# WAVELET SCATTERING REPRESENTATIONS FOR CLASSIFICATION OF HISTOLOGY IMAGES

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#### **ABSTRACT**

We use the Wavelet Scattering Convolution Network (Scat-Net), as defined in Bruna and Mallat (2013), to build invariant representations of microscopic images of breast tissue. We then classify the image segments as containing primarily epithelium or stroma. We evaluate multiple systems, and achieve results comparable to a strong baseline. Some of our systems surpass the accuracy of a best-effort classification of the data by our group members.

*Index Terms*— image invariants, scattering transform, convolution network, medical image analysis

#### 1. INTRODUCTION

Breast cancer is one of the leading diseases in the world, and it leads to nearly a million deaths a year across the world. The most common diagnosis method for breast cancer involves microscopic examination of the breast tissues. Currently, this is a labor-intensive manual process. Motivated both by costsaving and by potential elimination of human error, research efforts are ongoing to automate as well as improve the diagnosis process. An important sub-problem in automatic diagnosis pipelines is classifying portions of images as mostly *epithe-lium*, lining and surface cells, or *stroma*, connective cells [1].

## 2. DATASET

Microscope slide images are often large and contain many regions of both epithelium and stroma, and also regions that are distorted and not of interest. As part of the diagnosis pipeline, the digitized slide is segmented into small images, also called *super-pixels*. We address the task of classifying whether these super-pixels predominantly contain stroma or epithelium. Segmented images were graciously provided by Raghu Machirajus research group at the Ohio State University. The images had been hand-labeled by researchers trained in the task. Clear examples of epithelium and stroma can be found in **Figure 1**, as well as an example that has a well-balanced proportion of both epithelium and stroma, and is therefore difficult to classify, even for humans.

We evaluated two datasets, one set containing smaller largely unambiguous 80 x 80 pixel images and the other set

containing larger images of varying sizes. We had 600 of the 80 x 80 images, and 734 of the larger ones, with an equal distribution of images labelled epithelium or stroma in both. Near perfect classification on the smaller images was easily obtained and lot of the work focused on the dataset with the 734 variably sized superpixels. We held out 100 randomly-selected images from the super-pixels as the test set, and used the remaining 634 for training and cross-validation tuning of hyper-parameters.

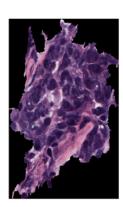
#### 3. SCATNET

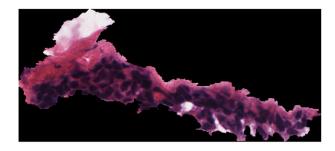
Each image contains a large amount of data, across all the pixels. Often, we care only about certain "important" aspects of the information in the image, which can be represented compactly. These features should be invariant to transformations such as translation, rotation, scale, and so on. For example, in this application, the position at which epithelial cells are located in the image is irrelevant, and their orientation is also irrelevant.

Transformations such as SIFT [2] provide some invariant properties and a compact representation of key information. However, in a hypothetical scenario, we may also want to control the degree to which we want our output to be invariant to such transformations. The transform described by Beck et. al. [1], generates a compact representation of the image, and provides control for the degree of invariance desired. Bruna and Mallat [3] suggest that for complex structures over large domains, locally invariant information from SIFT, etc. is insufficient. They say that the Scattering Transform provides not only SIFT-like descriptors at the output of Layer 1, but also higher order information at the further layers, which can help discriminate such complex structures.

The scattering transform is similar to the form of a deep convolution network with wavelets being the convolving functions at each node. As in **Figure 2**, at each layer, the input from the previous layer is convolved with wavelets at different scales and orientations. This is essentially doing a complex wavelet transform, where the information from the Image signal is divided in the complex frequency plane. [4] The number of divisions along the radial axis, J, and L, the number of divisions along the rotational axis, are both configurable. Since the outputs are taken through averaging







**Fig. 1**. Clockwise from top left – super-pixels containing stroma, epithelium, and a mixture of both (labeled stroma)

filters, in order to get the desired invariant properties, some high frequency information is lost. To recover the same, the un-averaged information is passed to the next layer, where it is again convolved with the wavelet filters. All the outputs are finally combined to form a feature vector. Since, the feature vector contains correlated values, a classifier based on affine principal component analysis (PCA) is used. It computes a transformation to de-correlate the values and reduce the dimensionality to N, for each class, in this case epithelium and stroma. The classification is done by looking at which class' transformation minimizes the approximation error. More details about the scattering transform, as well as the affine PCA classification technique are available in Bruna and Mallat [3].

ScatNet also allows configurability of the number of layers, M, the number of scales per octave in the wavelet transform; Q, the amount of oversampling carried out before each layer; and the dimensionality of the affine PCA classifier N. After exploratory preliminary testing, we tried all configurations of J=4, L=3 to 6, Q=1 to 10, M=1 to 4, oversampling=1 to 2, and N=2 to 20. Of the two best performing configurations, we randomly re-partitioned the training 10 times and took the average, finding the highest-performing configuration to be J=4, L=6, Q=1, M=3, oversampling=1, N=2.

ScatNet works with grayscale images and expects the images to be uniformly sized and rectangular. Our data, however, is colored and has wide variation in image sizes and aspect ratios, and the segments aren't rectangular. Further,

Hyperparameters	Performance
J = 4, L = 5, Q = 1, M = 3, o.s. = 1, N = 15	84.2%
J = 4, $L = 6$ , $Q = 1$ , $M = 3$ , $o.s. = 1$ , $N = 2$	87.2%

**Table 1**. Results for hyper-parameter configuration that achieve optimal results in exhaustive search

the staining process causes color to contain valuable information: stromata stain a dull pink, while epithelial cells stain a deeper purple. Our solution involves transforming the image from RGB space to HSV (Hue-Saturation-Value) space, and compute the feature vector independently for each color plane, eventually concatenating the 3 vectors to make the final feature vector. To account for the varying image sizes and non-rectangular segments, we normalize the final feature vector, dividing it by the number of non-zero pixels in the input image.

#### 4. BASELINE SYSTEM

As a baseline, we applied a standard computer vision technique referred to as *bag-of-visual-words* [5] [6]. This technique involves finding points of interest (called *keypoints*) and some useful information about them (called *descriptors*) and classifying based on the similarity of these descriptors. We evaluated two well-known algorithms to detect and describe keypoints: Speeded-Up Robust Features (SURF) [7] and Scale-Invariant Feature Transform (SIFT) [2].

Both SIFT and SURF keypoint detection algorithms generate a few hundredkeypoints for each image. These usually occur in areas of high contrast, such as edges and corners. Then, for each keypoint, these algorithms calculate a descriptor that is designed to be invariant to scale, rotations, deformations, and other distortions. For each keypoint, SURF gives a vector of 64 values, whereas SIFT gives a vector of 128 values.

If we directly used the SIFT and SURF descriptors as features, the feature vector length would be variable, since these algorithms generatea variable number of keypoints. Todeal with this, we clustered the keypointsto create a vocabulary of visual words. For this step we used the most significant 20% of all the keypoints (based on the magnitudes of the descriptors) to find distinctive visual words. The number of clusters 'k' was empirically chosen to be 500, with each cluster corresponding to one visual word in our vocabulary. These two parameters are stable over a large range of values: varying the number of means (from 30 to 10,000) and the percentage of descriptors kept (from 1% to 80%) showed no significant differences in accuracy.

Finally, we calculated a bag-of-words feature vector for each image in both the training set and the test set by counting the number of occurrences of each visual word. This was done by finding all keypoints in each image and mapping the

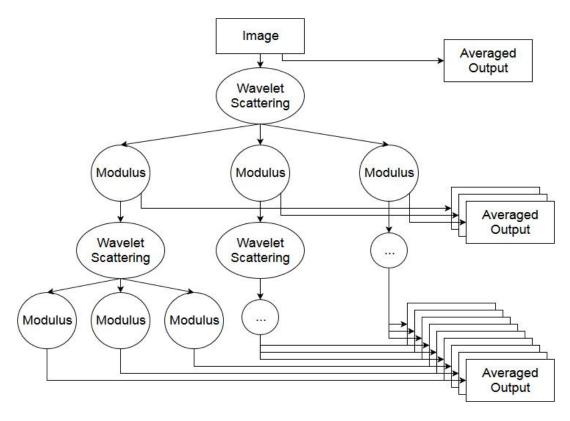


Fig. 2. ScatNet process. The output is the concatenation of all averaged moduli.

descriptors to the closest mean. Each descriptor counts as an occurrence of the associated word. Note that all descriptors are used at this stage, unlike at the clustering stage. The bag-of-words features are then used to train an SVM classifier to predict the class of the image.

Since the vanilla SIFT and SURF operate only on grayscale images, we tried two methods to improve accuracy by using color information. First, we tried decomposing the color image into three layers based on the HSV model of color, and we generated keypoints for all three layers, clustering and classifying as before. Second, we tried appending average hue and saturation to the feature vector for the grayscale version of each image. There was no significant difference in accuracy between these two methods.

## 5. ALTERNATIVE BASELINES

We also present human evaluation and a simple 2-feature system as alternative baselines. For the first, without formal instruction, we trained ourselves to identify epithelium and stroma by inspecting images in the training set. We then evaluated our ability to classify 50 randomly-selected images of the 100 in the test set.

While we are not experts at identifying stromata or epithelial cells, there are two reasons that this baseline is informative. First, we are training ourselves using exactly the same

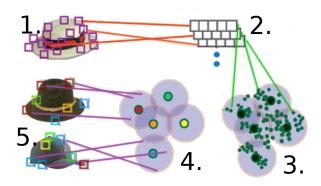
information that the computer has available to it: the images in the training set. Second, we suspect that the images are at best very hard to identify and at worst mislabeled. This baseline gives an impression of how good we should expect the best systems to perform.

Our second alternative baseline calculates two features for each image: average hue and average saturation. Then we train a gaussian mixture model (GMM) using expectation maximization. The resulting system performs surprisingly well, indicating that a great deal of information is contained within the color from the staining process.

## 6. RESULTS AND DISCUSSION

Results for each system can be found in **Table 2**. On the unambiguous 80x80 data set, near-perfect results are observed by all systems. (Occasionally, an "unlucky' training/test split would cause a single error in a trial.) For the more complicated non-uniform super-pixels, ScatNet is outperformed by all three baseline systems as well has human evaluation.

We are unsure about the exact reason for ScatNets poorer performance in the latter case. ScatNet is originally configured to use for uniformly sized rectangular images, which ours aren't, so perhaps the approximate normalization method used, needs more improvement. That ScatNet in general is suitable to the problem is evidenced by the high performance



**Fig. 3**. Visual depiction of the baseline system. Steps: 1. Find SIFT/SURF keypoints. 2. Extract descriptors. 3. Cluster with K-means. 4. Create visual words vocabulary. 5. Calculate bag-of-words feature vectors and classify with SVM.

on the uniformly sized 80x80 images. Another possibility is that, because we only ran each configuration once in the initial filtering, our hyper-parameter evaluation failed to identify the optimal set of parameters well. ScatNet is very sensitive to hyper-parameter values, so it is possible that a more optimal configuration would improve significantly.

On the other hand, our SIFT baseline performed better than we expected, outperforming even our human evaluation in the dataset with non-uniformly sized images. We think both the SIFT as well as the ScatNet systems be worth comparing more directly to the accuracy of a system with features manually designed by experts [1], which was evaluated on a different dataset.

Finally, it is worth mentioning that every system seems to improve from using HSV values over just the intensity values except for the SIFT baseline. Perhaps this is due to reaching the limit of what can be classified in this data set. The remaining images may simply be too mixed to be classified accurately by any system.

#### 7. FUTURE WORK

If the problem leading to poor performance with ScatNet is the normalization method, some of the following approaches could be adopted. One method could be to divide the input image into portions of a predetermined fixed size (80 x 80, say), run ScatNet independently on them and then generate a combined result based on the results for each portion. Another method could be to run ScatNet on a white image of the same size and shape (found by detecting the boundary of the input image), and then normalize based on the sum of all the feature values.

If on the other hand the problem is our hyper-parameter choices, we could do K-fold cross-validation instead of randomly generating the validation set for each case. To reduce the computational burden resulting from this, we could do a

Features + Algorithm	80x80	Gray-SP	Color-SP
SURF Descriptors + SVM	98.2	86.4	90.2
SIFT Descriptors + SVM	99.6	92.4	92.6
ScatNet + Affine PCA	99.9	75.0	87.2
Hue v. Saturation + GMM	98.5	_	91.0
Human Evaluation	100.0	84.0	89.0

**Table 2**. Accuracies of all tested systems. Best (non-buman) performers for each category are in bold.

coarse-exploration of the space, followed by a hill-climbing algorithm to find a (locally) optimal set of parameters.

Our method of handling color did help most of the systems tested, though not the accuracy of the SIFT baseline. Perhaps for this system it would be helpful to use a variant of SIFT that inherently takes color into account, such as the algorithms mentioned in [8] or [9]. It is possible that the techniques developed in these papers could also apply to our Scat-Net system.

One more difference between the ScatNet system and the baseline is the classifier used. It would be informative to try the PCA affine classifier on the baseline system, and the SVM classifier on the ScatNet system to see what kind of impact the classifier has on the result.

Since we found that the dataset had many images that were ambiguous even to humans, it may be helpful to either add a third category representing a mixture and re-label the confusing images, or to add some sort of score to represent how mixed the image is, and change this into a regression problem rather than a classification problem.

#### 8. REFERENCES

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