Dopant spatial distributions: Sample-independent response function and maximum-entropy reconstruction

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We demonstrate the use of maximum entropy based deconvolution to reconstruct boron spatial distribution from the secondary ion mass spectrometry (SIMS) depth profiles on a system of variously spaced boron δ layers grown in silicon. Sample-independent response functions are obtained using a new method that reduces the danger of incorporating real sample behavior in the response. Although the original profiles of different primary ion energies appear quite differently, the reconstructed distributions agree well with each other. The depth resolution in the reconstructed data is increased significantly and segregation of boron at the near surface side of the δ layers is clearly shown. [S0163-1829(97)05147-3]

The improvement of the depth resolution achieved in sputter profiling in general and secondary ion mass spectrometry (SIMS) in particular has continued over the past ten years despite occasional predictions about the limit having been reached.¹ Nevertheless, the resolution achieved directly may not be adequate for future generations of semiconductor device material, even if ultralow energies^{2,3} or large cluster ions⁴ are employed in the primary beam.

The SIMS mass transport effects due to energy deposition and probe beam incorporation from the primary beam are well known—the measured profile is broadened and shifted from its true position.^{5–7} Although the use of low beam energies or high mass clusters can greatly alleviate the effects, such mass transport effect is *intrinsic* to the SIMS depth profiling process and can still be observed even when the primary ion beam energy is as low as 250 eV (Ref. 8) or when using SF₅⁺ ions at 600 eV.¹⁴ The true spatial distributions remain to be reconstructed especially when there is an abrupt interface or δ doping present.

In the concentration range of common practical interest in SIMS, e.g., the dilute limit where dopant concentration $\leq 1\%$, there is a strictly proportional relation between the signal and the instantaneous surface concentration as well as between the primary ion flux density and erosion rate outside the transient region for a single matrix.⁹ Where the essential physics of the analytical process is linear, deconvolution is the mathematically correct method to recover and quantify the depth profiles.¹⁰ The ideal SIMS signal at the depth *z*, $Y_0(z)$, can be expressed as a convolution of the true concentration distribution C(z) with the SIMS instrumental response function R(z),

$$Y_0(z) = \int C(z') R(z - z') dz',$$
 (1)

since primary ion flux must be also proportional to the elapsed time (otherwise depth calibration becomes problematic). Here we define $Y_0(z)$ and C(z) with the same dimension of concentration per unit length and R(z) normalized over the depth to simplify the equation and ensure that sample mass is conserved. A slightly more sophisticated model $Y_0(z) = \int C(z')R(z-z',z)dz'$ might be used if the depth resolution was depth dependent.

It would be a straightforward inverse problem to determine the true spatial distribution C(z) if the corresponding ideal SIMS signal $Y_0(z)$ could be measured. In fact, the measured SIMS signal Y(z) is as usual a combination of the ideal signal and associated non-negligible noise component $Y_N(z)$,

$$Y(z) = Y_0(z) + Y_N(z).$$
 (2)

There is no obvious way to find C(z) from Y(z). Various methods have been used to obtain C(z).³ Yet, the lack of objective evidence makes it very difficult to reconstruct the real features and separate them from the SIMS effect justifiably, e.g., to distinguish the segregation and diffusion occurring during growth at the interface of two different materials from the SIMS atomic mixing. Moreover, the peculiar character of the SIMS depth profile data everywhere positive, and large dynamic range (may span 10 orders of magnitude overall and 4-6 orders for a particular species), requires a very careful and unbiased treatment. Manipulating data arbitrarily, such as placing a subjective penalty on each change in slope or simply filtering certain range of frequency components, could seriously distort the final results and/or easily lead to unphysical negative values. We believe that only the features with statistical evidence in the original data should be extracted and a empirical deconvolution method¹¹ based on the maximum-entropy (MaxEnt) principle^{12,13} fulfils such a requirement.

The success of a MaxEnt deconvolution method relies on the finding of a sample-independent response function and a suitable noise model. Neither of them can be obtained for the SIMS process with sufficient accuracy from first-principles calculations because of incomplete knowledge. Our recent investigation shows that the noise in the SIMS depth profiling follows Poissonian statistics universally¹⁴ and the relation between the mean counts $s_m(z)$ and its corresponding standard deviation $\sigma_s(z)$ is $\sigma_s(z) = s_m(z)^{1/2}$.

The response function should contain only the SIMS related information, i.e., the broadening, the shift and possibly the ion yield, but must not contain sample dependent fea-

15 167

tures. Ideally, the response function is the transient measured from an infinitesimally thin layer, often known as a δ layer. However, such a δ layer is an abstraction and even if it were not, we have no means to recognise its existence, other than very locally. Moreover, as the intrinsic resolution of SIMS is improved by using sub-keV probes,⁸ it is readily apparent that the real approximations to such structures that can be grown leave a measurable sample-related shape content in the SIMS profile due to statistical placement of atoms, segregation, and diffusion at the growth temperature, etc. Simply using such data as response function, deconvolution will suppress real features in the depth profile, and produce artificial concentration slopes and unrealistically small feature widths.

In the following, we outline a method for extracting a sample-independent SIMS response function for the case of boron in silicon sampled by oxygen beam for various primary beam energies. Subsequently, we demonstrate a Max-Ent deconvolution to reconstruct the dopant distribution from SIMS depth profiles using the corresponding response function and noise model. The results are then discussed.

Experimental and theoretical studies explored the mass transport in the SIMS process.^{15–18} It has been shown that the normalised response function can be represented by the following form in several orders of magnitude:¹⁸

$$R(z) = \frac{1}{2(\lambda_g + \lambda_d)} \left\{ \left[1 - \operatorname{erf}(\xi_g) \right] \exp[z/\lambda_g + (\sigma/\lambda_g)^2/2] \right. \\ \left. + \left[1 + \operatorname{erf}(\xi_d) \right] \exp[-z/\lambda_d + (\sigma/\lambda_d)^2/2] \right\},$$
(3)

where $\xi_g = (z/\sigma + \sigma/\lambda_g)^{1/2}$ and $\xi_d = (z/\sigma - \sigma/\lambda_d)^{1/2}$, σ is the primitive standard deviation, λ_g and λ_d the growth and decay lengths. The smaller the λ_g and λ_d are, the sharper the R(z) will be. When they approach zero, the R(z) will degenerate to a Gaussian distribution with the deviation σ . All these parameters of the response function apparently depend on primary ion beam energy E_p and should be monotonically increasing functions of it. If we could find a perfect δ layer, we might be able to fit the measured data with R(z) to obtain these parameters for a certain E_p and use them directly to deconvolve other SIMS profiles. Therefore, the difficulty here is how to determine the σ , λ_g , and λ_d from the measured data justifiably, since there are no other techniques that we can use to check the results. We have to substantiate our choices through statistical and trend analyses.

Before we study the R(z) of a specimen in a crystalline substrate, we first look at the R(z) of the same specimen in a corresponding amorphous substrate. This is because that the amorphous δ layers are grown at very low temperature where segregation is negligible and only broadening due to diffusion is present. Hence the amorphous R(z) should be qualitatively the same as the true crystalline one. We measured an amorphous boron δ layer in amorphous silicon grown at room temperature by molecular-beam epitaxy (MBE) with the range of E_p from 335 to 11 keV and fitted the measured profiles with the R(z) in Eq. (3). The primary ions used are normally incident oxygen. The obtained parameters are plotted against E_p in Fig. 1. Within the error of measurements, we have $\sigma^{\text{amph}}=0.86+0.27$ $E_p^{0.82}$, $\lambda_g^{\text{amph}}=0$ and λ_d^{amph} = 1.60 $E_p^{0.54}$, where the lengths are in nm and E_p in keV. This reveals that the SIMS mass transport effect itself has no contribution to λ_g^{amph} . We believe the same is true in crystalline case. Moreover, as the SIMS mass transport effect will be minimised at zero beam energy, it is reasonable to think that the λ_d^{amph} is only due to the SIMS effect and the residual σ^{amph} at $E_p = 0$ comes entirely from the boron amorphous δ layer itself. This is consistent with the well-known fact that a conserved diffusive point source has a Gaussian spatial distribution. Consequently, the SIMS contribution to the amorphous primitive deviation σ_s^{amph} can be obtained from $\sigma_s^{amph}(E_p)^2 = \sigma^{amph}(E_p)^2 - \sigma^{amph}(0)^2$.

Crystalline R(z) is built up by fitting the SIMS data of a MBE-grown boron δ layer in a crystalline silicon substrate with the range of E_p from 250 to 11 keV under the same experimental condition as for the amorphous study. Considering that the bonding energy of atoms in a crystalline material is usually larger than its amorphous counterpart, we expect that both σ^{cryst} and λ_d^{cryst} are smaller than the amorphous ones and λ_{p}^{cryst} should remain zero. The boron δ -layer structure grown by MBE normally has a segregated interface at the near surface side and an almost ideally abrupt interface at the other side. This will enable us to get a reliable λ_d^{cryst} from the fitting. For λ_g^{cryst} , the fit shows no significant energy dependence and we, therefore, take it as intrinsic to the sample. The results are also shown in Fig. 1. We obtain that $\sigma^{\text{cryst}} = 0.27 + 0.39 E_p^{0.75} \text{ and } \lambda_d^{\text{cryst}} = 1.39 E_p^{0.56}$. These parameters are indeed smaller than the amorphous ones, and again λ_d^{cryst} vanishes as E_p approaches zero. The σ^{cryst} can be partly affected by the sample structure in use and, indeed, shows a finite intercept. Therefore, we take the intercept away and calculate the SIMS part, σ_s^{cryst} , as $\sigma_s^{\text{cryst}}(E_p)^2$ $= \sigma^{\operatorname{cryst}}(E_n)^2 - \sigma^{\operatorname{cryst}}(0)^2.$



FIG. 1. The response function parameters, σ , λ_g , and λ_d , vs primary beam energy E_p . The open and solid symbols as well as the dashed and dotted lines are for the amorphous and crystalline parameters, respectively. The lines are fitted with form $a + bE_p^c$ for each of the parameters.



FIG. 2. (a) The SIMS depth profiles at different primary beam energy E_p for the boron δ layers in crystalline silicon substrate and (b) the corresponding spatial distributions reconstructed with the SIMS response function through a MaxEnt deconvolution.

Using the obtained parameters, we are now able to calculate sample-independent SIMS response functions for various E_p and use them with our noise model to deconvolve some measured SIMS depth profiles by MaxEnt method. A multiple boron δ -layer structure in silicon was grown at a constant 450 °C by MBE with the arrangement of, in turn from the surface, a pair of δ layers 2 nm apart, then another pair of δ layers 5 nm apart, etc. The intended boron concentration was 1×10^{20} atoms/cm³ for all the layers, which is well above the bulk solid solubility limit.^{19,20} We profiled the sample using a normally incident oxygen beam at E_p =0.25, 0.50, 1.0, 2.0, 4.0, and 6.0 keV, respectively. The depth calibration of profiles in silicon using oxygen primary ions has been discussed and clarified recently.²¹ In our case, since the part we study is outside the transient region and much longer than the transition depth we can align centroids of the profiles based on the principle that distance between centroids will not be changed by any convolution process. The MaxEnt algorithm used in our calculation is from Ref. 22. The measured profiles after conventional calibration and corresponding deconvolved spatial distributions for the first two pairs of boron δ layers are shown in Figs. 2(a) and 2(b), respectively.

From Figs. 2(a) and 2(b), we can see that although the original SIMS profiles appear quite differently, the reconstructed spatial distributions agree well with each other. The area under each feature is the same before and after deconvolution, i.e., the number of the boron atoms for each feature is conserved. It is obvious that the depth resolution of the

reconstructed spatial distributions has increased significantly. The δ layers at z = 87 nm which were separated by 2 nm can just be distinguished at $E_p = 2$ keV while the 5-nm pair at z= 135 nm are well separated at the maximum $E_p = 6$ keV we used. Note that the deconvolved features become smoother as E_p increases, i.e., better depth resolution can be achieved as E_p gets lower. From the reconstructed dopant distribution in Fig. 2(b), it is clearly shown that there is considerable boron segregation on the near surface side of the layers. This kind of feature would be automatically eliminated if a simplistic R(z) were used. There are some unexpected small concentration spikes in the reconstructed data at a level of two to three orders lower than the peak height. We have no objective criteria to confirm their existence or take them away so they remain whether one likes them or not.

Using Eqs. (1) and (2), we can easily work out the corresponding generalized Rayleigh limit of depth resolution Δz for two adjacent ideal δ layers after deconvolution if we assume that Δz is limited only by the SIMS noise:

$$\Delta z(E_p) \gg \frac{2}{Y_C^{1/2}} \lambda_d(E_p), \tag{4}$$

where we take $\lambda_g = 0$, $\lambda_d \gg \sigma_s$, and Y_C is the original measured counts. Estimating with the typical peak counts $Y_C \sim 10^4$ in our experiment and the λ_d from Fig. 1, we find the resolution we have achieved in the above deconvolution is within the limit.

Maximizing the entropy of spatial distribution gives us the most likely deconvolved solution. It has a *global* tendency to spread the solution onto the whole space range within a given noise deviation when total concentration is fixed. However, there is no constraint on the local changes of the distribution. For example, two spikes in the distribution contribute the same entropy no matter whether they are next to each other or not. This, unlike local constrains such as limiting the derivative of spatial distribution, makes possible the reconstruction of some abrupt features, e.g., δ layers, superlattice structures, step doping materials, etc.

To have some further understanding on the MaxEnt deconvolution method, we compare the frequency components of the SIMS measured profile and the corresponding convolved one calculated from the reconstructed distribution. Figure 3 shows the power spectra for the $E_p = 0.25$ keV case. Although the MaxEnt method only manipulates data in real space, the background noise frequency components in the convolved profile are clearly suppressed by 6-7 orders. It is clearly shown that the high-frequency features in the range of $0.75-1.50 \text{ nm}^{-1}$ are retained rather than eliminated if a conventional 1/f noise subtraction method were used. Profiles of other E_p have the similar results. This is consistent with Shannon's entropy loss theorem for signal passed through linear filters.²³ Consequently, the MaxEnt method is not only able to reduce the background noise level without using any artificial windows in real space or filters in frequency space but also capable of retaining sharp features that are statistically significant.

We should emphasis that although the MaxEnt method does improve the depth resolution and recovery sharp features, it *only* provides us with the statistically evident information which the original SIMS profile contains. The Max-



FIG. 3. The Fourier power spectrum $|Y(k)|^2$ of the measured SIMS signal Y(z) vs wave number k. The data are calculated from the 0.25-keV SIMS boron profile and the convolved one from the corresponding reconstructed distribution.

Ent deconvolution is not an alternative to instrumental improvement, such as achieving lower primary beam energy and higher corresponding beam current. For example, one cannot distinguish whether there are two close δ layers or a single δ layer of higher concentration if the profile is measured with very high beam energies. This is because highenergy profiling will not only have large λ_d but also lead to large depth increment and low sampling density, which will

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limit the information in the data at the first place. To reduce the depth increment for higher sampling density in this situation requires significant reductions in the primary beam current because of the high sputter yield at high beam energy. This will usually lead to lower values for Y_C . Therefore, the signal-noise ratio will decrease, which limits the potential increase of depth resolution from the deconvolution.

We describe a procedure to obtain sample-independent response function which is suitable for use even in the situation where the intrinsic SIMS effect is so small that samplerelated features can be seen. Using this together with an empirically determined noise model, MaxEnt deconvolution is performed to reconstruct self-consistent dopant distributions from the SIMS depth profiles obtained at different beam energies. There are no adjustable parameters involved in obtaining the response function and deconvolving the profiles. The depth resolution of the reconstructed distributions has been greatly improved and the segregation on the near surface sides is clearly demonstrated. With reconstructed true spatial distributions, further quantitative investigations on interface segregation, atomic diffusion, and related problems will be straightforward. Our method can be used to study other dopants, and the MaxEnt formalism may be extended to models other than convolution.

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