Laser Applications for Estimation of Microbial Activity and Investigation of NIR Skin Autofluorescence

Alexey Lihachev1, Dmitrijs Bļizņuks², Jānis Liepiņš3, and Ilze Lihacova1

¹ University of Latvia, Institute of Atomic Physics and Spectroscopy, Biophotonics Laboratory, 3 Jelgavas str., Riga LV-1004, Latvia

2 Riga Technical University, Faculty of Computer Science and Information Technology, 10 Zunda krastmala, Riga LV-1050, Latvia

3 University of Latvia, Institute of Microbiology and Biotechnology, 1 Jelgavas str., Riga LV-1004, Latvia

*** e-mail:** aleksejs.lihacovs@lu.lv

Abstract. The presented paper demonstrates the ongoing results of the joint research of University of Latvia and Riga Technical University aimed at the development of new laser based optical methods and technologies for fast evaluation of microbial activity in solid media. The second part of the paper represents the results of joint research conducted together with Katya Borisova during our visit to the Institute of Electronics, Bulgarian Academy of Sciences (Bulgaria, Sofia). © 2021 Journal of Biomedical Photonics & Engineering.

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1 Introduction

1.1 Laser speckle imaging for microorganisms' growth analysis in solid media

Enumeration of live microbial cells in the sample is one of the basic quantitative measurements in microbiology. Currently counting microorganisms in the early stages of colony formation is possible only by microscope, which is limited to small sample areas. Also, determination of colony size increase dynamics (microorganism growth speed estimation) is an extremely labor-intensive task. Automated counting is possible only by using regular cameras that are limited in optical resolution and therefore are capable of detecting signs of microbial colonies only after a certain size threshold is reached (minimal diameter 0.2 mm and / or 8–22 h). Therefore, the development of new cost-effective methods for *in situ* estimation of microorganisms' activity to reduce detection time and labor expenses raise a great practical interest for the scientists and technology developers. One of the options for the fast and early evaluation of microorganisms' activity is a non-contact optical technique called laser speckle contrast imaging. Laser speckle is an interference pattern produced by coherent light reflected or scattered from different parts of the illuminated surface. In the case of non-moving, static scatters, the scattered light generates

constant laser speckle patterns. However, if scatterers are spontaneously moving (e.g., Brownian motion, bacteria doubling and chemotaxis) such as particles in a fluid, the individual speckle appears to be "twinkle" or "boiling". This phenomenon is called a "time-varying speckle". Regarding this phenomenon, the laser speckle imaging technique has been proved to monitor moving particles in optically inhomogeneous media by analyzing timevarying laser speckle patterns. Thereby, the analysis of laser speckles patterns provides a powerful opportunity for quantifying the movement activity at the micro and macroscale. Literature data points to the significant progress and potential applications of laser speckle imaging technique to evaluate the dynamic processes in microbiological media. Technology using speckle decorrelation time maps has been demonstrated for the detection of *E. coli* and *B. cereus* on a meat (chicken breast) [1]. Speckle analysis has also been applied to biomass growth kinetic measurements in liquid culture, characterization of Colony Forming Units (CFU) morphology, determination of antibiotic susceptibility [2–5]. As well as laser speckle imaging techniques in combination with Deep Learning (DL) and Artificial Neural Networks (ANN) have demonstrated a fast system response of antibacterial susceptibility evaluation in MIC (Minimum Inhibitory Concentration) tests [6]. In the presented paper, we will prove that dynamic laser speckle

image analysis methods can be used to detect and predict the CFU of different microbes in solid media.

1.2 Investigation of Near Infrared Autofluorescence under continuous laser excitation

The phenomenon of autofluorescence in the near infrared region (NIR AF) under continuous laser excitation was described by many authors. The first mention of NIR AF was described by Z. Huang et al. [8] where the NIR AF was analyzed during Raman spectroscopy investigations under 785 nm excitation. Authors demonstrate promising sensitivity of NIR AF intensity analysis for evaluation of pigmented lesions, as well as, demonstrated applications for melanoma diagnostics. The physical explanation of the NIR AF and its origin are still under discussion, mainly it can be described by absence of absorption spectra of main tissue fluorophores in the NIR region. Several hypotheses describing this phenomenon are discussed in literature, mainly attributing NIR AF to the melanin, endogenous porphyrins and lipofuscin [9–11]. Analysis and detection of NIR AF is a challenging task, since the NIR AF intensity usually is very weak and requires a high sensitivity detecting system equipped with high >8 OD filters. The artefacts caused by reflected laser light significantly complicate the adequate detection of the NIR AF from living tissues. In presented research we have tried to investigate the phenomenon of NIR AF. The experiments were conducted in close collaboration with Ekaterina Borisova during our secondment to the Institute of Electronics, Bulgarian Academy of Sciences. The obtained results and discussion are described below.

2 Material and methods

2.1 Laser speckle imaging system

The system was assembled for burst capturing of macro scale images under white light and laser illumination. The optical measurement system consists of a laser source, white light source LED, 35mm CS lens @F18, optical attenuator, a testing agar plate (with inoculated bacteria) and a CMOS camera (Fig. 1). The laser speckles were generated by a linearly polarized 635 nm diode pumped solid state laser (output power 50 mW). To achieve optimal exposure for image capturing as well as to avoid heating effects of illuminated plate the optical attenuator was used enabling 3–5 mW/cm2 power density of the scattered laser light on the whole agar plate surface. Main components of the system are presented in Fig. 1. The speckle images with 10 MPix resolution were captured by a CMOS camera with different frame rates, for different experiments: 1 frame per 2, 5, 6, 10, 20, or 30 sec, and different durations of experiments about 10–25 h. The white light LED illumination was used for acquisition of RGB images of growing bacteria colonies. Further RGB images was used for referential visual determination of bacteria colonies.

Fig. 1 Setup scheme for burst image capturing of bacteria growing process under 635 nm laser and white LED illumination.

2.1.1 Evaluation of colony growth under white light illumination

Microbial growth on the agar surface and formation of colony is a self-limiting process which starts with the maximum rate, which is gradually inhibited by exhaustion of necessary nutrients, accumulation of end products of microbial metabolism. The most common mathematical model describing microbial colony growth is the Gompertz model, which mathematical expression is presented in Eq. (1).

$$
\mu(t) = \mu \cdot e^{-\mu_i(t - t_0)},\tag{1}
$$

where μ is specific growth rate (h^{-1}) , μ_0 – specific growth rate at beginning of colony formation (h^{-1}) , μ_i – specific growth rate at infliction (or specific inhibitory rate), t_0 represents time at beginning of the colony growth and t is the actual time (h).

$$
r(t) = r_{max} \cdot e^{-\left(ln\left(\frac{r_{max}}{r_0}\right) \cdot e^{-\left(\mu_i \cdot (t - t_0)\right)}\right)}.
$$
 (2)

Eq. (2) shows the growth of the colony radius, where r_{max} is the maximal radius that the colony can reach in the given environmental conditions, r_0 is the initial radius at t_0 , μ_i is the specific growth rate at the inflexion point of the area curve. Over time gradual inhibition of colony growth occurs, which can be described by μ_i . To use these equations to simulate growth of real CFU, μ_0 and μ_i should be found. A specific set of parameters should be found to describe microbial colony growth using the Gompertz model. It was done experimentally for each microbial species and cultivation conditions (temperature and media). Fast growing sea bacteria *Vibrio natriegens* were chosen for the experiments. Bacteria were grown on salt rich, agar plates in room temperature (see the materials and methods section). First, the series of micro images were used to find the growth rate at the beginning of CFU formation (when colony growth rate is practically equal to μ_0). Thus, μ_0 for *V. natriegens* in the room temperature was determined approximately 3.875 ± 0.25 h⁻¹. Also, a series of CFU images under white light were taken to experimentally find the specific *V. natriegens* CFU inhibitory rate (μ_i) . For this purpose,

the growth of the *V. natriegens* CFU was recorded (one image in 15 min, from 0 to 20 h or more) to estimate the growth of the colony radius under white light illumination. The diameter of each CFU was determined from images using the ImageJ program. The CFU was identified from images by visual inspection (without photo enhancing). Due to the low cell concentrations, the contrast of RGB images under white light illumination was low and CFU diameters were determined not earlier than after 8–12 h. Colony diameters were determined from the image series over time. Using Eq. (2) the μ_i for *V. natriegens* were calculated at given temperature and media conditions. Example of obtained results under white light illumination presented in Fig. 4 (white light experiments curves 1–4).

$$
RMSE = \sqrt{\frac{1}{n} \sum_{i=1}^{n} (R_{Gompertz}[i] - R_{measured}[i])^2}.
$$
 (3)

According to the experimentally obtained colony results, we set maximum colony radius to be 170 pix and initial colony radius (r_0) 0.1 pix (1 pix equals 5 μm). When simulating μ_i for *V. natriegens* in the room temperature for the given colony growth, the μ_i was $0.11 h^{-1}$. However, μ_i is not a fixed number. This coefficient characterizes decreasing of the colony growth rate over the time. The colony growth rate is dependent on many factors such as: temperature, number of surrounding colonies, and local concentration gradient of nutrients. Usually, the μ_i can vary by 20–50% among the same organism. To extract parameters from the given experimental data set, parameter scan was used. As a criterion we used Root Mean Square Error (RMSE) to minimum Eq. (3) to find μ_i and t₀. Where $R_{Gompertz}$ is the radius estimated from the Gompertz model, $R_{measured}$ is the radius determined experimentally, *n* is the number of measurement points. RMSE values can vary in each experiment as they represent a match between theoretical and experimental curves.

2.1.2 Evaluation of colony growth by the processing of laser speckle time series images

For the analysis of speckle images, the growth of the *V. natriegens* colony under laser light illumination was recorded. The time interval between subsequent frames was 30 sec, which corresponds to a sampling frequency of 33.3 mHz. The frequency domain of the observed signal is up to 16.7 mHz. According to our previous research described in paper [12, 13] "signal" (or time signal) represents the result of the correlation and interpolation algorithm between each pair of consecutive speckle image frames. Further, the scatter of the specific growth rate μ_i , with the average experimental value of 0.17 h^{-1} (Fig. 3) was calculated. To obtain a curve representing growth of colony radius the following steps on processing of speckle images were performed: 1) the location of the colonies was manually marked and automatically divided into 20×20 pix sections; 2) converting each 20×20 pix section into a time series; 3) determination of signal

threshold and rising time at low frequencies according to Eq. (4):

$$
spectrogram[k, f] = |STFT[k, f]|^2,
$$
 (4)

where $STFT[k, f]$ is the Fourier transform of $sig[n] \cdot w[n-k]$, a complex function representing the phase and magnitude of the signal over time and frequency; 4) marking and determination of the times and spatial locations of detected signals; 5) marking the growth start times of each 20×20 pix square, knowing the location of each square in space and its distance from the colony center (marked manually in step 1), finally the colony radius was calculated as a function of time (Fig. 3, red line). In general, the above-mentioned processing approach determines time events that characterize dynamic changes in speckle signals for all spatial sections $(20 \times 20 \text{ pix each})$ throughout the whole experimental data set.

2.2 Detection and processing of NIR autofluorescence under 785 nm laser excitation

The spectral measurements were conducted by an experimental spectroscopic system for analysis of AF emission under 785 nm laser excitation. The experimental setup comprises a diode laser (785 nm, 100 mW, CW), a USB 4000 portable spectrometer (3.5 nm resolution, Ocean Optics, Inc., Dunedin, USA), 400 µm fiber-optic probe, and a computer (Fig. 2). To block the reflected laser light in-line filter holder with 810 nm long pass filter was used. The diameter of laser spot on the measurement surface (skin, cuvette, samples) was approximately 4 mm². Experimental setup was used for detection of AF emission from cells cultures, *ex vivo* formalized skin lesion samples and *in vivo* skin.

Fig. 2 Experimental setup for detection and analysis of NIR AF under 785 nm continuous laser excitation. Setup comprises CW laser with output power 100 mW, optical probe with 810 nm long pass filter (OD > 4), optical fiber, spectrometer, and computer.

For cells experiments each cell culture (b16f10 and DC-3F) line was cultivated in growth DMEM (Sigma-Aldrich, St. Louis, USA) medium containing 10% fetal bovine serum (Sigma-Aldrich, St. Louis, MO, USA), 1% L-glutamine solution (Invitrogen Inc., Carlsbad, CA,

USA). The cells were grown in 96 mm culture dish (TPP, Trasadingen, Switzerland) at 37 °C in humidified 5% CO2 atmosphere in the incubator. After trypsinization, cells were suspended in medium (0.25 M glucose, 1 mM MgCl₂, 10 mM Na₂HPO₄ (pH 7.4)) at a concentration of \sim 3 Mio cells/ml. After the preparation, the cells suspension was filled to the quartz cuvette (filling volume 3.5 ml, pathlength 1 cm) for further detection of NIR AF emission. The detection of NIR AF emission was performed separately for each cell culture suspension. Spectrometer integration time for collection of AF and dark signal was set to 10 sec. Dark signal was acquired from the media (0.25 M glucose, 1 mM $MgCl₂$, 10 mM Na2HPO4 (pH 7.4)) containing no cells. The distance between fiber probe and measurement cuvette was set to 4 mm, the diameter of the laser spot on cuvette surface was 4 mm^2 .

For the detection of AF spectra from formalinized *ex vivo* skin lesion samples the 5 sec integration time was used. For each *ex vivo* sample, three spectra from the lesion area and three spectra from the surrounding healthy skin were collected. Further, all collected spectra were averaged and divided to the two groups: pigmented lesions and healthy skin. As result, the averaged spectra of pigmented lesions and healthy skin were depicted in the graph. In total, 10 *ex vivo* samples were included in the study: 3 melanomas, 3 dysplastic nevi and 4 dermal nevi.

For the *in vivo* experiments the same experimental setup was used. The spectra were collected from the outside part of fingertips where melanin concentration is higher. In total, 3 healthy 30–40 years old persons with different skin phototypes were participated in the study. To minimize intensity fluctuations three AF spectra were collected on the outside of fingertips and further averaged. Averaged spectra were normalized to the AF maximum value at 825 nm. According to the geometry of experimental setup the laser spot diameter on the skin surface was approximately 4 mm^2 , providing optical density ~250 mW/cm2 . To avoid painful effects caused by heating the laser exposition time was minimized to 1 sec.

3 Results and discussion

3.1 Microbial growth analysis

Fig. 3 shows the colony radius growth as a function of time for speckle image obtained according to the Gompertz model (Eg. 2). Using the scanning method for several parameters (growth rate, growth start time, maximum radius of colony) it is possible to get the most optimal suitable Gompertz curve in which the Root Mean Square Error value will be minimal. The numerous experiments were performed using speckle imaging technique and image analysis under white light illumination (see results in Fig. 4) to understand the typical signal dynamics of *V. natriegens* colony growth.

Fig. 3 *V. natriegens* colony radius obtained by speckle image experiment (red) and theoretical Gompertz curve (blue).

Fig. 4 Comparison of growth curves of a colony of bacteria *V. natriegens* measured by analysis of laser speckle images and visual examination under white light illumination.

Comparing these two (laser speckle and white light imaging) analyses techniques, it was obtained that with the speckle imaging technique it is possible to detect microbial activity earlier than using white light imaging. The colony formation on the agar media was observed 3 h after bacterial inoculation, while the CFU growth of the colony in white light could be detected after 8–13 h. All signals depicting CFU growth (obtained from speckle and white light images) demonstrate similar dynamics (the graphs of radius increase over time are parallel to each other). However, the growth signal graphs are scattered across *OX*-axes, which can be attributed to fluctuated room temperature across the experiments thus affecting growth speed and colony detection. To conclude, the proposed method based on analysis of laser speckle time series images provides earlier responses of growing bacteria in comparison with classical CFU growth detection under white light illumination. The results (growing curves) demonstrate wide distribution which can be attributed to unstable ambient temperature, which fluctuated across the experiments and therefore altered growth speed. Undoubtedly, further additional

studies performed in a controlled microbiological environment are necessary.

3.2 Analysis of NIR autofluorescence under 785 nm laser excitation

The practical use of NIR AF for tissue characterization and diagnostics has a great scientific interest. Moreover, the exact determination of endogenous fluorophores responsible for NIR AF emission is still under discussion. In this paper, in collaboration with Ekaterina Borisova, we have conducted series of experiments in different conditions to find spectral features suitable for skin diagnostics. NIR AF experiments were performed at invitro cells cultures, at *in vivo* skin and formalinized skin lesion *ex vivo* samples.

Fig. 5 NIR AF spectra obtained from melanocytic (murine melanoma cells) and melanocytic (Chinese hamster lung cells) cells under 785 nm laser excitation.

NIR AF spectra obtained from two different types of cells is presented in Fig. 5. As shown, all obtained AF spectra has a wide intensity distribution, and can be characterized by main maxima at ~825 nm.

Fig. 6 Normalized NIR AF spectra of murine melanoma and Chinese hamster lung cells under 785 nm laser excitation.

In general, the AF spectra of highly pigmented melanoma cells can be described by the lower AF relative intensity in comparison with lung cells. After the spectra normalization (Fig. 6) the significant differences in shape of the spectra become more visible. In particular, the normalized averaged AF spectrum of melanoma cells demonstrates higher integral square in comparison with lung cells spectrum. Further we have examined AF *in vivo* spectra acquired from healthy volunteers with different skin photo-types. The Fig. 7 represents NIR AF spectra collected on the top of the finger where the melanin concentration is higher. The obtained spectra demonstrate higher AF intensity in volunteers with phototype III in comparison with skin phototype II. Remarkably, that spectra acquired from cell cultures (Fig. 5) demonstrates an opposite trend melanocytic cells had a lower AF intensity in comparison with melanocytic cells.

Fig. 7 Averaged autofluorescence spectra for 3 different volunteer's fingertips *in vivo*.

Moreover, all collected *in vivo* NIR AF (Fig. 7) spectra can be characterized by three emission maxima: main peak at ~825 nm, and two smaller peaks located at \sim 840 nm and \sim 850 nm. The local peak at \sim 910 nm most probably is attributed to the CCD etaloning effects and could be neglected for the analysis. Obtained AF spectral features (peaks at 840 nm and 850 nm) are in a good agreement with available literature data, fully described by I. Bratchenko et al. [7] Normalization (Fig. 8) of AF spectra acquired from healthy *in vivo* skin demonstrates that all processed spectra are nearly equal. No spectral shift of emission maxima appeared. However, the intensity variations of AF peaks at 840 nm and 850 nm can be observed. In particular, the intensity of skin phototype II has a lowest AF intensity at 840 nm. Undoubtedly, the number of collected signals is statistically very low, as well as relative concentration of melanin content in skin phototypes III and II does not provide strong scientific evidence to draw any conclusions. However, the obtained data is in quantitative agreement with earlier published papers available in literature [7–11].

Fig. 8 Normalized AF spectra for 3 different volunteer's fingertips *in vivo*.

Further in collaboration with Ekaterina Borisova at the Institute of Electronics, Bulgarian Academy of Sciences we have collected NIR AF signals from formalinized *ex vivo* skin lesion samples (Fig. 6). The obtained spectra were divided into two groups: 1) NIR AF spectra from melanocytic lesions and 2) spectra obtained from surrounding healthy skin.

Fig. 9 NIR AF spectra collected from *ex vivo* skin lesion samples under 785 nm laser excitation.

Obtained averaged spectra also demonstrates (Fig. 9) the same trend acquired in *in vivo* skin the highly pigmented melanocytic lesions have a higher AF intensity in comparison with healthy skin. The shape of the formalinized spectra differs in comparison with spectra collected *in vivo*. In particular, the emission peaks of AF at ~840 nm and ~850 nm were not observed in *ex vivo* samples. Probably, it can be attributed to the dehydration of the skin samples due to its formalization. Also, the destruction of the responsible fluorophores under influence of formalin is not excluded. Notably, the integral spectral square of the melanocytic lesions has is higher in comparison with healthy skin. The same tendency has been acquired in cell cultures AF spectra (Fig. 6).

To summarize, in presented paper we have analyzed NIR AF spectra obtained at different conditions. The acquired spectral shapes of melanocytic cells and *ex vivo* lesions have a similar spectral feature. In particular, the integral square of the melanoma cells and melanocytic lesions is higher in comparison with healthy skin and amelanocytic lung cells. The relative intensities at AF emission maxima (820 nm) does not provide strong correlation with tissue and cells experiments, and most probably could not be used for diagnostics purpose. However, the analysis of AF spectra at the spectral regions between 840–900 nm seems to be potential for assessment of melanocytic lesions and melanoma diagnostics. Moreover, the development of imaging techniques intended for visualization of suspicious areas of the skin should be implemented at spectral bands including AF maxima (825 nm) and regions between 840 nm and 900 nm. Undoubtedly, the future studies complemented with hypotheses and theoretical models are necessary to understand the exact phenomenon of the NIR AF.

4 Conclusions

To conclude the presented paper demonstrates a potential of laser applications for evaluation of microbial activity and skin lesion diagnostics. Analysis of laser speckle time series images provides an earlier response of growing bacteria in comparison with classical CFU growth detection under white light illumination.

Obtained AF spectra under 785 nm laser excitation deliver a quantitative correlation between *in vitro* melanoma cells and *ex vivo* melanocytic lesions. The analysis of AF intensity at spectral maxima does not provide strong correlation with tissue and cells melanin content. Further studies should be focused on investigation of *in vivo* tissue and native histology samples in order to identify key morphological features responsible for NIR AF spectral variations.

Disclosures

All authors declare that there is no conflict of interests in this paper.

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