

## APPENDIX I: Used MatLab code

The same code was used for three groups of species, with exception of the species name which was named in “yNAME”.

```
yName = 'DIATOMO';
%yName = 'FILAMENT';
%yName = 'MACROVEG';

% fish species end variables are
%   GACMAX40   GACMIN40   PUNG   WHIBIN   VENBIN

% download data
% -----
data = importdata('data/majorlarvae_4matlab.txt');
rasterCovarNames = {'SHOPROFILE' 'DEPTH' 'BOTTOM' 'BOTTOMCOV'
'FETCH300W' 'FE300ME' ...
'DIST20M' 'LINED' 'ISLANDNUM' 'DIS_SAND' 'DIS_SHALLO' 'PE900'
'PE3000' 'SAL09SPRS' ... %'SAL10SPRS'
'SAL910WIN' 'ICELAST09' 'ICEWIN09' 'EUTSTAT' ... %'ICELAST10'
'ICEFIRST09' 'ICEFIRST10' 'ICEWIN10'
'EKOSTAT' 'PHOSP' 'NITROG' 'CHL_A' 'SECCHI' 'RIVERS' 'BOTTOMCLS'
'SHAREA'};
spatNames = {'N_etr89' 'E_etr89'};

% --- TAKE x -----
% Raster variables
indRaster = false(1,size(data.data,2));
for il=1:length(rasterCovarNames)
    indRaster(strcmp(data.textdata,rasterCovarNames{il})) = true;
end
% Spatial coordinates
indSpat = false(1,size(data.data,2));
for il=1:length(spatNames)
    indSpat(strcmp(data.textdata,spatNames{il})) = true;
end
% Take environmental variables
% order [Raster Spatial ]
x = [data.data(:,indRaster) data.data(:,indSpat)];
xlabel = [data.textdata(indRaster) data.textdata(indSpat)];

% rescale the covariates
standind = 1:size(x,2)-2; % columns of x to be standardized (all
other except co-ordinates)
mx = mean(x(:,standind)); stdx = std(x(:,standind));
x(:,standind) = (x(:,standind)-
repmat(mx,size(x,1),1))./repmat(stdx,size(x,1),1);
% scale co-ordinates to kilometers
x(:,end-1:end) = x(:,end-1:end)./1000;

% ----- TAKE y and Z -----
y = data.data(:, strcmp(data.textdata,yName)); % the number of
count locations where species is present
z = data.data(:, strcmp(data.textdata,'TOTAL_ED')); % the total
number of count locations

% ----- define model and inference method-----
lik = lik_binomial; % probit likelihood
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cfc = gpcf_constant('constSigma2', 10, 'constSigma2_prior',
prior_fixed); % The covariance function of the intercept term
cfs = gpcf_exp('magnSigma2', 2, 'lengthScale', [3 6],
'selectedVariables', size(x,2)-[1 0] ,...
'lengthScale_prior', prior_t('s2',100));
% The covariance function of the spatial component
cf_nn = gpcf_neuralnetwork('selectedVariables', 1:size(x,2)-2, ...
% The neural network covariance function for
'weightSigma2', 10*ones(1,size(x,2)-2));
% the nonlinear effect of covariates
%%cf_se = gpcf_sexp('selectedVariables', 1:size(x,2)-2, ...
% The squared exponential covariance function for
%% 'lengthScale', ones(1,size(x,2)-2));
% the nonlinear effect of covariates
cf_li = gpcf_linear('selectedVariables', 1:size(x,2)-2);
% covariance function corresponding to linear model of covariates

% use neural network covariance, that is non-linear response of
covariates
gp = gp_set('lik', lik, 'cf', {cfc cf_nn cfs}, 'latent_method', 'EP');
% use squared exponential covariance, that is non-linear response of
covariates
%gp = gp_set('lik', lik, 'cf', {cfc cf_se cfs}, 'latent_method',
'EP');
% use linear covariance function, that is linear response of
covariates
%gp = gp_set('lik', lik, 'cf', {cfc cf_li cfs}, 'latent_method',
'EP');

% Infer the model
% optimize the covariance function parameters and find Gaussian
% approximation (EP) for the latent variables
opt=optimset('TolFun',1e-3,'TolX',1e-3,'Display','iter'); % Options
for optimization
gp=gp_optim(gp,x,y, 'z', z,'opt',opt); % run optimization

%%
apcs = gp_avpredcomp(gp,x,y,'z',z)

figure; hold on
for i1=1:size(apcs.fs,2)
prc = prctile(apcs.fsa(:,i1),[5 50 95]);
plot([prc(1) prc(3)], [i1 i1])
plot(prc(2),i1, 'kx', 'linewidth', 5)
end
plot([0 0],[0 25])
set(gca,'yTick',1:28,'yticklabel',xlabels)
%% Analyze results

% in case of linear model you can calculate the posterior of beta
% parameters
if strcmp(gp.cf{2}, 'gpcf_linear')
Efl =[]; Varfl = [];
fprintf(' The posterior mean and 95% credible intervals of
betas:\n')
for i1 = 1:size(x,2)-2
xtmp = zeros(1,size(x,2)); xtmp(i1)=1;
[Eft, Vart] = gp_pred(gp,x,y, 'z',z, xtmp, 'predcf', 2);
Efl(i1) = Eft; Varfl(i1) = Vart;

```

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        fprintf('    %s: \t %.2f (%.2f, %.2f) \n', xlabel{il}, Eft,
Eft-2*sqrt(Vart), Eft+2*sqrt(Vart))
    end
end
figure, hold on
set(gcf, 'units', 'centimeters');
set(gcf, 'defaultaxesfontsize', 10)    %6 8
set(gcf, 'defaulttextfontsize', 10)    %6 8
set(gcf, 'defaultlinelength', 1.5)

% ind defines the locations at which we look at the response curve
ind = resampdet(ones(size(x,1),1),100,1);
npoints=20;
for il=1:size(x,2)-2
    x_il = [linspace(min(x(:,il)),max(x(:,il)),npoints)' ; 0];
    xtemp = x(ind,:);
    xtemp = reshape( repmat(xtemp(:)',npoints+1,1),
(npoints+1)*length(ind),size(x,2));
    xtemp(:,il) = repmat(x_il,length(ind),1);
    xtemp(npoints+1:npoints+1:size(xtemp,1),il) = x(ind,il);
    %xtemp(npoints+1:npoints+1:size(xtemp,1),il) =
x_il(floor(npoints/2));
    [Ef,Varf] = gp_jpred(gp,x,y,xtemp, 'z',z,'predcf',
2:length(gp.cf)-1);
    Eformat = reshape(Ef,npoints+1,length(ind));
    %
    Eformat = Eformat(1:npoints,:) - repmat(
Eformat(npoints+1,:), npoints,1);
    %
    Eformat = Eformat - repmat( mean(Eformat,1), npoints,1);
    Eformat = Eformat(1:npoints,:);
    Eformat = Eformat - repmat( mean(Eformat,1), npoints,1);
    EF = mean(Eformat,2);
    Varf2=[];
    for v1=1:length(ind)
        vind = (1:npoints+1) + (v1-1)*(npoints+1);
        vartemp = Varf(vind,vind);
        %Varf2(:,v1) = diag(vartemp) + vartemp(end,end) -
2*vartemp(end,:);
        wa=ones(length(x_il),1)./npoints;wa(end)=0;
        Varf2(:,v1) = diag(vartemp(1:end,1:end)) + wa'*vartemp*wa -
2*vartemp(1:end,:)*wa;
    end
    Varf2=Varf2(1:end-1,:);
    VARF = mean(Varf2,2) + var(Eformat,[],2);
    x_il=x_il(1:end-1);

    %subplot(3,3,il), hold on
    sp(il) = subplot(ceil(sqrt(size(x,2)-2)),ceil(sqrt(size(x,2)-
2)),il); hold on
    plot(repmat(x_il*stdx(il)+mx(il),1,length(ind)), Eformat, 'k',
'color', [0.7 0.7 0.7], 'linewidth', 0.5), hold on
    plot(x_il*stdx(il)+mx(il), EF, 'k')
    plot(x_il*stdx(il)+mx(il), EF-2*sqrt(VARF), 'k--'), hold on
    plot(x_il*stdx(il)+mx(il), EF+2*sqrt(VARF), 'k--'), hold on
    xlabel(xlabel{il})
    xlim([min(x_il*stdx(il)+mx(il)) max(x_il*stdx(il)+mx(il))])
    ylim([min(EF-2*sqrt(VARF)) max(EF+2*sqrt(VARF))])
    grid on
    %
    % define the prediction covariates (xtemp) so that one
covariate (il)
%
% changes at ind locations and all other covariates are kept
constant.

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%     x_il = linspace(min(x(:,il)),max(x(:,il)),npoints)';
%     xtemp = x(ind,:);
%     xtemp = reshape( repmat(xtemp(:)',npoints,1),
npoints*length(ind),size(x,2));
%     xtemp(:,il) = repmat(x_il,length(ind),1);
%
%     % Predict the response along the covariate il
%     [ef,varf] = gp_pred(gp,x,y,xtemp, 'predcf', 2:length(gp.cf)-1);
%
%     % Reshape ef and subtract the mean at each prediction point
%     efmat = reshape(ef,npoints,length(ind));
%     efmat = efmat - repmat( mean(efmat,1), npoints,1);
%     ef = mean(efmat,2);
%     varf = mean(reshape(varf,npoints,length(ind)),2) +
var(efmat,[],2);
%     %varf = var(efmat,[],2);
%
%     %subplot(3,3,il), hold on
%     subplot(ceil(sqrt(size(x,2)-2)),ceil(sqrt(size(x,2)-2)),il),
hold on
%     plot(repmat(x_il*stdx(il)+mx(il),1,length(ind)), efmat, 'k',
'color',[0.7 0.7 0.7], 'linewidth', 0.5), hold on
%     plot(x_il*stdx(il)+mx(il), ef, 'k')
%     plot(x_il*stdx(il)+mx(il), ef-2*sqrt(varf), 'k--'), hold on
%     plot(x_il*stdx(il)+mx(il), ef+2*sqrt(varf), 'k--'), hold on
%     xlabel(xlabels{il})
%     xlim([min(x_il*stdx(il)+mx(il)) max(x_il*stdx(il)+mx(il))])
%     ylim([min(ef-2*sqrt(varf)) max(ef+2*sqrt(varf))])
%     grid on
%     if il == 1
%         text(1.5*max(x_il*stdx(il)+mx(il)),1.2*max(ef+2*sqrt(varf)),
['The species modeled:' yName])
%     end
end
ylimits = [];
for il=1:length(sp)
    ylimits(il,:) = get(sp(il),'ylim');
end
for il=1:length(sp)
    set(sp(il),'ylim', [min(ylimits(:,1)) max(ylimits(:,2))]);
end
subplot(ceil(sqrt(size(x,2)-2)),ceil(sqrt(size(x,2)-2)),2)
text(min(x(:,2)*stdx(2)+mx(2)),1.5*max(ylimits(:,2)), ['The species
modeled:' yName])

set(gcf, 'pos', [0 0 25 25])
set(gcf, 'paperposition', get(gcf,'pos'))

%% Train model without in-situ variables
% Define the name of end variable (Species name)
% Most interesting options are:
%     FILAMENT%, DIATOMO%, MACROVEG (charop, phaeoph)
yName = 'DIATOMO';
%yName = 'FILAMENT'

% fish species end variables are
%     GACMAX40     GACMIN40     PUNG     WHIBIN     VENBIN

```

```

% download data
% -----
data = importdata('data/majorlarvae_4matlab.txt');
rasterCovarNames = {'FETCH300W' 'FE300ME' 'DIST20M' 'LINED'
'ISLANDNUM' 'DIS_SAND' ...
'DIS_SHALLO' 'PE900' 'PE3000' 'SAL09SPRS' 'SAL910WIN' 'ICELAST09'
'ICEWIN09' 'EUTSTAT' ...
'EKOSTAT' 'PHOSP' 'NITROG' 'CHL_A' 'SECCHI' 'RIVERS' 'BOTTOMCLS'
'SHAREA'};
spatNames = {'N_etr89' 'E_etr89'};

% --- TAKE x -----
% Raster variables
indRaster = false(1,size(data.data,2));
for i1=1:length(rasterCovarNames)
    indRaster(strcmp(data.textdata,rasterCovarNames{i1})) = true;
end
% Spatial coordinates
indSpat = false(1,size(data.data,2));
for i1=1:length(spatNames)
    indSpat(strcmp(data.textdata,spatNames{i1})) = true;
end
% Take environmental variables
% order [Raster Spatial ]
x = [data.data(:,indRaster) data.data(:,indSpat)];
xlabel = [data.textdata(indRaster) data.textdata(indSpat)];

% rescale the covariates
standind = 1:size(x,2)-2; % columns of x to be standardized (all
other except co-ordinates)
mx = mean(x(:,standind)); stdx = std(x(:,standind));
x(:,standind) = (x(:,standind)-
repmat(mx,size(x,1),1))./repmat(stdx,size(x,1),1);
% scale co-ordinates to kilometers
x(:,end-1:end) = x(:,end-1:end)./1000;

% ----- TAKE y -----
% Try different tresholds (e.g. under 5%, 10% coverage is zero)
y = data.data(:, strcmp(data.textdata,yName));
y = 2*(y>0)-1; % -1 = absence, 1 = presence

% ----- define model and inference method-----
lik = lik_probit; % probit likelihood
cfc = gpcf_constant('constSigma2', 10, 'constSigma2_prior',
prior_fixed); % The intercept term
cfs = gpcf_exp('magnSigma2', 2, 'lengthScale', [3 6],
'selectedVariables', size(x,2)-[1 0] ,...
'lengthScale_prior', prior_t('s2',100));
% The spatial covariance function
cf_nn = gpcf_neuralnetwork('selectedVariables', 1:size(x,2)-2, ...
% The nonlinear effect of covariates
'weightSigma2', 10*ones(1,size(x,2)-2));
gp = gp_set('lik', lik, 'cf', {cfc cf_nn cfs}, 'latent_method', 'EP');

% Infer the model
% optimize the covariance function parameters and find Gaussian
% approximation (EP) for the latent variables
opt=optimset('TolFun',1e-3,'TolX',1e-3,'Display','iter'); % Options
for optimization
gp=gp_optim(gp,x,y,'z',z,'opt',opt); % run optimization

```

```

%% Do the prediction
data_pred = importdata('data/G300_mod.txt');

% Rename few columns in the prediction data so that column names match
with
% training data column names
data_pred.textdata{strcmp(data_pred.textdata, 'FE300W')} = 'FETCH300W';
data_pred.textdata{strcmp(data_pred.textdata, 'SAL09SPR')} =
'SAL09SPRS';
data_pred.textdata{strcmp(data_pred.textdata, 'X')} = 'E_etr89';
data_pred.textdata{strcmp(data_pred.textdata, 'Y')} = 'N_etr89';
% pick up the right columns in right order from the prediction data
for il=1:length(xlabels)
    otaInd(il) = find(strcmp(data_pred.textdata, xlabels{il}));
end
% create the prediction inputs
x_pred = data_pred.data(:, otaInd);
% rescale the covariates
standind = 1:size(x, 2)-2; % columns of x to be standardized (all
other except co-ordinates)
x_pred(:, standind) = (x_pred(:, standind) -
repmat(mx, size(x_pred, 1), 1)) ./ repmat(stdx, size(x_pred, 1), 1);
% scale co-ordinates to kilometers
x_pred(:, end-1:end) = x_pred(:, end-1:end) ./ 1000;
[Ef, Varf, lpy] =
gp_pred(gp, x, y, x_pred, 'z', z, 'yt', ones(size(x_pred, 1), 1), 'zt', ones(size
(x_pred, 1), 1));

% Write the prediction into the file
pred = [x_pred(:, end) .* 1000 x_pred(:, end-1) .* 1000 exp(lpy)]; % scale
to intensity per 100m^3

fid = fopen([yName '_prediction.txt'], 'w');
fprintf(fid, 'X\tY\tZ\n');
for tmpi = 1:size(pred, 1)
    fprintf(fid, '%f\t%f\t%f\n',
pred(tmpi, 1), pred(tmpi, 2), pred(tmpi, 3));
end
fclose(fid);
%save([yName '_prediction.txt'], 'pred', '-ascii', '-double', '-tabs')

```

### APPENDIX I: Response curves for all the variables

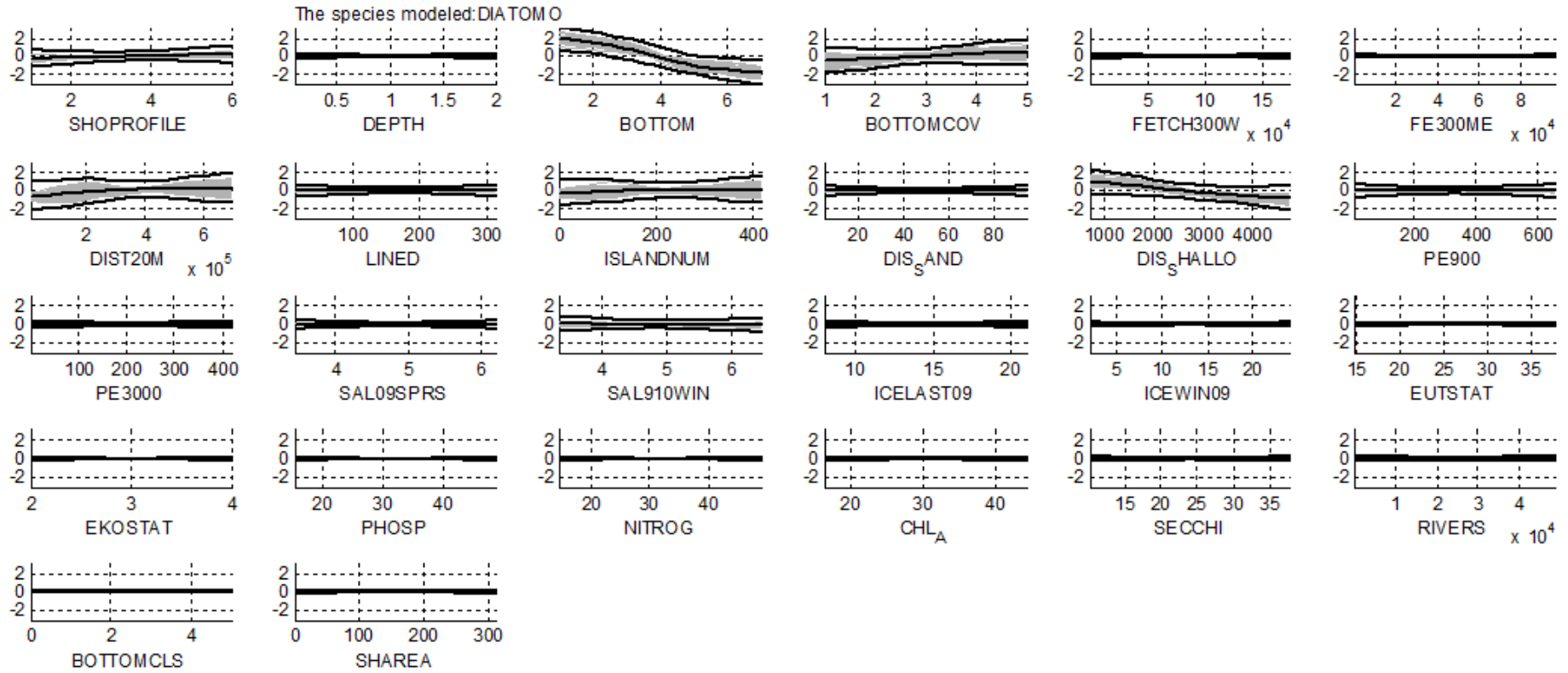


Figure 10: DIATOM response curves of all the variables.

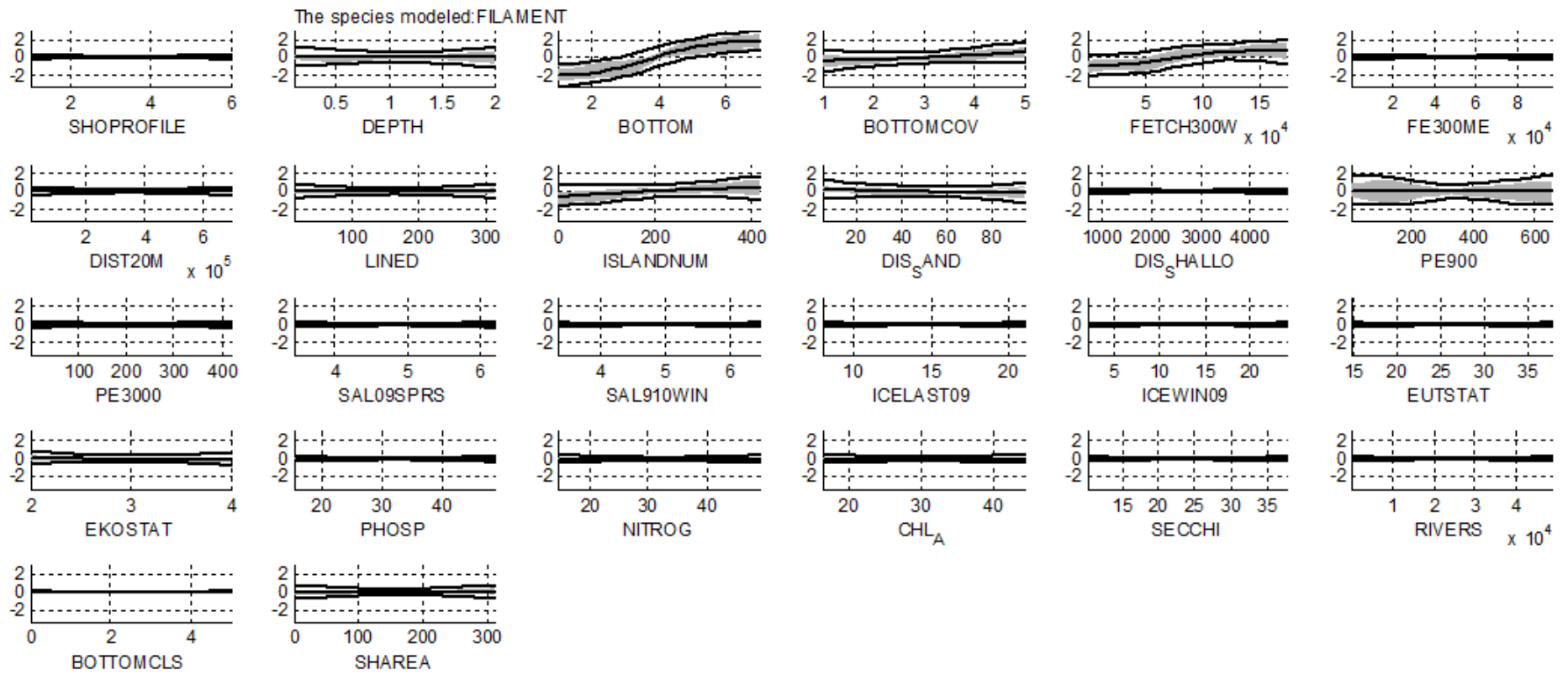


Figure 11: FILAMENT response curves of all the variables.



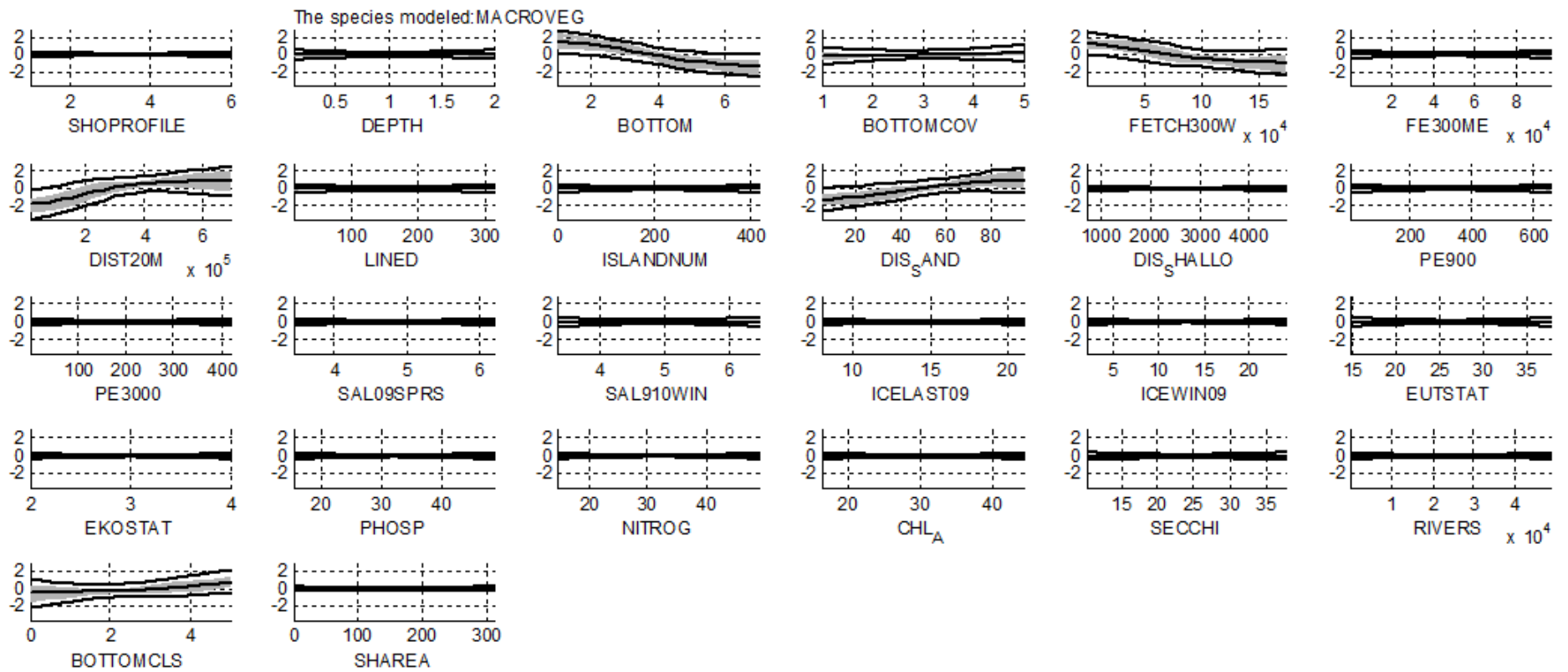


Figure 12: MACROVEG response curves of all the variables.