

# Animation ice cover

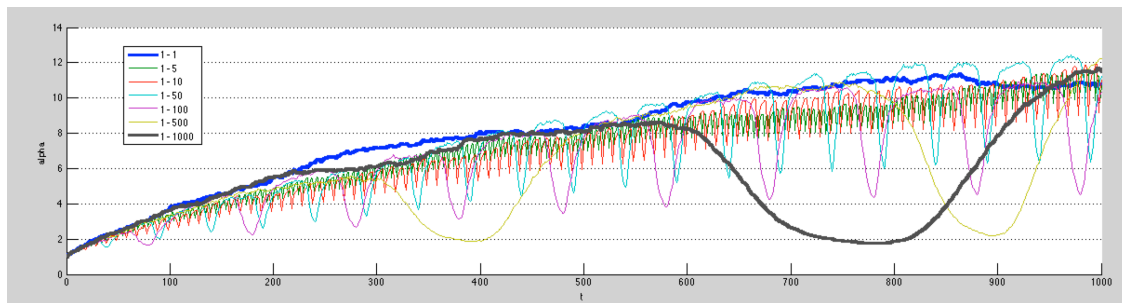
February 6, 2018

```
In [1]: from IPython.display import HTML
        HTML('<iframe width=\"1280\" height=\"720\" src=\"https://www.youtube.com/embed/VIXciS
```

```
Out[1]: <IPython.core.display.HTML object>
```

```
In [2]: from IPython.display import Image
        Image("circadianclock?.png")
```

Out[2]:



# 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 =
```

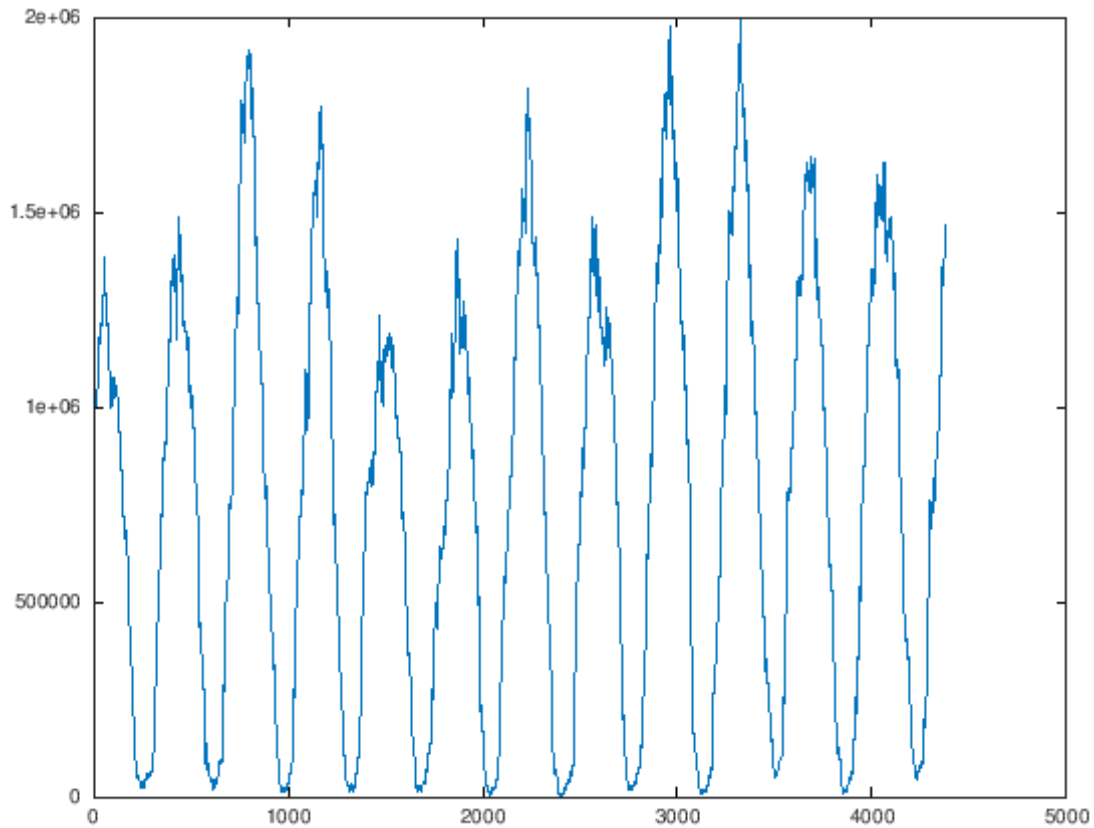
```
0.0000e+00
0.0000e+00
1.0697e+06
1.0697e+06
1.0697e+06
1.0697e+06
1.0697e+06
1.0697e+06
1.0697e+06
1.0697e+06
```

```
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;
                 end
             end
         end
```

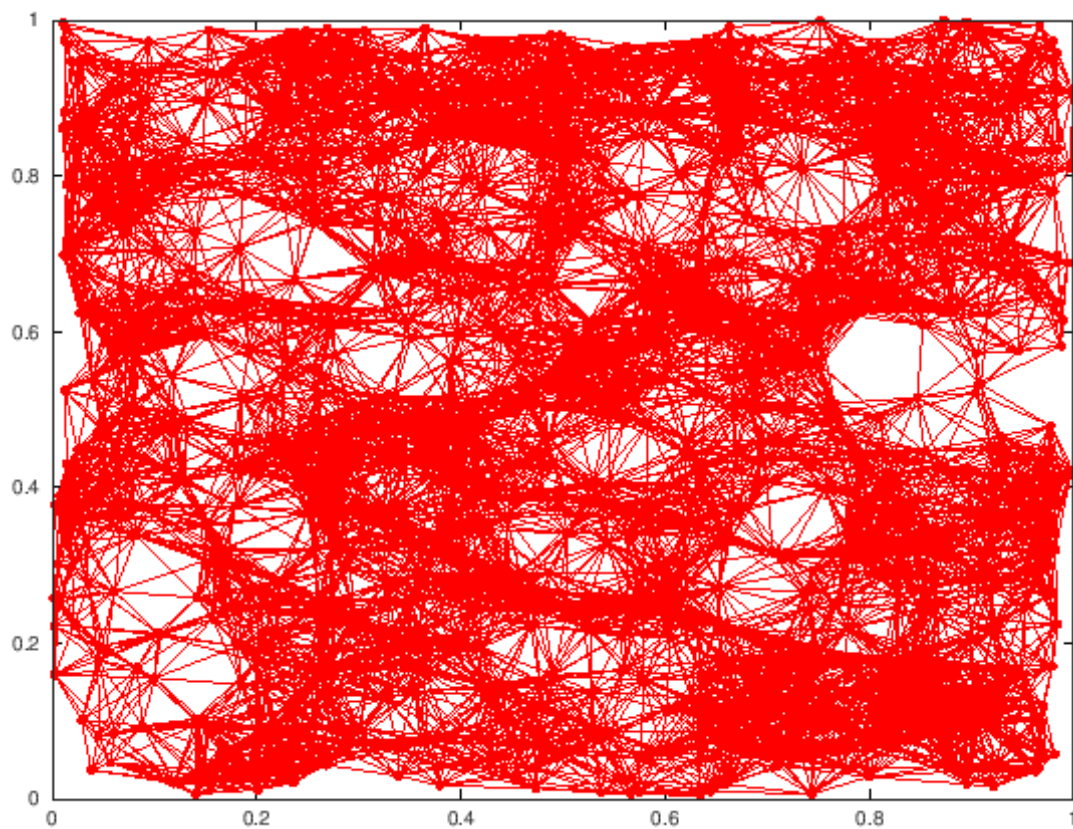
```

        else
            D(i,j) = 0;
        end
    end
end
end
D1=D+D';

%plot network
gplot(D1,n, "r.-")
set (get (gca, ("children")), "markersize", 12);

mu = 2.2711e-11

```



```

In [10]: show=true;
        showEach = 1;
        for ri = 1:1,
            S = 10; J = 100; %S sites and J inds. per site

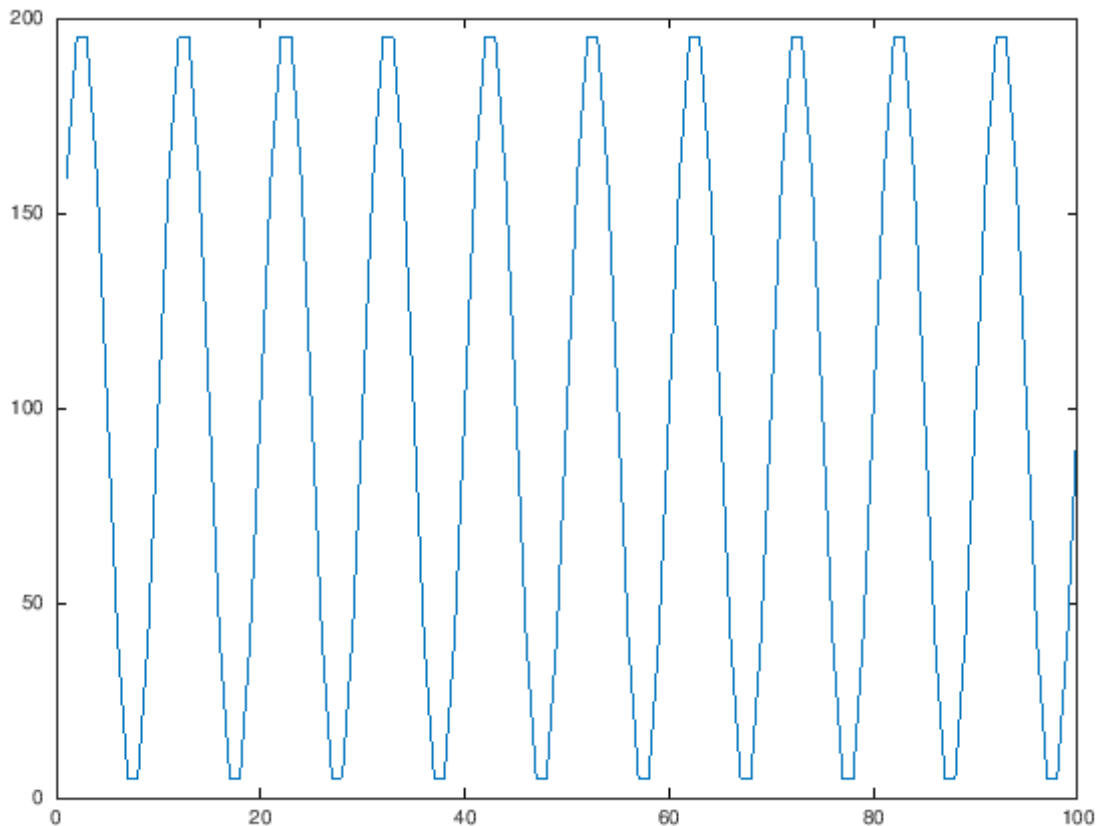
            %1. Implement a general case with zero-sum dynamics
            %combining static-dynamic vs. symmetric-asymmetric scenarios (non-stationary Gill

```

```

%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(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 sites and J 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; %Euclidean distance
                B = (n(i,2) - n(j,2))^2;
                d(i,j) = sqrt(A + B);
                Di(i,j) = 1/d(i,j);
            end
        end
    end
end

```

```

%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)^x

    if d(i,j) < r;%threshold
        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?
l=1-(m+v);

for j = 1:J*S, %MonteCarlo Time
    KillHab = unidrnd(S);
    KillInd = unidrnd(J);
    ep=unifrnd(0,1,1); %event probability
    if ep < m, %Migration
        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
        newSp = newSp +1;
        R(KillHab,KillInd) = newSp;
    else %birth
        BirthLocalInd = unidrnd(J);
        while BirthLocalInd == KillInd,
            BirthLocalInd = unidrnd(J);
        end
    end
end

```

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        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);
        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,:));
    end
end

```



```

    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
end
pause(0.001);

```

