DEEP LEARNING OF MICROSTRUCTURES

by

Amir Abbas Kazemzadeh Farizhandi

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Amir Abbas Kazemzadeh Farizhandi

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The following individuals read and discussed the dissertation submitted by student Amir Abbas Kazemzadeh Farizhandi, and they evaluated their presentation and response to questions during the final oral examination. They found that the student passed the final oral examination.

Mahmood Mamivand, Ph.D. Chair, Supervisory Committee
Edoardo Serra, Ph.D. Member, Supervisory Committee
Eric Jankowski, Ph.D. Member, Supervisory Committee

The final reading approval of the dissertation was granted by Mahmood Mamivand, Ph.D., Chair of the Supervisory Committee. The dissertation was approved by the Graduate College.
I dedicate my dissertation work to my family. A special feeling of gratitude to my loving wife and parents, whose words of encouragement and push for tenacity ring in my ears.
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ABSTRACT

The internal structure of materials also called the microstructure plays a critical role in the properties and performance of materials. The chemical element composition is one of the most critical factors in changing the structure of materials. However, the chemical composition alone is not the determining factor, and a change in the production process can also significantly alter the materials' structure. Therefore, many efforts have been made to discover and improve production methods to optimize the functional properties of materials. The most critical challenge in finding materials with enhanced properties is to understand and define the salient features of the structure of materials that have the most significant impact on the desired property. In other words, by process, structure, and property (PSP) linkages, the effect of changing process variables on material structure and, consequently, the property can be examined and used as a powerful tool in material design with desirable characteristics. In particular, forward PSP linkages construction has received considerable attention thanks to the sophisticated physics-based models. Recently, machine learning (ML), and data science have also been used as powerful tools to find PSP linkages in materials science. One key advantage of the ML-based models is their ability to construct both forward and inverse PSP linkages. Early ML models in materials science were primarily focused on process-property linkages construction. Recently, more microstructures are included in the materials design ML models. However, the inverse design of microstructures, i.e., the prediction of
process and chemistry from a microstructure morphology image have received limited attention. This is a critical knowledge gap to address specifically for the problems that the ideal microstructure or morphology with the specific chemistry associated with the morphological domains are known, but the chemistry and processing which would lead to that ideal morphology are unknown.

In this study, first, we propose a framework based on a deep learning approach that enables us to predict the chemistry and processing history just by reading the morphological distribution of one element. As a case study, we used a dataset from spinodal decomposition simulation of Fe-Cr-Co alloy created by the phase-field method. The mixed dataset, which includes both images, i.e., the morphology of Fe distribution, and continuous data, i.e., the Fe minimum and maximum concentration in the microstructures, are used as input data, and the spinodal temperature and initial chemical composition are utilized as the output data to train the proposed deep neural network. The proposed convolutional layers were compared with pretrained EfficientNet convolutional layers as transfer learning in microstructure feature extraction. The results show that the trained shallow network is effective for chemistry prediction. However, accurate prediction of processing temperature requires more complex feature extraction from the morphology of the microstructure. We benchmarked the model predictive accuracy for real alloy systems with a Fe-Cr-Co transmission electron microscopy micrograph. The predicted chemistry and heat treatment temperature were in good agreement with the ground truth. The treatment time was considered to be constant in the first study.

In the second work, we propose a fused-data deep learning framework that can predict the heat treatment time as well as temperature and initial chemical compositions
by reading the morphology of Fe distribution and its concentration. The results show that
the trained deep neural network has the highest accuracy for chemistry and then time and
temperature. We identified two scenarios for inaccurate predictions; 1) There are several
paths for an identical microstructure, and 2) Microstructures reach steady-state
morphologies after a long time of aging. The error analysis shows that most of the wrong
predictions are not wrong, but the other right answers. We validated the model
successfully with an experimental Fe-Cr-Co transmission electron microscopy
micrograph.

Finally, since the data generation by simulation is computationally expensive, we
propose a quick and accurate Predictive Recurrent Neural Network (PredRNN) model for
the microstructure evolution prediction. Essentially, microstructure evolution prediction
is a spatiotemporal sequence prediction problem, where the prediction of material
microstructure is difficult due to different process histories and chemistry. As a case
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<td>ML</td>
<td>Machine Learning</td>
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<tr>
<td>3D</td>
<td>Three-Dimensional</td>
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<td>2D</td>
<td>Two-Dimensional</td>
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<td>PF</td>
<td>Phase Field</td>
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<td>AI</td>
<td>Artificial Intelligence</td>
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<td>Recurrent Neural Network</td>
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<td>CNN</td>
<td>Convolutional Neural Network</td>
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<tr>
<td>LSTM</td>
<td>Long Short-Term Memory</td>
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<td>PCA</td>
<td>Principal Component Analysis</td>
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<td>RVE</td>
<td>Representative Volume Element</td>
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<tr>
<td>TEM</td>
<td>Transmission Electron Microscopes</td>
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<tr>
<td>MSE</td>
<td>Mean Squared Error</td>
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<td>PSNR</td>
<td>Peak Signal-to-Noise Ratio</td>
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CHAPTER ONE: INTRODUCTION

This Ph.D. dissertation aims to develop a framework based on deep learning (DL) to enable chemistry and process history prediction behind a microstructure as well as microstructure morphology prediction through microstructure evolution. The developed models enable the prediction of processing history and chemical compositions from microstructure images and predict microstructure morphologies without expensive and time-consuming simulations and experiments. Doing so will provide the materials science community with knowledge and algorithms that can be used for new materials development with the desired properties. In this way, we reviewed the previous studies for using machine learning in the construction of PSP linkages and material microstructure evolution prediction.

Process-Structure-Property Linkages

Heterogeneous materials are widely used in various industries, such as aerospace, automotive, and construction. These materials’ properties greatly depend on their microstructure, which is a function of the chemical composition and operational process of materials production. To accelerate the novel materials design process, the construction of process-structure-property (PSP) linkages is necessary. Establishing PSP linkages with sole experiments is not practical as the process is costly and time-consuming. Therefore, computational methods are used to study the structure of materials and their properties. A basic assumption for computational modeling of materials is that they are periodic on the microscopic scale and can be approximated by representative
elements (RVE) [1]. Finding the effects of process conditions and the chemical composition on the characteristics of the RVE, such as volume fraction, microstructure, grain size, and, consequently, the materials’ properties, will lead to the development of PSP linkages. In the past two decades, the phase-field (PF) method has been increasingly used as a robust method for studying the spatiotemporal evolution of the materials’ microstructure and physical properties [2]. It has been widely used to simulate different evolutionary phenomena, including grain growth and coarsening [3], solidification [4], thin-film deposition [5], dislocation dynamics [6], vesicle formation in biological membranes [7], and crack propagation [8]. PF models solve a system of partial differential equations (PDEs) for a set of continuous variables of the processes. However, solving high-fidelity PF equations is inherently computationally expensive because it requires solving several coupled PDEs simultaneously [9]. Therefore, PSP construction, particularly for complex materials, only based on the PF method is inefficient. To address this challenge, machine learning (ML) methods have recently been proposed as an alternative for creating PSP linkages based on the limited experimental/simulation data or both [10]. Artificial intelligence (AI), ML, and data science are beneficial in speeding up and simplifying the process of discovering new materials [11]. Developing and deploying an appropriate support data infrastructure that efficiently integrates closed-loop iterations between experimentation and multi-scale modeling/simulation efforts is climacteric. This need is addressed by a new interdisciplinary field called Materials Data Science and Informatics [12-18].

A fundamental element of the data science approach is a multi-faceted framework that enables the research community to collect, aggregate, nurture, disseminate, and reuse
valuable knowledge. In materials innovation efforts, this knowledge is primarily desired in the form of length and time scale PSP linkages associated with the material system of interest [19-24]. In a multi-scale materials modeling effort, this means developing a formal data science approach to extract reusable PSP linkages from an ensemble of simulation and experiment datasets, as depicted in Figure 1.1. The top arrow in Figure 1.1, forward design philosophy, shows a typical workflow that materials scientists historically have used in developing PSP linkages. In forward design philosophy, we loop through the ordered connection of process-structure-property. Forward design usually involves the use of experiments and advanced physics in combination with numerical algorithms. Generally, since material discovery requires exploration of big space, the forward design is prone to result in high costs and time. This cost can be a significant obstacle to materials innovation efforts, even in the realm of simulations, as these simulations are often expensive, and the design space is huge. This is precisely where the data science approaches offer many benefits. As shown in Figure 1.1, the data science tools and algorithms can enable us to perform inverse design, i.e., start from the desired properties and find the required processing. With the full advantage of advanced statistics and machine learning techniques, data science can provide a mathematically rigorous framework for PSP linkage in multi-scale material design. As depicted in Figure 1.1, one of the main benefits of adding data science components to the entire workflow is that it is very practical to solve the inverse problem, which is the ultimate goal of materials innovation efforts. In fact, materials informatics provides a low computational approach for materials design. This is mainly because the PSP linkages are cast as metamodels or
surrogate models. These models can be easily used to find the optimum conditions for making materials with desired properties.

In recent years, using data science in various fields of materials science has increased significantly [25-30]. For instance, data science is applied to help density functional theory calculations to establish a relationship between atoms’ interaction with the properties of materials based on the quantum mechanics [31-34]. AI is also utilized to establish PSP linkages in the context of materials mechanics. In this case, ML can be used to design new materials with desired properties or employed to optimize the production process of the existing materials for properties improvement. Through data science, researchers will be able to examine the complex and nonlinear behavior of a materials production process that directly affects the materials’ properties [35]. Many studies have focused on solving cause-effect design, i.e., finding the material properties from the microstructure or processing history. These studies have attempted to predict the structure of the materials from processing parameters or materials properties from the microstructure and processing history [10, 25, 36-43]. A less addressed but essential problem is a goal-driven design that tries to find the processing history of the materials from their microstructures. In these cases, the optimal microstructure that provides the optimal properties is known, e.g., via physics-based models, and it is desirable to find the chemistry and processing routes that would lead to the desirable microstructure.
The use of microstructure images in ML modeling is challenging. The microstructure quantification has been reported as the central nucleus in the PSP linkages construction [37]. Microstructure quantification is important from two perspectives. First, it can increase the accuracy of the developed data-driven model. Second, an in-depth understanding of the microstructures can improve the comprehension of the effects of process variables and chemical composition on the properties of materials [37]. In recent years, deep learning (DL) methods have been successfully used in other fields, such as computer vision. Their limited applications in materials science have also proven them as reliable and promising methods [38]. The main advantages of DL methods are their simplicity, flexibility, and applicability for all types of microstructures. Furthermore, DL
has been broadly applied in materials science to improve the targeted properties [34, 39-46]. One form of DL models that has been extensively used for feature extraction in various applications such as image, video, voice, and natural language processing is Convolutional Neural Networks (CNN) [47-50]. In materials science, CNN has been used for various image-related problems. Cang et al. used CNN to achieve a 1000-fold dimension reduction from the microstructure space [51]. DeCost et al. [52] applied CNN for microstructure segmentation. Xie and Grossman [53] used CNN to quantify the crystal graphs to predict the material properties. Their developed framework was able to predict eight different material properties such as formation energy, bandgap, and shear moduli with high accuracy. CNN has also been employed to index the electron backscatter diffraction patterns and determine the crystalline materials’ crystal orientation [54]. The stiffness in two-phase composites has been predicted successfully by the deep learning approach, including convolutional and fully-connected layers [55]. In a comparative study, the CNN and the materials knowledge systems (MKS), proposed in the Kalidindi group based on the idea of using the n-point correlation method for microstructures quantification [56-58], were used for microstructure quantification and then, the produced data were employed to predict the strain in the microstructural volume elements. The comparison showed that the extracted features by CNN could provide more accurate predictions [59]. Cecen et al. [20] proposed CNN to find the salient features of a collection of 5900 microstructures. The results showed that the obtained features from CNN could predict the properties more accurately than the 2-point correlation, while the computation cost was also significantly reduced. Comparing DL approaches, including CNN, with the MKS method, single-agent, and multi-agent
methods shows that DL always performs more accurately [59-61]. Zhao et al. utilized the
electronic charge density (ECD) as a generic unified 3D descriptor for elasticity
prediction. The results showed a better prediction power for bulk modulus than for shear
modulus [62]. CNN has also been applied for finding universal 3D voxel descriptors to
predict the target properties of the solid-state material [63]. The introduced descriptors
outperformed the other descriptors in the prediction of Hartree energies for solid-state
materials.

Training a deep CNN usually requires an extensive training dataset that is not always
available in many applications. Therefore, a transfer learning method that uses a
pretrained network can be applied for new applications. In transfer learning, all or a part
of the pretrained networks such as VGG16, VGG19 [64], Xception [65], ResNet [66],
and Inception [67], which were trained by the computer vision research community with
lots of open source image datasets such as ImageNet, MS, CoCo, and Pascal, can be used
for the desired application. In particular, in materials science which generally the image-
based data are not greatly abundant, transfer learning could be beneficial. DeCost et al.
[68] adopted VGG16 to classify the microstructures based on their annealing conditions.
Ling et al. [25] applied VGG16 to extract the feature from scanning electron microscope
(SEM) images and classify them. Lubbers et al. [69] used the VGG 19 pretrained model
to identify the physical meaningful descriptors in microstructures. Li et al. [70] proposed
a framework based on VGG19 for microstructure reconstruction and structure-property
predictions. The pretrained VGG19 network was also utilized to reconstruct the 3D
microstructures from 2D microstructures by Bostanabad [71].
Review provided above shows that the majority of the ML-microstructure related works in the materials science community were primarily focused on using ML techniques for microstructure classification [72-74], recognition [75], microstructure reconstruction [70, 71], or as a feature-engineering-free framework to connect microstructure to the properties of the materials [55, 76, 77]. However, the process and chemistry prediction from a microstructure morphology image has received limited attention. This is a critical knowledge gap to address specifically for the problems in them the ideal microstructure or morphology with the specific chemistry associated with the morphology domains are known, but the chemistry and processing which would lead to that ideal morphology is unknown. The problem becomes much more challenging for multicomponent alloys with complex processing steps. Recently, Kautz et al. [77] have used the CNN for microstructure classification and segmentation on Uranium alloyed with 10 wt% molybdenum (U-10Mo). They used the segmentation algorithm to calculate the area fraction of the lamellar transformation products of $\alpha-U + \gamma$-UMo, and by feeding the total area fraction into the Johnson-Mehl-Avrami-Kolmogorov equation, they were able to predict the annealing parameters, i.e., time and temperature. However, Kautz et al.’s [77] work for aging time prediction did not consider the morphology and particle distribution, and also, no chemistry was involved in the model. To address the knowledge gap, in this work, we develop a mixed-data deep neural network that is capable to predict the chemistry and processing history of a micrograph. The model alloy used in this work is Fe-Cr-Co permanent magnets.
Materials Microstructure Evolution Prediction

The processing-structure-property relationship of engineered materials is directly impacted by material microstructures, which are mesoscale structural elements that operate as an essential link between atomistic building components and macroscopic qualities. One of the pillars of contemporary materials research is the ability to manage the evolution of the material's microstructure while it is being processed or used, including common phenomena like solidification, solid-state phase transitions, and grain growth. Therefore, a key objective of computational materials design has been comprehending and forecasting of microstructure evolution. Simulations of microstructure evolution frequently rely on phase separation or coarse-grained models, such as partial differential equations (PDEs), which are used in the phase-field techniques [2, 78] because they can represent time and length scales that are far larger than those that can be captured by molecular dynamics. A wide range of significant evolutionary mesoscale processes, including grain development and coarsening, solidification, thin-film deposition, dislocation dynamics, vesicle formation in biological membranes, and crack propagation, have all been fully described using the phase-field method [3-8]. However, there are some significant problems with this strategy as well. First off, PDE-based microstructure simulations are still relatively expensive. The stability of numerical techniques that use explicit time integration for nonlinear PDEs sets stringent upper bounds on the smallest time-step size in the temporal dimension.

Similarly, implicit time-integration techniques manage longer time steps by adding more inner iteration loops at each step. Furthermore, despite the fact that in theory controlling PDEs can be inferred from the underlying thermodynamic and kinetic
considerations, actual PDE identification, parametrization, and validation take a significant amount of work. The evolution principles may not be fully understood or may be too complex to be characterized by tractable PDEs for difficult or less well-studied materials. Currently, the efforts to reduce computational costs have mostly concentrated on utilizing high-performance computer architectures [79-82] and sophisticated numerical techniques [83, 84], or on merging machine learning algorithms with simulations based on microstructures [43, 85-91]. Leading studies, for instance, have developed surrogate models using a variety of techniques, such as Green's function solution [85], Bayesian optimization [43, 86], a combination of dimensionality reduction and autoregressive Gaussian processes [88], convolutional autoencoder and decoder [92], or integrating a history-dependent machine-learning method with a statistically representative, low dimensional description of the microstructure evolution generated directly from phase-field simulations that can quickly predict the evolution of the microstructure from phase-field simulations [9]. The main problem, however, has been to strike a balance between accuracy and computing efficiency, even for these successful systems. For complex, multi-variable phase-field models, for example, precise answers cannot be guaranteed by the computationally effective Green's function solution. In contrast, complex, coupled phase-field equations can be solved using Bayesian optimization techniques, however, at a higher computational cost (although the number of simulations required is kept to a minimum because the Bayesian optimization protocol determines the parameter settings for each subsequent simulation). The capacity of this class of models to predict future values outside of the training set is constrained by the fact that autoregressive models can only forecast microstructural evolution for the values
for which they were trained. In other models based on dimensionality reduction methods like principle component analysis (PCA), a large amount of information is ignored, which will sacrifice accuracy. This study uses Predictive Recurrent Neural Network (PredRNN) [93] to forecast how the microstructure represented by 2D image sequences will change over time.

**Dissertation Structure**

The dissertation outline is as follows. In chapter two, we describe the methods used in this study, like PF, deep learning methods, train and test data generation, and error metrics.

In chapter three, we develop a mixed-data deep neural network capable of predicting a micrograph's chemistry and processing history. The model alloy used in this work is Fe-Cr-Co permanent magnets. These alloys experience spinodal decomposition at temperatures around 853 – 963 K. We use the PF method to create the training and test dataset for the DL network. CNN will quantify the produced microstructures by the PF method; then the salient features will be used by another deep neural network to predict the temperature and chemical composition.

In chapter four, we explain a model based on a deep neural network to predict a complete set of processing parameters, including temperature, time, and chemistry from a microstructure micrograph. As a case study, we focused on the spinodal decomposition process, and to prove the model applicability for realistic alloys, we picked the Fe-Cr-Co permanent magnets as the model alloy. We used the PF method to create the training and test datasets for deep network training. A fused dataset including material microstructure
as well as minimum and maximum iron concentration in the microstructure is used as the input data. We quantified the generated microstructures with the CNN and then combined the extracted salient features from the microstructures with iron composition to predict the processing history, i.e., annealing time and temperature, and chemical compositions of the micrograph.

In chapter five, we describe the microstructure evolution prediction by PredRNN. In this study, spinodal decomposition is used as a case study. Spinodal decomposition occurs in two separate phases: a quick composition modulation growth phase, followed by a slower coarsening phase, during which the Gibbs-Thomson effect causes a progressive rise in the length scale of the phase-separation pattern. We demonstrate that PredRNN can precisely capture all the required features from earlier microstructures to predict long-term microstructures. This result is particularly important because it can predict morphology evolution in both phases.

Finally, the key findings in this thesis are summarized and possible opportunities for future works are discussed.

This dissertation leads to the following peer-reviewed and conference papers;

2. Amir Abbas Kazemzadeh Farizhandi, Omar Betancourt, Mahmood Mamivand. Di
Deep learning approach for chemistry and processing history prediction from m
https://doi.org/10.1038/s41598-022-08484-7.

3. Amir Abbas Kazemzadeh Farizhandi, Mahmood Mamivand, Spatiotemporal P
Prediction of Microstructure Evolution with Predictive Recurrent Neural N
Network, Submitted to Materials & Design.

4. Amir Abbas Kazemzadeh Farizhandi, Mahmood Mamivand, Chemistry and P
Processing History Prediction from Materials Microstructure by Deep Learning, 2
2022 TMS Annual Meeting & Exhibition, Symposium: Algorithm Development i
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5. Amir Abbas Kazemzadeh Farizhandi, Mahmood Mamivand, Spatiotemporal P

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   2023
CHAPTER TWO: METHOD

In this chapter, we describe the models and algorithms that we have used in this dissertation. First, we briefly describe the Phase Field (PF) model for spinodal decomposition simulation and the dataset generation process. Then, we provide the details of the proposed fused-data deep neural network for process history and chemistry prediction from materials microstructure morphologies. Finally, we explain the LSTM model that has been used to predict the evolution of material microstructures along with an overview of methods to quantify the similarity between microstructure morphologies.

**Phase Field Method**

With the enormous increase in computational power and advances in numerical methods, the PF approach has become a powerful tool for the quantitative modeling of microstructures' temporal and spatial evolution. Some applications of this method include modeling materials undergoing martensitic transformation [94], crack propagation [95], grain growth [96], and materials microstructure prediction for optimization of their properties [97].

The PF method eliminates the need for the system to track each moving boundary by having the interfaces to be of finite width where they gradually transform from one composition or phase to another [2]. This essentially causes the system to be modeled as a diffusivity problem, which can be solved by using the continuum nonlinear PDEs. There are two main PF PDEs for representing the evolution of various PF variables. One being the Allen-Cahn equation [98] for solving non-conserved order parameters (e.g.,
phase regions and grains), and the other one being the Cahn-Hilliard equation [99] for solving conserved order parameters (e.g., concentrations).

Since the diffusion of constituent elements controls the process of phase separation, we only need to track the conserved variables, i.e., Fe, Cr, and Co concentration, during isothermal spinodal phase decomposition. Thus, our model will be governed by Cahn-Hilliard equations. The PF model in this work is primarily adopted from [100]. For the spinodal decomposition of the Fe-Cr-Co ternary system, the Cahn-Hilliard equations are,

\[
\frac{\partial c_{Cr}}{\partial t} = \nabla \cdot M_{Cr,Cr} \nabla \frac{\delta F_{tot}}{\delta c_{Cr}} + \nabla \cdot M_{Cr,Co} \nabla \frac{\delta F_{tot}}{\delta c_{Co}},
\]

(1)

\[
\frac{\partial c_{Co}}{\partial t} = \nabla \cdot M_{Co,Cr} \nabla \frac{\delta F_{tot}}{\delta c_{Cr}} + \nabla \cdot M_{Co,Co} \nabla \frac{\delta F_{tot}}{\delta c_{Co}}.
\]

(2)

The microstructure evolution is primarily driven by the minimization of the total free energy \( F_{tot} \) of the system. The free energy functional, using \( N \) conserved variables \( c_i \) at the location \( \vec{r} \) is described by:

\[
F_{tot} = \int \left( f_{loc}(c_1, \ldots, c_N, T) + f_{gr}(c_1, \ldots, c_N) \right) d\vec{r} + E_{el}.
\]

(3)

In this model, \( N=3 \) conserved variables are \( c_{Fe}, c_{Cr}, \) and \( c_{Co} \), and they denote the composition of Fe, Cr, and Co, respectively. \( f_{gr} \) is the gradient energy density and is described by

\[
f_{gr} = \frac{\kappa}{2} \sum_{l} |\nabla c_l|^2,
\]

(4)

where \( \kappa \) is the gradient energy coefficient. In this case, \( \kappa \) is considered a constant value. \( f_{loc} \) is the local Gibbs free energy density as a function of all concentrations, \( c_i \), and temperature, \( T \). For this work, we will model the body-centered cubic phase of Fe-Cr-Co, where the Gibbs free energy of the system is described as [100],
\[ f_{\text{loc}} = f_{\text{Fe}}^0 c_{\text{Fe}} + f_{\text{Cr}}^0 c_{\text{Cr}} + f_{\text{Co}}^0 c_{\text{Co}} + RT(c_{\text{Fe}} \ln c_{\text{Fe}} + c_{\text{Cr}} \ln c_{\text{Cr}} + c_{\text{Co}} \ln c_{\text{Co}}) + f^E + f^{mg}, \]  

where \( f_i^0 \) is the Gibbs free energy of the pure element \( i \) and \( f^E \) is the excess free energy defined by

\[ f^E = L_{\text{Fe,Cr}} c_{\text{Fe}} c_{\text{Cr}} + L_{\text{Fe,Co}} c_{\text{Fe}} c_{\text{Co}} + L_{\text{Cr,Co}} c_{\text{Cr}} c_{\text{Co}}, \]

where \( L_{\text{Fe,Cr}}, L_{\text{Fe,Co}}, \) and \( L_{\text{Cr,Co}} \) are interaction parameters. \( f^{mg} \) is the magnetic energy contribution and can be expressed as

\[ f^{mg} = RT \ln(\beta + 1) f(\tau), \]

where \( \beta \) is the atomic magnetic moment, \( f(\tau) \) is a function of \( \tau \equiv T/T_C \). \( T_C \) is the Curie temperature. \( E_{\text{el}} \) in Eq. (3) is the elastic strain energy added to the system and is expressed as

\[ E_{\text{el}} = \frac{1}{2} \int_{\mathbb{F}} C_{ijkl} \varepsilon_{ij}^{\text{el}}(\vec{r}, t) \varepsilon_{kl}^{\text{el}}(\vec{r}, t) d\vec{r}, \]

\[ \varepsilon_{ij}^{\text{el}}(\vec{r}, t) = \varepsilon_{ij}^\ell(\vec{r}, t) - \varepsilon_{ij}^0(\vec{r}, t), \]

where \( \varepsilon_{ij}^\ell(\vec{r}, t) \) is the elastic strain and \( C_{ijkl} \) are the elastic coefficients of the stiffness tensor. \( \varepsilon_{ij}^0(\vec{r}, t) \) is the eigen-strain and is expressed by

\[ \varepsilon_{ij}^0(\vec{r}, t) = [\varepsilon_{\text{Cr}}(c_{\text{Cr}}(\vec{r}, t) - c_{\text{Cr}}^0) + \varepsilon_{\text{Co}}(c_{\text{Co}}(\vec{r}, t) - c_{\text{Co}}^0)] \delta_{ij}, \]

where \( \varepsilon_{\text{Cr}} \) and \( \varepsilon_{\text{Co}} \) are lattice mismatches between Cr with Fe and Co with Fe, respectively. \( c_{\text{Cr}}^0 \) and \( c_{\text{Co}}^0 \) are the initial concentrations of Cr and Co, respectively and \( \delta_{ij} \) is the Kronecker delta. The constrained strain, \( \varepsilon_{ij}^c(\vec{r}, t) \), is solved using the finite element method.

\( M_{ij} \) in Eq. (2) are Onsager coefficients and are scalar mobilities from the coupled system involving the concentrations. They can be determined by [100],
\[ M_{Cr,Cr} = [c_{Fe}c_{Cr}M_{Fe} + (1 - c_{Cr})^2M_{Cr} + c_{Cr}c_{Co}M_{Co}] \frac{c_{Cr}}{RT}, \]  
\[ M_{Co,Co} = [c_{Fe}c_{Co}M_{Fe} + c_{Cr}c_{Co}M_{Cr} + (1 - c_{Co})^2M_{Co}] \frac{c_{Co}}{RT}, \]  
\[ M_{Cr,Co} = M_{Co,Cr} = [c_{Fe}M_{Fe} - (1 - c_{Cr})M_{Cr} - (1 - c_{Co})M_{Co}] \frac{c_{Cr}c_{Co}}{RT}. \]

The mobility \( M_i \) of each element \( i \) is determined by

\[ M_i = D_i^0 \exp \left( -\frac{Q_i}{k_B T} \right), \]

where \( D_i^0 \) is the self-diffusion coefficient and \( Q_i \) is the diffusion activation energy.

We parametrized the model with the calculation of phase diagram (CALPHAD) data [100]. To solve the non-linear CH partial differential equations (PDEs), we used the Multi-physics Object-Oriented Simulation Environment (MOOSE). MOOSE is an open-source finite element package developed at Idaho National Laboratory and efficient for parallel computation on supercomputers [101]. The coupled CH equations were solved with the help of MOOSE's prebuilt series of weak form residuals of CH PDEs with the input parameters given in Table 2.1.
Table 2.1 Phase-field model input parameters [100, 102, 103]

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Gas constant, $R$ $[J/(K\cdot mol)]$</td>
<td>8.31446261815324</td>
</tr>
<tr>
<td>Gradient energy coefficient, $\kappa$ $[m^2/mol]$</td>
<td>$1.0 \times 10^{-14}$</td>
</tr>
<tr>
<td>Gibbs free energy of the pure element $i$, $f_i^0$ $[J/mol]$</td>
<td>$f_{Fe}^0 = f_{Cr}^0 = f_{Co}^0 = 0$</td>
</tr>
<tr>
<td>Interaction parameters, $L_{ij}$ $[J/mol]$</td>
<td>$L_{Fe,Cr} = 20500-9.68T$</td>
</tr>
<tr>
<td></td>
<td>$L_{Fe,Co} = -23669 + 103.96277T - 12.78867T \ln T$</td>
</tr>
<tr>
<td></td>
<td>$L_{Cr,Co} = (24357 - 19.7977T) - 2010(c_3 - c_2)$</td>
</tr>
<tr>
<td>Curie temperature, $T_C [K]$</td>
<td>$T_C = 1043c_{Fe} - 311.5c_{Cr} + 1450c_{Co}$</td>
</tr>
<tr>
<td></td>
<td>+ ${1650 + 550(c_{Cr} - c_{Fe})}c_{Fe}c_{Cr} + 590c_{Fe}c_{Co}$</td>
</tr>
<tr>
<td>Atomic magnetic moment, $\beta$ $[J/T]$</td>
<td>$\beta = 2.22c_{Fe} - 0.01c_{Cr} + 1.35c_{Co} - 0.85c_{Fe}c_{Cr}$</td>
</tr>
<tr>
<td></td>
<td>+ ${2.4127 + 0.2418(c_{Cr} - c_{Fe})}c_{Fe}c_{Co}$</td>
</tr>
</tbody>
</table>
| Magnetic contribution’s $\tau$ function, $f(\tau) [-]$                  | $f(\tau) = \begin{cases} \tau^2 - \frac{1}{A} + \frac{1}{140p} & \text{if } \tau < 1 \\ \\
|                                                                          | \frac{1}{2} \left( \frac{\tau^2 - \frac{1}{A} + \frac{1}{140p}}{10} + \frac{1}{315} + \frac{1}{1500} \right) & \text{if } \tau \geq 1 \\
|                                                                          | A = \left( \frac{518}{1125} + \frac{11692}{15975} \right) (1 - \frac{1}{p}) & \text{if } \tau < 1 \\
|                                                                          | \text{for } p = 0.4 \text{ for bcc}                                 |
| Elastic coefficients, $C_{ijkl}$ $[GPa]$                                  | $C_{11}^{Fe} = C_{11}^{Co} = 233.1$                                |
|                                                                          | $C_{12}^{Fe} = C_{12}^{Co} = 135.44$                                |
|                                                                          | $C_{44}^{Fe} = C_{44}^{Co} = 117.83$                                |
|                                                                          | $C_{11}^{Cr} = 350$                                                 |
|                                                                          | $C_{12}^{Cr} = 67.8$                                                |
|                                                                          | $C_{44}^{Cr} = 100.8$                                               |
| Lattice mismatch, $\varepsilon_i [-]$                                    | $\varepsilon_{Cr} = 6.1 \times 10^{-3}$                            |
|                                                                          | $\varepsilon_{Co} = -7.1 \times 10^{-3}$                            |
| Self-diffusion coefficient of element $i$, $D_i^0$ $[m^2/s]$             | $D_{Fe}^0 = D_{Co}^0 = 1.0 \times 10^{-4}$                           |
|                                                                          | $D_{Cr}^0 = 2.0 \times 10^{-5}$                                     |
| Diffusion activation energy of element $i$, $Q_i$ $[kJ/mol]$              | $Q_{Fe} = Q_{Co} = 294$                                             |
|                                                                          | $Q_{Cr} = 308$                                                      |
Dataset Generation

Dataset Generation for Steady State Case Study

Since the compositions are subject to the constraint that they must sum to one, the dataset was produced based on the mixture design as a design of experiments method [104]. The Simplex-Lattice [105] designs were adopted to provide the data for simulation. The simulation variables and their range of values are given in Table 2.2. The simulations were run on Boise State University R2 cluster computers [106] using the MOOSE framework [101].

Table 2.2 Simulation variables and their range of values for database generation of steady state case study.

<table>
<thead>
<tr>
<th>Simulation variable</th>
<th>Range of values</th>
<th>Grid</th>
</tr>
</thead>
<tbody>
<tr>
<td>Temperature (K)</td>
<td>853 - 963</td>
<td>10</td>
</tr>
<tr>
<td>Chromium composition</td>
<td>0.05 - 0.9</td>
<td>0.05</td>
</tr>
<tr>
<td>Cobalt composition</td>
<td>0.05 - 0.9</td>
<td>0.05</td>
</tr>
</tbody>
</table>

After running the simulations, the microstructures were collected from the results showing the phase separation. The extracted microstructures for Fe, i.e., the morphology of Fe distribution, from the PF simulations, along with the minimum and maximum compositions of Fe in each microstructure, are utilized as the inputs to predict spinodal temperature, Cr, and Co compositions as processing history parameters. Indeed, the input data is a mixed dataset combined of microstructures, as image data, and Fe composition, as numerical or continuous data. Since these values constitute different data types, the machine learning model must be able to ingest the mixed data. In general, handling the
mixed data is challenging because each data type may require separate preprocessing steps, including scaling, normalization, and feature engineering [107].

**Dataset Generation for Unsteady State Case Study**

To develop proper training and test datasets, we need to span the possible ranges of input variables, i.e., time, temperature, and chemical compositions. For the temperature, we are bonded to the range of 850 – 970 K, as spinodal decomposition in Fe-Cr-Co happens in this window. For chemistry, we explore the range of 0.05-0.9 at. % for both Cr and Co. Since the chemistry is subjected to the conservation of mass constrain, i.e., \( c_{Fe} + c_{Cr} + c_{Co} = 1 \), we used the Simplex-Lattice [105] as a mixture design method to generate the chemistry space to explore. Finally, we bounded the dataset to 300 hours for the time, as our study showed most microstructures would reach equilibrium to some extent by this time. Unlike temperature and chemistry, we did not grid the time domain linearly because the microstructure is very sensitive to aging time in the early stages of annealing, but this sensitivity drops dramatically as time passes. Therefore, we picked a fine grid at the beginning, 50 s, and increased it exponentially, to 100000 s, with time. The variables and their ranges are given in Table 2.3. To cover all the range of input variables, the dataset was generated based on the design of the experiment (DOE). We generated the microstructures by solving the CH PDEs using the MOOSE framework [101]. The simulations were run on different clusters including Boise State University R2 cluster computers [106], Boise State University BORAH [108], and the Extreme Science and Engineering Discovery Environment (XSEDE) (Jetstream2 cluster), which is supported by National Science Foundation (NSF) [109] using the MOOSE framework.
We note that because of the deterministic nature of the PF technique, i.e., not being stochastic, and the physics of the spinodal decomposition, we only need to run each condition once.

**Table 2.3 Simulation variables and their range of values for database generation of unsteady state case study.**

<table>
<thead>
<tr>
<th>Simulation variable</th>
<th>Range of values</th>
<th>Grid</th>
</tr>
</thead>
<tbody>
<tr>
<td>Time (S)</td>
<td>10 - 1080000</td>
<td>10</td>
</tr>
<tr>
<td></td>
<td>10 - 3600</td>
<td>50</td>
</tr>
<tr>
<td></td>
<td>3600 - 36000</td>
<td>500</td>
</tr>
<tr>
<td></td>
<td>36000 - 360000</td>
<td>5000</td>
</tr>
<tr>
<td></td>
<td>360000 - 1080000</td>
<td>10000</td>
</tr>
<tr>
<td>Temperature (K)</td>
<td>850 - 970</td>
<td>10</td>
</tr>
<tr>
<td>Chromium composition</td>
<td>0.05 - 0.9</td>
<td>0.05</td>
</tr>
<tr>
<td>Cobalt composition</td>
<td>0.05 - 0.9</td>
<td>0.05</td>
</tr>
</tbody>
</table>

After simulations, we collected the morphology of Fe distribution, which represents the Fe-rich and Fe-depleted, i.e., Cr-rich, regions, as image data. In addition, we used the minimum and maximum compositions of Fe in each microstructure as numeric data. The deep network uses images and numeric data as input to predict the time, temperature, and chemical compositions. Therefore, different types of deep networks like convolutional and fully-connected layers are required to process the input data. We note that the accuracy of the model will increase for real materials if some experimental data is added to the training dataset. However, even having the experimental dataset to be just a few percent of the whole dataset, requires hundreds of tailored transmission electron microscopy (TEM) images. Generating such a big experimental dataset is time-
consuming and costly. Therefore, in this work, we limit the model to synthetic data. However, as we will show in the validation part, the model predicts the history of an experimental TEM image pretty well, because we are using a CALPHAD-informed phase field model to generate the training and test dataset, and CALPHAD is inherently informed by some experimental data.

Dataset Generation for Microstructure Evolution Case Study

The produced microstructures in the unsteady state case study are also used as training, validation, and test data in this study. The Fe-based composition microstructure morphologies sequences are utilized to construct the dataset. The length of each sequence is 20 microstructures; the first 10 microstructures until 30 hr of the process are used to predict the future 10 microstructures until 300 hr.

Deep Learning Methodology

Deep learning (DL), as an artificial intelligence (AI) tool, is usually used for image and natural language processing as well as object and speech recognition based on human brain mimicking [49, 110]. Indeed, DL is a deep neural network that can be applied for supervised, e.g., classification and regression tasks, and unsupervised, e.g., clustering, learning. In this work, since we have two different data types as input, two various networks are needed for data processing. The numerical data is fed into fully-connected layers while image features are extracted through the convolutional layers. For images involving a large number of pixel values, it is often not feasible to directly utilize all the pixel values for fully-connected layers because it can cause overfitting, increased
complexity, and difficulty in model convergence. Hence, convolutional layers are applied to reduce the dimensionality of the image data by finding the image features [73, 111].

**Fully-Connected Layers**

Fully-connected layers are hidden layers consist of hidden neurons and activation function [112]. The number of hidden neurons is usually selected based on trial and error. The neural networks can predict complex nonlinear behaviors of systems through activation functions. Any nonlinear function that is differentiable can be used as an activation function. However, there are some activation functions such as rectified linear (ReLU), leaky rectified linear, hyperbolic tangent (Tanh), sigmoid, Swish, and softmax that have been successfully used in different applications in neural networks [113]. In particular, ReLU ($f(x) = \max (0, x)$) and Swish ($f(x) = x \ \text{sigmoid}(x)$) activation functions have been recommended for hidden layers in deep neural networks [114].

**Convolutional Neural Networks (CNN)**

A convolutional neural network (CNN) is a deep network that is applied for image processing and computer vision tasks. For the first time, LeCun et al. proposed using CNN for image recognition [115]. CNN, like other deep neural networks, consists of input, output, and hidden layers. But the main difference lies in the use of hidden layers consisting of convolutional, pooling, and fully-connected layers that follow each other. Several convolutional and pooling layers can be designed in the CNN architectures.

Convolutional layers can extract the salient features of images without losing the information. At the same time, the dimensionality of the generated data gets reduced and
then fed as input to the fully-connected layer. Two significant advantages of CNN are parameter sharing and sparsity of the connections. A schematic diagram for CNN is given in Figure 2.1. The convolutional layer consists of filters that pass over the image and scanning the pixel values to make a feature map. The produced map proceeds through the activation function to add nonlinearity property. The pooling layer involves a pooling operation, e.g., maximum or average, which acts as a filter on the feature map. The pooling layer reduces the size of the feature map by pooling operation. Different combinations of convolutional and pooling layers are usually used in various CNN architectures. Finally, the fully-connected layers are added to train on image extracted features for a particular task such as classification or regression.

![Figure 2.1 Schematic of a typical convolutional neural network.](image)

Similar to other neural networks, a cost function is used to train a CNN and update the weights and biases by backpropagation. There are many hyperparameters such as the number of filters, size of filters, regularization values, dropout values, optimizer parameters, initial weights, and biases that must be initialized before training. Training a
CNN usually needs an extensive training dataset that is not always available for all applications. In this situation, transfer learning can be helpful in developing a CNN. In transfer learning, all or part of a pretrained network like VGG16, VGG19 [64], Xception [65], ResNet [66], and Inception [67], which were trained by computer vision research community with lots of open source image datasets such as ImageNet, MS, CoCo, and Pascal, can be used for the desired application. The state-of-the-art pretrained network is EfficientNet which was proposed by Tan and Le [116]. This method is based on the idea that scaling up the CNN can increase its accuracy [117]. Since there was no complete understanding of the effect of network enlargement on the accuracy, Tan and Le proposed a systematic approach for scaling up the CNNs. There are different ways to scale up the CNNs by their depth [117], width [118], and resolution [119]. Tan and Le proposed to scale up all the depth, width, and resolution factors for the CNN with fixed scaling coefficients. [116]. The results demonstrated that their proposed network, EfficientNet-B7, had better accuracy than the best-existing networks while using 8.4 times fewer parameters and performing 6.1 times faster. In addition, they provided other EfficientNet-B0 to -B6, which can overcome the models with the corresponding scale such as ResNet-152 [117] and AmoebaNet-C [120] in terms of accuracy with much fewer parameters. Due to the outstanding performance of EfficientNet, although it is trained based on the ImageNet dataset which is completely different from materials microstructures, it seems the EfficientNets convolutional layers have the potential to extract the features of images from other sources like materials microstructures.
Proposed Model for Steady State Case Study

The training and test datasets are produced using the PF method. In this work, two different algorithms, including CNN and transfer learning, were proposed to extract the salient features of the microstructure morphologies. We applied a proposed CNN (Figure 2.2) or part of pretrained EfficientNet B-6 and B-7 convolutional layers (Figure 2.3) to find the features of the microstructures. The architecture of the proposed CNN was found by testing different combinations of convolutional layers and their parameters based on the best accuracy. In the transfer learning part, different layers of the pretrained convolutional layers were tested to find the best convolutional layers for feature extraction.

On the other hand, the minimum and maximum Fe composition in the microstructure, as numerical data, is fed into the fully-connected layers. The extracted features from microstructures and the output of the fully-connected layers are combined to feed other fully-connected layers to predict the processing temperature and initial Cr and Co compositions. Different hyperparameters such as network architecture, cost function, and optimizer are tested to find the model with the highest accuracy. The model specifications, compilations (here loss function, optimizer, and metrics), and cross-validation parameters are listed in Table 2.4.
Figure 2.2 The flowchart of the developed model for chemistry and processing history prediction from microstructure images (FC: fully-connected layer)
Table 2.4  Parameters selected for model specification, compilation, and cross-validation.

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Selected value or option</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Model Specification</strong></td>
<td></td>
</tr>
<tr>
<td>Learning Rate</td>
<td>1.00E-0.3</td>
</tr>
<tr>
<td>Body activation</td>
<td>Swish, ReLU</td>
</tr>
<tr>
<td>Output activation</td>
<td>Linear</td>
</tr>
<tr>
<td>Input dimension</td>
<td>(224, 224, 1)</td>
</tr>
<tr>
<td>Output dimension</td>
<td>(3)</td>
</tr>
<tr>
<td><strong>Compilation</strong></td>
<td></td>
</tr>
<tr>
<td>Loss</td>
<td>Mean absolute percentage error</td>
</tr>
<tr>
<td>Optimizer</td>
<td>Adam</td>
</tr>
<tr>
<td>Metric</td>
<td>Root Mean square error (RMSE), R squared</td>
</tr>
<tr>
<td><strong>Cross-Validation</strong></td>
<td></td>
</tr>
<tr>
<td>Fold</td>
<td>5</td>
</tr>
<tr>
<td>Training data</td>
<td>80%</td>
</tr>
<tr>
<td>Testing data</td>
<td>20%</td>
</tr>
<tr>
<td>Batch size</td>
<td>8</td>
</tr>
<tr>
<td>Epochs</td>
<td>750</td>
</tr>
</tbody>
</table>

![Figure 2.3](image)

**Figure 2.3  The flowchart of the developed model for chemistry and processing history prediction from microstructure images (FC: fully-connected layer)**
**Proposed Model for Unsteady State Case Study**

In this study, different in-house CNNs or different layers of pretrained convolutional layers of EfficientNet have been adopted to extract microstructure features. The proposed deep network in the framework includes different in-house CNNs or pretrained convolutional layers from EfficientNet-B7 (transfer learning) for microstructure feature extraction and fully-connected layers for processing of the extracted features and numeric data (Iron minimum and maximum composition in the micrographs). CNNs with different convolutional layers are applied for microstructure feature extraction in the in-house CNNs. In transfer learning, different layers of pretrained convolutional networks are tested to find the optimum number of layers based on the overall accuracy. The architecture of the proposed network is found by testing different combinations of convolutional, fully-connected layers and their parameters based on the best accuracy. A schematic flowchart of the proposed framework is given in Figure 2.4. The extracted features of microstructures are passed through fully-connected layers to get combined with the output of the fully-connected layers that proceed the numeric data. The network is trained by the end-to-end method to find the optimum hyperparameters. The model parameters and specifications are the same as in Table 2.4, and only the output dimension will be 4.
Proposed Model for Microstructure Evolution Prediction

Prediction of microstructure evolution is a spatiotemporal problem. Different network architectures, which can generally be grouped into three categories: feed-forward models based on CNNs, recurrent models, and others such as the combinations of convolution and recurrent networks, as well as the Transformer-based and flow-based methods, are used to encode different inductive biases into neural networks for spatiotemporal predictive learning [121]. The inductive bias of group invariance over space has been brought to spatiotemporal predictive learning through the use of convolutional layers. For next frame prediction in Atari games, Oh et al. [122] defined an action-conditioned autoencoder with convolutions. The Cross Convolutional Network, developed by Xue et al. [123], is a probabilistic model that stores motion data as convolutional kernels and learns to predict a likely set of future frames by understanding their conditional distribution. In order to complete the crowd flow prediction challenge, Zhang et al. [124] suggested using CNNs with residual connections. It specifically takes into account the proximity, duration, trend, and external elements that affect how population flows move. Additionally, the convolutional architectures are employed in tandem with the generative
adversarial networks (GANs) [125], which successfully lowered the learning process’ uncertainty and enhanced the sharpness of the generated frames. Most feed-forward models demonstrate greater parallel computing efficiency on large-scale GPUs compared to recurrent models [126-128]. However, these models generally fail to represent long-term reliance across distant frames since they learn complex state transition functions as combinations of simpler ones by stacking convolutional layers.

Some helpful insights into how to forecast upcoming visual sequences based on historical observations are provided by recent developments in RNNs. In order to forecast future frames in a discrete space of patch clusters, Ranzato et al. [129] built an RNN architecture that was influenced by language modeling. As a remedy for video prediction, Srivastava et al. [130] used a sequence-to-sequence LSTM model from neural machine translation [131]. Later, other approaches to describe temporal uncertainty or the multimodal distribution of future frames conditioned on historical observations have been presented, by integrating variational inference with 2D recurrence [132-135]. By arranging 2D recurrent states in hierarchical designs, certain additional techniques successfully increased the forecast time horizon [136]. The factorization of video information and motion is another area of research, typically using sequence-level characteristics and temporally updated RNN states [137]. The use of optical flows, new adversarial training schemes, relational reasoning between object-centric content and pose vectors, differentiable clustering techniques, amortized inference enlightened by unsupervised image decomposition, and new types of recurrent units constrained by partial differential equations are typical approaches [138-143]. The aforementioned techniques work well for breaking down dynamic visual scenes or understanding the
conditional distribution of upcoming frames. To describe the spatiotemporal dynamics in low-dimensional space, they primarily use 2D recurrent networks, which inadvertently results in the loss of visual information in actual circumstances.

Shi et al. [144] created the Convolutional LSTM (ConvLSTM), which substitutes convolutions for matrix multiplication in the recurrent transitions of the original LSTM to combine the benefits of convolutional and recurrent architectures. An action-conditioned ConvLSTM network was created by Finn et al. [145] for visual planning and control. Shi et al. [146] coupled convolutions with GRUs and used non-local neural connections to expand the receptive fields of state-to-state transitions. Wang et al. [147] introduced a higher-order convolutional RNN that uses 3D convolutions and temporal self-attention to describe the dynamics and includes a time dimension in each hidden state. Su et al. [148] increased the low-rank tensor factorization-based higher-order ConvLSTMs' computational effectiveness. Convolutional recurrence provides a platform for further research by simultaneously modeling visual appearances and temporal dynamics [149-154]. The spatiotemporal memory flow, a novel convolutional recurrent unit with a pair of decoupled memory cells, and a new training method for sequence-to-sequence predictive learning are all used to enhance the existing architectures for action-free and action-conditioned video prediction in Predictive Recurrent Neural Network (PredRNN) [93].

A network component known as a memory cell is crucial in helping stacked LSTMs solve the vanishing gradient issue seen by RNNs. It can latch the gradients of hidden states inside each LSTM unit during training, preserving important information about the underlying temporal dynamics, according to strong theoretical and empirical evidence.
However, the spatiotemporal predictive learning task necessitates a distinct focus on the learned representations in many areas from other tasks of sequential data; therefore, the state transition pathway of LSTM memory cells may not be optimum. First, rather than capturing spatial deformations of visual appearance, most predictive networks for language or speech modeling concentrate on capturing the long-term, non-Markovian features of sequential data [155, 156]. However, both space-time data structures are essential and must be carefully considered in order to forecast future frames. Second, low-level features are less significant to outputs in other supervised tasks using video data, such as action recognition, where high-level semantical features may be informative enough. The stacked LSTMs don't have to maintain fine-grained representations from the bottom up because there are no complex structures of supervision signals. Although the current inner-layer memory transition-based recurrent architecture can be sufficient to capture temporal variations at each level of the network, it might not be the best option for predictive learning, where low-level specifics and high-level semantics of spatiotemporal data are both significant to generating future frames. Wang et al. [93] proposed a new memory prediction framework called PredRNN, which extends the inner-layer transition function of memory states in LSTMs to spatiotemporal memory flow. This framework aims to jointly model the spatial correlations and temporal dynamics at different levels of RNNs. All PredRNN nodes are traversed by the spatiotemporal memory flow in a zigzag pattern of bi-directional hierarchies: A newly created memory cell is used to deliver low-level information from the input to the output at each timestep, and at the top layer, the spatiotemporal memory flow transports the high-level memory state to the bottom layer at the following timestep. The Spatiotemporal LSTM (ST-
LSTM), in which the proposed spatiotemporal memory flow interacts with the original, unidirectional memory state of LSTMs, was therefore established as the fundamental building element of PredRNN. It seems that they would require a unified memory mechanism to handle both short-term deformations of spatial details and long-term dynamics if they anticipated a vivid imagination of numerous future images: On the one hand, the network may learn complex transition functions within brief neighborhoods of subsequent frames thanks to the new spatiotemporal memory cell architecture, which also increases the depth of nonlinear neurons across time-adjacent RNN states. Thus, it considerably raises ST-modeling LSTM's capacity for short-term dynamics. To achieve both long-term coherence of concealed states and their fast reaction to short-term dynamics, ST-LSTM, on the other hand, still uses the temporal memory cell of LSTMs and closely combines it with the suggested spatiotemporal memory cell. Schematics of the PredRNN architecture and the ST-LSTM unit with twisted memory states are given in Figure 2.5.

On five datasets—the Moving MNIST dataset, the KTH action dataset, a radar echo dataset for precipitation forecasting, the Traffic4Cast dataset of high-resolution traffic flows, and the action-conditioned BAIR dataset with robot-object interactions—the proposed methodology demonstrated state-of-the-art performance. The original paper [93] contains information about the investigation in detail. This dissertation adopts the PredRNN to predict the microstructure evolution in a split second.
Figure 2.5  Left: the main architecture of PredRNN, in which the orange arrows denote the state transition paths of $M_l$, namely the spatiotemporal memory flow. Right: the ST-LSTM unit with twisted memory states serves as the building block of the proposed PredRNN, where the orange circles denote the unique structures compared with ConvLSTM (the figure was adopted from the original study [157]).

Measure the Similarity Between Images

Perhaps the most fundamental process underpinning all of computing is the capability to compare data elements. It is not particularly challenging in many fields of computer science; for example, binary patterns may be compared using the Hamming distance, text files can be compared using the edit distance, vectors can be compared using the Euclidean distance, etc. Even the seemingly straightforward operation of comparing visual patterns is still an open problem, which makes computer vision a particularly difficult field to study. Visual patterns are not just exceedingly high-dimensional and strongly correlated, but the idea of visual similarity itself is frequently arbitrary and intended to emulate human visual perception [158]. In order to compare the various outcomes of the experiments when working on computer vision tasks, we must select a method for measuring the similarity between two images. Objective quality or distortion assessment techniques can be divided into two main categories. The first category includes metrics that may be expressed quantitatively, such as the frequently used mean
square error (MSE), peak signal to noise ratio (PSNR), root mean square error (RMSE),
mean absolute error (MAE), and signal-to-noise ratio (SNR). In an effort to include
measurements of perceptual quality, the second class of measurement techniques takes
into account the properties of the human visual system (HVS) [159].

The mean-square error estimator is the most common. The average squared difference
between the anticipated values (estimated values) and the actual value is measured by
MSE (ground truth). Therefore, we just square the differences between each pixel.
However, this only works well if we want to create a picture with the best pixel colors
consistent with the real-world image. We occasionally like to focus on the picture's
structure or relief [158].

The second conventional estimator is PSNR (Peak Signal to Noise Ratio). All pixel
representation values must be converted to bit form to utilize this estimator. The values of
the pixel channels must range from 0 to 255 if we are using 8-bit pixels. By the way, the
RGB color model, sometimes known as red, green, and blue, suits the PSNR the best.
The PSNR metric displays the relationship between a signal's maximum achievable
power and the power of corrupting noise that compromises the accuracy of its
representation [159].

However, PSNR, a variant of MSE, continues to focus on the pixel-by-pixel
comparison. Another technique for image similarity quantification is the structural
similarity approach (SSIM). SSIM and the effectiveness and perception of the human
visual system are connected (HVS color model). The SSIM represents picture distortion
as a combination of three elements, namely loss of correlation, luminance distortion, and
contrast distortion, as opposed to utilizing conventional error summation techniques [159].

Convolutional neural networks’ hidden variables have recently been demonstrated to be an effective measure of perceptual similarity that accurately predicts human perception of relative picture similarity. The perceptual similarity between two images is assessed using the Learned Perceptual Image Patch Similarity (LPIPS). In essence, LPIPS determines how comparable two picture patches’ activations are for a given network. This measurement has been demonstrated to reflect human perception closely. Image patches with a low LPIPS score are perceptually similar [158].

In material science, there are other methods for microstructure assessment, including two-point correlation function, chord length distribution, etc. In addition to index values such as MSE, PSNR, SSIM, and LPIPS, two-point correlation function [160] and chord length [161] are used for distribution comparison between two images in this dissertation.

In two-point correlation, the local state and local state space can be used to digitize the microstructure images [162]. Local space (h) is the attributes that are needed to completely identify all relevant material properties for the selected length scale and can be defined as follows:

\[ h = (\rho, c_i) \]  

Where \( \rho \) is a phase identifier (\( \alpha, \beta, \gamma, \ldots \)) and \( c_i \) represents chemical composition. The complete set of all theoretically possible local states in a selected material system is the local state space (\( H \)) [163].
\[ H = \{(\rho, c_i) | \rho \in \{\alpha, \beta, \gamma, \ldots\}, c_i \in C_i^f\} \] (16)

Representation of the microstructure as a function \( h(x, t) \) specifies the local state present at every spatial position \( x \) and time \( t \). In practice, all microstructure characterization techniques probe the local state in the materials over a finite volume and a finite time interval. It is impractical to implement this function in practice due to the resolution limits and uncertainty inherent to the characterization techniques used. In addition, the local state can only be characterized as an average measure over a finite probe volume and finite time step. The problem raises is the fact that the local state in any particular pixel or voxel at any particular time step may not be unique [164]. To solve the mentioned issues, microstructure function \( m(h, x) \) is defined as the probability density associated with finding local state \( h \) at the spatial location \( x \) at time \( t \). It captures the probability of finding one of the local states that lie within a small interval \( dh \) centered around \( h \) at a selected \( x \). \( m(h, x) \) \( dh \) \( dx \) would represent the probability and \( m(h, x) \) the corresponding probability density [165]. The desired information for the evaluation of \( m(h, x) \) is usually discrete values.

The microstructure image on a square lattice can be represented by pixel in two-dimensional (2D) images and voxel in three-dimensional (3D) images. In this case, the microstructure images are expressed by arrays that each element of the array has a value based on that pixel or voxel brightness. Then, enough sampling grid is needed to capture rich attributes from the material internal structure. The different phases in the material microstructure can be represented by special values. For example, Figure 2.6a shows a real two-phase microstructure that can be depicted by a binary image (black and white).
Figure 2.6 (a) A real two-phase microstructure, (b) and (c) a simple checkerboard microstructure for presenting $X_{uv}$ and two-point correlation (white color is phase 1 and black color is phase 2)

If $X$ indicates the microstructure on a square lattice, it can be displayed mathematically as follows:

$$X_{uv} = \begin{cases} 1, & \text{if } uv \in \text{phase 1} \\ 0, & \text{Otherwise} \end{cases}$$  \hspace{1cm} (17)$$

$uv$ is the pixel index and represents the pixel location in the microstructure image.

In the two-point correlation function, as a simple n-point correlation method, the correlation between two random points in the microstructure that can be specified by vector $r$ are evaluated as follows:

$$f^{np}_{r,uv} = \langle X^p_{uv}, X^p_{uv+r} \rangle$$  \hspace{1cm} (18)$$

Where $\langle \cdot \rangle$ is the expectation operator. A simple example for $X_{uv}$ and two-point correlation has been presented in Figures 1b and 1c. Since we are dealing with discrete values, the expectation can be defined as:
\[ f_{r, uv}^{np} = \langle X_{uv}^n, X_{uv+r}^p \rangle = \sum_{uv} X_{uv}^n X_{uv+r}^p \] (19)

\[ f_{r, uv}^{np} \] is the conditional probability of finding local state \( n \) at bin \( uv \) given finding local state \( p \) at bin \( uv+r \). This definition can be extended for three, four, or \( n \)-point correlation function. If there is a periodic microstructure, \( f_{r, uv}^{np} \) is independent from \( uv \).

For a two-phase material, there are \( f_r^{11}, f_r^{12}, f_r^{21}, \) and \( f_r^{22} \).

\[ f_r^{np} = \begin{bmatrix} f_r^{11} & f_r^{12} \\ f_r^{21} & f_r^{22} \end{bmatrix} \] (20)

The lineal-path function is an additional statistical function that can help for microstructure characterization. The lineal-path function quantifies the clusteredness of the straight lines in the microstructure. In fact, from probabilistic point of view, the probability that a line drawn on the microstructure will be completely in one phase is calculated [166, 167]. It can be calculated by different methods like chords distribution [166] or Monte Carlo simulation [168]. The lineal-path function in a microstructure is linearly independent unlike two-point correlation which is more effective for phases recognition. The second derivative of lineal-path function is chord-length distribution (CLD) which is also used for microstructure quantification [169]. The lineal-path function cannot show the connectivity of the phases accurately because just linear connections are considered in this method. In addition, these linear connections are measured in the certain directions. Some studies tried to apply different methods to evaluate it in the multiple directions [170-172]. Despite these weaknesses, the lineal-path function has been applied for microstructure characterization in different studies [166,
The CLD function which can be derived from lineal-path function was used by Popova et al. [174] to quantify the material structure in additive manufacturing. Some researchers have reported that the lineal-path function and the two-point cluster correlation function is useful for finding clusters in the microstructures [166, 173, 175].
CHAPTER THREE: DEEP LEARNING APPROACH FOR CHEMISTRY AND PROCESSING HISTORY PREDICTION FROM MATERIALS MICROSTRUCTURE

In this chapter, we develop a deep neural network to predict chemistry and processing history prediction from materials microstructure. All the microstructures in this chapter belong to a heat treatment time of 100 hr.

The result provided in this chapter is published as a research paper [176] in the Scientific Report Journal (Volume 12, 4552 (2022), https://doi.org/10.1038/s41598-022-08484-7).

Phase-Field Modeling and Dataset Generation

Different microstructures are produced by PF modeling for different chemical compositions and temperatures. The chemical compositions and temperature were designed based on the design of experiment method. Since the chemical compositions are subject to the constraint that they must sum to one, the Simplex-Lattice design as a standard mixture design was adopted to produce the samples. In this regard, the compositions start from 0.05 and increase to 0.90 at 0.05 intervals, and the temperature rises from 853 K to 963 K at 10 K increment, see Table 2.2. Therefore, 2053 different samples were simulated by the PF method, and the microstructures were constructed for different chemical compositions and temperatures. All the proposed operating conditions were simulated for the 100 hr spinodal decomposition process. Figure 3.1 depicts three sample results of the PF simulation. The MOOSE-generated data can be presented in different color formats. In most transmission electron microscopy (TEM) images in
literature, the Fe-rich and Cr-rich phases have been shown by bright and dark contrasts, respectively. We followed the same coloring for the extracted microstructures from the MOOSE. The Chigger python library in MOOSE has been used for microstructures extraction.

![Figure 3.1 Fe-Cr-Co alloys microstructure generated by the phase-field method for: a) Fe-20%, Cr-40%, Co-40% at 873K, b) Fe-20%, Cr-40%, Co-40% at 963K, c) Fe-25%, Cr-30%, Co-45% at 933K. (Composition are in atomic percent).](image)

Since decomposition does not occur in all the proposed operating conditions and chemistries, the microstructures showing the 0.05 difference in Fe composition between Cr-rich and Fe-rich phases were considered spinodally decomposed results. Hence, 454
samples in which decomposition has taken place are used to create the database. 80% of 454 samples were used for training and 20% for testing. The training was validated by 5-fold cross-validation. The Fe-based composition microstructure morphologies, as well as minimum and maximum of Fe compositions in the microstructure along with corresponding chemical compositions and temperatures, form the dataset. A sample workflow on the dataset construction is given in Figure 3.2.

![Microstructure morphology](image)

**Figure 3.2** A sample workflow of dataset construction.

### Convolutional Layers for Feature Extraction

The overreaching goal of the convolutional layers is feature extraction from the images. First, we train a proposed CNN, which includes three convolutional layers, batch normalization, max pooling, and ReLU activation function. Filters in each convolutional layer encode the salient features of images. Once the input images are fed into the network, the filters in the convolutional layers are activated to produce the response maps as an output of the filters. Some response maps of each convolutional layer in the

<table>
<thead>
<tr>
<th>Input</th>
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<th></th>
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<tbody>
<tr>
<td>Microstructure morphology</td>
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<td>Fe maximum concentration</td>
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<td>0.339211</td>
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</table>

<table>
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<th>Output</th>
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<th></th>
</tr>
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<tr>
<td>( C_{Fe} )</td>
<td>( C_{Cr} )</td>
<td>( C_{Co} )</td>
</tr>
<tr>
<td>0.33</td>
<td>0.4</td>
<td>0.4</td>
</tr>
</tbody>
</table>

...
proposed CNN are given in Figure 3.3. Then, as a comparison, the EfficientNet-B6 and EfficientNet-B7 convolutional layers were also applied to extract the salient features of produced microstructure by the PF method. The EfficientNet-B6 and EfficientNet-B7 have 43 and 66 million parameters which are less than other network parameters with similar accuracy. The trained weights and biases of the EfficientNet models on the ImageNet dataset for classification tasks are loaded for convolutional layers without top fully-connected layers. EfficientNet-B6 and EfficientNet-B7 have 668 and 815 layers, including 139 and 168 convolutional layers, respectively. The response maps for some layers are given in Figure 3.4 and Figure 3.5 for EfficientNetB7 and EfficientNetB6, respectively. They represent the locations of the encoded features by the filters on the input image.
Figure 3.3 Sample response maps in developed CNN for 2D microstructure morphology inputs. The response map of the first four filters of three convolutional layers is illustrated for three input images. The layer numbers are presented at the top of the images.

Figure 3.4 Sample response maps in EfficientNetB7 for 2D microstructure morphology inputs. The response map of the first four filters of some convolutional layers is illustrated for three input images. The layer numbers are presented at the top of the images.
Figure 3.5  Sample response maps in EfficientNetB6 with 2D microstructure inputs. The response map of the first four filters of some convolutional layers is illustrated for three input images. The layer number is presented at the top of the figure.

The response maps for both trained CNN and pretrained EfficientNet show that the first layers capture the simple features like edges, colors, and orientations, while the deeper layers extract more complicated features that are less visually interpretable, see Figure 3.4; similar observations are reported in other studies [65, 67, 177]. The filters from the first layers can extensively detect the edges; hence the microstructures are segmented by the borders of two different phases. By going into deeper layers, understanding the extracted information by the filters becomes more difficult and can only be analyzed by their effects on the accuracy of the final model. Since the pretrained EfficientNet has deeper layers, they can extract more complicated features from the microstructure morphologies. Indeed, we can use different layers for microstructure information extraction and test them to predict the processing history and find the most optimum network.
Temperature and Chemical Compositions Prediction

The mixed dataset contains microstructure morphologies as image data and the minimum and maximum of Fe composition in the microstructures as numeric data. The most common reported experimental images in literature for the spinodally decomposed microstructures are greyscale TEM images. To enable the model to predict the chemistry and processing history of the experimental microstructures, we have used the greyscale images in the network training. The proposed CNN, as well as EfficientNet-B6 and EfficientNet-B7 pretrained networks, were used for microstructures’ feature extraction. Then, the extracted features are passed through the fully-connected layers with batch normalization, Swish activation function, and dropout. The numeric data was proceeded by fully-connected layers with the ReLU activation function. The output of both layers was combined with other fully-connected layers to predict temperature and chemical compositions through the linear activation in the last fully-connected layer. After testing different fully-connected layer sizes, the best architecture was selected based on prediction accuracy and stability, which is shown in Figure 2.2 for the proposed CNN and Figure 3.6 for pretrained networks. The models were trained on XSEDE resources [178].

As a starting point, the proposed CNN network with fully-connected layers was trained to predict the processing history parameters. After testing different CNN architectures, the presented network in Figure 2.2, provided the best results that are given in Figure 3.7. The results show that the proposed network can predict the chemical compositions reasonably well, but the temperature accuracy is poor. Temperature is a key parameter in the spinodal decomposition process and developing a model with higher accuracy is required. To increase the accuracy, we need to extract more subtle features
from the morphologies. However, training a CNN with more layers requires numerous training data. A pretrained network can extract more valuable features from images and consequently can be helpful for accuracy improvement. Therefore, after fixing the architecture of fully-connected layers, different layers of EfficientNet-B6 and EfficientNet-B7 were tested to find the best layer for microstructures’ feature extraction. Herein, layers 96, 111, 142, 231, 304, 319, 362, 392, 496, 556, 631, 659, and 663 from EfficientNet-B6 and layers 25, 108, 212, 286, 346, 406, 464, 509, 613, 673, 806, and 810 from EfficientNet-B7 were selected to quantify the microstructures. The models were run, based on the given parameters in Table 2.4, for different layers. The model training was repeated five times. The average R Squares and mean square error (MSE) for cross-validation and test set are given in supplementary materials, Table 3.1 and Table 3.2, for EfficientNet-B6 and EfficientNet-B7, respectively. Indeed, the models were validated by 5-fold cross-validation during training, and the test set contains the data that the model never sees in the training process. According to the results, both trained models based on EfficientNet-B6 and EfficientNet-B7 can predict the Co composition very well and while the prediction of temperature and Cr composition is good, they are more challenging. Accordingly, the most accurate prediction belongs to the models that use up to layer 319 of the EfficientNet-B6 and layer 806 of EfficientNet-B7 for microstructures’ quantification.
In addition to cross-validation and test set accuracy, which can be used for overfitting identification, tracking the loss change in each epoch during the training process can also help in overfitting detection. Figure 3.8a depicts the loss change in each epoch for the developed model based on EfficientNet-B7, a corresponding plot for EfficientNet-B6 is available in supplementary materials, Figure 3.9a. Figure 3.8a shows that both training and validation losses reduce smoothly with the epoch increase. The insignificant gap between the train and validation losses proves that the models’ parameters converge to the optimal values without overfitting. To better understand the application of the developed models, the models were tested by a sample from the test set; the microstructure belongs to the spinodal decomposition of 20% Fe, 40% Cr, and 40% Co at 913 K after 100 hr. The model predictions for temperature and chemical compositions...
compositions are given in Figure 3.8b, for EfficientNet-B7, and Figure 3.9b, for EfficientNet-B6. The comparison between the ground truth and prediction demonstrates that the models can predict the chemistry and processing history reasonably well. To quantify the models’ predictive accuracy on all test data points, we have used the parity plots in which the models’ predictions are compared with ground truth in an $x$-$y$ coordinate system. For an ideal 100% accurate model all data points will overlap on a 45-degree line. The parity plots of the models, i.e., EfficientNet-B7 and EfficientNet-B6, for temperature, Cr composition, and Co composition along with their accuracy parameters are given in Figure 3.8c and Figure 3.9c. The results show that the models can predict the Co composition with the highest accuracy. It seems that temperature prediction is the most challenging variable for the models, but still, there is a good agreement between the models’ prediction and ground truth.
Table 3.1  R-squared and MSE of model predictions for training and testing dataset when different layers of EfficientNet-B6 are used for microstructures’ feature extraction.

<table>
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<th>Test</th>
<th>Validation</th>
<th>Test</th>
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<td>MSE</td>
<td>R squared</td>
<td>MSE</td>
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<tr>
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<td>T</td>
<td>C_cr</td>
<td>C_co</td>
<td>T</td>
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Table 3.2  R-squared and MSE of model predictions for training and testing dataset when different layers of EfficientNet-B7 are used for microstructures’ feature extraction.

| Layer | R squared Validation | | R squared Test | | MSE Validation | | MSE Test |
|-------|----------------------|---|----------------|---|----------------|---|
|       | T   | Ccr | Cco | T   | Ccr | Cco | T   | Ccr | Cco | T   | Ccr | Cco |
| 25    | 0.9407 | 0.9906 | 0.9878 | 0.9248 | 0.9112 | 0.9499 | 6E-05 | 5E-05 | 3E-04 | 7E-05 | 0.0003 | 0.001 |
| 108   | 0.9587 | 0.9944 | 0.9988 | 0.9324 | 0.9325 | 0.9956 | 4E-05 | 3E-05 | 3E-05 | 7E-05 | 0.0003 | 1E-04 |
| 212   | 0.9700 | 0.9944 | 0.9982 | 0.9488 | 0.9079 | 0.9935 | 3E-05 | 3E-05 | 4E-05 | 6E-05 | 0.0004 | 1E-04 |
| 286   | 0.9622 | 0.9749 | 0.9980 | 0.9528 | 0.9253 | 0.996 | 4E-05 | 0.0001 | 5E-05 | 5E-05 | 0.0003 | 9E-05 |
| 346   | 0.9687 | 0.9911 | 0.9878 | 0.9451 | 0.8977 | 0.9298 | 3E-05 | 5E-05 | 3E-04 | 6E-05 | 0.0005 | 0.002 |
| 406   | 0.9729 | 0.9959 | 0.998 | 0.9375 | 0.9116 | 0.9964 | 3E-05 | 2E-05 | 5E-05 | 7E-05 | 0.0004 | 9E-05 |
| 464   | 0.9663 | 0.9939 | 0.9983 | 0.9491 | 0.9248 | 0.9972 | 4E-05 | 3E-05 | 4E-05 | 5E-05 | 0.0003 | 8E-05 |
| 509   | 0.967 | 0.9959 | 0.9984 | 0.9461 | 0.9215 | 0.9952 | 4E-05 | 2E-05 | 4E-05 | 6E-05 | 0.0004 | 1E-04 |
| 613   | 0.9110 | 0.9853 | 0.9696 | 0.8782 | 0.8421 | 0.7446 | 7E-05 | 8E-05 | 7E-04 | 1E-04 | 0.0007 | 0.007 |
| 673   | 0.9625 | 0.9945 | 0.9976 | 0.9484 | 0.9248 | 0.9937 | 4E-05 | 3E-05 | 5E-05 | 6E-05 | 0.0003 | 1E-04 |
| 806   | 0.9599 | 0.9887 | 0.9978 | 0.9319 | 0.955 | 0.9962 | 4E-05 | 6E-05 | 5E-05 | 8E-05 | 0.0003 | 1E-04 |
| 810   | 0.9422 | 0.993 | 0.9976 | 0.9095 | 0.9412 | 0.9963 | 6E-05 | 4E-05 | 6E-05 | 8E-05 | 0.0002 | 1E-04 |
The results include two important points. First, while the extracted features from the shallow trained CNN can predict the compositions well, we need deep CNN to precisely predict the temperature. For this reason, the deep pretrained EfficientNet networks were used, which could predict temperature with higher accuracy. This observation indicates that the compositions are more relevant to simple extracted features of the microstructure morphology, however, more complicated extracted features are required to estimate the temperature. The physical concepts of the problem can also explain this. A small change in compositions would alter the microstructure morphology.
much more dramatically than a small change in temperature. The differences among the microstructures with different compositions and the same processing temperature are easily recognizable. For example, with a slight change in chemistry the volume fraction of the decomposed phases would vary and this information, i.e., change in the number of white and black pixels, can easily get extracted from the very first layers of the network. However, there are subtle differences between the microstructure morphologies when we slightly change the processing temperature. Therefore, much more complex features are needed to distinguish the differences among the morphologies with small processing temperature variations. Extraction of these complex features requires deeper convolutional layers. In addition, with convolutional layers increasing, the receptive field size would improve. And that ensures no important information is left out from the microstructure when making predictions. Therefore, more information is extracted from the microstructures, and it would also increase the temperature prediction accuracy. On the other hand, training a deep CNN with limited training and test dataset is not practical. To overcome this challenge, transfer learning can be helpful, and some other studies have shown that pretrained networks are effective in feature extraction in materials science-related micrographs [25, 36, 52, 72, 179-181].
Figure 3.8  

(a) Training and validation loss per each epoch,  

(b) prediction of temperature and chemical compositions for a random test dataset, and  

(c) the parity plots of temperature and chemical compositions for the testing dataset from the proposed model when first 806 layers of EfficientNetB7 are used for microstructures’ feature extraction (The size of the input images are 224 × 224 pixels)
Validation of The Proposed Model with The Experimental Data

The model accuracy against the test dataset, i.e., the data that the model has never seen in the training process, is good, but the test dataset is still from phase-field simulation. Since the ultimate goal of the developed framework is to facilitate the microstructure mediated materials design via predicting chemistry and processing history for experimental microstructures, it is valuable to test the model accuracy on the real microstructures. For this purpose, we have tested the model against an experimental TEM image for spinodal decomposition of Fe-Cr-Co with initial composition 46% Fe, 31% Cr,
and 23% Co after 100 hr heat treatment at 873 K from Okada et al. [182]. Since the Fe composition of the micrograph was not reported in Okada et al.’s paper, we selected the Fe composition by interpolating between the adjacent simulation points in our database. Figure 3.10 shows the predictions of the proposed network for an experimental TEM microstructure.

While Co composition and processing temperature prediction is very good, we see a 16% error in Cr composition prediction. We believe the error could stem from several factors. Firstly, the TEM micrograph that we used does not have the image quality of the training dataset. Secondly, the Fe composition associated with the micrograph was not reported in the original paper [182], and we used a phase-field-informed Fe composition. Thirdly, the dimension of the experimental image was larger than the simulated data, and it was cropped to be at the same size as the required input microstructure size. Despite all these limitations, the proposed model based on the first 806 convolutional layers of EfficientNetB7 predicts the chemistry and processing temperature of an experimental TEM image reasonably well. And it demonstrates that the developed model in this work is suitable for finding the process history behind the experimental microstructures.
Beyond the specific model alloy that we used in this work, the developed model can also be generalized to other materials by considering the material production processes. The developed framework can be used for other ternary alloys that are produced by spinodal decomposition. The model performance in the process history and chemistry prediction should be considered for other spinodal decomposed alloys with less or more elements. The domain adaptation methods such as unsupervised domain adaptation [183] can provide the ability to use the developed model for other spinodal decomposed alloys. In practice, the proposed model needs two experimental inputs, 1) a TEM micrograph that shows the morphology and, 2) X-ray fluorescence spectroscopy (XRF) that provides the corresponding compositions.

Conclusion

We introduced a framework based on a deep neural network to predict the chemistry and processing history from the materials’ microstructure morphologies in this chapter. As a case study, we generated the training and test dataset from phase-field modeling of the spinodal decomposition process of Fe-Cr-Co alloy. We considered a mixed input dataset by combining the image data, the produced microstructure morphologies based on Fe composition, with numeric data, the minimum and maximum of Fe composition in the microstructure. The temperature and chemical compositions were predicted as processing history. We quantified the microstructures by a proposed CNN and different convolutional layers of EfficientNet-B6 and EfficientNet-B7 pretrained networks. Then, the produced features were combined with the output of a fully-connected layer for numeric data processing by other fully-connected layers to
predict processing history. After testing different architectures, the best network was found based on the model’s accuracy. A detailed analysis of the model’s performance indicated that the model parameters were optimized based on training and validation loss reduction. The results show that while the simple extracted features from the microstructure morphology by the first convolutional layers are enough for the chemistry prediction, the temperature needs more complicated features that can be extracted by deeper layers. The model benchmark against an experimental TEM micrograph indicates the model’s well predictive accuracy for real alloy systems. We demonstrated that the pretrained convolutional layers of EfficientNet networks could be used to extract the meaningful features relevant to the compositions and temperature from the microstructure morphology. In general, the proposed models were able to predict the processing history based on the materials’ microstructure reasonably well.

**Data availability**

The raw/processed data and codes required to reproduce these findings are available at https://github.com/Amir1361/Materials_Design_by_ML_DL.
CHAPTER FOUR: PROCESSING TIME, TEMPERATURE, AND INITIAL CHEMICAL COMPOSITION PREDICTION FROM MATERIALS MICROSTRUCTURE BY DEEP NETWORK FOR MULTIPLE INPUTS AND FUSED DATA

In this chapter, we consider heat treatment time as another processing parameter for process history and chemistry prediction from materials microstructure.


Phase-Field Modeling and Dataset Generation

We ran the PF model for the different combinations of time, temperature, and chemical compositions informed by the Simplex-Lattice design. Within the ranges given in Table 2.3, 125,233 different samples were simulated by the PF method, and the microstructures were extracted for different chemical compositions, temperatures, and time. Figure 4.1 depicts the sample results of the PF simulation. MOOSE simulations of the 2D domains take approximately 120 service units (SU) per run on a 24 Core CPU. Therefore, screening the proposed range of different temperatures and chemical compositions for microstructure evolution required approximately 505k SU to complete.
Since decomposition does not occur for all proposed operating conditions and chemistries, the microstructures showing the 0.1 difference in Fe composition between Cr-rich and Fe-rich phases and at least 15 % volume fraction for each phase were considered as spinodally decomposed results. Hence, only 14,376 samples in which decomposition has taken place are used to create the database. 80% of samples were used for training and 20% for testing. The training was validated by 5-fold cross-validation. The Fe-based composition microstructure morphologies, as well as minimum and maximum of Fe compositions in the microstructure along with corresponding time, temperature, and chemical compositions, form the dataset. A sample workflow of the dataset construction is given in Figure 4.2.
Deep Network Training

First, the in-house CNNs with different convolutional layers have been tested to find the best architecture. The results are given in Table 4.1. The results indicate that the CNNs can predict the chemistry reasonably well. The accuracy of time prediction increases proportionally with the number of filters. However, the temperature accuracy is poor for all networks. According to previous study findings [185], the temperature is related to complicated microstructure features that can only be extracted by deep convolutional layers. Training such a deep network needs a very large training dataset, which is not available. Therefore, we adopted the transfer learning method to check the network accuracy. We used the EfficientNet-B7 convolutional layers to extract the salient features of the produced microstructure by the PF method as transfer learning. The EfficientNet-B7 has 66 million parameters which are less than other networks with similar accuracies, such as VGG16, etc. Similar to other studies [65, 67, 177], the first
layers capture simple features like edges, colors, and orientations. In contrast, the deeper layers extract more complicated, less visually interpretable features (see Figure 4.3). The fused data, including microstructure morphology and Fe minimum and maximum concentration in morphology, are used for network training. Different pretrained convolutional layers of EfficientNet-B7 are applied to extract microstructures salient features while numeric data is proceeded by fully-connected layers. After passing fully-connected layers, the extracted features by convolutional layers are combined with the numeric data through the fully-connected layers to predict the outputs by linear activation function in the last layer, see Figure 4.4. Convolutional layers 25, 108, 212, 286, 346, 406, 464, 509, 613, 673, 806, and 810 from EfficientNet-B7 are used to extract the microstructures features. For different convolutional layers, the model is trained based on the given parameters in Table 2.4. The model training is based on 5-fold cross-validation and dividing the dataset into training (80%) and testing (20%) datasets. The average R Squares and mean square error (MSE) for cross-validation and test set that the model never sees in the training process are given in Table 4.2. According to the results, the prediction of time and temperature is more challenging than compositions. Almost all the models can predict the compositions very well. The whole training process was repeated three times to check the models' stability. Finally, the most accurate prediction belongs to the model that uses up to layer 286 of the EfficientNet-B7 for microstructures' quantification. The error distribution of this model, which shows a normal distribution, is given in Figure 4.5.
<table>
<thead>
<tr>
<th>Filters size</th>
<th>R squared Validation</th>
<th></th>
<th>R squared Test</th>
<th></th>
<th>MSE Validation</th>
<th></th>
<th>MSE Test</th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Time</td>
<td>T</td>
<td>(C_r)</td>
<td>(C_\alpha)</td>
<td>Time</td>
<td>T</td>
<td>(C_r)</td>
<td>(C_\alpha)</td>
</tr>
<tr>
<td>(128, 64, 32)</td>
<td>0.948</td>
<td>0.660</td>
<td>0.968</td>
<td>0.998</td>
<td>0.932</td>
<td>0.651</td>
<td>0.967</td>
<td>0.998</td>
</tr>
<tr>
<td>(256, 128, 64, 32, 16)</td>
<td>0.920</td>
<td>0.703</td>
<td>0.974</td>
<td>0.998</td>
<td>0.894</td>
<td>0.678</td>
<td>0.973</td>
<td>0.998</td>
</tr>
<tr>
<td>(128, 128, 64, 64, 32, 16)</td>
<td>0.964</td>
<td>0.663</td>
<td>0.973</td>
<td>0.998</td>
<td>0.952</td>
<td>0.644</td>
<td>0.970</td>
<td>0.998</td>
</tr>
<tr>
<td>(224, 112, 56, 28, 14)</td>
<td>0.956</td>
<td>0.681</td>
<td>0.964</td>
<td>0.998</td>
<td>0.941</td>
<td>0.652</td>
<td>0.964</td>
<td>0.998</td>
</tr>
<tr>
<td>(512, 256, 256, 128, 128, 64, 64)</td>
<td>0.978</td>
<td>0.716</td>
<td>0.977</td>
<td>0.999</td>
<td>0.964</td>
<td>0.702</td>
<td>0.976</td>
<td>0.998</td>
</tr>
</tbody>
</table>
Figure 4.3 Sample response maps in EfficientNetB7 with 2D microstructure inputs. The response map of the first four filters of some convolutional layers is illustrated for four input images. The layer number is presented at the top of the figure.

Figure 4.4 The architecture of the proposed model (input image size is $224 \times 224$ pixels).
In addition to cross-validation, testing data has also been used for overfitting detection. The training and validation losses diminish smoothly with epoch, as shown in Figure 4.6a, and it is an indication that the model parameters converge to the global optimum without overfitting. A sample from the test set is given in Figure 4.6b to show the developed model performance. The presented microstructure is for 15% Fe, 25% Cr, and 60% Co (all in atomic percent) after 195000 seconds of heat treatment at 950 K. The model's predictions have good agreement with ground truth values for time, temperature, and chemical compositions. The parity plots with accuracy metrics for comparing the model prediction with ground truth for all testing data are shown in Figure 4.6c. The results show that the model can predict the chemical compositions with the highest accuracy. The prediction accuracy for time and temperature is not as good as chemical compositions. But the model can still predict them reasonably well.
Table 4.2  R-squared and MSE of model predictions for training and testing datasets when different layers of EfficientNet-B7 were used for microstructures' feature extraction.

<table>
<thead>
<tr>
<th>Layer</th>
<th>R squared</th>
<th>MSE</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Validation</td>
<td>Test</td>
</tr>
<tr>
<td></td>
<td>Time</td>
<td>T</td>
</tr>
<tr>
<td>25</td>
<td>0.958</td>
<td>0.966</td>
</tr>
<tr>
<td>108</td>
<td>0.988</td>
<td>0.960</td>
</tr>
<tr>
<td>212</td>
<td>0.982</td>
<td>0.943</td>
</tr>
<tr>
<td>286</td>
<td>0.987</td>
<td>0.987</td>
</tr>
<tr>
<td>346</td>
<td>0.972</td>
<td>0.960</td>
</tr>
<tr>
<td>406</td>
<td>0.959</td>
<td>0.936</td>
</tr>
<tr>
<td>464</td>
<td>0.957</td>
<td>0.964</td>
</tr>
<tr>
<td>509</td>
<td>0.971</td>
<td>0.965</td>
</tr>
<tr>
<td>Layer</td>
<td>Validation</td>
<td>Test</td>
</tr>
<tr>
<td>-------</td>
<td>------------</td>
<td>------</td>
</tr>
<tr>
<td></td>
<td>R squared</td>
<td></td>
</tr>
<tr>
<td></td>
<td>Time</td>
<td>T</td>
</tr>
<tr>
<td>613</td>
<td>0.940</td>
<td>0.944</td>
</tr>
<tr>
<td>673</td>
<td>0.977</td>
<td>0.984</td>
</tr>
<tr>
<td>806</td>
<td>0.967</td>
<td>0.981</td>
</tr>
<tr>
<td>810</td>
<td>0.978</td>
<td>0.980</td>
</tr>
</tbody>
</table>
The results indicate that the time and temperature prediction is more challenging than chemical compositions, which is explainable by physical concepts. According to our simulation results (see Figure 4.7) and reported studies [182, 186], a slight change in initial chemical compositions can lead to a sensible change in microstructure morphology which is even recognizable by human eyes. Therefore, it will be uncomplicated for the model to realize the chemical composition changes. However, small changes in temperature will hardly lead to noticeable changes in the microstructure morphologies.
when the time and chemical composition are fixed. Therefore, finding these differences is hard and makes temperature prediction challenging. In the case of time, there are two different conditions. The morphology changes rapidly with time at the early stages of heat treatment, but the rate of change drops dramatically after some time, i.e., when the morphology reaches some stability, and changes will be minimal over time. This insensitivity will make the identification strenuous for the model. According to Table 4.2, the models can predict the chemical composition, temperature, and time in order of maximum to least accuracy. However, as we will discuss in the next section, most errors in time and temperatures, are not actually real errors, but just other right answers.

![Figure 4.6](image)

**Figure 4.6**  a) Training and validation loss per each epoch, b) prediction of time, temperature, and chemical compositions for a random test dataset, and c) the parity plots for time, temperature, and chemical compositions for the testing dataset based on the transfer learning model when the first 286 layers of EfficientNetB7 are used for microstructures' feature extraction (The size of the input images are $224 \times 224$ pixels)
Model Performance Analysis

The R-square and RSME results for testing points show that the model can predict time, temperature, and chemical compositions well. However, the model's reliability depends on knowing the sources of the errors. Therefore, in this section, we will do a more in-depth study on some low-accuracy cases to find out the source of errors. As was mentioned earlier and according to the parity plots in Figure 4.6c, the lowest accuracy belongs to time and temperature predictions. Some worst cases in time and temperature prediction are given in Figure 4.8.

After studying some random cases, among the predictions with high errors, we concluded that two scenarios are possible for the sources of errors. One is achieving stability in the microstructure morphology after a certain time, and the second is achieving an identical microstructure from two different paths. Based on the observations
and physical concepts, the microstructure morphologies change very sluggishly with time after passing the early stages of separation and coarsening, and reach some sort of stability. As mentioned earlier, once the stability is achieved, it is hard for the model to distinguish the differences between the microstructures due to the subtle or no changes between two considerable time steps. Therefore, we hypothesize that the errors that we observe in time predictions for high heat treatment times, i.e., times above 100 hrs, are associated with morphology stability. To test this hypothesis, we compared the simulated microstructures based on the model's predictions with the microstructure given as the input, i.e., ground truth, to the model. Quantitative comparison of different images can be made either by evaluating specific metrics or by observing the distribution of defined parameters in the images. We adopted some evaluation metrics that were widely used in the computer vision community including the Root Mean Squared Error (RMSE), Peak Signal-to-Noise Ratio (PSNR) [187, 188], the Structural Similarity Index Measure (SSIM) [189], and the Learned Perceptual Image Patch Similarity (LPIPS) [158]. In these metrics, smaller RMSE and LPIPS, and higher PSNR and SSIM indicate more similarity between images. For distributions comparison between two images, two-point correlation function [160] and chord length [161] are standard techniques and we used them in this study. Figure 4.9 shows the comparison between the ground truth and simulated microstructure for the first row of Figure 4.8. We note that the simulated microstructures in Figure 4.9 are informed by the DL-predicted chemistry, temperature, and time, i.e., the prediction values in Figure 4.8. The evaluation metrics and distributions demonstrate that the two microstructures are similar, while there is about 70 hrs differences in their heat treatment times. These quantitative comparisons endorsed our hypothesis that the errors
that we observe in time predictions for high heat treatment times are associated with the morphology stability.

Figure 4.8 Some worst cases for time (first row of images) and temperature (second row of images) predictions.
Figure 4.9  Comparison of the ground truth microstructures with the simulated microstructures from model predictions for four random cases with high errors in time.
Another source of error that we observed in predictions stems from the interplay between time and temperature. For these types of errors, we hypothesize that the predicted processing conditions, while being different from the ground truth, are indeed another path to reach a similar microstructure. To test this hypothesis, we ran the PF model with the predicted chemistry and processing parameters and compared quantitatively the simulated microstructures with the ground truth microstructures in Figure 4.10. Again, the metrics and distributions show that the microstructures are very similar, and in fact, we can generate similar microstructures from two separate paths, i.e., higher time/lower temperature and lower time/higher temperature. Therefore, in these cases, the model does not predict wrong processing but just discovers a new path. Therefore, according to the model review results, the primary sources of errors, primarily in heat treatment time and temperature, root in the physical concepts behind the spinodal decomposition and are not inherently wrong predictions but just another right answer.
Figure 4.10  Comparison of the ground truth microstructures with the simulated microstructures from model predictions for four random cases with errors in time and temperature.
Validation of The Proposed Model with the Experimental Data

The main motivation of the proposed model is to enable the chemistry and processing history prediction of a micrograph. This makes the model a unique tool that enables, for the first-time, microstructure inverse design possible with no lost information, i.e., reducing the complexity of microstructure to just average grain size, etc. Since ultimately, the predicted chemistry and processing parameters are going to feed into the experiment, the model validation is crucial. In this section, we validate the model's predictability against an experimental transmission electron microscopy (TEM) image for spinodal decomposition of Fe-Cr-Co with the initial composition of 46% Fe, 31% Cr, and 23% Co after 100 hrs of heat treatment at 873 K from Okada et al. [182]. The original TEM image was larger than the model's required input size, so it was cropped to meet the 224x224 pixels size. Also, the Fe composition minimum and maximum in the micrograph were not given, and we selected these values by interpolating between the adjacent simulation points in the database. Figure 4.11 shows the predictions of the model for the experimental TEM microstructure along with the ground truth.

<table>
<thead>
<tr>
<th>Ground truth</th>
<th>Prediction</th>
</tr>
</thead>
<tbody>
<tr>
<td>Time (Sec.)</td>
<td>360000</td>
</tr>
<tr>
<td>Temperature (K)</td>
<td>873</td>
</tr>
<tr>
<td>Cr composition</td>
<td>0.31</td>
</tr>
<tr>
<td>Co composition</td>
<td>0.23</td>
</tr>
</tbody>
</table>

Figure 4.11 Prediction of processing time, temperature, and chemistry for an experimental TEM image adopted from Okada et al. [182]. The original image was cropped to be in the desired size of 224 × 224 pixels.
Comparison between the predicted and ground truth shows that the model performs very well in terms of Co composition and temperature predictions with just 0.6% and 0.9% error, respectively. The predictions show 10% and 15% errors for annealing time and Cr composition, respectively. While all computational models naturally have some errors, we identify five key sources for the uncertainties in using the model for experimental micrographs, 1) the TEM micrograph does not have the image quality of the simulation microstructures, i.e., the training data, 2) the TEM image size was larger than the model's input, and we cropped it to 224x224 pixels, 3) the Fe composition was not reported for the TEM image and we used the PF input, 4) the model was trained with synthetic data and not TEM micrographs, 5) the PF model was parameterized with CALPHAD, and some errors correlate with uncertainty in CALPHAD data. These uncertainties can be reduced if there are enough experimental images to be used in the training dataset. Despite all these shortcomings, the model's predictions for chemistry and processing history for the TEM micrograph were reasonably well.

**Conclusion**

In this chapter, we have developed a computational framework that enables the microstructure inverse design. As a model material, we studied the Fe-Cr-Co based permanent magnet alloys. The developed deep neural network is able to read a micrograph of one element distribution and predicts the chemistry and processing parameters that would lead to that micrograph. The model integrates the physics-based and data-driven modeling. The training and testing data were generated from the phase-field modeling of the spinodal decomposition process in Fe-Cr-Co alloys. The fused input
data, including the microstructure morphologies and the associated minimum and maximum Fe composition, were used to train the proposed network to predict the heat treatment time and temperature as well as the initial chemical composition, i.e., the Cr and Co. We used different CNN layers as well as different convolutional layers of EfficientNet-B7 pretrained networks to quantify the microstructure morphologies. The accuracy metrics, parity plots, and error distribution demonstrate that the model with the EfficientNet-B7 pretrained network performs well on the training data. We found that temperature is the most challenging parameter to predict and it requires deeper layers and more complicated extracted features from microstructures. The error analysis showed that some wrong predictions, in particular the ones with high errors in time and temperature predictions, are not actually wrong but just other correct answers. We identified that the errors are associated with either the microstructure morphology stability or the possibility of having one microstructure with two processing paths. Finally, we validated the model with an experimental TEM microstructure and the model was able to predict the processing history and chemistry of the TEM micrograph reasonably well. The process parameters and chemistry prediction for experimental micrographs can improve significantly if we have the right size, and high-resolution microstructures, and also add some experimental data to the training dataset.

**Data availability**

The raw/processed data and codes required to reproduce these findings are available at

https://github.com/Amir1361/time_temperature_composition_prediction
https://github.com/Amir1361/Materials_Design_by_ML_DL.
CHAPTER FIVE: SPATIOTEMPORAL PREDICTION OF MICROSTRUCTURE EVOLUTION WITH PREDICTIVE RECURRENT NEURAL NETWORK

In this chapter, we propose a Predictive Recurrent Neural Network (PredRNN) model for microstructure evolution prediction, which extends the inner-layer transition function of memory states in LSTMs to spatiotemporal memory flow.

The result provided in this chapter is submitted as a research paper in the Materials & Design Journal (October 2022).

Phase-Field Modeling for Microstructure Sequences Generation

Following the Simplex-Lattice design, the microstructure sequences are produced by PF modeling Fe-Cr-Cr spinodal decomposition for the different times, chemical compositions, and temperatures. The microstructures were retrieved for 125,233 different samples that were simulated using the PF approach within the parameters listed in Table 2.3 for various chemical compositions, temperatures, and times. The sample microstructure sequences from PF simulation results are shown in Figure 4.1. On a 24 Core CPU, a MOOSE simulation of the 2D domains uses about 120 service units (SU) every run. Therefore, it took around 505k SU to screen the suggested range of various temperatures and chemical compositions for microstructure evolution. In other words, each MOOSE simulation of a 200 nanometer 2D domains takes approximately 120 hrs per run for 100 mesh grid on a 24 Core CPU. Figure 4.1 indicates that the microstructure evolution process differs in various chemical compositions and temperatures.
The training dataset can be generated from simulated microstructures. The length of each sequence is 20 microstructures. The first 10 microstructures are for the first 30 hr of heat treatment time and are used to predict the future 10 microstructures, which belong to 50 hr to 300 hr, as output sequence. There are 20,000 sequences for training and 4,000 sequences as testing data. Three different Fe-composition-based microstructure morphology sequences are presented in Figure 5.1.

As can be seen, the dataset contains very different evolution sequences in terms of structure. In addition, since the microstructures are selected from both distinct stages of spinodal decomposition, a fast composition modulation growth stage and a slower coarsening stage, the difference between the input and output sequence is significant,
which can be easily recognized in Figure 5.1. In this case, the model has a more difficult task in predicting the output sequence.

**Microstructure Evolution Prediction by PredRNN**

20,000 sequences trained the PredRNN to predict the output microstructures. With a mini-batch of 8 sequences, we trained the models using the ADAM optimizer. After 80,000 iterations, the training process is terminated with a learning rate of $10^{-4}$. PredRNN typically employs four ST-LSTM layers to balance training effectiveness with prediction quality. We set the size of the convolutional kernels inside the ST-LSTM unit to $5 \times 5$ and the number of channels of each hidden state to 128.

As illustrated in Figure 5.2, the training loss decreases smoothly with iteration, which indicates that the model's parameters have reached their optimal value globally. In addition, we employ evaluation measures that are frequently used to determine how similar two images are. The predicted and ground truth microstructures are compared using the Mean Squared Error (MSE), the peak signal to-noise ratio (PSNR), the Structural Similarity Index Measure (SSIM), and the Learned Perceptual Image Patch Similarity (LPIPS). The distinction between these metrics is that PSNR compares image compression quality, SSIM measures the similarity of structural information within the spatial neighborhoods, and LPIPS is based on deep features and is more in line with human perceptions. MSE estimates the absolute pixel-wise errors. Smaller MSE and LPIPS, and higher PSNR and SSIM indicate more similarity between images.
After training, test sequences are used to compute MSE, LPIPS, PSNR, and SSIM; the average values for each iteration are given in Figure 5.3. The results demonstrate that all the metrics improve with iteration to reach almost stability. It proves that the model learns from the data and can train the hyperparameters.
Figure 5.3  Average MSE, PSNR, SSIM, and LPIPS for test sequences during training per each iteration.

Figure 5.4  Frame-wise results on the three randomly selected samples from the test set produced by the final PredRNN model (predictions (P) vs. ground truth (G)).
Figure 5.4 displays three randomly selected samples from the test set for a qualitative comparison. The left microstructures of the dashed line are the input frames, the right ones in the top row are the ground truth of output microstructures, and the bottom row shows the PredRNN prediction. The microstructures produced by PredRNN predict clear images, meaning it can be confident of future variations. In addition, we can see that the predicted sequence is close to the ground truth sequence.

**Trained Model Performance on The Microstructure Evolution Prediction During Time**

Model performance frames prediction during time is one of the key parameters in spatiotemporal models' evaluation [190, 191]. Basically, the prediction of earlier frames because of similarity with input sequence is easier than long-term prediction. Figure 5.5 provides the corresponding frame-wise comparisons between the final PredRNN model predictions and ground truth microstructures for test sequences. The average values of metrics show that the model can predict all the microstructures with reasonable accuracy. On the other hand, the model is more powerful in predicting the first frames than the last as MSE and LPIPS increase and PSNR and SSIM decrease from time step 1 to 10.

For qualitative comparison of long-term and short-term prediction, three randomly selected samples from the test set produced by the final PredRNN model are given in Figure 5.6. The results show that PredRNN prediction for short-term cases is more accurate than long-term prediction. These results seem reasonable because there is a stronger relationship between the first microstructure from the output sequence and the input sequence. However, in general, the predictions for long-term cases also have good
agreement with the ground truth. It proves that the PredRNN can predict the microstructure evolution reasonably well.

Figure 5.5 Frame-wise results on the test set produced by final PredRNN model
Figure 5.6 Trained PredRNN model performance on short and long-term prediction for three randomly selected samples from the test set

Trained Model Inference Performance in Future Microstructures Prediction

The time it takes to calculate the model's outputs as a function of the inputs is known as the inference speed. The model's response time is crucial in many applications, especially those requiring real-time data [192]. Since this study aims to develop a deep network to predict the microstructure evolution quickly and accurately, the model inference performance is a principal factor. Therefore, the trained model performance is compared with the simulation on a reference computer. Since MOOSE can only run with the CPU, we used the same resource for the trained model. The result for randomly
selected test data is given in Figure 5.7. While the simulation of rest microstructures takes
more than 75 hr by PF modeling, the trained model can predict the future sequence
quickly by just having earlier microstructures. The error metrics indicate that this prediction is robust and reliable compared to simulated microstructures.

<table>
<thead>
<tr>
<th>MSE</th>
<th>SSIM</th>
<th>PSNR</th>
<th>LPIPS</th>
</tr>
</thead>
<tbody>
<tr>
<td>20.7820</td>
<td>0.9988</td>
<td>35.2164</td>
<td>0.0002</td>
</tr>
</tbody>
</table>

Figure 5.7  Comparison of trained PredRNN model speed with PF simulation on a randomly selected sample from the test set

Conclusion

We introduced a framework based on a deep neural network to predict the material microstructure evolution. As a case study, we generated the training and test dataset from phase-field modeling of the spinodal decomposition process of Fe-Cr-Co alloy. We considered the microstructure morphologies evolution based on Fe composition. The future microstructure sequence was predicted by knowing the earlier sequence by PredRNN. A detailed analysis of the model's performance indicated that the model parameters were optimized based on training loss reduction and error metrics improvement. The quantitative and qualitative comparisons show that the trained PredRNN model can predict the output sequence accurately. Although the model
accuracy for the short-term prediction is better than the long-term prediction, the model still shows reliable performance in the long-term forecasting. The model inference test demonstrates that it can predict the microstructure evolution quickly and accurately. In general, the proposed models could reasonably predict the materials' microstructure evolution.

**Data availability**

The trained model parameters and dataset to reproduce these findings are available at https://doi.org/10.24435/materialscloud:es-a4.
CONCLUSION AND FUTURE WORKS

This dissertation aims to present using the deep neural network on the materials’ microstructure that plays a critical role in the properties and performance of materials. This dissertation attempts to achieve these research goals based on the following steps.

A deep neural network for chemical compositions and process history in steady-state processes was developed at the first step. While the simulation methods based on physical concepts, such as the PF method, can predict the spatiotemporal evolution of the materials’ microstructures, they are not efficient techniques for predicting processing and chemistry if a specific morphology is desired. The model alloy used in this work is Fe-Cr-Co permanent magnets. These alloys experience spinodal decomposition at temperatures around 850 – 970 K. We used the PF method to create the training and test dataset for the DL network. The PF results are extracted after the 100 hr spinodal decomposition process, and all the training data are independent of time. The mixed dataset, which includes both images, i.e., the morphology of Fe distribution, and continuous data, i.e., the Fe minimum and maximum concentration in the microstructures, are used as input data, and the spinodal temperature and initial chemical composition are utilized as the output data to train the proposed deep neural network. A CNN will quantify the produced microstructures by the PF method; then, another deep neural network will use the salient features to predict the temperature and chemical composition. The proposed convolutional layers were compared with pretrained EfficientNet convolutional layers as transfer learning in microstructure feature extraction.
We quantified the microstructures by using a suggested CNN and various convolutional layers of the pretrained EfficientNet-B6 and EfficientNet-B7 networks. Then, further fully-connected layers integrated the generated features with the output of a fully-connected layer for processing numerical data to forecast processing history. The most accurate network was discovered after evaluating various microstructures. A thorough examination of the model's performance revealed that the model's parameters were chosen to minimize loss during training and validation. The findings demonstrate that while the chemistry prediction may be made with just the basic elements that were derived from the microstructure morphology by the first convolutional layers, the temperature prediction requires more sophisticated data that deeper layers can extract. The model's comparison to an experimental TEM micrograph shows that it is highly accurate in predicting the behavior of real alloy systems. We showed that the meaningful information pertinent to the compositions and temperature could be extracted from the microstructure morphology using the pretrained convolutional layers of EfficientNet networks. Generally speaking, the proposed models were able to fairly accurately predict the processing history based on the microstructure of the materials.

As mentioned, prediction of the chemical composition and processing history from microstructure morphology can help optimize processing conditions and discover possible processing paths for a targeted microstructure. But we did not consider the process treatment time effect on the microstructures in the first step. In the second step, we proposed a deep learning framework that can predict the treatment time, temperature, and chemistry of a microstructure just by knowing the morphological distribution of one element. We again used the Fe-Cr-Co-based permanent magnet alloy as model material.
We generated a dataset by simulating the spinodal decomposition process in Fe-Cr-Co alloys using the PF method. In this case study, the time, temperature, and initial chemical compositions are used as output, i.e., processing history, while the mixed dataset of microstructure morphology, as image data, and minimum/maximum of iron concentration in the morphology as numeric data are input.

To characterize the microstructure morphologies, we used several CNN layers as well as various convolutional layers of EfficientNet-B7 pretrained networks. The model with the EfficientNet-B7 pretrained network works well on the training data, as shown by the accuracy metrics, parity plots, and error distribution. We discovered that the most difficult characteristic to predict is temperature, which calls for deeper layers and more intricately derived features from microstructures. The error analysis revealed that some incorrect predictions—particularly those with significant errors in time and temperature predictions—were simply other right responses. We identified that the inaccuracies are related to either the stability of the microstructure morphology or the potential for a single microstructure to have two processing routes. Finally, we tested the model using an experimental TEM microstructure, and the results showed that the model was reasonably accurate in predicting the chemistry and processing history of the TEM micrograph. If we have the appropriate size and high-resolution microstructures and include some experimental data in the training dataset, the process parameters and chemical prediction for experimental micrographs can be much enhanced.

Data set generation in the first two parts of the thesis was very expensive. With the aim to expand the current model to more complex alloys, the data set generation will become a bottleneck. Therefore, in the third step, we presented a deep neural network-
based framework to predict the materials microstructure evolution, which is a
spatiotemporal prediction problem. In this case study, we used PF modeling to create the
training and test datasets for the spinodal decomposition of a Fe-Cr-Co alloy. We took
into account the evolution of microstructure morphologies dependent on Fe composition.
Knowing the earlier sequence through PredRNN allowed us to anticipate the future
microstructure sequence. According to a thorough review of the model's performance, the
model parameters were improved based on training loss reduction and improved error
metrics. The trained PredRNN model is capable of properly predicting the output
sequence, as shown by the quantitative and qualitative comparisons. Although the
model's accuracy for short-term forecasting is higher than that for long-term forecasting,
it nevertheless exhibits dependable performance in the latter.

In summary, the developed models in this dissertation will be able to find the
process conditions and chemical compositions from an ideal microstructure and predict
microstructures without expensive and time-consuming simulations and experiments.
Doing so provides the materials science community with knowledge and algorithms that
can be used for new materials development with the desired properties.

Future Works

Material informatics is one of the rapidly developing fields. With the development
of more powerful models, new rooms are opened for AI use in materials science. This
work is the first step in our group toward using deep learning and data science in
materials design. In the following, we provide some information about how it is possible
to expand this study in the future.
• Expand the model to consider 3D microstructures and predict the process history and chemistry behind them. Data generation will be the first challenge in the 3D model. At the same time, training a deep network that can digest the 3D microstructures will be another interesting problem.

• Unknown parameters in microstructure modeling are another challenge in materials design. Several complex parameters, particularly for multicomponent alloys, such as interfacial energies, diffusion coefficients, and coefficients of the Onsager diffusion matrix, are usually very difficult to measure accurately, either experimentally or computationally, and therefore not available for many materials. We can hypothesize that these parameters could be predicted from a few sets of experimental microstructures with known processing history by a machine learning model that has been trained by physics-based simulations. Our team has recently succeeded in developing a model for the prediction of Onsager and gradient energy coefficients from microstructure images with machine learning.

• Knowledge of the microstructure of the materials during the manufacturing processes, such as additive manufacturing, can greatly improve the final product quality. Live microstructure prediction based on common simulation techniques is not practical because of computational costs. The PredRNN model can be improved to predict the materials’ microstructure based on chemical compositions, processing conditions, and earlier microstructures quickly and accurately.
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