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# **PyHistopathology**

***Release 1.0***

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PyHistopathology is a Python API for pre-processing Whole-Slide Images for use in a machine learning algorithm.



# CHAPTER 1

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## Features

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Reading Whole-Slide Images with `WSI_Scanning.readWSI()`

- Input: WSI path or directory
- Output: Numpy array of WSI with data type int32

Denoising Whole-Slide Images with `Denoising.denoising()`

- Input: WSI Path or directory
- Output: Numpy array of WSI Image (After denoising) with dtype int32

Patch Extraction for Whole-Slide Images with `Extractingpatches.extractingPatches()`

- Input: WSI Path or directory
- Output: Fills up the outputpath with patches extracted from the WSI



# CHAPTER 2

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## Table of Contents

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### 2.1 Reading

Use WSI\_Scanning.readWSI() to read Whole-Slide Images

### 2.2 Usage

```
readWSI(WSI_path, magnification_level, annotation_file, annotated_level)
```

### 2.3 Arguments

#### 2.3.1 WSI\_path

Path or location of WSI.

#### 2.3.2 magnification\_level

Level of zoom, for example 40, 20, 10, or 5. Default magnification level is 20. - Note: if magnification 40x for max zoom level of 20x image an error will be raised.

#### 2.3.3 annotation\_file

Default annotation = None. If annotation are available in a xml file, set annotation\_file to be the xml file path.

### 2.3.4 annotated\_level

If annotation is given then set annotated\_level equal to the z-axis of the annotations. Default annotatedlevel is 0.

## 2.4 Return Type

Numpy array of WSI Image (After denoising) with dtype int32

## 2.5 Example

```
from WSI_Preprocessing.Preprocessing import WSI_Scanning
import cv2
img,slide_dim = WSI_Scanning.readWSI("example.svs")
cv2.imwrite("example.png",img)
```

## 2.6 Denoising

Use Denoising.denoising() to remove stains, folds and other background noise in Whole-Slide Images

## 2.7 Usage

```
denoising(inputsvs, magnification, filtering, patch_size, upperlimit, lowerlimit, red_value, green_value, blue_value)
```

## 2.8 Arguments

### 2.8.1 inputsvs

Path or location of WSI.

### 2.8.2 magnification

Level of zoom, for example 40, 20, 10, or 5. Default magnification level is 20. - Note: if magnification 40x for max zoom level of 20x image an error will be raised.

### 2.8.3 filtering

GaussianBlur, RGBThersholding, or None

GaussianBlur: Homogeneity calculations based on image smoothing and Gaussian blur equations. We compute sum of square differences between two consecutive Gaussian blurred images as score for homogeneity.

- Upper limit: Upper threshold of homogeneity score. Default value is 9500 with kernel size of 1111
- Lower limit: lower threshold of homogeneity score. default value is 1500 with kernel size of 1111

- Patch size: Not significant parameters for GaussianBlur filtering

RGBThersholding: Validated patches based on RGB values of patches

- red\_value: Red threshold
- green\_value: Green threshold
- blue\_value: Blue Threshold

None: Only removes Background

Note that our default is GaussianBlur technique. GaussianBlur is highly effective and requires more computational power (RAM). RGBThersholding is less effective which needs less computational power

## 2.9 Return Type

Numpy array of WSI Image (After denoising) with dtype int32

## 2.10 Extracting

Use Extractingpatches.extractingPatches() to extract patches from Whole-Slide Images

## 2.11 Usage

```
extractingPatches(inputsvs, outputpath, magnification, patch_extraction_creatia, number_of_patches, filtering,
patch_size, upperlimit, lowerlimit, red_value, green_value, blue_value, Annotation, Annotationlevel, Requiredlevel,
reconstructionimagepath)
```

## 2.12 Arguments

### 2.12.1 inputsvs

Path or location of WSI.

### 2.12.2 magnification

Level of zoom, for example 40, 20, 10, or 5. Default magnification level is 20. - Note: if magnification 40x for max zoom level of 20x image an error will be raised.

### 2.12.3 filtering

GaussianBlur, RGBThersholding, or None

GaussianBlur: Homogeneity calculations based on image smoothing and Gaussian blur equations. We compute sum of square differences between two consecutive Gaussian blurred images as score for homogeneity.

- Upper limit: Upper threshold of homogeneity score. Default value is 9500 with kernel size of 1111
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- Patch size: Not significant parameters for GaussianBlur filtering

RGBThersholding: Validated patches based on RGB values of patches

- red\_value: Red threshold
- green-value: Green threshold
- blue\_value: Blue Threshold

None: Only removes Background

Note that our default is GaussianBlur technique. GaussianBlur is highly effective and requires more computational power (RAM). RGBThersholding is less effective which needs less computational power

#### **2.12.4 patch\_extraction\_creatia**

random, or None. Default is None. For extracting a fixed number of patches for WSI we can use random.

#### **2.12.5 number\_of\_patches**

Default number of patches is 2000

#### **2.12.6 outputpath**

Folder to store the extracted patches

#### **2.12.7 reconstructionimagepath**

If you want to compare the patches with WSI we can mention the reconstructionimagepath. Default is None. Note: it only works with patch\_extraction\_creatia = None.

### **2.13 Return Type**

None, fills up output path with images directly instead of returning a Numpy array.