Simulation:

* -two patches, 2 genotypes (assumed that each one has replacement >1 for one patch, <1 for the other, the favored patch for one genotype was the unfavored patch for the other genotype)
* -takes user input for carrying capacities, lambdas
* -simulates stepwise operations of reproduction [defined by equations in PDF], combination into the common offspring pool (“eggland”), dispersal [into the patch that favors the more common genotype and then the other, defined by equations in PDF] for 1000 years
* Recorded relative frequencies of the two genotypes, and an analog to relative migration rates [need to get specific mathematical definition from Jaime]
* Use analog to relative migration rates to define an actual inversion
* Ran 1000 simulations with KD varying from .01KE to 4KE, looking at the relative migration rates at the final year to show what ratios of KD:KE cause source-sink inversion.