Optimal experimental hyperspectral image acquisition conditions of biomedical structures

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Abstract—In recent decades there has been a great interest in using hyperspectral imaging (HSI) for biomedical applications. HSI provides a three dimensional dataset from which one can obtain both the spatial and spectral information of biological samples being imaged. From the acquired hyperspectral images, it is possible to extract diagnostic information about tissue constituents and morphology. This paper focuses on determining the experimental conditions for an optimal hyperspectral acquisition of biological samples.

Index Terms—Biomedical applications, diagnostic information, hyperspectral imaging, spectral curves, spatial resolution, tissue composition.

1 INTRODUCTION

Spectral imaging is imaging in several wavelength bands. The common RGB camera makes use of a filter that separates the light into three wavelengths- red, green, and blue. In multispectral imaging, the number of wavelength bands is about ten while in hyperspectral imaging, the number of distinct bands could be as high as several hundreds. Apart from a high number of wavelength bands, the hyperspectral camera can detect a higher wavelength range (200nm – 2500nm) and offers a wide range of applications in remote sensing, archaeology and art conservation, [1, 2] vegetation and water resource control [3,4], food quality and safety control [5,6] forensic medicine, [7,8] crime scene detection, [9,10] biomedicine [11,12] etc.

For biomedical applications, hyperspectral imaging shows great potential especially in the areas of disease diagnosis. The biological and pathological changes in tissues and organs have a close relationship with the spectra [13]. Thus different pathological conditions can be identified from the spectral information of each pixel in the hyperspectral images.

A hyperspectral imager mainly consists of a light source, a dispersive element (prism, grating, etc) and a 2D detector array. Such imagers use the 'push-broom' method to obtain hyperspectral data of a scene. This method captures a single line of spatial information containing full-spectrum data for every spatial pixel in the line [14]. In the basic design of a hyperspectral camera (Fig. 1), the light from the tissue sample passes through the entrance aperture of the camera and some other aberration correcting elements.

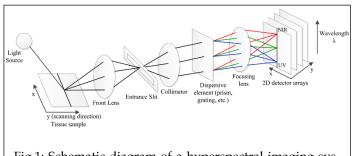
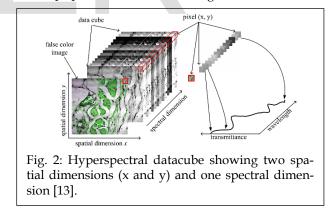


Fig.1: Schematic diagram of a hyperspectral imaging system [15].

The incoming light beam is then focused onto a slit by a fo-

cusing mirror which transforms it into a collimated beam and then focuses on a dispersive element (grating) that separates the beam into its different wavelength components. The dispersed light beam emerging from the grating is now passed through the lens optics system with the effect that for each pixel interval along the line defined by the slit (spatial axis), a corresponding spectrum is projected on a column of detectors on the array. The data read out from the array thus contains a slice of a hyperspectral image, with spectral information in one direction and spatial (image) information in the other. By scanning over the tissue sample, the camera collects slice from adjacent lines, forming a three dimensional data set called "hypercube" with two spatial dimensions and one spectral dimension [15]. This is illustrated in Fig. 2.



The main focus of this paper is to determine the conditions and specifications of the imaging system needed for an optimal hyperspectral acquisition of biological samples. For this purpose, a VNIR scientific CMOS camera and a photospectrometer has been used together with a transmissive light microscope to acquire hyperspectral images. The acquired hyperspectral images using varying amount of the intensity of the incident light, at different wavelength bands have been compared to that seen with the microscope eye-piece and also with that taken by another CCD camera. This allowed the determination of the amount of incident light needed to optimize the hyperspectral image acquisition process. Furthermore, the field of view, magnification, spatial and spectral resolutions of the hyperspectral imaging system have also been computed.

2 MATERIALSAND METHODS

The experimental setup is shown in Fig. 3. The system consists of a photo-spectrometer SpecimV10 (SpecIm Ltd) connected to a high sensitivity sCMOS camera from PCO. The overall system is operative in the 400-1000 nm spectral range, with a spectral resolution of about 0.89nm. The hyperspectral camera system is coupled to an optical microscope, Nikon Eclipse TE 2000-U (see technical details in Fig. 4). The HSI system uses the Camware software for acquisition and visualisation of the images. The camera can also be controlled using this software. The user-friendly graphical interface can provide images at different chosen wavelengths and spectral graphs can be obtained for a particular selected position on the images.

In order to calculate the field of view of the hyperspectral camera system, a grating with known dimensions is imaged using the optical microscope coupled with the HSI system.

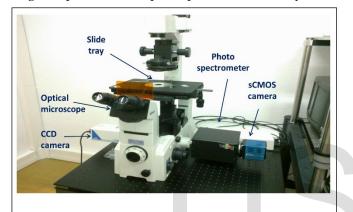


Fig. 3: Experimental setup of the hyperspectral imaging system.

Camera : pco.edge sCMOS		Spectrometer: SPECIM V10		Optical Microscope: Nikon Eclipse TE 2000-U	
	2560 x 2160 pixel	Spectral region	400-1000nm	Numerical aperture	1.05
Pixel size	6.5µm x 6.5 µm	F-number	F/ 2.8	Objective magnification	10 X
Full well Scapacity	30 000e-	Spectral sampling	0.63 – 5.0nm per pixel	Eye piece lens	10X (FOV 22mm)
Readout noise	<1.3rms	Spectral resolution	6.8nm	Tube magnification	1 or 1.5
	27000: 1 (88.6dB)	Spatial resolution	< 40 µm	Microscope configuration	Transmissive
Quantum : efficiency	>54% @peak	Slit width	30µm x 14 µm		
	370 nm – 1100nm	Lens mount	C-mount		
Frame rate	100fps				

3 RESULTS AND DISCUSSION

3.1 Evaluation of the field of view and spatial resolution of the HSI system

The field of view has been evaluated by imaging a grating with known dimensions. On the grating, 100 lines equals to 10 mm. Therefore, for 1 line corresponds to 0.1mm.Number of vertical lines observed on the computer screen = 15 lines. Number of horizontal lines observed on the computer screen = 3 lines.

Estimated field of view = 1.5mm on the horizontal axis and 0.3mm on the vertical axis. Effective linear magnification of the camera = (size of the sensor used)/(field of view) = $(2560 \times 6.5 \text{um})/1.5 \text{mm} = 11$.

It has been mentioned before that the hyperspectral imaging system works as a scanner taking a slice of the image at a time. The different wavelength components of each slice are dispersed on the pixels along one of its axis. Thus, for each slice of the observed object, a 2D image is obtained with one spatial and one spectral axis. By taking the entire FOV, a 3D hypercube is formed with two of the dimensions being the spatial axis and the last being the spectral axis. The number of slices is fixed by setting the number of frames to capture on the hyperspectral camera. The maximum number of frames is 1000 which corresponds to the entire field of view along the scanning direction. Using this, the time to capture an entire field of view is calculated to be,

Capture time = number of frames \div frame rate = 500 \div 100fps = 5s.

The speed of the micro-translator needed to translate the specimen= (field of view in the scanning axis)/(capture time) = 1.5mm/5s = 0.3 mm/s.

Spatial resolution = (field of view in the scanning axis)/ (number of pixels) = 1.5mm/2560 = 0.6µm.

It is noted that the resolution of the image is not diffraction limited but limited by the size of each pixel.For a diffraction limited system, the resolution should be,

 $R_{diffraction} = 0.61 \lambda / NA = 0.32 \mu m$ (assuming an incident light of wavelength 550nm).

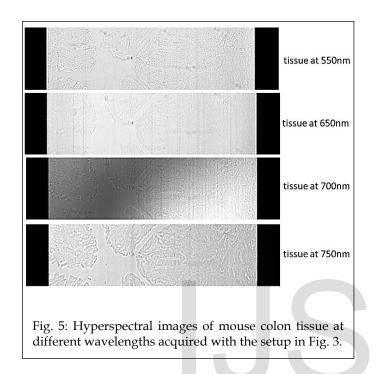
The size of the diffraction resolvable unit projected on the sensors is about 3mm, using a magnification of 10. For the image seen on the sensor to be diffraction limited, using the Nyquist sampling theorem, the pixel size should be half of the resolvable unit. This corresponds to a pixel size of 1.5mm.

The spectral resolution is equal to the amount of dispersion for each pixel. This as seen on the technical documentation of the hyperspectral camera is equal to 0.89 nm.

3.2 Hyperspectral imaging of a mouse colon tissue:

The observed images of a biological tissue (colon slice of a mouse) at different wavelength bands acquired with the hyperspectral imaging camera are shown in Fig 5. The spectral curve shown in Fig.6(b) is obtained for different positions on the tissueas shown in Fig.6(a).

From the acquisitions of the tissue samples, it has been observed that at different wavelengths some features of a sample become easier to observe. Also due to the large amount of spectral bands, a careful selection has to be made of bands which contained the most useful information about the specimen. Doing this in practice often depends on what application the scientist is interested in, for example an unusual cell in the midst of many identical cells. Image processing softwares are used to perform this selection based on different hyperspectral image processing methods [16,17].

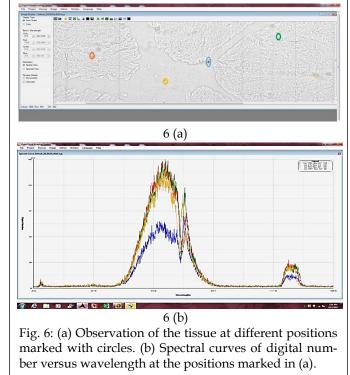


The graph of the digital number against the wavelength in Fig. 6b) shows that for positions on the image (Fig 6a)) with similar constituents the spectral curves does not vary much compared to positions with a different constituent. This means that the hyperspectral imaging data can be used to characterize different constituents of the specimen being imaged.

From the acquisition of hyperspectral images, it has also been observed that the intensity of the incident light source is required to be increased compared to the level needed for conventional microscopy with a CCD camera. This is because in case of hyperspectral imaging, the light scattered by the object reaching the sensors is dispersed across several pixels in the spectral direction. This reduces the amount of light detected at each pixel location and explains the need for more sensitive cameras (e.gsCMOS or EMCCDs) for hyperspectral imaging.

4 CONCLUSION

The aim of this work has been to understand and determine the conditions for acquiring good hyperspectral images of biological samples. For this purpose, a VNIR sCMOS camera together with a photo-spectrometer has been used to form a hyperspectral imaging system. The field of view and the spatial resolution of the HSI system are then evaluated by using gratings of known dimensions which are imaged by coupling the HSI system with an optical microscope.



Then hyperspectral images of mouse colon tissues are acquired. From the images obtained it has been observed that certain features of the tissue samples are visible only at certain wavelength bands. However, due to the high number of spectral information available from an HSI system, the wavelength bands have to be selected carefully depending on the intended application. Spectral graphs of selected positions on the acquired images are then obtained. The regions with similar constituents showed no change in the spectral curves compared to those with a different constituent. Therefore, specimen constituents can be characterized by comparing the spectral curves at different locations. Thus, hyperspectral imaging can be used in obtaining information about tissue composition and shows much potential to be used in many other biomedical applications.

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