Operations

x + y addition x - y substraction

 $x \cdot y$ scalar product

x * y vectorial product

x. y power

Vectors

 $x = [1 \ 2 \ 3]$ create a row vector x = [2; 1; 5] create a column vector x = 1 : 10 : 1000 vector from 1 to 1000, by 10 Matrix

 $A = \begin{bmatrix} 1 & 2 & 3; 4 & 5 & 6; 7 & 8 & 9 \end{bmatrix} 3x3 \text{ matrix}$ B = zeros(2, 3) & 2 rows and 3 colums of 0 C = ones(2, 3) & 2 rows and 3 colums of 1 D = A' transpose of the matrixeig(A) eigenvalues of A

Plot

plot(x, y, '*') plot y as a function of x with * hold on, hold of f open or not a new figure xlabel('Myxlabel')ylabel('Myylabel')

Control flow

 $\begin{array}{l} for \ dummy = first: last \\ x[dummy] = dummy + 1; \\ end \\ if \ a > b \\ y = a + b; \\ else \\ y = a - b; \\ end \\ while \ i < j \\ i = i + 1; \\ end \end{array}$

A generic function

[out1, out2...] = function(in1, in2, ...)Leslie

rate = leslie(leslieMatrix, popVectorInit)

Density independent growth $rate = discrete(\lambda, n)$

Density dependent growth popSize = ricker(r, Ninit, K)

Competition

global r k alpha Only once $r = [r_1r_2]$; Growth rate $k = [k_1k_2]$; Carrying capacity $alpha = [\alpha \ \beta]$; Competition $n = [N_1N_2]$; Initial population size plotlv(n) Make graph

Competition 2 resources

global s j cA cB m Only once $s = [s_1s_2]$; Supply point cA = [c1c2]; Consumption rate of A cB = [c1c2]; Consumption rate of B $m = [m_Am_B]$; Mortality $j = [j_1j_2j_3j_4]$; Half saturation $n = [N_AN_B]$; Initial population size $R = [R_1R_2]$; Initial resource available tilman(n, R) Make graph

Predator-prey

global r a p m Only once r = Prey growth rate p = Capture effiency a = Conversion of biomass m = Predator mortality H Number of prey P Number of predator lotka(H, P) Make graph

Rosenzweig-MacArthur

global r a q c m K Only once r = Prey growth rate a = Conversion of biomass q = Feeding rate c = Half saturation m = Predator mortality K = Prey carrying capacity H Number of prey P Number of predator RM2(H, P) Make graph type II RM3(H, P) Make graph type III

SIR

global d D beta gamma v Only once d = Death rate non-infected D = Death rate infected beta = Transmission rate gamma = Recovery rate v = Vaccination S Susceptible I Infected R Recovered SIR(S, I, R) SIR model SIRv(S, I, R) SIRv model

Food chain

gurneychain Enter values phi = Photosynthetic primary production d* = Mortality of * a* = Predator * efficiency P = Primary producer H = Herbivore C = Predator