Inhomogeneous Poisson Point Process Nucleation: Comparison of Analytical Solution with Cellular Automata Simulation

Paulo Rangel Rios^a*, Douglas Jardim^a, Weslley Luiz da Silva Assis^a,

Tatiana Caneda Salazar^a, Elena Villa^b

^aEscola de Engenharia Industrial Metalúrgica de Volta Redonda, Universidade Federal Fluminense, Av. dos Trabalhadores, 420, 27255-125 Volta Redonda - RJ, Brasil ^bDepartment of Mathematics, University of Milan, via Saldini 50, 20133 Milano, Italy

Received: April 10, 2009; Revised: May 6, 2009

Microstructural evolution in three dimensions of nucleation and growth transformations is simulated by means of cellular automata (CA). In the simulation, nuclei are located in space according to a heterogeneous Poisson point processes. The simulation is compared with exact analytical solution recently obtained by Rios and Villa supposing that the intensity is a harmonic function of the spatial coordinate. The simulated data gives very good agreement with the analytical solution provided that the correct shape factor for the growing CA grains is used. This good agreement is auspicious because the analytical expressions were derived and thus are exact only if the shape of the growing regions is spherical.

Keywords: microstructure, analytical methods, phase transformations, recrystallization, cellular automata

1. Introduction

Transformation kinetics is often described by an expression of the form:

$$\mathbf{V}_{\mathbf{V}}(t) = 1 - \exp(-kt^n) \tag{1}$$

where, $\mathbf{V}_{\mathbf{v}}$ is the volume fraction, *t* is time, *k* and *n* are adjustable parameters. Equation 1 is often called "Avrami" equation. In fact, the use of adjustable parameters is in some sense the opposite that Kolmogorov¹, Johnson-Mehl² and Avrami³⁻⁵ (KJMA) did. KJMA obtained an exact solution to the problem of nucleation and growth of a new phase. Their solution is not general. They assumed uniform randomly nucleation, either site-saturated or to taking place with a constant nucleation rate. Site-saturated nucleation implies that all nucleation sites are exhausted early in the transformation, or more informally, all nuclei are already present at the time origin, *t* = 0. Moreover, they further assumed the growth rate, *G*, to be constant and the growing regions to have a spherical shape. They solved the problem of impingement and obtained an exact expression connecting $\mathbf{V}_{\mathbf{v}}$ to the extended volume fraction, $\mathbf{V}_{\mathbf{F}}$

$$\mathbf{V}_{\mathbf{V}} = 1 - \exp(-\mathbf{V}_{\mathbf{E}}) \tag{2}$$

Specifically, their exact analytical expressions for site-saturated nucleation is the well-known expression:

$$\mathbf{V}_{\mathbf{V}} = 1 - \exp\left(-\frac{4}{3}\pi N_V G^3 t^3\right) \tag{3}$$

where N_v is the number of nuclei per unit of volume.

Their theory, and its developments and extensions^{6,7} can be applied to a diversity of transformations in both metallic and nonmetallic materials. Such transformations include austenite to pearlite transformation², recrystallization⁸, abnormal grain growth in^{9,10} BaTiO₃, martensite spread^{11,12} and polymer crystallization^{12,14}. Thus, KJMA theory is useful even when nucleation and growth retain solely an operational meaning.

In a recent work, Rios and Villa¹⁴ resorted to recent developments on stochastic geometry¹⁵⁻¹⁷ to revisit the classical KJMA theory and generalize it for situations in which nucleation took place both for homogeneous and for inhomogeneous Poisson point processes. "Nuclei being located in space according to a homogeneous Poisson point process" simply means that they are uniform randomly located in space. Poisson point processes will be more rigorously defined in section 2 below.

For that matter, $\mathbf{V}_{\mathbf{v}}$ and $\mathbf{V}_{\mathbf{E}}$ were replaced by a position-dependent mean volume density, $V_{V}(t, x)$, mean extended volume density $V_{E}(t, x)$, where $x = (x_1, x_2, x_3)$ is the spatial coordinate. Thus, the relationship between volume fraction and extended volume fraction, Equation 1, became a relationship between the mean volume density and the mean extended volume density:

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$$V_V(t,x) = 1 - \exp(-V_E(t,x)) \tag{4}$$

Using this methodology, it was possible to significantly broaden the scope of analytical solutions available. One example of the results obtained by Rios and Villa is an equation for site-saturated nucleation, which closely parallels Equation 3,

$$V_V(t,x) = 1 - \exp\left(-\frac{4}{3}\pi\lambda(x)G^3t^3\right)$$
(5)

where $\lambda(x)$ is called the *intensity* of the inhomogeneous Poisson point process. For Equation 5 to be valid $\lambda(x)$ must be a harmonic function, i.e must satisfy Laplace's partial differential equation. For the particular case of uniform randomly located nuclei or synonymously nuclei located according to a homogeneous Poisson point process the intensity is independent of the position and coincides with the number of nuclei per unit of volume. Further distinction between homogeneous and inhomogeneous Poisson point process and the definition of $V_{v}(t, x)$ and $V_{r}(t, x)$ are clarified in the Mathematical background section below.

In previous papers¹⁸⁻²⁴, cellular automata (CA) simulation of phase transformation/recrystallization in 2-d and 3-d were carried out and compared with KJMA analytical solutions. The purpose of the present paper is to simulate phase transformations/recrystallization in by cellular automata (CA) in order to compare the simulation with the analytical solution by Rios and Villa. The simulation is restricted to site-saturated inhomogeneous Poisson point process nucleation with the intensity, $\lambda(x)$, varying linearly along the x, coordinate but remaining constant for x_2 and x_3 . In order to make such a comparison feasible it is necessary to adapt Equation 5, derived for spherical growth, to CA where growth is not spherical.

Thus, this work presents not only a comparison but in fact it shows that it is possible to use Equation 5 to describe CA even though Equation 5 is not exact for CA, as will become clear in what follows.

Noteworthy, to our best knowledge, this is the first time a computer simulation is carried out in which nuclei are located in space according to an inhomogeneous Poisson process.

2. Mathematical Background

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A rigorous definition of $V_{v}(t, x)$ and of point processes entail a minimum background in stochastic geometry. Nonetheless, for the present work an intuitive notion of those suffices. Such may be apprehended from Figure 1a and 1b, in which a homogeneous and an inhomogeneous Poisson point process were simulated with the help of the "R" software²⁷. Details that are more mathematical can be found in Rios and Villa¹⁴.

Figure 1a and 1b show nuclei, i.e. points, represented by small circles located according to a homogeneous and a heterogeneous Poisson point process, respectively. The word "process" does not necessarily

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implies time dependence. In fact Figure 1a and 1b represent sitesaturated nucleation, or, informally, the situation in which all nuclei are already present at t = 0. Figure 1a depicts nuclei located in what may be intuitively described as "randomly" or uniform randomly. Conversely in Figure 1b the number of nuclei changes from $x_1 = 0$ to $x_1 = 1, x_1$ -horizontal axis, following $\lambda(x) = 350 x_1 + 25$. Therefore, in this case, as x_1 changes from 0 to 1 the number of nuclei increases.

More generally, when A is a 3-d region, N(A), is the random number of points/nuclei located within A. If these are Poissonian distributed:

$$P(N(A) = k) = \frac{\lambda^{k}(x)}{k!} \exp(-\lambda(x))$$
(6)

where, P(N(A) = k) is the probability of k nuclei falling within the unit volume A and $\lambda(x)$ is the intensity. In the homogeneous case, the intensity is independent of the spatial coordinate, x. In contrast, in the inhomogeneous case the intensity does depend on the spatial coordinate, $\lambda(x)$, $\lambda(x)$ can be interpreted as the mean number of nuclei in the infinitesimal volume dx. Notice that in the homogeneous case λ is constant; therefore in such a case it represents the mean number of nuclei per unit of volume.

For the volume transformed originating from grains that grow from heterogeneously located nuclei, one cannot talk of a volume fraction as was done above. It is necessary to define a mean volume *density*, which takes different values depending on x: $V_{y}(t, x)$ and $V_{\rm F}(t, x)$ corresponding to $V_{\rm V}(t)$ and $V_{\rm E}(t)$, respectively. Of course, one can still find the volume fraction by integration of the mean volume density but now it depends on where the integration volume is located within the specimen. Notice that for a "homogeneous specimen" the volume fraction, $\mathbf{V}_{\mathbf{v}}(t)$, represents the probability that an arbitrary point will be transformed regardless of the location of this point in space. For a "heterogeneous specimen" this probability, the mean volume density, $V_{v}(t,x)$, depends on the spatial coordinates of the point considered.

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homogeneous process. Simulations were performed using the R software.



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3. Description of the Cellular Automata Simulation

Cellular automata simulation used a 3-d von Neumann neighborhood^{26,27}. The matrix consisted of a cubic lattice with 300 x 300 x 300 cells. One cell edge was considered to have a length equal to 1/300 so that the simulated domain is effectively a unit cube: [0,1] x [0,1] x [0,1]. Units of all quantities reported here follow from this. Site saturated nucleation was assumed. The nuclei were located in the matrix according to an inhomogeneous Poisson point process with an intensity equal to $\lambda(x) = \lambda(x_x, x_2, x_3) = 596 x_1 + 2$. Consequently, 300 nuclei were present at the simulated volume.

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 updates^{26,27}. From the simulated matrices, all the desired quantities could be extracted.

4. Analytical Expressions

4.1. Exact solutions for spherical growth

The exact analytical solutions from Rios and Villa¹⁴ for sitesaturated nucleation according to an inhomogeneous Poisson process and constant growth velocity, *G*, are repeated here for convenience. Notice that in what follows $\alpha = \pi$.

$$V_V(t,x) = 1 - \exp(-\frac{4\alpha}{3}\lambda(x)G^3t^3)$$
(7)
and

$$S_V(t,x) = 12\alpha\lambda(x)G^2t^2\exp(-\frac{4\alpha}{3}\lambda(x)G^3t^3)$$

 $S_v(t, x)$ is the mean interfacial area density between the new phase and the parent matrix. The microstructural path is:

$$S_V(t,x) = 3\left(36\alpha\lambda(x)\right)^{\frac{1}{3}} \left(1 - V_V(t,x)\right) \left[\ln\left(\frac{1}{1 - V_V(t,x)}\right)\right]^{\frac{2}{3}}$$
(9)

If

$$\lambda(x) = mx_1 + n \tag{10}$$

and supposing $\lambda(x) > 0$ and that x belongs to the unit cube, $A = [0,1] \times [0,1] \times [0,1] = [0,1]^3$, then using Equation 7.

$$\mathbf{V}_{\mathbf{V}}\left(t,[0,1]^{3}\right) = 1 - \frac{-\exp\left(-\frac{4\alpha}{3}(mx_{1}+n)G^{3}t^{3}\right) + \exp\left(-\frac{4\alpha}{3}nG^{3}t^{3}\right)}{\exp\left(-\frac{4\alpha}{3}mG^{3}t^{3}\right)}$$
(11)

$$\mathbf{S}_{\mathbf{V}}(t,[0,1]^{3}) = \frac{3\left(-1 - \exp\left(-\frac{4\alpha}{3}(m+n)G^{3}t^{3}\right) + \left(1 + \frac{4\alpha}{3}nG^{3}t^{3}\right)\right)}{\exp\left(\frac{4\alpha}{3}mG^{3}t^{3}\right)}$$

$$\frac{\exp\left(-\frac{4\alpha}{3}(m+n)G^{3}t^{3}\right)}{\frac{4\alpha}{3}mG^{3}t^{3}}$$
(12)

Moreover, the growth velocity can be obtained from:

$$G = \frac{1}{S_E} \frac{dV_E}{dt} = \frac{1}{S_V} \frac{dV_V}{dt} = \frac{1}{\mathbf{S_V}} \frac{d\mathbf{V_V}}{dt}$$
(13)

The relationship between mean and extended mean area density, $S_{F}(t,x)$ is:

$$S_V(t,x) = (1 - V_V(t,x))S_E(t,x)$$
(14)

but

(8)

$$\mathbf{S}_{\mathbf{V}}(t) \neq \left(1 - \mathbf{V}_{\mathbf{V}}(t)\right) \mathbf{S}_{\mathbf{E}}(t) \tag{15}$$

It is worthy pointing out again that for all the equations in this subsection $\alpha = \pi$.

4.2. Analytical expressions for cellular automata

Two important assumptions were made by Rios and Villa¹⁴ in the derivation of the above relationships:

- a) The growing regions are spherical
- b) λ (x) is an harmonic function. In fact, these assumptions may be relaxed¹⁴ but in this case $V_v(t, x)$ would not have the simple form shown in Equations 8-10.

Unfortunately, CA grains are *not* spherical. The shape of CA is compared with a sphere in Figure 2. The analytical expression equivalent to Equation 3 for CA was derived by Rios et al.²².

$$V_V = 1 - \exp(-\frac{4}{3}N_V G^3 t^3)$$
(16)

Comparison between Equations 3 and 16 shows that a factor of π is "missing" from Equation 16. It is suggested here that, by analogy, one may obtain simple approximate analytical expressions to CA from Equations 7-9 and Equations 11-14, replacing $\alpha = \pi$ by $\alpha = 1$. May be, in fact, this is it is not a matter of shape factor α itself, but of the volume of the growing grains. It is as if the theorem for harmonic functions (which holds for spheres) used by Rios and Villa¹⁴ in the derivation of Equation 7 holds also for growing grains such as the one depicted in Figure 2a. Strict mathematical proof of this cannot be given at this time.

Thus, Equations 7-9 and 11-14 with $\alpha = 1$ are compared with the CA simulation in the next section.

5. Results and Discussion

In this section, the expressions derived in the previous section for single grain evolution are compared with CA simulation.

First, it is interesting to show the microstructure resulting from the simulation. Figure 3 depicts a 3-d view of the simulated microstructures for a) $\mathbf{V}_{\mathbf{V}} = 0$; b) $\mathbf{V}_{\mathbf{V}} = 0.1$; c) $\mathbf{V}_{\mathbf{V}} = 0.5$ and d) $\mathbf{V}_{\mathbf{V}} = 1$. The grains are slightly elongated towards the lower plane, $x_1 = 0$. This is more apparent for values of $x_1 < 0.5$. Figure 4 complements Figure 3. It displays the microstructure of the fully transformed specimens of cross-sections taken at planes $x_1 = 0.1$, $x_1 = 0.5$ and $x_1 = 0.9$. It is clear that, as expected, the grain size decreases as the nuclei intensity increases from $x_1 = 0.1$ to $x_1 = 0.9$.

Figure 5 exhibits the volume fraction, the mean volume density, $V_v(t, x)$ as a function of time at planes $x_1 = 0.3$, 0.5, and 0.9. It is worth mentioning that for nuclei too close to $x_1 = 0$ or $x_1 = 1$ one has to use expressions that take into account the influence of the specimen surface given by Rios and Villa¹⁴. Figure 6 shows $\mathbf{V}_v(t)$ as a function of time. Good agreement is obvious in both cases. The microstructural path, \mathbf{S}_v vs. \mathbf{V}_v plot, is depicted in Figure 7. Again,

Rios et al.



Figure 2. Comparison between the shape of a growing region in cellular automata, (a), and spherical growth, (b).



Figure 3. 3-d view of the simulated microstructures a) $\mathbf{V}_{\mathbf{v}} = 0$; b) $\mathbf{V}_{\mathbf{v}} = 0.1$; c) $\mathbf{V}_{\mathbf{v}} = 0.5$; and d) $\mathbf{V}_{\mathbf{v}} = 1$

good agreement is obtained but a small deviation can be observed for V_v close to 1.

A significant change occurred in the way phase transformations have been studied. Formerly, one had to rely on experiments dealing almost exclusively with measurements on 2-d single planar sections. Those experiments were then modeled by theories that were developed bearing in mind the availability of 2-d quantities such as the volume fraction and interfacial area per unit of volume. Moreover, using 2-d sections it is very difficult to detect departures of nucleation from the usual uniform randomly assumption. This is so because the number of nuclei per unit of volume and the non-uniform randomly located nuclei are inherently 3-d properties. Considerable change took place in recent years with the advent of 3-d computer simulation, such as that carried out here, and 3-d experimental techniques like serial sectioning coupled with computer reconstruction. With such techniques, it is now possible to measure and simulate *3-d microstructures*. Now, intrinsic 3-d quantities like number of nuclei per unit of volume or even the position of nuclei in space may be measured/simulated. Nonetheless, this advance has not yet been matched by the development of 3-d analytical treatments such as that put forward by Rios and Villa¹⁴. All three approaches, i. e. experimental, simulation and analytical, must converge if one is to have full 3-d understanding of



Figure 4. Fully transformed microstructure on cross-sections taken at planes a) $x_1 = 0.1$; b) $x_1 = 0.5$; and c) $x_1 = 0.9$.



Figure 5. Mean volume density, $V_V(t, x)$ as a function of time at planes $x_1 = 0.03, 0.5, \text{ and } 0.9$.



Figure 6. The volume fraction, V_v , against time: CA simulation compared with analytical expression.



Figure 7. Microstructural path: area of the interface between transformed and untransformed regions per unit of volume, S_v , against volume fraction, V_v . There is good agreement between theory and simulation.

the problems. Therefore, the current result is of particular importance as it shows that 3-d analytical and 3-d simulation may be combined to analyze an eventual 3-d experimental dataset of nucleation and growth transformations. Noteworthy, 3-d experimental techniques are very demanding in terms of equipment and labor and therefore it is of considerable interest to gather all possible knowledge of the problem by analytical or computer simulation prior to the experiments itself. Thus, one should not see the theoretical approach just as an useful "after the fact" calculation. On the contrary, these techniques may be pro-actively used from the onset to plan experiments and perhaps locate the experimental range within which it might be more profitable to invest time and effort.

6. Summary and Conclusions

CA simulation showed very good agreement with exact mathematical expressions proposed by Rios and Villa even though the shape of CA growing grains is not spherical. However, this good agreement can only be obtained if the appropriate shape factor is used, namely, $\alpha = 1$ (CA shape) instead of $\alpha = \pi$ (spherical shape) in Equations 7-12. It is worth mentioning that a full derivation and discussion of the shape factor, $\alpha = 1$, was carried out in a previous work²¹.

Acknowledgements

This work was supported by Conselho Nacional de Desenvolvimento Científico e Tecnológico, CNPq, Coordenação de Aperfeiçoamento de Pessoal de Nível Superior, CAPES, and Fundação de Amparo à Pesquisa do Estado do Rio de Janeiro, FAPERJ.

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