```
AMATH 422 HW 2
In [1]: #imports
        import matplotlib.pylab as plt # That gives plotting, and the next line makes plots appear inl
        %matplotlib inline
        import numpy as np # That gives numerical arrays and tools for manipulating them
        import scipy.optimize as opt
        import scipy.linalg as la
        QUESTION I (a)
In [2]: #intial conditions
        Ia_arr=np.array([3,0.5,0.9,0.95])
        fa arr=np.array([0,1,5,0.5])
        initial_population = np.array([100, 100, 100, 100]) # initial population for each age
        n_zero=np.array([100,100,100,100])
        A_mat=np.array([[0,1,5,0.5],[0.5, 0, 0, 0],[0, 0.9, 0,0],[0, 0.0, 0.95,0]])
        print(A mat)
        print(n_zero)
      [[0.
                      0.5 1
             1.
                  5.
                      0. ]
       [0.5 0.
                  0.
       [0. 0.9 0.
                  0.95 0. ]]
             0.
      [100 100 100 100]
In [3]: Tmax=50
        n_vs_t=np.zeros([4,Tmax])
        n_vs_t[:,0]=n_zero
        #Print out our quanties so far
        print(n_vs_t[:,0])
        print(A_mat)
        print(np.dot(A_mat,n_vs_t[:,0]))
      [100. 100. 100. 100.]
      [[0.
            1. 5.
                      0.5 ]
       [0.5 0. 0.
                      0. ]
       [0. 0.9 0.
                      0. ]
             0. 0.95 0. ]]
       [0.
      [650. 50. 90. 95.]
```

```
In [4]: #Generation of Values
   iter_arr=np.arange(Tmax-1)

for t in iter_arr:
        n_vs_t[:,t+1]=np.dot(A_mat,n_vs_t[:,t])

#calculate the log of n(t)
   log_n_vs_t = np.log(n_vs_t)

# Calculate total population size N(t) at each time step
   N_t = np.zeros(Tmax)
   for t in range(Tmax):
```

```
N_t[t] = np.sum(n_vs_t[:, t])

print(n_vs_t.shape)
print(log_n_vs_t.shape)
print(N_t.shape)
print(iter_arr.shape)

(4, 50)
(4, 50)
(50,)
(49,)
```

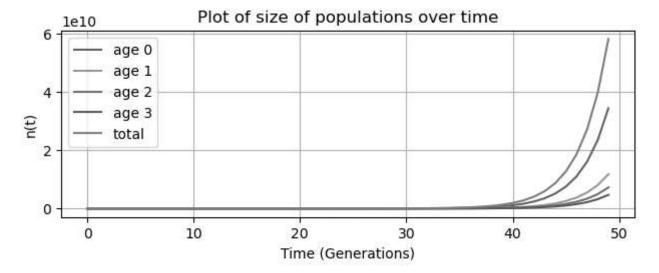
Generate Graphs \

```
Plot of each N(t) \
Plot of each log N(t) \
Plot of each fraction of N(t)
```

```
In [5]: #Plot of each fraction of N(t)

generation_arr=np.arange(Tmax)
plt.subplot(2, 1, 2)
plt.plot(generation_arr,n_vs_t[0,:],label="age 0")
plt.plot(generation_arr,n_vs_t[1,:],label="age 1")
plt.plot(generation_arr,n_vs_t[2,:],label="age 2")
plt.plot(generation_arr,n_vs_t[3,:],label="age 3")
plt.plot(generation_arr,N_t,label="total")

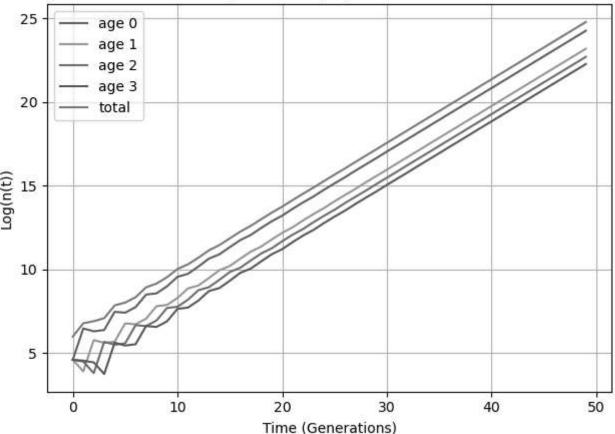
plt.xlabel('Time (Generations)')
plt.ylabel('n(t)')
plt.title('Plot of size of populations over time')
plt.legend()
plt.grid(True)
plt.tight_layout()
plt.show()
```



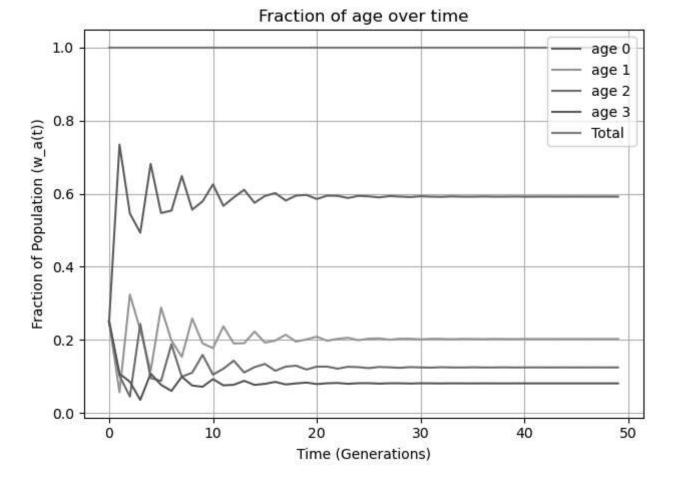
```
In [6]: #plot of log(n) vs t
    plt.plot(generation_arr,np.log(n_vs_t[0,:]),label="age 0")
    plt.plot(generation_arr,np.log(n_vs_t[1,:]),label="age 1")
    plt.plot(generation_arr,np.log(n_vs_t[2,:]),label="age 2")
    plt.plot(generation_arr,np.log(n_vs_t[3,:]),label="age 3")
    plt.plot(generation_arr,np.log(N_t),label="total")
```

```
plt.xlabel('Time (Generations)')
plt.ylabel('Log(n(t))')
plt.title('Plot of log of size of populations over time')
plt.legend()
plt.grid(True)
plt.tight_layout()
plt.show()
```

Plot of log of size of populations over time



```
In [7]: # Calculate the fraction of individuals in each age group w_a(t) = n_a(t) / N(t)
        w vs t = np.zeros((4, Tmax))
        for t in range(Tmax):
            w_vs_t[:, t] = n_vs_t[:, t] / N_t[t]
        print(w_vs_t.shape) #should be a (4, 50 matrix)
        plt.plot(np.arange(Tmax), w_vs_t[0, :], label='age 0')
        plt.plot(np.arange(Tmax), w_vs_t[1, :], label='age 1')
        plt.plot(np.arange(Tmax), w_vs_t[2, :], label='age 2')
        plt.plot(np.arange(Tmax), w_vs_t[3, :], label='age 3')
        w_t_sum = np.zeros(Tmax)
        for t in range(Tmax):
            w_t_sum[t] = np.sum(w_vs_t[:, t])
        plt.plot(np.arange(Tmax), w t sum, label='Total')
        plt.xlabel('Time (Generations)')
        plt.ylabel('Fraction of Population (w a(t))')
        plt.title('Fraction of age over time')
        plt.legend(loc='upper right')
        plt.grid(True)
        plt.tight_layout()
```



Use the numpy polyfit function to fit a first order polynomial to the log N(t) and report the growth rate lambda.

```
In [8]: p=np.polyfit(x=generation_arr,y=np.log(N_t),deg=1)
    lambda_estimate=np.exp(p[0])
    print(p)
    print(lambda_estimate)
```

[0.38019638 6.15837247] 1.4625717774380425

Lambda Estimate without the Squiggles

```
In [9]: generation_arr_after_squiggles=np.arange(25,Tmax)

p=np.polyfit(generation_arr_after_squiggles,np.log(n_vs_t[0,generation_arr_after_squiggles]),1)
lambda_estimate=np.exp(p[0])

print(p)
print(lambda_estimate)
```

[0.38006369 5.63874003] 1.4623777242600995

Question I (b)

Write down the Euler-lotka formula for this example, and solve it numerically or the population growth rate.

Define Function for the Euler-Lotka sum. $\sum_{a=0}^n \lambda^{-(a+1)} I_a f_a - 1$ where I_a and f_a are two 1-D arrays and λ is a scalar.

```
#Write down the Euler-lotka formula for this example, and solve it numerically or the population
         #growth rate lambda: How close are your predictions of lambda from the Euler-lotka formulas and j
         #simulations above? Turn in the code you used for this.
         import numpy as np
         import matplotlib.pyplot as plt
         from scipy import optimize as opt
         def eulot func(lam,Ia arr,fa arr):
                 """compute the Euler-Lotka sum, taking as arguments a scalar and two 1-D numpy arrays"""
                 length_of_array=Ia_arr.size
                 age_arr=np.arange(0,length_of_array)
                 temp_arr=lam**(-(age_arr+1))*Ia_arr*fa_arr
                 return sum(temp arr) -1
         Ia_arr=np.array([3,0.5,0.9,0.95])
         fa_arr=np.array([0,1,5,0.5])
         #Lambda estimate?
In [11]: # Range of Lambda values to plot over
         lambda_min = 0.1
         lambda max = 5
         lambda_arr = np.linspace(lambda_min, lambda_max, 100)
         # Compute G(lambda) for each value in the range
         G arr = np.zeros(lambda arr.size)
         for j in range(lambda_arr.size):
             G_arr[j] = eulot_func(lambda_arr[j], Ia_arr, fa_arr)
         # Plotting G(lambda)
         plt.plot(lambda arr, G arr)
```

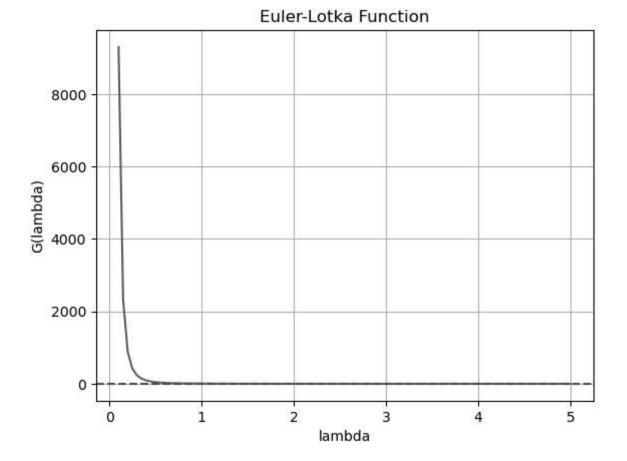
plt.axhline(0, color='r', linestyle='--') # Line at G=0 for reference

In [10]: #Euler Lokta Formulation

plt.xlabel('lambda')
plt.ylabel('G(lambda)')

plt.grid(True)
plt.show()

plt.title('Euler-Lotka Function')



```
In [12]: #Brent's method to find the root of the function between the given brackets
left_bracket = 0.5
right_bracket = 2
args = (Ia_arr, fa_arr)

#Solve for lambda using brentq
lambda_solution = opt.brentq(eulot_func, left_bracket, right_bracket, args=args)
print("Estimated population growth rate (lambda):", lambda_solution)
```

Estimated population growth rate (lambda): 1.7818660191092137

First Order Polynomial Estimate: 1.4623777242600995

Euler-Lokta Estimate: 1.7818660191092137

The difference between the estimates is because parts of the matrix population model has transient dynamics before the population reaches a stable age distribution, whereas the Euler-Lotka equation assumes a stable age distribution from the beginning. By extending the simulation to a longer time period, the population should approach a stable age structure, and the estimate of λ from the simulation will converge to the value obtained from the Euler-Lotka equation.

Question II (a)

Population Growth Rate Depends on Overall Survival: The long-term population growth rate λ . λ depends on the overall survival to reproductive age and reproductive output (i.e., the number of offspring produced by individuals who reach breeding age). Since survival to age 3 is represented by I3, the precise distribution of survival across earlier ages does not change the total survival to age 3 as long as the product remains constant.

Question II (b)

```
In [13]: import numpy as np
         # Define parameters
         p0, p1, p2 = 0.3, 0.4, 0.6 # Example survival probabilities with p0 * p1 * p2 = 0.0722
         pa = 0.942 # Adult survival probability for ages 3 to 49
         fa = 0.240 # Fecundity rate for ages 3 and above
         max_age = 50 # Max age Size of A.
         # Construct the projection matrix A
         A = np.zeros((max_age, max_age))
         # Set fecundities in the first row for ages 3 and above
         A[0, 3:] = fa
         # Set survival probabilities in the subdiagonal
         A[1, 0] = p0
         A[2, 1] = p1
         A[3, 2] = p2
         A[4:, 3:-1] = np.eye(max_age - 4) * pa
         print(A)
         print(A.shape)
        [[0.
               0.
                           ... 0.24 0.24 0.24 ]
        [0.3
                           ... 0.
               0.
                     0.
                                     0.
                                           0.
                                               ]
        [0.
               0.4
                     0.
                           ... 0.
                                     0.
                                           0.
        . . .
        [0.
               0.
                     0. ... 0.
                                     0.
                                           0.
        [0.
                     0.
                          ... 0.942 0.
                                           0.
                                                1
                           ... 0.
        [0.
                     0.
                                     0.942 0.
                                               ]]
        (50, 50)
```

Question II (c)

```
In [14]: # Compute the dominant eigenvalue (long-term growth rate lambda)
    eigenvalues = np.linalg.eigvals(A)
    lambda_long_term = max(eigenvalues.real)

print(f"Long-term growth rate (lambda): {lambda_long_term:.4f}")
```

Long-term growth rate (lambda): 0.9431