Current assumptions:

* State matrix initialization:
  + Assume that the input csv file already has all the people that will ever be alive in the simulation, and the features have been populated based on some joint distribution of some of the features
* Determining birth of people:
  + The people eligible to be born are those who are not in the simulation
* The format of the csv that contains the probability distributions for each of the population groups will be age, race, MSM, parameter1, parameter 2, etc.
  + Because currently we are using the Gaussian distribution for transition probabilities, we are only specifying two parameters (sigma and mu)