```
%clears all variables to ensure accuracy between runs.
clearvars
%-----%
                     Predator Prey Model 2.a
                    Dakota Ewing
<u>%______</u>
                    Variables
§_______
% To is the start month.
To = 1;
% Tf is the end month.
Tf = 12;
% dt is the time step in months.
dt = .0001;
% steps is the total number of timesteps.
Steps = (Tf - To + 1)/dt;
% T is the array in which we hold all the time values. It starts at To,
% increments by the time step, and ends at the target year, Tf.
T = To:dt:Tf;
% Yo is the starting population of Tuna.
Yo = 10000;
% Po is the starting population of Sharks.
% Ho is the population of humans, which remains constant.
Ho = 500;
% Y is the array in which we hold the number of Tuna at a given time.
% We initialize it to all zeros, except for the first month
% being simulated, which is initialized to Yo.
Y(To:Steps) = 0;
Y(To) = Yo;
% P is the array in which we hold the number of Sharks at a given time.
% We initialize it to all zeros, except for the first month
% being simulated, which is initialized to Po.
P(To:Steps) = 0;
P(To) = Po;
% H is the array in which we hold the number of Humans at a given time.
% We initialize every value to Ho, because the number of humans remains
% constant throughout the simulation.
% NOTE: This array is created to simplify the process of expanding the
% simulation to run with non-constant human populations.
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H(To:Steps) = Ho;
%Ky is the birth rate fraction of Tuna.
%Kp is the death rate fraction of Sharks.
Kp = 0.8;
%Khb is the birth rate fraction of Humans.
Khb = 0.000001;
%Khd is the death rate constant of Humans.
Khd = 0.0001;
%Kyp is the death proportionality constant of the Tuna.
Kyp = 0.0001;
%Kpy is the birth proportionality constant of the Sharks.
Kpy = 0.0001;
%Khf is the fishing rate of the Humans towards both Sharks and Tuna.
Khf = 0.000;
                             Simulation
<u>%______</u>%
t = 1;
%while the itterator is less than or equal to the total number of steps:
while t <= Steps
   %We calculate the change in Y, P and H for a dt by subtracting the
   % number of births from the number of deaths for the species.
   %The number of births of Y is proportional to the population of Y at t.
   Ybirths = Ky * Y(t);
   %The number of deaths of Y is proportional to the number of
   % interactions between Y and P at t.
   Ydeaths = Kyp * Y(t) * P(t);
   dy = Ybirths - Ydeaths;
   %The number of births of P is proportional to the number of
   % interactions between Y and P at t.
   Pbirths = Kpy * P(t) * Y(t);
   %The number of deaths of P is proportional to the population of P at t,
   %plus a proportion of interactions between P and H at t.
   Pdeaths = Kp * P(t) + Khf * P(t) * H(t);
   dp = Pbirths - Pdeaths;
   %The number of births of H is proportional to the number of
   % interactions between H and P at t.
   Hbirths = Khb * P(t) * H(t);
   %The number of deaths of H is proportional to the population of H at t.
   Hdeaths = Khd * H(t);
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dh = Hbirths - Hdeaths;

%The populations at t + 1 are equal to the populations at t, plus the
%change in the given population times the time step.
Y(t + 1) = Y(t) + dy * dt;
P(t + 1) = P(t) + dp * dt;
H(t + 1) = H(t) + dh * dt;

%t is incremented by 1.
t = t + 1;
end

* Plots

*plots the Y, P and H species over time.
plot(1:Steps, P(1:Steps),1:Steps, Y(1:Steps),1:Steps, H(1:Steps));

*plots the Y and P species in relation to each other.
plot(Y(1:Steps), P(1:Steps));
```