Project proposal

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Abstract

Abstract goes here if we need it.

Introduction

Background Research

Many fields require materials with properties that often conflict or trade off with one another. Examples of such properties are low density, high strength, and high flexibility. In most cases, it is difficult or impossible for traditional materials to exhibit combinations of these properties.

Methodology

The overall design process can be seen in the following figure.

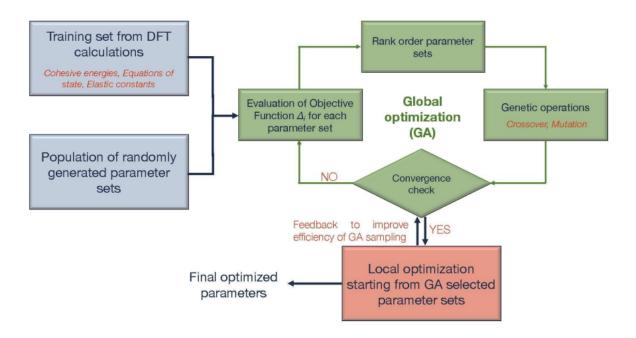


Figure 1: Overall Flow of Genetic Algorithm Process. Adapted from Narayanan et al., 2017

The main optimizations will be taking place within the global optimization loop and the local optimization section, as seen above. The loop will contain the majority of our implementation. The local optimization section will be carried out primarily with LAMMPS.

A global single-population master-slave genetic algorithm was chosen as the best method of parallelization. The master node receives a population and divides individuals between the slave nodes. Slave nodes will evaluate the fitness of individuals they receive and send the results back to the master node. A visualization of this technique can be seen below.

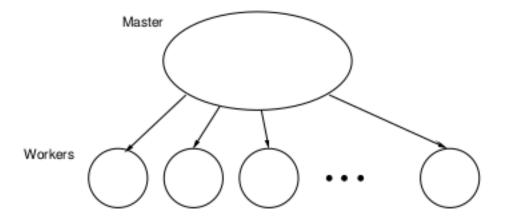


Figure 2: Master-Slave Distributed Genetic Algorithm. Adapted from Cantú-Paz, 1998

We will realize this solution using several nodes on the Beocat supercomputing cluster. One computing node will be designated as the master node, while several other nodes will be the slave nodes. The master node will receive the population of randomly generated parameter sets and the training set provided. It will partition the population into subsets and send the sets to slave nodes for evaluation. Each slave node will evaluate the fitness of each individual within their subset and transfer the results to the master node for modification via trait crossover and mutation. The best one hundred individuals are retained for subsequent generations.

This process will continue until the genetic algorithm fails to return better individuals than the previous run. At this convergence step, the twenty best individuals are selected to be locally minimized by using the Simplex method. Finally, the most optimized individual results in the best set of parameters for this generation.

The Large-scale Atomic/Molecular Massively Parallel Simulator (LAMMPS) package will be used to evaluate individuals' fitness. This software is a classical molecular dynamics code developed by Sandia National Laboratories. It has optimized integrations with MPI, which will fit into our overall design very well.

Conclusion

References

- Cantú-Paz, E. (1998). A survey of parallel genetic algorithms. Calculateurs paralleles, reseaux et systems repartis, 10(2), 141-171.
- Narayanan, B., Chan, H., Kinaci, A., Sen, F. G., Gray, S. K., Chan, M. K. Y., & Sankaranarayanan, S. K. R. S. (2017). Machine learnt bond order potential to model metal-organic (co-c) heterostructures. *Nanoscale*, doi:10.1039/C7NR06038F