

Code used to generate simulation results for MS: *Predation can select for later and more synchronous arrival times in migrating species*. Harts, Kristensen & Kokko. In *Oikos*

This code was designed for use in Matlab. The code generates a few files based on last generation data, it can be modified to generate data for each generation (not shown).

Words in blue and black are code, annotations are in green.

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function[]=arrival_time(X,tmax,P0,initarriv,stdarriv,winterMort,nFocal,nTwo,Nterr,a,b,z,alpha,betaP,  
    betaG,gen,ms)  
  
% to start a simulation call e.g. arrival_time(850,5,0.05,0.5,0.5,0.1,1000,1000,500,0.5,5,1,0.5,1,2,5000,999)  
  
% X = daily handling time predators have on breeding grounds (a measure of predator abundance)  
  
% tmax = number of arrival days, number of days an individual can arrive on  
  
% P0 = daily mortality during migration period if not arrived yet  
  
% initarriv & stdarriv are for the initial populations arrival time allele, see below for calculation  
  
% migrMort = mortality during migration  
  
% nFocal = total number of individuals of the focal species (initial generation only)  
  
% nTwo = number of alternative prey (i.e. S2) alive at beginning of each spring  
  
% Nterr = total number of territories  
  
% a = predator preference, 1 = S1 and 0 = S2, at 0.5 there is no preference  
  
% b = handling time of S1 (has to be < X)  
  
% z = handling time factor such that bz=handling time of S2  
  
% alpha = proportion good territories (e.g. 0.5 means half is good and half is poor)  
  
% betaP and betaG: number of S1 offspring in poor and good territories respectively (beta gives the mean of the  
    poisson distribution which is #offspring per individual)  
  
% gen = number of generations the simulation runs for  
  
% ms = seed number, allows for replicating a specific simulation  
  
rng(ms) % setting seed for repeatability and independent simulations  
  
mutProb=0.1; mutStep=0.01; % mutation probability and determinant of size of mutation.
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%creation of initial population, each individual gets an allele for arrival time (Pop(:,1)) and initial %territorial state (Pop(:,2)) NaN means not arrived; 0 means floater, -1 means poor terr owner, 1 %means good terr owner

Pop=max(0,min(1,initarriv+stdarriv\*(rand(nFocal,1)-rand(nFocal,1))));

Pop(:,2)=NaN\*ones([nFocal 1]);

%data collected

meanArriv=NaN\*ones([1 gen]); % mean of arrival alleles in focal population (S1)

S1spring=NaN\*ones([1 gen]); % number of S1 at start of each generation

nof\_arrived=NaN\*ones([tmax gen]); % number of S1 arrived this time within each generation

dead\_arrived=NaN\*ones([tmax gen]); % actual number of S1 predated on breeding grounds / gen.

dead\_nonarrived=NaN\*ones([tmax gen]); % actual number of S1 predated on nonbreeding grounds /gen.

prodOff=NaN\*ones([1 gen]); % number of offspring produced this gen

track\_S2=NaN\*ones([tmax gen]); % number of S2 on each day t after predation during arrival S1

arrivHist=NaN\*ones([1 tmax]); % distribution of arrival allele in the last generation

for g=1:gen % start of new gen.

S2=nTwo; % each gen starts with the same number of individuals of S2

RF=0; % RF is number of floaters that have arrived, at start of each gen it is set to 0

meanArriv(g)=mean(Pop(:,1)); % calculate the mean arrival allele within the population for gen=g

S1spring(g)=length(Pop(:,1)); % calculate number of S1at start of gen=g

for t=1:tmax % arrival day within a gen.

% territory availability update

aG=alpha\*Nterr-sum(Pop(:,2)>0); % good territories still available

aP=(1-alpha)\*Nterr-sum(Pop(:,2)<0); % poor territories still available

% find those who are arriving now OR are floating, and

% randomize the order in which they are allowed to take territories

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if t<tmax

    f=vectperm(find(((Pop(:,1)>=(t-1)/tmax) & (Pop(:,1)<t/tmax)) | (Pop(:,2)==0))));

else % on t=tmax

    f=vectperm(find(((Pop(:,1)>=(t-1)/tmax) & (Pop(:,1)<(t+1)/tmax)) | (Pop(:,2)==0))));

end

%arrival and territory assignment (within each time t)

for j=1:length(f)

    if aG>0 % if there are good territories available

        aG=aG-1; % one good territory less available

        Pop(f(j),2)=1; % this indiv. is now the owner of a good terr

    elseif aP>0 % else if there are poor terr available

        aP=aP-1; % one less poor territory available

        Pop(f(j),2)=-1; % this indiv. is now the owner of a poor terr

    else

        RF=RF+1; % one more floater

        Pop(f(j),2)=0; % this indiv. is now a floater

    end

end

% predation of all individuals that have arrived at breeding site

% S1 denotes the number of indiv. arrived at time t

arrived=find(Pop(:,1)<=(t/tmax)); S1=length(arrived);

nof_arrived(t,g)=S1; % data collection, individuals at breeding ground after arrival

% number of eaten individuals of species 1

est_dead1=min(S1,((a*S1*X)/(1+(a*b*S1)+((1-a)*(b*z*S2))))); % unrounded # of indiv. to die

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prob_dead1=est_dead1-round(est_dead1); % difference between two integers becomes the prob.
if rand(1)<prob_dead1 % if random number is smaller than difference (probability)
    dead1=round(est_dead1)+1; % than it will become the larger integer
else
    dead1=round(est_dead1); % else the smaller integer
end

dead_arrived(t,g)=dead1; % data collection, S1 indiv. died per t from predation
doomed=picksurv(ones([1 S1]),dead1); % pick the indiv. that will be predated
Pop(arrived(doomed),:)=[]; % remove these individuals from population

% the 2nd species dies in a simpler manner as identities don't have to be tracked
dead2=min(S2,round((((1-a)*S2*X)/(1+(a*b*S1)+((1-a)*(b*z*S2))))));
S2=S2-dead2; % mortality updates the number of second species present
track_S2(t,g)=S2; % data collection, tracking S2

% finally, daily mortality of individuals that have not yet arrived on breeding grounds
if t<tmax % because everybody has arrived on day tmax
    notarrived_and_dead=find(Pop(:,1)>=(t/tmax) & rand([size(Pop,1) 1])<P0);
    dead_nonarrived(t,g)=length(notarrived_and_dead); % data collection
    Pop(notarrived_and_dead,:)=[]; % remove
end

if t==tmax % last time step t, fill remaining territories with floaters
    bG=alpha*Nterr-sum(Pop(:,2)>0); % good terr. available
    bP=(1-alpha)*Nterr-sum(Pop(:,2)<0); % poor terr. available
    fl_tmax=vectperm(find(Pop(:,2)==0)); % to create random selection of floaters to good and poor

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for k=1:length(fl_tmax)

    if bG>0 % if there are good territories available

        bG=bG-1; % one good territory less available

        Pop(fl_tmax(k),2)=1; % this ind is now the owner of a good terr

    elseif bP>0 %else if there are poor terr available

        bP=bP-1; %one less poor territory available

        Pop(fl_tmax(k),2)=-1; % this nFocal is now the owner of a poor terr

    end

end

end

end

% Post arrival:

% reproduction & mutation

allNew=[]; % vector to collect new indiv. in

for q=1:length(Pop(:,1))

    nOff=0; New=[];

    if Pop(q,2)>0 % individuals with a good breeding territory

        nOff=poissrnd(betaG); %number of offspring produced at a good site, average=betaG

    elseif Pop(q,2)<0 % individuals with a poor breeding territory

        nOff=poissrnd(betaP); %number of offspring produced at a poor site, average=betaP

    end

    if nOff>0

        New=max(0,min(1,((Pop(q,1)*ones([nOff 1]))+((rand(nOff,1)<mutProb).*(randn(nOff,1)*mutStep)))));
        % offspring inherit parent's arrival time & mutation may occur

        New(:,2)=zeros([nOff 1]); % offspring start as floater

        allNew=[allNew;New];

    end

end

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end

prodOff(g)=length(allNew);

Pop=[Pop;allNew]; % offspring join adult population

% mortality in the migration & wintering season

Pop(rand(size(Pop,1),1)<winterMort,:)=[]; %mortality during migration & overwintering

% get the distribution of arrival time allele in the last generation
if g==gen
    for p=1:tmax
        if p<tmax
            arrivDoc=find((Pop(:,1)>=(p-1)/tmax) & (Pop(:,1)<p/tmax));
        else
            arrivDoc=find((Pop(:,1)>=(p-1)/tmax) & (Pop(:,1)<(p+1)/tmax));
        end
        arrivHist(p)=length(arrivDoc);
    end
end

% the annual cycle ends by all individuals getting status NaN, i.e. they have not arrived yet
Pop(:,2)=NaN;

% save datafiles

save('meanArriv','meanArriv'); save('S1spring','S1spring');

save('nof_arrived','nof_arrived'); save('dead_arrived','dead_arrived');

save('dead_nonarrived','dead_nonarrived'); save('prodOff','prodOff');

save('track_S2','track_S2'); save('arrivHist','arrivHist');

end

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