

MDP.jl: The Julia Library of Molecular Dynamics Potentials

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Classical Molecular Dynamics

- **MD simulations require a potential energy function to describe force field governing the interaction among atoms**
 - Determining a suitable potential is the most crucial task in molecular dynamics.
- **Empirical potentials**
 - Lennard-Jones, Tersoff, van der Waals, Coulomb potentials
 - AMBER, CHARMM potentials
 - Reactive potentials: ReaxFF and COMB potentials
 - LAMMPS provides a comprehensive library of empirical potentials
- **Machine learning potentials**
 - Neural network potentials (NNP)
 - Gaussian approximation potentials (GAP)
 - Spectral neighborhood analysis potentials (SNAP)
 - Moment tensor potentials (MTP)

Machine Learning Potentials

- **There are a number of open-source packages for training ML potentials:**
 - Neural network potentials: DeepMD, PyXtal_FF
 - Gaussian approximation potentials: GAP, FLARE
 - Spectral neighborhood analysis potentials: FitSNAP, MAML
 - Moment tensor potentials: MLIP
- **Pros and cons in adapting existing packages:**
 - As these packages has plugins for LAMMPS, they can be used to train ML potentials and run MD simulations through LAMMPS.
 - As most of the existing packages are written in Python, it is difficult to integrate them with our Julia-based software stack.
 - Most of the packages do not quantify uncertainties in the trained ML potential.
 - Several packages do not include an empirical potential in the training.

The Julia Library of MD Potentials

- **Motivation for coupled empirical/ML potentials**
 - Existing empirical potentials (ReaxFF, COMB) have not produced satisfactory results for our target problem. Need to retrain ReaxFF and COMB.
 - ML potentials may be able to fill the gap.
 - ML potentials require significant amount of DFT data for training. Coupled ML/empirical potentials may reduce the amount of data needed to train the model.
- **MDP.jl The Julia library of MD potentials:**
 - Explore a variety of atomistic descriptors and ML methods
 - Exploit automatic differentiation to accelerate training and force calculation
 - Identify near optimal configurations to include in the training data
 - **Provide a comprehensive library of coupled empirical and ML potentials**
 - **Quantify uncertainties for the trained potentials**
 - **Provide fast and accurate potentials for classical MD**
 - **Provide LAMMPS plugins to accelerate MD simulations on exascale supercomputers**

Potential Energy Representation

- Hybrid characterization of the potential energy surface

$$E(\mathbf{r}^N) = E_{\text{EP}}(\mathbf{r}^N) + E_{\text{ML}}(\mathbf{r}^N)$$

- Machine learning potential

$$E_{\text{ML}}(\mathbf{r}^N) = \mathcal{G}(\mathbf{d}(\mathbf{r}^N), \mathbf{w})$$

$$\mathbf{d}(\mathbf{r}^N) = \{d_1(\mathbf{r}^N), \dots, d_M(\mathbf{r}^N)\} \quad (\text{descriptors})$$

$$\mathbf{w} = (w_1, \dots, w_K) \quad (\text{weights})$$

- The weights are determined by minimizing a loss function.

Thompson, A. P., Swiler, L. P., Trott, C. R., Foiles, S. M., & Tucker, G. J. (2015). **Spectral neighbor analysis method for automated generation of quantum-accurate interatomic potentials.** *Journal of Computational Physics*, 285, 316-330.

Regression Methods

- **Linear regression (SNAP, MLIP):**

$$E_{\text{ML}}(\mathbf{r}^N) = \sum_{m=1}^M w_m d_m(\mathbf{r}^N)$$

- **Gaussian process regression (GAP, FLARE):**

$$E_{\text{ML}}(\mathbf{r}^N) = \sum_{k=1}^K w_k \mathcal{K}(\mathbf{d}(\mathbf{r}^N), \mathbf{d}(\chi_k))$$

- **Neural network (nonlinear) regression (DeepMD, MAML):**

$$E_{\text{ML}}(\mathbf{r}^N) = \sum_{i=1}^N \left(b_i + \sum_{k=1}^{N_h} w_{ik} \mathcal{H}(\mathbf{d}(\mathbf{r}^N), \mathbf{u}_{ik}) \right)$$

Descriptors

- **Descriptors are crucial to the performance of ML potentials**
 - Atom centered symmetry functions (ACSF): neural network potentials
 - SO(3) descriptors: Smooth overlap of atomic position (SOAP) potentials, moment tensor potentials, DeepMD potentials
 - SO(4) descriptors: GAP and SNAP potentials
- **They key requirements of the descriptors:**
 - The descriptors are invariant to permutation, translation, and rotation
 - The descriptors and their derivatives are fast to evaluate
 - The descriptors are able to capture various atomic interaction phenomena
- **In addition to existing descriptors, we consider a new SO(3) descriptor.**

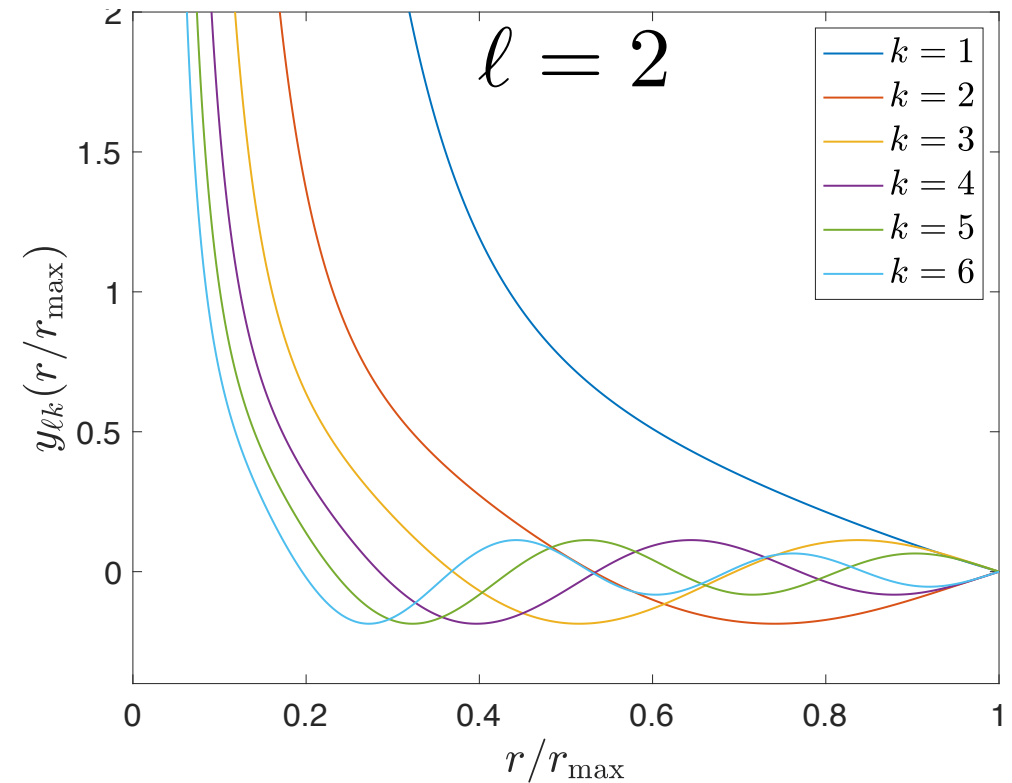
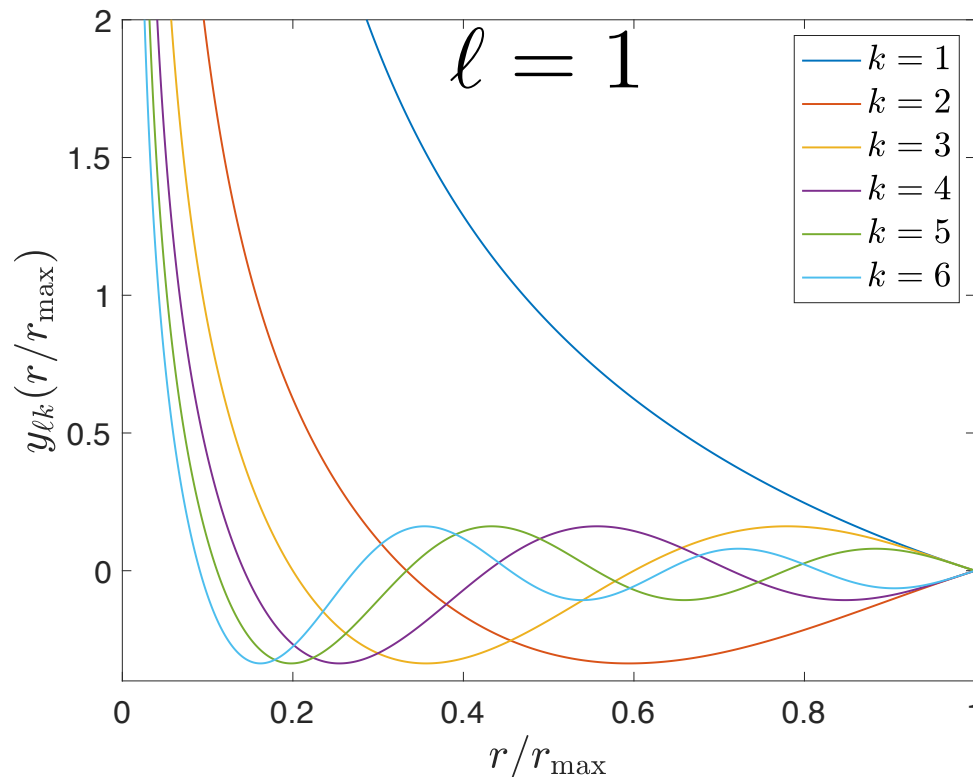
Bartók, Albert P., Risi Kondor, and Gábor Csányi. "*On representing chemical environments.*" *Physical Review B* 87.18 (2013): 184115.

Behler, Jörg. "*Perspective: Machine learning potentials for atomistic simulations.*" *The Journal of chemical physics* 145.17 (2016): 170901

New SO(3) Descriptors

- The descriptors are based on products of spherical harmonics $Y_{lm}(\theta, \phi)$ and spherical Bessel functions $y_{lk}(r)$

$$u_{klm}(\mathbf{r}) = y_{lk}(r)Y_{lm}(\theta, \phi), \quad 1 \leq k \leq K, 0 \leq l \leq L, -l \leq m \leq l$$



New SO(3) Descriptors

- Introduce local atomic functions

$$a_{iklm}(\mathbf{r}^N) = \sum_j^{|\mathbf{r}_j - \mathbf{r}_i| \leq r_c} u_{klm}(\mathbf{r}_j - \mathbf{r}_i), \quad 1 \leq i \leq N$$

- Define the power spectrum components of the descriptors:

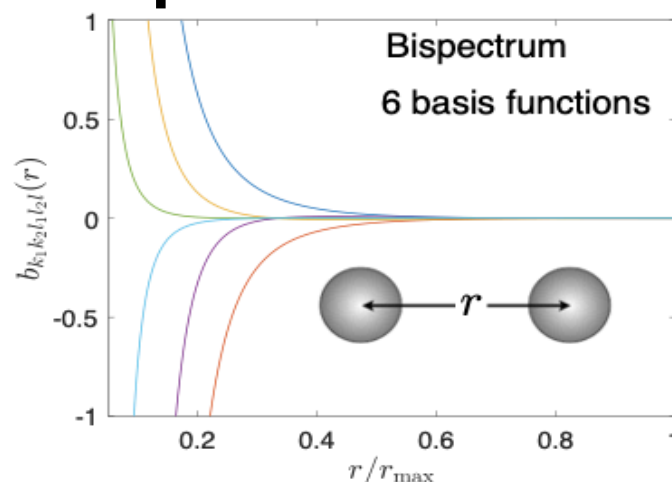
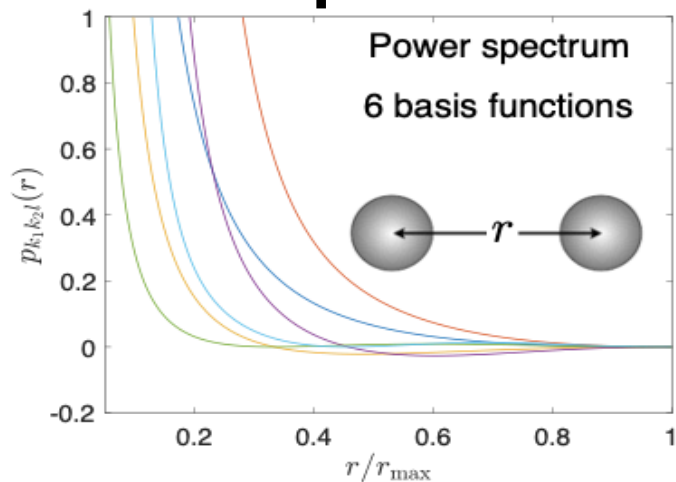
$$p_{ikk'l}(\mathbf{r}^N) = \sum_{m=-l}^l a_{iklm}^*(\mathbf{r}^N) a_{ik'l m}(\mathbf{r}^N), \quad 1 \leq i \leq N$$

- Define the bispectrum components of the descriptors:

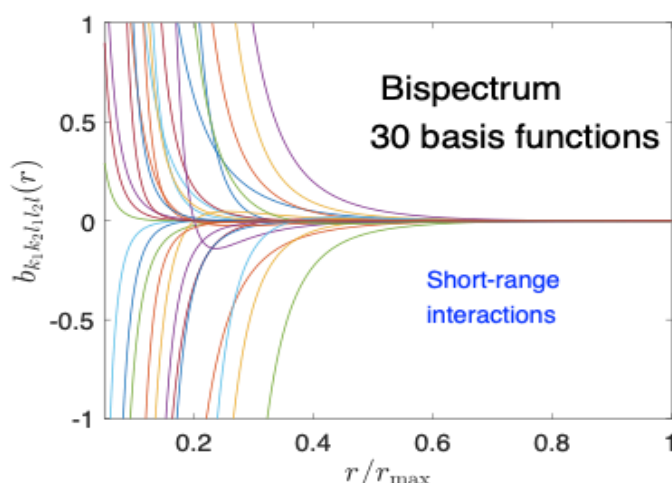
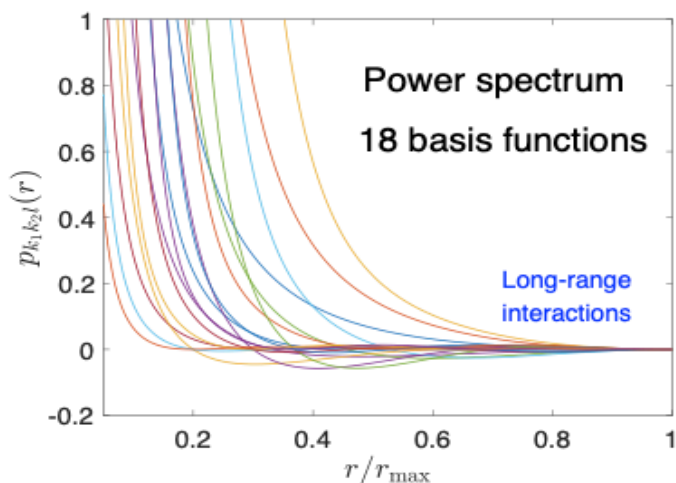
$$b_{ikk' ll_1 l_2}(\mathbf{r}^N) = \sum_{m=-l}^l \sum_{m_1=-l_1}^{l_1} \sum_{m_2=-l_2}^{l_2} a_{iklm}^*(\mathbf{r}^N) C_{m_1 m_2 m}^{l_1 l_2 l} a_{ik'l_1 m_1}(\mathbf{r}^N) a_{ik'l_2 m_2}(\mathbf{r}^N)$$

New SO(3) Descriptors

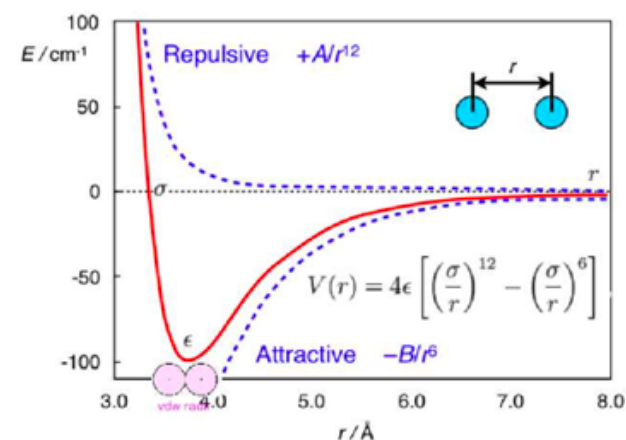
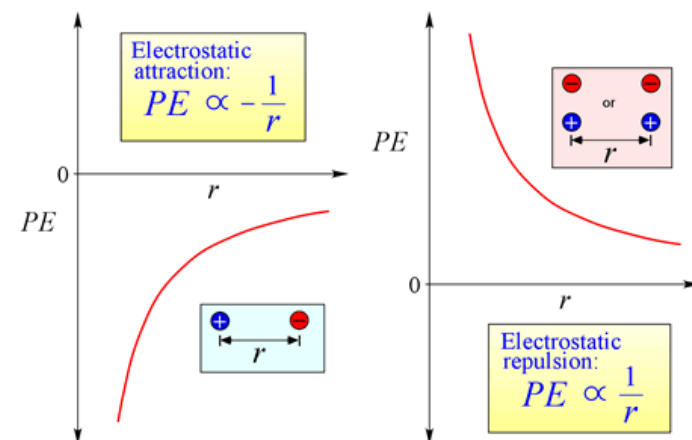
- Radial components of the descriptors



Radial basis functions for $K = 2$ and $L = 1$ (first-order)



Radial basis functions for $K = 3$ and $L = 2$ (second-order)



The Lennard-Jones potential

Automatic Differentiation

- Force calculation needs the partial derivatives of the descriptors
- For SO(3) and SO(4) descriptors, their analytical derivatives are an order of magnitude more expensive than to compute the descriptors.
- We will use automatic differentiation to calculate the derivatives to reduce the cost of force calculation.
- **Enzyme** is a high-performance AD software developed by our CS group.

C Source

```
double relu3(double x) {  
    double result;  
    if (x > 0)  
        result = pow(x, 3);  
    else  
        result = 0;  
    return result;  
}
```

<https://enzyme.mit.edu/>

Enzyme Usage

```
double diffe_relu3(double x) {  
    return __enzyme_autodiff(relu3, x);  
}
```

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Potential Fitting

- Find the weights to minimize the following loss function

$$\boldsymbol{w}^* = \arg \min_{\boldsymbol{w}} \sum_{j=1}^J \alpha_j (E_{\text{ML}}(\boldsymbol{\chi}_j, \boldsymbol{w}) - \bar{E}(\boldsymbol{\chi}_j))^2 + \beta_j \|\boldsymbol{F}_{\text{ML}}(\boldsymbol{\chi}_j, \boldsymbol{w}) - \bar{\boldsymbol{F}}(\boldsymbol{\chi}_j)\|^2,$$

where

$$\bar{E}(\boldsymbol{\chi}_j) = E_{\text{DFT}}(\boldsymbol{\chi}_j) - E_{\text{EP}}(\boldsymbol{\chi}_j)$$

$$\bar{\boldsymbol{F}}(\boldsymbol{\chi}_j) = \boldsymbol{F}_{\text{DFT}}(\boldsymbol{\chi}_j) - \boldsymbol{F}_{\text{EP}}(\boldsymbol{\chi}_j)$$

- How do we choose the configurations $\{\boldsymbol{\chi}_1, \dots, \boldsymbol{\chi}_J\}$?
- How do we ensure the accuracy of the ML potential?
- How do we choose new configurations if necessary?

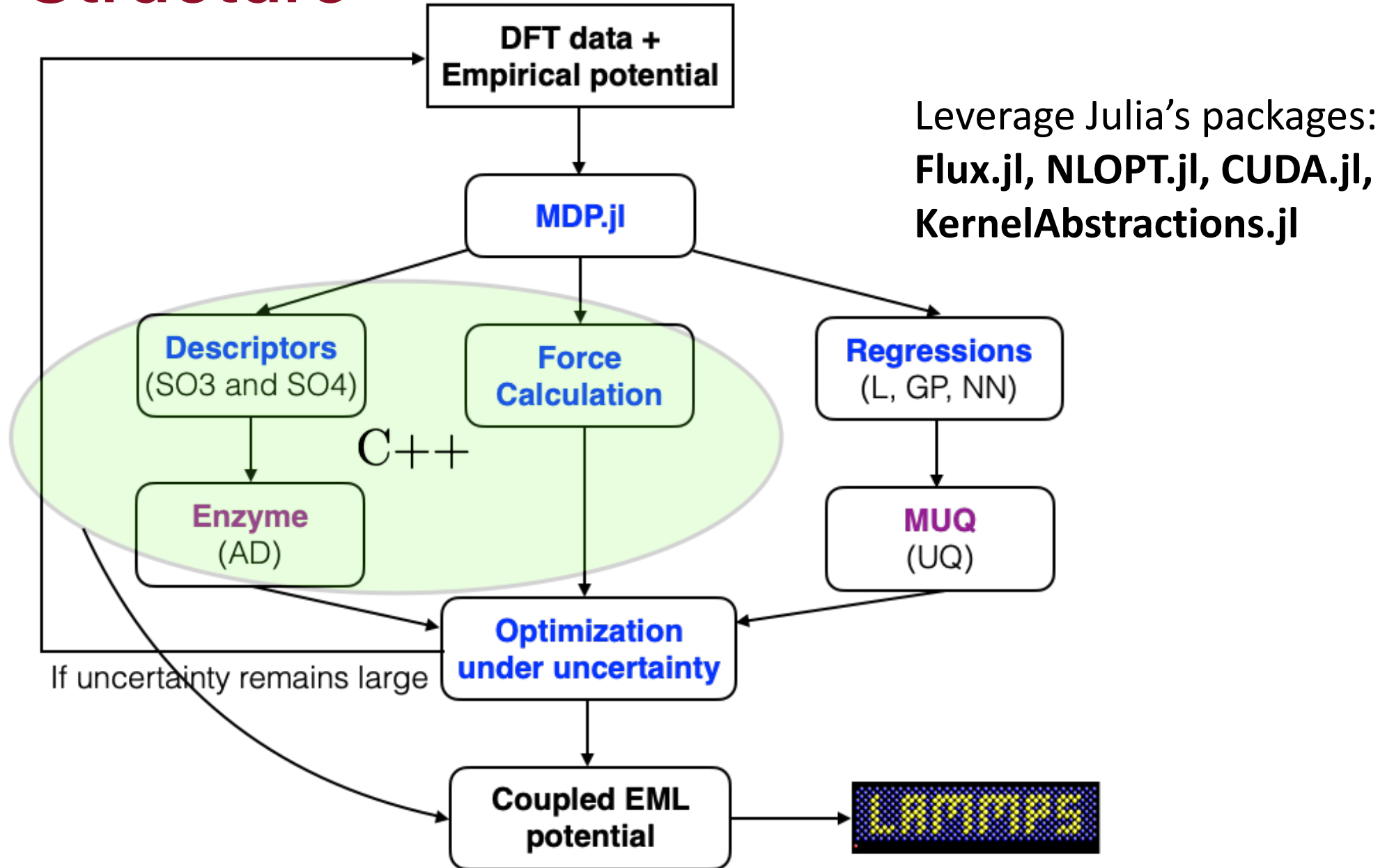
Uncertainty Quantification

- A **key advantage** of developing our own potential-fitting code (besides flexibility, performance, integration) is the ability to **introduce UQ** from the ground up!
- **UQ strategy:**
 - Recast deterministic/optimization-driven potential fitting (point estimation) as *fully Bayesian inference*
 - Characterize posterior on potential parameters either using *sampling* algorithms (MCMC) or fast/approximate *variational inference*
 - Gradients of the misfit (log-likelihood) will drive either of these approaches
 - Gradient-based dimension reduction and sensitivity analysis also useful
 - Use posterior uncertainty in predictions to drive *active learning* approaches that select/demand new DFT data
 - Calculations much easier in the Gaussian process case (cf. GAP and FLARE), but we will seek greater generality

Uncertainty Quantification

- Longer-term: start thinking about *structural error* in potentials, and adaptation of model complexity
 - Model discrepancy terms
 - Adaptive multi-fidelity potentials
 - Robust Bayesian inference under misspecification
- Stand-up of our initial UQ strategy is a priority for latter half of Year 1
- Make use of experience/algorithms from MUQ (<http://muq.mit.edu>), but write new Julia libraries linked to MDP.jl

MDP.jl Structure



Questions/Comments?

- **MDP.jl: The Julia library of MD potentials**
 - Implement a variety of atomistic descriptors and ML methods
 - Identify near optimal configurations to include in the training data via UQ methods
 - Employ automatic differentiation to accelerate training and force calculation
 - **Provide a comprehensive library of coupled empirical and ML potentials**
 - **Quantify uncertainties for the trained potentials**
 - **Enable fast and accurate MD simulations by implementing and auto-tuning new algorithms for force calculation (tomorrow talk)**