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Anisotropic solid–liquid interface kinetics in silicon: an atomistically informed phase-field model

S Bergmann¹, K Albe², E Flegel², D A Barragan-Yani² and B Wagner¹

¹Weierstrass Institute for Applied Analysis and Stochastics, Mohrenstr. 39, D-10117 Berlin, Germany

²Technische Universität Darmstadt, FB 11, Institut für Materialwissenschaft, Jovanka-Bontschits-Str. 2, D-64287 Darmstadt, Germany

E-mail: sibylle.bergmann@wias-berlin.de

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Abstract

We present an atomistically informed parametrization of a phase-field model for describing the anisotropic mobility of liquid–solid interfaces in silicon. The model is derived from a consistent set of atomistic data and thus allows to directly link molecular dynamics and phase field simulations. Expressions for the free energy density, the interfacial energy and the temperature and orientation dependent interface mobility are systematically fitted to data from molecular dynamics simulations based on the Stillinger–Weber interatomic potential. The temperature-dependent interface velocity follows a Vogel–Fulcher type behavior and allows to properly account for the dynamics in the undercooled melt.

Keywords: phase-field model, molecular dynamics simulation, interface kinetics, silicon recrystallization

(Some figures may appear in colour only in the online journal)

1. Introduction

The growth of silicon is relevant for a wide range of technological processes in semiconductor industry, including the production of polycrystalline silicon for photovoltaics by



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electromagnetic casting, edge-defined film feed methods, ingot directional solidification techniques, and also liquid phase crystallization. Currently, over 90% of the commercial solar cells are made from single- or multi-crystalline silicon. The production volume of solar cells using the multi-crystalline silicon is higher than that of single-crystalline silicon. In order to obtain a detailed understanding of the interplay of process parameters and the resulting microstructure, computer simulations have become an increasingly important tool. However, modeling of nucleation processes and growth morphologies requires a quantitatively correct description of anisotropic interface energies and mobilities of the crystal-melt interface.

Simulations of the solidification of multi-crystalline Si including the evolution of grains can be divided in macroscopic, microscopic and atomistic methods. On the macroscopic scale, cellular automata and geometric models were proposed, which are most efficient, but lack some physical details. Atomistic molecular dynamics simulations have been successfully applied to simulate solidification of silicon and thus offer a route for revealing details of the growth kinetics [8, 9, 36]. Because of the enormous computational effort, however, these models are restricted to relatively small system sizes of typically not more than a couple of million atoms. This is why for modeling phenomena on the microscopic scale, phase field models (PFM) have emerged as a promising and powerful tool for simulating free boundary problems with complex morphological evolution. Since the transport equations for heat and mass and the phase field are solved simultaneously, the effects of surface tension, nonequilibrium, and anisotropy can be directly included. PF models are based on physical parameters and can take into account anisotropies of interface energies and mobilities.

In the context of silicon grain growth PFM face, however, several challenges. The large anisotropy of interface energies and directional dependent mobilities determine in a delicate way the combination of occurring facets. Moreover, the solidification process is in general much slower than for metallic systems and thus there is smaller thermal gradients. A technical drawback of PFM lies in the fact that the minimum mesh size has to be smaller than the interface thickness, while a realistic interface thickness is only on the order of the capillary length approximately several Ångström. The large body of literature on phase-field models for transitions between liquid and solid phases has been reviewed, for example in Boettinger *et al* [5], Wheeler *et al* [46] and more recently in Moelans *et al* [32] and in the context of solidification and dendritic growth by Steinbach [40]. Especially, for the problem of excimer laser annealing of a Si layer on an amorphous substrate Magna *et al* [30] and Shih *et al* [39] developed specific phase-field models based on coupled equations describing the thermal, phase and impurity redistribution during the annealing process. A recent review on liquid thermal annealing was published by Fiscaro *et al* [15]. However, in the PFM existing studies dealing with silicon mostly qualitative assumptions on free energy densities, anisotropic interface energies and mobilities were used. On the other hand, detailed information on melting points, interface velocity and formation of defects during crystal growth are in principle available from molecular dynamics simulations and can directly be used. Therefore, it seems natural to ask if both modeling approaches can be combined to yield quantitative accurate models, that are amenable to large scale simulations. This has been the concern of a number of studies in recent years, where it has been shown how atomistic molecular dynamics computations can be used to obtain quantitative information for kinetic and thermodynamic properties to correctly predict the dynamics of the corresponding multi-phase systems using phase-field models. In the context of dendritic solidification, for example, Hoyt *et al* [21] developed a method for extracting anisotropic interface energies from atomistic molecular dynamics simulations and used them in a phase-field description with weak anisotropy of the solid-liquid interface. Similarly, Bragard *et al* [6], derived PF-parameters for predicting the dendrite growth velocity as a function of undercooling in pure Ni. A more

detailed overview on these problems can be found in Hoyt *et al* [22]. For the solidification of the alloy systems of NiZr, Danilov *et al* [10] and Guerdane *et al* [20] addressed the more fundamental question if molecular dynamics simulations and the phase-field approach can give quantitative equivalent results. At least for these specific alloy systems they found good agreement in quantities such as the melting rates by comparing their numerical results.

Interestingly, there is no published study in which the thermodynamic parameters of a phase-field model for solidification of silicon are extracted from atomistic simulations, although some relevant data are available [4, 7, 13, 19]. Thus, the focus of this study is to establish a phase-field model, where the complete set of necessary parameters is derived from molecular dynamics simulations based on the Stillinger–Weber (SW) interatomic potential for Si [42]. In particular, we incorporate a consistent description of the Vogel–Fulcher-type temperature dependence of the interface velocity of Si [18, 41, 44]. In order to establish the necessary phase-field parameters we investigate three distinct planar interface orientations.

2. From an atomistic to a phase-field description

In the diffuse interface description the transition between a liquid and crystalline phase is introduced by a phase-field variable $p(\mathbf{x}, t)$. It is a function in space and time that varies from $p(\mathbf{x}, t) = 0$ in the liquid state to $p(\mathbf{x}, t) = 1$ in the crystalline state. In the simplest setting for a pure melt in the isothermal case a free energy functional

$$\Phi(p, T) = \int_V F(p, T) + \frac{\sigma^2}{2} |\nabla p|^2 dV, \quad (1)$$

can be derived from thermodynamic considerations [5]. The bulk free energy density $F(p, T)$ is a function of the phase-field variable and temperature. The gradient energy coefficient σ is related to the steepness or width of the transition from the liquid to the solid phase. For anisotropic interfacial energies, this also depends on the orientation angle normal to the contours of constant p as shown in [26, 31]. Since we consider a one-dimensional model, different σ parameters are chosen depending on the given interface orientation $\{hkl\}$, denoted by $\sigma_{\{hkl\}}$. Upon minimization of the decreasing free energy functional $\Phi(p, T)$ the evolution equation for $p(\mathbf{x}, t)$ is obtained as

$$\frac{\partial p}{\partial t} = -M_{\{hkl\}}^{\text{PF}}(T) \left(\frac{\partial F}{\partial p}(p, T) - \sigma_{\{hkl\}}^2 \Delta p \right), \quad (2)$$

where $M_{\{hkl\}}^{\text{PF}}$ denotes the interfacial mobility parameter of the phase field describing the relaxation dynamics of the interface. The mobility parameter $M_{\{hkl\}}^{\text{PF}}$ depends on temperature and also on interface orientation.

As schematically sketched in figure 1, the free energy density F is conveniently chosen as double-well potential with minima at $p = 0$ for the liquid and $p = 1$ for the crystalline phase and with a maximum in between. We choose a fourth order polynomial in p with temperature dependent coefficients. The temperature dependent minima correspond to the free energies of the liquid and solid phase and can be calculated from atomistic simulations by thermodynamic integration (see section 3.2).

By measuring the normal velocity

$$v = MF \quad (3)$$

of a moving flat interface in a molecular dynamics simulation (see section 3.4) one can directly determine the orientation dependent mobility M of a solid–liquid interface. The

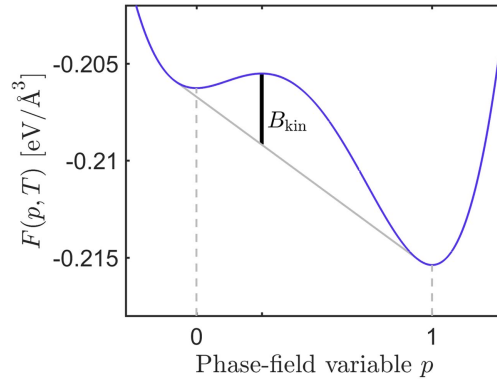


Figure 1. Example of the bulk free energy density F and its kinetic barrier B_{kin} at temperature 400 K (less than T_m^V). B_{kin} is the kinetic barrier, that has to be overcome to pass from one phase into the other.

driving force F is determined by the free energy difference of the liquid and solid phase at a given temperature and can be independently calculated. However, the thermodynamic mobility M obtained in such a manner, cannot be used as parameter in a phase-field model. The reason is that in a phase-field simulation the system has to overcome the barrier B_{kin} dividing the two potential wells of the free energy density F as sketched in figure 1. As we show in section 4.1 in equations (11) and (14), the kinetic barrier B_{kin} has a direct effect on the free energy density. This contributes to the mobility of the phase-field and thus M^{PF} cannot be identical to M . This is often disregarded in parametrizations for recrystallization phenomena as e.g. in [15, 39, 46].

The second term of the integrand in equation (1) describes the crystal-melt interfacial energy, which scales with the coefficient $\sigma_{\{hkl\}}$. For a given interfacial energy $\gamma_{\{hkl\}}$, which can again be obtained from molecular dynamics simulations, and a fixed interface width $\varepsilon_{\{hkl\}}$, one can adjust B_{kin} and $\sigma_{\{hkl\}}$. The relations between B_{kin} , $\sigma_{\{hkl\}}$, $\gamma_{\{hkl\}}$ and $\varepsilon_{\{hkl\}}$ are

$$B_{\text{kin}} T_m^V = \frac{\gamma_{\{hkl\}}}{\varepsilon_{\{hkl\}}}, \quad \sigma_{\{hkl\}} = \sqrt{2\gamma_{\{hkl\}} \varepsilon_{\{hkl\}}},$$

as given in [1].

Again, it should be noted that the parameter σ is often considered as freely adjustable. Thus, for a given interface energy, rescaling the σ parameter for numerical reasons is equivalent to rescaling of the interface width, which in turn means that the parameter B_{kin} needs to be readjusted, if the interface energy shall remain unaffected.

In this paper, we choose $\varepsilon_{\{111\}} = 1$ nm as a physical parameter describing the width of the interface for the $\{111\}$ growth plane and use interface velocities, interfacial energies and free energy densities obtained from molecular dynamics simulations.

The velocity versus temperature relationship is fitted to an equation describing the competition between kinetics and thermodynamics of the crystallization process. In section 4.3, the remaining parameter of the phase-field model, the mobility M^{PF} , is then obtained by a shooting method applied to (2) in one dimension, such that the crystallization velocity of the phase-field simulation agrees with the growth velocity calculated by means of molecular dynamics, which we prove numerically in section 4.4.

Table 1. Heat capacity, latent heat and melting point of silicon. T_m^{MD} is the melting point obtained from moving interface simulations, T_m^{V} is the melting point obtained at constant volume from the adiabatic switching method (see section 3.2).

	$c_{p,\text{solid}}$ (J mol ⁻¹ K ⁻¹)	$c_{p,\text{liquid}}$ (J mol ⁻¹ K ⁻¹)	ΔH_m (kJ mol ⁻¹)	T_m^{MD} (K)	T_m^{V} (K)
SW-pot.	26.9	28.7	31.7	1683	1697.12
other sim.			30.9 [7]	1682 [17]	
Exp.	26–29 [12]	27.2 ± 1.5 [12]	50.25 ± 0.6 [12]	1683 [42]	
Data	(900–1687 K)				

3. Atomistic model and parameter calculation

3.1. Method

Molecular dynamic calculations are performed using the SW interatomic potential for silicon [42], which describes the structure of the molten phase realistically, and reproduces the experimental melting point [25, 28, 29, 38]. We calculate thermal properties and interfacial velocities with the widely used LAMMPS code [35] and obtain free energy densities via using the MD++ code [37]. In table 1 we present a summary of the thermal properties obtained using the SW potential along with results from previous simulations and experiments.

We initialize a simulation box containing 4096 atoms in the diamond structure and heat it up. For doing so, we apply a Nosé–Hoover thermostat with a rate of 10^{13} K s⁻¹. After melting occurs, we cool it down with the same rate and calculate the specific heat and latent heat from the average total energy. Clearly, the specific heat and melting point are in good agreement with the experimental finding. The latent heat, in contrast, is low compared to experimental measurements. The reason is that we describe the solid phase and the liquid phase by a single empirical model despite their different bonding mechanisms [37]. The melting point T_m^{MD} calculated from a simulation cell with solid–liquid phases co-existence is almost exact compared to the experimental value of 1683 K and in good agreement with earlier simulations [14, 17]. If the intersect of free energies calculated by the adiabatic switching method at constant volume is used (see section 3.2) the calculated melting point $T_m^{\text{V}} = 1697.12$ K is slightly higher. In order to be consistent with the free energy data, we use T_m^{V} as melting temperature for the phase-field model.

3.2. Free energy densities

For calculating the free energy density, we use a supercell with 512 atoms and periodic boundary conditions in all three directions. The actual calculation of the free energy is performed using both adiabatic switching and reverse scaling [11, 45] as implemented in the MD++ software package by Ryu and Cai [37]. In order to be consistent with our phase-field model, we modify the code such that an NVT ensemble is used, the initial volume for both phases—solid and liquid—is equal to the equilibrium volume of the crystalline silicon supercell at 0 K. The free energy density is calculated then by dividing the obtained free energies by that same volume. Furthermore, in our approach we include the free energies of the amorphous state calculated by Broughton and Li [7]. The results are presented in figure 2. The melting point used in the phase-field model is the one obtained from this free energy calculation, $T_m^{\text{V}} = 1697.12$ K, which is the temperature at which the amorphous/liquid and

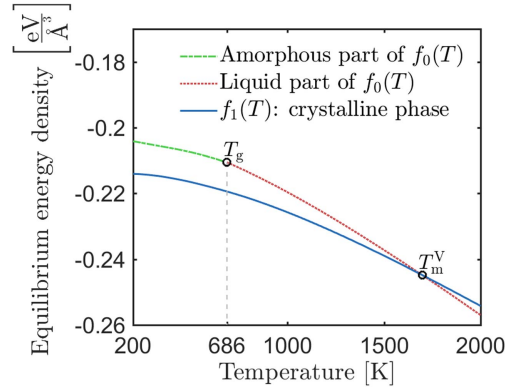


Figure 2. Minimum values for the free energy density for liquid/amorphous ($f_0(T)$) and crystalline ($f_1(T)$) equilibrium to the SW potential for a temperature range of 200–2000 K. The free energy equilibrium values are calculated with the codes provided by Ruy and Cai [37]. For the liquid branch, T_g points out the glass transition. The free energies for the amorphous phase were calculated by Broughton and Li [7].

Table 2. The crystal-melt interface free energies γ at the melting point T_m^V for the orientations {100}, {110} and {111} calculated by Apte and Zeng in [2].

Orientation	{100}	{110}	{111}
$\gamma(T_m^V)$ in $\text{eV } \text{\AA}^{-2}$	0.026	0.0218	0.0212

solid free energies intersect. We point out that this value does not correspond to the experimental value or to the values obtained from direct interface calculations (see table 1). The reasons for this discrepancy are the NVT ensemble we used for the free energy calculation and a numerical error in the adiabatic switching and reverse scaling methods, used for a large temperature intervals like the one we analyze.

3.3. Interfacial free energies

Interfacial free energies calculated with the SW potential are available in literature from Apte and Zeng [2], who used molecular dynamics to determine $\gamma_{\{100\}}$, $\gamma_{\{110\}}$ and $\gamma_{\{111\}}$ at the melting point. Their mean values are given in table 2. The most densely packed (111) orientation has the lowest interface energy, while the (110) direction exhibits nearly the same excess energy. This is obviously at odds with the equilibrium shape of Si grains embedded in a melt, which show a Wulff shape with (111) and (100) facets only. Thus, $\gamma_{\{110\}}$ is obviously underestimated by the SW-potential. Since the purpose of this study is to devise model parameter, which allows us to directly combine phase-field and molecular dynamics simulations with consistent model parameters corresponding to the SW potential, we adopt these values for our parametrization.

3.4. Interface velocities

We calculate interface velocities from molecular dynamics simulations of moving planar liquid–solid interfaces with different crystallographic orientations at constant temperature

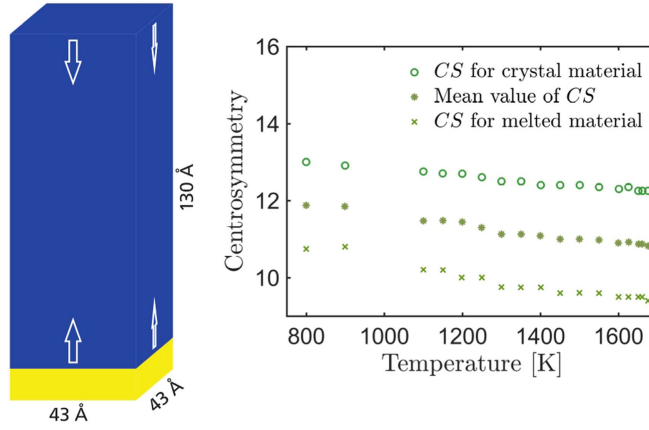


Figure 3. The schematic of the simulation box used to calculate the interface velocities is presented in the left figure. For temperatures below the melting point the yellow region corresponds to the crystalline silicon and the blue region to the melt. In the case of temperatures above the melting point, the color code is reversed. In the right figure we show the dependence of the average centrosymmetry parameter of melt and solid on temperature. The mean value $CS_{crit}(T)$ is used to determine the position of the crystallization front.

$T \in [800, 2000 \text{ K}]$. For this, we initialize a simulation box of about $43 \times 43 \times 130 \text{ Å}$, like the one shown in the left-hand side of figure 3. Such box contains 12 000–13 000 atoms, depending on the crystal orientation and has periodic boundary conditions in all three directions. The box sizes in x - and y -direction are adjusted in order to obtain a single crystal without lattice defects near the boundaries of the box. We start our simulation with an equilibration phase using a Nosé–Hoover thermo- and barostat for 10 ps at the desired temperature in order to consider thermal expansion of the box. One timestep corresponds to 1 fs. The box dimensions are left free to vary independently of each other. For the case of temperatures below the melting point, as shown in the left-hand side of figure 3, the crystalline part of the box (about 1/12) is equilibrated at the desired temperature. The remaining atoms are melted at 1000 K above the melting point and then cooled to a temperature near T_m^{MD} . While melting, the box dimensions in x - and y -direction are fixed, but in z -direction the box is allowed to shrink or expand. Finally, we run the crystallization for some nanoseconds with a global thermostat at the desired temperature. The x - and y -dimensions are fixed again, but not in the growth direction. For the case of temperatures above the melting point the procedure is analogous. In this case, the lower part of the box (about 1/12) is heated up to the desired temperature and, therefore, is melted, while the upper part of the box is kept crystalline with a temperature near T_m^{MD} . Then, for some nanoseconds, the complete simulation box is connected to global thermostat at the desired temperature above T_m^{MD} .

To extract the velocity of the interface, one first has to determine its position at each timestep. There are numerous ways to do so (see [28]), for instance, by monitoring the particle density or the atomic potential energy. The observed parameter only has to fulfill the condition that it is sufficiently different in the solid and liquid phase. In this study, we choose the centrosymmetry parameter [24], which can be calculated for each atom within LAMMPS. It is zero for an atom in a perfect lattice, and gives a positive value for disturbed atomic environments. The average of the centrosymmetry parameter over one atomic layer perpendicular to the growth direction (corresponding to some hundreds atoms) results in a

centrosymmetry of 8.5–11 for a liquid layer and 12–13.5 for a crystalline layer. We find that the centrosymmetry of melt and crystal is dependent on temperature, as shown in the right-hand of figure 3.

Therefore, we take the mean value of the centrosymmetry of crystal and liquid as the critical value CS_{crit} in order to distinguish liquid and solid atoms. Using the centrosymmetry method, the isothermal interface velocities are finally determined for certain temperatures in the range 800–2000 K for the SW potential.

The latent heat and the heat capacity determine (together with heat conductivity) how much heat is generated in a crystallizing sample at the moving interface and how fast it is conducted away. It was shown by Monk *et al* [33] that due to the release of latent heat, the actual interface temperature can differ from the one which is set by the thermostat. Thus, Monk *et al* proposed to use multiple thermostats, from which each one only sets the temperature for a volume element smaller than 20 Å in thickness. They simulated the scenario for pure Ni. Our temperature calculations during crystallization show a flat temperature profile over the whole simulation box. This indicates that heat is taken away fast enough by one global thermostat and did not influence the crystallization velocity.

Another important feature of silicon is the presence of an amorphous phase, if there is a significant undercooling. This is captured by our isothermal conditions for the moving interface and results in a Vogel–Fulcher type dependence of the interface velocity on temperature [18, 41, 44].

However, in order to feed the PFM with these information, we need an analytical expression for the growth velocity. The growth velocity is described by the product of driving force P and mobility M , which is formulated for an atomically flat solid–liquid interface by Jackson [23] as

$$v = M \cdot P = f \cdot A \cdot \nu \cdot P, \quad (4)$$

where f represents the percentage of favorable growth sites (i. e. steps) on the crystal surface, $A = \sqrt[3]{\Omega}$ the cube root of the atomic volume, ν the attempt frequency of atom jumps over the interface and P the driving force for crystallization. Within transition state theory [23] the interface velocity is given by the difference between the velocities of crystallization and melting:

$$v = v_0 \cdot \exp\left(-\frac{Q}{kT}\right) \cdot \left[1 - \exp\left(-\frac{\Delta G}{kT}\right)\right], \quad (5)$$

where Q is activation energy associated with atomic mobility and ΔG is the Gibbs free energy difference between the two phases. The last term in brackets is the thermodynamic driving force for crystallization F , and can be approximated by a series expansion, which we develop upto second order to obtain

$$F = \left(1 - \exp\left(-\frac{\Delta G}{kT}\right)\right) \approx \frac{\Delta G}{kT}. \quad (6)$$

In [47], Wilson derives the term $d \cdot \nu$ in equation (4) as $6D/d$ with the diffusion coefficient D . Frenkel [16] refines this expression further by replacing the diffusion coefficient with the Stokes–Einstein relation

$$D = \frac{kT}{3\pi\eta r}, \quad (7)$$

which describes the diffusion of a spherical particle with radius r in a liquid with viscosity η . As a first approximation, the mobility of a particle in the liquid follows an Arrhenius function.

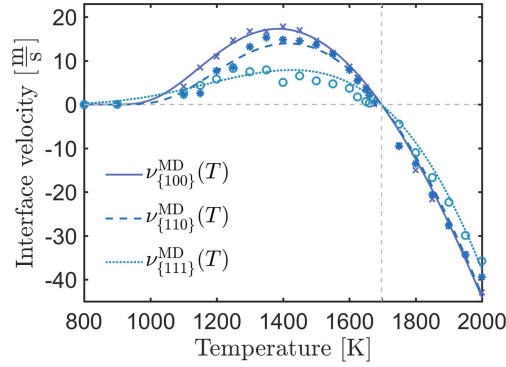


Figure 4. Crystallization velocity over temperature for silicon with the Stillinger–Weber potential, fitted by a Vogel–Fulcher equation (9).

However, from the enthalpy as a function of temperature from the section 3.1, we noted the occurrence of a glass transition. Specifically, when approaching and crossing T_g the mobility of the atoms in the melt is reduced and diffusion is slowed down drastically. The Arrhenius description does not describe this behavior of glass-forming melts. To overcome this problem, Vogel [44] and Fulcher [18] introduced an empirical relation allowing the increase in viscosity when approaching the glass transition

$$\eta = \eta_0 \cdot \exp\left(\frac{A}{T - T_{VF}}\right), \quad (8)$$

where η_0 and A are constants and the Vogel temperature, T_{VF} , lies about 50 K below the glass transition temperature. By replacing the Arrhenius- with the Vogel–Fulcher-expression, we finally obtain

$$v = f \cdot \frac{\Delta G}{kT} \cdot \frac{6D}{d} = \frac{2f \cdot \Delta H_m}{3\pi\eta_0 r d \cdot T_m^V} \cdot (T_m^V - T) \cdot \exp\left(-\frac{A}{T - T_{VF}}\right). \quad (9)$$

The resulting velocity–temperature relationships are depicted in figure 4, where we fit our measurements with the Vogel–Fulcher expression (9).

The small velocity of the $\{111\}$ interface is related to its dense packed structure and low energy, which does not provide favorable sites for the attachment of atoms. Therefore, a nucleation step has to take place before a new $\{111\}$ layer can grow.

On the contrary, the $\{110\}$ and $\{100\}$ interfaces are rough because of the formation of $\{111\}$ facets. Since the growth in these directions is not nucleation limited, it is faster, which is in agreement with results reported in previous studies [3, 43]. In figure 5 the atomic configuration of an $\{110\}$ interface under growth conditions is shown. The formation of crystalline $\{111\}$ can be clearly seen as well as the formation of faulted planes on these facets. As a rule of thumb, the relation between the maximum velocities $v_{\{100\}}$; $v_{\{110\}}$; $v_{\{111\}}$ of the fitted curves is about 1:0.8:0.4 for this potential. The difference in growth velocity of $\{100\}$ and $\{110\}$ is related to the factor f in equation (4) since all other terms are bulk properties and not orientation dependent. To approach the factor f theoretically, we calculated the density of favorable sites by the density of $\{111\}$ planes ending at a $\{100\}$ surface. This is given by

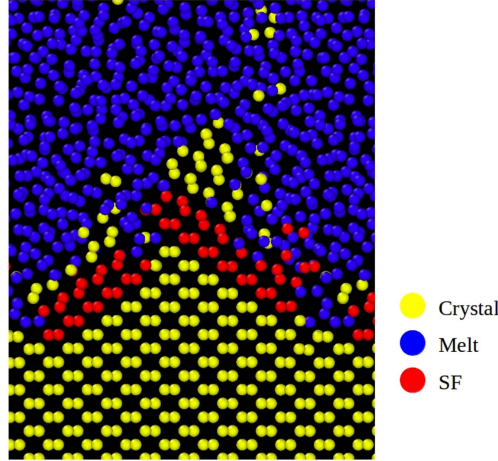


Figure 5. The atomic configuration of the $\{110\}$ interface while crystal growth occurs. It is possible to see the rough character of the interface evidenced by the presence of $\{111\}$ facets. Furthermore, stacking faults (SF) occur on these facets (red atoms). A similar behavior is observed for the $\{100\}$ interface.

$$\rho_{\{100\}} = \frac{\sin \alpha_{\{100\}}}{d},$$

where d is the distance of $\{111\}$ planes and $\alpha_{\{100\}}$ the angle between the $\{111\}$ and $\{100\}$ plane. If one compares $\rho_{\{100\}}$ and $\rho_{\{110\}}$ a relation of 1: $1/\sqrt{2}$ or 1:0.7 is found, which is in rough agreement with the relation derived from the simulation. Alternatively one can count the number of broken bonds per area at the surface, which gives an identical result.

Compared to literature values for MD simulations, we find values of 18–20 m s^{-1} for Stillinger $\{100\}$ and 9–14 m s^{-1} for $\{111\}$ [25, 28, 29], which is in good agreement with the above measured values. From experiments, velocities of 1.6 m s^{-1} are reported by Kuo [27] and 14 m s^{-1} by Ohdaira [34], so that we conclude that the results for the SW potential are a good representation of the anisotropic growth velocity of silicon crystals.

4. Atomistically informed phase field parameters

In this section, we derive the parameters for the phase-field model

$$\frac{\partial p}{\partial t} = M_{\{hkl\}}^{\text{PF}}(T) \left(\sigma_{\{hkl\}} \frac{\partial^2 p}{\partial x^2} - \frac{\partial F}{\partial p}(p, T) \right), \quad (10)$$

where the one dimensional phase-field variable $p : \mathbb{R} \times (0, \tau) \rightarrow [0, 1]$ varies between 0 and 1 to describe the two bulk states: liquid ($p = 0$) and solid ($p = 1$) and the interface region between the bulk states ($0 < p < 1$), as already mentioned in section 2.

Our main focus is that (10) reproduces the interface velocities calculated in section 3.4, while all parameters are carefully chosen, such that they are consistent with molecular dynamics with the SW potential. Since we have from molecular dynamics information about the three crystallographic orientations $\{100\}$, $\{110\}$ and $\{111\}$, we also derive the model parameters for this orientations, which are indicated with the indices hkl in (10).

At the first step we derive the bulk free energy density F with the help of our molecular dynamical results described in section 3.2. In section 4.2, we incorporate interface energies from literature, which are also calculated via molecular dynamics with the SW potential. Finally, in section 4.3, we adapt the mobility parameter M^{PF} , such that the model reproduces interface velocities from section 3.4, which we prove numerically in section 4.4.

4.1. Polynomial describing the bulk free energy density

A free energy density $F(p, T)$, which has the form of a double-well potential in p , can be established as a polynomial of fourth degree, which is one of the common forms for F . Here, the coefficients may depend on the temperature and we assume the expression

$$F(p, T) = a_0(T) + a_1(T)p + a_2(T)p^2 + a_3(T)p^3 + a_4(T)p^4.$$

Since the equilibrium states for the bulk free energy density are represented by the two minima of the double-well polynomial, we choose the coefficients a_0, \dots, a_4 , such that $(0, F(0, T))$ and $(1, F(1, T))$ are the minima points of F . Then, we equip the free energy density F of the phase-field model with the equilibrium values of the atomistic free energy and via polynomial interpolation, such that we obtain two polynomials, $f_1(T)$ for the crystalline values in figure 2 at $p = 1$ and $f_0(T)$ for the amorphous/liquid values at $p = 0$. With respect to the condition, that f_1 and f_0 represent the minima of F , the free energy density has the form

$$F(p, T) = f_0(T) + a(T)p^2 - 2(a(T) + 2H(T))p^3 + (a(T) + 3H(T))p^4, \quad (11)$$

where $H(T) = f_0(T) - f_1(T)$. Note here, that $H(T < T_m^V) > 0$. For the remaining degree of freedom $a(T)$ in (11) yields:

$$a(T) > \begin{cases} 0 & T \in [0, T_m^V] \\ -6H(T) & T > T_m^V. \end{cases} \quad (12)$$

This expression for F fulfills $F(0, T) = f_0(T)$ and $F(1, T) = f_1(T)$ for the minima. Details of the derivation of (11) and (12) are given in the appendix A.1.

Furthermore, the maximum point of F is $(\mu(T), F(\mu(T), T))$ with

$$\mu(T) = \frac{a(T)}{2a(T) + 6H(T)} \in (0, 1). \quad (13)$$

The calculation of (13) is described in appendix A.2. Hence, as we indicated in section 2, for the line $g(p)$ which is tangent to both of the minima of F , the energy barrier B_{kin} , that has to be exceeded to get from one equilibrium phase to the other, is the difference of the function values of the maximum $F(\mu)$ and $g(\mu)$. At the melting point T_m^V , the two minima of F have the same function value and thus $B_{\text{kin}}(T_m^V)$ is the difference of the function value of the maximum of F and the function value of the minima. Since the function values of the minima are then $f_0(T_m^V) = f_1(T_m^V)$, with equation (13) the maximum point at T_m^V has the simple expression

$$(\mu(T_m^V), F(\mu(T_m^V))) = (1/2, f_0(T_m^V) + a(T_m^V)/16).$$

Hence, the kinetic barrier at the melting point has the form

$$B_{\text{kin}}(T_m^V) = \frac{a(T_m^V)}{16}. \quad (14)$$

Table 3. The crystal-melt interface free energies γ at the melting point T_m^V for the orientations $\{100\}$, $\{110\}$ and $\{111\}$ calculated by Apte and Zeng in [2]. With the help of γ and (15), we calculate the model constants a and σ for the three orientations. For that we choose an interface thickness of $\varepsilon = 1$ nm for orientation $\{111\}$.

Orientation	$\{100\}$	$\{110\}$	$\{111\}$
$\gamma_{\{hkl\}}(T_m^V)$ in $\text{eV } \text{\AA}^{-2}$	0.0262	0.0218	0.0212
$\varepsilon_{\{hkl\}}$ in \AA	12.35	10.29	10
$\sigma_{\{hkl\}}^2$ in $\text{eV } \text{\AA}^{-1}$	0.804 ²	0.670 ²	0.651 ²
a in $\text{eV } \text{\AA}^{-3}$	0.0339		

The kinetic barrier closely relates to the interface energy γ , which we discuss in section 4.2, where we also determine B_{kin} and hence with (14) the remaining degree of freedom for the bulk energy.

4.2. Interfacial free energy and width

We derive the gradient energy coefficient σ and the degree of freedom a in the free energy density (11) consistent with the interfacial energies obtained in Apte and Zeng [2], who used molecular dynamic simulations. We note that for the calculation of σ and a , we presently use only the values for the equilibrium state and hence make a independent on the temperature, since there is no literature with values of temperature dependent interface free energy calculated by means of molecular dynamics with SW potential for Si. But after all, the mobility parameter M^{PF} will compensate the missing temperature dependence at this point.

Since Allen and Cahn [1], we know the relations between the interfacial energy coefficient σ in the PFM, the modeled interface thickness ε , the interfacial free energy γ and the modeled kinetic barrier B_{kin} .

At the melting point $T = T_m^V$ the relations are

$$\gamma_{\{hkl\}}(T_m^V) \approx \sigma_{\{hkl\}} \sqrt{\frac{1}{2} B_{\text{kin}}(T_m^V)}, \quad \varepsilon_{\{hkl\}} \approx \frac{\sigma_{\{hkl\}}}{\sqrt{2 B_{\text{kin}}(T_m^V)}}.$$

For convenience we set

$$\frac{a}{16} = B_{\text{kin}}(T_m^V) = \frac{\gamma_{\{111\}}(T_m^V)}{\varepsilon_{\{111\}}}, \quad \sigma_{\{hkl\}} = \sqrt{2 \gamma_{\{hkl\}}(T_m^V) \varepsilon_{\{hkl\}}}. \quad (15)$$

For the crystal orientations $\{100\}$, $\{110\}$ and $\{111\}$, Apte and Zeng [2] obtained the crystal-melt interfacial free energy γ , see table 3. We assume for orientation $\{111\}$ an interface thickness of $\varepsilon_{\{111\}} = 1$ nm and find a kinetic barrier of $B_{\text{kin}} \approx 0.002 \text{ eV } \text{\AA}^{-3}$. Hence, the interface thickness of orientation $\{100\}$ results in $\varepsilon_{\{100\}} \approx 12.35 \text{ \AA}$ and of $\{110\}$ in $\varepsilon_{\{110\}} \approx 10.29 \text{ \AA}$. In table 3, the values γ , ε , and the resulting model parameters σ and a are given for all three considered orientations. Figure 6 shows the resulting double-well potential at different temperatures.

4.3. Realization of interface velocity in the phase-field model via the mobility parameter

In the previous subsections, we derived the bulk energy and the gradient energy of the phase-field model (10) by explicit use of the results from molecular dynamics. In this section, we calculate the mobility parameter $M_{\{hkl\}}^{\text{PF}}(T)$ of the phase-field model (10), such that the model

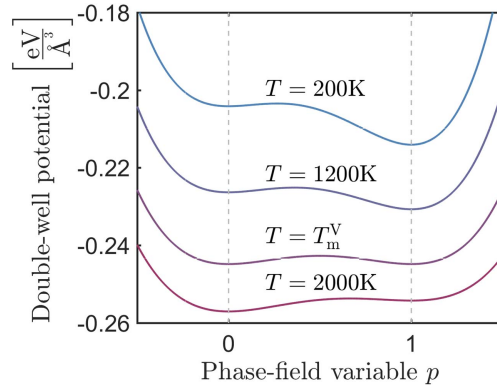


Figure 6. The double-well potential at different temperatures. At the melting point T_m^V , the two minima of the bulk free energy density have the same value.

reproduces the atomistic velocities $v_{\{hkl\}}^{\text{MD}}(T)$ from section 3.4. But in the phase field, the interface velocity is not a parameter which can be directly incorporated as the interface energy $\gamma_{\{hkl\}}$ in $\sigma_{\{hkl\}}$ or the free energy minima in F . Hence, we implement a shooting method, where we vary $M_{\{hkl\}}^{\text{PF}}(T)$ until all required conditions are fulfilled for a fixed temperature and orientation. This procedure is repeated for all three considered directions (100), (110) and (111), where we make measurements for the temperatures $T = 800, 850, 900, \dots, 1950, 2000$ K.

For the shooting method, we first define boundary values for (10), such that we have crystal material at the left boundary, and liquid material at the right:

$$0 = M_{\{hkl\}}^{\text{PF}}(T) \sigma_{\{hkl\}}^2 \frac{\partial^2 p}{\partial x^2} - M_{\{hkl\}}^{\text{PF}}(T) \frac{\partial F}{\partial p} - \frac{\partial p}{\partial t}, p(-\infty, t) = 1, p(+\infty, t) = 0. \quad (16)$$

We now fix the temperature and the orientation arbitrarily. Hence, M^{PF} , σ and also the velocity ν^{MD} are constants. The phase field is a traveling wave solution and can be expressed as

$$p(x, t) = \phi(x - \nu^{\text{MD}}t) =: \phi(\xi).$$

Substitution of ϕ in (16) leads to the boundary value problem:

$$0 = M^{\text{PF}} \sigma^2 \phi'' - M^{\text{PF}} F' + \nu^{\text{MD}} \phi', \phi(-\infty) = 1, \phi(\infty) = 0, \quad (17)$$

where $\phi' = \frac{d\phi}{d\xi}$ and $F' = \frac{\partial F}{\partial \phi}$.

To receive an initial-value problem, we integrate the ordinary differential equation in (17) with the left boundary $\phi(-\infty) = 1$, which holds as our first initial condition, and the right boundary s , which represents the new space-dependence variable for the shooting method. Furthermore, we define $\tilde{F}'(\xi) := F'(\phi(\xi))$, thereby we are able to integrate the derivative of the potential F in ξ . For the calculation of the integral we use the second initial conditions $\phi'(-\infty) = 0$. The both initial conditions guarantee, that we have crystal material at the left boundary. Applying all described conditions, we receive

$$c_1 = M^{\text{PF}} \sigma^2 \int_{-\infty}^s \phi'' d\xi - M^{\text{PF}} \int_{-\infty}^s \tilde{F}'(\xi) d\xi + \nu^{\text{MD}} \int_{-\infty}^s \phi' d\xi, \quad (18)$$

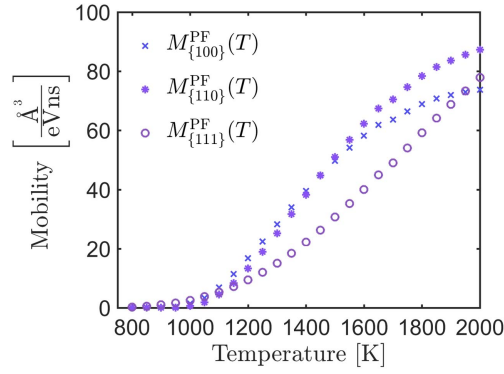


Figure 7. The extracted mobilities of the phase-field model as a function of temperature and orientation. The concrete values are listed in appendix A.3 in table A1.

$$c_2 = M^{\text{PF}} \sigma^2 (\phi'(s) - 0) - M^{\text{PF}} (\tilde{F}(s) - \tilde{F}(-\infty)) + \nu^{\text{MD}} (\phi(s) - 1). \quad (19)$$

By choosing $c_2 = M^{\text{PF}} \tilde{F}(-\infty)$, our initial value problem has the form:

$$\begin{pmatrix} \tilde{F}'(s) \\ \phi'(s) \end{pmatrix} = \begin{pmatrix} F'(\phi(s)) \\ \frac{\nu^{\text{MD}}}{M^{\text{PF}} \sigma^2} (1 - \phi(s)) + \frac{1}{\sigma^2} \tilde{F}(s) \end{pmatrix}, \quad (20)$$

$$\phi(-\infty) = 1, \quad (21)$$

$$\phi'(-\infty) = 0. \quad (22)$$

We calculate the mobility M^{PF} via a bisection method. Therefore we vary M^{PF} until the right boundary condition $\phi(\infty) = 0$ is fulfilled for the numerical solution of (20)–(22). Thereby we solve the initial value problem with Runge-Kutta 4/5. We apply this method for each of the three orientations $\{100\}$, $\{110\}$ and $\{111\}$ and use the respective value of $\sigma_{\{hkl\}}$ listed in table 3. Thereby we calculate ν^{MD} for the respective temperature and orientation with the fits shown in figure 4. Our results are shown in figure 7 and listed in appendix A.3.

4.4. Numerical solution of the phase-field model and velocity reproduction tests

We solve model (10) numerically at fixed temperatures $T = 1050, 1100, \dots, 2000$ K for the three crystallographic orientations. During the simulation, we measure the velocity of the interface region by interpolating the position of $p = 0.5$. In fact, the model reproduces the velocities $v_{\{hkl\}}^{\text{MD}}(T)$ from molecular dynamics, see figure 8.

As for the molecular dynamical simulations, we also have periodic boundary conditions for the phase-field model for $p(x, t)$, which we solve numerically using a Fourier spectral method. Our equidistant grid guarantees that enough grid points are located on the interface to secure an accurate solution. For velocities close to zero, we lower the time step for a better result.

As initial condition we simply define a jump function $J(x)$: for $T < T_m^{\text{V}}$ we set $J(x)$ equal to one close to the boundaries and zero in between. For $T > T_m^{\text{V}}$ we define $J(x)$ the other way around. In both cases, one has to take care that the intervals where $J(x) = 1$ or $J(x) = 0$ are

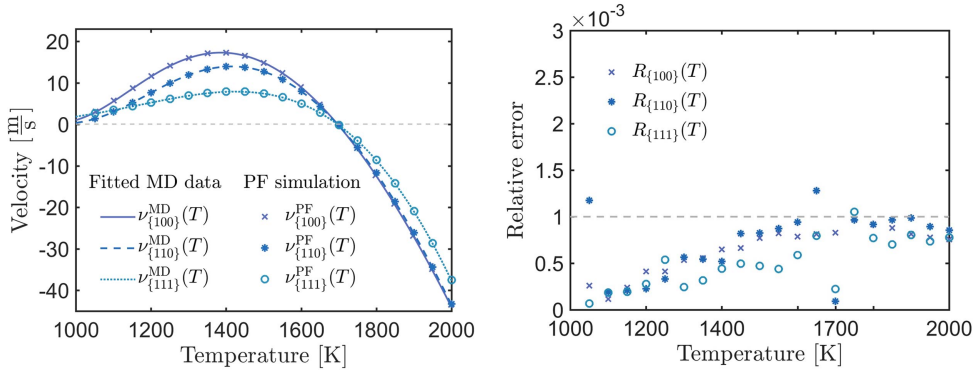


Figure 8. The points in the left figure are the measured velocities from the numerical solution of equation (10) which we calculate with the Fourier-spectral method. The solid lines are the fitted velocities from molecular dynamics. Clearly, model (10) with the atomistic informed parameters reproduces exactly the kinetic interface-behavior. As shown in the right figure, only 5% of the relative errors of the velocities are greater than 10^{-3} . The highest error is about 0.0013 for the {110} orientation at $T = 1650$ K.

wider than the interface thickness, else the system evolves to an equilibrium state before the traveling wave is established.

Our results of the numerical velocity measurement match accurately with the results from molecular dynamics. On the left-hand side of figure 8, the velocities from the phase-field simulation of model (10) are located directly on the line of the Vogel–Fulcher fit of the molecular dynamical data. For a better comparison, we calculate the relative error, which is

$$R_{\{hkl\}}(T) = \left| \frac{v_{\{hkl\}}^{\text{MD}}(T) - v_{\{hkl\}}^{\text{PF}}(T)}{v_{\{hkl\}}^{\text{MD}}(T)} \right|. \quad (23)$$

The relative error is shown on the right-hand side of figure 8. We observe, that the maximal relative error is 0.0013, which is at $T = 1650$ K for orientation {110}. 95% of the errors are even lower than 10^{-3} .

5. Conclusion and outlook

In this study we have extracted the necessary parameters to obtain a phase-field model that can accurately describe the solid–liquid interface kinetics. In particular, using molecular dynamics simulations with the interatomic potential by SW, we derived an expression for the bulk free energy, the interfacial width of the liquid-crystal interface and the crystallization velocity and hence the corresponding anisotropic mobility parameter for three different orientations in silicon as a function of temperature. To properly capture the behavior of the temperature-dependent viscosity near the glass transition a Vogel–Fulcher fit is used for the SW potential. We show that these results are essential to obtain an accurate temperature dependence of the mobility parameter in the corresponding phase-field model for liquid-phase crystallization.

Our approach is presently being extended to two- and three-dimensional setting. Further extensions include the amorphous and poly-crystalline structure as well as defects such as stacking faults, in the free energy density and are expected to prove useful for validation against experimental results of Si recrystallization in the future.

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Appendix

A.1. Determination of the bulk free energy density F

Our approach for the free energy density is a polynomial of fourth degree in p

$$F(p, T) = a_0(T) + a_1(T)p + a_2(T)p^2 + a_3(T)p^3 + a_4(T)p^4, \quad (\text{A.1})$$

which attains the values in figure 2 at the equilibrium states $p = 0$ (liquid phase) and $p = 1$ (crystalline phase). So we need to consider the derivatives of F with respect to p :

$$\frac{\partial F}{\partial p}(p, T) = a_1(T) + 2a_2(T)p + 3a_3(T)p^2 + 4a_4(T)p^3,$$

$$\frac{\partial^2 F}{\partial p^2}(p, T) = 2a_2(T) + 6a_3(T)p + 12a_4(T)p^2.$$

From the conditions for the existence of minima at $p = 0$ and $p = 1$ follows

$$a_1(T) \equiv 0, \quad (\text{A.2})$$

$$a_2(T) > 0, \quad (\text{A.3})$$

$$a_3(T) = -\frac{2}{3}a_2(T) - \frac{4}{3}a_4(T), \quad (\text{A.4})$$

$$a_2(T) < 2a_4(T). \quad (\text{A.5})$$

The polynomials $f_0(T)$ and $f_1(T)$ pass the equilibrium values in figure 2 for liquid/amorphous and crystalline Si, respectively. For the minima, we need for (A.1) the equalities $F(0, T) = f_0(T)$ and $F(1, T) = f_1(T)$ and hence

$$f_0(T) = F(0, T) = a_0(T), \quad (\text{A.6})$$

$$f_1(T) = F(1, T) \stackrel{(\text{A.2}, \text{A.4}, \text{A.6})}{=} f_0(T) + \frac{1}{3}a_2(T) - \frac{1}{3}a_4(T). \quad (\text{A.7})$$

With (A.4) and (A.7) the two coefficients a_3 and a_4 have the form

$$a_3(T) = -2a_2(T) - 4(f_0(T) - f_1(T)) = -2a_2(T) - 4H(T), \quad (\text{A.8})$$

$$a_4(T) = a_2(T) + 3(f_0(T) - f_1(T)) = a_2(T) + 3H(T). \quad (\text{A.9})$$

Together with (A.9) we can verify the inequality (A.5):

$$a_2(T) > -6H(T), \quad (\text{A.10})$$

where the right-hand side of (A.10) is negative if and only if $T < T_m^V$, as one can observe in figure 2. Hence, for temperatures below the melting point, a_2 has to fulfill (A.3). Finally, together with (A.8), (A.9), the double-well potential (A.1) has the form

$$F(p, T) = f_0(T) + a_2(T)p^2 - 2(a_2(T) + 2H(T))p^3 + (a_2(T) + 3H(T))p^4, \quad (\text{A.11})$$

where

$$a_2(T) = \begin{cases} 0 & T \leq T_m^V \\ -6H(T) & T > T_m^V. \end{cases} \quad (\text{A.12})$$

The formulations (A.11) and (A.12) are equivalent to (11) and (12), respectively, by renaming $a_2(T)$ to $a(T)$.

A.2. Derivation of the maximum argument μ

In the following, the calculation of μ is described, where the notation of (11) and (12) is used. We first note that with (12) we have

$$a(T) + 3H(T) > 0, \quad (\text{A.13})$$

which can be seen easily by using the restrictions on a in (12):

First of all we note that for $T > T_m^V$

$$a(T) + 3H(T) \stackrel{(12)}{>} -6H(T) + 3H(T) = -3H(T) \stackrel{T > T_m^V}{>} 0$$

and for $T \leq T_m^V$,

$$a(T) + 3H(T) > 0 + 3H(T) \geq 0.$$

Besides that $p = 0$ and $p = 1$, the first derivative of F

$$\frac{\partial F}{\partial p}(p, T) = 2p(1 - p)[a(T) - (2a(T) + 6H(T))p]$$

has a third root $\mu(T)$

$$\mu(T) = \frac{a(T)}{2a(T) + 6H(T)}.$$

Since (A.13) holds, the denominator of $\mu(T)$ can not become zero. Hence, with (12) we have $\mu(T) > 0$. Further, the condition

$$\mu(T) = \frac{a(T)}{2a(T) + 6H(T)} < 1 \quad (\text{A.14})$$

is equivalent to

$$a(T) > -6H(T) \quad (\text{A.15})$$

and together with (12) we have $\mu(T) \in (0, 1)$ for all $T \geq 0$. In addition, for a maximum in $\mu(T)$, the second derivative of F in $\mu(T)$ has to be less than zero, i.e.

$$\frac{\partial^2 F}{\partial p^2}(\mu(T), T) = \frac{a(T)(a(T) + 6H(T))}{-a(T) - 3H(T)} \stackrel{!}{<} 0. \quad (\text{A.16})$$

Since (A.13) holds, the denominator of (A.16) is negative and we get, since $a(T) > 0$

$$a(T) > -6H(T). \quad (\text{A.17})$$

With (12), the last condition is always true. Hence, the second derivative of F in $\mu(T)$ is less than zero, so $(\mu(T), F(\mu(T), T))$ is a maximum point of F .

Table A1. The calculated mobilities via a shooting method for orientations {100} {110} and {111}. They are shown in figure 7.

Temperature T in K	Mobility $M_{\{100\}}^{\text{PF}}$ in $\text{\AA}^3 (\text{eV ns})^{-1}$	Mobility $M_{\{110\}}^{\text{PF}}$ in $\text{\AA}^3 (\text{eV ns})^{-1}$	Mobility $M_{\{111\}}^{\text{PF}}$ in $\text{\AA}^3 (\text{eV ns})^{-1}$
<800	0	0	0
800	8.8818×10^{-14}	8.8818×10^{-14}	2.93×10^{-1}
850	9.7105×10^{-8}	1.7795×10^{-9}	5.6933×10^{-1}
900	5.8741×10^{-3}	8.4051×10^{-4}	1.0082
950	2.2923×10^{-1}	6.8938×10^{-2}	1.658
1000	1.425	6.2816×10^{-1}	2.5636
1050	3.3476	1.8627	3.761
1100	6.8805	4.4784	5.2861
1150	11.4558	8.3461	7.1611
1200	16.737	13.2755	9.4112
1250	22.4071	18.9898	12.0447
1300	28.2093	25.213	15.06
1350	33.9597	31.7057	18.4506
1400	39.5291	38.274	22.2025
1450	44.8397	44.7766	26.3002
1500	49.7514	51.0148	30.6662
1550	54.1681	56.8348	35.2213
1600	58.1903	62.2964	39.9958
1650	61.8892	67.4445	45.0044
1700	63.6398	70.4575	48.9622
1750	66.429	74.5941	54.0443
1800	68.8907	78.3509	59.2121
1850	70.7543	81.4046	64.1717
1900	71.9325	83.6335	68.749
1950	72.9413	85.6235	73.3545
2000	73.6987	87.2768	77.8687

A.3. Listed mobility values

In section 4.3 we describe the calculation of the mobility parameters for the phase-field model (10). The mobility values are shown in figure 7 and listed in table A1.

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