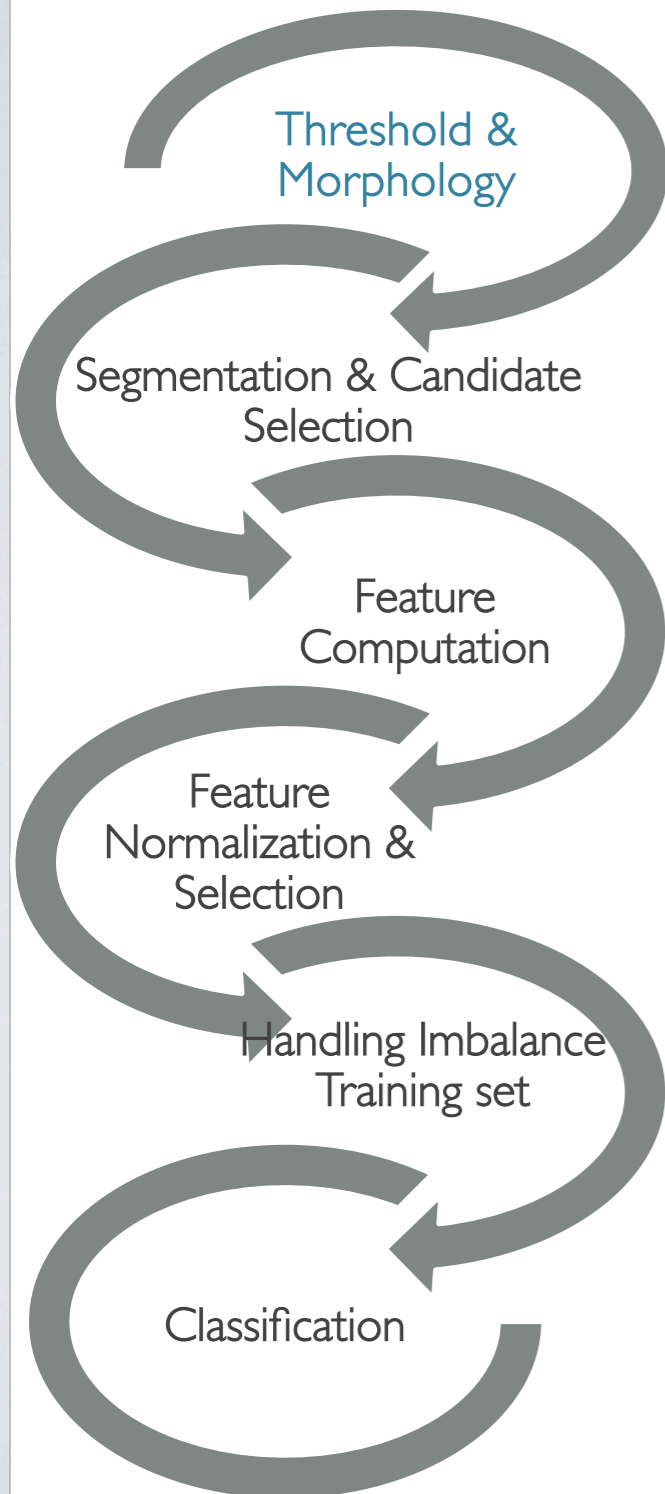


## Histogram of Selected Channels on Hamamatsu Dataset

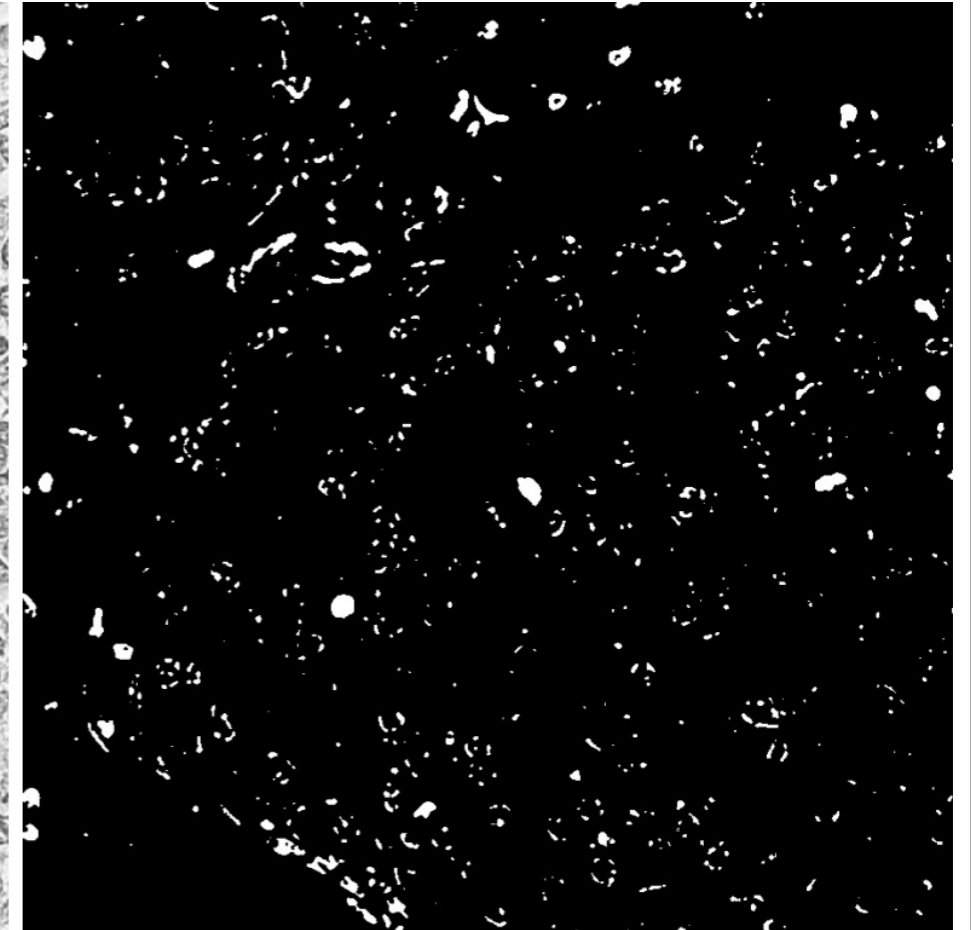
$$BlueRatio = \frac{100 \times B}{1 + R + G} \times \frac{256}{1 + R + G} + B^H \text{ (H\&E)}$$



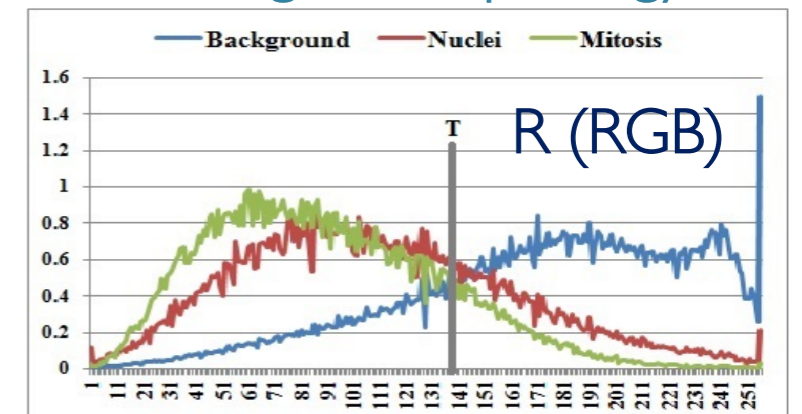
## ITM<sup>2</sup>C Framework Step 1



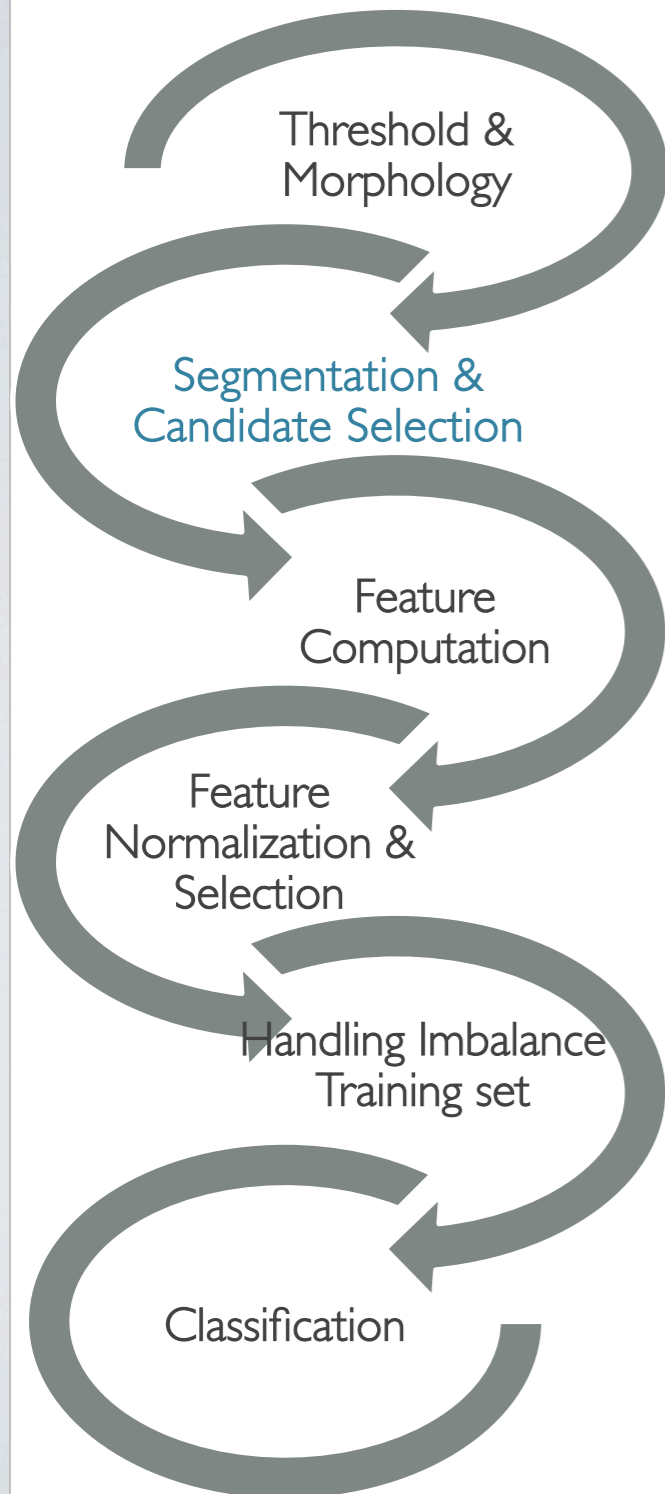
Red (RGB)



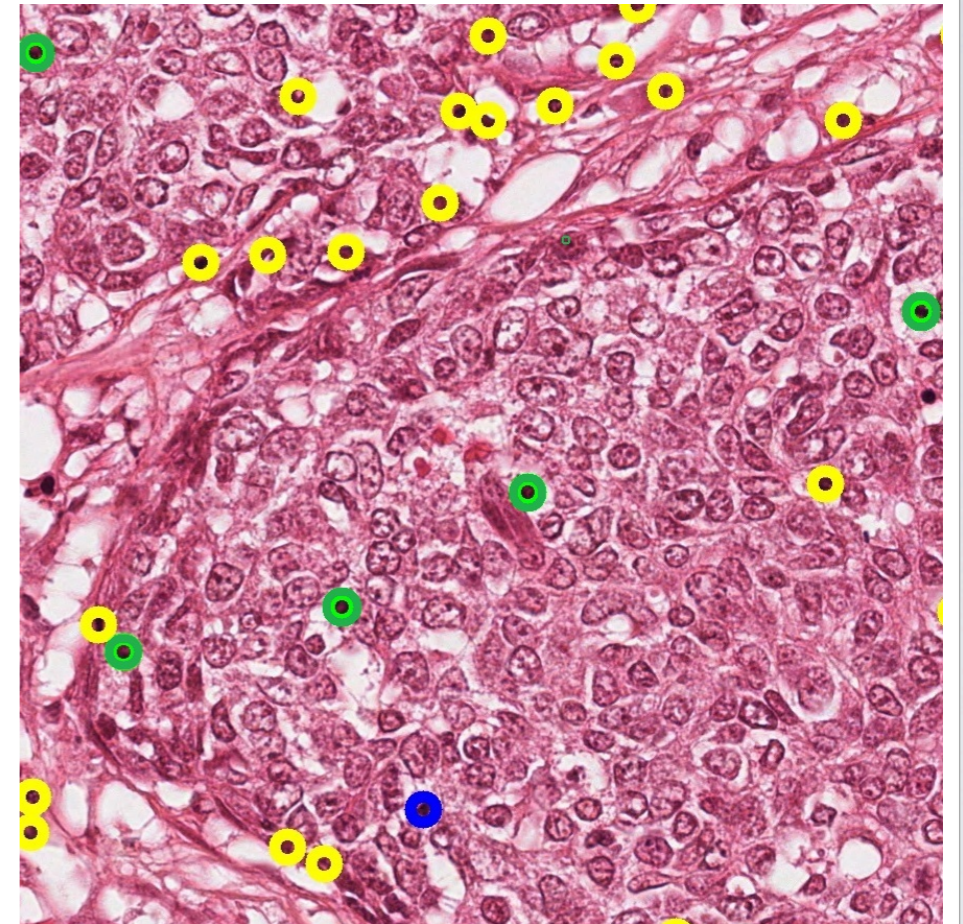
Thresholding & Morphology



## ITM<sup>2</sup>C Framework Step 2



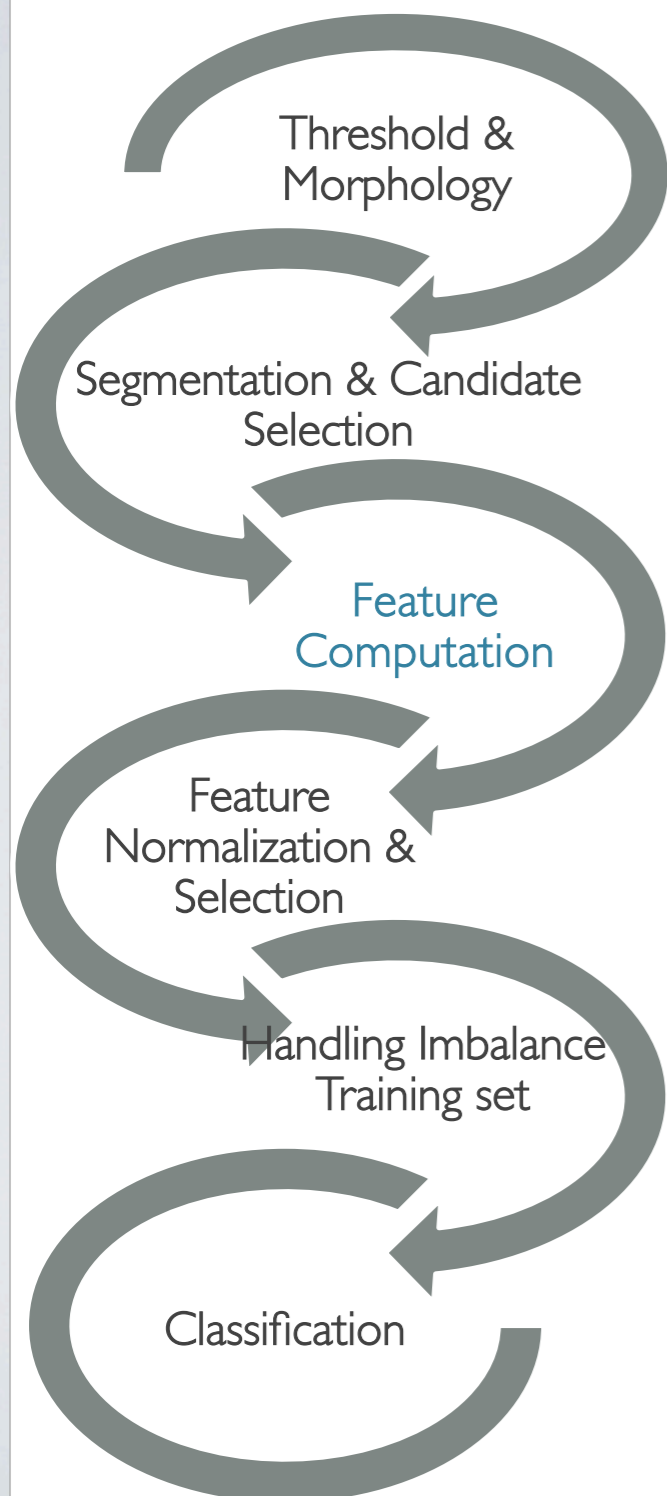
Segmentation & Candidate Selection



Selected Candidates  
(TP=Green, FP=Yellow, FN=Blue)

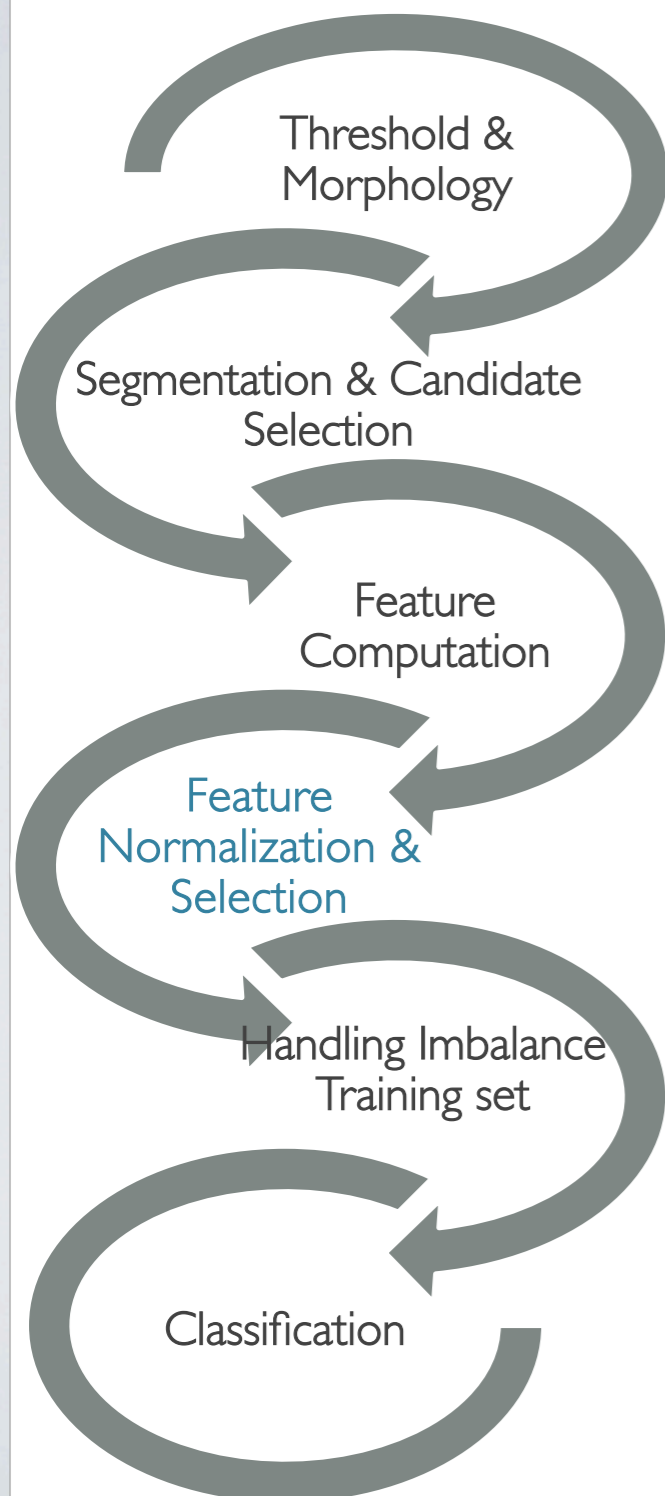


## ITM<sup>2</sup>C Framework Step 3



- **Morphology Features** (5 Features)
  - Area, Perimeter, Roundness, Elongation, Equivalent spherical perimeter
- **Intensity Features** (5 Features)
  - Mean, Median, Standard Deviation, Skewness, Kurtosis
- **Texture Features** (18 Features)
  - Co-occurrence Features (8 Features)
    - Correction, cluster shade, cluster prominence, energy, entropy, Hara-correlation, inertia, difference moment
  - Run-Length Features (10 Features)
    - SRE, LRE, GLN, RLN, LGRE, HGRE, SRLGE, SRHGE, LRLGE, LRHGE
- Compute intensity and texture features for each color channel (total eight channels)
- **Total Features = 5 + 8 ( 5 \* 18 ) = 189**

## ITM<sup>2</sup>C Framework Step 4



- **Feature Normalization**

$$f' = \frac{f - f_{\min}}{f_{\max} - f_{\min}}$$

where  $f'$  is normalized feature,  $f$  is actual feature

- **Feature Selection using Consistency subset evaluation method**

$$Consistency_s = 1 - \frac{\sum_{j=0}^{|s|} |D_j| - |M_j|}{N}$$

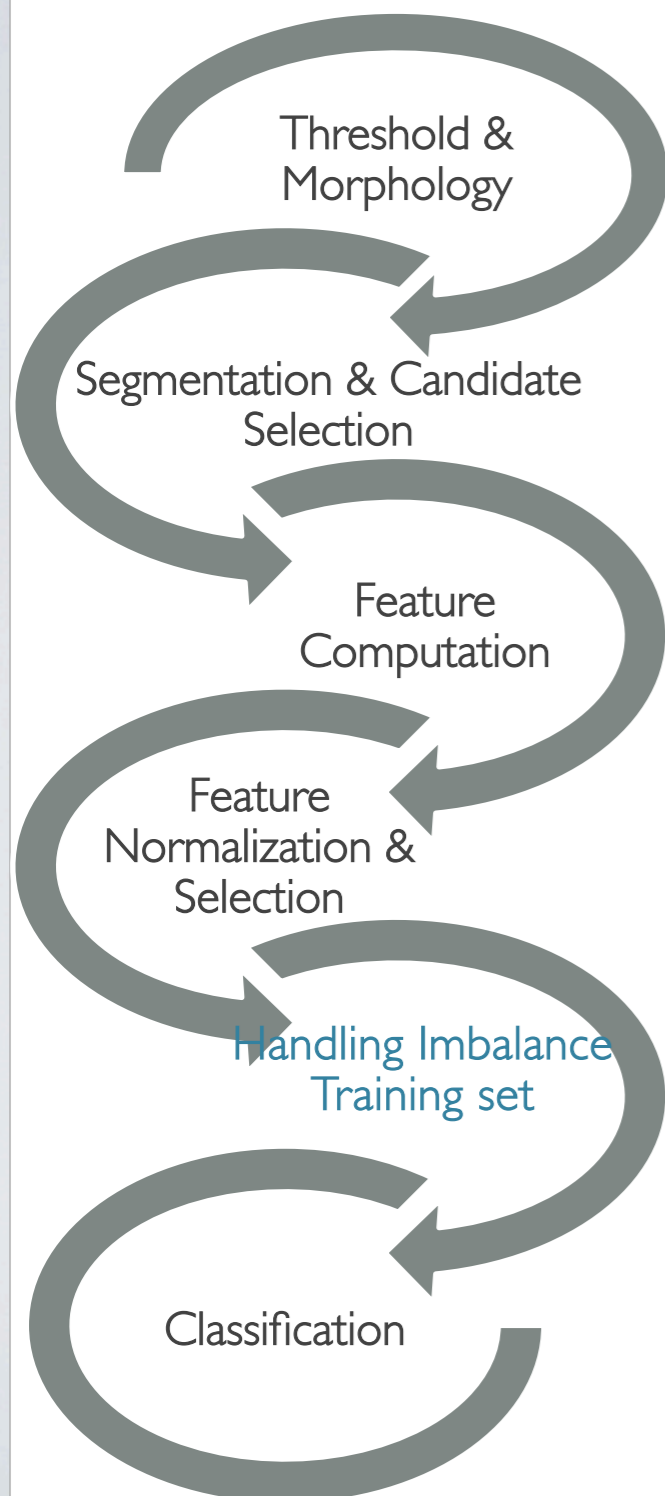
where  $s$  is a feature subset,  $j$  is the number of distinct combination of features for  $s$ ,  $|D_j|$  is the number of occurrences of the  $j^{\text{th}}$  feature combination,  $|M_j|$  is the cardinality of the majority class for the  $j^{\text{th}}$  feature combination and  $N$  is the total number of instances

- Use these subsets in conjunction with a hill climbing search method, augmented with backtracking

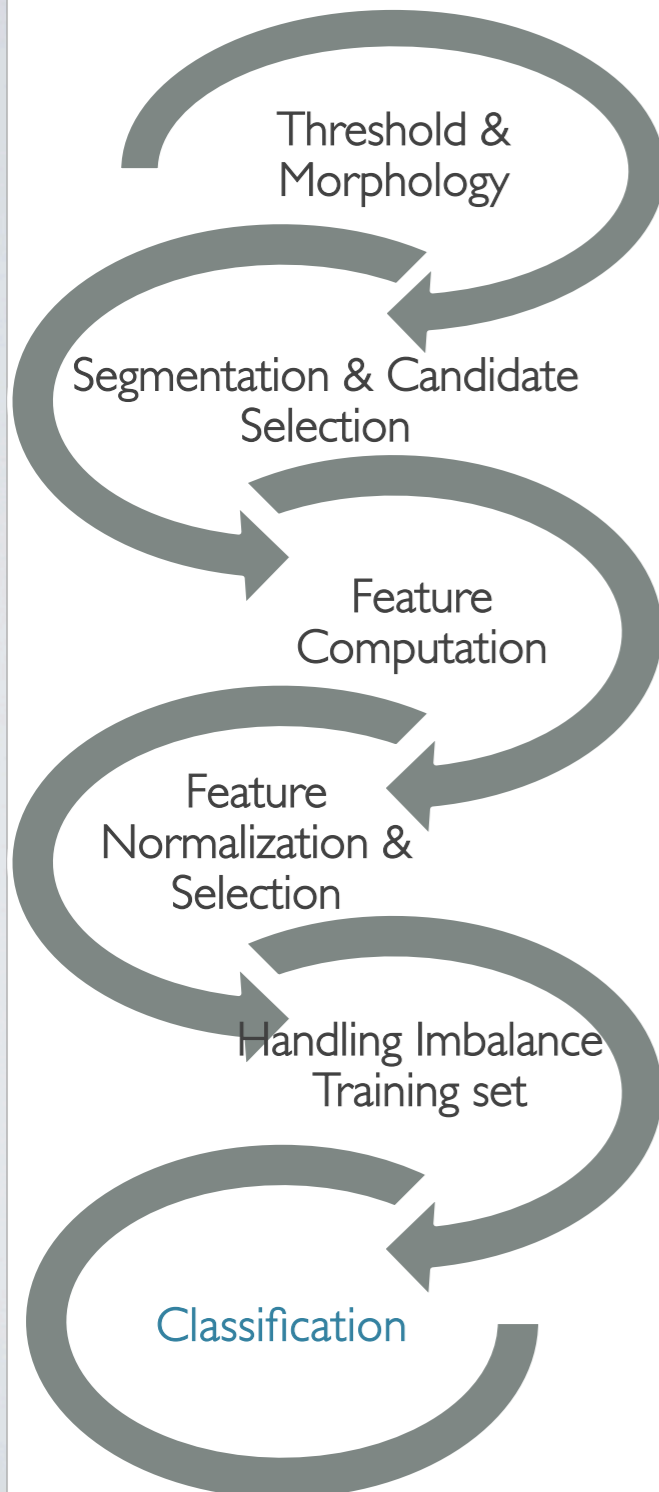
## ITM<sup>2</sup>C Framework Step 5

### Handling Imbalanced Training Set

- High degree of imbalance in training set (mitosis vs non-mitosis)
- Down-sampling of non-mitosis
- Over-sampling of mitosis using **Synthetic Minority Over-sampling Technique (SMOTE)**
  - 2 neighbors are selected from 5-nearest neighbors
  - New instance is generated in the direction of selected 2 neighbors



## ITM<sup>2</sup>C Framework Step 6



### 1. Decision Tree (DT)

$$P(C=c, X=x) = e^{F_c(x)} / \sum_{i=1}^{|C|} e^{F_i(x)}$$

$C$  = a label set,  $X$  = an instance set,  $F_c(x)$  = Functions of input variables

### 2. Multilayer Perceptron (MLP)

$$y(j) = 1 / (1 + e^{-S(j)})$$

$$S(j) = \sum_{k=1}^K y(k) w(k, j)$$

$$w(k, j)^{t+1} = w(k, j)^t + \alpha \varepsilon(j) y(j) + \beta (w(k, j)^t - w(k, j)^{t-1}),$$

$y(j)$  = output of node  $j$

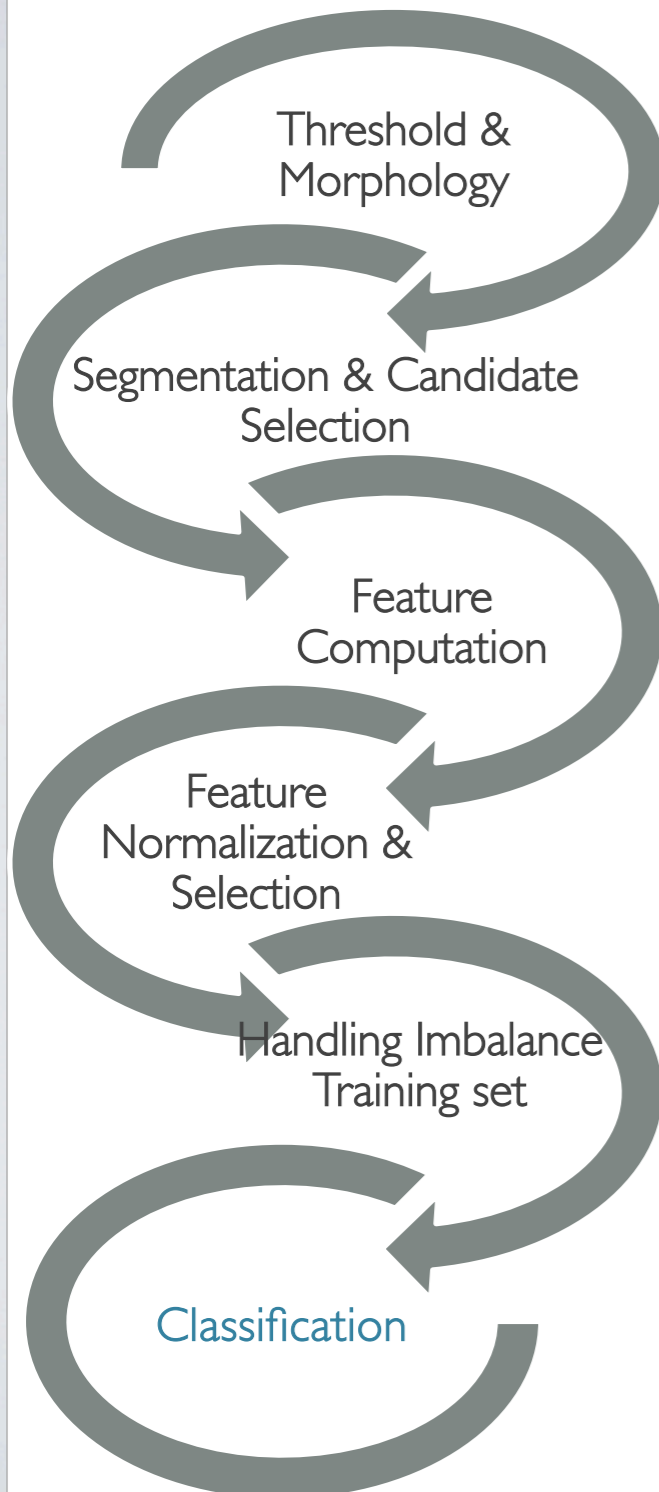
$S(j)$  = sum of all inputs weighted inputs of previous layer to node  $j$

$y(k) w(k, j)$  = weighted output of the previous node  $k$  to input node  $j$ ,

$K$  = number of inputs to node  $j$ ,

$w(k, j)$  = connections weights between previous node  $k$  and current node  $j$ ,

## ITM<sup>2</sup>C Framework Step 6



### 3. Linear Support Vector Machine (LSVM)

$$\min_{w, c} \frac{1}{2} \|w\|^2 + \alpha \sum_{k=1}^K \xi_k \varepsilon(w, x_k, c_k),$$

$$\varepsilon(w, x_k, c_k) = (\max(0, 1 - c_k w^T x_k))^2$$

$C$  = a label set,  $X$  = an instance set,  $\alpha$  = penalty parameter

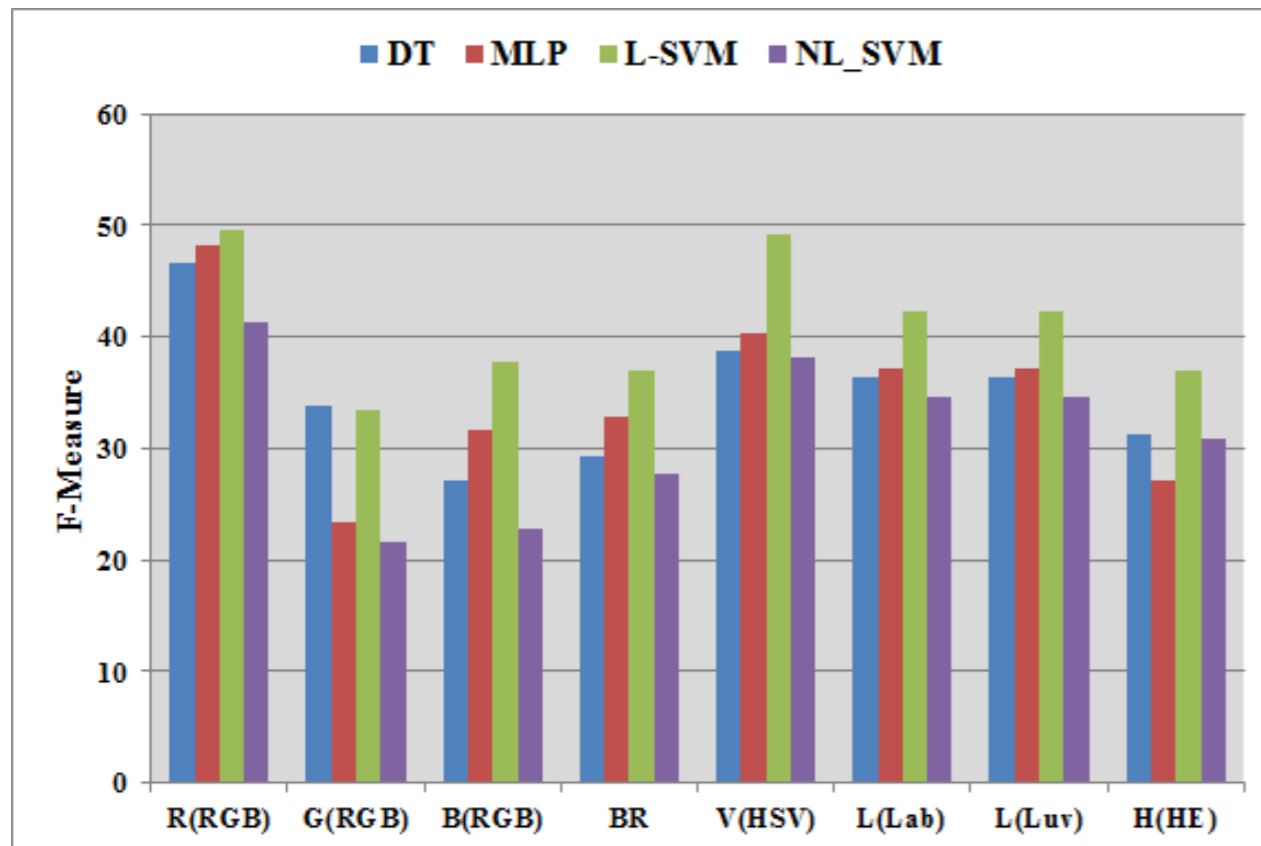
### 4. Non-Linear Support Vector Machine (NLSVM)

$$\min_{z, b, \xi} \frac{1}{2} \|z\|^2 + \alpha \sum_{k=1}^K \xi_k$$

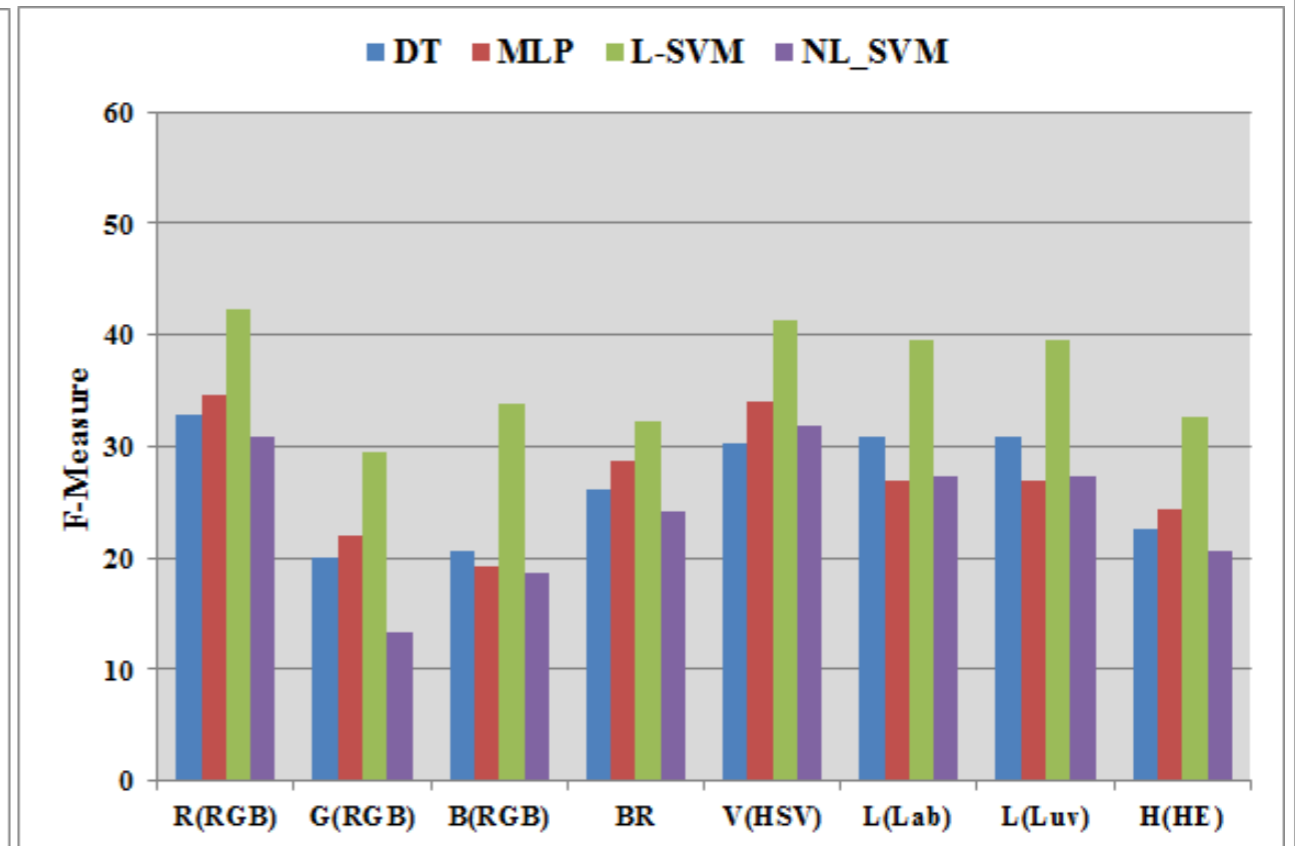
$z$  = normal to the hyper planes

$b$  = bias which describes the distance of the decision hyper plane from origin

## ITM<sup>2</sup>C Classification Results using Single Channel Features



On Aperio Dataset



On Hamamatsu Dataset

## ITM<sup>2</sup>C Result using All vs Selected Features on Evaluation set

		Aperio Dataset			Hamamatsu Dataset		
Features	Classifiers	TPR	PPV	FM	TPR	PPV	FM
All Features	DT	65%	71%	67.71%	60%	62%	60.91%
	MLP	68%	69%	68.34%	60%	61%	60.61%
	LSVM	72%	66%	68.57%	61%	62%	61.31%
	NLSVM	58%	83%	68.24%	53%	73%	61.27%
Selected Features	DT	67%	73%	69.79%	61%	64%	62.56%
	MLP	66%	74%	69.84%	60%	66%	62.83%
	LSVM	74%	71%	<u>72.55%</u>	63%	66%	<u>64.62%</u>
	NLSVM	59%	84%	69.41%	55%	74%	63.22%

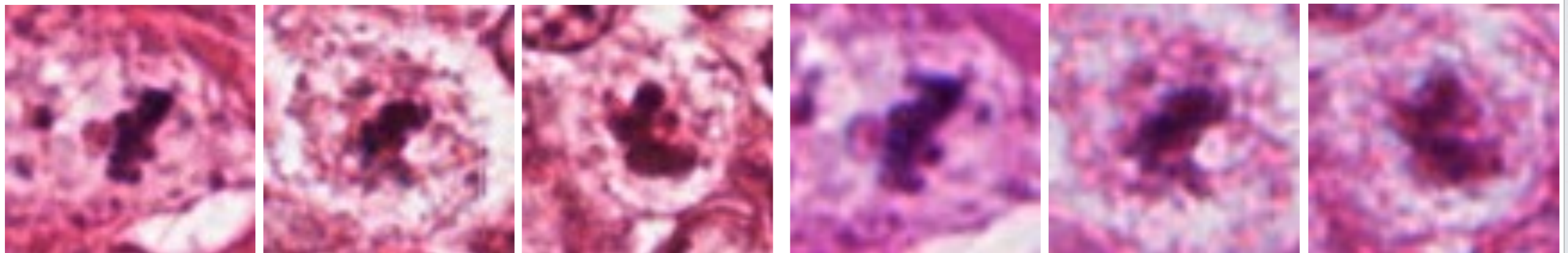
TPR = True Positive Rate

PPV = Predictive Positive Value

FM = F-Measure

## Different Patch Sizes for Feature Computation

Patch Size in pixels	Aperio Dataset ( $\mu\text{m}$ )	Hamamatsu Dataset ( $\mu\text{m}$ )
Patch $80 \times 80$	$19.648 \times 19.648$	$18.184 \times 18.202$
Patch $70 \times 70$	$17.192 \times 17.192$	$15.911 \times 15.927$
Patch $60 \times 60$	$14.736 \times 14.736$	$13.638 \times 13.652$
Patch $50 \times 50$	$12.28 \times 12.28$	$11.365 \times 11.377$
Patch $40 \times 40$	$9.824 \times 9.824$	$9.092 \times 9.101$



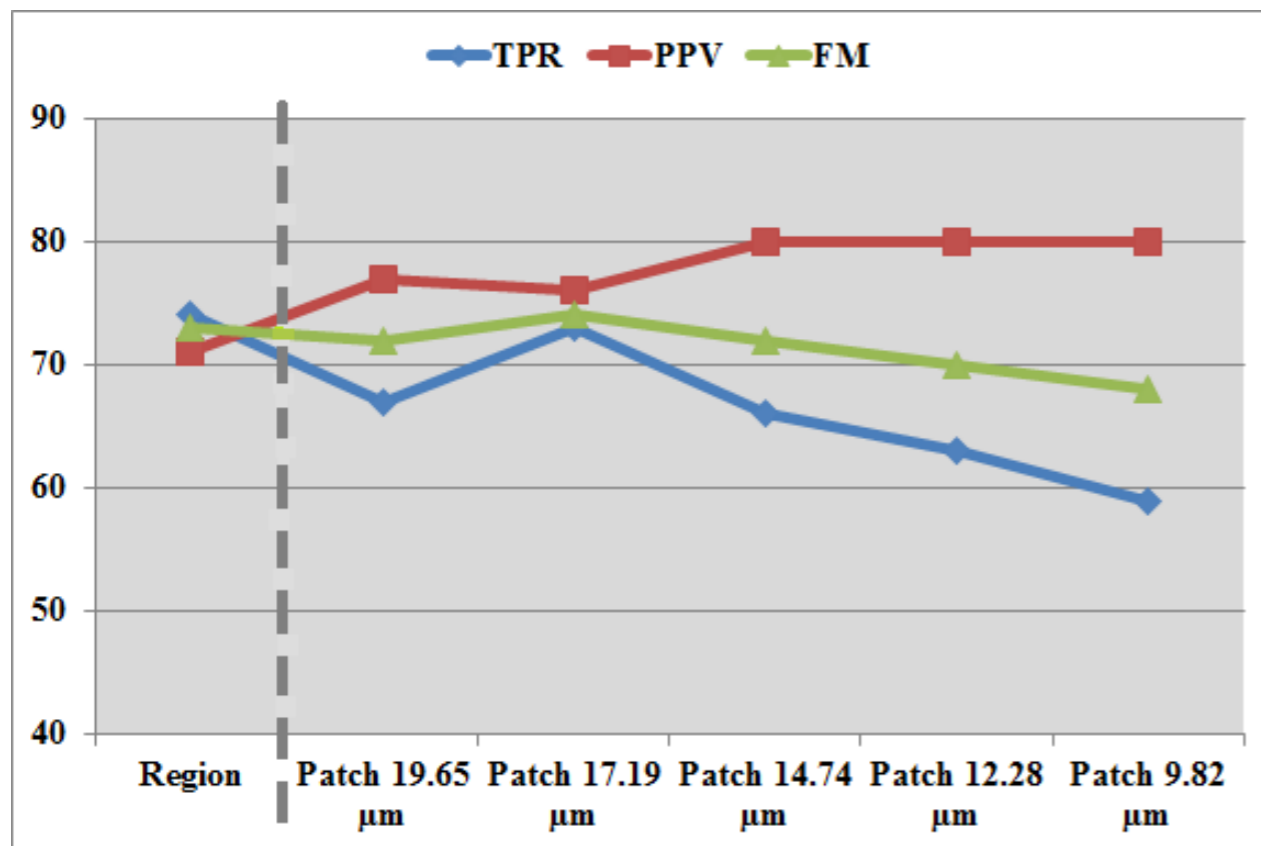
Mitosis Patch from Aperio Dataset

Mitosis Patch from Hamamatsu Dataset

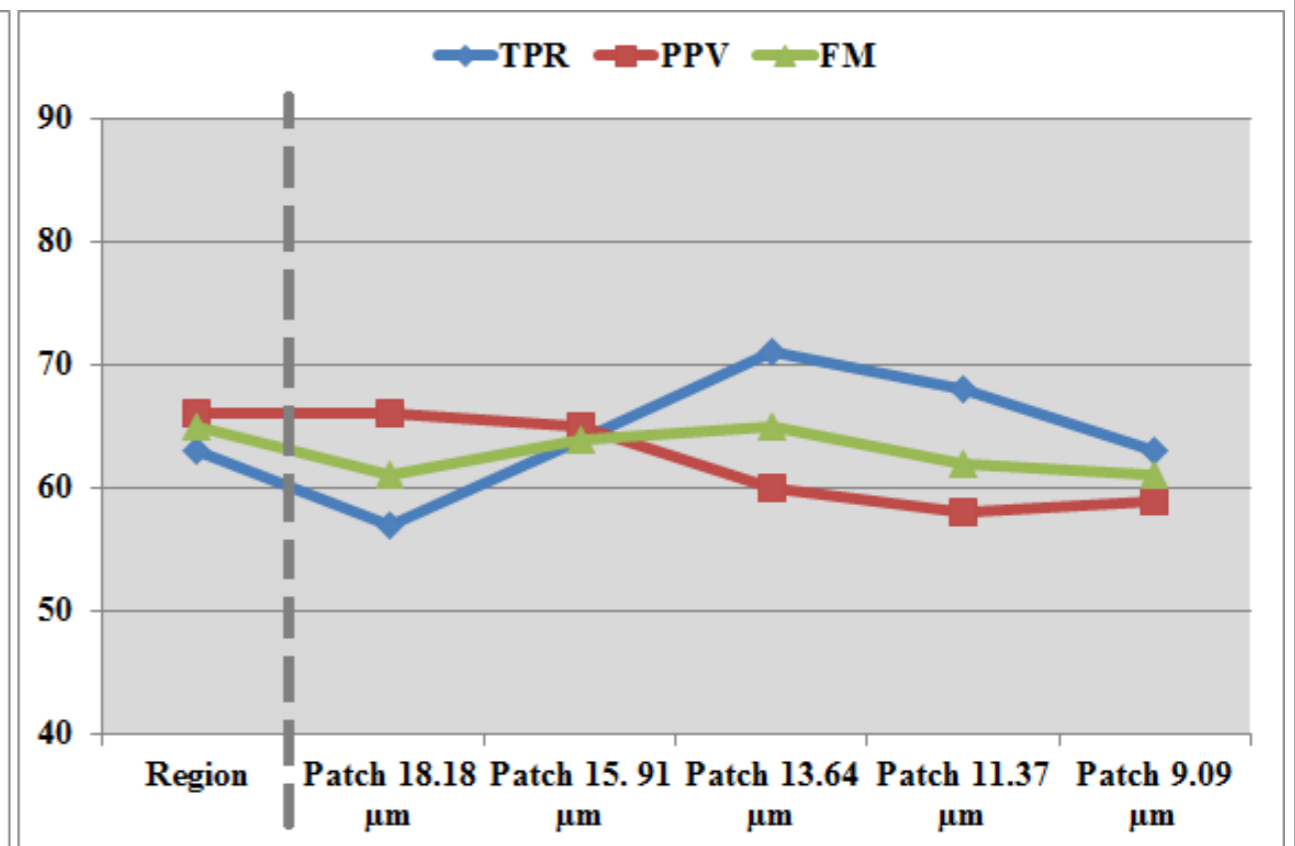


## Region vs Patch Features based Classification using LSVM Classifier

- Both scanners have different information on same patch size.

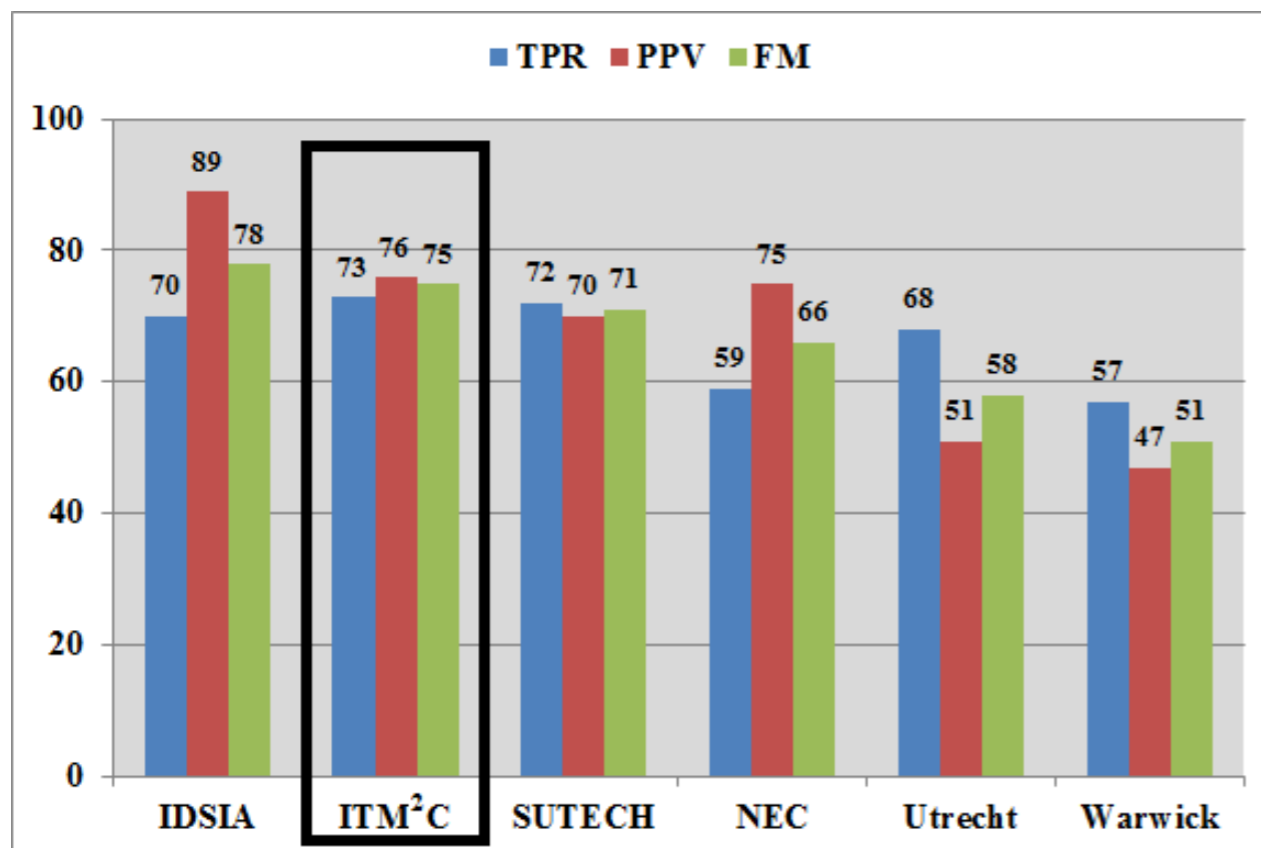


On Aperio Dataset

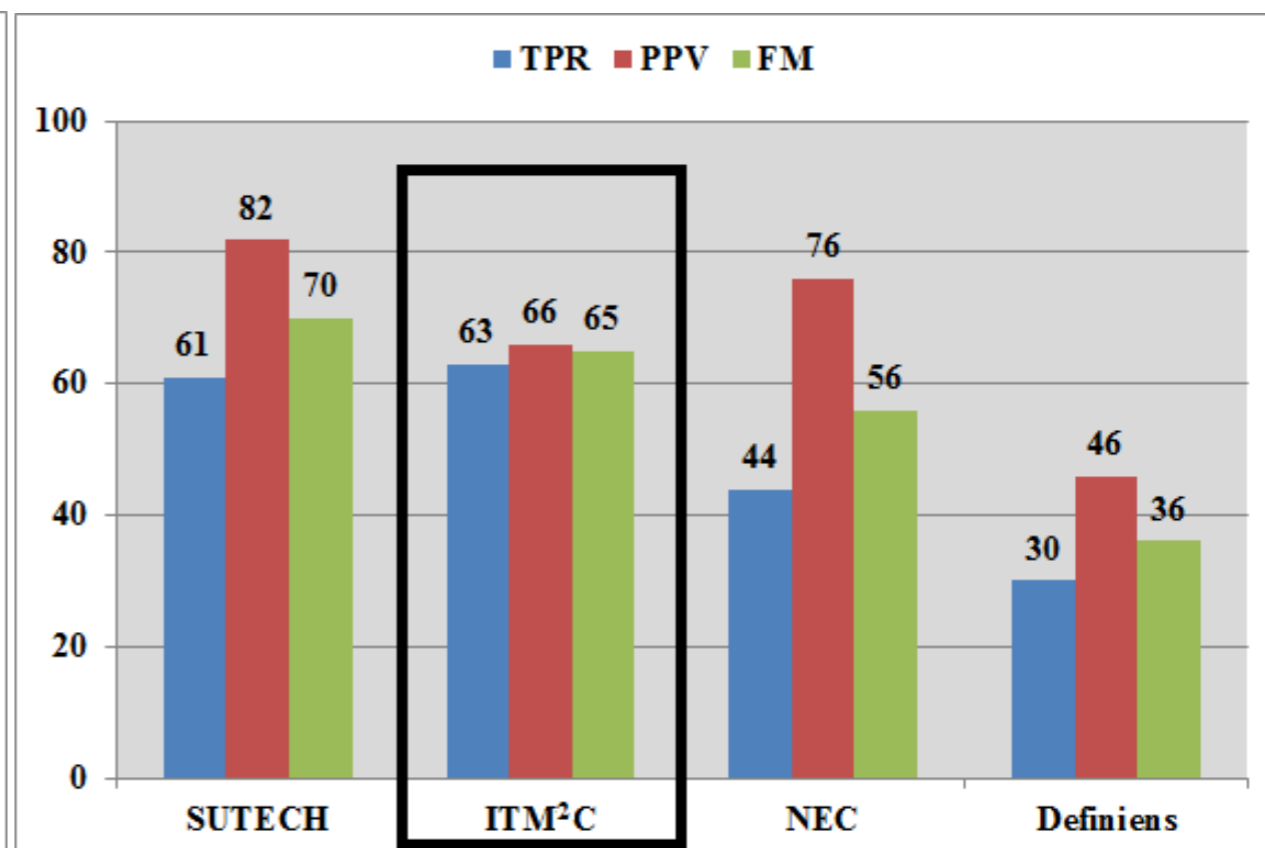


On Hamamatsu Dataset

## Comparison of Results with ICPR MITOS Contest 2012

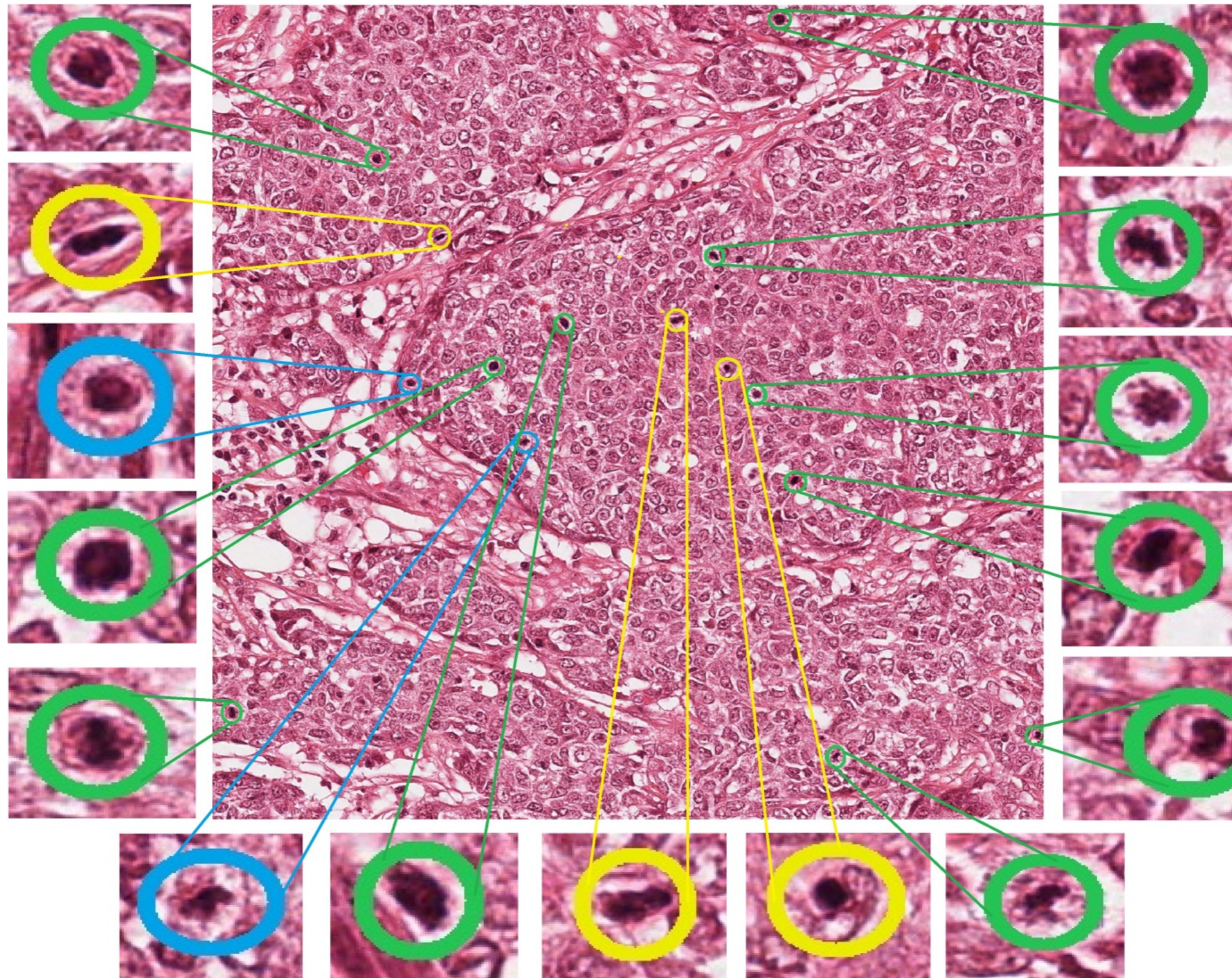


Comparison on Aperio Dataset



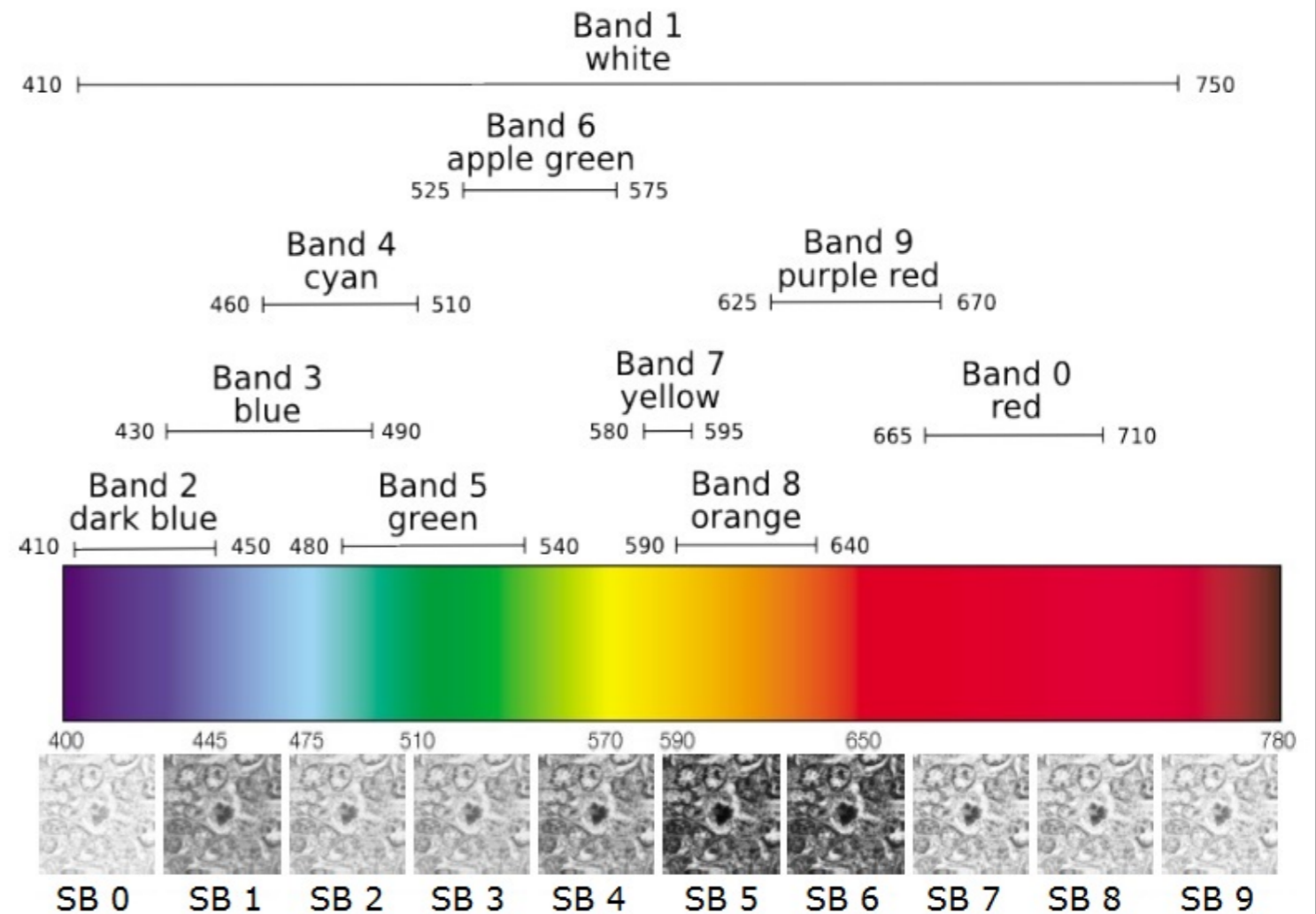
Comparison on Hamamatsu Dataset

Candidate Classification on Aperio Dataset (TP=Green, FP=Yellow, FN-Blue)



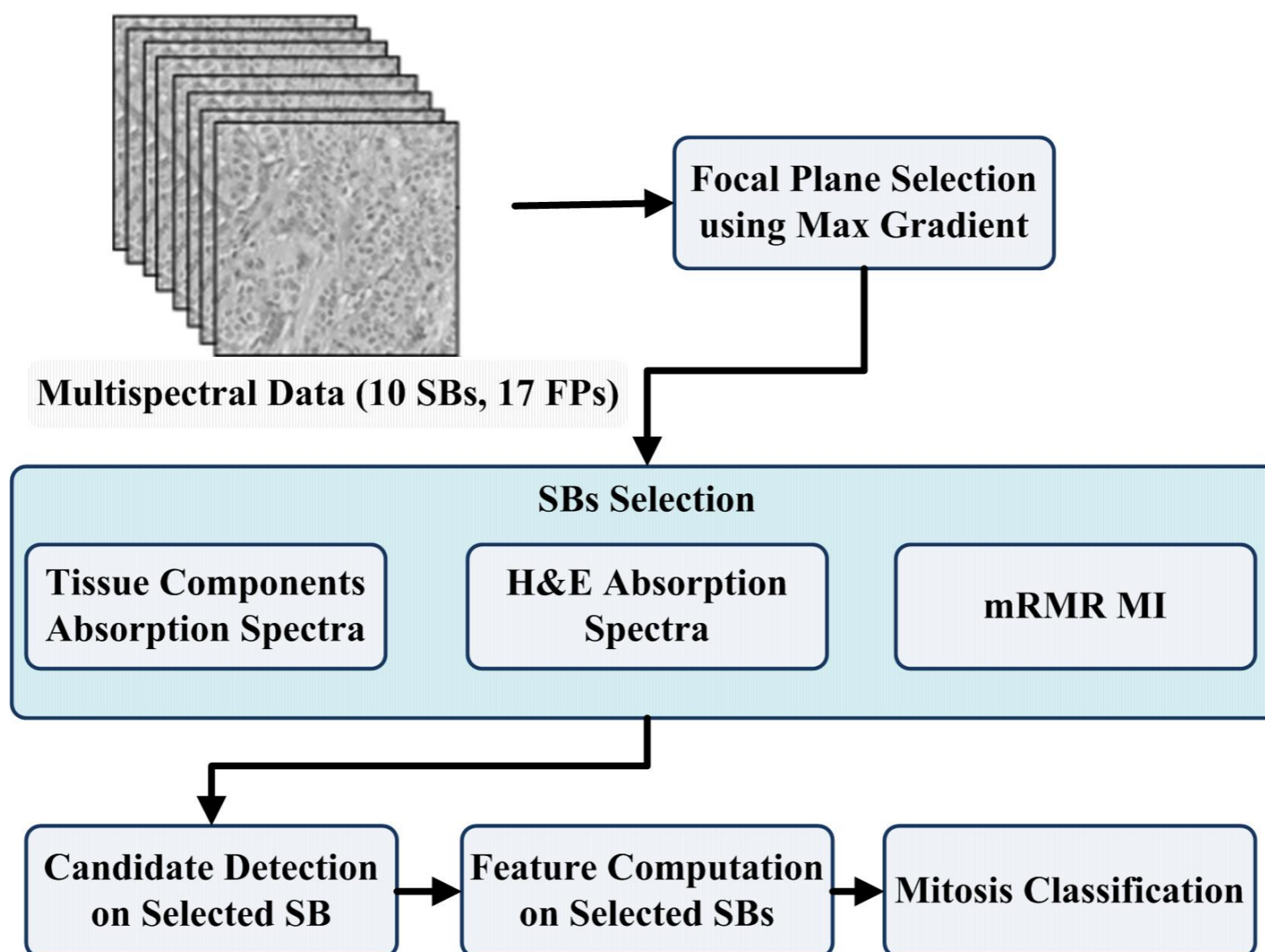
## Multispectral Dataset

- 10 Spectral bands
- 17 Focal Planes (layer Z-stack)
- 4 images per HPF
- 1 HPF =  $4 * 17 * 10 = 680$  images
- Image resolution =  $251.6 \times 251.6 \mu m^2$   
an area of  $0.063 mm^2$
- Total 50 HPF (322 mitosis)
  - 35 Training set (244 mitosis)
  - 15 Evaluation set (98 mitosis)



Spectral bands (SBs) of multispectral microscope and example of each SB

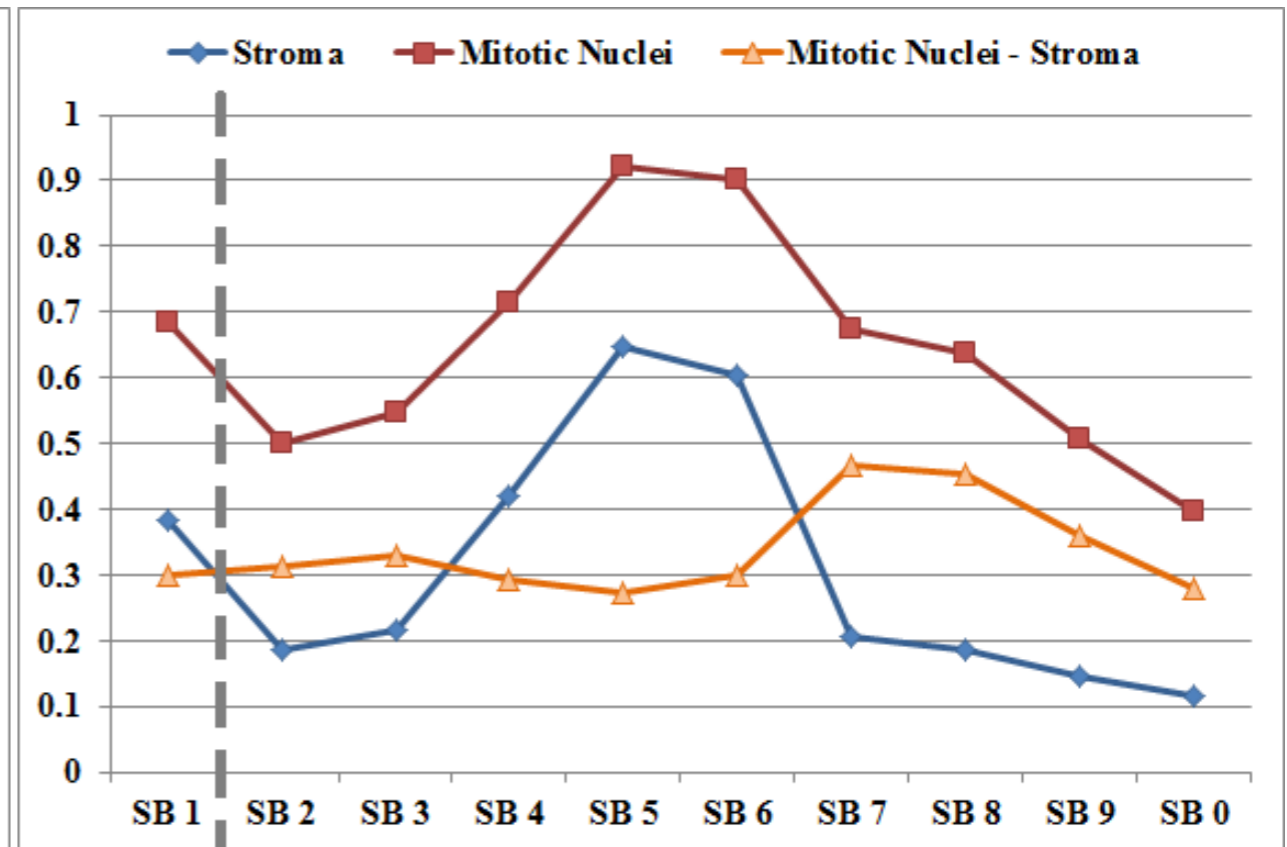
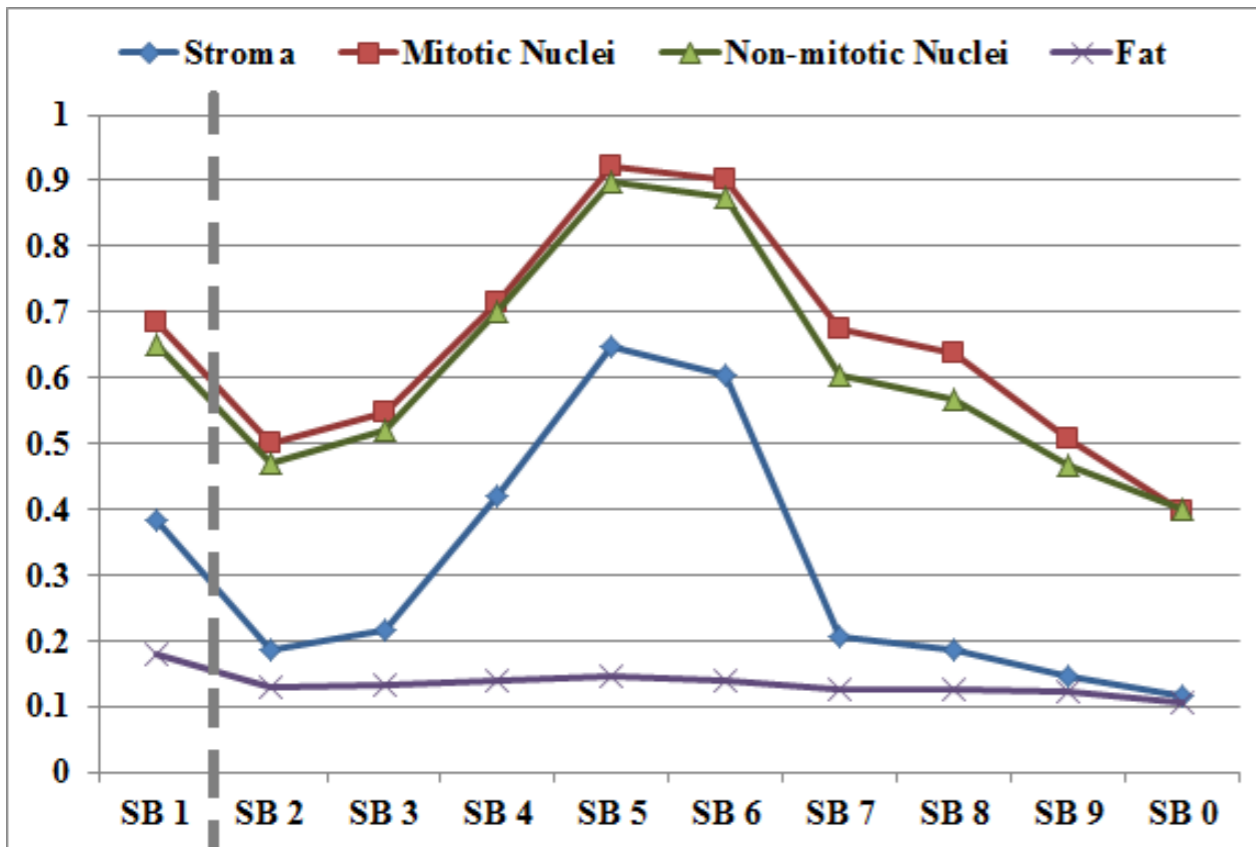
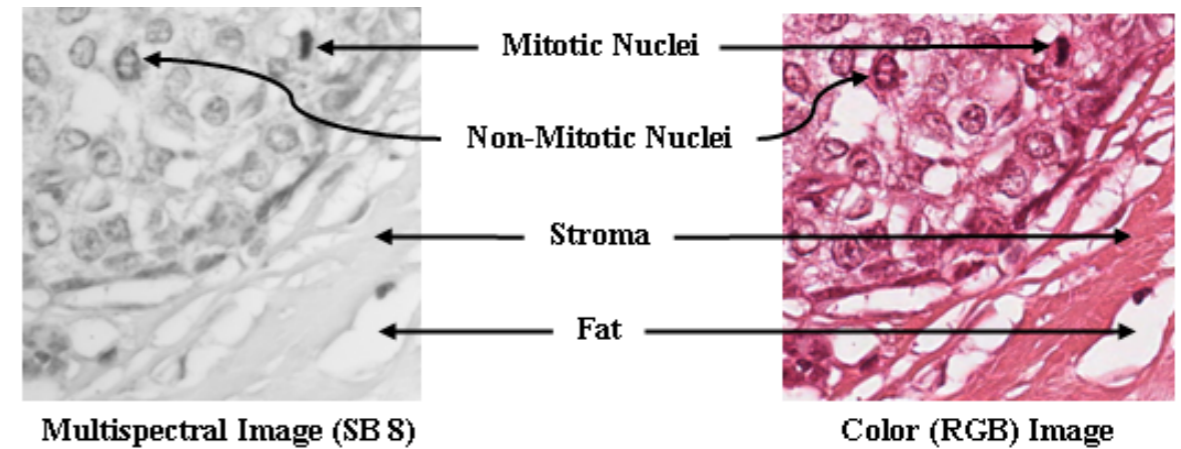
## Multispectral Intensity, Texture & Morphology-based Mitosis detection in Multispectral images (MITM<sup>3</sup>) Framework



MITM<sup>3</sup> Framework

## Spectral Band Selection

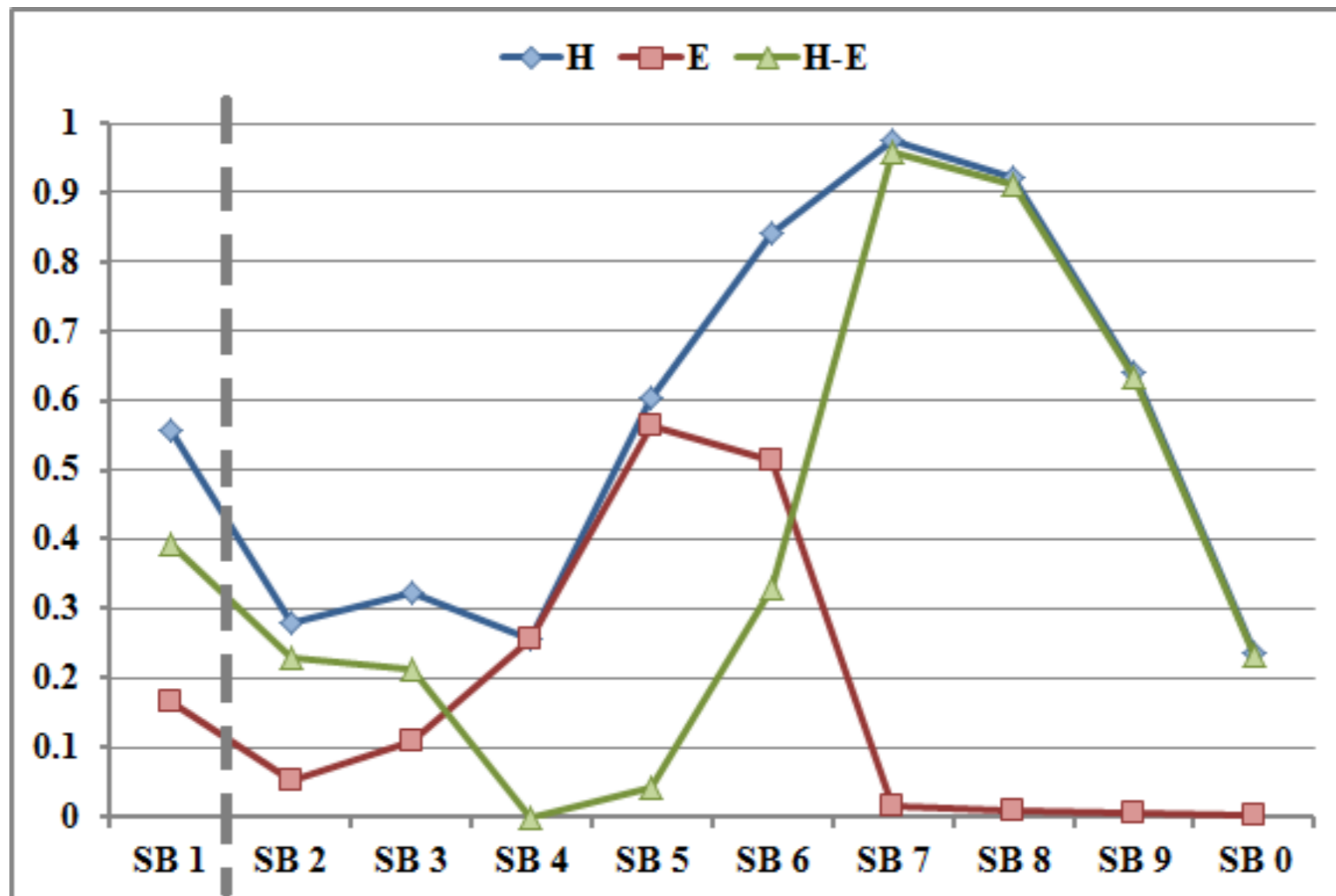
### Method I – Tissue Spectral Absorption



Normalized average gradient spectra of four tissue components

## Spectral Band Selection

### Method 2 – H & E Spectral Absorption



Normalized plot of the Hematoxylin (Blue line) and Eosin (Red line) dye absorption spectra and difference of Hematoxylin and Eosin (green line)

## Spectral Band Selection

### Method 3 – SBs Selection using Minimum redundancy and Maximum Relevance (mRMR)

- Relevance  $D = 1/|S| \sum_{s \downarrow i \in S} MI(s \downarrow i; c \downarrow j)$
- Redundancy  $R = 1/|S|^2 \sum_{s \downarrow i, s \downarrow j \in S} MI(s \downarrow i; s \downarrow j)$
- Mutual Information

$$MI(S; C) = - \sum_{s \downarrow i \in S} p(s \downarrow i) \log_2(p(s \downarrow i)) + \sum_{s \downarrow i \in S} \sum_{c \downarrow j \in C} p(s \downarrow i, c \downarrow j) \log_2(p(s \downarrow i, c \downarrow j))$$

- Incremental search method is used to find the  $n$  SBs from the set  $\{S \downarrow T - S \downarrow (n-1)\}$  by maximizing  $\max_{s \downarrow i \in S \downarrow T - S \downarrow (n-1)} [MI(s \downarrow i; c) - 1/n-1 \sum_{s \downarrow j \in S \downarrow (n-1)} MI(s \downarrow i; s \downarrow j)]$

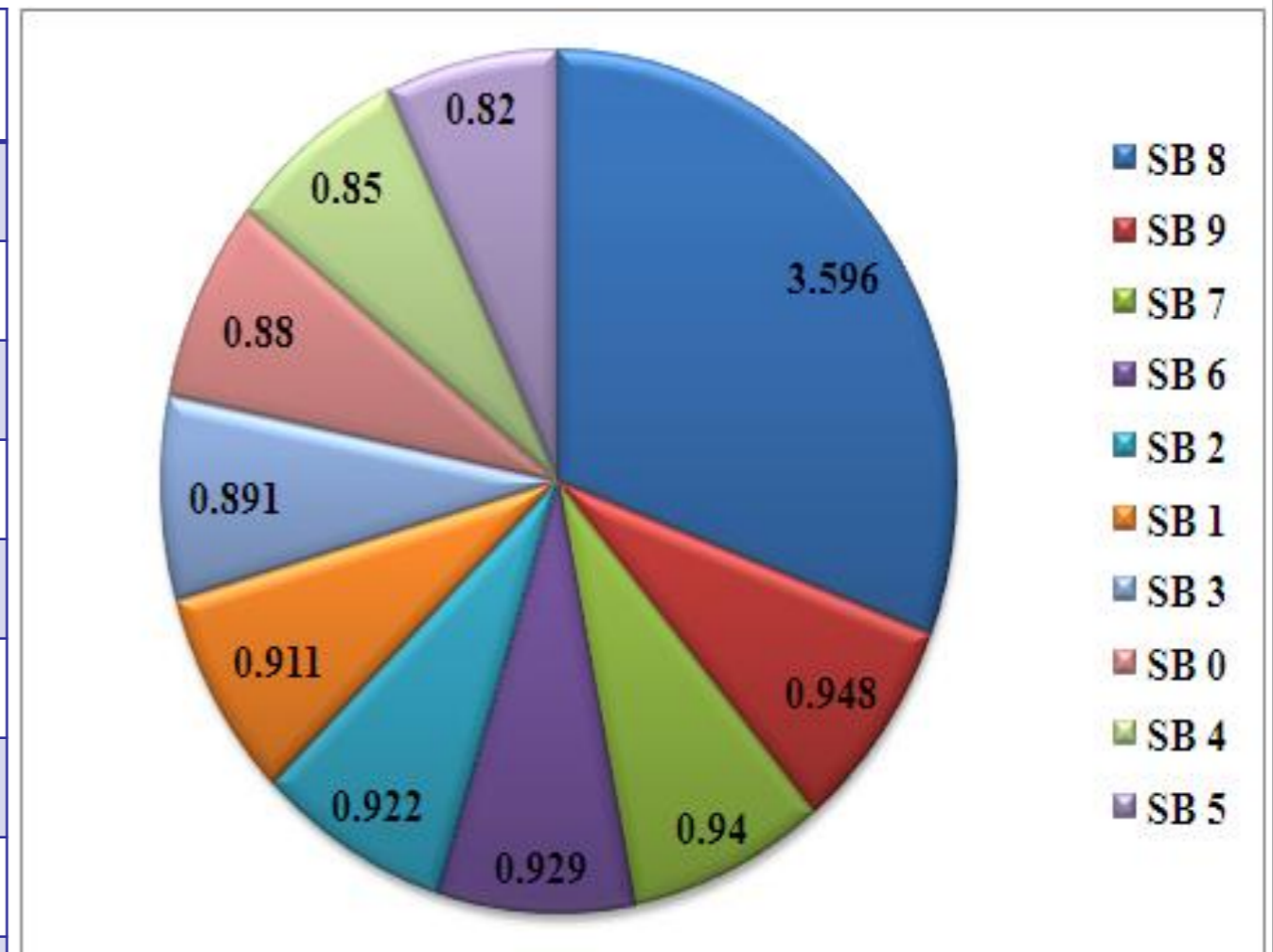
$S$  = SBs set,  $C$  = class label,  $p(s \downarrow i)$  = probability density function of SB  $s \downarrow i$ ,  $p(s \downarrow i, c \downarrow j)$  = conditional probability density function of SB  $s \downarrow i$  and class label  $c \downarrow j$



## Spectral Band Selection

Method 3 – SBs Selection using Minimum redundancy and Maximum Relevance (mRMR)

SBs	MI	Accumulated MI	Accumulated MI%
SB 8	3.60	3.60	33%
SB 9	3.59	0.95	42%
SB 7	3.58	0.94	51%
SB 6	3.18	0.93	60%
SB 2	3.16	0.92	69%
SB 1	3.11	0.91	78%
SB 3	3.05	0.89	86%
SB 0	2.99	0.88	91%
SB 4	2.94	0.85	95%
SB 5	2.85	0.82	100%



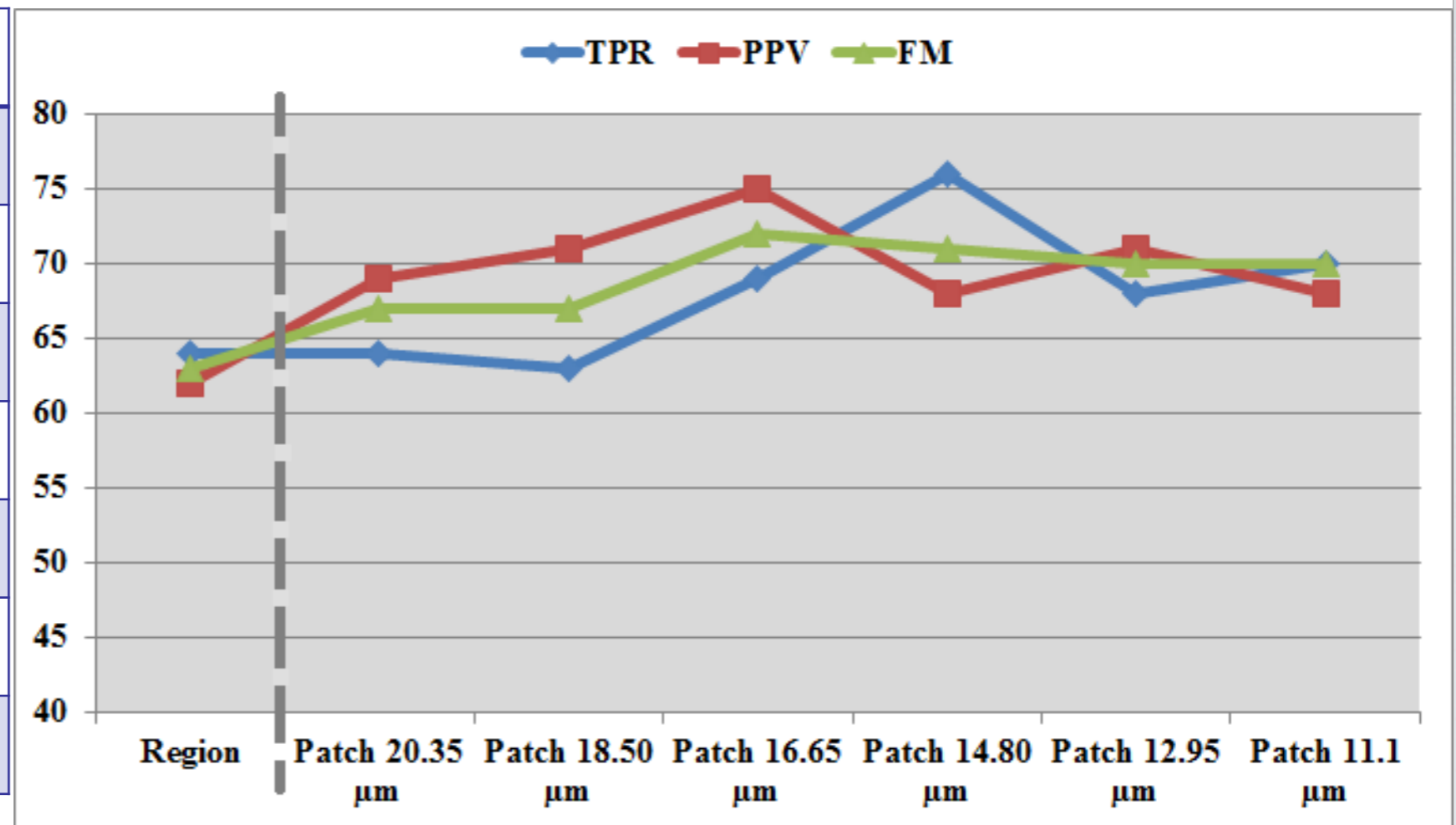
## 3 Rankings of Spectral Bands

Method 1 – Tissue Absorption Spectra		Method 2 – H&E Absorption Spectral		Method 3 – mRMR	
SBs	Mitosis-Cytoplasm	SBs	H-E	SBs	MI
SB 7	0.47	SB 7	0.96	SB 8	3.60
SB 8	0.45	SB 8	0.91	SB 9	3.59
SB 9	0.36	SB 9	0.64	SB 7	3.58
SB 3	0.33	SB 1	0.39	SB 6	3.18
SB 2	0.31	SB 6	0.33	SB 2	3.16
SB 6	0.30	SB 0	0.23	SB 1	3.11
SB 1	0.30	SB 2	0.23	SB 3	3.05
SB 4	0.29	SB 3	0.21	SB 0	2.99
SB 0	0.28	SB 5	0.04	SB 4	2.94
SB 5	0.27	SB 4	0	SB 5	2.85

- Selected SBs are 8,9,7,6,2,1,3 and 0.

## Region vs Patch Features based Classification Results

Different Patch Sizes			
Sizes in pixels		Sizes in $\mu\text{m}$	
110	110	20.35	20.35
100	100	18.50	18.50
90	90	16.65	16.65
80	80	14.82	14.80
70	70	12.95	12.95
60	60	11.1	11.1

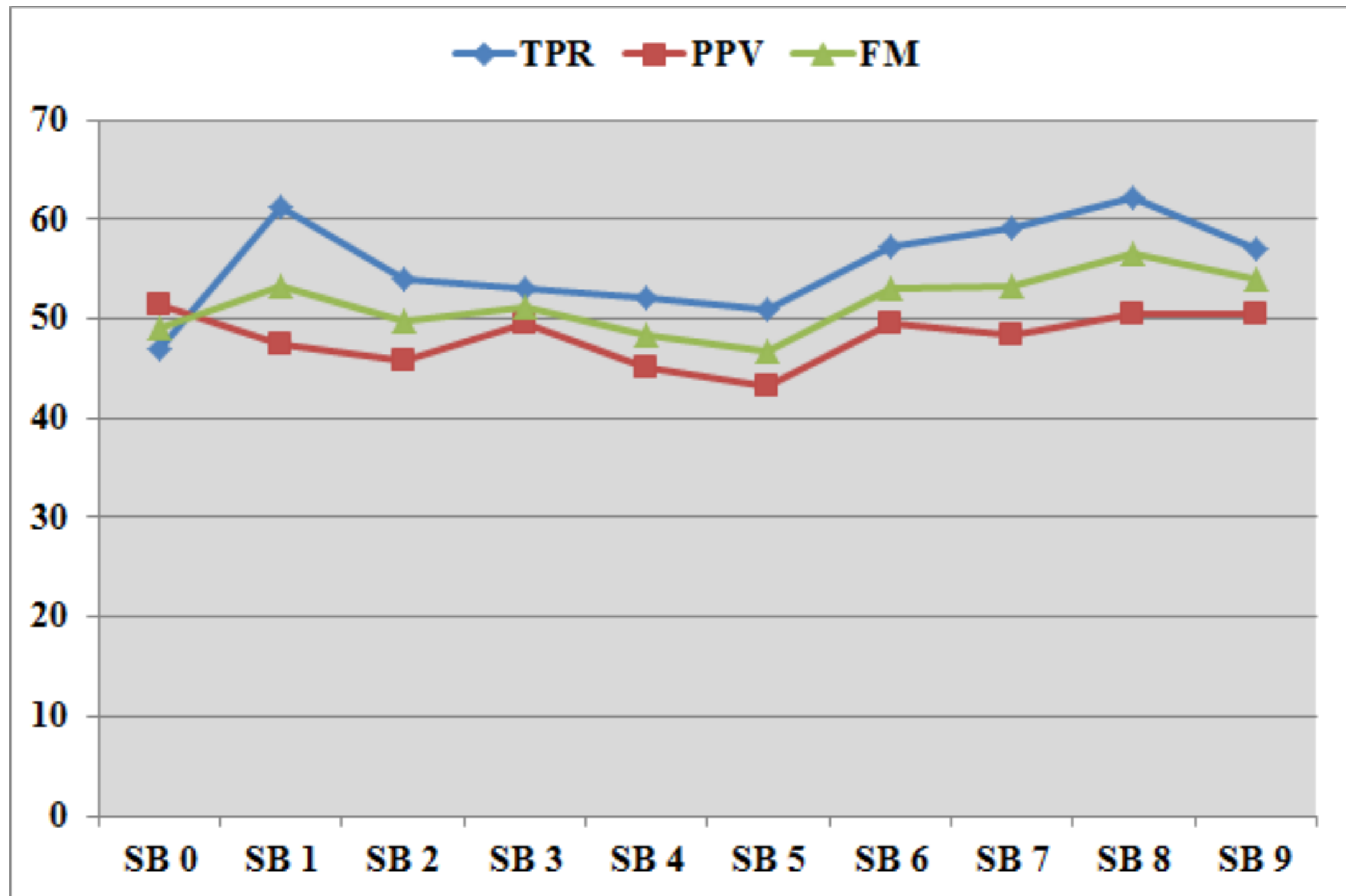


Region vs Patch features based classification results using LSVM classifiers

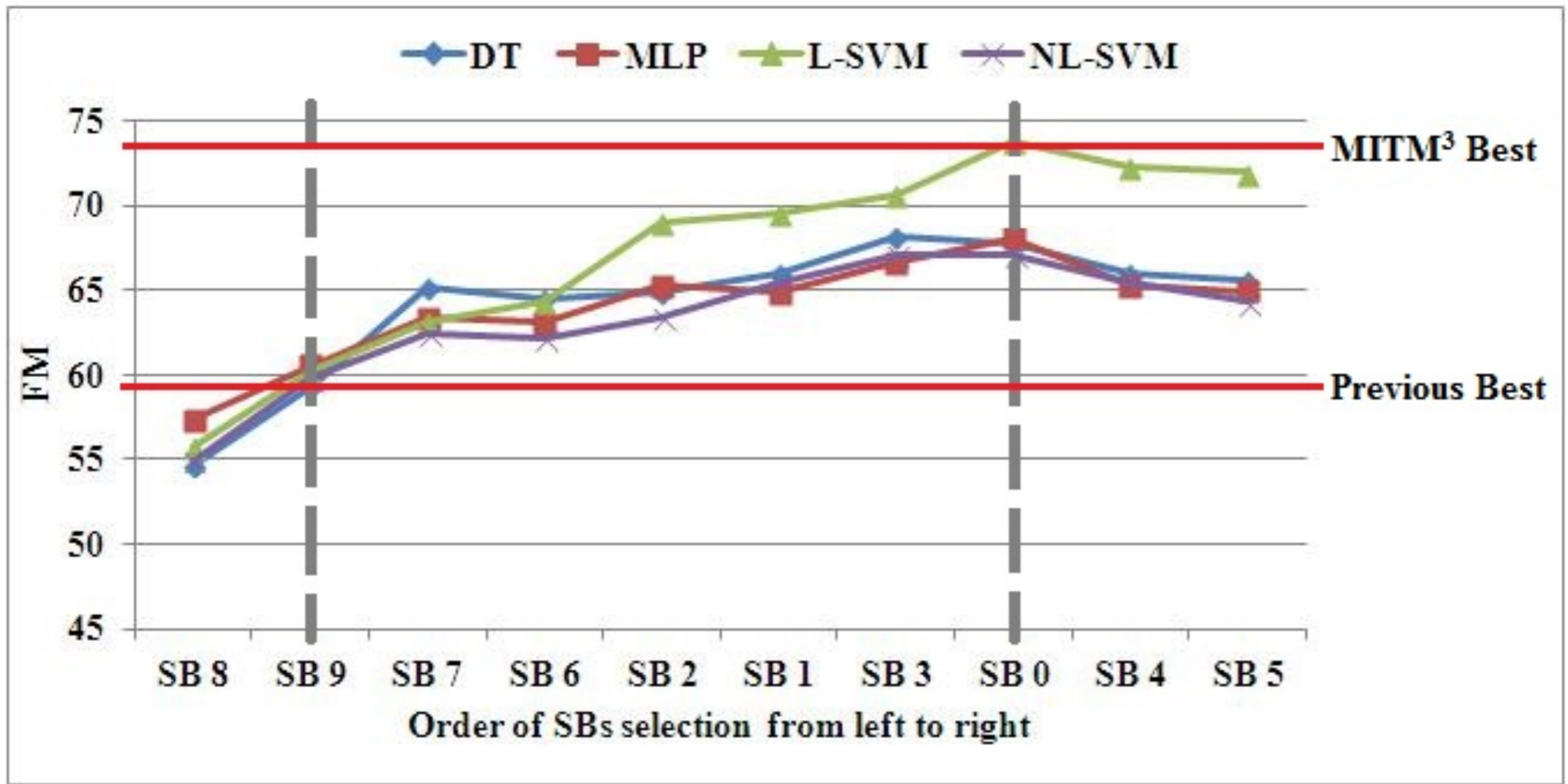
## MITM<sup>3</sup> Classification on Evaluation Set

		All SBs Features			Selected 8 SBs Features		
Features	Classifiers	TPR	PPV	FM	TPR	PPV	FM
Region Features	DT	67%	53%	59.1%	62%	62%	62.24%
	MLP	64%	56%	59.72%	62%	66%	63.10%
	LSVM	63%	60%	61.69%	64%	62%	63.32%
	NLSVM	54%	68%	60.23%	59%	69%	63.74%
Patch size 16.65 $\mu$ m Features	DT	61%	71%	65.57%	65%	70%	67.72%
	MLP	63%	67%	64.92%	66%	70%	68.06%
	LSVM	69%	75%	<u>71.96%</u>	74%	73%	<u>73.74%</u>
	NLSVM	55%	77%	64.29%	59%	77%	67.05

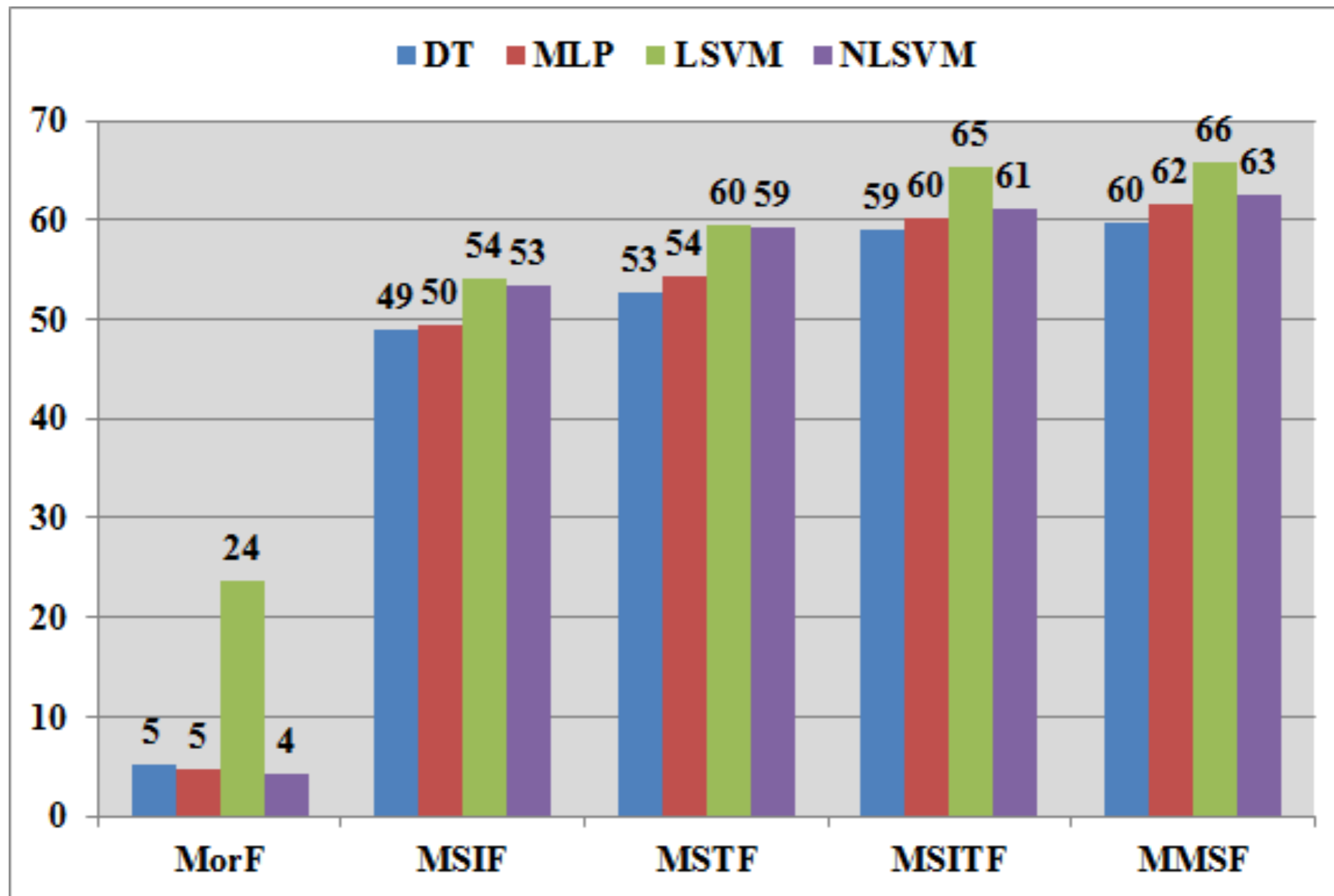
Plot of TPR, PPV and FM using Single SB Features with LSVM classifier



Plot of FM using Accumulated Features from the order of mRMR Selection



## Results on different subsets of Features using 5-Fold Cross Validation



MorF = Morphology Features

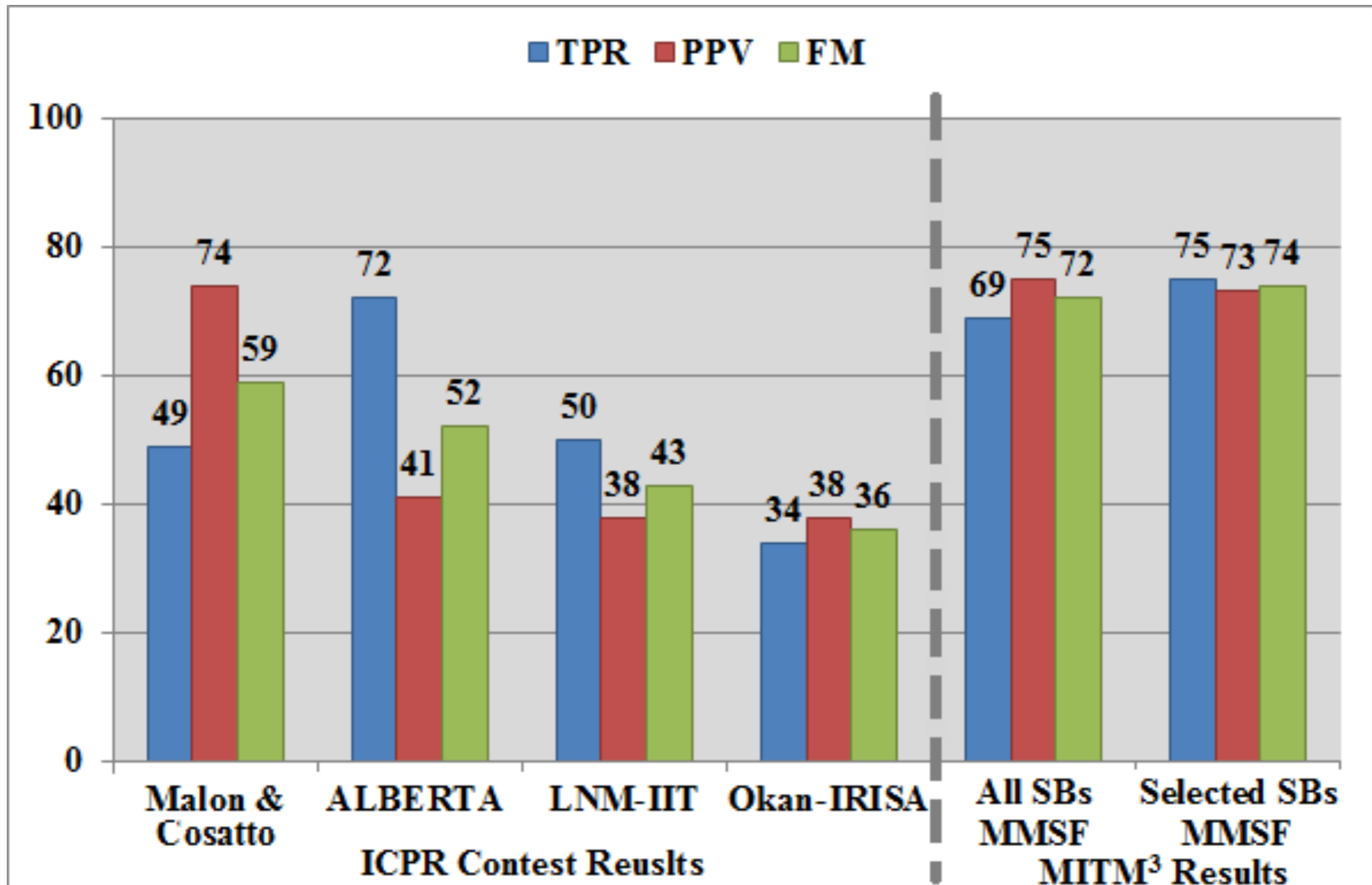
MSIF = Multi-Spectral Intensity Features

MSTF = Multi-Spectral Texture Features

MSITF = Multi-Spectral Intensity & Texture Features

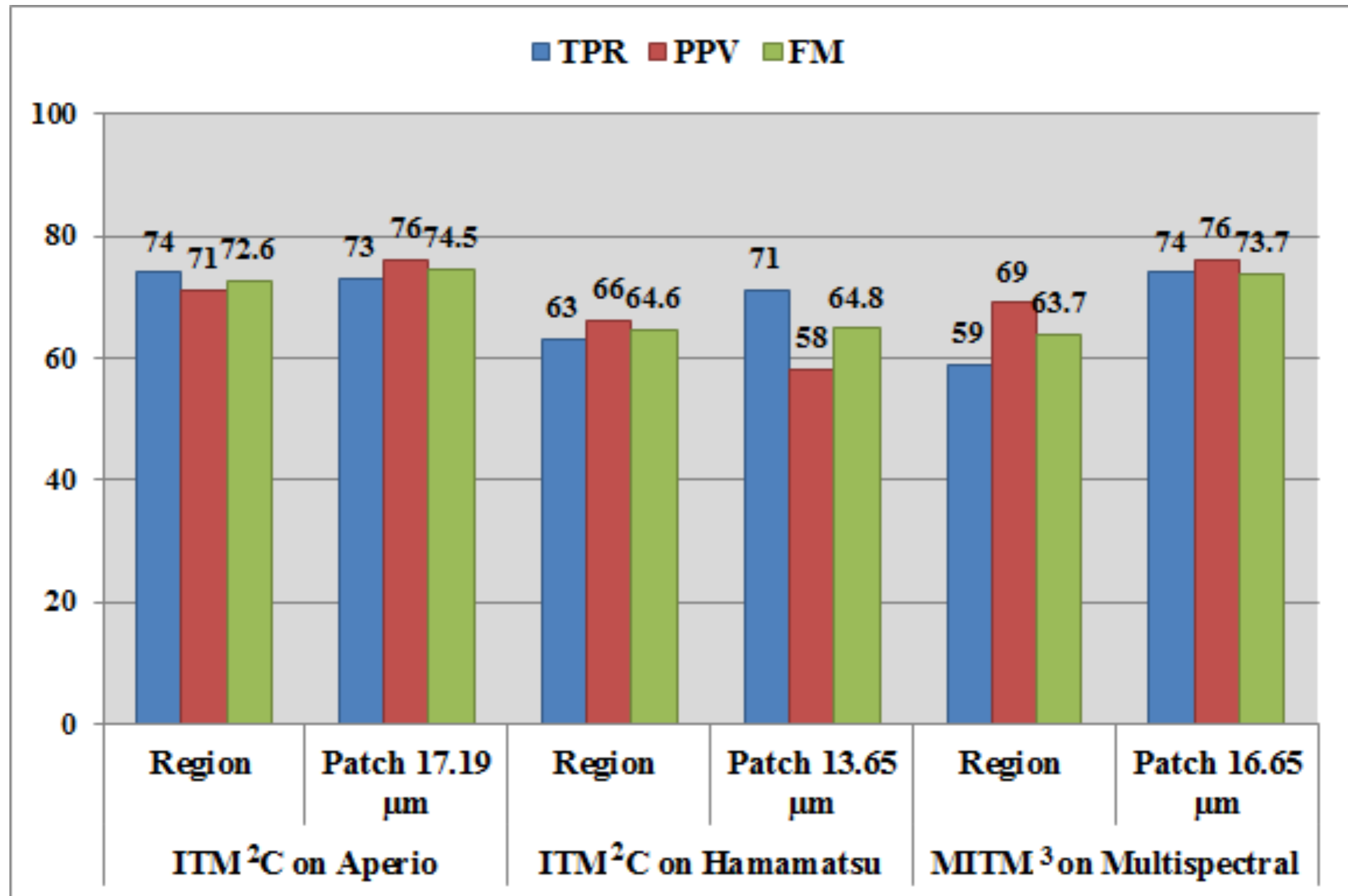
MMSF = Morphological & Multispectral Statistical Features

## Comparison of MITM<sup>3</sup> Framework with ICPR 2012 MITOS Contest

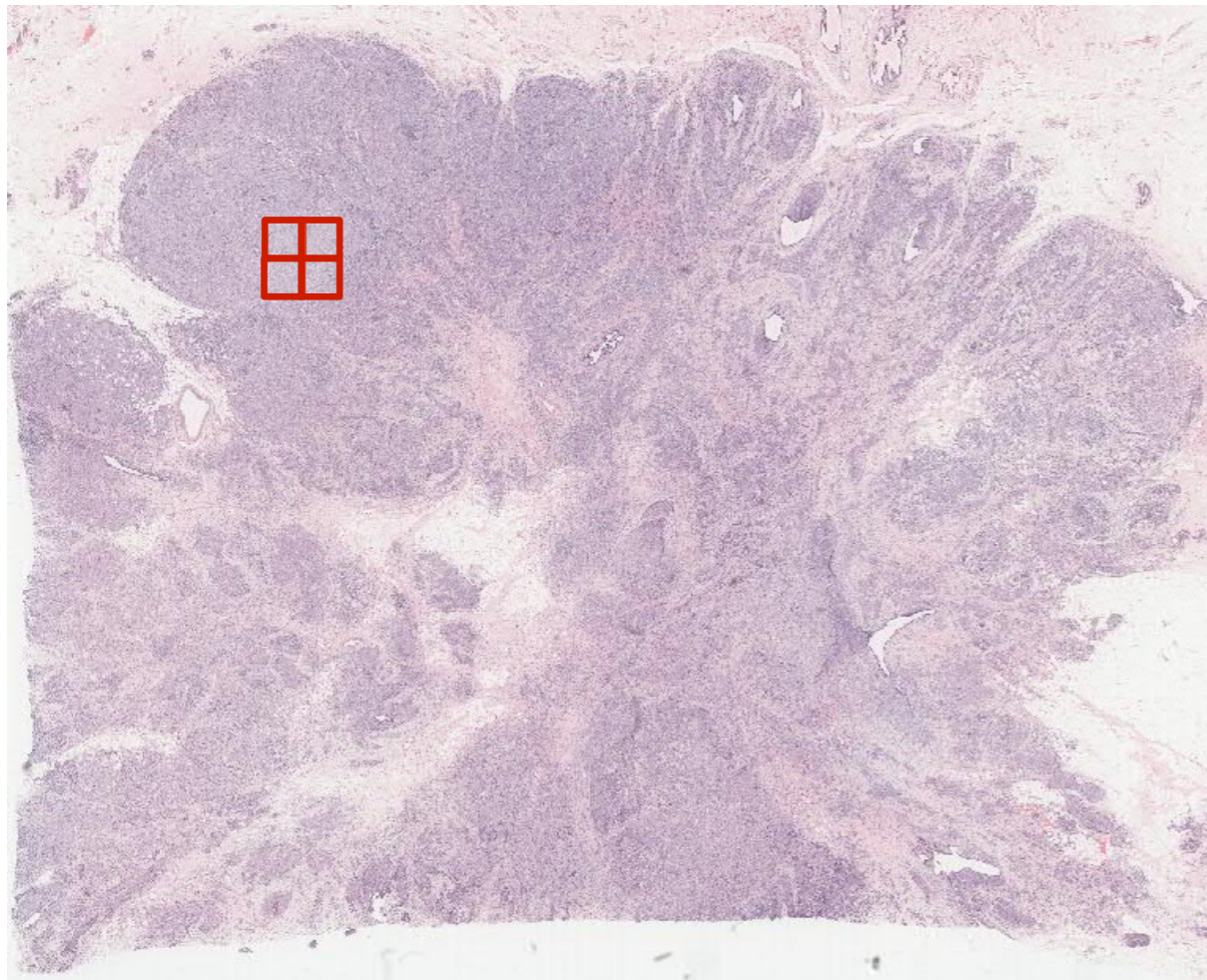




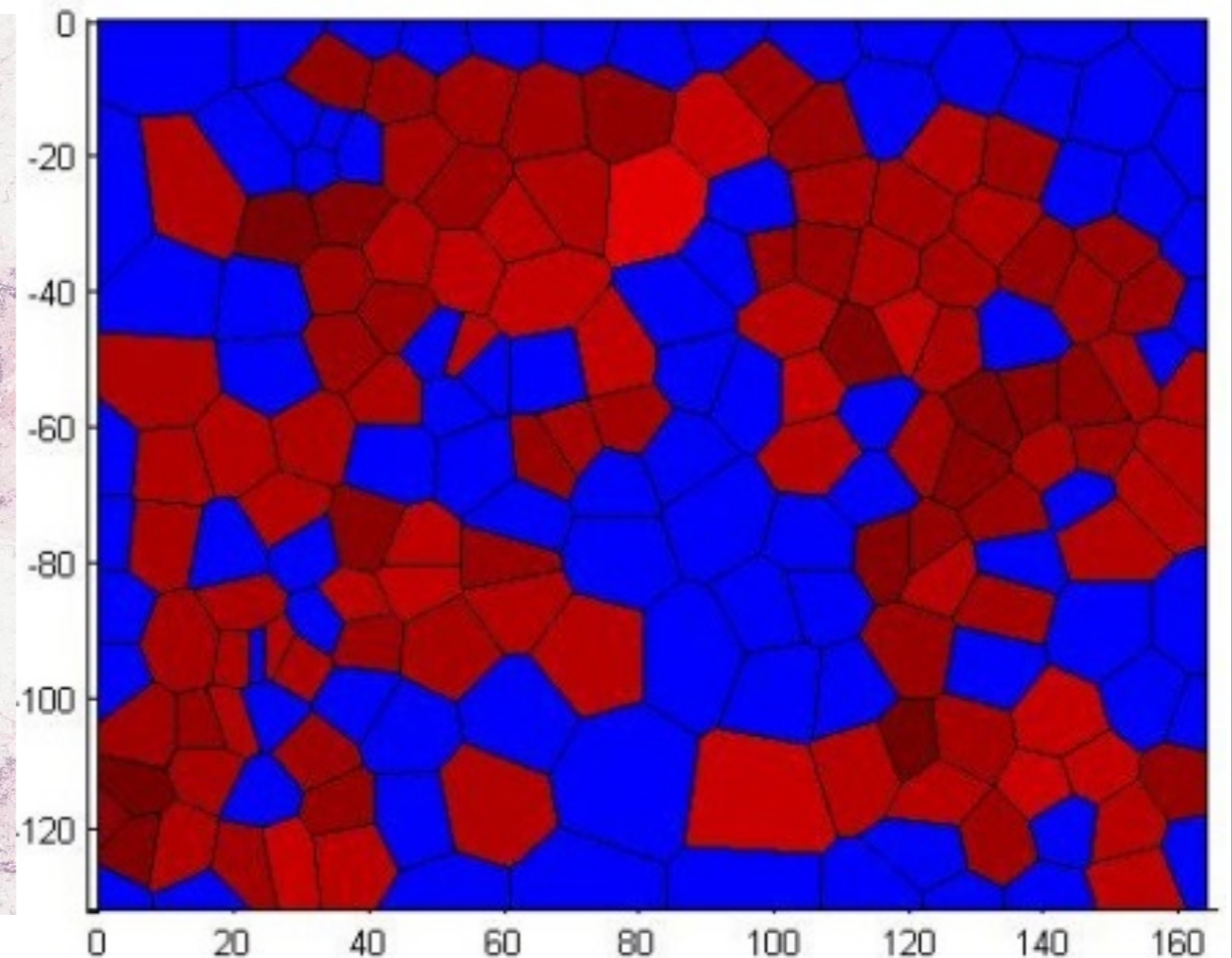
## Comparison of ITM<sup>2</sup>C and MITM<sup>3</sup> Frameworks



## Switching from HPF to WSI Analysis



HPF analysis for mitotic count / nuclei atypia in WSI

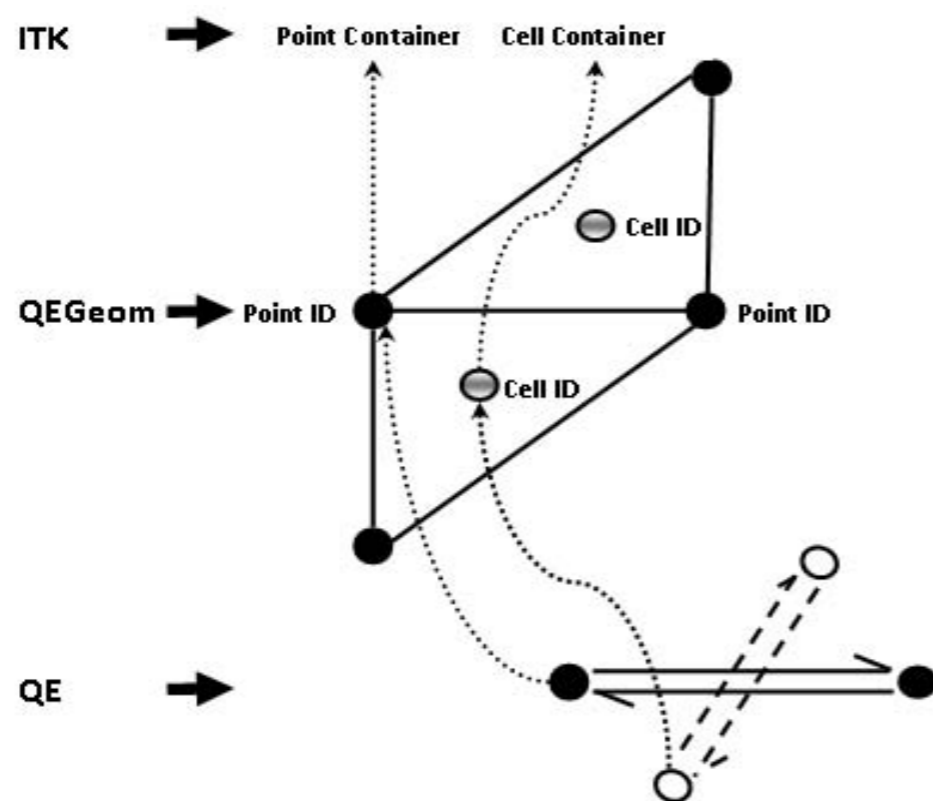


Voronoi Diagram computed using analyzed HPF of WSI

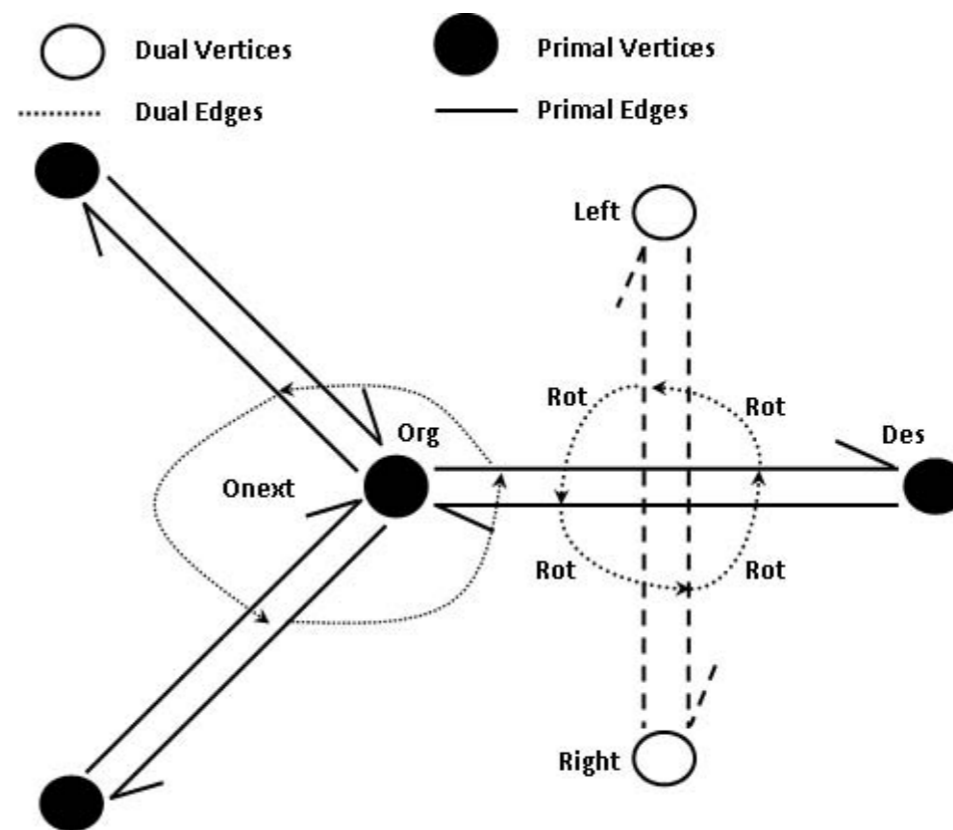
## Orientable 2-Manifold Meshes and Existing Data Structure

- *itk::QuadEdgeMesh* existing data structure in ITK can handle discrete 2-manifold surfaces
- A constant complexity local access on modifications

	Primal	Dual
Geometry	Yes	<u>No</u>
Topology	Yes	Yes



QuadEdgeMesh Structure

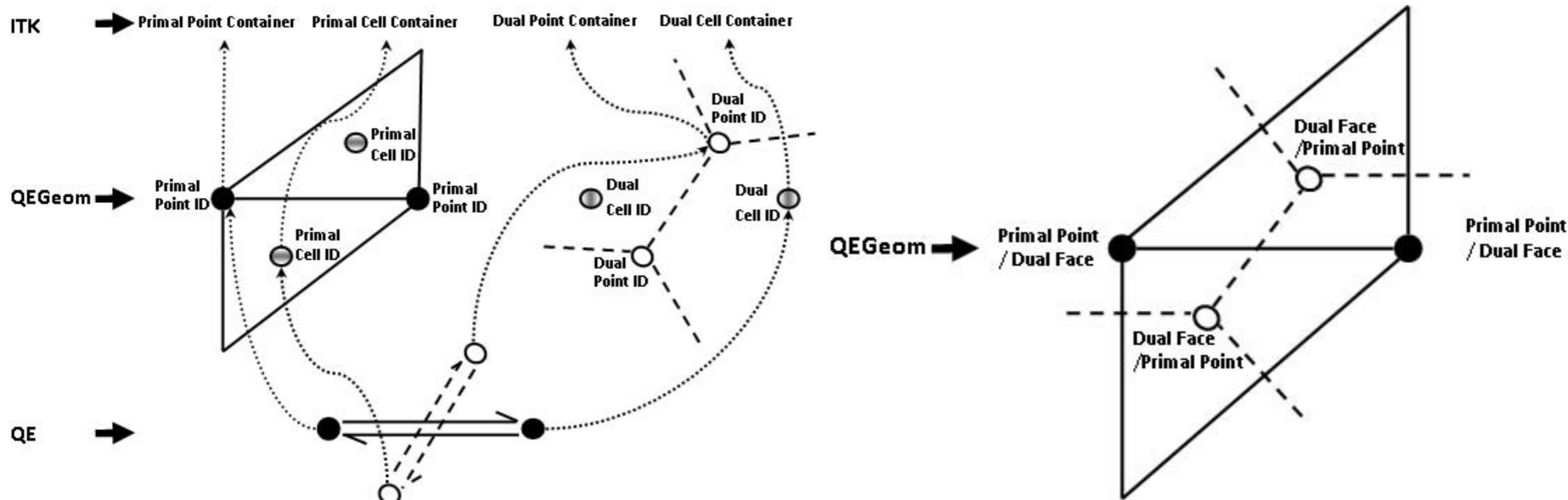


QuadEdge Structure

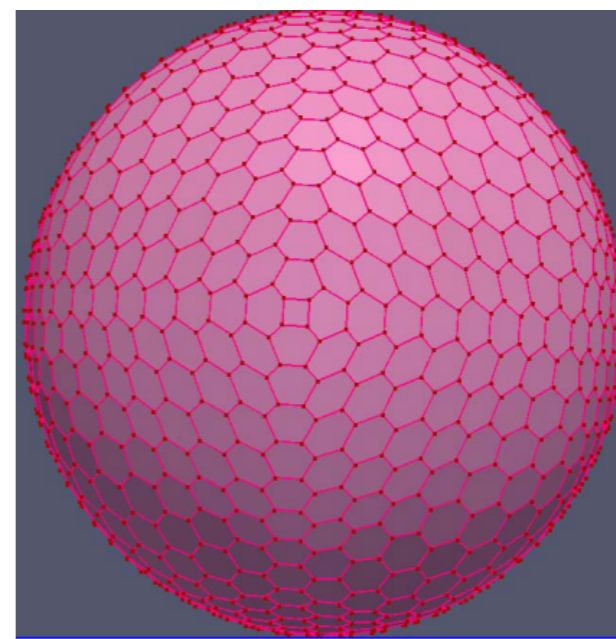
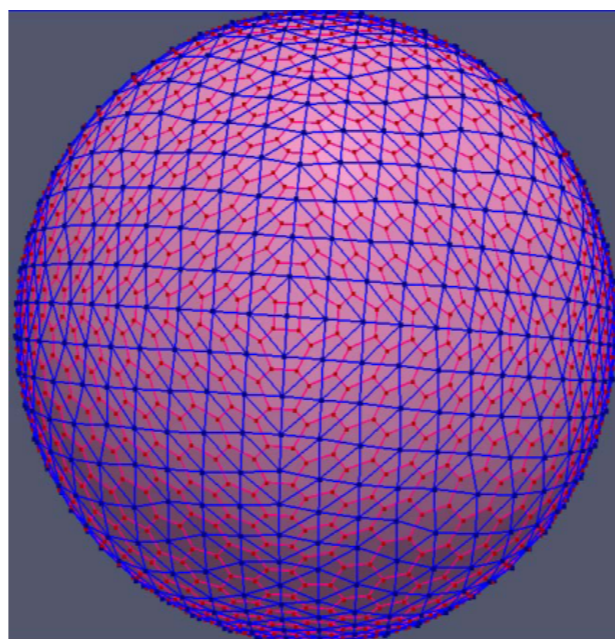
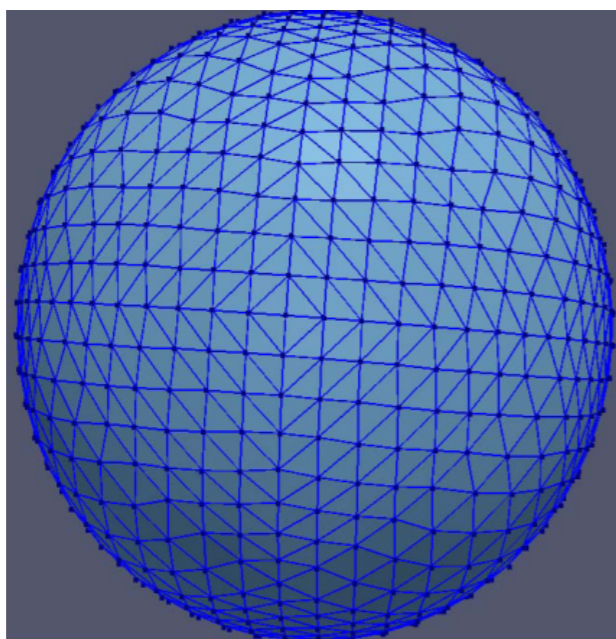
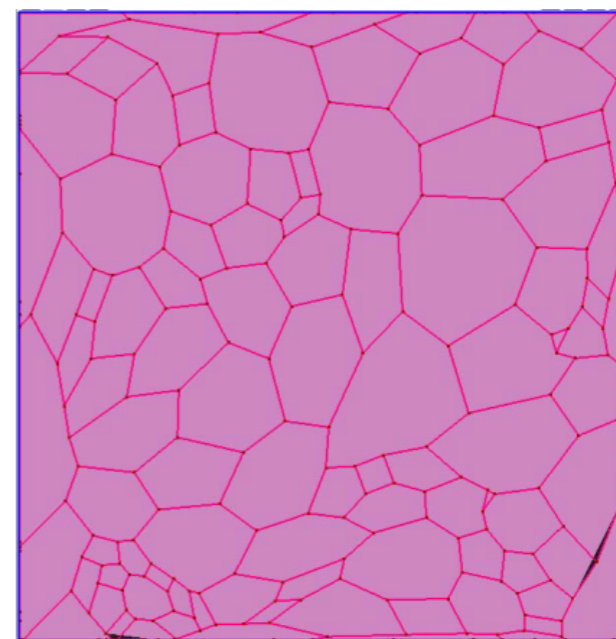
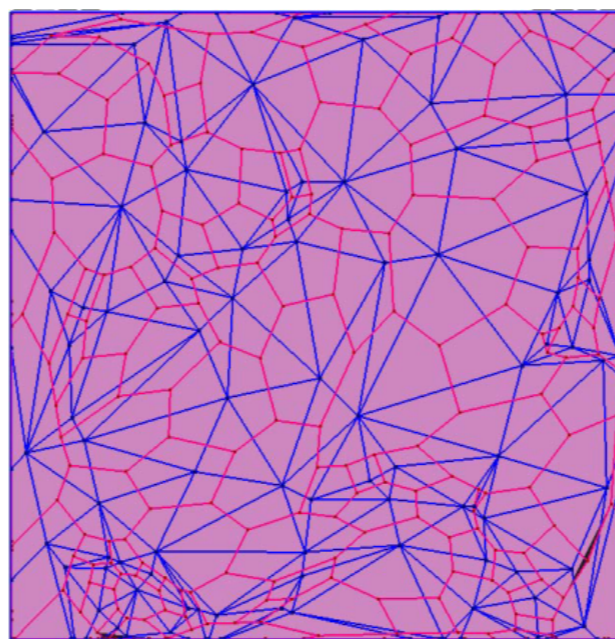
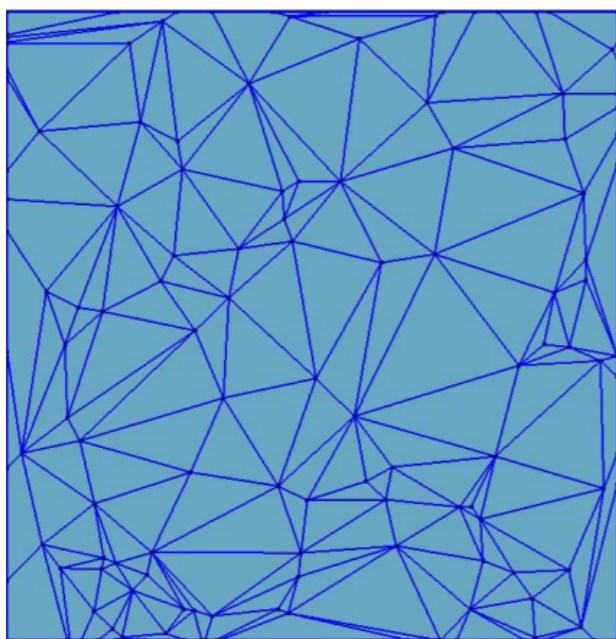
## Orientable 2-Manifold Meshes and NewData Structure

- Proposed an extension of existing ITK data structure for Orientable 2-manifold meshes to handle duality
- itk::QuadEdgeMeshWithDual* new data structure and a filter that transform primal mesh to primal/dual mesh

		Old Structure	New Structure
Changes	OriginRefType	Point ID, Cell ID	Pair< Point, Cell >, Pair< Cell, Point >
Additions	Dual Containers	-	Dual Pointers, Cells and EdgeCells Containers



## Planer Delaunay/Voronoi Mesh and Non-Planer Triangulation/Simplex Mesh



Primal Mesh

Primal with dual Mesh

Dual Mesh

## MICO Project (ANR TecSan)

- COgnitive virtual Microscope for Breast Cancer Grading (MICO) Project
  - Funded by French National Research Agency (ANR)
  - Launched in Feb, 2011 – Jul, 2014 (3.5 Years)



**IPAL — University Joseph Fourier**  
Image Processing, Ontologies, Reasoning



**Pitié-Salpêtrière Hospital**  
Pathologists, Expertise on breast cancer



**University Pierre and Marie Curie**  
Contextual graphs, GPU



**TRIBVN**  
User interface



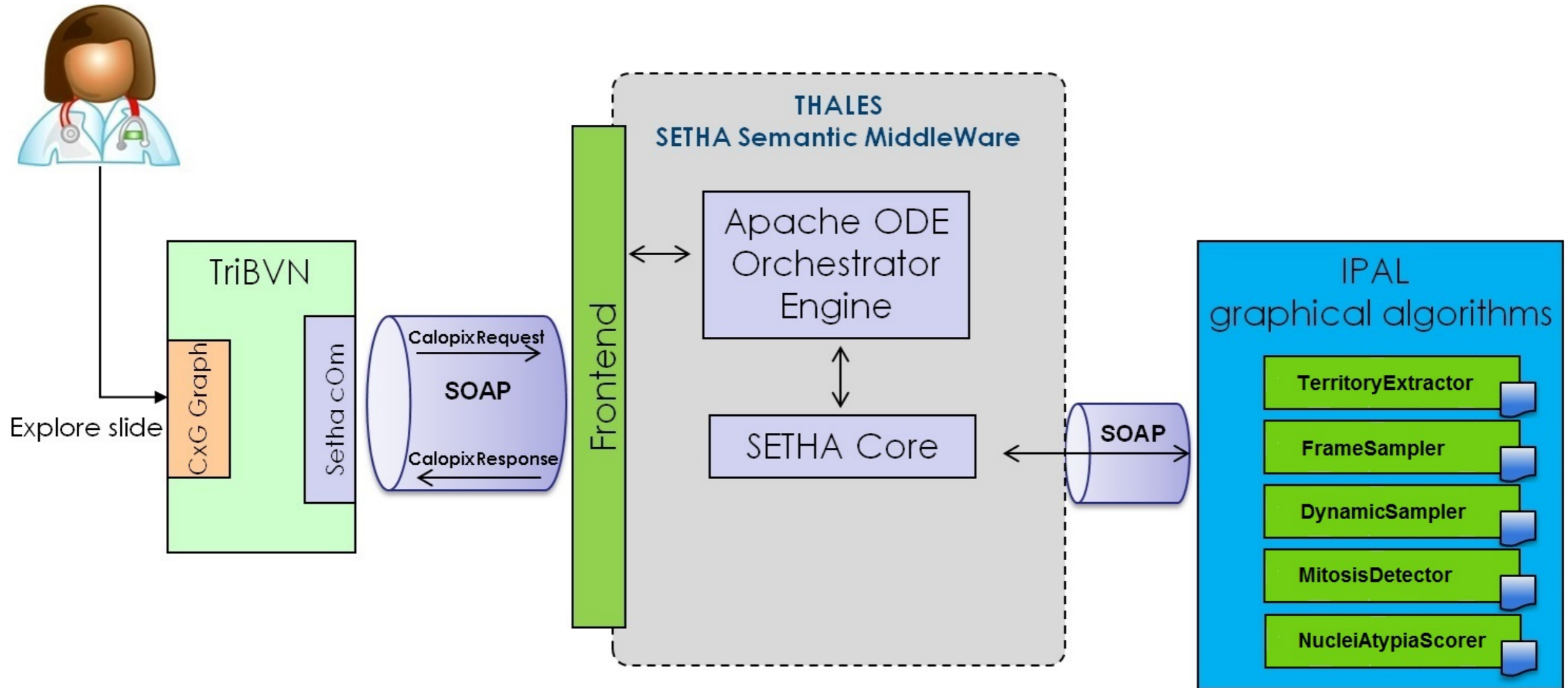
**Agfa HealthCare**  
Ontologies, Reasoning



**Thales**  
Semantic middleware

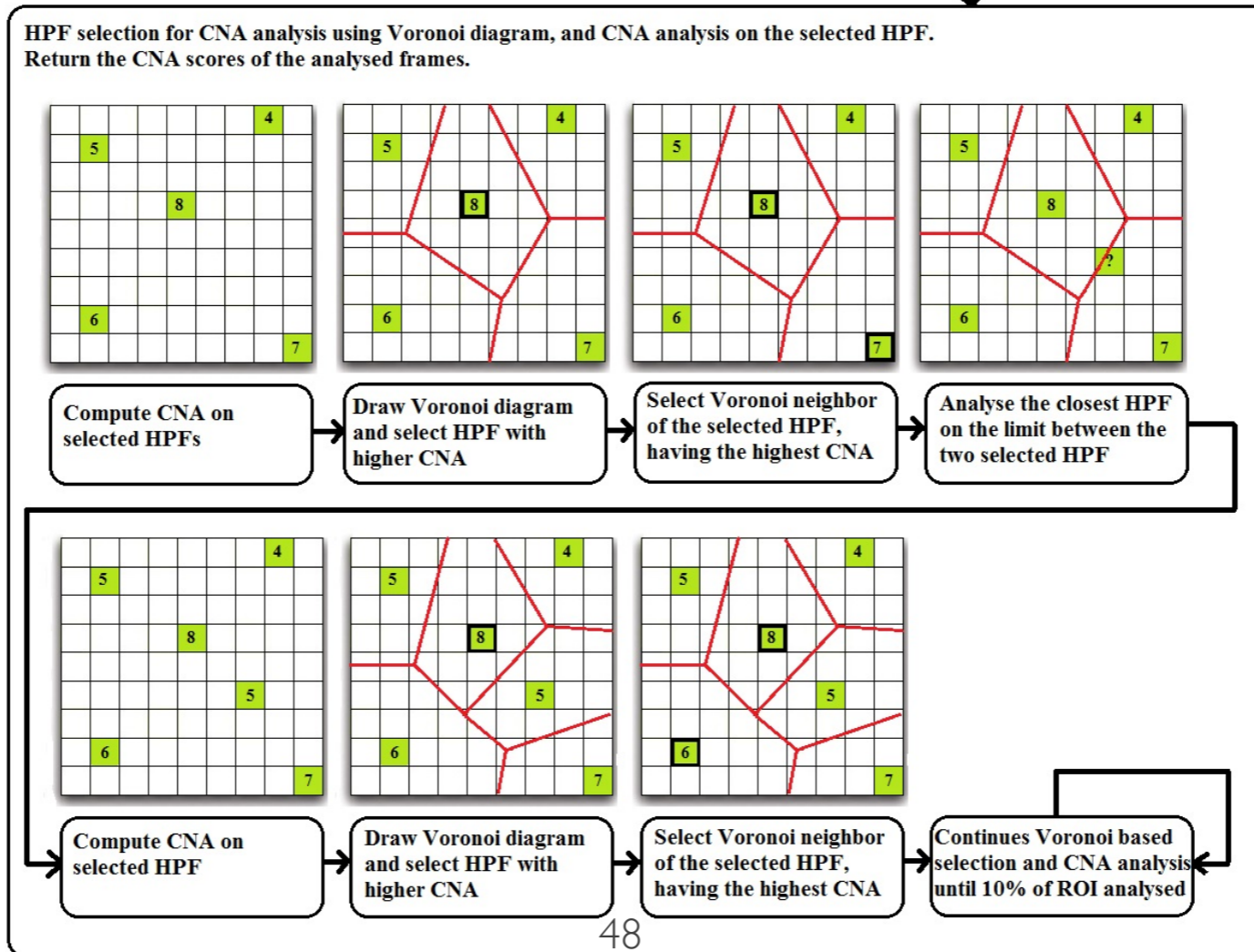
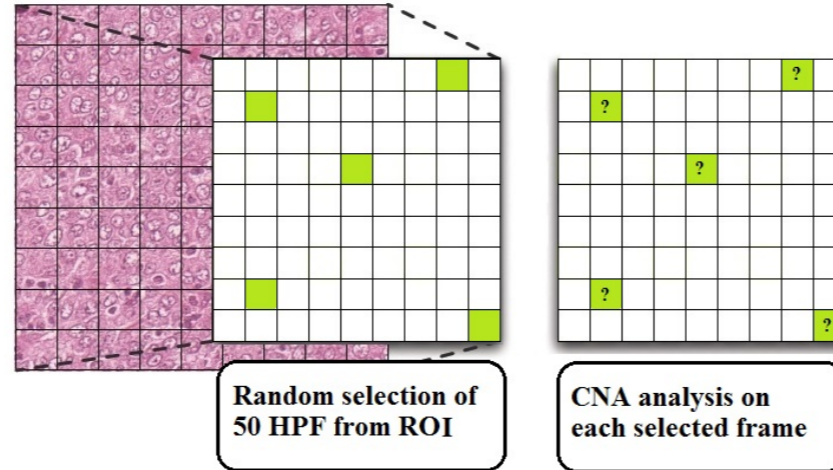
MICO ANR TecSan Project Partners

## MICO Architecture



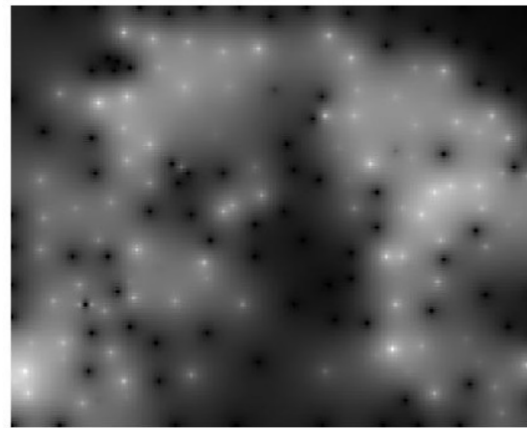
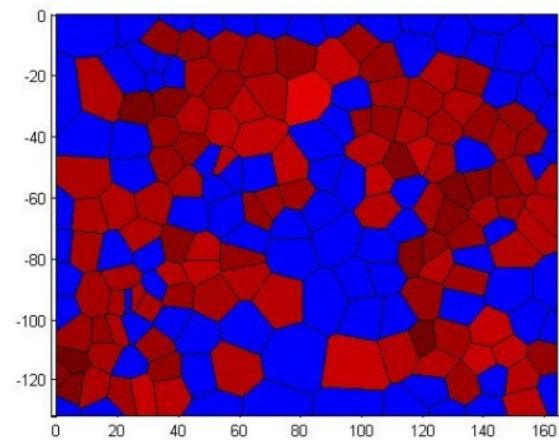
MICO 2.0 Architecture

## Dynamic Sampling applied over ROI for CNA Evaluation

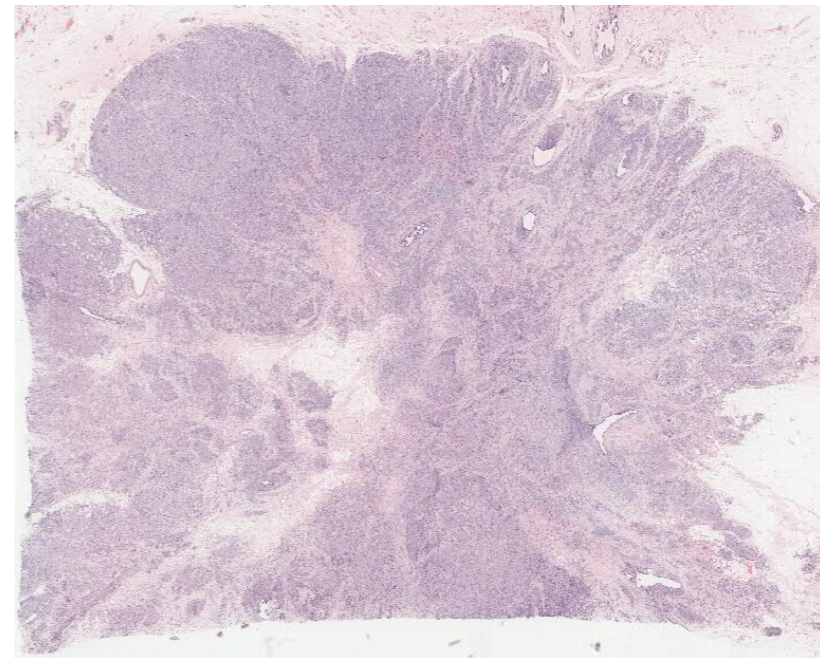




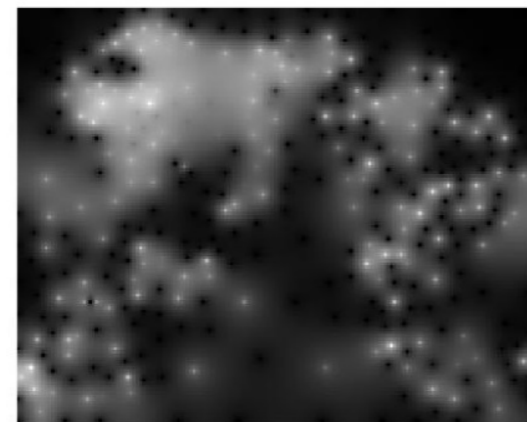
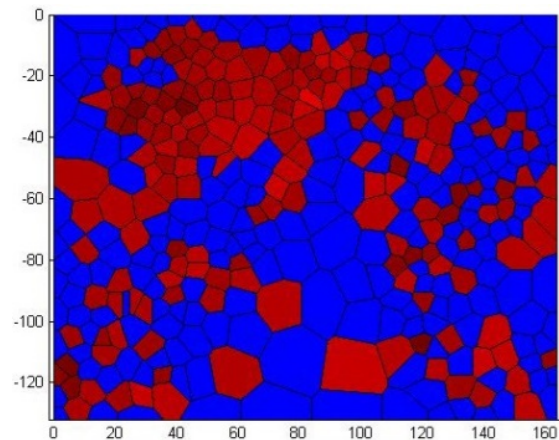
## Dynamic Sampling applied over WSI: Incrementally Voronoi Diagram



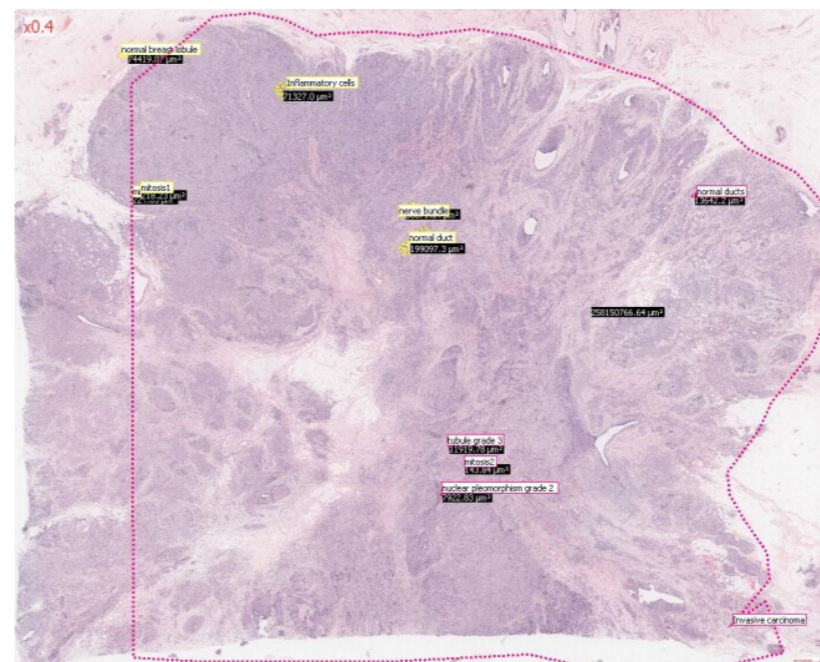
After 200  
Iteration



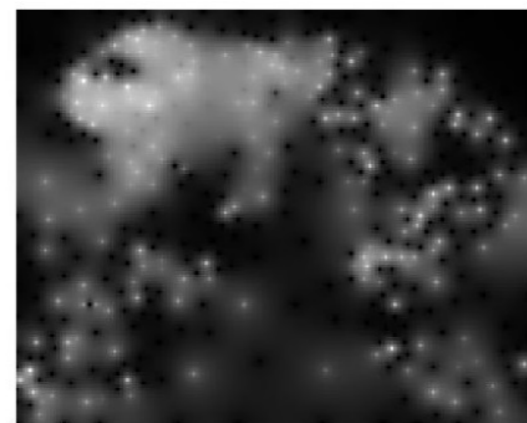
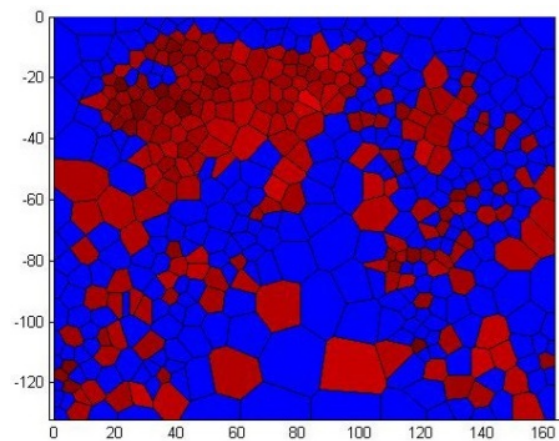
Actual WSI



After 300  
Iteration



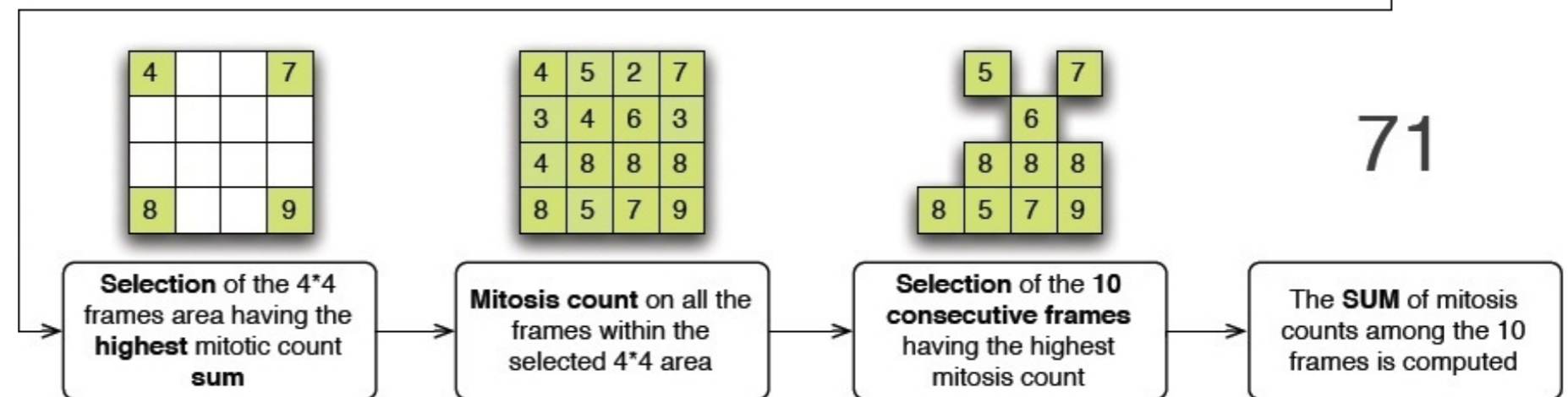
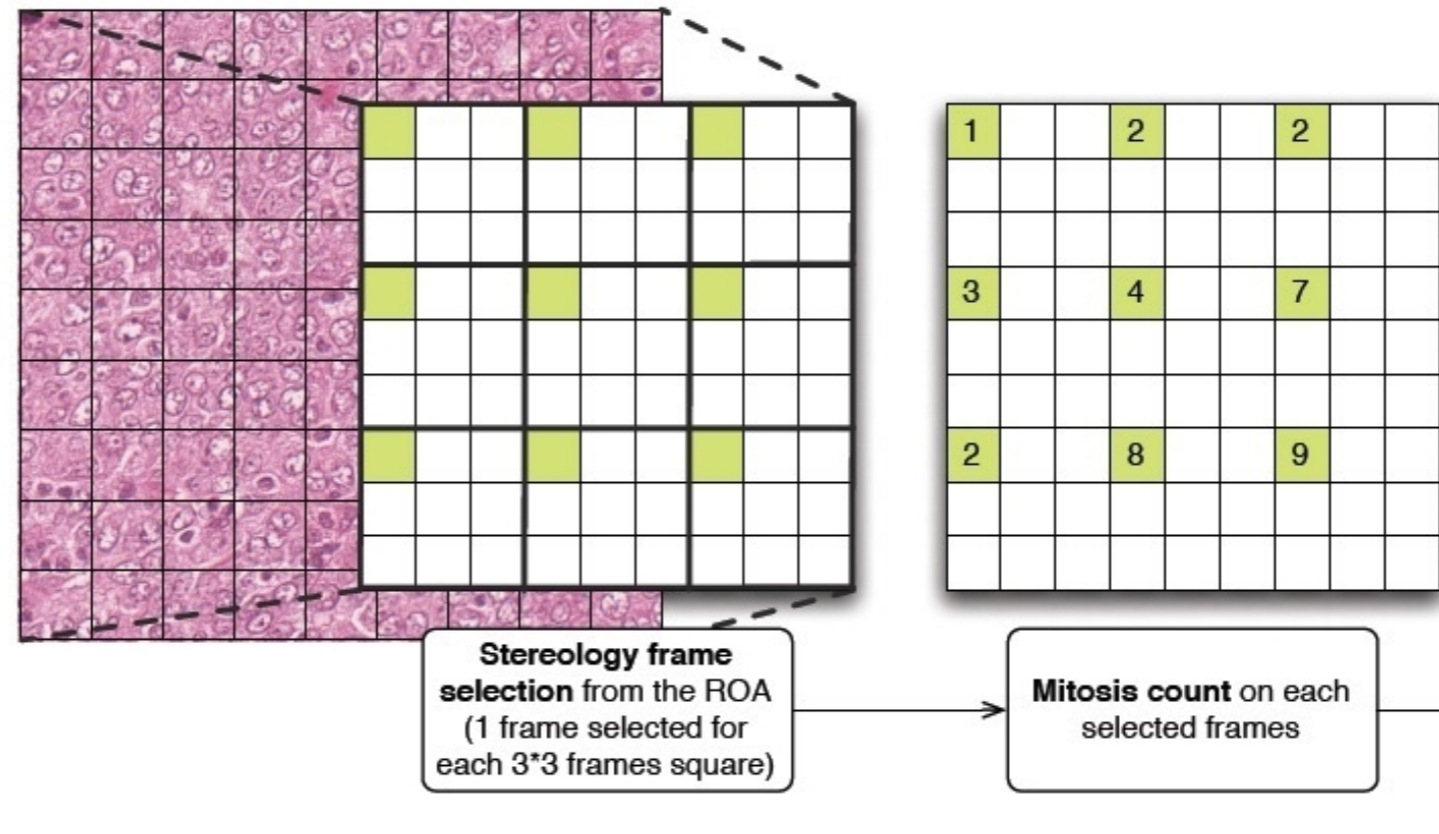
Annotated WSI



After 500  
Iteration

## Stereology Framework for evaluation of ITM<sup>2</sup>C framework in MICO

- TerritoryExtractor
- FrameGenerator
- FrameSampler 3x3
- FrameSampler 4x4
- ITM<sup>2</sup>C Framework
- MitosisScorer



Stereology Flow used for Mitosis Score over a ROI

## WSI analyzed by ITM<sup>2</sup>C Framework are displayed on Calopix platform

The color code is based on the number of mitosis detected in the frame (from blue for zero mitosis to red for 10 or more mitosis).

The screenshot displays the Calopix software interface. The main window shows a histological image with a grid of colored squares overlaid on it. The color of each square represents the number of mitoses detected in that frame, ranging from blue (zero mitoses) to red (10 or more mitoses). The image is labeled 'x0.3' and has a scale bar of 5000 μm.

The interface includes a menu bar (File, Services, Viewer, Help) and a toolbar. The Explorer panel on the left shows the file '09C17270-2-HE.wfml'. The right-hand panel contains a table with analysis data and several control buttons.

Global data	Analysis	Annotations	Label	ROI
<input type="checkbox"/>	Title	Wide field	Numb...	
<input type="checkbox"/>		1 09C17270-2-HE	2	

Below the table, there are several control panels:

- Title:** A text input field with a 'Modify' button.
- Ipal mitosis analysis:** A section with a 'Number of mitoses' input field set to '60' and a 'Display results' button.
- Date:** A section with a 'Date' input field set to 'Thu Jun 21 17:00:52 CEST 2012' and a 'Delete analysis result' button.
- Ipal mitosis analysis (repeated):** A second section with a 'Number of mitoses' input field set to '60' and a 'New analysis' button.

The status bar at the bottom shows 'Zoom : 0%', 'Lens : x40.0', and 'x = 40639 μm y = 16989 μm'.

## Mitosis Detector Integration in Calopix

The screenshot displays the CaloPix software interface. The main window shows a histological image with a central tumor region (Territoire de la tumeur) and surrounding adipose tissue (Tissu Adipeux). The tumor region is densely packed with blue labels 'Gx40' and a green 'Mitose' label. A scale bar at the bottom indicates 10000 µm. The interface includes a menu bar (File, Services, Viewer, Help), a toolbar, and a right-hand panel with a table of analysis data.

Global data	Analysis	Annotations	Label	ROI
<input checked="" type="checkbox"/>	Title	Wide field		Numb...
<input type="checkbox"/>	test1	1 08C18251-9-HE-0...		0

Additional interface elements include a 'Title' field with 'test1' and a 'Modify' button, and a 'New analysis' button at the bottom right. The status bar at the bottom shows 'Zoom : 0%', 'Lens : x40.0', and 'x = 52064 ... = 6118 µm'.

## Summary

- ▶ Proposed automated mitosis detection framework for different scanners and multispectral microscope
- ▶ Efficient and generic strategies (Stereology & Dynamic Sampling) to explore WSI
- ▶ Evaluation of these frameworks in MICO platform

## Future work

- ▶ Expand proposed frameworks from two-class problem to multi-class problem and classify other microscopic objects like lymphocytes, apoptosis, normal nuclei, cancer nuclei
- ▶ Main area of Interests:
  - Machine Learning
  - Computer Vision
  - Pattern Recognition
  - Medical Image Analysis

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  - Computer Vision
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  - Medical Image Analysis

## Intellectual Property

- “*MitosisDetector – Mitosis Detector for Histopathology*”, H. Irshad, L. Roux, D. Racoceanu, *Copyright CNRS (CNRS Statement Software) No. DL 05963-01 for 2955 IPAL UMI, 2013.*

## Journals

1. H. Irshad, A. Gouaillard, L. Roux, D. Racoceanu, "Multispectral Band Selection and Spatial Characterization: Application to Mitosis Detection in Breast Cancer Histopathology", in *Computerized Medical Imaging and Graphics (CMIG)*, (Submitted).
2. H. Irshad, A. Veillard, L. Roux, D. Racoceanu, "Methods for Nuclei Detection, Segmentation and Classification in Digital Histopathology: A Review. Current Status and Future Potential", in *IEEE Reviews on Biomedical Engineering (RBME)*, 2013, vol. PP, issue 99, pp. 1.
3. H. Irshad, I. Hassan, J. Iqbal, A. R. Aghdam, M. Kamalpour, "m-Health System Support For LHWs Working in Rural Areas", in *Journal Science International-Lahore*, July-Sept., 2013, Vol. 25, issue 3, pp. 653-655.
4. H. Irshad, "Automated Mitosis Detection in Histopathology using Morphological and Multi-channel Statistics Features", in *Journal of Pathology Informatics*, May, 2013, vol. 4, issue 1, pp. 10.
5. L. Roux, D. Racoceanu, N. Loménie, M. Kulikova, H. Irshad, J. Klossa, F. Capron, C. Genestie, G. L. Naour, M. N. Gurcan, "Mitosis detection in breast cancer histological images An ICPR 2012 contest", in *Journal of Pathology Informatics*, May, 2013, vol. 4, issue 1, pp. 8.
6. H. Irshad, S. Jalali, L. Roux, D. Racoceanu, L. J. Hwee, G. L. Naour, F. Capron, "Automated Mitosis Detection using Texture, SIFT Features and HMAX Biologically Inspired Approach", in *Journal of Pathology Informatics*, March, 2013, vol. 4, issue 2, pp. 12.

## Technical White Paper (pubmed Indexed)

7. H. Irshad, S. Rigaud, A. Gouaillard, "Primal/Dual Mesh with Application to Triangular / Simplex Mesh and Delaunay / Voronoi", in *Insight Journal*, January-December, 2012.

## Peer-reviewed International Conference

8. H. Irshad, A. Gouaillard, L. Roux, D. Racoceanu, "Spectral Band Selection for Mitosis Detection in Histopathology", in *11th International Symposium on Biomedical Imaging (ISBI)*, Beijing China, 2014.
9. H. Irshad, L. Roux, D. Racoceanu, "Multi-channels Statistical and Morphological Features based Mitosis Detection in Breast Cancer Histopathology", in *Proc. of 35th Inter. Conf. of the IEEE Engineering in Medicine and Biology Society (EMBC)*, Osaka, Japan, Jul., 2013, pp. 6091-6094.
10. H. Irshad, L. Roux, O. Morère, D. Racoceanu, G. L. Naour, and F. Capron, "Détection automatique et calcul du compte de mitoses sur lames H&E", in *Recherche en Imagerie et Technologies pour la Santé (RITS)*, Bordeaux, France, Apr., 2013.
11. H. Irshad, L. Roux, D. Racoceanu,, "Multi-channel Statistics Features based Mitosis Detection in Histopathology", in *International Workshop on Pattern Recognition and Healthcare Analytics, 21st Inter. Conf. on Pattern Recognition (ICPR)*, Tsukuba, Japan, Nov., 2012.
12. H. Irshad, S. Jalali, L. Roux, D. Racoceanu, L. J. Hwee, G. L. Naour, F. Capron, "Automated Mitosis Detection Using Texture, SIFT Features and HMAX Biologically Inspired Approach", in *Workshop on Histopathology Image Analysis, 15th Inter. Conf. on MICCAI*, Nice, France, Oct., 2012.
13. S. Naz, H. Irshad, H. Majeed, "Image Segmentation using Fuzzy Clustering: A Survey", in *Proc. of IEEE Inter. Conf. on Emerging Technologies*, Islamabad, Pakistan, Oct., 2010, pp. 181-186.
14. H. Irshad, S. Athar, F. Shahzad, M. Farooq, "M-Health System with focus on Antenatal Care for Rural Areas", in *First Inter. Conf. on eHealth (e-HAP)*, Karachi, Pakistan, Jan., 2010.
15. H. Irshad, S. Athar, F. Shahzad, A. Bashir, F. Jehan, "On The Move Ultrasound Diagnosis on Mobile", in *First Inter. Conf. on eHealth (e-HAP)*, Karachi Pakistan, Jan., 2010.
16. J. Afridi, M. Kamran, H. Irshad, S. Khan, M. Farooq, "Use of CDSS on the Personal Digital Assistant of the Medical Expert", in *First Inter. Conf. on eHealth (e-HAP)*, Karachi, Pakistan, Jan., 2010.
17. H. Irshad, M. Kamran, A. B. Siddiqui, A. Hussain, "Image Fusion using Computational Intelligence: A Survey", in *Proc. of IEEE Second Inter. Conf. on Environment and Computer Science*, Dubai, UAE, Dec., 2009, pp. 128-132.



Questions ?



Humayun Irshad

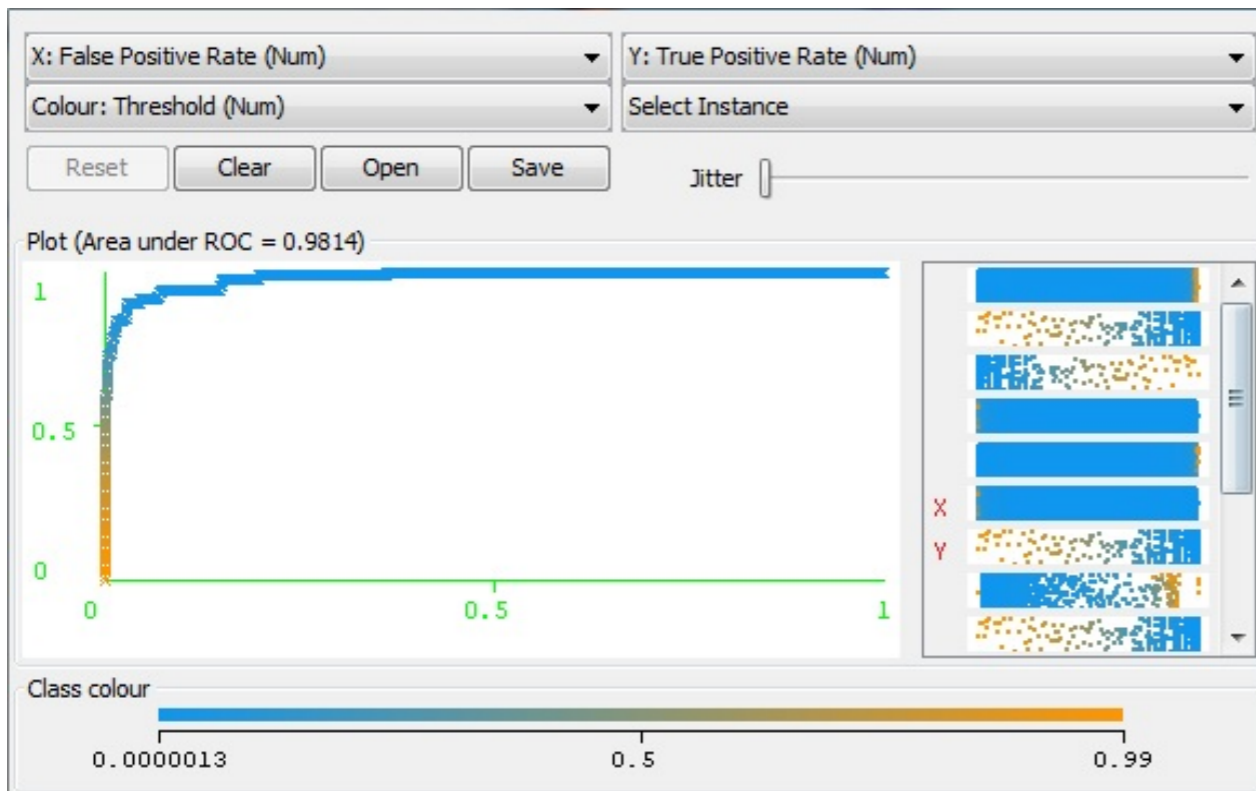
[Humayun.irshad@ipal.cnrs.fr](mailto:Humayun.irshad@ipal.cnrs.fr)

Image & Pervasive Access Lab

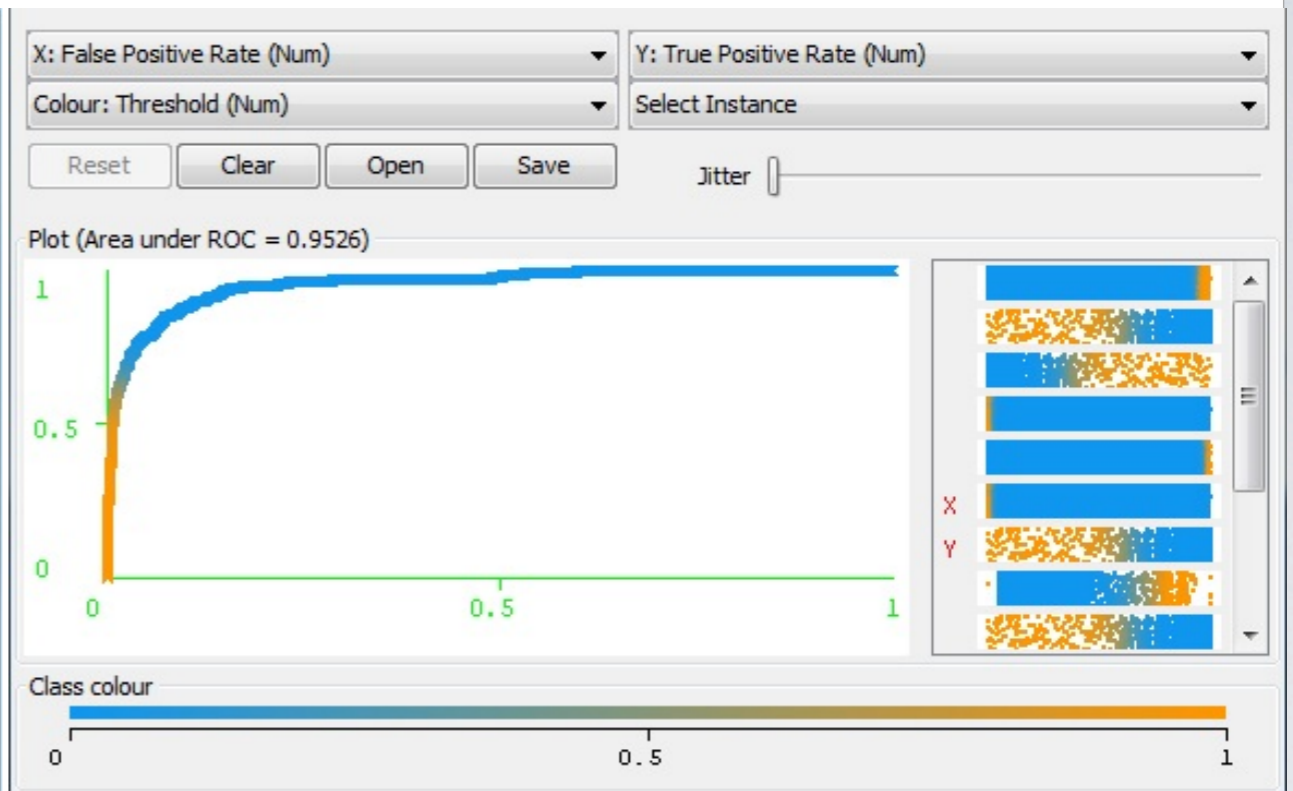
International joint research unit - UMI CNRS 2955

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## Receiver Operating Characteristic (ROC) curve of patch based features with LSVM Classifier

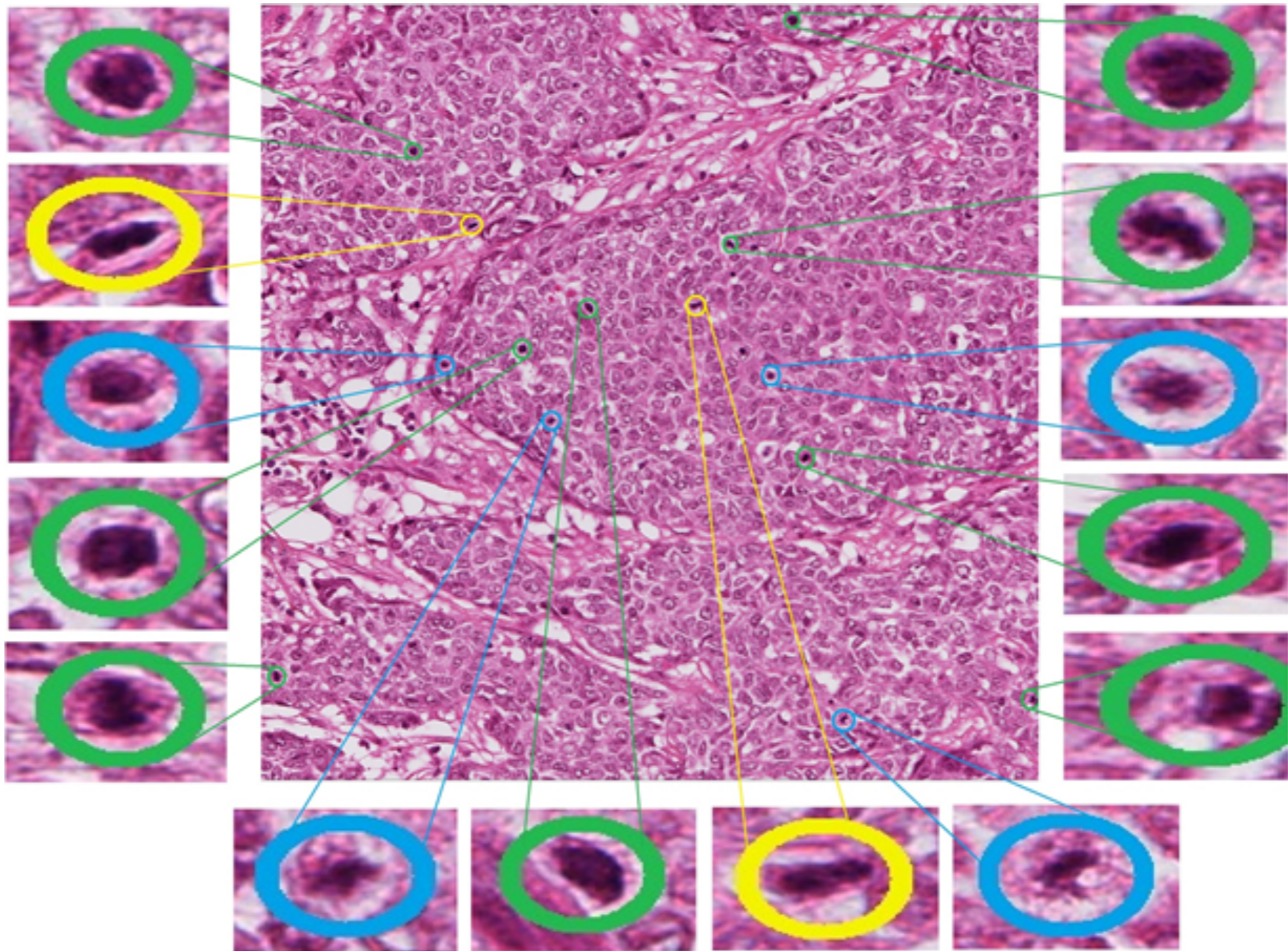


On Aperio Dataset



On Hamamatsu Dataset

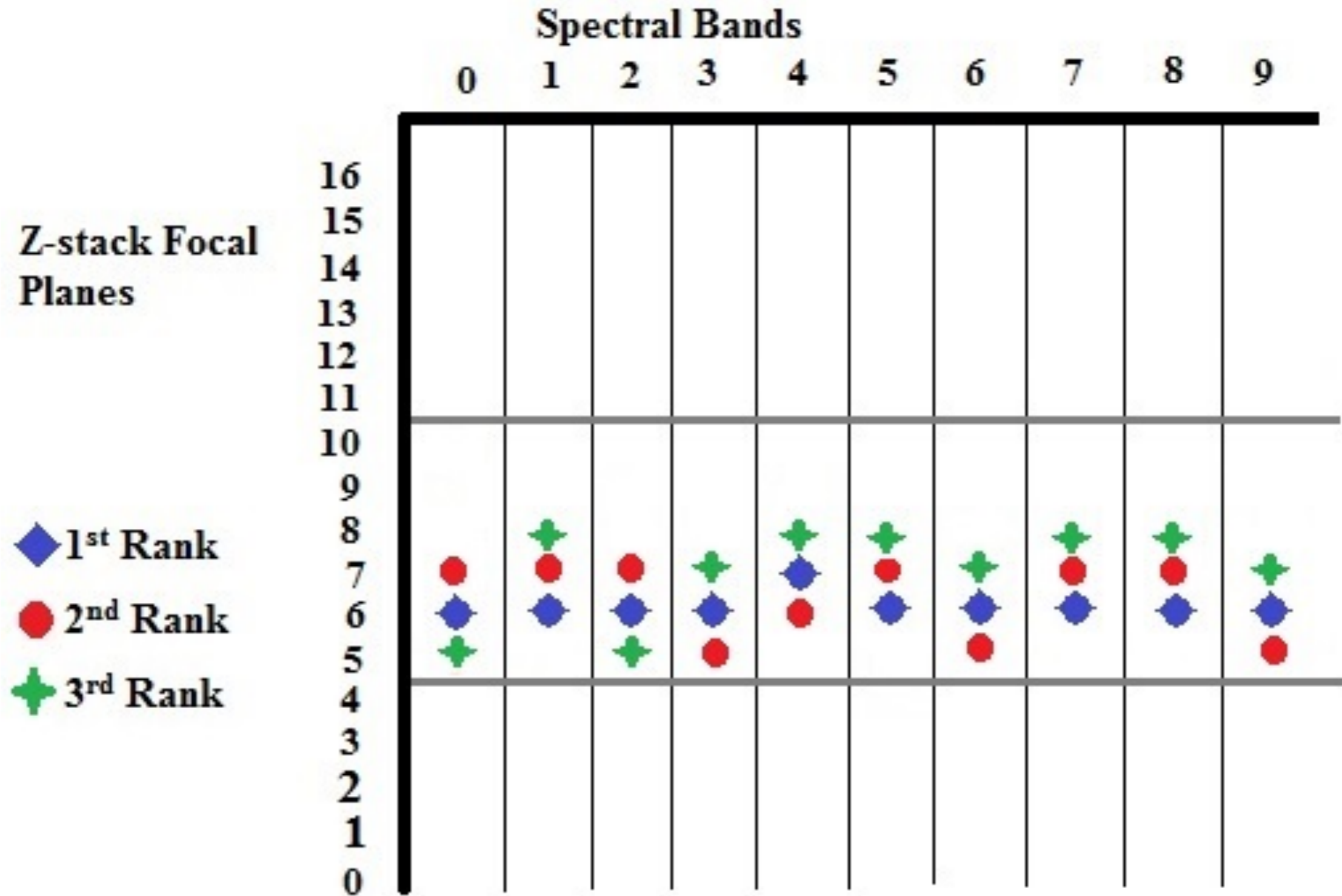
Candidate Classification on Hamamatsu Dataset (TP=Green, FP=Yellow, FN-Blue)



## Classification Results with White, Red, Green and Blue SBs using 5-Fold Cross Validation

Features	Classifiers	TPR	PPV	FM
Red SBs (SB 0,8,9)	DT	51%	63%	56.11%
	MLP	48%	71%	57.46%
	LSVM	67%	56%	61.19%
	NLSVM	49%	75%	59.19%
Green SBs (SB 5,6,7)	DT	50%	68%	57.55%
	MLP	50%	65%	56.84%
	LSVM	65%	58%	61.14%
	NLSVM	48%	78%	59.39%
Blue SBs (SB 2,3,4)	DT	43%	59%	49.82%
	MLP	46%	69%	55.49%
	LSVM	54%	60%	56.81%
	NLSVM	46%	75%	56.65%
White SB (SB 1)	DT	42%	65%	51.32%
	MLP	44%	74%	55.15%
	LSVM	56%	52%	54.11%
	NLSVM	44%	77%	55.84%

## Top 3 Ranked Focal Planes



## Dynamic Sampling for Cyto-Nuclear Atypia Score

- A dynamic sampling framework was developed based on computational geometry for Cyto-Nuclear Atypia (CAN) evaluation to avoid exhaustive analysis on WSI
  - Main steps of method are:
    1. Pathologist annotated territories by observing WSI using Calopix user interface
    2. Territories are extracted from WSI and split into several HPF frames
    3. 50 HPF are randomly selected for computation CNA scores using Christophe and Maria method [5]
    4. These scores are used for initialization of Voronoi diagram
    5. Next HPF is selected based on two criteria
      1. At least one of its neighboring Voronoi cells has a high score that control the convergence of method towards areas with high score
      2. The distance between the new sample and its neighbors is not too short that prevents oversampling
    6. The final overall CNA score is the grade of the most atypia frame
- 5) Christophe & Maria, Marked point processes with simple and complex shape objects for cell nuclei extraction from breast cancer H&E images, SPIE Medical Imaging, 2013.

## Dynamic Sampling Algorithm

**Input:** Current frames  $E$ , Voronoi Diagram  $VD_E$ ,  $p$ ,  $d$ ,  $\max_E$

**Output:** updated frames  $E$ , Voronoi Diagram  $VD_E$ ,  $\max_E$

Compute  $V_E$

Sort  $V_E$  according to decreasing distance to  $E$

**for** every  $x \in V_E$  **do**

**if** Distance( $x$ ,  $E$ ) >  $d$  **then**

**if** MaxScore( $x$ ) >  $p \times \max_E$  **then**

$E = E \cup \{x\}$

            Update  $VD_E$

$\max_E = \max(S(x), \max_E)$

            break loop

**end if**

**Else**

        Break loop

**End if**

**End for**