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% -----
% Eulerian phytoplankton water column model
% (based on Huisman et al.(2006))
% -----
%Programmed by C Lindemann & DL Aksnes (02102018)

clear all
close all

%% Parameter

% Temporal and spatial discretization
nZ=100;           % Number of depth cells
DZ=1;            % Length of a depth cell (m)
nT=10;           % Number of time steps
DT=60*10;        % Length of a time step(s)'
t= (0:nT)/(60*60*24/DT); % for plotting

%Water column properties
kappa=10/(60*60*24) ; % Turbulent diffusivity (m2 s-1)
I0=100;           % Irradiance at surface (mmol photons m-2 s-1)
Ksw=0.15;         % Background light attenuation (m-1)
NB=5;             % Fixed nutrient concentration in last depth cell (mmol N m-3)

%Phytoplankton traits
um=1/(60*60*24); % Max growth rate (s-1)
m=0.25/(60*60*24); % loss (death)rate (s-1)
Hi=20;           % Half saturation irradiance (micromol photons m-2 s-1)
Hn=0.025;        % Half saturation nutrient (mmol m-3)
v=1/(60*60*24); % Sinking velocity of phytoplankton (m s-1)
alpha=0.5;        % Nutrient recycling coefficient

%% testing variable sensitivity analysis

varKsw=[0,0,0,0,0];

Nall=zeros(length(varKsw),nZ,nT+1);
Pall=zeros(length(varKsw),nZ,nT+1);

for Ktest=1:length(varKsw)

    Ksw=varKsw(Ktest);

%% Initial conditions

% define initial conditions
phy1=linspace(0.3,0.05,nZ);
nit1=linspace(0.1,NB,nZ);

% create storage structures
P=zeros(nZ,nT+1);
N=zeros(nZ,nT+1);

% save initial conditions to storage
P(:,1)=phy1;
N(:,1)=nit1;

%% Discretisation & Constant rates

umax = um*DT;           % constant maximum growth rate
Mort=m*DT;             % constant Mortality
Diff=kappa*DT/(DZ*DZ); % the fraction of a depth cell that is exchanged
                        % with the neighbour cells due to turbulent diffusivity

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Sink=v*DT/DZ;          % The fraction of phytoplankton in a depth cell that
sinks                  % to the cell below during DT

% time-independent light in the water column
z = 0.5*DZ:DZ:DZ*nZ-0.5*DZ; % depth array
Iz=I0*exp(-Ksw.*z);      % water column irradiance
Ilim=Iz./(Iz+Hi);       % Light limitation

%% Time loop
for tstep=1:nT

    %Phytoplankton growth
    Nlim=nit1./(nit1+Hn); %nutrient limitation of depth cell
    u=umax*min(Ilim,Nlim); %phytoplankton growth rate

    %Update first (surface) depth cell - no exchange through surface
    nit2(1)=phy1(1)*(-u(1)+Mort*alpha) + nit1(1)*(1-Diff)
+nit1(2)*Diff;
    phy2(1)=phy1(1)*(u(1)-Mort) -phy1(1)*Sink + phy1(1)*(1-Diff)
+phy1(2)*Diff;

    %Update last depth cell. Nutrient concentration is fixed in last cell!
    %Phytoplankton is not transported (by diffusion/sinking) below the last
depth cell.
    nit2(nZ)=NB;
    phy2(nZ)=phy1(end)*(u(end)-Mort) +phy1(end-1)*Sink + phy1(end)*(1-Diff)
+phy1(end-1)*(Diff);

    % Update all the intermediate depth cells
    for n=2:nZ-1
        nit2(n)= phy1(n)*(-u(n)+Mort*alpha) +
nit1(n)*(1-Diff-Diff) +nit1(n-1)*Diff +nit1(n+1)*Diff;
        phy2(n)= phy1(n)*(u(n)-Mort) + phy1(n-1)*Sink - phy1(n)*Sink +
phy1(n)*(1-Diff-Diff) +phy1(n-1)*Diff +phy1(n+1)*Diff;
    end

    %Store results
    N(:,tstep+1)=nit2(:);
    P(:,tstep+1)=phy2(:);

    %Proceed to next time step with updated state variables
    nit1=nit2;
    phy1=phy2;
end

Nall(Ktest, :, :)=N;
Pall(Ktest, :, :)=P;

%% plot output

figure;
subplot(2,1,1)
contourf(t,z,N);
title(['Nutrient concentration (mmol m-3) for K=' num2str(Ksw)]);
xlabel('Time (days)');
ylabel('Depth (m)');
colorbar
set(gca, 'YDir', 'reverse');

subplot(2,1,2)
contourf(t,z,P);
title(['Phytoplankton concentration (mmol m-3) for K=' num2str(Ksw)]);

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xlabel('Time (days)');  
ylabel('Depth (m)');  
colorbar  
set(gca, 'YDir', 'reverse');
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%clear N P  
end
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Nfinal=squeeze(Nall(:, :, end));  
Pfinal=squeeze(Pall(:, :, end));
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