Diagnosing breast cancer using diffuse reflectance spectroscopy and intrinsic fluorescence spectroscopy

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1 Introduction

Screening procedures such as x-ray mammography, ultrasound, and/or palpation are the first step in diagnosing breast cancer. Screening procedures do not accurately identify malignant tissue. Therefore, to establish a diagnosis, the patient must undergo breast biopsy followed by histopathological diagnosis. Due to the lack of specificity of mammography, a high percentage of suspected abnormal lesions (70 to 90%) are found to be benign or normal upon biopsy,¹ resulting in unnecessary cost, patient trauma, and time delay to obtain histopathological diagnosis. Imaging and spectroscopic optical diagnostic techniques are currently being explored to reduce the number of unnecessary breast biopsies in a minimally invasive manner.

Diffuse optical tomography (DOT), an optical imaging technique to identify suspect lesions, is currently under study.² Similar to current screening techniques, it is noninvasive and can detect lesions deep within the breast. It employs light of a number of near-IR wavelengths. Contrast, provided by scattering and absorption, measures relative concentrations of oxy- and deoxyhemoglobin, as well as the presence of lipids and water content.³ However, this technique has a low resolution,⁴ due to which lesions of small size go undetected; a resolution of 4 to 13 mm has been reported, and the sources of contrast are limited.³,⁵,⁶ Despite these challenges, DOT has been utilized in several reported breast cancer studies.⁷,⁸

A number of promising optical spectroscopy techniques for diagnosing breast cancer are also under study. These techniques currently employ point contact probes, but have the potential to be extended to imaging modes.⁹ The most frequently utilized methods are diffuse reflectance spectroscopy¹⁰–¹² (DRS) and fluorescence spectroscopy.¹³–¹⁵ As of now, intrinsic fluorescence spectroscopy¹⁶ (IFS), a technique that uses diffuse reflectance to suppress the effects of scattering and absorption in fluorescence, has not been implemented. The objective of this study is to explore the use of DRS and IFS, in combination, to identify and diagnose suspect lesions.

Briefly, DRS measures tissue scattering and absorption properties, which provide information about the morphology and biochemistry of the breast tissue epithelium and stroma. DRS typically utilizes broadband UV-visible excitation in the wavelength range 300 to 800 nm, spanning the prominent hemoglobin absorption features. Fluorescence spectroscopy pro-
vides information about the endogenous tissue fluorophores excited with UV-visible light in the same wavelength range 300 to 800 nm. Fluorophores excited in this range have the most diagnostic significance. However, fluorescence spectra recorded from tissue are distorted by absorption and scattering, thereby limiting the accuracy of the extracted information. To overcome this problem, DRS can be used in conjunction with the measured fluorescence spectrum to extract the intrinsic (undistorted) fluorescence. This process is called intrinsic fluorescence spectroscopy (IFS).

The DRS and fluorescence studies published up to now have focused on differentiating malignant from nonmalignant tissues. However, it is also clinically important to discriminate benign lesions that may carry a risk of breast cancer or that need to be surgically excised from normal breast tissue. For example, fibrocystic change is a benign condition that does not require excision, whereas fibroadenoma proliferates rapidly and must be excised. As our data will show, DRS and fluorescence spectroscopy have the potential to identify these lesions in real time at the bedside when performed percutaneously using a fiber optic probe inserted through a breast biopsy needle.

Furthermore, previously used diagnostic algorithms have not been based on parameters that can be identified as pathophysiological features of disease. While doing so is not a requirement for success, the use of parameters that may be directly compared to histopathology is likely to lead to more robust algorithms and to aid clinicians in interpreting spectroscopic results. It also represents a step toward quantifiable, objective tissue diagnosis, as compared to subjective histopathology analysis.

We report an ex vivo spectroscopic study of freshly excised human breast tissue. We use DRS and fluorescence, combined in a particular manner, to discriminate breast cancer from both benign breast lesions and normal breast tissue. Our technique, which we refer to subsequently as DRS-IFS, provides a more complete biochemical, morphological, and metabolical tissue profile than either modality alone. Our analysis extracts parameters that optimize classification and, at the same time, provide physical insight. This added information should enable the development of a more robust diagnostic algorithm.

2 Instrumentation

A clinical instrument for DRS-IFS studies, the FastEEM [Fig. 1(a)], was developed at the G.R. Harrison Spectroscopy Laboratory at the Massachusetts Institute of Technology (MIT). This instrument collects white light reflectance and fluorescence excitation-emission matrices (EEMs) in a fraction of a second. It delivers a sequence of 10 excitation laser pulses (308 to 480 nm) and two white light pulses (300 to 800 nm) to the tissue via an optical fiber probe, which is in the form of a flexible catheter with an overall length of 3 m and a diameter of approximately 1.2 mm [Fig. 1(b)]. The distal tip consists of a single delivery fiber surrounded by six collection fibers, all with core diameters of 200 μm, and terminated with a transparent, protective optical shield. The tip is brought into gentle contact with the tissue when data are taken. The diameter of tissue sampled is approximately 600 μm, with depth sensing of approximately 100 μm. The same probe delivers the excitation light and collects the diffusely reflected white light and fluorescence. At the proximal end of the probe, the collection fibers are confined in a single line and imaged onto the slit of an intensified CCD detector. Ten laser-induced fluorescence emission spectra and the two white light reflectance spectra are collected in approximately 0.3 s. Sufficient SNR is obtained by averaging five measurements, resulting in a typical acquisition time of 1.5 s. The SNR, determined as the ratio of the peak of the signal to the standard deviation of the noise, was found to be 100.

The instrument was calibrated daily prior to data collection. Calibration consisted of four steps: (1) a background spectrum of water was collected and subtracted from all subsequent measurements to correct for dark current and stray light; (2) the excitation intensities at each of 10 wavelengths were then normalized by measuring a rhodamine B dye mixture; (3) a Spectralon standard with 20% reflectance was then measured to account for the wavelength-dependent system response; (4) and finally, a mercury spectrum was acquired to calibrate wavelength.

3 Patients and Breast Tissue Handling

This ex vivo study was conducted at the University Hospitals-Case Medical Center (UHC) under approval from the UHC.
Institutional Review Board and the MIT Committee on the Use of Humans as Experimental Subjects. Spectra were collected from fresh breast surgical specimens within 30 min after excisional biopsy (n=16) and partial mastectomy (lumpectomy; n=1) procedures. Overall, we have obtained 202 spectra from 104 sites, providing approximately 2 spectra per site. Most of the 30-min delay was due to gross examination, inking, and sectioning of the specimen performed as part of the routine pathology intraoperative consultation. For each surgical specimen examined, data were collected from multiple tissue sites of interest identified by the pathologist. Spectra were also acquired from the colloidal inks used to mark the surgical margins of each specimen, and it was confirmed that no artifacts were introduced in the data as a result of this processing step.

Some of the data collected were subsequently excluded from the analysis. Specifically, DRS data with overall reflectance of less than 1% were excluded because of the inability to use reflectance information of that data to process the fluorescence data to obtain the intrinsic fluorescence. Sites whose spectra failed this criterion (11 sites) were found to be contaminated by the presence of surface blood that absorbed too much of the reflected light. Additionally, specimens from patients with preoperative chemotherapy or who underwent repeat excisional biopsy were excluded from analysis (7 specimens). Sites exhibiting other lesions [ductal epithelial hyperplasia, n=2; fat necrosis, n=1; and ductal carcinoma in situ (DCIS), n=6] were also excluded, as their limited numbers precluded development of a robust diagnostic algorithm. Following these exclusions, 202 IFS and DRS spectra acquired from a total of 104 breast tissue sites (1 and often 2 spectra per site) in specimens from 17 consecutive patients remained and were used for further analysis.

After acquisition of the spectra, the exact site of probe placement was marked with colloidal ink for registration with histopathology. The breast specimens were then fixed in formalin and submitted for routine pathology evaluation, performed by an experienced breast pathologist blinded to the spectroscopy results. The histopathology diagnoses were19 31 obtained from the surgical margins of each specimen, and it was confirmed that no artifacts were introduced in the data as a result of this processing step.

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Table 1 Expected fluorophores for various excitation wavelengths.

<table>
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<tr>
<th>Excitation Wavelength (nm)</th>
<th>308</th>
<th>340</th>
<th>360</th>
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<td>NADH</td>
<td>NADH</td>
<td>collagen</td>
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<td>collagen</td>
<td>collagen</td>
<td>collagen</td>
<td>elastin</td>
</tr>
<tr>
<td>Tryptophan</td>
<td>elastin</td>
<td>FAD</td>
<td>FAD</td>
<td>porphyrins</td>
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*aOptimal excitation wavelength for each fluorophore.*

Data Analysis

We analyzed the DRS spectrum from each breast tissue site using a mathematical model based on the diffusion approximation of light propagation in tissue to determine the values of the absorption coefficient $\mu_a(\lambda)$ and the reduced scattering coefficient $\mu_s'(\lambda)$, previously described in detail.17 The model requires index of refraction $n$, which for soft biological tissue has a typical value of 1.35 to 1.45. Two absorbers, oxyhemoglobin and $\beta$-carotene,21 were used to model the extracted absorption coefficient $\mu_a(\lambda)$. During our study, the tissue was exposed to room air and hemoglobin was oxygenated; therefore, deoxyhemoglobin was not included. For the reduced scattering coefficient $\mu_s'(\lambda)$, wavelength dependence of the form $\alpha \lambda^{-\beta}$ was employed.22

IFS spectra were extracted from the combined fluorescence and DRS spectra, as previously described.23 An IFS spectrum is composed of a linear combination of the spectra of the individual fluorophores present in the tissue. However, differences in local chemical environments, such as in tissue, may alter the shape of a fluorophore spectrum from that of its pure component.24 Therefore, pure component spectra were not directly fit to a given IFS spectrum. Instead, the IFS spectra were analyzed using multivariate curve resolution (MCR) with nonnegativity constraints, a standard chemometric method.25 MCR generates spectral components by adjusting the spectral shape of initial input parameters to minimize the fitting error to a given tissue spectrum. The initial input spectra in our case were the pure component spectra of fluorophores known to be present in breast tissue25 (Table 1). The maximum number of the initial input spectra was not constrained; however, on analysis, it was found that only two spectra were required to produce good fits to the data. The resulting MCR-generated spectral components at 340 nm are shown in Figs. 2(a) and 2(b). The combined basis spectra and fit to a representative IFS spectrum are shown in Fig. 2(c). The MCR-generated spectral components are similar to the spectra of NADH and collagen-like, respectively, but are broader and red-shifted, as expected of fluorophores in a tissue environment.26

As stated previously, the FastEEM instrument provides emission spectra at 10 different excitation wavelengths. Collagen, NADH, elastin, tryptophan, FAD, and porphyrins, the native fluorophores of breast tissue, have their maximum emission at 308, 340, 360, and 425 nm excitation wavelengths26 (Table 1). Thus, those four fluorescence excitation wavelengths were employed in the initial analysis. However, it was found that not all of these fluorophores could be detected, presumably because they were not present in our specimens at high enough levels. These include elastin excited at 340 nm and FAD and porphyrins excited at 360 nm. The absence of these fluorophores was confirmed by attempting to extract their basis spectra using MCR and finding that the contributions were negligible.

Of the measured fluorophores, only NADH and collagen-like excited at 340 and 360 nm were required to obtain good fits to the IFS spectra. However, no substantial differences in the fluorophore contributions at those two wavelengths were observed. Due to the proximity in wavelength of 340 and 360 nm, the difference in the penetration depth of the light (<5 μm) evidently was not sufficient to reveal these differences. Averaging the information at these two wavelengths...
did not improve the fits. To reduce complexity, we opted to utilize only one excitation wavelength, with the added benefit that as a result only one UV source will be required in future clinical instruments. Therefore, contributions of NADH and collagen-like excited at 340 nm were chosen for algorithm development, as this is the wavelength for maximum emission of NADH. Representative DRS and IFS spectra collected with 340-nm excitation are shown in Fig. 3.

Thus, the data analysis of each tissue site was performed using two spectra (one DRS and one IFS). Spectra were fur-
ther characterized by six parameters: $A$ and $B$—scattering parameters, fit coefficients of oxyhemoglobin and $\beta$-carotene, as well as the fit coefficients of collagen-like and NADH.

5 Algorithm Development

Following data analysis, all of the fit parameters, except $B$, were found to be diagnostically relevant. Boxplots for these parameters are shown in Fig. 4. (A boxplot is a graphical representation of data spread, with the first, second, and third quartiles “contained” in the box; the median for the data set is indicated by the black center line. The interquartile range is defined as the distance between the first and third quartiles. The extreme values are defined as 1.5 times the interquartile range, measured from the median. Data points outside of this range, indicated by crosses, are defined as outliers.) This form of data depiction is useful for visually examining the diagnostic capability of each parameter. For example, the boxplot for collagen-like indicates that normal breast sites may be partially separated from other sites using this parameter. Note that diagnostic parameters in a particular tissue type may be correlated. Hence, one parameter may provide only partial separation, but the use of a second parameter in combination with it may provide full separation. Visualizing this behavior requires a 2-D scatter plot and is not revealed by the boxplots.

The desired algorithm must be able to classify invasive breast cancer (in this case, infiltrating ductal carcinoma), benign breast lesions such as fibroadenoma and fibrocystic change, and normal breast tissue. Because each of these tissue types can likely be characterized using different diagnostic parameters, we performed the analysis in a stepwise fashion, as shown in Fig. 5. Initially, normal specimens are separated from the rest of the specimens (step 1), followed by the separation of fibroadenoma from fibrocystic change and infiltrating ductal carcinoma (step 2), and finally, the separation of infiltrating ductal carcinoma from fibrocystic change lesions (step 3). The diagnostic algorithm was developed using logistic regression and leave-one-out cross-validation. Histopathology diagnoses were used as the standard against which the spectroscopic diagnoses were compared. By maximizing the sensitivity of each stage of the algorithm we were able to identify the diagnostic parameters that can best discriminate pathologies of breast tissue.

The scatter plots and decision lines for each step of the diagnostic algorithm are depicted in Fig. 6. In step 1, normal sites are classified using the collagen-like and $\beta$-carotene fit coefficients extracted from IFS at 340-nm excitation and DRS, respectively [Fig. 6(a)]. In step 2, fibroadenoma sites are discriminated from fibrocystic change and infiltrating ductal carcinoma using the DRS scattering parameter $A$ and the IFS NADH fit coefficient [Fig. 6(b)]. Finally, in step 3, infiltrating ductal carcinoma sites are distinguished from fibrocystic change using the DRS oxyhemoglobin and a weaker dependence of IFS collagen-like (excited at 340 nm) fit coefficients [Fig. 6(c)].

Fig. 4 Boxplots of normal tissue (Normal), fibrocystic change (FCC), fibroadenoma (FA), and infiltrating ductal carcinoma (IDC) for diagnostically relevant parameters: $A$, scattering parameter extracted from DRS; $Hb$ (oxyhemoglobin) and $\beta$-carotene, absorption fit coefficients extracted from DRS; and NADH and collagen-like, fit coefficients from IFS spectra excited at 340 nm.

Fig. 5 Stepwise diagnostic algorithm.
6 Results and Discussion

Table 2 compares the DRS-IFS spectral diagnoses and the histopathology diagnoses. The diagnostic algorithm achieves the goal of distinguishing normal breast, fibrocystic change, fibroadenoma, and infiltrating ductal carcinoma using physically meaningful parameters extracted from DRS and IFS. All of the invasive carcinomas are correctly classified and only three normal specimens and one fibrocystic change lesion are misclassified as infiltrating ductal carcinoma. The total efficiency (overall accuracy) is 91% (95/104). The sensitivity and specificity are 100 and 96%, respectively, with sensitivity defined as the fraction of cancers correctly identified by the algorithm, and specificity defined as the fraction of correctly identified noncancerous lesions. The positive and negative predictive values are 69 and 100%, respectively. The positive predictive value (PPV) is the probability that a positive result is accurate, i.e., that the patient actually has the specified disease. The negative predictive value (NPV) is the probability that a negative result is accurate.\(^2\) NPV is the most important value for the clinical test we envision because it is less adverse to the health of a patient to excise normal or benign lesions (owing to a low PPV for a given technique) than to leave a cancerous lesion in place (owing to a low NPV). Therefore, it is important that a technique have a high NPV to be effective in clinical practice. Note that, unlike sensitivity and specificity, predictive values depend on the characteristics of the population and, in particular, the prevalence of disease. In our study, the disease prevalence of the data set is similar to the usual prevalence encountered in breast cancer; predictive values achieved by our technique are thus meaningful.

Our use of multiple parameters from multiple spectroscopic modalities enables discrimination of benign lesions from normal and malignant tissues. Furthermore, it allows for discrimination among different types of benign lesions in the majority of cases. This is difficult to accomplish using only one or two parameters. Gupta et al. utilized differences in the integrated fluorescence intensities to separately discriminate cancerous from fibroadenoma and cancerous from normal tissue types in an \textit{ex vivo} study.\(^\text{14}\) They were unsuccessful in their attempt to discriminate fibroadenoma from normal tissue owing to similar fluorescence intensities.\(^\text{14}\) Palmer et al. examined both fluorescence and diffuse reflectance spectroscopy during an \textit{ex vivo} study.\(^\text{28}\) Their study utilized principal component analysis followed by a support vector machine to discern differences in spectral components of the pathology groupings. It was found that multiexcitation fluorescence

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Table 2

<table>
<thead>
<tr>
<th>Pathology</th>
<th>Normal (31 Samples)</th>
<th>Fibrocystic Change (55 Samples)</th>
<th>Fibroadenoma (9 Samples)</th>
<th>Infiltrating Ductal Carcinoma (9 Samples)</th>
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<td>Fibrocystic change</td>
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<td>51</td>
<td>0</td>
<td>0</td>
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<tr>
<td>Fibroadenoma</td>
<td>0</td>
<td>0</td>
<td>9</td>
<td>0</td>
</tr>
<tr>
<td>Infiltrating ductal carcinoma</td>
<td>3</td>
<td>1</td>
<td>0</td>
<td>9</td>
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</table>
DRS and fluorescence spectroscopy was successful at discriminating malignant and nonmalignant tissues with a sensitivity and specificity of 70 and 92%, respectively. However, the sensitivity (30%) and specificity (78%) of diffuse reflectance spectroscopy alone were significantly lower. They found that the combination of fluorescence and diffuse reflectance spectroscopy did not improve the classification accuracy of an algorithm based on fluorescence spectra alone. Breslin et al.11 also explored the combination of fluorescence and diffuse reflectance spectroscopy in an ex vivo study. Using principle component analysis to discriminate malignant from nonmalignant tissue, they were unable to improve sensitivity and specificity with the combination of techniques from that achieved using only fluorescence spectroscopy. Table 3 provides a synopsis of published results for the application of DRS and fluorescence modalities to the study of breast cancer.

In contrast to these studies, we find that the majority of our diagnostic accuracy in discriminating benign from malignant lesions arises from parameters extracted from DRS. For example, generating a classification scheme from DRS parameters alone (scattering coefficient \( A \) and relative contributions of \( \beta \)-carotene and oxyhemoglobin), provides a sensitivity, specificity, PPV, NPV, and total efficiencies of 100, 100, 100, 100, and 81% (85/104), respectively. That is, all sites are correctly diagnosed as either benign or malignant. However, benign sites may be misclassified. A total of six normal and one fibroadenoma are misclassified as fibrocystic change, eight fibrocystic change lesions are misclassified as normal, and four fibrocystic change lesions are misclassified as fibroadenoma.

Somewhat surprisingly, an accurate classification scheme could not be generated from IFS parameters alone (relative contributions of NADH and collagen-like). Results for the sensitivity, specificity, PPV, NPV, and total efficiency are 89, 66, 20, 98, and 80%, respectively. Note that although the inclusion of IFS parameters into the combined algorithm does not improve the classification of benign versus malignant lesions, IFS significantly improves the total efficiency (increase of 12% from the DRS algorithm alone). The total efficiency is the only metric that considers the accuracy of the technique in subclassifying the benign lesions; specifically: normal, fibrocystic change, and fibroadenoma.

Although it is difficult to compare and contrast past studies with this current study given the wide range of experimental variables, differences in analysis methods, and differences in classification schemes and scope, we hypothesize that our model-based approach is the key advantage. We correct the observed fluorescence spectra using DRS and extract parameters that represent NADH and collagen-like fluorescence signatures. We also fit the DRS spectra to extract multiple physiological parameters. Further, our analysis better accounts for physical changes in tissue composition that contribute to changes in histopathology. Normal breast tissue consists mostly of adipocytes (fat), whereas most breast lesions are characterized by an increase in fibrous stroma and, as a result, an increase in the amount of collagen. Therefore, in the first stage of our diagnostic algorithm, we expect that normal tissue can be separated from the three types of breast lesions by the relative contributions of collagen-like and \( \beta \)-carotene, a lipid-soluble marker of adipocytes. In the second stage of the algorithm, fibroadenoma is separated from fibrocytic change and infiltrating ductal carcinoma with the aid of NADH and the scattering parameter \( A \). By histopathology, fibrocytic change and infiltrating ductal carcinoma are both cellular proliferative lesions, accounting for their relatively large contributions of NADH, a product of cellular metabolism. The parameter \( A \) is representative of the density of scatterers in the tissue.\(^{17}\) We expect infiltrating ductal carcinoma to have a relatively high value of the parameter \( A \), as a result of fragmentation and disorganization of the collagen fibers in the stroma of invasive cancers due to the action of matrix

### Table 3

Synopsis of previous works that utilizes DRS and/or fluorescence spectroscopy to diagnose breast cancer and reports sensitivity (Se) and specificity (Sp) values.

<table>
<thead>
<tr>
<th>Technique</th>
<th>Study</th>
<th>Se (%)</th>
<th>Sp (%)</th>
<th>Patient (number)</th>
<th>Site (number)</th>
<th>Discrimination</th>
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<td>85</td>
<td>31</td>
<td>126</td>
<td>Malignant versus nonmalignant</td>
<td>Bigio et al.(^{10a})</td>
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<td>Zhu et al.(^{12a})</td>
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<td>32</td>
<td>56</td>
<td>Malignant versus nonmalignant</td>
<td>Palmer et al.(^{28a})</td>
</tr>
<tr>
<td></td>
<td>Ex vivo</td>
<td>100</td>
<td>100</td>
<td>17</td>
<td>104</td>
<td>Normal, FCC, fibroadenoma, IDC</td>
<td>Volynskaya, 2008</td>
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<td>Fluorescence</td>
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<td>99.6</td>
<td>98.4</td>
<td>63</td>
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<td>Gupta et al.(^{14a})</td>
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<td>98.8</td>
<td>IDC versus normal</td>
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<td></td>
<td>Ex vivo</td>
<td>70</td>
<td>92</td>
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<td>Malignant versus nonmalignant</td>
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</tbody>
</table>

\(^{a}\)PPV, NPV, and total efficiency were not reported.
Volynskaya et al.: Diagnosing breast cancer using diffuse reflectance spectroscopy...

References


