Spring 2019: Advanced Topics in Numerical Analysis: High Performance Computing Final Project Proposal

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We will work on parallelizing code from an old research project that implements Kinetic Monte Carlo (KMC) in serial. KMC is an algorithm for modelling the dynamics of a continuous time jump Markov process. The process we will consider is atom movement on the surface of a crystal. In 2D we consider a lattice $\Omega=\{1,\ldots,L\}\times\{1,\ldots,L\}$. At each site $\vec{\ell}\in\Omega$ on the lattice we have atoms stacked on top of each other. The number of atoms at each site $\vec{\ell}$ will be denoted by the height of the stack $h_{\vec{\ell}}$. The surface of the crystal is then just the atoms at the top of each stack. KMC simulates a Markov jump process describing the evolution of the vector $(h_{\vec{\ell}}(t))_{\vec{\ell}\in\Omega}$ by randomly choosing one lattice site, at which the topmost atom will jump to the top of one of the neighboring stacks. The rates of this process are governed by the height difference between two neighboring stacks. In the serial implementation of KMC, only one jump occurs at a time, and all other atoms, even those far away, must wait before getting a chance to move.

To parallelize KMC we will divide the lattice into sections and then independently run Markov jump processes on each section. After every process has run for a while we will need to synchronize the sections since the jump times are random. We will also need to handle potential conflicts on the boundaries of the sections where atoms may be jumping across. This could potentially be handled with a layer of ghost lattice sites. We plan to transfer the computation to a GPU for the parallelization.

For large values of L, this Markov process can be seen as a discretized version of continuous dynamics (described by a PDE) governing the evolution of a smooth surface h(x,t). This PDE is known and solutions to it can be computed numerically. To test our implementation, we can evolve a continuous height profile forward for a certain amount of time using the PDE, evolve a discretization of that profile for the same amount of time using parallel KMC, and compare the two resulting profiles.