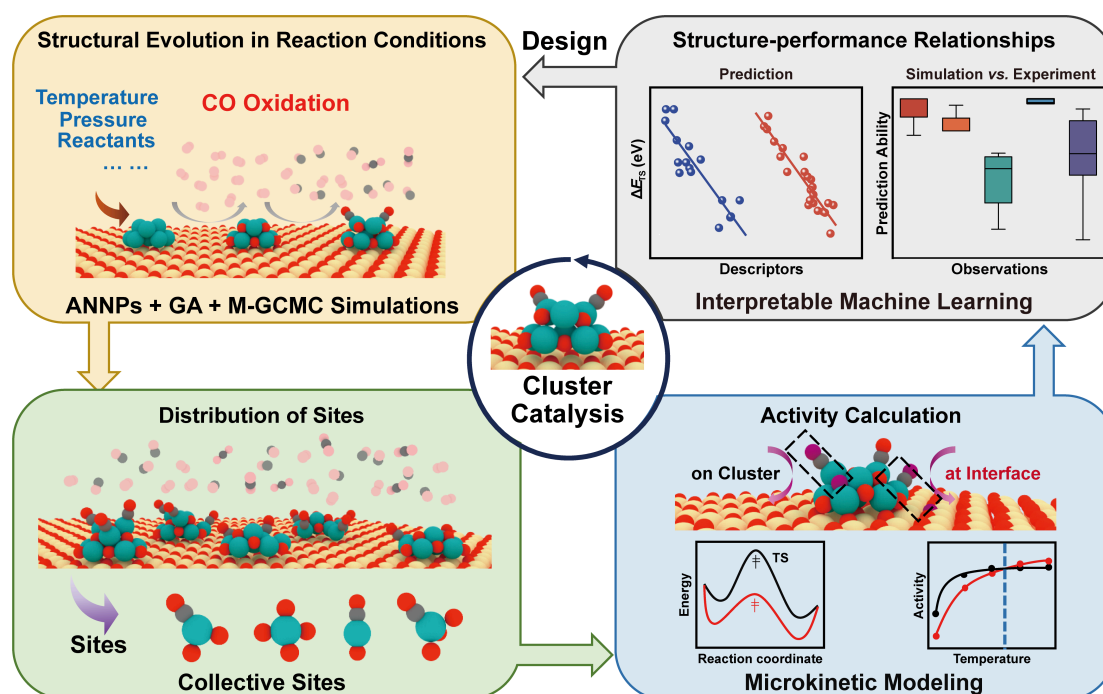


The Framework to Study Collectivity of Active Sites for Cluster Catalysis

This repository provides a computational framework to investigate the **collectivity of active sites** in cluster-based catalysis. It integrates several modules for modeling, simulation, and analysis of catalytic behavior using machine learning potentials and atomistic simulations.



Key Components

1. **Artificial Neural Network Potentials (ANNPs) Training:** For accurate energy and force predictions.
2. **Genetic Algorithm (GA):** Used for global optimization in catalyst design.
3. **Modified GCMC (M-GCMC) Simulations:** Simulates realistic catalytic environments under operational conditions.
4. **Reaction Mechanism Calculations:** For identifying transition states and elementary steps.
5. **Microkinetic Modeling (MKM):** For simulating reaction kinetics and understanding the rate-determining steps.
6. **Catalytic Descriptors:** Uses compressed sensing methods (e.g., SISSO) to uncover key descriptors.

Developers

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Methods

- **EANN (Embedded Atom Neural Network):** A simple yet efficient neural network framework for modeling atomic interactions.

- **Active learning to expand the configurations:** Iteratively expands configuration space to improve ANNP accuracy.
- **GA:** Integrated with ASE for global structure optimization.
- **M-GCMC simulations:** Modified GCMC (M-GCMC) simulations for simulating adsorption and reaction processes under *operational* conditions and search metastable structures.
- **SISSO (The sure independence screening and sparsifying operator):** A dimensionality reduction method to identify key descriptors for catalytic performance prediction.

Contents

Folder	Description
<code>nn/</code>	Scripts for training reliable ANNPs with active learning.
<code>str/</code>	Metastable structure search using GA and M-GCMC.
<code>sites/</code>	Identification and distribution analysis of active sites.
<code>ts/</code>	Transition state search using NEB and DyNEB with ANNPs.
<code>mkm/</code>	Microkinetic modeling input files.
<code>desp/</code>	SISSO input data and training sets for descriptor learning.

Module Details

✓ Training Reliable ANNPs (`nn/`)

This process includes generating the initial dataset and expanding the configuration space using an active learning algorithm. The ASE interface for the Embedded Atom Neural Network (EANN) is employed via `eann.py`.

To train the model and handle uncertain or poorly predicted structures:

```
1 python3 train_forces.py # Handle uncertain regions in the configuration space
2 python3 get_atoms # Identify structures with poor predictive accuracy
```

✓ Searching metastable structures under *operational* conditions

These simulations are used to search metastable structures under *operational* conditions.

All relevant scripts and configuration files can be found in the **str** folder.

- **Genetic algorithm (GA)**

Used to optimize catalyst structures via evolutionary principles.

```

1 python3 initial.py \
2   --poscar POSCAR_base \
3   --db gadb.db \
4   --atoms 8x10 29x8 \
5   --size 24 \
6   --zvac 2.0 # Initialize the GA search
7
8 python3 ga_test.py \
9   --db gadb.db \
10  --processes 4 # Run the GA optimization

```

- **Modified GCMC (M-GCMC) simulations**

A modified GCMC approach tailored for metastable structure database under varying chemical potentials of CO and O₂.

```

1 python3 main.py # Execute the GCMC simulation in gcmc folder
2 python3 main.py # Execute the M-GCMC simulation in m-gcmc folder

```

✓ Identification of Sites (sites/)

This step involves identifying metastable structures and analyzing the distribution of potential active sites.

The corresponding scripts and data are located in the **site** folder.

- **Metastable structures and their distribution**

Identifies metastable structures.

```

1 python3 meta_min_energy_Cu3.py \
2   --energy_files energy_meta energy_meta2 \
3   --poscar_dirs dataa111 dataa112 \
4   --temperature 400 \
5   --energy_cutoff 0.5 # metastable structures

```

- **Site distribution**

Determines the type and distribution of every site.

```

1 python3 meta_min_energy_next_sites.py \
2   --energy_files energy_meta energy_meta2 \
3   --poscar_dirs dataa111 dataa112 \
4   --temperature 400 \
5   --energy_cutoff 0.5 # every site and its distribution

```

✓ Transition State Search (ts/)

ANNPs are employed as an initial guess for transition states in catalytic reactions.

The automated scripts for transition state searches are located in the [ts](#) folder.

- **Dynamic NEB (DyNEB)**

- Executes a dynamic NEB search to refine the transition state.

```

1 python3 dyneb.py --poscar1 POSCAR1 --poscar2 POSCAR2 \
2   --pes EANN_PES_DOUBLE.pt --atomtype Cu O C \
3   --n_images 3 --traj myneb.traj --dyneb # Run DyNEB for more precise transition state
    search

```

`--poscar1`: Initial structure (POSCAR format)

`--poscar2`: Final structure (POSCAR format)

`--pes`: Path to the trained ANNs file

`--atomtype`: List of atom types in the system

`--n_images`: Number of intermediate NEB images

`--traj`: Output trajectory file

`--dyneb`: Enables the dynamic NEB mode for improved TS localization

To visualize the NEB potential energy surface (PES), run:

```

1 python3 neb_pos.py

```

This will generate a file named `barrier.png`, which shows the energy profile along the reaction coordinate.

✓ Microkinetic Modeling (`mkm/`)

Microkinetic modeling is performed by MKMCXX to simulate reaction kinetics and identify key rate-determining steps.

The relevant scripts can be found in the **mkm** folder.

✓ Descriptor Discovery (`desp/`)

The transition state energies are predicted by SISO. This helps in identifying key catalytic descriptors for further analysis.

The input files are located in the **desp** folder.

Dependencies

Package	Purpose	Version
ASE	Structure handling & GA/M-GCMC interface	3.22.1
MKMCXX	Microkinetic modeling	—
PyTorch	ANNP model training	1.8.0
NumPy	Numerical operations	1.20.3
Matplotlib	Plotting	3.4.3

Package	Purpose	Version
Pandas	Data handling & export	1.3.3
SciPy	Polynomial fitting	1.7.1

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