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Materials Knowledge Systems in Python - A Data Science Framework for Accelerated Development of Hierarchical Materials

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Abstract

To Do

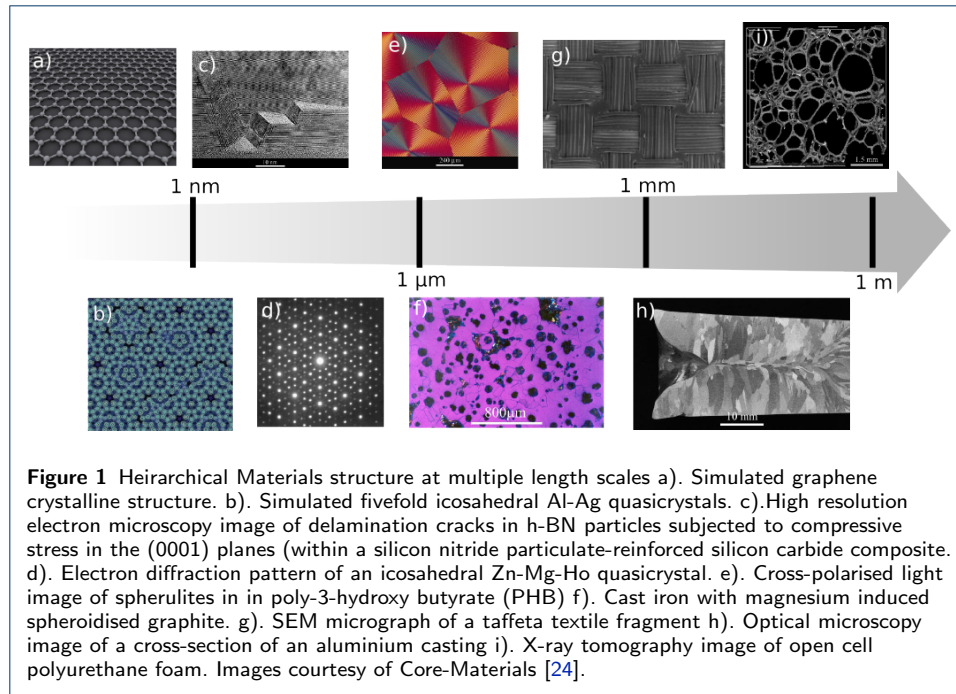
Keywords: TBD

1 Introduction

Current practices for developing tools and infrastructure used in multiscale materials design, development, and deployment are generally highly localized (sometimes even within a single organization) resulting in major inefficiencies (duplication of effort, lack of code review, not engaging the right talent for the right task, etc.). Although it is well known that the pace of discovery and innovation significantly increases with effective collaboration [1–4], scaling such efforts to large heterogeneous communities such as those engaged in materials innovation has been very difficult.

The advent of information technology has facilitated massive electronic collaborations (generally referred to as e-collaborations) that have led to significant advances in several domains including the discovery of the Higgs boson [5], the sequencing of the human genome [6], the Polymath project [7], the monitoring of species migration [8, 9] and numerous open source software projects. E-collaborations allow experts from complementary domains to create highly productive collaborations that transcend geographical, temporal, cultural, and organizational distances. E-collaborations require a supporting cyber-infrastructure that allows team members to generate, analyze, disseminate, access, and consume information at dramatically increased pace and/or quantity [10]. A key element of this emerging cyber-infrastructure is open source software as it eliminates collaboration hurdles due to software licenses and can help foster truly massive e-collaborations. In other words, even with collaborations involving proprietary data, open source cyber-infrastructure provide a common language that can facilitate e-collaborations with large numbers of team members (could even become a community effort).

Several recent national and international initiatives [11–13] have been launched with the premise that the adoption and utilization of modern data science and informatics toolsets offers a new opportunity to accelerate dramatically the design and deployment cycle of new advanced materials in commercial products. More specifically, it has been recognized that innovation cyber-ecosystems are needed to allow experts from the materials science and engineering, design and manufacturing, and data science domains to collaborate effectively. The challenge in integrating



these traditionally disconnected communities comes from the vast differences in how knowledge is captured, curated, and disseminated in these communities [14]. More specifically, knowledge systems in the materials field are rarely captured in a digital form. In order to create a modern materials innovation ecosystem, it is imperative that we design, develop, and launch novel collaboration platforms that allow automated distilling of materials knowledge from large amounts of heterogeneous data acquired through customized protocols that are necessarily diverse (elaborated next). It is also imperative that this curated materials knowledge is presented to the design and manufacturing experts in highly accessible (open) formats.

Customized materials design has great potential for impacting virtually all emerging technologies, with significant economic consequences [12, 13, 15–23]. However, materials design (including the design of a manufacturing process route) resulting in the combination of properties desired for a specific application is a highly challenging inverse problem due to the hierarchical nature of materials internal structure. Material properties are controlled by the hierarchical internal structure (over multiple length scales which spans from atomic to macroscopic) as well as coupled physical phenomena which can occur at different timescales at each of the hierarchical length scales. Characterization of the structure at each of these different length scales is often in the form of images which come from different experimental/computational techniques resulting in highly heterogeneous data. As a result, tailoring the material hierarchical structure to yield desired combinations of properties or performance characteristics is enormously difficult. Figure 1 provides a collection of materials images depicting material structures at different length scales, which are generally acquired using diverse protocols and are captured in equally diverse formats.

While the generation (from experiments and computer simulations) and dissemination of datasets consisting of heterogeneous images are necessary elements in

a modern materials innovation ecosystem, there is an equally critical need for customized analytics that take into account the stochastic nature of these data at multiple length scales in order to extract high value, transferable, knowledge. Data-driven Process-Structure-Property (PSP) linkages [25] provides a systemic, modular, and hierarchical framework for community engagement (i.e., several people making complementary or overlapping contributions to the overall curation of materials knowledge). Computationally cheap PSP linkages also communicate effectively the curated materials knowledge to design and manufacturing experts in highly accessible formats.

The Materials Knowledge Systems in Python project (PyMKS) is the first open source materials data analytics toolkit that can be used to create high value PSP linkages for hierarchical materials in large scale efforts driven and directed by an entire community of users. In this regard, it could be a foundational element of the cyber-infrastructure needed to realize a modern materials innovation ecosystem.

2 Current Materials Innovation Ecosystem

Open access materials databases and computational tools are critical components of the cyber-infrastructure needed to curate materials knowledge through effective e-collaborations [26]. Several materials science open source computational toolsets and databases have emerged in recent years to help realize the vision outlined in the Materials Genome Initiative (MGI) and the Integrated Computational Materials Engineering (ICME) paradigm [12, 13, 15–23]. Yet, the creation and adoption of a standard materials taxonomy and database schema has not been established due to the unwieldy size of material descriptors and heterogeneous data. Additionally, the coupled physical phenomena that govern material properties is too complex to model all aspects of a material simultaneously using a single computational tool. Consequently, current practices have resulted in the development of computation tools and databases with a narrow focus on specific length/structure scales, material classes, or properties.

NIST Data Gateway contains over 100 free and paid query-able web-based materials databases. These databases contain atomic structure, thermodynamics, kinetics, fundamental physical constants, x-ray spectroscopy, among other features [27]. NIST DSpace provides a curation of links to several materials community databases [28]. NIST Materials Data Curation Systems (MDCS) is a general online database that aims to facilitate the capturing, sharing, and transforming of materials data [29]. Open Quantum Materials Database (OQMD) is an open source data repository for phase diagrams and electronic ground states computed using density functional theory [30]. MatWeb is a database containing materials properties for over 100,000 materials [31]. Atomic FLOW of Materials Discovery (AFLOW) databases millions of materials and properties and hosts computational tools that can be used for atomic simulations [32]. The Materials Project (and the tool pyMatgen) [33, 34] provides open web-based access to computed information on known and predicted materials as well as analysis tools for electronic band structures. The Knowledgebase of Interatomic Models (OpenKIM) hosts open source tools for potentials for molecular simulation of materials [35]. PRedictive Integrated Structural Materials Science (PRISMS) hosts a suite of ICME tools and datastorage for

the metals community focused on microstructure evolution and mechanical properties [36].

SPPARKS Kinetic Monte Carlo Simulator (SPPARKS) is a parallel Monte Carlo code for on-lattice and off-lattice models [37]. MOOSE is a parallel computational framework for coupled systems of nonlinear equations [38]. Dream3D is a tool used for synthetic microstructures generation, image processing and mesh creation for finite element [39].

While there exists a sizable number of standard analytics tools [40–49], none of them are tailored to create PSP linkages from materials structure image data and their associated properties. PyMKS aims to seed and nurture an emergent user group in the materials data analytics for establishing homogenization and localization (PSP) linkages by leveraging open source signal processing and machine learning packages in Python. An overview of the PyMKS project accompanied with several examples is presented here. This paper is a call to others interested in participating in this open science activity.

3 Theoretical Foundations of Materials Knowledge Systems

Material properties are controlled by their internal structure and the diverse physical phenomena occurring at multiple time and length scales. Generalized composite theories [50, 51] have been developed for hierarchical materials exhibiting well separated length scales in their internal structure. Generally speaking, these theories either address homogenization (i.e., communication of effective properties associated with the structure at a given length scale to a higher length scale) or localization (i.e., spatiotemporal distribution of the imposed macroscale loading conditions to the lower length scale). Consequently, homogenization and localization are the essential building blocks in communicating the salient information in both directions between hierarchical length/structure scales in multiscale materials modeling. It is also pointed out that localization is significantly more difficult to establish, and implicitly provides a solution to homogenization.

The most sophisticated composite theory available today that explicitly accounts for the full details of the material internal structure (also simply referred as microstructure) comes from the use of perturbation theories and Green's functions [50, 52–63]. In this formalism, one usually arrives at a series expansion for both homogenization and localization, where the individual terms in the series involve convolution integrals with kernels based on Green's functions. This series expansion was refined and generalized by Adams and co-workers [62, 64, 65] through the introduction of the concept of a microstructure function, which conveniently separates each term in the series into a physics-dependent kernel (based on Green's functions) and a microstructure-dependent function (based on the formalism of n -point spatial correlations [56–61]).

Materials Knowledge Systems (MKS) [66–72] complements these sophisticated physics-based materials composite theories with a modern data science approach to create a versatile framework for extracting and curating multiscale PSP linkages. More specifically, MKS employs a discretized version of the composite theories mentioned earlier to gain major computational advantages. As a result, highly adaptable and templatable protocols have been created and used successfully to extract robust

and versatile homogenization and localization metamodells with impressive accuracy and broad applicability over large microstructure spaces.

The MKS framework is based on the notion of a microstructure function that describes many attributes and combinations of attributes in materials systems including phase identifiers, lattice orientation, chemical composition, defect types and densities, among others. The microstructure function, $m(h; s)$, represents a probability distribution for the given local state, $h \in H$, at each position, $s \in S$, in the microstructure [73–76]. The MKS framework requires a discretized description of m , which is denoted here as $m[h; s]$, where the $[\cdot; \cdot]$ represent the discretized space (in contrast to $(\cdot; \cdot)$, which defines the continuous space). In most applications, S is simply tessellated into voxels on a regular (uniform) grid such that the position can be given by $s = i, j, k$ in three dimensions. On the other hand H often needs a more sophisticated decomposition than simple linear elements.

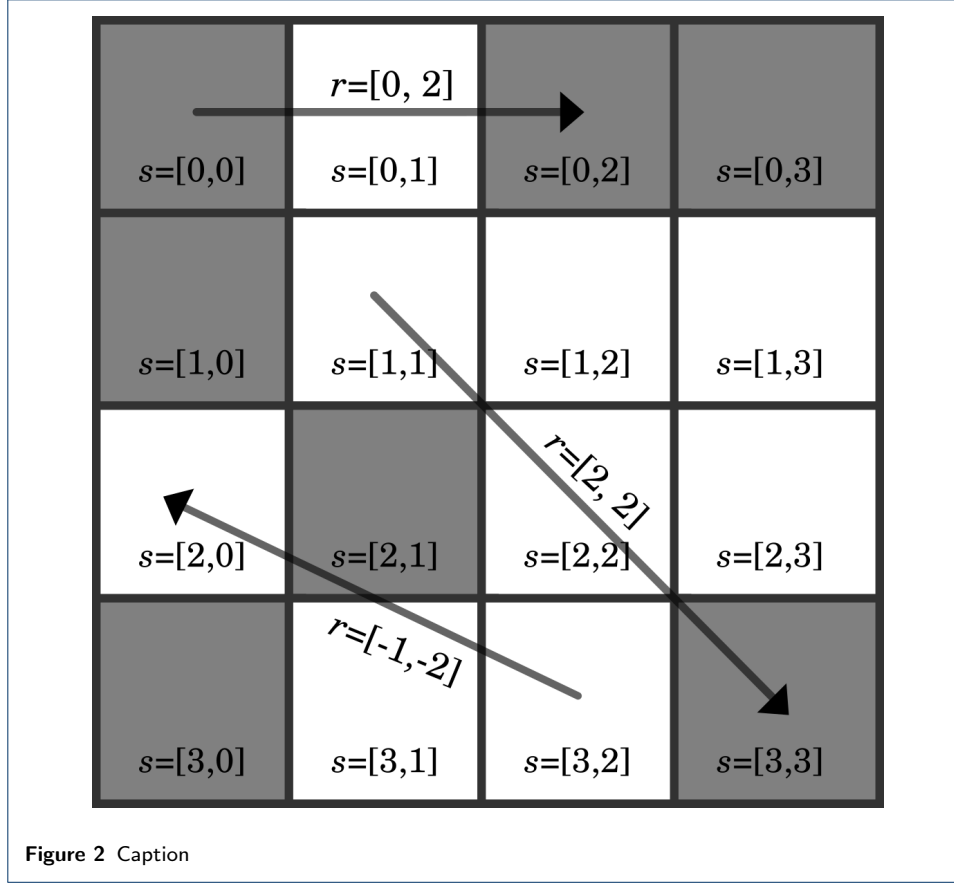
As noted earlier, the local state space in most advanced materials is likely to demand sophisticated representations. In prior work [68, 77, 78], it was found that spectral representations on functions on the local state space offered many advantages both in compact representation as well as in reducing the computational cost. In such cases, h indexes the spectral basis functions employed. The selection of these functions depends on the nature of local state descriptors. Examples include: (i) the primitive basis (or indicator functions) used to represent simple tessellation schemes [66, 67, 69–74, 79], (ii) generalized spherical harmonics used to represent functions over the orientation space [68, 77], and (iii) Legendre polynomials used to represent functions over the concentration space [78].

3.1 Homogenization

Comparing different microstructures is quite difficult even after expressing them in convenient discretized descriptions mainly due to the lack of a reference point or a natural origin for the index s in the tessellation of the microstructure volume. Yet the relative spatial distributions of the local states provide a valuable representation of the microstructure that can be used effectively to quantify the microstructure and compare it with other microstructures in robust and meaningful ways [72–74, 76, 79]. The lowest order of spatial correlations comes in the form of 2-point statistics and can be computed as a correlation of a microstructure function such that

$$f_j[h, h'; r] = \frac{1}{\Omega_j[r]} \sum_s m_j[h; s] m_j[h'; s + r] \quad (1)$$

where r is a discrete spatial vector within the voxelated domain specified by s , $f_j[h, h'; r]$ is one set of 2-point statistics for the local states h and h' and $\Omega_j[r]$ is a normalization factor that depends on r [79]. The subscript j refers to independent ensembles of sample microstructures used for analysis (each j could refer to a separate microstructure image). The physical interpretation of the 2-point statistics is explained in Fig. 2 with a highly simplified two-phase microstructure (the two phases are colored white and gray). Using the primitive basis for both the spatial domain and the local state space, $f_j[h, h'; r]$ can be interpreted as the probability of finding local states h and h' at the tail and head of the vector r , respectively.



2-Point statistics provide a meaningful representation of the microstructure, but create an extremely large feature space that often contains redundant information. Dimensionality reduction can be used to create low dimensional microstructure descriptors from the sets of spatial correlations (based on different selections of h and h') with principal component analysis (PCA). As a convenience for PCA, the $f_j[h, h'; r]$ indices are expressed as a single index given by $f_j[k]$ where k is an unraveling, such that $k = k(h, h', r)$ is a unique integer for every combination of h , h' and r . The PCA dimensionality reduction can be mathematically expressed as

$$f_j[k] \approx \sum_{k' \in K'} \mu_j[k'] \phi[k', k] + \overline{f_j[k]} \quad (2)$$

In Eq. 2, $f_j[k]$ are the reconstructed 2-point statistics. The $\mu_j[k']$ are low dimensional microstructure descriptors (the transformed 2-point statistics) or principal component scores (PC scores). The $\phi[k', k]$ are the calibrated principle components (PCs) and the $\overline{f_j[k]}$ are the mean values for each j and independent of k . The $k' \in K'$ indices refer to the $\mu_j[k']$ in decreasing order of significance and are independent of l , l' and r . The main advantage of this approach is that the $f_j[k]$ can be reconstructed to sufficient fidelity with only a small subset of the K' indices [ADDREFS].

After obtaining the needed dimensionality reduction in the representation of the material structure, machine learning models can be used to create homogenization

or localization PSP linkages of interest. As an example, a generic homogenization linkage can be expressed as

$$p_j^{\text{eff}} = \mathcal{F}(\mu_j[k']) \quad (3)$$

In Eq. 3, p_j^{eff} is the effective materials response (reflecting an effective property in structure-property linkages or an evolved low dimensional microstructure descriptor in process-structure linkages), and \mathcal{F} is a machine learning function that links $\mu_j[k]$ to p_j^{eff} .

3.2 Localization

MKS Localization linkages are significantly more complex than the homogenization linkages. These are usually expressed in the same series forms that are derived in the general composite theories, while employing discretized kernels based on Green's functions [50, 52–63]. Mathematically, the MKS localization linkages are expressed as

$$p_i[s] = \sum_{h;r} \alpha[h;r] m_i[h;s-r] + \sum_{h,h';r,r'} \alpha[h,h';r,r'] m_i[h;s-r] m_i[h';s-r'] + \dots \quad (4)$$

In Eq. 4, $p_i[s]$ is the spatially resolved (localized) response field (e.g. a response variable such as stress or strain rate, or an evolved microstructure function), and $\alpha[h;r]$ are the Green's function based discretized influence kernels. These digital kernels are calibrated using regression methods [66–69, 77, 78].

Figs. 4 and 3 provide schematic overviews of the MKS homogenization and localization workflows. THE CURRENT IMAGES ARE JUST PLACE HOLDERS. More detailed explanations on the MKS homogenization and localization linkages can be found in prior literature [66–74, 79].

4 Materials Knowledge Systems in Python

PyMKS is an object-oriented numerical implementation to the MKS theory developed in the literature [67]. It provides a high-level, computational efficient framework to implement data pipelines for classification, cataloging and quantifying materials structures for PSP relationships. PyMKS is written in Python, a natural choice for scientific computing due to its ubiquitous use among the data science community as well as many other favorable attributes [80]. PyMKS is licensed under the permissive MIT license [81] which allows for unrestricted distribution in commercial and non-commercial systems.

4.1 Core Functionality

PyMKS consists of four main components including a set of tools to compute 2-point statistics, tools to obtain low-dimensional microstructure descriptors and tools for both homogenization and localization. In addition, PyMKS has modules for generating data sets using conventional numerical simulations and a module for plotting and visualizing customized for microstructures.

Figure 3 Localization Figure To Do - Place holder

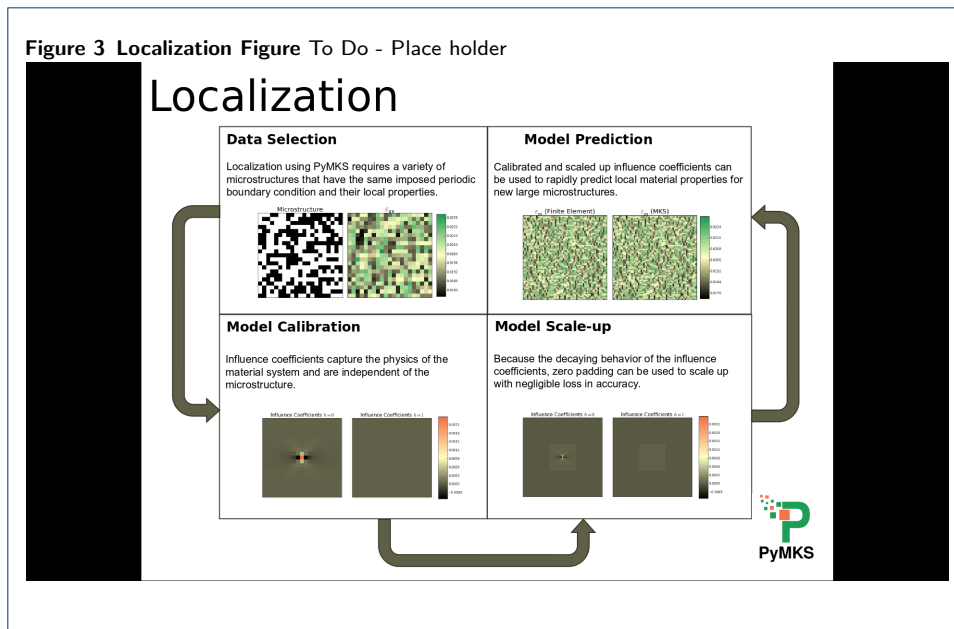
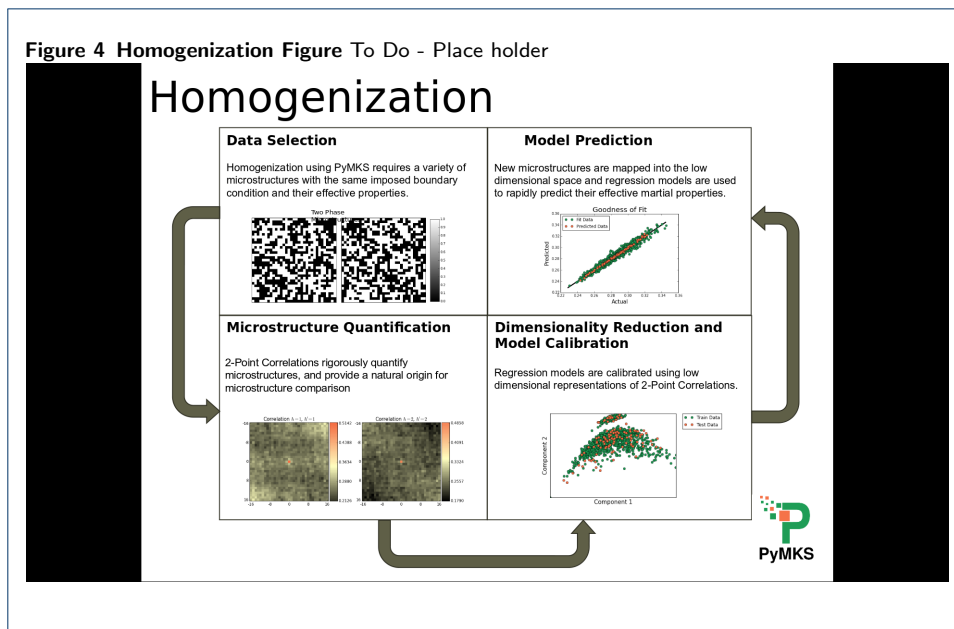


Figure 4 Homogenization Figure To Do - Place holder



The functions `autocorrelate`, `crosscorrelate`, and `correlate` provides APIs to compute the 2-point statistics for a given microstructure as outlined in Eq. 1. The `MKSStructureAnalysis` class provides access to the objective low dimensional structure descriptors, $\mu[k]$. While the default dimensionality reduction technique is PCA, any model from Scikit-learn that has a `transform_fit` method can be used. The `MKSHomogenizationModel` creates a linkage between $\mu[k]$ and a effective material response, p_{eff} as indicated in Eq. 3. The default machine learning function is a polynomial regression, but any estimator from Scikit-learn that has `fit` and `predict` methods can be used to predict effective material responses or microstructure classes. The `MKSLocalizationModel` provides the API to calibrate

the first order influence kernels $\alpha[r, l]$ in order to predict local materials responses $p[s]$ as indicated by Eq. 4. The calibration of the influence kernels is done using the multiple linear regression techniques outlined in previous studies [66–68, 78]. The localization model has `fit` and `predict` methods to follow the standard API for an estimator.

Classes in the module `bases` have methods to discretize the raw structure data and introduce the local state variable l . The four basis classes are designed to efficiently discretize different types of local state variables [66–72, 78]. The `PrimitiveBasis` class uses indicator (or hat) functions and it is well suited for microstructures that have discrete local states (e.g., distinct thermodynamic phases encountered in describing the microstructure). The `LegendreBasis` and `FourierBasis` create spectral representations of functions defined on nonperiodic and periodic continuous local state spaces, respectively. For example, functions over a range of chemical compositions can be described using `LegendreBasis`, while functions over orientations in two-dimensional space can be described using `FourierBasis`. As another option, `GSHBasis` creates compact spectral representations for functions over lattice orientation space (such as those needed to describe polycrystalline microstructures) [82–93].

PyMKS also contains modest data generation tools that are used in both examples and in unit tests, and can be found in the module `datasets`. The `MicrostructureGenerator` class creates stochastic microstructures using digital filters. A phase field and finite element simulation are implemented with the `CahnHilliardSimulation` and `ElasticFESimulation` classes. The microstructures and simulations can be accessed using the functions which have names starting with `make`. Lastly PyMKS contains some plotting tools that are used throughout the examples which can be found in the module `tools`.

4.2 Code Design

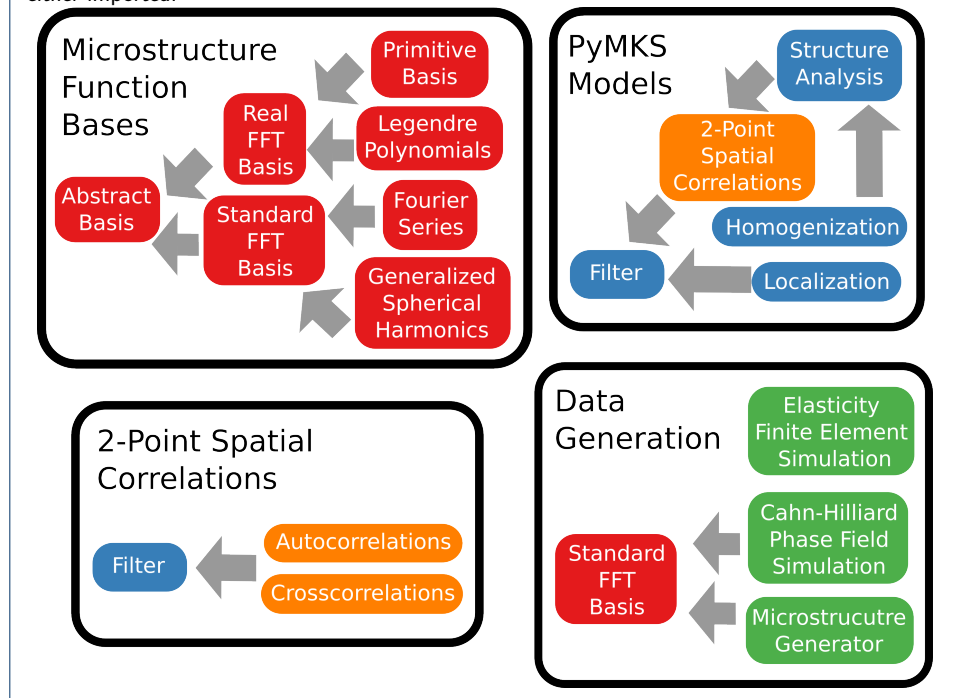
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4.3 Underlying Technologies

PyMKS is built on highly optimized Python packages NumPy [94], SciPy [95], and Scikit-learn [42]. NumPy arrays are the primary data structure used throughout PyMKS and provide basic algorithmic functionality. SciPy’s signal processing and numerical linear algebra functions are used to calibrate models and generate synthetic data. PyMKS is highly integrated with Scikit-learn and mimics its simple API in order to leverage models for dimensionality reduction, regression, classification, and model selection. In addition, PyMKS uses the Python testing framework nose for unit-tests and doc-tests.

Additional packages that can be used with PyMKS include Simple Finite Element in Python (SfePy) [96], the python wrapper for the FFTW library (pyFFTW) [97] and the plotting package Matplotlib [98]. SfePy is used for data generation to simulate the linear elastic response of composite materials, and is not necessary for analysis using external data. PyFFTW is an optional package that can be used to reduce run time of Fast Fourier Transforms (FFTs) through the optimized FFTW library and allows for the computation to be run in parallel with simple high level API arguments. If pyFFTW is not installed, NumPy’s `fft` module is used. Matplotlib is used to generate plots found throughout the examples.

Figure 5 PyMKS Design AGAIN A PLACE HOLDER - An illustration of the design for PyMKS. The four blocks represent the main functionalists in PyMKS. The modules listed on the far left are exposed in the API while the other plots are private. The arrows indicate module from that is either imported.



4.4 Development Practices

The development team for PyMKS is an open community that uses Github for pull-requests, code review, issue tracking and release management -

<https://github.com/materialsinnovation/pymks>. Additionally a Google group is used as a public forum to discuss the project development, support and announcements pymks-general@googlegroups.com.

ADD CONTENT ABOUT TRAVIS CI, etc.

PyMKS follows PEP8 standards and uses pull request to ensure code integrity, and combines overlapping functionality between classes using abstractions. Detailed administrative guidelines are outlined in the `ADMINISTRATA.md` document on Github, and potential developers are encouraged to follow them.

5 Examples of Homogenization and Localization with PyMKS

5.1 Prediction of Effective Stiffness using Homogenization

5.1.1 Data Generation

A set of periodic microstructures and their volume averaged elastic stress values $\bar{\sigma}_{xx}$ can be generated by importing the `make_elastic_stress_random` function from `pymks.datasets`. This function has several arguments. `n.samples` is the number of samples that will be generated, `size` specifies the dimensions of the microstructures, `grain_size` controls the effective microstructure feature size, `elastic_modulus` and `poissons_ratio` are used to indicate the material property for each of the phases, `macro_strain` is the value of the applied uniaxial strain, and the `seed` can be used to change the the random number generator seed.

Let's go ahead and create 6 different types of microstructures each with 200 samples with dimensions 21 x 21. Each of the 6 samples will have a different microstructure feature size. The function will return the microstructures and their associated volume averaged stress values.

```
import numpy as np
from pymks.datasets import make_elastic_stress_random

sample_size = 200
grain_size = [(15, 2), (2, 15), (7, 7), (8, 3), (3, 9), (2, 2)]
n_samples = [sample_size] * 6
elastic_modulus = (310, 200)
poissons_ratio = (0.28, 0.3)
macro_strain = 0.001
size = (21, 21)

X, y = make_elastic_stress_random(
    n_samples=n_samples, size=size, grain_size=grain_size,
    elastic_modulus=elastic_modulus, poissons_ratio=poissons_ratio,
    macro_strain=macro_strain, seed=0)
```

Lets take a look at 3 types the microstructures to get an idea of what they look like. We can do this by importing `draw_microstructures`.

```
from pymks.tools import draw_microstructures
```

```
X_examples = X[::sample_size]
draw_microstructures(X_examples[:3])
```



5.1.2 Calibration of Homogenization Model

In order to make an instance of the `MKSHomogenizationModel`, we need to pass an instance of a basis class. For this particular example, there are only 2 discrete phases, so we will use the `PrimitiveBasis` from `pymks.bases`. We only have two phases denoted by 0 and 1, therefore we have two local states and our domain is 0 to 1. Let's make an instance of the `MKSHomogenizationModel`.

```
from pymks import MKSHomogenizationModel
```

```

from pymks.bases import PrimitiveBasis

p_basis = PrimitiveBasis(n_states=2, domain=[0, 1])
model = MKSHomogenizationModel(basis=p_basis, periodic_axes=[0, 1],
                                correlations=[(0, 0), (1, 1)])

```

The default machine model used to create the homogenization linkage is polynomial regression. We need to optimize the number of principal components and the degree of polynomial. To do this we are going to split the data into test and training sets. This can be done using the `train_test_split` function from `sklearn`.

```

from sklearn.cross_validation import train_test_split

flat_shape = (X.shape[0],) + (X[0].size,)
X_train, X_test, y_train, y_test = train_test_split(
    X.reshape(flat_shape), y, test_size=0.2, random_state=3)
print(X_train.shape)
print(X_test.shape)

(960, 441)
(240, 441)

```

We will use cross validation with the testing data to find the optimal value for our model parameters using `GridSearchCV` from `sklearn`. We will pass a dictionary `params_to_tune` with the range of the degree of the polynomial and the number of principal components we want to search. Let's vary `n_components` from 1 to 11 and degree from 1 to 3.

```

from sklearn.grid_search import GridSearchCV

params_to_tune = {'degree': np.arange(1,4),
                  'n_components': np.arange(2, 12)}
fit_params = {'size': X[0].shape}
gs = GridSearchCV(model, params_to_tune,
                  fit_params=fit_params).fit(X_train, y_train)

```

The default score method for the `MKSHomogenizationModel` is R-squared.

```

print('Order of Polynomial'), (gs.best_estimator_.degree)
print('Number of Components'), (gs.best_estimator_.n_components)
print('R-squared Value'), (gs.score(X_test, y_test))

```

```

Order of Polynomial 2
Number of Components 11
R-squared Value 0.999808062591

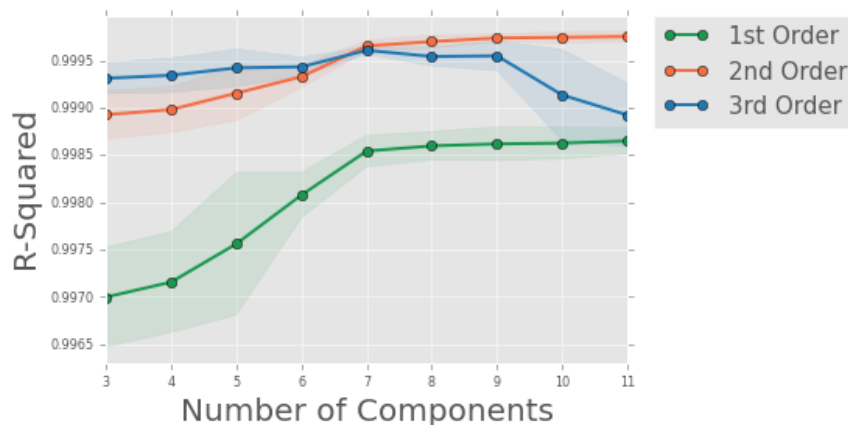
```

For the parameter range that we searched, we have found that a model with 2nd order polynomial and 11 components had the best R-squared value. Let's look at the results using `draw_grid_scores`, and set the model to have those parameter values.

```
from pymks.tools import draw_grid_scores

gs_deg_1 = [x for x in gs.grid_scores_
             if x.parameters['degree'] == 1][1:]
gs_deg_2 = [x for x in gs.grid_scores_
             if x.parameters['degree'] == 2][1:]
gs_deg_3 = [x for x in gs.grid_scores_
             if x.parameters['degree'] == 3][1:]

draw_grid_scores([gs_deg_1, gs_deg_2, gs_deg_3], 'n_components',
                 data_labels=['1st Order',
                              '2nd Order', '3rd Order'],
                 param_label='Number of Components',
                 score_label='R-Squared')
```



```
model = gs.best_estimator_
```

Now that we have found optimal values for the parameters `n_components` and `degree`, let's fit the model with the data.

```
model.fit(X, y)
```

5.1.3 Prediction of Effective Stiffness

Let's generate some more data to validate our model. We are going to generate 20 samples of all six different types of microstructures using the same `make_elastic_stress_random` function.

```
test_sample_size = 20
```

```
n_samples = [test_sample_size] * 6
X_new, y_new = make_elastic_stress_random(
    n_samples=n_samples, size=size, grain_size=grain_size,
    elastic_modulus=elastic_modulus, poissons_ratio=poissons_ratio,
    macro_strain=macro_strain, seed=1)
```

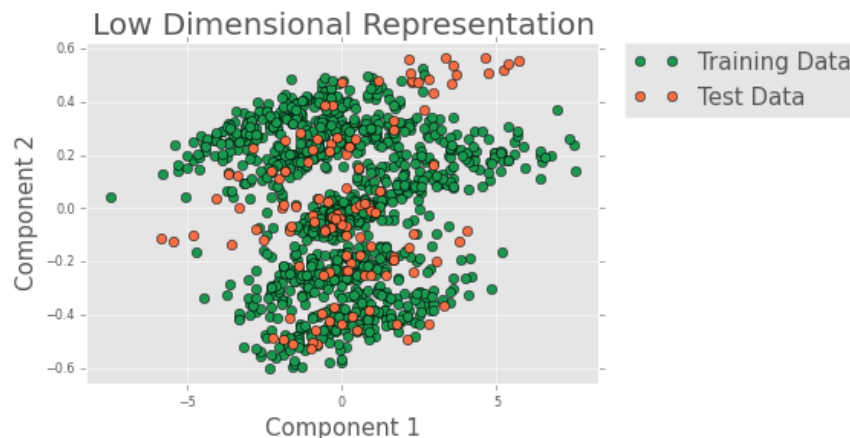
Now let's predict the stress values for the new microstructures using the `predict` method.

```
y_predict = model.predict(X_new)
```

We can look to see, if the low-dimensional representation of the new data is similar to the low-dimensional representation of the data we used to fit the model using `draw_components_scatter` from `pymks.tools`.

```
from pymks.tools import draw_components_scatter
```

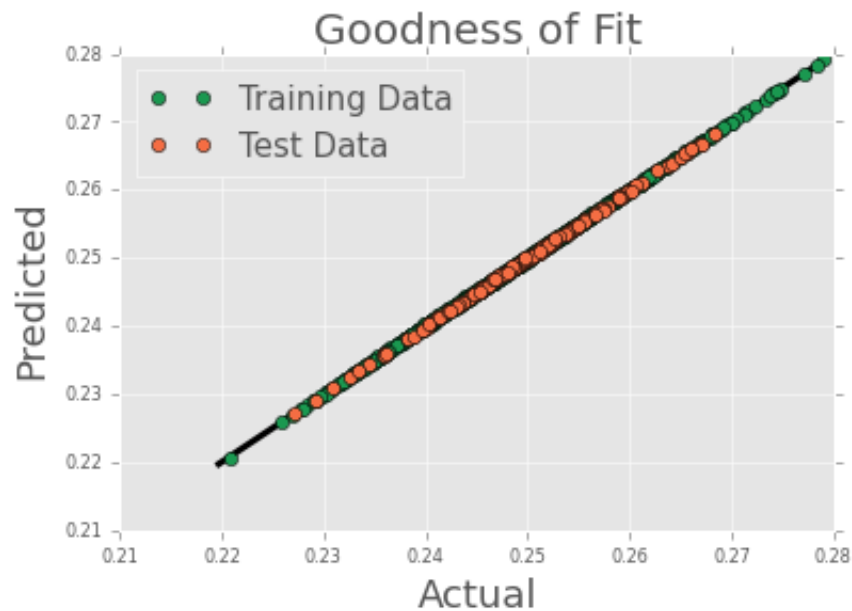
```
draw_components_scatter([model.reduced_fit_data[:, :2],
                        model.reduced_predict_data[:, :2]],
                        ['Training Data', 'Test Data'])
```



The predicted data seems to be reasonably similar to the data we used to fit the model with. Now let's look at the score value for the predicted data. We can evaluate our prediction by looking at a goodness-of-fit plot. We can do this by importing `draw_goodness_of_fit` from `pymks.tools`.

```
from pymks.tools import draw_goodness_of_fit
```

```
fit_data = np.array([y, model.predict(X)])
pred_data = np.array([y_new, y_predict])
draw_goodness_of_fit(fit_data, pred_data,
                    ['Training Data', 'Test Data'])
```



5.2 Prediction of Local Strain Field with Localization

5.2.1 Generating Calibration Data

In this example, let's look at a three phase microstructure with elastic moduli values of 80, 100 and 120 and Poisson's ratio values all equal to 0.3. Let's also set the macroscopic imposed strain equal to 0.02. All of these parameters used in the simulation must be passed into the `make_elasticFEstrain_delta` function from `pymks.datasets`. The number of Poisson's ratio values and elastic moduli values indicates the number of phases.

```
import numpy as np
from pymks.tools import draw_microstructures
from pymks.datasets import make_delta_microstructures

n = 21
n_phases = 3
from pymks.datasets import make_elastic_FE_strain_delta
from pymks.tools import draw_microstructure_strain

elastic_modulus = (80, 100, 120)
poissons_ratio = (0.3, 0.3, 0.3)
macro_strain = 0.02
size = (n, n)

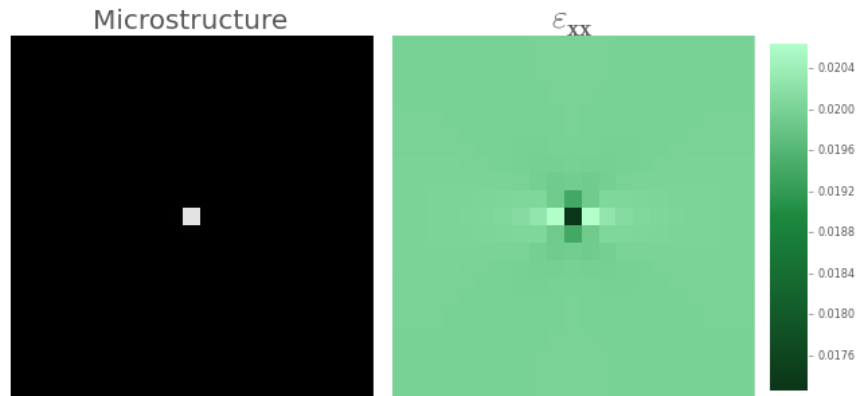
X_delta, strains_delta = make_elastic_FE_strain_delta(
    elastic_modulus=elastic_modulus,
    poissons_ratio=poissons_ratio,
```



```
size=size, macro_strain=macro_strain)
```

Let's take a look at one of the delta microstructures and the ε_{xx} strain field.

```
draw_microstructure_strain(X_delta[0], strains_delta[0])
```



5.2.2 Calibrating Localization Model

Now that we have the delta microstructures and their strain fields, we will calibrate the influence coefficients by creating an instance of the `MKSLocalizationModel` class. Because we are going to calibrate the influence coefficients with microstructures that contain discrete local states (in this case phases), we can create an instance of `PrimitiveBasis` with `n_states` equal to 3, and use it to create an instance of `MKSLocalizationModel`.

```
from pymks import MKSLocalizationModel
from pymks import PrimitiveBasis

p_basis = PrimitiveBasis(n_states=3, domain=[0, 2])
model = MKSLocalizationModel(basis=p_basis)
```

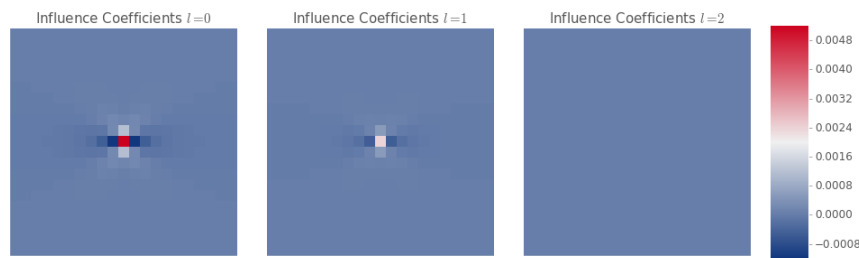
Now, pass the delta microstructures and their strain fields into the `fit` method to calibrate the first-order influence coefficients.

```
model.fit(X_delta, strains_delta)
```

That's it, the influence coefficient have been calibrated. Let's take a look at them.

```
from pymks.tools import draw_coeff

draw_coeff(model.coef_)
```



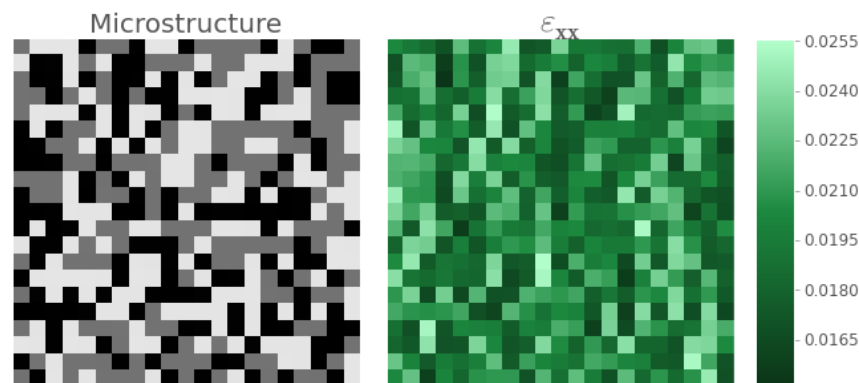
5.2.3 Prediction of the Strain Field for a Random Microstructure

Let's now use our instance of the `MKSLocalizationModel` class with calibrated influence coefficients to compute the strain field for a random two-phase microstructure and compare it with the results from a finite element simulation.

The `make_elasticFEstrain_random` function from `pymks.datasets` is an easy way to generate a random microstructure and its strain field results from finite element analysis.

```
from pymks.datasets import make_elastic_FE_strain_random

np.random.seed(101)
X, strain = make_elastic_FE_strain_random(
    n_samples=1, elastic_modulus=elastic_modulus,
    poissons_ratio=poissons_ratio, size=size,
    macro_strain=macro_strain)
draw_microstructure_strain(X[0], strain[0])
```



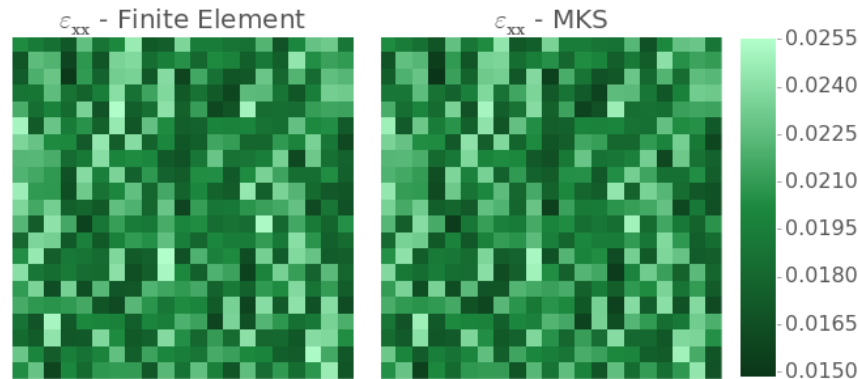
Now, to get the strain field from the `MKSLocalizationModel`, just pass the same microstructure to the `predict` method.

```
strain_pred = model.predict(X)
```

Finally let's compare the results from finite element simulation and the MKS model.

```
from pymks.tools import draw_strains_compare
```

```
draw_strains_compare(strain[0], strain_pred[0])
```



6 Conclusion

To Do

Author details

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