Cellular Automata Simulation of Site-saturated and Constant Nucleation Rate Transformations in Three Dimensions

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Microstructural evolution in three dimensions of nucleation and growth transformations is simulated by means of cellular automata. Two types of nucleation are considered: site-saturated nucleation and constant nucleation rate. The simulated microstrutural evolution agrees very well with exact analytical expressions. The simulated data also gives very good agreement with expressions derived to describe the evolution of the interfaces between transformed grains.

Keywords: microstructure, kinetics, recrystallization, computer simulation, cellular automata

1. Introduction

Formal kinetics is a powerful technique to model nucleation and growth solid state transformations. In these transformations, a parent phase, α , also called "matrix" or untransformed region", transforms into a product phase, β , also called "transformed region". In this approach, nucleation and growth are regarded as purely "operational" concepts meaning that no assumption is made regarding their mechanism but only their geometrical and kinematical characteristics. Recrystallization kinetics in particular has been mainly analyzed with the help of formal kinetics methodology!-4.

This approach has two facets: one experimental another theoretical. The experimental facet specifies the microstrutural descriptors that may be experimentally obtained, preferentially from planar sections, such as: volume fraction transformed, V_{ν} ; interface area between transformed and untransformed regions, S_{ν} , and interface area between transformed regions, $S_{\nu\beta}$. The theoretical facet aims at developing expressions that involve these measurable quantities. Today, one attempts not to rely on three-dimensional quantities that cannot be measured by standard stereological methods but require tedious and time consuming serial sectioning. With the advent of 3DXRD techniques experimental determination of three-dimensional quantities like the number of nuclei per unit of volume, N_{ν} , or even nuclei location in space might well become viable on a routine basis.

The basis of formal kinetic modeling is the early work of Johnson-Mehl⁷, Avrami⁸⁻¹⁰ and Kolmogorov¹¹ (JMAK), which used only a single microstructural descriptor, V_V . JMAK's work was subsequently extended by DeHoff¹², who proposed the use of an additional microstructural descriptor, S_V , and the associated concept of microstructural path. Vandermeer and coworkers¹³ extended DeHoff's microstructural path concept and crystallized it in an all round theoretical treatment covering variable nucleation and growth rates as well as non-spherical regions.

Nevertheless, even general analytical treatments have significant limitations owing to unavoidable simplifying assumptions. For instance, Vandermeer et al.¹³ assume that nuclei are randomly located in space. However, when one considers, for example, austenite to ferrite transformation, ferrite nucleates at grain boundaries^{14,15}. Another example is recrystallization, where the deformed state is,

as a rule, highly heterogeneous ¹⁶ casting some doubt on the assumption that nuclei are randomly located in space. Relaxing this and other assumptions may lead to situations too complex for analytical methods. In this regard, computer simulation of microstructural evolution can be an invaluable tool since it can simulate situations beyond analytical treatment. Compared to other simulation methods, cellular automata¹⁷⁻³⁰ is a convenient choice for phase transformations in general and recrystallization in particular²³.

Computer simulation aims at describing transformation in complex situations for which there is no analytical theory. In order to accomplish this, one must first make sure that the simulation is able to describe accurately the simplest cases for which *there are* exact mathematical descriptions. Interestingly, this basic or minimum validation of the simulation has been rarely reported in published papers²³.

In previous papers^{28,29}, present authors carried out cellular automata (CA) simulation of phase transformation/recrystallization in two dimensions (2D). Earliest computer simulations of microstructural evolution were usually performed in two dimensions. Even in recent papers some authors have chosen to work in two dimensions probably because the computer cost of three-dimensional (3D) simulations is significantly higher than that of 2D simulations. Although there is a certain interest in the study of two-dimensional transformations, for example, transformations taking place in thin films or thin sheets, the greatest interest lye in three dimensional transformations.

The purpose of this paper is to simulate phase transformations/ recrystallization in 3D by cellular automata (CA). Two basic nucleation types were assumed:

- a) site-saturation: all nucleation occurs at the beginning of the transformation; and
- b) constant nucleation rate: the number of new nuclei per unit of time per unit of volume is constant.

Simulated microstrutural evolution was compared in detail with predictions of mathematically exact analytical theories²⁸. Moreover, simulated data was used to test new expressions recently derived by Rios et al.³¹ to describe the evolution of the interfaces between transformed grains as a function of time or volume fraction transformed.

2. Description of the Cellular Automata Simulation

Cellular automata simulation used a 3D von Neumann neighborhood¹⁷. The matrix consisted of a cubic lattice with 304 x 304 x 304 cells. One cell edge was considered to have a unit length and consequently of a single cell had unit volume. Units of all quantities reported here follow from this. The nuclei were randomly dispersed on the matrix. For site-saturated nucleation, the number of nuclei per unit of volume, N_v , was equal to $125/304^3$. For constant nucleation rate, the nucleation rate per unit of time per unit of volume, I, was equal to $2/304^3$. This nucleation rate resulted in 141 nuclei per unit of volume when the transformation reached $V_v = 1$.

The simulation produced a sequence of matrices as a function of time. Time is discrete in CA, it takes integer values starting from t = 0. One time unit corresponds to the interval between two consecutive matrix updates¹⁷. From the simulated matrices, all the desired quantities could be extracted. Oliveira³⁰ gives further details of the present simulation.

3. Growth of a Single Grain

When a single grain grows in isolation, it is possible to find analytical expressions for its growth. The following expressions are specific for the 3D von Neumann neighborhood adopted in this simulation. The time origin, t=0, corresponds to a volume fraction equal to zero, $V_v=0$. Therefore, the volume of a single grain, v, and its area, s, at t=0 are equal to:

$$v(0) = 0 \tag{1a}$$

$$s(0) = 0 \tag{1b}$$

For $t \ge 1$ the volume of a single grain as a function of time is:

$$v(t) = \frac{4}{3}t^3 - 2t^2 + \frac{8}{3}t - 1 \tag{2}$$

This equation is not valid for t = 0. Neglecting the lower order powers of time:

$$v(t) \cong \frac{4}{3}t^3$$
 units of volume (3)

The area of a single grain, s, for $t \ge 1$ is:

$$s(t) = 12t^2 - 12t + 6 (4)$$

Again, neglecting early time behavior:

$$s(t) \cong 12t^2$$
 area units (5)

Equations 3 and 5 will be used below in section 4 to represent the growth of a single grain.

The average interface velocity, $\langle G \rangle$, can be calculated with the help of Cahn-Hagel³² equation:

$$\langle G \rangle = \frac{\iint G ds}{\iint ds} = \frac{1}{s} \frac{dv}{dt}$$
 (6)

Inserting Equations 3 and 5 gives:

$$\langle G \rangle \cong \frac{1}{3} \text{ units of length/unit time}$$
 (7)

The units will be omitted from now on.

The average interface velocity is constant. Therefore, for a single grain the simulation was able to produce constant growth rate.

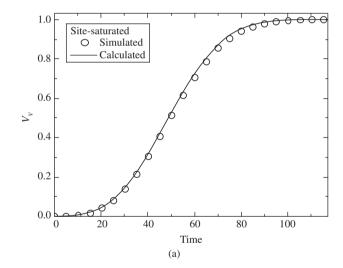
It is worthy mentioning that the volume and the area of a single grain showed time dependencies different from what would be expected from constant velocity growth. In this case it was possible to use only the leading term, with the highest power of time. A similar situation was found in a previous work²⁸ on 2D cellular automata

simulation. It is not easy to say what effect a different choice of neighborhood or a different transition criterion would have on this time dependence. So, CA simulations should check this. This is rarely reported in published papers on this subject¹⁷⁻³⁰.

4. Microstructural Evolution: Volume Fraction, Interface Area Per Unit of Volume and Microstructural Path

In this section, the expressions derived in the previous section for single grain evolution were used to obtain expressions containing "global" measurements, that is, average measurements such as volume fraction and interface area per unit of volume. The calculated expressions were compared with the results from the simulation. This section mainly deals with quantities related to the mobile interface: the interface between transformed and untransformed regions. In each case, an analytical expression was derived for site-saturation nucleation and constant nucleation rate. Section 5 will compare simulations results with new expressions derived for the evolution of the interfaces between transformed regions.

Figures 1-3 show the results obtained from the simulation compared with the theoretical expressions obtained in this section. One



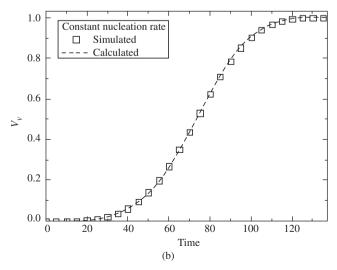


Figure 1. Volume fraction, V_{y} , against time. There is good agreement between theory and simulation: a) site-saturated; and b) constant nucleation rate.

cannot directly compare Figures 1a-3a with Figures 1b-3b because the number of nuclei per unit of volume is different in each case. In order to avoid overloading the figures not all simulation points were plotted.

4.1. Volume fraction as a function of time

The extended volume fraction, $V_{\it VE}$, is the sum of the volume of all individual grains supposing that they grow without impingement. The subscript "E" is reserved for "extended quantities" defined in a manner similar to the extended volume fraction. Using Equation 3 one obtains:

$$V_{VE} = \frac{4}{3} N_V t^3 \tag{8a}$$

$$V_{VE} = \frac{1}{3} I t^4$$
 (8b)

where N_{V} is the number of nuclei per unit of volume and I is the nucleation rate per unit of volume; Equations 8a and 8b correspond to site-saturated nucleation and constant nucleation rate, respectively.

For randomly located nuclei, the extended volume fraction, $V_{\rm VE}$, and the real volume fraction, $V_{\rm V}$, are related by the JMAK expression:

$$V_{v} = 1 - \exp(-V_{vx}) \tag{9}$$

As a result:

$$V_{\nu} = 1 - \exp\left(-\frac{4}{3} N_{\nu} t^{3}\right)$$
 (10a)

$$V_{\nu} = 1 - \exp\left(-\frac{1}{3}H^4\right) \tag{10b}$$

where Equations 10a and 10b correspond to site-saturated nucleation and constant nucleation rate, respectively.

Figures 1a and 1b show a comparison between $V_{\scriptscriptstyle V}$ calculated from the analytical solution and obtained from the CA simulation. The agreement is very good.

4.2. Interface area between transformed and untransformed regions as a function of time and volume fraction

The extended interface area per unit of volume between transformed grains and untransformed regions, S_{vE} , is the sum of the interface area of all individual grains supposing that they grow without impingement. Using Equation 5 it can be shown that:

$$S_{VE} = 12N_V t^2 \tag{11a}$$

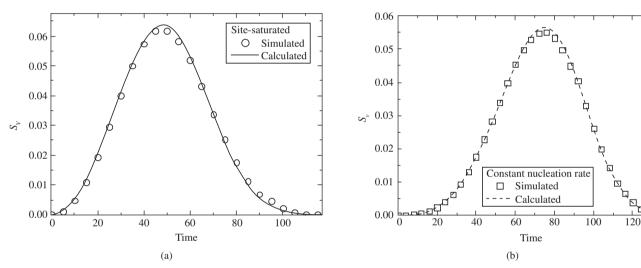


Figure 2. Interface area between transformed and untransformed region per unit of volume, S_{v} , against time. There is good agreement between theory and simulation: a) site-saturated; and b) constant nucleation rate.

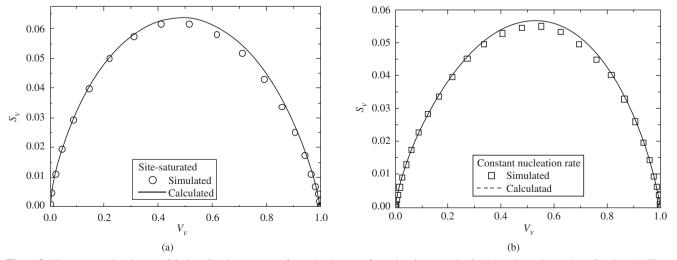


Figure 3. Microstructural path: area of the interface between transformed and untransformed region per unit of volume, S_v , against volume fraction, V_v . There is good agreement between theory and simulation: a) site-saturated; and b) constant nucleation rate.

$$S_{VE} = 4It^3 \tag{11b}$$

where Equations 11a and 11b correspond to site-saturated nucleation and constant nucleation rate, respectively.

The extended interface area and real interface area, S_{ν} , are related by the DeHoff's 12 expression:

$$S_{v} = (1 - V_{v}) S_{vF} \tag{12}$$

Equation 12 together with Equations 1a and 1b give:

$$S_{\nu} = 12N_{\nu}t^{2}\exp\left(-\frac{4}{3}N_{\nu}t^{3}\right)$$
 (13a)

$$S_{v} = 4It^{3} \exp\left(-\frac{1}{3}It^{4}\right) \tag{13b}$$

where Equation 13a and 13b correspond to site-saturated nucleation and constant nucleation rate, respectively.

Figures 2a and 2b show a comparison between the S_{ν} , calculated from the analytical solution, Equations 13a-b, and obtained from the CA simulation. The agreement is very good.

In three dimensions the microstructural path can be seen as a curve on the (V_V, S_V) plane. From Equations 8a,b and 11a,b one can find a relationship between S_{VE} and V_{VE} :

$$S_{VE} = 3 \left(36N_V\right)^{\frac{1}{3}} V_{VE}^{\frac{2}{3}} \tag{14a}$$

$$S_{vE} = 4 (27I)^{\frac{1}{4}} V_{vE}^{\frac{3}{4}} \tag{14b}$$

where Equation 14a and 14b correspond to site-saturated nucleation and constant nucleation rate, respectively. From Equations 9, 12 and Equations 14 one can obtain S_v as a function of volume fraction transformed:

$$S_{\nu} = 3 (36N_{\nu})^{\frac{1}{3}} (1 - V_{\nu}) \left[\ln \left(\frac{1}{1 - V_{\nu}} \right) \right]^{\frac{2}{3}}$$
 (15a)

$$S_{\nu} = 4 (27I_0)^{\frac{1}{4}} (1 - V_{\nu}) \left[\ln \left(\frac{1}{1 - V_{\nu}} \right) \right]^{\frac{3}{4}}$$
 (15b)

where Equation 15a and 15b correspond to site-saturated nucleation and constant nucleation rate, respectively.

Figures 3a and 3b show a comparison between the values calculated using the theoretical expression for the microstructural path obtained above and those obtained from the simulation. The agreement is very good.

The average grain boundary velocity is given by³²:

$$\langle G \rangle = \frac{1}{s} \frac{dv}{dt} = \frac{1}{S_{vE}} \frac{dV_{vE}}{dt} = \frac{1}{S_v} \frac{dV_v}{dt} = \frac{1}{3}$$
 (16)

The grain boundary velocity calculated from the simulated values was 0.34 + 0.01, very close to the theoretical prediction for both site-saturated nucleation and constant nucleation rate. This means that the interface velocity in real space is the same as that in extended space, as expected.

In summary CA simulation showed very good agreement with exact mathematical expressions.

5. Comparison of the Simulation with New Analytical Expressions for the Impinged Interface Area Per Unit of Volume, Total Interface Area Per Unit of Volume and the Contiguity

There are two kinds of interface during an heterogeneous phase transformation: mobile and immobile. The mobile interface is the interface between the transformed region and the untransformed matrix. This interface migrates and its migration determines the transformation progress. The immobile interface is a consequence of impingement of transformed grains. The transformed grain size is a function of the interface area between transformed grains when the transformation reaches $V_{\nu} = 1$.

Traditionally, experimental and theoretical studies focus on the mobile interface. This could be seen in the previous section where all equations involved only S_V . However, the amount of interface between transformed grains, $S_{V\beta}$, is an important quantity and contains valuable information on transformation evolution. CA simulation is able to provide accurate data on these interfaces and so gives a good basis for comparison with analytical models. Combining $S_{V\beta}$ with S_V two useful quantities may be defined. One is the total interface area per unit of volume, S_{VT} :

$$S_{VT} = S_V + 2S_{VB} \tag{17}$$

Another important quantity, derived from $S_{\nu\beta}$ with S_{ν} is the contiguity parameter, C_{β} . The contiguity is the ratio of immobile interface area to the total interface area and is quite sensitive to deviations of nuclei location from randomness, as recently shown by Vandermeer and Jensen^{5,33}; see also Rios et al.²⁹. It can be defined as⁵:

$$C_{\beta} = \frac{2S_{\nu\beta}}{S_{\nu} + 2S_{\nu\sigma}} \tag{18}$$

Recently Rios et al. 31 have derived analytical expressions for $S_{V\beta}$ as a function of volume fraction transformed for time dependent nucleation rate and time dependent velocity. They followed up the earlier work by Meijring 34 , Gokhale 35,36 and more recently an elegant mathematical derivation by Tong et al. 37 . All these authors solved this problem for spherical growth. Rios et al. 31 derivation has the advantage of being mathematically simpler and not restricted to spherical growing regions. The disadvantage of their method is that, as will be shown below, they do not obtain an explicit expression for $S_{V\beta F}$, the value of $S_{V\beta}$ for a fully transformed matrix. Their expressions can be used to obtain analytical expressions for S_{VT} and C_{β} . Rios et al. 31 derivation is repeated here for convenience. The starting point of their derivation was DeHoff's relationship between the extended and the real interface area, Equation 12, rewritten below in a slightly different way:

$$S_{V}(V_{V}, S_{VF}) = S_{VF}(1 - V_{V})$$
(19)

One can consider that a small change in S_v , δS_v , can be seen as a sum of changes in each variable, δV_v and δS_{vE} :

$$\delta S_{v} \left(\frac{\partial S_{v}}{\partial S_{vE}} \right)_{v_{e}} \delta S_{vE} + \left(\frac{\partial S_{v}}{\partial V_{v}} \right)_{S_{ve}} \delta V_{v}$$
(20)

Or

$$\delta S_{v} = (1 - V_{v}) \, \delta S_{vF} - (S_{vF}) \, \delta V_{v} \tag{21}$$

The above has two terms:

- a) The first is always positive, it corresponds to the increase in S_v when the interface length increases δS_E in extended space, at a constant V_v , corrected for impingement by the factor $(1 V_v)$; and
- b) The second term, $S_{VE} \delta V_V$, is negative because S_{VE} and δV_V are always positive. It corresponds to the decrease in S_V , when the area fraction increases for a constant S_{VE} , this decrease corresponds to a "loss" in S_V due to the "creation" of new impinged interface. Two extended interfaces meet to make one impinged interface, S_{VB} , so:

$$\frac{dS_{V\beta}}{dV_{\nu}} = \alpha \frac{S_E}{2} \tag{22}$$

where α is a normalizing factor to be determined below.

Integrating Equation 22 gives:

$$S_{\nu\beta} = \frac{\alpha}{2} \int_0^{\nu_{\nu}} S_{\nu E} dV_{\nu} \tag{23}$$

It is more convenient to write Equation 23 in terms of extended quantities with the help of a relationship obtained from Equation 9:

$$dV_{V} = \exp(-V_{VF})dV_{VF} \tag{24}$$

where V_{VE} , the extended volume fraction, is the sum of the volume of all growing grains divided by the total volume assuming that the growing grains do not impinge. Equation 23 then becomes:

$$S_{V\beta} = \frac{\alpha}{2} \int_{0}^{V_E} S_{VE} \exp(-V_{VE}) dV_{VE}$$
 (25)

In order to integrate Equation 25 it is necessary to know the relationship between S_{VF} and V_{VF} which is supposed to be of the form¹³:

$$S_{VF} = C(V_{VF})^q \tag{26}$$

Resulting in:

$$S_{\nu\beta} = \frac{\alpha C}{2} \int_{0}^{V_{\nu E}} (V_{\nu E})^{q} \exp(-V_{\nu E}) dV_{\nu E}$$
 (27)

This integral can be written in terms of the incomplete Gamma function, Γ :

$$S_{V\beta} = \frac{\alpha C}{2} \left(\Gamma \left(q + 1 \right) - \Gamma \left(q + 1, V_{VE} \right) \right) \tag{28}$$

The factor α can be found by taking the limit $V_{VE} \to \infty$. For $V_{VE} \to \infty$, $V_V \to 1$, the specimen is fully transformed and $S_{V\beta}$ tends to a constant value $S_{V\beta F}$:

$$\alpha = \frac{2S_{V\beta F}}{C\Gamma(\alpha + 1)} \tag{29}$$

Inserting Equation 29 into 26:

$$\frac{S_{\nu\beta}}{S_{\nu\beta F}} = \left(1 - \frac{\Gamma(q+1, V_{\nu E})}{\Gamma(q+1)}\right) =$$

$$\left(1 - \frac{\Gamma(q+1, -\ln(1-V_{\nu}))}{\Gamma(q+1)}\right)$$
(30)

where $S_{V\beta F}$ is the amount of interface area between transformed regions when the transformation has reached $V_V = 1$. Using the definition of contiguity, Equation 18:

$$C_{\beta} = \frac{2S_{V\beta F}\left(1 - \frac{\Gamma\left(q+1, \ln\left(\frac{1}{1-V_{V}}\right)\right)}{\Gamma\left(q+1\right)}\right)}{C\left(1-V_{V}\right)\left(\ln\left(\frac{1}{1-V_{V}}\right)\right)^{q} + 2S_{V\beta F}\left(1 - \frac{\Gamma\left(q+1, \ln\left(\frac{1}{1-V_{V}}\right)\right)}{\Gamma\left(q+1\right)}\right)}$$
(31)

For the cases considered here, C and q can be found from the relationships between S_{VE} and V_{VE} given by Equations 14a-b:

- a) site-saturation: q = 2/3 and $C = 3(36N_v)^{1/3}$; and
- b) constant nucleation rate: q = 3/4 and $C = 4(27I)^{1/4}$.

More general expressions for C and q can be found in Rios et al.³¹.

Equations 30 and 31 can be written as a function of time with the help of Equations 10a and b. The total interface area per unit of volume, S_{VT} , can be found by means of Equations 15a,b and Equations 30 and Equation 17.

Figures 4-7 show the comparison of the values calculated by means of Equations 17, 30 and 31 and those obtained from the simulation. The agreement is very good. For $S_{\nu\beta}$ and $S_{\nu T}$, Figures 4-7, the normalized quantities $S_{\nu\beta}/S_{\nu\beta}$ and $S_{\nu T}/(2S_{\nu\beta})$ were plotted. In order to avoid overloading the figures not all simulation points were plotted.

An interesting aspect of the present comparison is that the growing regions were not spherical in this simulation. This shows that Rios et al.³¹ expression applies when the shape of the growing regions is not spherical³⁴⁻³⁶.

Figures 5c and 7c show that the normalized quantities $S_{V\beta}/S_{V\beta F}$ and C_{β} are practically the same for both nucleation types. Randomness in nuclei distribution might have an underlying effect that is stronger than the details of the kinetics³¹.

It is important to mention that no adjustable parameters were used in this work. Only exact analytical solutions were compared with the simulations. The good agreement between theory and simulation shows that CA is able to reproduce situations for which there are mathematically exact solutions. This good agreement suggests that CA may be applied to complex situations for which there are no analytical solutions.

6. Summary and Conclusions

First it is worthy mentioning that no adjustable parameters were used throughout this work. Moreover, this work was motivated by recrystallization studies and it is perhaps more useful in that context.

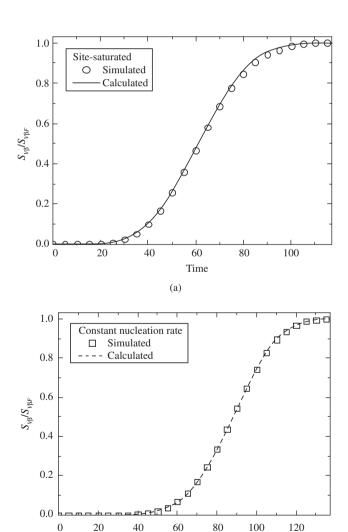
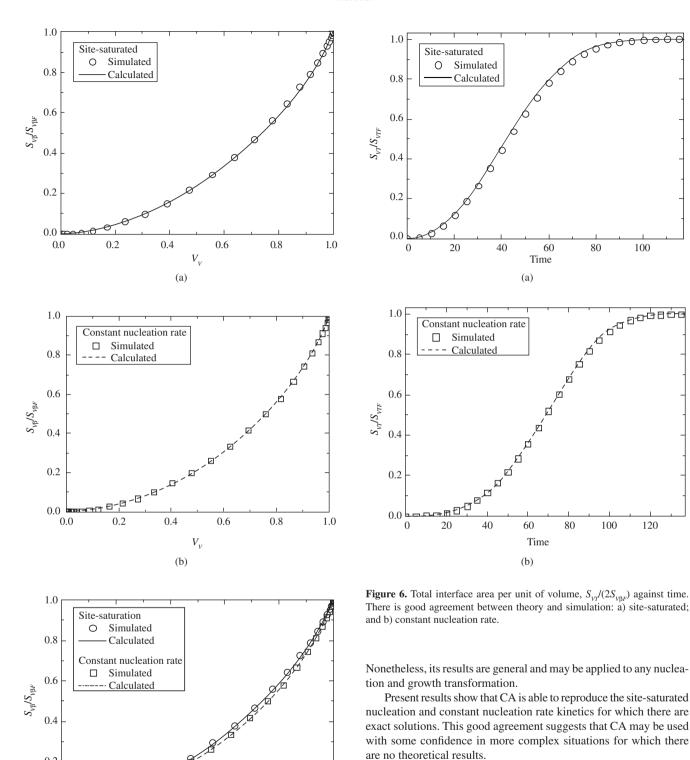


Figure 4. Area of the interface between transformed and transformed region per unit of volume, $S_{V\beta}/S_{V\beta,F}$, against time. There is good agreement between theory and simulation: a) site-saturated; and b) constant nucleation rate.

(b)

Time



interface between the transformed regions. The good agreement 0.0 0.0 0.2 0.8 0.6 1.0 0.4 observed is a reconfirmation of the validity of these new expressions in the situations examined here. (c) Acknowledgments

Figure 5. Area of the interface between transformed and transformed region per unit of volume, S_{vR}/S_{vRF} , against volume fraction. There is good agreement between theory and simulation: a) site-saturated; b) constant nucleation rate; and c) Figures 5a and 5b plotted together; S_{VB}/S_{VBF} as a function of volume fraction is practically the same for both nucleation types.

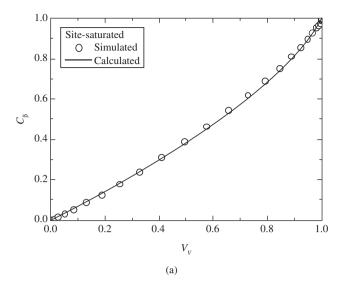
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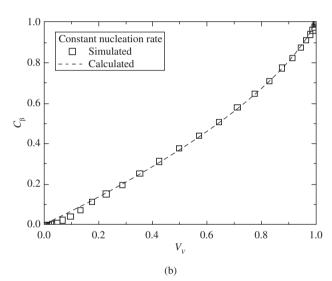
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CA simulation showed to be useful to test expressions developed by Rios et al.31 for the evolution of area per unit of volume of the

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120





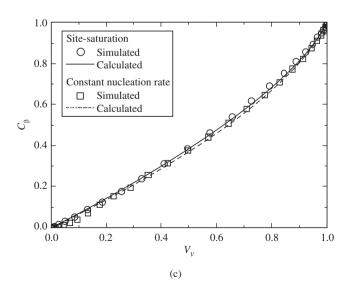


Figure 7. Contiguity, C_{β} , against volume fraction. There is good agreement between theory and simulation: a) site-saturated nucleation; b) constant nucleation rate; and c) Figures 7a and 7b plotted together; C_{β} as a function of volume fraction is practically the same for both nucleation types.

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