pHandDyingRoughDraft

September 24, 2018

1 pH and Dying

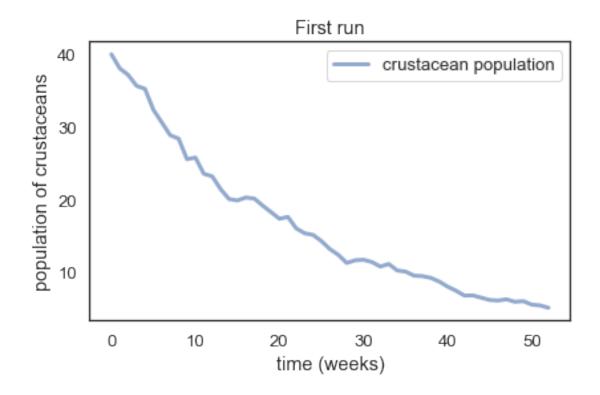
Meagan Rittmanic & Dieter Brehm - Project 1 - Fall 2018 A project examining how to model an assumed relation between crustaceans and pH levels

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In [2]: # Configure Jupyter so figures appear in the noebook
        %matplotlib inline
        # import functions from the modsim.py module
        from modsim import *
        import random
        import matplotlib.pyplot as plt
In [8]: # Plot our data points for pH levels and crustacean birth rates
        plt.plot([9,9.5,10,10.5], [11.1, 9.7, 9, 7.6], '*')
        # Create a best-fit line
        def run_match():
            match = TimeSeries()
            match[9] = 11.1
            for p in linrange(9, 11, 1):
                match [p+1] = match[p] - 2.33
            return match
        # overlay the fit line with our data
        p2 = run_match()
        plot (p2)
        plt.show()
```

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In [5]: def run_match2():
            match = TimeSeries()
            match[9] = 11.1
            for p in linrange(9, 11, 1):
                match [p+1] = match[p] - 2.33
            return match[11]
        run_match2()
Out[5]: 6.439999999999995
In [80]: # set our constants
        birth_rate_normal = 0.12
         birth = run_match2()
         birth1 = birth_rate_normal*(birth/11.1)
         values = System(death=0.1, birth_lb = birth1-0.07, birth_ub = birth1+0.07, pop_0 = 40
In [81]: def run_ph_sim(system):
             11 11 11
             Runs the simulation
             system: a System object with our initial constants, relies on:
                     - pop_O: initial population
                     - birthi: lower range deviation of birthrate due to pH
                     - birthii: upper range deviation of birthrate due to pH
                     - death: death rate
```

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Return TimeSeries with results
             11 11 11
             run = TimeSeries()
             run[0] = system.pop_0
             for t in linrange (0, values.time_period, 1):
                 for x in range(1):
                     b = random.uniform(system.birth_lb, system.birth_ub)
                 run[t+1] = run[t] + run[t]*(b)-run[t]*(system.death)
             return run
         def plot_rps(variable, title):
             Plots data
             variable: the data we want to graph
             title: the title of the graph
             directly plots data with no return statement
             plot (variable, label = 'crustacean population')
             decorate(xlabel = 'time (weeks)',
                     ylabel = 'population of crustaceans',
                     title = title)
In [82]: x = run_ph_sim(values)
        plot_rps(x, 'First run')
```

- time period: how many weeks to run the simulation for

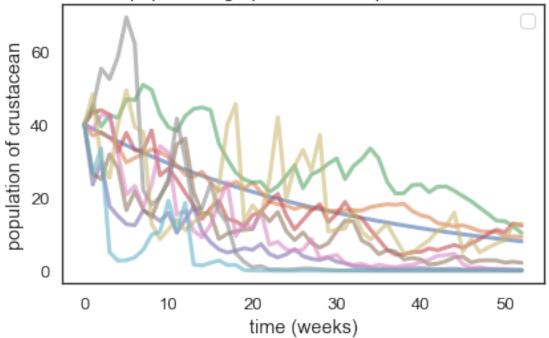


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In [112]: # sweep for a few different birthrate / pH ranges
    birth_ub_array = linspace(0, 0.9, 10);

for rate in birth_ub_array:
    # for now, let's keep the upper and lower limits absolute value the same
    values.birth_lb = birth1 - rate
    values.birth_ub = birth1 + rate
    x = run_ph_sim(values)
    plot(x)

decorate(xlabel = 'time (weeks)',
    ylabel = 'population of crustacean',
    title = "population graph with varied pH/birthrate")
```





```
In [113]: def run_ph_sim2(system):
              Runs the simulation
              system: a System object with our initial constants, relies on:
                      - pop_0: initial population
                      - birthi: lower range deviation of birthrate due to pH
                      - birthii: upper range deviation of birthrate due to pH
                      - death: death rate
                      - time period: how many weeks to run the simulation for
              Return TimeSeries with 52nd result
              n n n
              run = TimeSeries()
              run[0] = system.pop_0
              for t in linrange (0, values.time_period, 1):
                  for x in range(1):
                      # randomly select a decimal number between
                      # the lower and upper birth rate(pH) bounds
                      b = random.uniform(system.birth_lb, system.birth_ub)
                  run[t+1] = run[t] + run[t]*(b)-run[t]*(system.death)
              return run[52]
```

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In [114]: def run_program(number_of_runs, system, state):
              Runs the simulation multiple times
              number_of_runs: how many times we want to run the simulation
              system: a System object with our initial constants, relies on:
                      - extinct: tracks number of runs in which the population went extinct
              state: a State object with the number of extinctions that have occured
              11 11 11
              for i in range (number_of_runs):
                  x = run_ph_sim2(system)
                  if x<=8:
                      extinct.extinct +=1
              return extinct.extinct
In [115]: # run the simulation 100 times
          extinct = State(extinct = 0)
          extinctions = run_program(12, values, extinct)
          print(extinctions)
```

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