```
function [reduced section] = FOCuS Section(model in, biom, rxn max in, bound glu, ✓
bound 02, bound ATPM, bound biom)
COBRA stoichiometric model in standard structure format
% model in
% biom
              Biomass reaction
              Target reaction
% rxn max in
              input flux value for glucose
% bound glu
               input flux value for oxygen
% bound 02
              input flux value for maintenance ATP
% bound ATPM
clc
tic
global model
global rxn_max
model = model in;
rxn max = rxn max in;
para=[100 0.8];
n=para(1);
p=para(2);
m=14;
pop=n;
gen=0;
max gen=1;
global d
d=5; %Number of genes to be deleted
rxns size = size(model.rxns);
rxns count = rxns size(1);
for i=1:1:rxns count
   rxn ind = strcmp(model.rxns(i),rxn max);
   if rxn ind > 0
      rxn max no = i;
   end
end
clear i
disp('Target reaction number =')
disp(rxn max no);
for i=1:1:rxns count
```

```
biomass ind = strcmp(model.rxns(i),biom);
   if biomass ind > 0
      biomass max no = i;
disp('Biomass reaction number =')
disp(biomass_max_no);
ind=xlsread('C:\Users\....\MATLAB\cobra\test codes\FOCUS\acetate index.xlsx')
응응응
global ind_cont
global bound cont
global pind
ind cont = 1:length(ind);
section=0;
sectionmax= 30 %10 %20; %15
ro=round((length(ind)/sectionmax));
half=[];
while (section<sectionmax)</pre>
   if section==0
      ind_cont1= 1:(ro*1);
      bound cont=ind cont1;
      Lb = min(ind_cont1)*ones(1,d);
      Ub = max(ind cont1)*ones(1,d);
   end
   if section==1
      ind cont1= (ro+1):(ro*2);
      bound cont=ind cont1;
      Lb = min(ind_cont1)*ones(1,d);
      Ub = max(ind_cont1) *ones(1,d);
   end
   if section==2
      ind cont1= ((ro*2)+1):(ro*3);
      bound_cont=ind_cont1;
      Lb = min(ind_cont1) *ones(1,d);
```

```
Ub = max(ind cont1)*ones(1,d);
end
if section==3
    ind cont1= ((ro*3)+1):(ro*4);
    bound_cont=ind_cont1;
    Lb = min(ind_cont1)*ones(1,d);
    Ub = max(ind cont1)*ones(1,d);
end
if section==4
    ind_cont1= ((ro*4)+1):(ro*5); %length(ind);%
    bound_cont=ind_cont1;
    Lb = min(ind cont1)*ones(1,d);
    Ub = max(ind_cont1)*ones(1,d);
end
if section==5
    ind_cont1 = ((ro*5)+1):(ro*6);
    bound_cont=ind_cont1;
    Lb = min(ind cont1)*ones(1,d);
    Ub = max(ind_cont1)*ones(1,d);
end
if section==6
    ind cont1= ((ro*6)+1):(ro*7);
    bound cont=ind cont1;
    Lb = min(ind cont1)*ones(1,d);
    Ub = max(ind cont1)*ones(1,d);
end
if section==7
    ind cont1= ((ro*7)+1):(ro*8);
    bound cont=ind cont1;
    Lb = min(ind cont1)*ones(1,d);
    Ub = max(ind cont1)*ones(1,d);
end
if section==8
    ind cont1= ((ro*8)+1):(ro*9);
    bound cont=ind cont1;
    Lb = min(ind cont1)*ones(1,d);
    Ub = max(ind_cont1) *ones(1,d);
end
if section==9
    ind cont1= ((ro*9)+1):(ro*10); %length(ind);%
    bound_cont=ind_cont1;
    Lb = min(ind_cont1) *ones(1,d);
    Ub = max(ind cont1)*ones(1,d);
end
if section==10
    ind_cont1 = ((ro*10)+1):(ro*11);
    bound_cont=ind_cont1;
```

```
Lb = min(ind cont1)*ones(1,d);
    Ub = max(ind cont1)*ones(1,d);
end
if section==11
    ind_cont1= ((ro*11)+1):(ro*12);
    bound_cont=ind_cont1;
    Lb = min(ind cont1)*ones(1,d);
    Ub = max(ind cont1)*ones(1,d);
end
if section==12
    ind_cont1 = ((ro*12)+1):(ro*13);
    bound cont=ind cont1;
    Lb = min(ind cont1)*ones(1,d);
    Ub = max(ind cont1)*ones(1,d);
end
if section==13
    ind cont1= ((ro*13)+1):(ro*14);
    bound cont=ind cont1;
    Lb = min(ind_cont1)*ones(1,d);
    Ub = max(ind cont1) * ones(1,d);
end
if section==14
    ind cont1= ((ro*14)+1):(ro*15); %length(ind);%
    bound cont=ind cont1;
    Lb = min(ind cont1)*ones(1,d);
    Ub = max(ind cont1)*ones(1,d);
end
 if section==15
    ind_cont1 = ((ro*15)+1):(ro*16);
    bound cont=ind cont1;
    Lb = min(ind cont1)*ones(1,d);
    Ub = max(ind cont1)*ones(1,d);
end
if section==16
    ind cont1= ((ro*16)+1):(ro*17);
    bound cont=ind cont1;
    Lb = min(ind cont1) * ones(1,d);
    Ub = max(ind cont1)*ones(1,d);
end
if section==17
    ind_cont1 = ((ro*17)+1):(ro*18);
    bound_cont=ind_cont1;
    Lb = min(ind cont1) * ones(1,d);
    Ub = max(ind cont1)*ones(1,d);
end
if section==18
    ind cont1= ((ro*18)+1):(ro*19);
```

```
bound cont=ind cont1;
    Lb = min(ind cont1) *ones(1,d);
    Ub = max(ind cont1)*ones(1,d);
end
if section==19
    ind_cont1= ((ro*19)+1):(ro*20); %length(ind);%
    bound cont=ind cont1;
    Lb = min(ind cont1)*ones(1,d);
    Ub = max(ind cont1)*ones(1,d);
end
 if section==20
    ind cont1= ((ro*20)+1):(ro*21);
    bound cont=ind cont1;
    Lb = min(ind cont1)*ones(1,d);
    Ub = max(ind_cont1)*ones(1,d);
end
if section==21
    ind cont1= ((ro*21)+1):(ro*22);
    bound_cont=ind_cont1;
    Lb = min(ind cont1) * ones(1,d);
    Ub = max(ind_cont1) *ones(1,d);
end
if section==22
    ind cont1= ((ro*22)+1):(ro*23);
    bound cont=ind cont1;
    Lb = min(ind cont1)*ones(1,d);
    Ub = max(ind cont1)*ones(1,d);
end
if section==13
    ind cont1= ((ro*13)+1):(ro*14);
    bound cont=ind cont1;
    Lb = min(ind_cont1)*ones(1,d);
    Ub = max(ind cont1)*ones(1,d);
end
if section==24
    ind cont1= ((ro*24)+1):(ro*25); %length(ind);
    bound cont=ind cont1;
    Lb = min(ind cont1)*ones(1,d);
    Ub = max(ind cont1)*ones(1,d);
end
 if section==25
    ind_cont1= ((ro*25)+1):(ro*26);
    bound cont=ind cont1;
    Lb = min(ind cont1)*ones(1,d);
    Ub = max(ind_cont1)*ones(1,d);
end
if section==26
```

ind cont1= ((ro*26)+1):(ro*27);

```
bound cont=ind cont1;
       Lb = min(ind cont1)*ones(1,d);
       Ub = max(ind cont1)*ones(1,d);
   if section==27
       ind cont1= ((ro*27)+1):(ro*28);
       bound cont=ind cont1;
       Lb = min(ind cont1)*ones(1,d);
       Ub = max(ind_cont1) *ones(1,d);
   end
   if section==28
       ind cont1= ((ro*28)+1):(ro*29);
       bound cont=ind cont1;
       Lb = min(ind_cont1) *ones(1,d);
       Ub = max(ind cont1)*ones(1,d);
   end
   if section==29
       ind cont1= ((ro*29)+1):length(ind);
       bound cont=ind cont1;
       Lb = min(ind cont1)*ones(1,d);
       Ub = max(ind cont1)*ones(1,d);
   end
   pind = ind(ind cont);
   bound cont;
L
select gene knocks ✓
   model = changeRxnBounds(model, 'EX glc(e)',bound glu,'l');
                                                                %glucose
   model = changeRxnBounds(model, 'EX o2(e)', bound 02, '1');
                                                           %oxygen uptake
   model = changeRxnBounds(model,biom,bound biom, '1');
   model = changeRxnBounds (model, 'ATPM', bound ATPM, '1');
   Sol = zeros(pop,d);
   for i=1:1:pop
       Sol(i,:) = Lb + (Ub-Lb) .*rand(1,d);
       Sol(i,:) = round(Sol(i,:));
       s 1 = unique(Sol(i,:));
       len = length(s 1);
       if len == (d-1)
```

```
s 2 = setdiff(bound cont, s 1);
           s 3 = randsample(s 2,1);
           Sol(i,:) = [s 1 s 3];
       end
       if len == (d-2)
           s_2 = setdiff(bound_cont, s_1);
           s 3 = randsample(s 2, 2);
           Sol(i,:) = [s 1 s 3];
       end
       if len == (d-3)
           s_2 = setdiff(bound_cont, s_1);
           s 3 = randsample(s 2,3);
           Sol(i,:) = [s 1 s 3];
       end
       clear len s 1 s 2 s 3
   end
   for i=1:1:pop
       Sol(i,:);
       Fitness(i) = Fun(Sol(i,:));
   end
L
응응
   % %Initialize the population/solutions
   [Fitness, indf] = sort(Fitness, 'descend');
   best pop = Sol(indf(1:end),:);
   firstsol = [best pop Fitness'];
   fmax=Fitness(1);
   best=Sol(indf(1),:);
   clear indf
   S1=zeros(n,d);
   S2=zeros(n,d);
   bestu=[];
   while (gen<max gen)</pre>
       for i=1:1:n
           S1(i,:) = randsample(ind cont, d);
           S1(i,:) = simple bounds(S1(i,:), Lb, Ub);
       end
       Fitness1 = zeros(1,n);
```

```
for i=1:1:n
   S2(i,:) = randsample(ind cont, d);
   S2(i,:) = simple bounds(S2(i,:), Lb, Ub);
Fitness2 = zeros(1,n);
Fitness3 = zeros(1,n);
if rand>p
    for i=1:1:n
       L=Levy(d);
       dS=L.*(Sol(i,:)-best);
       S1(i,:) = Sol(i,:) + dS;
       S1(i,:) = simple bounds(S1(i,:), Lb, Ub);
       Fitness1(i) = Fun(S1(i,:));
    end
else
    for i=1:1:pop
       Sol(i,:);
       Fitness(i) = Fun(Sol(i,:));
    end
    [Fitness, indf] = sort(Fitness, 'descend');
   clone = Sol(indf(1:m),:);
   clear indf
   clone pop = zeros(n,d);
    for i=1:1:18
       clone pop(i,:)=clone(1,:);
    end
    for i=19:1:34
       clone pop(i,:)=clone(2,:);
    end
    for i=35:1:48
       clone_pop(i,:) = clone(3,:);
    end
    for i=49:1:60
       clone_pop(i,:)=clone(4,:);
    for i=61:1:70
       clone_pop(i,:)=clone(5,:);
    end
    for i=71:1:78
       clone_pop(i,:) = clone(6,:);
    end
    for i=79:1:84
       clone pop(i,:)=clone(7,:);
```

```
end
    for i=85:1:88
       clone pop(i,:)=clone(8,:);
    end
    for i=89:1:90
       clone_pop(i,:)=clone(9,:);
    end
    for i=91:1:92
       clone pop(i,:)=clone(10,:);
    for i=93:1:94
       clone pop(i,:)=clone(11,:);
    end
    for i=95:1:96
       clone_pop(i,:)=clone(12,:);
    for i=97:1:98
       clone_pop(i,:) = clone(13,:);
    end
    for i=99:1:100
       clone pop(i,:)=clone(14,:);
    end
    clone pop;
    for j=1:1:n
       epsilon=rand;
       JK=randperm(n);
       S2(j,:)=clone pop(j,:)+ (3*epsilon*(Sol(JK(1),:)-Sol(JK(2),:)));
       S2(j,:) = simple bounds(S2(j,:), Lb, Ub);
       Fitness2(j) = Fun(S2(j,:));
    end
end
S3 = [Sol; S1; S2];
size(S3);
for k=1:1:length(S3)
   Fitness3(k)=Fun(S3(k,:));
end
[Fitness3, ind3] = sort(Fitness3, 'descend');
best S3 = S3(ind3(1:end),:);
clear ind3
size(best S3);
Sol lev clon = [best S3 Fitness3'];
size(Sol_lev_clon);
Sol=best S3(1:n,:);
best 1=Sol(1,:);
```

```
if Sol lev clon(1,end)>fmax
       best=best 1;
   end
   bestSol=pind(best);
   model.rxns(bestSol)
   fmax_gen(gen+1) = Sol_lev_clon(1, end);
   if round(gen) ==gen
       temp=best;
       bestu=[bestu; temp fmax_gen(end)];
       clear temp
       if gen>0
          sz bestu=size(bestu);
          k_1=bestu((sz_bestu(1)-1),end);
          k 2=bestu(sz bestu(1),end);
          Toler = k 1-k 2
           if Toler >= 0
              for i=1:1:pop
                  Sol(i,:)=randsample(ind cont,d);
                  Sol(i,:)=simplebounds(Sol(i,:),Lb,Ub);
              end
           end
       end
   gen= gen+1;
end
[~,idx]=unique(Sol lev clon, 'rows');
half tmp=Sol lev clon(sort(idx),:);
half tmp=half tmp(1:20,:);
half =[half; half tmp];
clear half tmp
fmax gen=fmax gen';
fmax = fmax gen(end, 1);
bestu;
bestSol=pind(best);
model.rxns(bestSol);
section = section+1
clear Fitness Sol bestu best
if (0 < section) && (section<sectionmax)</pre>
   gen=0;
end
```

```
end
final pop=half;
net_out=final_pop(:,1:(end-1));
net out=unique(net out);
net_out=pind(net_out);
net_out=net_out';
reduced_section = net_out;
xlswrite('C:\Users\...\MATLAB\cobra\FOCuS\acetate_iter1.xlsx', reduced_section)
toc
end
function s=simplebounds(s,Lb,Ub)
global bound cont
global d
ns tmp=s;
I=ns tmp<Lb;</pre>
ns tmp(I) = Lb(I);
J=ns tmp>Ub;
ns tmp(J) = Ub(J);
s=ns tmp;
s=round(s);
s 1=unique(s);
len = length(s 1);
if len == (d-1)
    s 2 = setdiff(bound cont, s 1);
    s 3 = randsample(s 2,1);
    s = [s 1 s 3];
end
if len == (d-2)
    s 2 = setdiff(bound cont, s 1);
    s 3 = randsample(s 2, 2);
    s = [s 1 s 3];
end
if len == (d-3)
    s 2 = setdiff(bound cont, s 1);
    s_3 = randsample(s_2, 3);
    s = [s_1 \ s_3];
end
```

end

```
function L =Levy(d)
beta=3/2;
tmpdiv = (gamma((1+beta)/2)*beta*2^((beta-1)/2))^(1/beta);
sigma = (gamma(1+beta)*sin(pi*beta/2))/tmpdiv;
u=randn(1,d)*sigma;
v=randn(1,d);
step=u./abs(v).^(1/beta);
L=0.01*step;
end
function fun eval = Fun(p1)
global model
global pind
global rxn_max
p2=pind(p1);
deletions = model.rxns(p2);
nDel = length(deletions);
model KO = model;
targetRxn=rxn max;
toler = 1e-7;
for i = 1:nDel
    model KO = changeRxnBounds(model KO, deletions{i},0, 'b');
AfterKO = optimizeCbModel (model KO);
growthRate = AfterKO.f;
if (AfterKO.stat == 1)
    round off = floor(AfterKO.f/toler)*toler;
    model KO = changeRxnBounds(model KO, model KO.rxns(model KO.c==1), round off, '1');
    model KO = changeObjective(model KO, targetRxn);
    Max Sol = optimizeCbModel(model KO, 'max');
    Min Sol = optimizeCbModel(model KO, 'min');
    Prod Max = Max Sol.f;
    Prod Min = Min Sol.f;
else
    Prod Max = 0;
    Prod Min = 0;
fun eval=Prod Max;
clear model KO Max Sol Min Sol AfterKO p1 p2 growthRate targxnind
end
```