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LIGHT SCATTERING FROM MOTILE BACTERIA

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Abstract. — The «Intermediate scattering function» \( I(K, t) \) of light scattered by motile bacteria is shown to be
\[
I(K, t) = \int_0^{\infty} \frac{\sin KVt}{KVt} P_s(V) \, dV , \tag{1}
\]
where \( P_s(V) \) is the distribution of swimming speeds of the bacteria. Scattered spectra from motile \( E. \ coli \, K_{12} \) bacteria have been investigated by laser light intensity correlation spectroscopy. \( [I(K, t)]^2 \) is obtained from the data and its angular dependence is shown to agree with that predicted theoretically by eq. (1). In contrast, bacteria whose motility has been arrested by addition of \( 10^{-2} \, M \, CuCl_2 \) are found to scatter light with a spectrum characteristic of large Brownian particles.

The swimming speed distribution \( P_s(V) \) may be determined by performing appropriate Fourier inversions of the data taken from the motile bacteria. The manner by which the distribution changes when environmental factors are varied is indicated.

We have been using laser light intensity correlation spectroscopy to determine various quantities related to the motility of bacteria. Particular attention has been given to the longitudinal swimming speed distribution of the bacteria \( P_s(V) \), which is especially easy to obtain. In the following we first review the theoretical relationships between \( P_s(V) \) and the intermediate scattering function \( I(K, t) \). The latter may be obtained directly from the light scattering data, and we discuss various measurements of bacterial motion based upon these considerations. We also report upon a preliminary experiment showing the possibility of studying the chemotactic responses of the bacteria.

Motile strains of \( E. \ coli \, K_{12} \) bacteria [1] are of specific interest to us. When these bacteria are observed under a microscope they appear to move at constant speed in straight lines. These motions persist for times of the order of seconds before the bacteria change directions. Such times are long when compared with typical decay times of bacterial scattering spectra. Consequently, for the purpose of calculating \( I(K, t) \), one may assert that the velocities of the individual
bacteria remain unchanged over measurement times relevant to the scattering experiments. It is then particularly easy to calculate \( I(K, t) \), since the effect of the bacteria simply is to cause a Doppler shift of the frequency of the laser beam [2], [3]. Indeed, for such a homogeneous sample of swimming bacteria, \( I(K, t) \) is given as

\[
I(K, t) = |A|^2 \left( \sum_{\text{bacteria}} \left( e^{-iK \cdot R_0(t)} e^{-iK \cdot R_1(t)} \right) \right)
\]

\( = N |A|^2 \left< e^{iK \cdot V_t} \right> \),

where \( V \) is the bacterial velocity, \( K \) is the Bragg wave vector, and \( t \) is the time. If no external forces are acting upon the bacteria, the velocity distribution is isotropic, and we have

\[
\left< e^{iK \cdot V_t} \right> = \int_0^\infty \int_0^{2\pi} e^{iKV_t \cos \theta} \sin \theta [V^2 P(V)] \times dV d\theta d\varphi
\]

\[
= \int_0^\infty \frac{\sin KV_t}{KV_t} \left[ 4 \pi V^2 P(V) \right] dV .
\]

Upon identifying the swimming speed distribution \( P_s(V) \) as

\[
P_s(V) = 4 \pi V^2 P(V),
\]

then \( I(K, t) \) may be written as

\[
I(K, t) = \int_0^\infty \frac{\sin KV_t}{KV_t} P_s(V) dV,
\]

in a form normalized to the value \( I(K, 0) = 1 \).

A laser light intensity correlation spectrometer, specially designed and constructed by one of us (S. H. Chen), was employed in this study. The source for this spectrometer is a well stabilized He-Ne laser (Spectra-Physics model 125) operating in the TEM\(_{00}\) mode (\( \lambda = 6328 \) Å). The sample rests in a temperature controlled bath (in these experiments the stability was \( \pm 0.1 ^\circ \text{C} \)). Scattered light at various scattering angles \( \theta \) is collimated by two pinholes of 0.2 mm diameter, approximately 50 cm apart, and is detected by an ITT F-4085 phototube having a 0.35 mm circular photocathode. The loss of spatial correlation of the light at the photodetector in this geometry is only about 5%. The output of the phototube is processed by a set of fast amplifiers and a fast discriminator to produce a series of standardized short pulses (50 ns wide, 5 V amplitude), each pulse corresponding to detection of a photon. These pulses are then fed into a digital autocorrelator which computes the «clipped autocorrelation function» on 128 delay time points simultaneously.

The correlator measures, essentially, the expected number of counts at the delay time \( \tau \), given that at \( \tau = 0 \) there are more than \( k \) counts within a counting period \( T \). This is achieved by splitting the input pulse trains into two routes, clipping the first into a binary sequence, and then performing a delayed coincidence measurement with the second. The schematic circuit diagram of a portion of our 128 channel correlator is given in figure 1. The correlator has buffer storage capacity of 4 bits per channel; the overflow pulse (corresponding to 16 counts) is transferred into the memory of a multichannel analyzer every 1.31 ms. (The control and timing circuits, and the input clipping register, are not shown in figure 1.) The detailed circuit and its operation are to be reported elsewhere [4].

The total counts accumulated in the \( j \)-th channel \( N_j \) in the multichannel analyzer is

\[
N_j = \eta_t < n_s(0; T) n(jT; T) >
\]

where \( n(\tau; T) \) is the number of counts at time \( \tau \) accumulated over the gating time \( T \), and the product \( \eta_t T \) is the total measurement time. \( n_s \) has a value of 1 when \( n(0, T) > k \) and is otherwise 0 (see Jakeman and Pike [5] for a precise discussion of notation). The uncorrelated d. c. background \( N_B \) can be obtained by letting \( \tau = jT \to \infty \), and observing that

\[
N_B = \eta_t < n_s > < n > = N_{\text{trigger}} < n > ,
\]

where \( N_{\text{trigger}} \) is the number of 1’s in the binary sequence and \( < n > \) is the average number of counts per gating time. Our correlator also registers these numbers during the experiment, so that the uncorrelated background may be unambiguously extracted. Foord et al. [6] have reported an 8-channel correlator built on the same principle. They further showed that, when the statistical nature of the scattered light is Gaussian, aside from a small spatial coherence correction factor one may write

\[
\frac{N_j - N_B}{N_B} = \frac{1 + k}{1 + < n >} \left| g^{(1)}(\tau) \right|^2 .
\]
In our case $g^{(1)}(\tau)^2 = I^2(K, \tau)$ [7]. For purposes of comparing spectra in these experiments it is useful to normalize $I(K, t)$ such that $I(K, 0) = 1$. To accomplish this we simply subtract the uncorrelated background from the total counts of each channel and then normalize the resultant data to unity at $\tau = 0$.

Eq. (4) implies that $I(K, t)$ should be a functional of the variable $X = Kt$, i.e., if the data taken at different angles are plotted as a function of $X$ rather than as a function of time, they should superimpose. Figure 2a shows scattering data from a motile sample, taken at three different angles. These curves have been obtained from the output of the correlator by subtracting the uniform background according to eq. (6) and then normalizing. The bacteria in this sample seemed to move independently (the sample contained approximately $10^7$ bacteria/cc), as indicated by the fact that dilution of the bacteria did not give rise to changes in the normalized scattering spectrum. Thus, although the spectrometer is operated in the homodyne mode in our experiments, the data given in figure 2a may be directly interpreted as $[I(K, t)]^2$. In figure 2b the data are plotted as a function of $X = Kt$, and we see that the parametric dependence indicated by eq. (4) is well substantiated.

The Fourier sine transform of eq. (4) yields the swimming speed distribution $P_s(V)$, viz.,

$$P_s(V) = \frac{2V}{\pi} \int_0^\infty \sin XV[I(X)]dX.$$  

(8)

In figure 3 we show the swimming speed distribution which corresponds to the spectrum presented in figure 2. Parenthetically, we remark that the swimming speed distribution of the bacteria decreases to zero at a much slower rate than does the Maxwellian distribution which has its maximum at the same speed.

The bacteria to which the data of figures 2 and 3 pertain all appeared to be moving when observed under the microscope. In general, a sample may also contain non-motile bacteria. To study the scattering properties of such bacteria, $10^{-2}$ M CuCl$_2$ was added to a motile sample. After several hours persistent movement had ceased and the bacteria assumed the characteristics of large Brownian particles. Scattering data for this motion-arrested sample are presented in figure 4, where they have been plotted as a function of the variable $Y = K^2 t$. The curves corresponding to different scattering angles overlap remarkably well.

If the non-motile sample were to consist solely of spherical particles of a single size, the intermediate scattering function would have the form

$$I(K, t) \sim e^{-DK^2 t}.$$  

However, as seen from the inset of figure 4, the data do not quite follow such a simple exponential decay.
The deviation probably is due to heterogeneity of the sample (e.g., clumping of the bacteria, dust, etc.), but the spectrum may also contain components arising from rotational diffusion. On the other hand, an effective value for the longitudinal diffusion coefficient of the non-motile bacteria may be obtained, for example, from the expression

$$D_{av} = -\lim_{r \to 0} \frac{d \ln I(Y)}{dY}.$$  

For the sample discussed in figure 4, such a procedure yields the value $D_{av} = 5 \times 10^{-9}$ cm$^2$/s corresponding to an equivalent Stokes diameter of 1.0 micron.

Of course, a sample will not necessarily be either entirely motile or entirely non-motile. Analysis of the scattering spectra is slightly more difficult when the sample contains both types of particles. However, by taking measurements at different scattering angles, an estimate of the fraction of each component in the sample may be made and, for example, data pertaining only to the motile particles may be extracted. A procedure for unscrambling the two types of spectra is discussed in Appendix A. It is important to take data always at more than a single angle, in order to check on the characteristics of the particles being studied.

Currently, we are gathering data on the manner by which the swimming speed distributions of the bacteria are affected by changes in such parameters as temperature, salinity of the media, and concentrations of metabolites. An example of the change which can be seen in the spectrum is shown in the oscilloscope displays presented in figure 5a. These data pertain to the effect of NaCl on motility. In figure 5b, we present a preliminary analysis of some data showing the inhibition of motility resulting from increasing NaCl concentration.

Finally, we should like to report upon some crude experiments which suggest that techniques similar to those discussed above can be applied to study directional responses of the bacteria. In figure 6 we show some photographs illustrating oxygen taxis [8], [9]. Bacteria which had been washed free of broth and resuspended in a medium consisting of $10^{-4}$ M EDTA, $10^{-2}$ M KPO$_4$ buffer pH 7.0, and $10^{-2}$ M glucose.
The bacteria move in response to the gradients in concentrations of various chemicals [9]. When the scattering cell lies on its side, the gradient in oxygen concentration occurs in the plane defined by the incident and detected beams; on the other hand, when the cell is upright the component of motion due to the oxygen taxis is perpendicular to the plane of movement. When the various spectra are compared, the modulation arising from the directional motion associated with the oxygen taxis is clearly seen.

**Appendix A. — Mixtures of Motile and Non-motile Scatterers.**

When the scattering sample is composed of a mixed population of motile and non-motile particles, estimates of the properties of each type may be obtained in the following manner. Let us first represent the portion of the spectrum associated with the non-motile particles as a single exponential (cf. Fig. 4). Then, the entire spectrum may be described as

\[
I(K, t) = \alpha I_{sw}(K, t) + (1 - \alpha) e^{-K_D t},
\]

(9)

where \(I_{sw}\) is the spectrum due to the swimmers, \(\alpha\) is the fraction of the scattered light due to the motile particles, and \(I(K, t)\) is normalized so that \(I(K, 0) = 1\). If we now rewrite eq. (9) in terms of the variable \(X = Kt\), we find

\[
I(X) = \alpha I_{sw}(X) + (1 - \alpha) e^{-KDX}.
\]

(10)

Therefore, if data pertaining to two scattering angles are expressed in this way and subtracted one from the other, we find the «difference spectrum» \(I_{diff}(X)\) to be given as

\[
I_{diff} \equiv I_1 - I_2 = (1 - \alpha) [e^{-K_1DX} - e^{-K_2DX}].
\]

(11)

When \(K_2 > K_1\), eq. (11) has the form shown in figure 7. The quantity \(\alpha\) may be determined from the
maximum of the curve, and $D$ may be determined from the point $X_m$ where the maximum occurs (see caption of Fig. 7). The spectrum due only to the swimming particles may be obtained by subtracting the portion arising from the non-motile particles and then renormalizing.

\[ I_{\text{diff}}(X) = (1 - \alpha) \left\{ \frac{K_1}{K_2} \right\} \left( \frac{K_2}{K_1} \right)^{K_3(K_1-K_2)} \]

\[ (\theta_1 = 20^\circ, \theta_2 = 40^\circ). \]

Fig. 7. — $I_{\text{diff}}(X)$ vs. $X$. Example of a mixed population of motile and non-motile scatterers. The diffusion coefficient may be determined according to $D = \frac{1}{X_M(K_1 - K_2)} \ln K_1/K_2$. $\alpha$ may be determined from the expression

\[ I_{\text{max}}^{\text{diff}} = (1 - \alpha) \left\{ \frac{K_2}{K_1} \right\}^{K_3(K_1-K_2)} \]

\[ \left( \frac{K_3}{K_2} \right)^{K_3(K_1-K_2)} \]

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