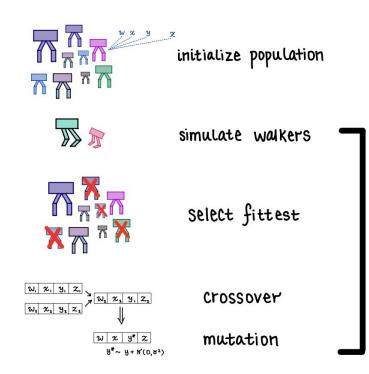


## Sequential Baseline

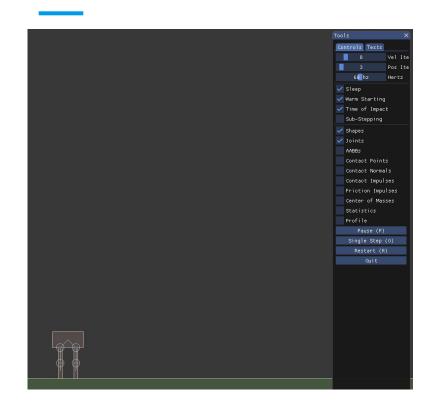
#### Algorithm

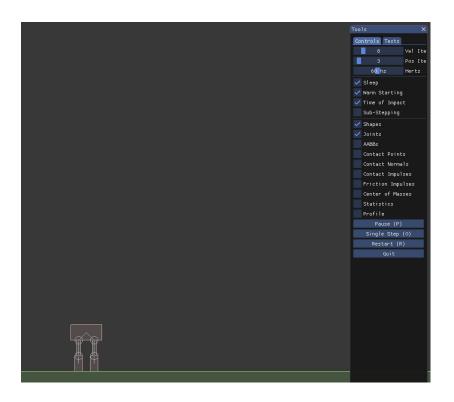
- 1. Create a population
- 2. Simulate walkers
- 3. Select the fittest walkers
- 4. Cross and mutation of the fittest walkers
- 5. Repeat steps 2-4



REPEAT

### **Initial Visualization**





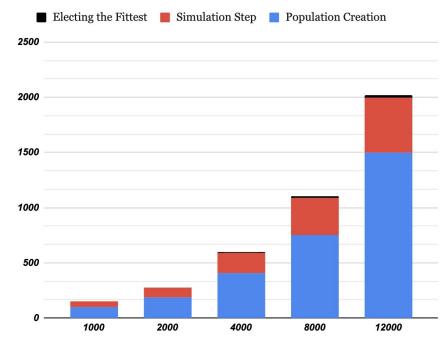
#### **Bottlenecks**

#### Bottlenecks after profiling:

- Creating a new population is the most expensive step
- 2. **Simulating a step** for each of the walkers
- 3. **Memory constraints** on sequential implementation. Opportunities for scaling and parallelizing (Cache locality and exploiting multiple processes can be explored).

The algorithm is synchronization- and not compartmentalized computation-heavy

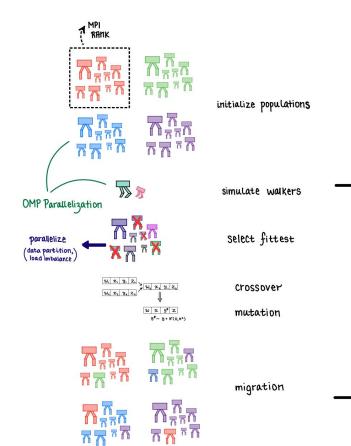




Number of Walkers

### Parallelism

- Create population with MPI
  - Each rank takes a subset of the population to be created
  - Every rank simulates the walkers of the subpopulation
  - Reports the fitnesses and the torque
- Simulation
  - Pass walkers to Box2D environment, calculate distance travelled
- Select the fittest walkers
  - Sorting algorithm
- Migration
  - Randomly select 5% of the walkers from population A → population B
  - o Communication between ranks



REPEAT

# Migration – Student Visas

