

Three-dimensional image of hepatocellular carcinoma under confocal laser scanning microscope

Wang Hai Zhang¹, Shi Neng Zhu², Shi Lun Lu², Ya Lin Huang³ and Peng Zhao³

Subject headings HCC; nucleus, three-dimensional reconstruction; microscopy, confocal laser scanning

Zhang WH, Zhu SN, Lu SL, Huang YL, Zhao P. Three-dimensional image of hepatocellular carcinoma under confocal laser scanning microscope. *World J Gastroentero*, 2000;6(3):344-347

Abstract

AIM To investigate the application of confocal laser scanning microscopy (CLSM) in tumor pathology and three-dimensional (3-D) reconstruction by CLSM in pathologic specimens of hepatocellular carcinoma (HCC).

METHODS The 30 μm thick sections were cut from the paraffin-embedded tissues of HCC, hyperplasia and normal liver, stained with DNA fluorescent probe YOYO-1 iodide and examined by CLSM to collect optical sections of nuclei and 3-D images reconstructed.

RESULTS HCC displayed chaotic arrangement of carcinoma cell nuclei, marked pleomorphism, indented and irregular nuclear surface, and irregular and coarse chromatin texture.

CONCLUSION The serial optical tomograms of CLSM can be used to create 3-D reconstruction of cancer cell nuclei. Such 3-D impressions might be helpful or even essential in making an accurate diagnosis.

INTRODUCTION

Under conventional light microscope, *histopathologists* often use plane image to evaluate the three-dimensional (3-D) cellular characteristics. Three-D configuration may be reconstructed by using serial mechanical sectioning, but its axis definition is not good, image is blur and tiny structure can't be shown clearly, and the specimen might be damaged. Using the serial optical tomograms and 3-D reconstruction function, confocal laser scanning microscopy (CLSM) can provide a much better quality 3-D image than conventional light microscope, and lead the observer into a brand-new 3-D world. Although CLSM has been used extensively in cell biology^[1], few applications were reported in routine clinical pathology such as three-dimensional DNA image cytometry by CLSM in thick tissue blocks of prostatic lesions and 3-D reconstruction by CLSM in routine pathologic specimens of benign and malignant lesions of human breast^[2-4]. In this study, 3-D reconstruction was performed on routine formalin-fixed, paraffin-embedded tissues of normal, and hyperplastic tissues of liver and hepatocellular carcinoma by using computer-assisted CLSM together with 3-D reconstruction. The goals of our study were to present 3-D morphologic characteristics of benign and malignant specimens of the liver and to attempt to demonstrate the usefulness of CLSM in routinely obtained surgical pathologic tissues.

MATERIALS AND METHODS

Two cases of normal liver tissue were selected from autopsy specimens collected in the Department of Pathology of Shanghai Medical University, six cases of hepatocellular carcinoma including adjacent liver tissue were routine clinical specimens collected in 1996 from the Liver Cancer Institute of Zhongshan Hospital affiliated to Shanghai Medical University.

All tissues were fixed in 10% formalin, embedded in paraffin, serial sections were cut at 5 μm and 30 μm . The 5 μm slices were stained with hematoxylin and eosin for conventional light microscopic observation. The 30 μm slices were stained with DNA fluorescent probe, YOYO-1 iodide (Molecular Probes, Eugene, Ore., USA).

¹Division of Pathology, Zhongshan Hospital, Shanghai Medical University, Shanghai 200032, China

²Department of Pathology, Shanghai Medical University, Shanghai 200032, China

³National Laboratory of Medical Neurobiology, Shanghai Medical University, Shanghai 200032, China

Dr. Wang Hai Zhang, graduated from Shanghai Medical University as a Ph.D. in 1999, majoring in surgical pathology and molecular pathology, having 10 papers published.

Correspondence to: Dr. Wang Hai Zhang, Division of Pathology, Zhongshan Hospital, Shanghai Medical University, Shanghai 200032, China

Tel. 0086-21-64041990 Ext.2732

Email. whzhangsh@hotmail.com

Received 2000-03-19 **Accepted** 2000-04-28

The sections were deparaffinized with xylene (10 min \times 2) and dehydrated with 100%, 95%, and 70% ethanol (5 min \times 2) and rinsed in distilled water for 2 min \times 5. The specimens were then fixed with 10% neutral buffered formalin for 30 min and washed with tap water. After rinsing with distilled water and 0.01M phosphate buffer 5 min \times 2, nuclear RNA was removed by incubating the sections for 30 min at 37°C in 200 μ L of ribonuclease A (RNAase; Sigma, USA) at a concentration of 160 g/L in PBS. DNA was next hydrolyzed with 2N HCl for 25 min at 27.5°C. After rinsing with distilled water for 2 min \times 5, the sections were covered with 200 μ L of YOYO-1 iodide diluted into 1:2000 with PBS. The PBS was diluted 1:5 with distilled water to reduce the salt concentration. To this 200 μ L working solution of YOYO-1 iodide, 20 μ L of 0.1N HCl was added and the final solution was stored in the dark at 4°C for use. Homogeneous fluorescence intensity of nuclei at different depths of the confocal slices was obtained by agitating the YOYO-1 iodide for 1 h in the dark. Afterwards, the sections were rinsed with distilled water, covered with buffered glycerol, and the glass cover slip were sealed with finger nail polish. Sections were stored at 4°C in the dark until CLSM examination.

A Leica TCS-NT confocal laser scanning microscope equipped with epifluorescence optics and an appropriate combination of filters to visualize and digitize the images of the different specimens. An argon laser with an excitation wavelength of 488 nm was used to activate the green fluorescence of the YOYO-1 iodide-stained nuclear DNA (maximal absorption 491 nm and emission 509 nm). A 16 \times objective (numerical aperture of 1.30) was used to observe the specificity of the staining, and a 100 \times water objective (numerical aperture of 1.30) was used to study the details of chromatin pattern. Optical sections were collected throughout the entire stained thickness of the paraffin sections with a Z-step interval of 0.3 μ m or 0.6 μ m. The 3-D image processing was performed on a Leica computer with the original 3-D interactive visualization software. The collected confocal optical sections and the 3-D reconstructed images were printed with a Panasonic color video dye-sublimation copy processor.

RESULTS

Three-D morphologic features of normal liver cells

A microscopic field of view was selected from the normal liver specimen and comparison of the images taken under transmission conventional light microscope and CLSM is illustrated in Figure 1. The transmission conventional light microscopic image

was shown on the right side and the out-of-focus signals were visible. The out-of-focus blur reduced the contrast and sharpness of the final image. In the confocal image (left) the out-of-focus signals were cut off and only signals in focus were clearly visible. Optical sections (planes 1-60) were taken from the surface to the bottom of normal liver specimen with a Z-step interval of 0.5 μ m. Figure 2 shows the 10th, 20th, 30th, 40th, 50th and 60th plane digital images of Z-series. The confocal images were taken at 5, 10, 15, 20, 25 and 30 μ m depths, respectively. In the confocal images some nuclei appeared or disappeared depending on their orientation in space. The 60 2-D optical sections were computer focused on a plane (deep-focusing) to analyze the fine structure of chromatin patterns inside the nucleus and reconstructed 3-D images to display the 3-D detailed arrangement of nucleus. Figure 3 shows normal liver cells with similar round or ovoid nuclei, homogeneous intensity of YOYO-1 iodide fluorescence as well. Three-dimensional view was shown in Figure 4. The nuclear surface appeared smooth with homogeneous fluorescence intensity.

Three-D morphologic features of atypical hyperplasia of liver cells

Optical sections (planes 1-50) were taken from the surface to the bottom of atypical hyperplasia of liver specimen with a Z-step interval of 0.6 μ m. The structure of chromatin patterns inside the nucleus with homogeneous fluorescence intensity is shown with deep-focusing in Figure 5. Three-dimensional view is shown in Figure 6. The nuclear surface appeared smooth with homogeneous fluorescence intensity. But the volume of nucleus of atypical hyperplasia liver cell was bigger than that of normal liver cell.

Three-D morphologic features of hepatocellular carcinoma cells

Optical sections (planes 1-50) were taken from the surface to the bottom of hepatocellular carcinoma specimen with a Z-step interval of 0.6 μ m. The deep-focusing images were shown in Figure 7 and Figure 8. The structure of chromatin patterns inside the nucleus with heterogeneous karyotheca thickness, irregular and coarse chromatin texture, chromatin underside the karyotheca mainly. Three-dimensional view is shown in Figure 9 and Figure 10. HCC displayed remarkably different features in 3-D morphology, including: indented, molding, and irregular nuclear surface; marked pleomorphism; chaotic arrangement of tumor cell nuclei.

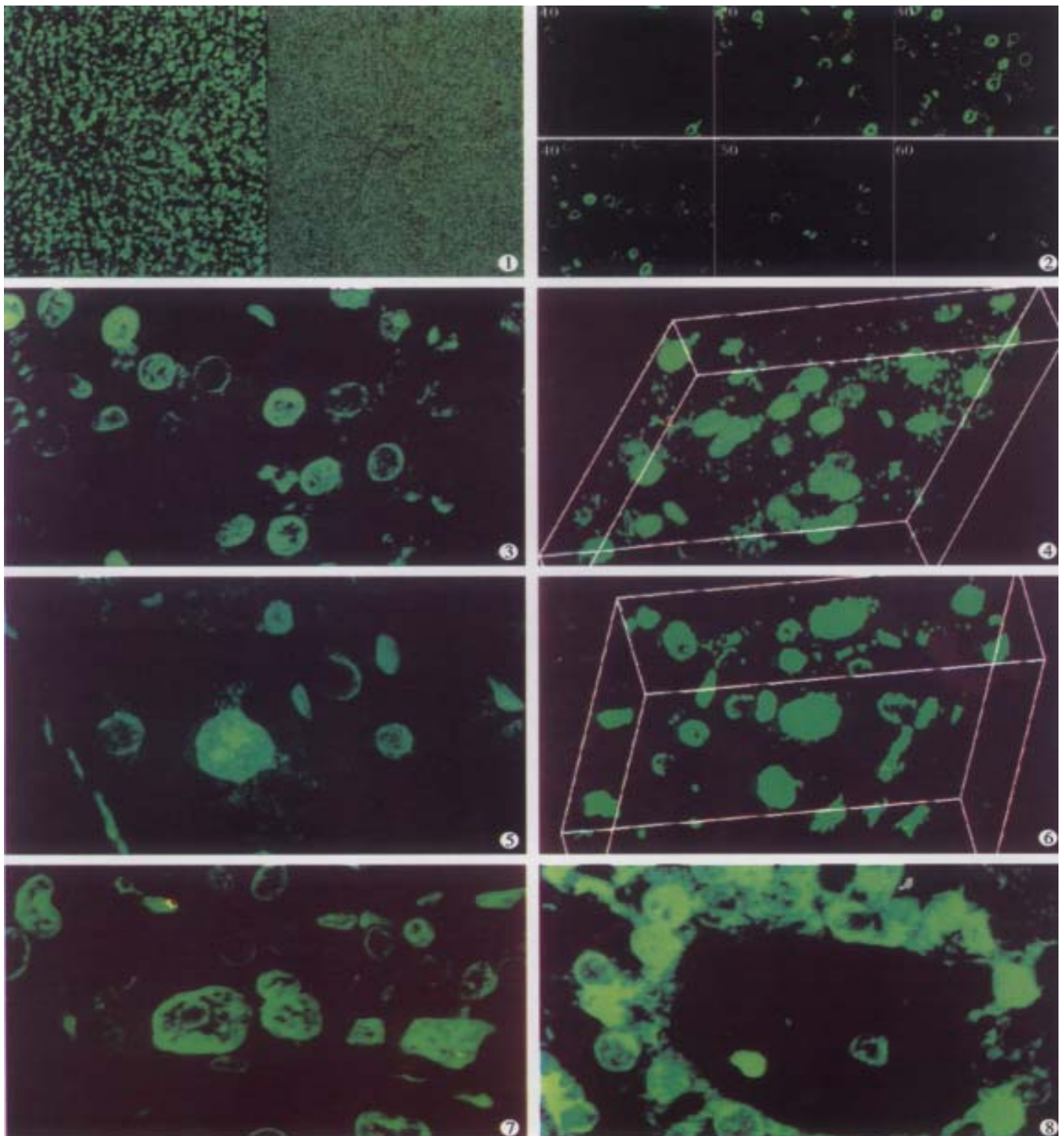


Figure 1 The transmission conventional light microscopical image was shown on the right side. The out-of-focus blur reduced the contrast and sharpness of the image. In the confocal image (left) the out-of-focus signals were cut off and only signals in focus were clearly visible. $\times 160$

Figure 2 The 10th, 20th, 30th, 40th, 50th and 60th plane digital images of Z-series. In the normal liver tissues of the confocal images some nuclei appeared or disappeared depending on their orientation in space. $\times 1000$

Figure 3 Deep-focusing image showed the normal liver cells with similar round or ovoid nuclei, similar in size and homogeneous intensity of YOYO-1 iodide fluorescence as well. $\times 1000$

Figure 4 Three-dimensional view of normal liver cells, the nuclear surface appeared smooth with homogeneous fluorescence intensity. $\times 1000$

Figure 5 Deep-focusing image of atypical hyperplasia liver cells, the structure of chromatin patterns inside the nucleus with homogeneous fluorescence intensity. $\times 1000$

Figure 6 Three-dimensional view of atypical hyperplasia liver cells. $\times 1000$

Figure 7 Deep-focusing image of HCC cells, the structure of chromatin patterns inside the nucleus with heterogeneous karyotheca thickness, irregular and coarse chromatin texture, chromatin underside the karyotheca mainly. $\times 1000$

Figure 8 Deep-focusing image of highly differentiated HCC chromatin texture. $\times 1000$

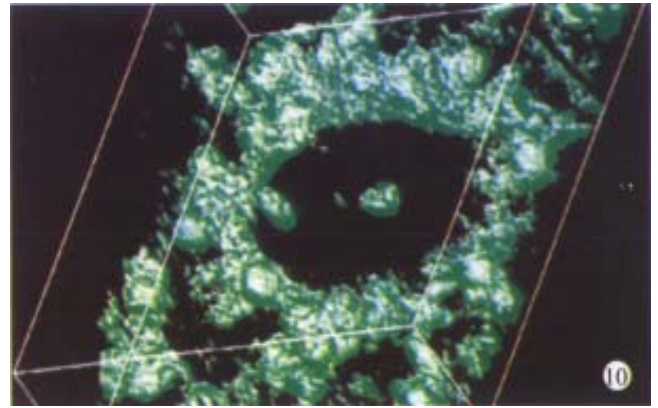
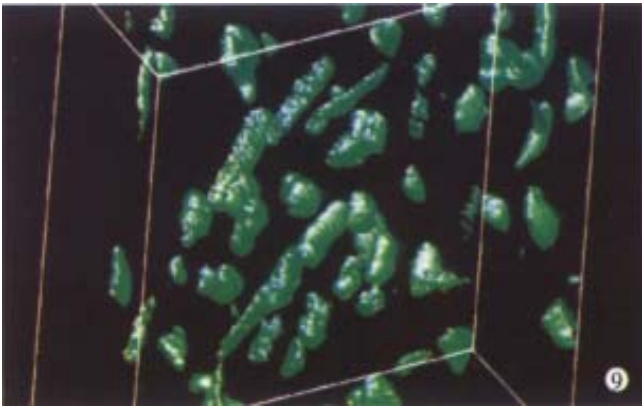


Figure 9 Three-dimensional view of spindle cell HCC. $\times 1000$

Figure 10 Three-dimensional view of tubular adenocarcinoma of HCC. $\times 1000$

DISCUSSION

The area, volume, shape, DNA content, and chromatin pattern of nuclei may be important for the diagnosis and prognosis of cancer. *Histopathologists* often use $4\text{ }\mu\text{m}$ to $6\text{ }\mu\text{m}$ thick paraffin sections to obtain representative and diagnostically relevant images. Due to the very limited section thickness in comparison with the size of the tissue, and the images are nearly two-dimensional, focusing up and down at high magnification provides a rough idea of the 3-D cellular characteristics, and such 3-D impression may be helpful or even essential in arriving at a certain diagnosis, especially for borderline lesions or tumors. However, in spite of the usefulness of such 3-D information about nuclei, conventional light microscopy is not always the ideal tool for 3D evaluation due to the interference of out-of-focus structures with the images of the focus plane studied. Much of the light emitted from the regions of specimen above and below the focal plane contributes to the out-of-focus blur, which seriously reduces the contrast and sharpness of the final image. Confocal laser scanning microscope allows the acquisition of optical sections from a thick specimen and out-of-focus blur can be reduced considerably and thus much sharper and clearer images will be obtained. CLSM has become an exciting new instrument in biomedical research because of its increased resolution over conventional light microscope and its utility for subsequent 3-D-reconstruction analysis^[5,6].

In this paper, the 3-D reconstruction have demonstrated 3-D contour of representative characteristics of normal liver cells, atypical hyperplasia liver cells, and HCC cells and the spatial relationship of nuclei, as well as the subtle structure of chromatin texture inside nuclei. This

paper emphasized the practical feasibility of CLSM and 3-D reconstruction from routine surgical histopathologic materials. To obtain desirable quality 3-D image from formalin-fixed, paraffin-embedded specimens, YOYO-1 iodide, a highly specific and sensitive (picogram sensitivity)^[7] DNA probe was utilized. Intense and homogeneous fluorescence was obtained by incubating the YOYO-1 iodide for 1h with agitation in the dark. To reveal subtle details of nuclear structure, RNA was removed by RNase predigestion, since nuclear RNA was stained by YOYO-1 iodide as well.

CLSM combines the three most advanced and important elements of our era, the microscope, the laser and the computer in one, moreover it is non-invasive and can be used on archival paraffin blocks. We anticipate that, in the future, pathologists may utilize these new techniques to make more precise diagnosis.

REFERENCES

- 1 Minsky M. Memoir on inventing the confocal scanning microscope. *Scanning*, 1988;10:128-138
- 2 Boon ME, Schut JJ, Suurmeijer AJ, Benita EM, Hut PK, Kok LP. Confocal microscopy of false-negative breast aspirates. *Diagn Cytopathol*, 1995;12:42-48
- 3 Liu S, Weaver DL, Taatjes DJ. Three-dimensional reconstruction by confocal laser scanning microscopy in routine pathologic specimens of benign and malignant lesions of the human breast. *Histochem Cell Biol*, 1997;107:267-278
- 4 Beil M, Irinopoulou T, Vassy J, Rigaut JP. Chromatin texture analysis in three-dimensional images from confocal scanning laser microscopy. *Anal Quant Cytol Histol*, 1995;17:323-331
- 5 White JG, Amos WB, Fordham M. An evaluation of confocal versus conventional imaging of biological structures by fluorescence light microscopy. *J Cell Biol*, 1987;105:41-48
- 6 Shotton DM. Electronic light microscopy: present capabilities and future prospects. *Histochem Cell Biol*, 1995;104:97-137
- 7 Tekola P, Baak JP, Belien JA, Brugghe J. Highly sensitive, specific, and stable new fluorescent DNA stains for confocal laser microscopy and image processing of normal paraffin sections. *Cytometry*, 1994;17:191-195