Computational Microscopy for Sample Analysis

by

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B.E., the University of Tokyo (2010) M.S., Kyoto University (2012)

Submitted to the the Program in Media Arts and Sciences, School of Architecture and Planning in partial fulfillment of the requirements for the degree of

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Abstract

Computational microscopy is an emerging technology which extends the capabilities of optical microscopy with the help of computation. One of the notable example is superresolution fluorescence microscopy which achieves sub-wavelength resolution. This thesis explores the novel application of computational imaging methods to fluorescence microscopy and oblique illumination microscopy. In fluorescence spectroscopy, we have developed a novel nonlinear matrix unmixing algorithm to separate fluorescence spectra distorted by absorption effect. By extending the method to tensor form, we have also demonstrated the performance of a nonlinear fluorescence tensor unmixing algorithm on spectral fluorescence imaging. In the future, this algorithm may be applied to fluorescence unmixing in deep tissue imaging. The performance of the two algorithms were examined on simulation and experiments. In another project, we applied switchable multiple oblique illuminations to reflected-light microscopy. While the proposed system is easily implemented compared to existing methods, we demonstrate that the microscope detects the direction of surface roughness whose height is as small as illumination wavelength.

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Chapter 1

Introduction

Optical microscopy is a fundamental research tool in engineering and science. Through its 400-year history, numerous optical systems have been proposed to visualize microstructures that cannot be seen with naked eyes [1]. Since the introduction of digital image acquisition and image processing, computers have been more and more integrated with optical microscopy [2].

As is happening in computational photography and display [3, 4, 5, 6, 7], the combination of optical microscopy and signal processing is creating new capabilities. This emerging field is called computational microscopy. Most computational microscopy techniques acquire several images under different illumination condition and computes a synthesized image. By combining active illumination and signal processing, optical microscopy can reveal structures which is invisible only with an optical imaging system. For example, LC-PolScope and orientation-independent differential interference microscope (OI-DIC) acquire images under differently-polarized light illumination and quantitatively compute birefringence and refractive index, respectively [8, 9]. Structured illumination microscopy (SIM) illuminates a sample with differently-rotated grid patterns and generate a superresolution image [10]. Single-molecule localization microscopy, such as STORM and PALM, also reconstructs a superresolution image from images each of in which a subgroup of fluorescent molecules are stochastically excited [11, 12].

In this thesis, we explore the potential of fluorescence spectroscopy and oblique

illumination for the application to computational optical microscopy. Chapter 2 describes the theory and performance of nonlinear fluorescence unmixing for fluorescence spectroscopy and microscopy. Chapter 3 reports the development of the switchable multiple oblique illumination microscopy and its application.

Chapter 2

Nonlinear Fluorescence Unmixing for Spectroscopy and Microscopy

2.1 Proposed Method and Related Work

Fluorescence spectroscopy is a widely-used technique in many fields, such as analytical chemistry, environmental sciences and medicine. Since fluorescence spectra reflect the state and type of molecules, the analysis of fluorescence spectra nondestructively provides molecular information of a sample. For example, the analysis of autofluorescence, which is naturally emitted by a sample, has been used for the detection of cancer, contamination in water and degradation of food [13]. In addition, fluorescence spectral imaging system, where each pixel has spectral information, has expanded the capability of microscopy.

In the analysis of fluorescence spectra, the separation of fluorescence is of great importance. Most analyzed samples for fluorescence spectroscopy, including blood, water and food, have multiple fluorophores. Furthermore, it is common for fluorescence microscopy to label biological samples with multiple fluorophores. The measured fluorescence spectrum from those samples is a sum of the spectrum of each fluorophore. If their individual spectra do not significantly overlap, the fluorescence light can be separated by optical filters. However, severely-overlapping spectra cannot be separated optically. In those cases, a mathematical technique, called linear unmixing (LU), is often used to separate them.

LU is an algorithm which decomposes the total fluorescence into each fluorophore's contribution [14]. This algorithm works on the assumption that a sample's absorbance is negligible. On this assumption, the measured signal can be considered to be a linear combination of each fluorescence spectrum. Therefore, by using premeasured emission spectrum for each fluorophore, these linear equations can be solved when the number of channels recorded is larger than the number of fluorescence species. This method has been successfully applied to fluorescence spectroscopy and multi-channel fluorescence light microscopy [15, 16].

In addition to LU, under the assumption of negligible absorption, blind unmixing methods have been developed with mathematical tools, such as nonnegative matrix factorization (NMF) and parallel factor analysis (PARAFAC), also known as CAN-DECOMP and CP decomposition [17]. In many cases, to conduct blind unmixing, excitation and emission matrix (EEM) of a sample is measured. EEM is composed of a series of emission spectra measured with different excitation wavelengths. In a single fluorophore case, its EEM can be described as an outer product of its excitation spectrum and emission spectrum, while its intensity is proportional to its concentration. Therefore, the EEM of a sample containing several fluorophores has trilinearity. NMF and PARAFAC exploit this multilinearity of fluorescence.

However, when the sample's absorption is not negligible, the measured fluorescence spectra are distorted by the wavelength-dependent absorption, which is called inner filter effect. This nonlinear effect disables the introduced linear unmixing algorithms. This inner filter effect has been a major problem for fluorescence spectroscopy, and numerous techniques have been proposed to recover the intrinsic fluorescence spectra from the distorted measurements [18]. All of the methods have their own advantages and disadvantages. For example, liquid samples can be diluted to the extent where absorption is negligible [19, 20]. Even though this procedure can provide accurate estimation of intrinsic fluorescence spectra, it cannot be applied to solid materials. On the other hand, Monte Carlo simulations for the inner filter effect correction with photon migration theory can be performed on solid materials [21, 22]. However, they require expensive computation.

In this thesis, we propose a novel nonlinear unmixing method to estimate abundance of fluorophores in a light-absorbing sample for fluorescence spectroscopy and microscopy. In particular, we make the following contributions:

- We introduce a model of fluorescence spectra from a light-absorbing sample as a matrix representation. We also model spectral fluorescence imaging of a light-absorbing sample as a tensor representation.
- Based on the matrix and tensor representations, we introduce a new nonlinear unmixing method to estimate abundances of fluorophores by using nonnegative matrix and tensor factorization. This method does not require any additional measurements and can be applied to both liquid and solid samples.
- We evaluate the performance of the unmixing method on simulated fluorescence spectra and spectral images.
- We show that the nonlinear unmixing method experimentaly outperforms the conventional linear unmixing method.

2.2 Theory

In this section, we model the spectra of mixture of fluorophores in a light-absorbing medium and propose the algorithms to estimate each fluorophore's contributions from the model. We consider a conventional fluorometer and a fluorescence microscope as a measurement system.

2.2.1 Model of Fluorescence Spectrophotometer Measurements

There have been many attempts to model fluorescence spectra affected by absorption effect in fluorometer measurements [18]. Based on the Beer-Lambert law, Luciani et al. mathematically described the absorption effect in a fluorescence spectroscopy



Figure 2-1: Noiseless EEM model affected by absorption effect. The top row is the visualization of Equation 2.2. The bottom row is the visualization of Equation 2.11.

measurement in detail [19]. Following their formulation, the measured fluorescence intensity $F(\lambda_{ex}, \lambda_{em})$ can be described as:

$$F(\lambda_{ex}, \lambda_{em}) = M_1 10^{-(A(\lambda_{ex}) + A(\lambda_{em}))/2} \sum_{n=1}^N I_0(\lambda_{ex}) \phi_n c_n \varepsilon_n(\lambda_{ex}) \gamma_n(\lambda_{em}) + e, \qquad (2.1)$$

where e is the measurement noise, λ_{ex} and λ_{em} are the excitation and emission wavelength, $I_0(\lambda)$ is the intensity of the illumination, $A(\lambda)$ is the absorption spectrum of the solution, M_1 is a wavelength-independent constant factor, N is the number of fluorophores in the sample, ϕ_n , c_n , ε_n and γ_n are the quantum yield, the concentration, the molar extinction coefficient and the emission spectrum of the fluorophore n.

Let N columns of the matrix $\mathbf{A} = [\mathbf{a}_1, \mathbf{a}_2, \dots, \mathbf{a}_N]$ and $\mathbf{B} = [\mathbf{b}_1, \mathbf{b}_2, \dots, \mathbf{b}_N]$ be defined as the emission and excitation spectrum of fluorophores, where $\mathbf{a}_{n(i)} = \phi_n \varepsilon_n(\lambda_i)$ and $\mathbf{b}_{n(j)} = I_0(\lambda_j)\gamma_n(\lambda_j)$. The equation (2.1) can be rewritten in discretized matrix representation as:

$$\mathbf{F} = M_1(\mathbf{w}_1 \mathbf{w}_2^\mathsf{T}) \circledast \mathbf{A} \mathbf{D} \mathbf{B}^\mathsf{T} + \mathbf{E}, \qquad (2.2)$$

where $\mathbf{F}_{i,j} = F(\lambda_i, \lambda_j)$, $\mathbf{w}_{1(i)} = 10^{-A(\lambda_i)/2}$, $\mathbf{w}_{2(j)} = 10^{-A(\lambda_j)/2}$, $\mathbf{D} = \text{diag}(c_1, c_2, \dots, c_N)$ and \mathbf{E} is a random noise matrix. $\mathbf{A} \circledast \mathbf{B}$ denotes the Hadamard product (elementwise product) of \mathbf{A} and \mathbf{B} . Equation 2.2 is illustrated in Figure 2-1.

2.2.2 Nonlinear Fluorescence Matrix Unmixing

Here, we describe the way to estimate \mathbf{D} from the measurement \mathbf{F} in the equation (2.2). This estimation problem can be described as the following minimization problem:

$$\min_{\widehat{\mathbf{w}}_{1}, \ \widehat{\mathbf{w}}_{2}, \ \widehat{\mathbf{D}}} \|\mathbf{F} - (\widehat{\mathbf{w}}_{1} \widehat{\mathbf{w}}_{2}^{\mathsf{T}}) \circledast \mathbf{A} \widehat{\mathbf{D}} \mathbf{B}^{\mathsf{T}} \|_{F}$$
subject to $0 \leq \widehat{\mathbf{w}}_{1}, \ \widehat{\mathbf{w}}_{2} \leq 1, \ \widehat{\mathbf{D}} \geq 0.$

$$(2.3)$$

 $\|\cdot\|_F$ denotes the Frobenius norm. $\widehat{\mathbf{w}}_r$ (r = 1, 2) and $\widehat{\mathbf{D}} = \operatorname{diag}(\widehat{c}_1, \widehat{c}_2, \ldots, \widehat{c}_N)$ are the estimations of \mathbf{w}_p and \mathbf{D} , respectively. Since this minimization problem has scale ambiguities, the estimated values $\widehat{\mathbf{w}}_r$ and $\widehat{\mathbf{D}}$ are not the absolute concentrations. However, this does not become a problem, because most chemical analysis demands only the ratio of components. As in LU, we assume the emission and excitation spectra of fluorophores can be used separately.

To solve this problem, we derive new update rules by modifying the multiplicative update rules employed for nonnegative matrix factorization [23]:

$$\widehat{\mathbf{c}} \leftarrow \widehat{\mathbf{c}} \circledast \frac{\left((\mathbf{B} \circledast \widehat{\mathbf{W}}_2) \odot (\mathbf{A} \circledast \widehat{\mathbf{W}}_1) \right)^{\mathsf{T}} \operatorname{vec}(\mathbf{F})}{\left((\mathbf{B} \circledast \widehat{\mathbf{W}}_2) \odot (\mathbf{A} \circledast \widehat{\mathbf{W}}_1) \right)^{\mathsf{T}} \left((\mathbf{B} \circledast \widehat{\mathbf{W}}_2) \odot (\mathbf{A} \circledast \widehat{\mathbf{W}}_1) \right) \widehat{\mathbf{c}}}, \qquad (2.4)$$

$$\widehat{\mathbf{w}}_{1} \leftarrow \frac{((\mathbf{A}\mathbf{D}\mathbf{B}^{\mathsf{T}}) \circledast \mathbf{F}) \mathbf{w}_{2}}{((\mathbf{A}\widehat{\mathbf{D}}\mathbf{B}^{\mathsf{T}}) \circledast (\mathbf{A}\widehat{\mathbf{D}}\mathbf{B}^{\mathsf{T}}))(\widehat{\mathbf{w}}_{2} \circledast \widehat{\mathbf{w}}_{2})},$$
(2.5)

$$\widehat{\mathbf{w}}_{2} \leftarrow \frac{((\mathbf{A}\mathbf{D}\mathbf{B}^{\mathsf{T}}) \circledast \mathbf{F})^{\mathsf{T}} \widehat{\mathbf{w}}_{1}}{((\mathbf{A}\widehat{\mathbf{D}}\mathbf{B}^{\mathsf{T}}) \circledast (\mathbf{A}\widehat{\mathbf{D}}\mathbf{B}^{\mathsf{T}}))^{\mathsf{T}} (\widehat{\mathbf{w}}_{1} \circledast \widehat{\mathbf{w}}_{1})},$$
(2.6)

where $\widehat{\mathbf{W}}_r$ (r = 1, 2) is a N-column matrix whose columns are identical horizontal copies of the column vector $\widehat{\mathbf{w}}_r$ and $\widehat{\mathbf{c}} = [\widehat{c}_1, \widehat{c}_2, \dots, \widehat{c}_N]^{\mathsf{T}}$. During the updates of $\widehat{\mathbf{c}}$ and $\widehat{\mathbf{w}}_r$, $\widehat{\mathbf{D}}$ and $\widehat{\mathbf{W}}_r$ are updated accordingly. \odot represents the Khatri-Rao product, defined as

$$\mathbf{P} \odot \mathbf{Q} = [\mathbf{p}_1 \otimes \mathbf{q}_1 \ \mathbf{p}_2 \otimes \mathbf{q}_2 \ \dots \ \mathbf{p}_K \otimes \mathbf{q}_K]$$
(2.7)

for matrices $\mathbf{P} \in \mathbb{R}^{I \times K}$ and $\mathbf{Q} \in \mathbb{R}^{J \times K}$. \otimes represents the Kronecker product operator, and \mathbf{p}_i and \mathbf{q}_j denote the *i*th and *j*th columns of \mathbf{P} and \mathbf{Q} . Here, we name this method nonlinear fluorescence matrix unmixing (NFMU), which decomposes a nonlinearlydistorted EEM.

2.2.3 Model of Fluorescence Spectra from Tissue in Depth

Recently, mainly in neuroscience, there is a growing need to capture images of neuronal activities through significant depths of the brain. To meet this demand, multiphoton microscopy is becoming a basic tool to visualize neurons [24, 25]. Furthermore, in addition to conventional multilabel approach, a stochastic expression of multiple fluorescence proteins is used to label neurons with approximately 100 colors [26, 27]. However, the model of fluorescence spectra from deep tissues has not been investigated in detail so far.

In fluorescence microscopy, a captured image is the convolution of the point spread function (PSF) of the imaging system and the distribution of fluorophores in a sample. The PSF depends on excitation and emission wavelength. Therefore, if the bandwidth of exploited excitation and emission wavelength are wide, deconvolution should be performed on the captured images beforehand to analyze spectra [28]. Hereafter, we base our discussion on deconvolved images and images where the convolution effect is negligible.

In a *p*-photon process, the intensity of fluorescence from N types of fluorophores contributing to the signal of pixel k is proportional to the sum of the product of the excitation spectrum $s_n(\lambda_{ex})$, the emission spectrum $t_n(\lambda_{em})$, the abundance u_n of the fluorophore n:

$$f_k(\lambda_{ex}, \lambda_{em}) = M_2 E_0(\lambda_{ex})^p \sum_{n=1}^N s_n(\lambda_{ex}) t_n(\lambda_{em}) u_{k,n}, \qquad (2.8)$$

where M_2 is a wavelength-independent constant, and $E_0(\lambda_{ex})$ is the intensity of excitation light. When the focal plane of the imaging system is inside the object, the emitted fluorescence light is attenuated by scattering and absorption of the object. In those cases, following the photon migration model of fluorescence light from tissue, $f_k(\lambda_{ex}, \lambda_{em})$ can be described as follows [29, 30]:

$$f_k(\lambda_{ex}, \lambda_{em}) = M_3 E_0(\lambda_{ex})^p v_1(\lambda_{ex}) v_2(\lambda_{em}) \sum_{n=1}^N s_n(\lambda_{ex}) t_n(\lambda_{em}) u_{n,k}, \qquad (2.9)$$

where M_3 is a wavelength-independent factor and $v_1(\lambda_{ex})$ and $v_2(\lambda_{em})$ are attenuation factors of excitation and emission light, respectively. The equation (2.8) can be considered as the case where $v_1(\lambda_{ex}) = v_2(\lambda_{em}) = 1$.

Considering an image has K pixels, the equation (2.9) can be rewritten in tensor representation as:

$$\mathbf{\mathcal{F}} = M_3 \mathbf{\mathcal{V}} \circledast \sum_{n=1}^N \mathbf{s}_n \circ \mathbf{t}_n \circ \mathbf{u}_n + \mathbf{\mathcal{E}}$$
(2.10)

$$= M_3 \mathcal{V} \circledast \llbracket \mathbf{S}, \mathbf{T}, \mathbf{U} \rrbracket + \mathcal{E}, \qquad (2.11)$$

where \mathcal{F} is a third-order tensor whose element $\mathcal{F}_{ijk} = f_k(\lambda_i, \lambda_j)$, $\mathbf{v}_{1(i)} = v_1(\lambda_i)$, $\mathbf{v}_{2(j)} = v_2(\lambda_j)$, \mathcal{V} is a third-order tensor all of whose third frontal slices $\mathcal{V}_{ij:} = \mathbf{v}_{1(i)}\mathbf{v}_{2(j)}$, $\mathbf{s}_{n(i)} = s_n(\lambda_i)$, $\mathbf{S} = [\mathbf{s}_1, \mathbf{s}_2, \dots, \mathbf{s}_N]$, $\mathbf{t}_{n(j)} = t_n(\lambda_j)$, $\mathbf{T} = [\mathbf{t}_1, \mathbf{t}_2, \dots, \mathbf{t}_N]$, $\mathbf{u}_{n(k)} = u_{n,k}$, $\mathbf{U} = [\mathbf{u}_1, \mathbf{u}_2, \dots, \mathbf{u}_N]$, and $\boldsymbol{\mathcal{E}}$ is a random noise tensor. \circ represents the outer product operator, and $[\cdot]$ represents the Kruskal operator, which performs the summation of the outer products of the columns of matrices [31]. Equation 2.11 is illustrated in Figure 2-1.

2.2.4 Nonlinear Fluorescence Tensor Unmixing

Similarly to the case for the fluorometer, we formulate the following nonlinear least squares problem to estimate the abundance matrix **U** from the fluorescence measurement \mathcal{F} in the equation (2.11):

$$\min_{\widehat{\boldsymbol{\mathcal{V}}, \widehat{\mathbf{T}}, \widehat{\mathbf{U}}}} \| \boldsymbol{\mathcal{F}} - \widehat{\boldsymbol{\mathcal{V}}} \circledast [\![\mathbf{S}, \widehat{\mathbf{T}}, \widehat{\mathbf{U}}]\!] \|_{F}$$
subject to $0 \leq \widehat{\boldsymbol{\mathcal{V}}} \leq 1, \ \widehat{\mathbf{T}} \geq 0, \ \widehat{\mathbf{U}} \geq 0.$

$$(2.12)$$

 $\hat{\boldsymbol{v}}, \hat{\mathbf{T}}$ and $\hat{\mathbf{U}}$ are the estimated values of $\boldsymbol{v}, \mathbf{T}$ and \mathbf{U} . As is the case in (2.3), (2.12) also has scaling ambiguities, which is also not a problem because absolute concentration is not demanded in fluorescence microscopy [32]. To solve this least-squares problem, we derive the following update rules by modifying the multiplicative update rules for the non-negative tensor decomposition [33]:

$$\widehat{\mathbf{T}} \leftarrow \widehat{\mathbf{T}} \circledast \frac{(\widehat{\mathbf{V}}_2 \circledast \widehat{\mathbf{T}}) \circledast \mathcal{F}_{(2)}(\widehat{\mathbf{U}} \odot (\widehat{\mathbf{V}}_1 \circledast \mathbf{S}))}{((\widehat{\mathbf{V}}_2 \circledast \widehat{\mathbf{T}})(\widehat{\mathbf{U}} \odot (\widehat{\mathbf{V}}_1 \circledast \mathbf{S}))^{\mathsf{T}})(\widehat{\mathbf{U}} \odot (\widehat{\mathbf{V}}_1 \circledast \mathbf{S}))}, \qquad (2.13)$$

$$\widehat{\mathbf{U}} \leftarrow \widehat{\mathbf{U}} \circledast \frac{\mathbf{U} \circledast \mathcal{F}_{(3)}((\mathbf{V}_2 \circledast \mathbf{T}) \odot (\mathbf{V}_1 \circledast \mathbf{S}))}{(\widehat{\mathbf{U}}((\widehat{\mathbf{V}}_2 \circledast \widehat{\mathbf{T}}) \odot (\widehat{\mathbf{V}}_1 \circledast \mathbf{S}))^{\mathsf{T}})((\widehat{\mathbf{V}}_2 \circledast \widehat{\mathbf{T}}) \odot (\widehat{\mathbf{V}}_1 \circledast \mathbf{S}))}, \qquad (2.14)$$

$$\widehat{\mathbf{v}}_{1} \leftarrow \frac{(\mathcal{F}_{(1)} \circledast (\widehat{\mathcal{V}} \circledast \llbracket \mathbf{S}, \widehat{\mathbf{T}}, \widehat{\mathbf{U}} \rrbracket)_{(1)}) \widetilde{\mathbf{v}}_{2}}{((\widehat{\mathcal{V}} \circledast \llbracket \mathbf{S}, \widehat{\mathbf{T}}, \widehat{\mathbf{U}} \rrbracket)_{(1)} \circledast (\widehat{\mathcal{V}} \circledast \llbracket \mathbf{S}, \widehat{\mathbf{T}}, \widehat{\mathbf{U}} \rrbracket)_{(1)}) (\widehat{\mathbf{V}}_{2} \circledast \widehat{\mathbf{V}}_{2})},$$
(2.15)

$$\widehat{\mathbf{v}}_{2} \leftarrow \frac{(\mathcal{F}_{(2)} \circledast (\widehat{\mathcal{V}} \circledast \llbracket \mathbf{S}, \widehat{\mathbf{T}}, \widehat{\mathbf{U}} \rrbracket)_{(2)}) \widetilde{\mathbf{v}}_{1}}{((\widehat{\mathcal{V}} \circledast \llbracket \mathbf{S}, \widehat{\mathbf{T}}, \widehat{\mathbf{U}} \rrbracket)_{(2)} \circledast (\widehat{\mathcal{V}} \circledast \llbracket \mathbf{S}, \widehat{\mathbf{T}}, \widehat{\mathbf{U}} \rrbracket)_{(2)}) (\widehat{\mathbf{V}}_{1} \circledast \widehat{\mathbf{V}}_{1})},$$
(2.16)

where \mathbf{v}_p is the estimate of \mathbf{v}_r , $\mathbf{\hat{V}}_r$ is a *N*-column matrix whose columns are identical horizontal copies of $\mathbf{\hat{v}}_r$ (r = 1, 2), $\mathbf{\tilde{v}}_i$ is a column vector which has *K* vertical copies of $\mathbf{\hat{v}}_r$. During the updates of $\mathbf{\hat{v}}_r$, $\mathbf{\hat{V}}_r$ and $\mathbf{\tilde{v}}_r$ are updated accordingly. Similarly to NFMU, we name this method nonlinear fluorescence tensor unmixing (NFTU). This method decomposes the three-way fluorescence data affected by the nonlinear optical process.

2.3 Simulation

Simulated datasets are used in this section to demonstrate the performance of our nonlinear unmixing algorithm, NFMU and NFTU, in case of fluorescence spectroscopy and spectral imaging. The limitations are discussed in the aspects of signal-to-noise ratio (SNR), the number of channels, the number of fluorophores and the degree of the overlap of the spectra.

2.3.1 Generation of Spectroscopic Data

Fluorescence EEMs of samples containing several fluorophores in a light-absorbing medium are generated based on Equation 2.2 in MATLAB. Fluorescence excitation and emission spectra are simulated as Gaussians with its peak position μ_n and variance σ_n^2 (n = 1...N), $\mathcal{N}(\mu_n, \sigma_n^2)$. Since the shape of fluorescence spectra is not important in our algorithm, all of the variances σ_n^2 are set to be equal. An absorption spectrum is also simulated as a Gaussian which has more variance than fluorescence spectra.

As a standard dataset for the performance evaluations, the following parameters are used for the generation of the spectroscopic data: the ground truth spectra are simulated with 1 nm resolution, the sampling bandwidth for excitation and emission spectra measurements is set to be 5 nm, the number of fluorophores N is set to be 2, $\Delta \mu_n = \mu_{n+1} - \mu_n = 12$ nm for both excitation and emission spectra and $\sigma_i = 20$. The variance of the absorption spectrum is set to be 120^2 . The standard dataset is shown in Figure 2-2.

2.3.2 Performance Analysis

To quantify the error of the estimated contributions from each fluorophore, we define the normalized mean squared error (NRMSE) for the estimation of the contributions:

$$NRMSE = \sqrt{\left\|\frac{\widehat{\mathbf{c}}}{\|\widehat{\mathbf{c}}\|_2} - \frac{\mathbf{c}}{\|\mathbf{c}\|_2}\right\|_2^2 / \left\|\frac{\mathbf{c}}{\|\mathbf{c}\|_2}\right\|_2^2}.$$
(2.17)

For each evaluation, the average NRMSE was calculated from thirty trials. The parameters for generating datasets are changed for the performance evaluations in the following ways; 1) White Gaussian noise is added to the standard dataset to generate a range of SNR. 2) Sampling resolution for both excitation and emission channels is changed from 1 nm to 250 nm. 3) $\Delta \mu_i$ is changed from 1 nm to 20 nm. 4) The number of fluorophores is increased from 2 to 10. All $\Delta \mu_i$ is set to be equal. Their spectra are set to be in the range of sampling range.



Figure 2-2: Simulated fluorescence emission and excitation spectra and an absorption spectrum (top) and an EEM distorted by the absorption (bottom). Continuous lines are emission spectra, and dotted lines are excitation spectra. The contribution from both fluorophores are simulated to be the same.



Figure 2-3: Performance evaluation of NFMU. Normalized root mean squared error of the estimated contributions. The robustness of NFMU is evaluated with SNR, sampling resolution, number of fluorophores and peak wavelength difference.

Figure 2-3 summarizes the performance of our proposed method. As SNR and sampling rate increases, the proposed method estimates the contribution ratio accurately. Since the standard dataset has 5 nm resolution for excitation and emission channels, the performance degrades when the peak wavelength difference is less than 5 nm. Remarkably, the number of fluorophores does not affect the estimation error.

2.3.3 Generation of Spectral Imaging Datasets

As in spectroscopy, excitation and emission spectra and an absorption spectrum are simulated in the same way. In a 32 by 32 pixel image, each fluorophore is simulated to have different spatial distributions. Their distribution patterns are generated by



Figure 2-4: Representative images of simulated fluorophore distributions. The bar shows each fluorophore's abundance.

using the real part of the two dimensional discrete Fourier component. The representative examples of the simulated distributions are shown in Figure 2-4. Commonly, a fluorescence laser scanning microscope is equipped with several excitation lasers whose wavelength are set to be 40 nm apart. To increase signal to noise ratio, the emission channel's bandwidth is set to be 10 nm in spectral imaging mode. The standard dataset for spectral imaging is simulated in such a way.

2.3.4 Performance Analysis

Similarly to the evaluation of the fluorescence spectroscopy, the NRMSE is defined as follows:

$$NRMSE = \sqrt{\left\| \mathbf{\tilde{T}} - \mathbf{\ddot{T}} \right\|_{2}^{2}} / \left\| \mathbf{\ddot{T}} \right\|_{2}^{2}.$$
(2.18)

Here, $\tilde{\mathbf{T}}$ and $\ddot{\mathbf{T}}$ are a row-wise normalized matrix of $\hat{\mathbf{T}}$ and \mathbf{T} , respectively. For each evaluation, the average NRMSE was calculated from ten trials. Similarly to the simulation of fluorometer, the parameters for generating datasets are changed from the standard spectral imaging dataset for the performance evaluations in the following ways: 1) White Gaussian noise is added to the standard dataset to generate a range of SNR. 2) Sampling resolution for both excitation and emission channels is



Figure 2-5: Performance evaluation of NFTU. Normalized root mean squared error of the estimated contributions in various conditions. The robustness of NFTU is evaluated with SNR, sampling resolution, number of fluorophores and peak wavelength difference.

changed from 1 nm to 50 nm. 3) $\Delta \mu_i$ is changed from 1 nm to 20 nm. 4) The number of fluorophores is increased from 2 to 10. $\Delta \mu_i$ is set to be all equal. Their spectra are set to be in the range of sampling range.

Figure 2-5 summarizes the performance of NTFU in terms of the estimation error of molecular fractions. As can be seen, NTFU is robust to all of the four parameters. Even though the spectral resolution of emission wavelength is 10 nm, NTFU can accurately estimate the fractions of fluorophores whose spectra are highly overlapping.

2.4 Experimental Results

2.4.1 Fluorometer

Fluorescence solution was prepared by dissolving two fluorophores, disodium fluorescein and 4-(Dicyanomethylene)-2-methyl-6-(4-dimethylaminostyryl)-4H-pyran (pyrene derivative), in ethanol. Green ink in a commercial pen was used as light absorbent. The EEMs of samples were measured with a conventional fluorescence spectrophotometer at room temperature. The reference excitation and emission spectra of each fluorophore were obtained separately. By using these reference spectra, NFMU and LU were applied to the EEMs of the mixtures to estimate the molecular fraction.

The proposed NFMU and the linear unmixing algorithm were applied to the measured fluorescence EEMs. The estimated contribution ratios of the two fluorophores were compared with the ground truth (Figure 2-6). As is clearly seen, NFMU accurately estimated the contribution ratio, as opposed to linear unmixing.

2.4.2 Fluorescence Microscopy

The FocalCheck fluorescence Microscope Test Slide #2 (Life Technologies), which is commonly used for testing linear unmixing algorithm, was used as a sample. The test slide contains microspheres. Their shell and core are stained with different fluorophores whose fluorescence spectra are significantly overlapping. The emission spectra are shown in Figure 2-7. Images were captured by a Zeiss LSM 710 laser scanning confocal microscope with 488 nm and 514 nm excitation lasers. The emission wavelength bandwidth was set to be 10 nm. A Plan-Apo 63x/1.4 NA oil immersion objective was used. The size of a pinhole was set to 1 airy unit.

To imitate the environment of deep tissue imaging, hemoglobin from bovine blood was used as fluorescence attenuator. Lyophilized powder of hemoglobin was dissolved in distilled water and placed on a coverslip. After evaporation of the solution, the coverslip was placed on each sample. Then, the image was captured through the two stacked coverslips. Although the objective is not designed to capture images



Figure 2-6: Experimental result of fluorometer. (a) Measured emission spectra of 4-(Dicyanomethylene)-2-methyl-6-(4-dimethylaminostyryl)-4H-pyran, Disodium Fluorescein, and the mixture of the two fluorophores with absorbent and the absorption spectrum of the absorbent. The fluorescence spectra are measured with the right-angle geometry. The excitation wavelength is 400 nm. Ethanol is used as a solvent. (b) Comparison of estimated relative concentrations and reference concentrations of each type of the fluorophores. The concentrations are estimated by the developed nonlinear unmixing and the conventional linear unmixing algorithms.

through two coverslips, oil immersion objectives can capture sharp images without suffering from spherical abberation because immersion oil is designed to have the same refractive index to glass coverslips. Then, NFTU and LU were applied to the



Figure 2-7: Emission spectra of the microsphere's fluorophores, measured with confocal microscopy.

spectral images.

Fig. 2-8 shows the images of the microspheres unmixed by LU and NFTU. Since the microspheres are stained with different fluorophores at its core and shell, the fluorescence light from Fluorophore 2 should dominate at the shell, which is seen in the LU-unmixed images captured without hemoglobin. However, the LU-unmixed images captured through hemoglogin do not show the dominance of Fluorophore 2 on the shell. On the other hand, the NFTU-unmixed images clearly show the dominance of the Fluorophore 2 on the shell. The signals both at the core and the shell has contribution from both fluorophores because of out-of-focus light.

2.5 Discussion

2.5.1 Benefits and Limitations

In summary, we have presented new nonlinear fluorescence unmixing algorithms, NFMU for fluorescence spectroscopy and NFTU for fluorescence microscopy, which use multiplicative update rules for nonnegative matrix and tensor factorization. From a EEM measurement of a sample, NFMU estimates the abundance of fluorophores by using each fluorophore's EEM. On the other hand, NFTU unmixes spectral images of a multiply-stained sample acquired with several excitation wavelengths by using only emission spectra of flurophores. To the best of our knowledge, NFTU is the only unmixing algorithm which separates fluorescence spectral images affected by absorption. Although we showed the performance of both NFMU and NFTU on samples containing two fluorophores in experiments, they can be applied to unmix spectra of samples containing more than two fluorophores, as we show in the simulation.

NFTU is a natural extension of NFMU. However, in contrast to NFMU, NFTU exploits the benefit of large number of samples in one image acquisition. By utilizing this advantage, NFTU is much more robust to noise compared to NFMU.

NFMU and NFTU have several limitations inherent to EEM measurements and fluorescence spectral imaging. Since both methods require scanning of wavelength with narrow bandwidth, there is an inevitable trade-off between scanning time, noise level and sampling resolution. Since this trade-off depends on the properties of fluorophores, an optimization framework for finding the best measurement parameters should be developed. In addition, as is the case with SU and PARAFAC, NFMU and NFTU cannot be applied when fluorescence spectral properties change due to other nonlinearities, such as quenching and pH dependence [17].

2.5.2 Future Work

While our algorithms, NFMU and NFTU, require spectral information of fluorophores as a priori knowledge, we would like to explore a new mathematical technique for blind unmixing of our fluorescence model 2.2 and 2.11. As is incorporated in several blind unmixing methods [34, 35], fluorescence lifetime would help to solve the problem. Furthermore, the evaluation of NFMU should be compared with existing inner-filtering correction methods. While we examined our algorithms on experimentally-controlled samples, we would like to apply our methods, NFMU and NFTU, to real world problems, such as the analysis of dissolved organic matter and deep tissue imaging.



Figure 2-8: Unmixed images of microspheres stained with two fluorophores. The left and middle column show the unmixed results on the same spectral image datasets captured through hemoglobin. The right column shows the unmixed results by LU on the datasets captured without hemoglobin. The top row shows the combined image of the second and third row. The bottom row shows the contribution from each fluorophore on the yellow line of the top-row images. Colors are pseudocolor.

Chapter 3

Switchable Multiple Oblique Illumination Microscopy

3.1 Proposed Method and Related Work

Surface roughness in microscopic scale is an important characteristic of materials. To characterize it, various optical methods have been developed, such as confocal microscopy [36], fringe projection [37] and interferometry [38]. There are also more accurate non-optical methods available, including scanning electron microscopy (SEM) and atomic force microscopy (AFM). Among them, microscopic photometric stereo is one of the easiest and cheapest methods for 3D measurement of microscopic structure [39]. With a conventional long-working distance microscope, this method captures several microscopic images with manually-changed illumination directions and applies the photometric stereo method to extract surface albedos, surface normals and height of microstructure. Even though it is easy-to-implement, this method only shows the structures more than several micrometers due to diffraction of light. However, there is a need in many fields, such as defect inspection and dermatology, for a method that can visualize the surface microstructure as small as the illumination wavelength with an easily-implementable device, instead of accurately measuring the roughness. We introduce such a method here.

Our technique applies the strategy of oblique illumination and switchable multi-

ple light sources to a reflected light compound microscope. Oblique illumination is a traditional technique in transmission light microscopy to enhance image contrast. It is also applied to endoscopy to obtain phase-gradient images [40]. Although this microscope, called oblique back-illumination microscopy, configures epi-illumination geometry, the light collected by an objective lens transmits through tissue like transillumination geometry. As often used in reflected-light stereomicroscopy, oblique illumination can reveal surface structures much smaller than the illumination wavelength [36]. In addition, the angle of illumination relative to both a sample and the optical axis is also of great importance to increase image contrast. However, to the best of our knowledge, the capability of oblique illumination with reflected-light compound microscopy has not been explored yet. In particular, we make the following contributions:

- We build the prototype of the switchable multiple oblique illumination compound microscope system in reflection mode.
- We examined its performance by acquiring images of microscopic features as small as the illumination wavelength.

3.2 Method

This section describes the switchable oblique reflected illumination microscopy which qualitatively assess the surface roughness.

3.2.1 Hardware Implementation

The hardware is composed of Canon's EOS Rebel T3 digital camera, Olympus's UPlanSApo 20x NA0.75 objective lens and tube lens, a No.1.5 coverslip, an iris diaphragm ($\emptyset 0.8$ to $\emptyset 20$ mm), Arduino Mega with a light-emissing diode (LED) driving circuit, a XYZ translation stage and 3D printed light blockers and sample stages and white LEDs (Figure. 3-1). The illumination angle is set to be nearly 90° to the optical



Figure 3-1: Prototype of the multiple oblique illumination microscope.

axis of the objective lens (Figure. 3-2). To increase the depth of field, the size of the iris is set to be the smallest size.

Our microscope system is controlled via MATLAB. The LEDs are switchable through serial communication, and the image acquisition is achieved with the crossplatform digital camera library, gPhoto2 [41]. Two images are sequentially acquired with each LED illumination.

3.2.2 Sample Preparation

Intact black hair and bleached and dyed black hair are used as samples. Hair is covered with cuticle cells which consist of the outermost layers. Each cuticle cell has about 500 nm thickness. Since bleaching and coloration dissolve the cell membrane complex that connects cuticles to each other, the cuticles of the bleached and dyed hair is partly ripped off from the surface [42]. The SEM images of intact and damaged hair surface is shown in Figure 3-3. For the image acquisition, a single strand of hair is placed on the custom-designed stage (Figure 3-2).

3.3 Result

Figure 3-5 and 3-6 show the images of the healthy hair and highly-damaged hair illuminated from tip and root side. In the case of black hair, light penetrating the surface is absorbed by the melanin pigment, and subsurface scattered light is negligible



Figure 3-2: A sample stage for a hair strand and a adapter for a translational stage (left) and a schematic illustration of illuminating direction (right). The sample holder is designed to be inserted to the adapter to the translational stage.



Figure 3-3: Scanning electron micrographs of healthy hair (left) and damaged hair (right), cited from [42].

for the image formation. Therefore, the image-forming light is surface-reflected light.

As is illustrated in Figure 3-4, the edge of cuticles of intact hair has higher scattering and reflection when illuminated from root. Since the effective roughness for illumination from tip is the thickness of cubicles, incident light from tip strongly scatters at the edge of cuticles. On the other hand, in the case of damaged hair, both incident illuminations from tip and root side scatter and reflect strongly at the edge of detached cuticles due to the ripped-off cuticles.

To extract the scattering effect in images, the variance map of the captured images are calculated and shown in Figure 3-5 and 3-6. As can be clearly seen in Figure 3-6,



Figure 3-4: Illustration of the interaction between the direction of illumination and hair surface. The condition of cuticles affects the reflection and scattering of light.

the computation of local variance enhances the contrast of cuticles and suppresses the undesirable contrast from out-of-focus light. While the healthy hair shows the shape of cuticles in the variance image with the illumination from tip side but not with the illumination from root, the shape of cuticles is revealed in the variance map with both illuminations.

3.4 Discussion

3.4.1 Benefits and Limitations

Taking intact and damaged hair as a representative sample, we show that the switchable oblique reflected illumination microscope system extracts the anisotropy of surface roughness whose height is about the same size as illumination wavelength. While one of the computational photography techniques enhances depth-edge with multiple flashes in macroscopic scale [43], our microscope extracts depth-edge in microscopic



Figure 3-5: Images of healthy black hair captured by the prototype and their processed images. The top row shows the original microscopic images. The bottom row shows their local variance map. Illumination from root side enhances the contrast at the edge of cuticles.



Figure 3-6: Images of damaged black hair captured by the prototype and their processed images. The top row shows the original microscopic images. The bottom row shows their local variance map. Illumination from both tip and root side enhances the contrast at the edge of cuticles.

Microscopic image

scale. Compared to the conventional surface probing technologies, our system is remarkably low-cost and small. This simple setup will allow surface diagnostics of materials in resource-limited environment.

Furthermore, our microscopic technique reveals the surface damage of hair. To the best of our knowledge, the detachment of cuticles has been undetectable with optical microscopy. Currently, hair researchers utilize SEM to analyze the surface of hair. However, SEM requires metal-coating for biological specimens, which prevents the observation of samples in intact state. Therefore, a method to characterize the surface structure without artificial procedures on samples has been awaited. Our microscope system has large potential to advance hair research and can be used as a quick hair diagnostic device for consumer applications.

The prototype has several limitations. While AFM and SEM can quantitatively measure the roughness and reveal three dimensional surface structure, our system does not provide quantitative roughness value and elucidate the structure of surface. In addition, subsurface light transport can contribute to the image formation, which may deteriorate the analysis. In this case, the removal of subsurface reflection with an advanced lighting system, such as structured illumination and polarized light, is required. A detailed analysis of the multiple oblique illumination system in comparison to the conventional probing technologies is left for future work.

3.4.2 Future Direction

Our current prototype utilizes a personal computer, a high-quality objective lens and a CCD image sensor of a digital SLR camera. As a next step, we would like to miniaturize it with the use of a smartphone. Since most smartphones have enough computation power and a high resolution image sensor, we plan to develop an attachment which modifies a smartphone to a multiple oblique illumination microscope. Furthermore, we currently compute local variance of images after capture. We would like to explore the strategy to quantitatively measure roughness by the co-design of optical system and computation. We believe that the quantitative measurements further expand the application of the switchable multiple oblique illumination.

Chapter 4

Conclusion

Computational control of illumination creates large potential in optical imaging [44, 45, 46]. As opposed to photography, optical microscopy has full control of illumination. This capability has allowed numerous optical designs to acquire different properties of microscopic samples. Even though spectral imaging was introduced to fluorescence microscopy about ten years ago and oblique illumination was invented more than three hundred years ago, integration with modern computational approaches had not been explored. In this thesis, we explored the possibilities of fluorescence spectroscopy and microscopy and oblique illumination microscopy.

In science and engineering, not limited to microscopy, a large number of advanced sensing methods have been proposed to measure various material properties. In most cases, however, computational and mathematical analysis of captured data have not been conducted enough to extract information as much as possible. We hope that this thesis inspires others to explore imaging and sensing technologies at the interface of material analysis and computing.

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