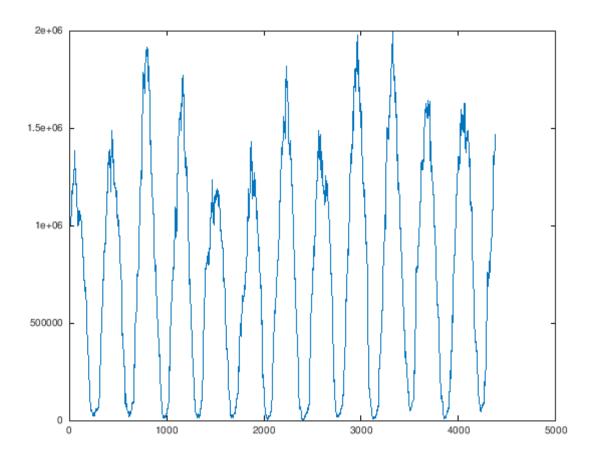
DynaLands1

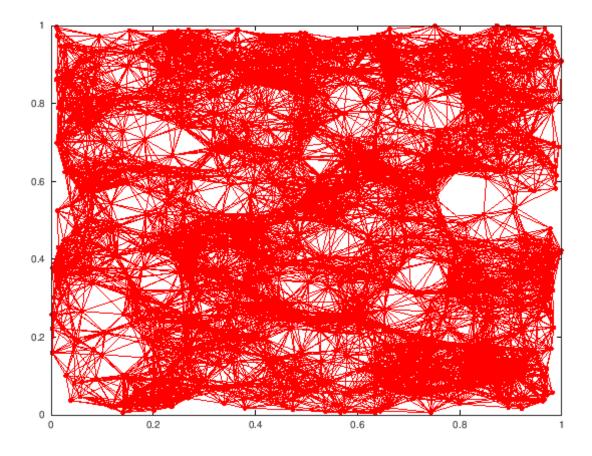
February 6, 2018

```
In [11]: %Importing icecover data to understand biodiversity dynamics --- data from http://nsi
         A = dlmread("masie_4km_allyears_extent_sqkm.csv",",");
In [12]: A(1:10,3)
ans =
  0.0000e+00
  0.0000e+00
   1.0697e+06
   1.0697e+06
   1.0697e+06
   1.0697e+06
   1.0697e+06
   1.0697e+06
   1.0697e+06
   1.0697e+06
In [13]: size(A)
ans =
  4377
            18
In [14]: plot(A(:,10))
```

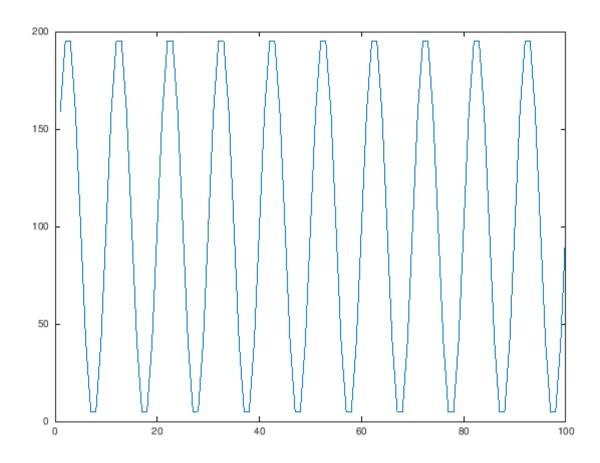


In [15]: "We will use the fluct in ice cover as a proxy of habitat and connectivity dynamics %RGG %Quick code Carlos J Melian %November 2013 J = 1000; r = 0.1; %r = unifrnd(0.01, 1);D = zeros(J,J);%Asymptotic behavior $mu = J*(e^(-pi * r^2 * J))$ MA = log(J) - log(mu);MB = pi*J;rc = sqrt(MA/MB); n = unifrnd(0,1,J,2);for i = 1:J-1;for j = i+1:J; $A = (n(i,1) - n(j,1))^2;$ "Euclidean distance" $B = (n(i,2) - n(j,2))^2;$ d(i,j) = sqrt(A + B);if d(i,j) < r;

D(i,j) = 1;



```
\%Be \ sure \ that \ the \ mij + lambda + nu == 1
n = unifrnd(0,1000,S,2); %sites!
R = ones(S,J);
countgen = 0;
Pairs = zeros(1,2);cevents = 0;
newSp = 1;
gamma=[];
A = 200;%amplitude, is the peak deviation:
   %350 to match simulations in random landscapes
   f = 0.1; %ordinary frequency, number of
    %cycles that occur each second of time
   sig = 0; %the phase
    countgen = countgen + 1;
   %r = A*sin(2*pi*f*countgen + sig) + A; %starting point with r approx.
   r(k,1) = A/2*(sin(2*pi*f*countgen + sig)+1);
end
v=1:100;
plot(v,r)
end
```



```
In [16]: %-----
         %General dynamic landscapes
         %Melian@KB May 2017
         %Palamara&Melian June 2017 version from scratch
         %Alex Rozenfeld June 2017
          show=true;
          showEach = 1;
          for ri = 1:50,
            S = 10; J = 100; \%S \text{ sites and } J \text{ inds. per site}
            %1. Implement a general case with zero-sum dynamics
             %combining static-dynamic vs. symmetric-asymmetric scenarios (non-stationary Gill
            %Be sure that the mij + lambda + nu == 1
            n = unifrnd(0,1000,S,2); %sites!
            R = ones(S,J);
            countgen = 0;
            Pairs = zeros(1,2);cevents = 0;
            newSp = 1;
            gamma=[];
            for k = 1:100, %Generations...
                A = 200;%amplitude, is the peak deviation:
                %350 to match simulations in random landscapes
                f = 0.1; %ordinary frequency, number of
                %cycles that occur each second of time
                sig = 0; %the phase
                countgen = countgen + 1;
                %r = A*sin(2*pi*f*countgen + sig) + A; %starting point with r approx.
                r = A/2*(sin(2*pi*f*countgen + sig)+1);
                 %2. Check sinusoidal with boundary conditions considering continuous A and f
                %Check\ r_min == 0\ and\ r_max == max\ distance\ ij
                D = zeros(S,S); %theshold matrix
                Di = zeros(S,S); %distance matrix
                mu = S*(exp((-pi * (r/1000)^2 * S))); %site connectivity
                for i = 1:S-1,
                     for j = i+1:S,
                        A = (n(i,1) - n(j,1))^2; \text{``Euclidean distance'}
                        B = (n(i,2) - n(j,2))^2;
                        d(i,j) = sqrt(A + B);
                        Di(i,j) = 1/d(i,j);
```

```
%3. This is the simplest kernel
        %Explore the asymmetry under 1/d(i,j)
        %Do we need to implement more asymmetric situations, like 1/(d(i,j) \hat{x}
        if d(i,j) < r; %threshold</pre>
          D(i,j) = 1;
        else
          D(i,j) = 0;
        end
   end
end
%DI=Di+Di';Dc=cumsum(DI,2);D1=D+D';
DI=Di+Di';
D1=D+D';
DI=DI.*D1; %<=====ALEX
Dc=cumsum(DI,2);
m = unifrnd(0.001,0.1,1); %migraion from the blocks
v = unifrnd(0.0001, 0.01, 1); % regional migration?
1=1-(m+v);
KillHab = unidrnd(S);
    KillInd = unidrnd(J);
    ep=unifrnd(0,1,1); %event probability
    if ep < m, %Migration</pre>
      MigrantHabProb = unifrnd(0,max(Dc(KillHab,:)));
      MigrantHab = find(Dc(KillHab,:) >= MigrantHabProb);
    %pause
      if D1(KillHab,MigrantHab) == 1;
       %4. Implement local birth dynamics and speciation dynamics
       MigrantInd = unidrnd(J);
       cevents = cevents + 1;
       Pairs(cevents,1) = KillHab;
       Pairs(cevents,2) = MigrantHab(1,1);
       R(KillHab,KillInd)=R(MigrantHab(1,1),MigrantInd);
      end
    elseif ep <= m+v, %mutation</pre>
       newSp = newSp +1;
       R(KillHab,KillInd) = newSp;
    else
                       %birth
       BirthLocalInd = unidrnd(J);
       while BirthLocalInd == KillInd,
         BirthLocalInd = unidrnd(J);
       end
```

```
R(KillHab,KillInd) = R(KillHab,BirthLocalInd);
             end
        end
        %Species at each site:
        Sp_eachSt=arrayfun(@(ix) unique(R(ix,:)), [1:size(R,1)], 'uniformoutput', false
        %alpha(g) Num of species at each site for present generation
        alpha = arrayfun(@(v) length(cell2mat(v)),Sp_eachSt);
        gamma(countgen) = numel(unique(R));
        Sim=CalcSim(Sp_eachSt,S);
        if show && (k==1 || mod(k,showEach)==0), %Show results
          ShowResults(ri,countgen,S,n,D1,d,alpha,gamma,Sim)
        end
    end
  end
end
function Sim = CalcSim(Sp_eachSt,S)
  Sim = zeros(S,S);
  for i = 1:S-1,
     for j = i+1:S,
        %CantSpEnComun_ij = \#(Sp_i \ n \ Sp_j)
        %Similaridad_ij = CantSpEnComun_ij / (\#Sp_i + \#Sp_j - CantSpEnComun_ij)
        CantSpEnComun_ij = length(intersect(Sp_eachSt{i},Sp_eachSt{j}));
        Sim(i,j) = CantSpEnComun_ij / (length(Sp_eachSt{i})+length(Sp_eachSt{j})-CantSpEnComun_ij / (length(Sp_eachSt{i})+length(Sp_eachSt{j}))
     end
  end
  Sim = Sim + Sim' + eye(S,S);
end
function ShowResults(ri,countgen,S,n,D1,d,alpha,gamma,Sim)
          sizeFactor=10;
          figure(ri)
          subplot(2,3,[1 2 4 5]) %alpha
          hold off
          for i=1:S,
             scatter(n(i,1),n(i,2),sizeFactor*alpha(i),'b','filled') %Sites
            hold on;
             hola=1;
             ixStConnected=find(D1(i,:));
```

```
for ix=ixStConnected,
   line([n(i,1) n(ix,1)]',[n(i,2) n(ix,2)]','color','r') %Links
  end
end
xlim([0 1000])
ylim([0 1000])
text(1,1050,['Realization: ' num2str(ri) ' Gen: ' num2str(countgen)])
subplot(2,3,3) %gamma
plot(gamma);
%hold on
%scatter(countgen, gamma(countgen), 5, 'k')
subplot(2,3,6) %Sim VS d (connected and non connected)
hold off
for i = 1:S-1,
   for j = i+1:S,
      scatter(d(i,j),Sim(i,j),5,'k')
     hold on
   end
end
pause(0.001);
```

