

# Extract a thermodynamically consistent subnetwork from a given model

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## Reviewer(s):

1. Identify the largest subset of a model that admits a thermodynamically consistent flux
2. Specify a random subset of active/inactive reactions and present/absent metabolites
3. Compute the smallest thermodynamically consistent subnetwork containing a list of present metabolites and active reactions, and not containing a list of absent metabolites and inactive reactions

```
[solverOK,solverInstalled]=changeCobraSolver('ibm_cplex','all');
```

```
> changeCobraSolver: IBM ILOG CPLEX interface added to MATLAB path.  
> ibm_cplex (version 1210) is compatible and fully tested with MATLAB R2021a on your operating system.  
> changeCobraSolver: Solver for LP problems has been set to ibm_cplex.  
  
> changeCobraSolver: IBM ILOG CPLEX interface added to MATLAB path.  
> ibm_cplex (version 1210) is compatible and fully tested with MATLAB R2021a on your operating system.  
> changeCobraSolver: Solver for MILP problems has been set to ibm_cplex.  
  
> changeCobraSolver: IBM ILOG CPLEX interface added to MATLAB path.  
> ibm_cplex (version 1210) is compatible and fully tested with MATLAB R2021a on your operating system.  
> changeCobraSolver: Solver for QP problems has been set to ibm_cplex.  
  
> changeCobraSolver: IBM ILOG CPLEX interface added to MATLAB path.  
> ibm_cplex (version 1210) is compatible and fully tested with MATLAB R2021a on your operating system.  
> changeCobraSolver: Solver for MIQP problems has been set to ibm_cplex.  
> changeCobraSolver: Solver ibm_cplex not supported for problems of type EP. No solver set for this problem.  
> changeCobraSolver: Solver ibm_cplex not supported for problems of type NLP. No solver set for this problem.
```

```
%[solverOK,solverInstalled]=changeCobraSolver('gurobi','all');  
%[solverOK,solverInstalled]=changeCobraSolver('ibm_cplex','QP');
```

## Load model

```
modelToLoad='circularToy';  
modelToLoad='ecoli_core';  
modelToLoad='modelRecon3MitoOpen';  
modelToLoad='Recon3DModel';  
%modelToLoad='iDopa';
```

## Load a model

```
driver_thermoModelLoad
```

```
Model loaded: Recon3DModel  
lower bounds greater than zero  
Internal stoichiometric nullspace computed in 4.5238 seconds.
```

## Stoichiometric consistency

```
if ~isfield(model,'SConsistentRxnBool') ||  
~isfield(model,'SConsistentMetBool')  
    massBalanceCheck=0;  
    %massBalanceCheck=1;  
    printLevel=2;  
    [SConsistentMetBool, SConsistentRxnBool,  
SInConsistentMetBool, SInConsistentRxnBool, unknownSConsistencyMetBool,  
unknownSConsistencyRxnBool, model,stoichConsistModel]...  
        = findStoichConsistentSubset(model, massBalanceCheck, printLevel);  
else  
    %Extract stoich consistent submodel  
    if any(~model.SConsistentMetBool)  
        rxnRemoveMethod='inclusive';%maintains stoichiometric consistency  
        [stoichConsistModel, rxnRemoveList] = removeMetabolites(model,  
model.mets(~model.SConsistentMetBool),rxnRemoveMethod);  
        SConsistentRxnBool2=~ismember(model.rxns,rxnRemoveList);  
        if ~all(model.SConsistentRxnBool==SConsistentRxnBool2)  
            error('inconsistent reaction removal')  
        end  
        try  
            stoichConsistModel = removeUnusedGenes(stoichConsistModel);  
        catch ME  
            disp(ME.message)  
        end  
    else  
        stoichConsistModel = model;  
    end  
end  
  
[nMet,nRxn]=size(stoichConsistModel.S)
```

```
nMet = 5835  
nRxn = 10600
```

## Flux consistency

```
fluxConsistentParam.method='fastcc';%can handle additional constraints  
fluxConsistentParam.printLevel=1;  
[~,~,~,~,stoichConsistModel]=  
findFluxConsistentSubset(stoichConsistModel,fluxConsistentParam);
```

Extract flux consistent submodel

```
if any(~stoichConsistModel.fluxConsistentRxnBool)  
    rxnRemoveList =  
stoichConsistModel.rxns(~stoichConsistModel.fluxConsistentRxnBool);
```

```

    stoichFluxConsistModel = removeRxns(stoichConsistModel,
rxnRemoveList, 'metRemoveMethod', 'exclusive', 'ctrRemoveMethod', 'inclusive');
    try
        stoichFluxConsistModel = removeUnusedGenes(stoichFluxConsistModel);
    catch ME
        disp(ME.message)
    end
else
    stoichFluxConsistModel = stoichConsistModel;
end
[nMet,nRxn]=size(stoichFluxConsistModel.S)

nMet = 5835
nRxn = 10600

```

## Thermodynamic consistency

```

%save('debug_prior_to_findThermoConsistentFluxSubset.mat')
%return
param.printLevel = 1;
param.relaxBounds=0;
param.acceptRepairedFlux=1;
[thermoFluxConsistentMetBool,thermoFluxConsistentRxnBool,stoichFluxConsistModel,stoichFluxThermoConsistModel] =
findThermoConsistentFluxSubset(stoichFluxConsistModel,param);

```

```

--- findThermoFluxConsistentSubset START ----
    printLevel: 1
    relaxBounds: 0
    acceptRepairedFlux: 1
    epsilon: 1.0000e-06
    formulation: 'pqzw'
    iterationMethod: 'random'
    nMax: 20
    warmStartMethod: 'random'
    thetaMultiplier: 1.5000
    theta: 0.5000
    regularizeOuter: 0
thermoConsistencyMethod: 'cycleFreeFlux'
    bigNum: 10000
    debug: 0

```

optCardThermo objective data:

```

0.1 = beta, the global weight on one-norm of internal reaction rate.
-5 = min(g0), the local weight on zero-norm of internal reaction rate.
-0 = max(g0), the local weight on zero-norm of internal reaction rate.
0 = min(h0), the local weight on zero-norm of metabolite production rate.
0 = max(h0), the local weight on zero-norm of metabolite production rate.

```

optimizeCardinality objective data:

0 min cardinality variables:

NaN mean(c(p))	NaN min(c(p))	NaN max(c(p))
1 lambda0	NaN min(k)	NaN max(k)
1 lambda1	NaN min(o(p))	NaN max(o(p))

5303 max cardinality variables:

```

-0 mean(c(q))          -0 min(c(q))          -0 max(c(q))
1 delta0                5 min(d)              5 max(d)
0 delta1                0 min(o(q))            0 max(o(q))

22879 cardinality free variables:
0.077 mean(c(r))        -0 min(c(r))          0.1 max(c(r))
0 alpha1                0 min(o(r))            0 max(o(r))

itn    theta    ||dx||    del_obj    obj    linear    ||x||0    a(x)    ||x||1    ||y||
1      0.50    7.6657e+05  -8.8e+06  -2.2e+04  1.6e+03      0      0      0     -236
2      0.75    184.22    -8.5e+02  -2.3e+04  1.1e+03      0      0      0     -238
3      1.12    98.541    -4.8e+02  -2.3e+04  7.3e+02      0      0      0     -239
4      1.69    61.895    -2.6e+02  -2.3e+04  4.9e+02      0      0      0     -239
5      2.53    40.001    -1.7e+02  -2.4e+04  3.3e+02      0      0      0     -239
6      3.80    26.092    -1.3e+02  -2.4e+04  2.2e+02      0      0      0     -239
7      5.70    18.183      -73    -2.4e+04  1.5e+02      0      0      0     -239
8      8.54    11.708     -68    -2.4e+04      97      0      0      0     -239
9     12.81      7.942     -34    -2.4e+04      65      0      0      0     -239
10    19.22     5.2426     -36    -2.4e+04      44      0      0      0     -239
11    28.83     3.5411     -48    -2.4e+04      29      0      0      0     -240
12    43.25     2.3919     -17    -2.4e+04      20      0      0      0     -240
13    64.87     1.6485     -13    -2.4e+04      14      0      0      0     -240
14    97.31     1.0952     -4.2    -2.4e+04      9.3      0      0      0     -240
15   145.96     0.76091     -2.9    -2.4e+04      6.5      0      0      0     -240
16   218.95     0.51853     -42    -2.4e+04      4.7      0      0      0     -240
17   328.42     0.53726     -14    -2.4e+04      3.5      0      0      0     -240
itn    theta    ||dx||    del_obj    obj    linear    ||x||0    a(x)    ||x||1    ||y||
Optimise cardinality reached the stopping criterion. Finished.

cycleFreeFlux: No solution found.
Debugging relaxation etc...
  full: []
  obj: []
  rcost: []
  dual: []
  slack: []
  solver: 'ibm_cplex'
  algorithm: 'Automatic'
  stat: 0
  origStat: 3
  origStatText: 'Model has been proved infeasible'
  time: 0.0300
  basis: []

itn    obj    obj_old    err(obj)    err(x)    card(v)    card(r)    card(p)    card(q)
0      14.072    66705    66691    165.16    3913      0      543      541
1      1391    14.072    1376.9    48.563    2734      0      692      717
2      944.42    1391    446.54    62.844    3799      0      652      666
3      1526.3    944.42    581.91    64.636    3026      0      712      702
4      828.59    1526.3    697.73    63.09    3157      0      574      605
5      992.49    828.59    163.9    55.391    3334      0      716      682
6      1074.4    992.49    81.866    59.058    3674      0      687      657
7      1300.5    1074.4    226.18    63.35    3238      0      700      708
8      1040.9    1300.5    259.63    62.911    3773      0      685      639
9      1327.8    1040.9    286.85    63.278    3013      0      703      667
10     814.59    1327.8    513.18    60.13    3064      0      614      603
11     1027.5    814.59    212.94    55.715    2717      0      628      614
12     677.72    1027.5    349.8    53.573    3435      0      602      618
13     1061.2    677.72    383.44    54.092    3505      0      734      693
14     1272.7    1061.2    211.54    62.794    3509      0      693      670
15     1197.8    1272.7    74.901    64.628    3152      0      669      689
16     971.72    1197.8    226.09    60.517    2983      0      585      622
17     903.71    971.72    68.006    56.208    2923      0      612      658
18     834.71    903.71      69    54.078    3806      0      671      642

```

19	1348.3	834.71	513.59	60.702	3075	0	720	659
itn	obj	obj_old	err(obj)	err(x)	card(v)	card(r)	card(p)	card(q)

Relaxed model is feasible.  
Statistics:  
599 lower bound relaxation(s)  
576 upper bound relaxation(s)  
0 steady state relaxation(s)  
... done.  
ans = 9.1346e-07  
ans = 0  
ans = 0  
solution = struct with fields:  
  stat: 1  
  v: [19391x1 double]  
  r: [23417x1 double]  
  p: [19391x1 double]  
  q: [19391x1 double]  
solutionRelaxed1 = struct with fields:  
  full: [19391x1 double]  
  obj: 26.3611  
  rcost: [19391x1 double]  
  dual: [23417x1 double]  
  slack: [23417x1 double]  
  solver: 'ibm\_cplex'  
  algorithm: 'Automatic'  
  stat: 1  
  origStat: 1  
  origStatText: 'Optimal solution found'  
  time: 1.2415  
  basis: []  
solutionRelaxed2 = struct with fields:  
  full: [19391x1 double]  
  obj: 0  
  rcost: [19391x1 double]  
  dual: [23417x1 double]  
  slack: [23417x1 double]  
  solver: 'ibm\_cplex'  
  algorithm: 'Automatic'  
  stat: 1  
  origStat: 1  
  origStatText: 'Optimal solution found'  
  time: 0.6092  
  basis: []  
solutionRelaxed3 = struct with fields:  
  full: [19391x1 double]  
  obj: 0  
  rcost: [19391x1 double]  
  dual: [23417x1 double]  
  slack: [23417x1 double]  
  solver: 'ibm\_cplex'  
  algorithm: 'Automatic'  
  stat: 1  
  origStat: 1  
  origStatText: 'Optimal solution found'  
  time: 0.3797  
  basis: []  
cycleFreeFlux: No solution found, try using a different solver.  
  
cycleFreeFlux: No solution found.  
Debugging relaxation etc...  
  full: []  
  obj: []  
  rcost: []  
  dual: []

```

    slack: []
    solver: 'ibm_cplex'
    algorithm: 'Automatic'
    stat: 0
    origStat: 3
    origStatText: 'Model has been proved infeasible'
    time: 0.0305
    basis: []

itn      obj      obj_old    err(obj)    err(x)    card(v)    card(r)    card(p)    card(q)
0        14.072      66355      66341      164.43      3913         0         543         541
1         1391      14.072     1376.9     48.563      2734         0         692         717
2        944.42      1391      446.54     62.844      3799         0         652         666
3        1526.3     944.42     581.91     64.636      3026         0         712         702
4        828.59     1526.3     697.73     63.09       3157         0         574         605
5        992.49     828.59     163.9      55.391      3334         0         716         682
6       1074.4     992.49     81.866     59.058      3674         0         687         657
7       1300.5     1074.4     226.18      63.35       3238         0         700         708
8       1040.9     1300.5     259.63     62.911      3773         0         685         639
9       1327.8     1040.9     286.85     63.278      3013         0         703         667
10      814.59     1327.8     513.18      60.13       3064         0         614         603
11      1027.5     814.59     212.94     55.715      2717         0         628         614
12      677.72     1027.5      349.8     53.573      3435         0         602         618
13      1061.2     677.72     383.44     54.092      3505         0         734         693
14      1272.7     1061.2     211.54     62.794      3509         0         693         670
15      1197.8     1272.7     74.901     64.628      3152         0         669         689
16      971.72     1197.8     226.09     60.517      2983         0         585         622
17      903.71     971.72     68.006     56.208      2923         0         612         658
18      834.71     903.71      69         54.078      3806         0         671         642
19     1348.3     834.71     513.59     60.702      3075         0         720         659
itn      obj      obj_old    err(obj)    err(x)    card(v)    card(r)    card(p)    card(q)
Relaxed model is feasible.
Statistics:
599 lower bound relaxation(s)
576 upper bound relaxation(s)
0 steady state relaxation(s)
... done.
ans = 9.1346e-07
ans = 0
ans = 0
solution = struct with fields:
    stat: 1
    v: [19391x1 double]
    r: [23417x1 double]
    p: [19391x1 double]
    q: [19391x1 double]
solutionRelaxed1 = struct with fields:
    full: [19391x1 double]
    obj: 26.3611
    rcost: [19391x1 double]
    dual: [23417x1 double]
    slack: [23417x1 double]
    solver: 'ibm_cplex'
    algorithm: 'Automatic'
    stat: 1
    origStat: 1
    origStatText: 'Optimal solution found'
    time: 1.3163
    basis: []
solutionRelaxed2 = struct with fields:
    full: [19391x1 double]
    obj: 0
    rcost: [19391x1 double]
    dual: [23417x1 double]

```

```

        slack: [23417x1 double]
        solver: 'ibm_cplex'
        algorithm: 'Automatic'
        stat: 1
        origStat: 1
        origStatText: 'Optimal solution found'
        time: 0.5761
        basis: []
solutionRelaxed3 = struct with fields:
        full: [19391x1 double]
        obj: 0
        rcost: [19391x1 double]
        dual: [23417x1 double]
        slack: [23417x1 double]
        solver: 'ibm_cplex'
        algorithm: 'Automatic'
        stat: 1
        origStat: 1
        origStatText: 'Optimal solution found'
        time: 0.3650
        basis: []

```

Warning: cycleFreeFlux did not solve, trying v2QNTy

76.81% thermodynamically feasible internal fluxes (checked by v2QNTy method).

iter	card(y)	nz	%feas	int.nz.	tot %feas	int.nz.	tot
1	5303	7882	0.68	0.50			
2	2376	3075	1.00	0.69			
3	1427	1708	1.00	0.77			
4	1022	1296	1.00	0.83			
5	769	794	1.00	0.86			
6	606	587	1.00	0.88			
7	506	432	1.00	0.90			
8	449	377	1.00	0.91			
9	389	324	1.00	0.92			
10	348	301	1.00	0.93			
11	298	332	1.00	0.94			
12	254	384	1.00	0.95			
13	216	209	1.00	0.96			
14	191	200	1.00	0.97			
15	153	201	1.00	0.97			
16	151	162	1.00	0.97			
17	106	139	1.00	0.98			
18	102	161	1.00	0.98			
19	78	127	1.00	0.98			
20	91	128	1.00	0.98			
iter	card(y)	nz	%feas	int.nz.	tot %feas	int.nz.	tot

findThermoConsistentFluxSubset terminating early: n = nMax = 20  
 --- findThermoFluxConsistentSubset END----

Size of the largest flux, stoich and thermo consistent submodel

```

[nMet,nRxn]=size(stoichFluxThermoConsistModel.S)
save(['~/work/sbgCloud/programModelling/projects/thermoModel/results/
thermoKernel/' modelToLoad
'_stoichFluxThermoConsistModel.mat'],'stoichFluxThermoConsistModel')

```

```

%modelToLoad='Recon3DModel';
load(['~/work/sbgCloud/programModelling/projects/thermoModel/results/
thermoKernel/' modelToLoad
'_stoichFluxThermoConsistModel.mat'],'stoichFluxThermoConsistModel')

```

## Nullspace

Nullspace is necessary for backup check of thermodynamic consistency using thermoFlux2QNty

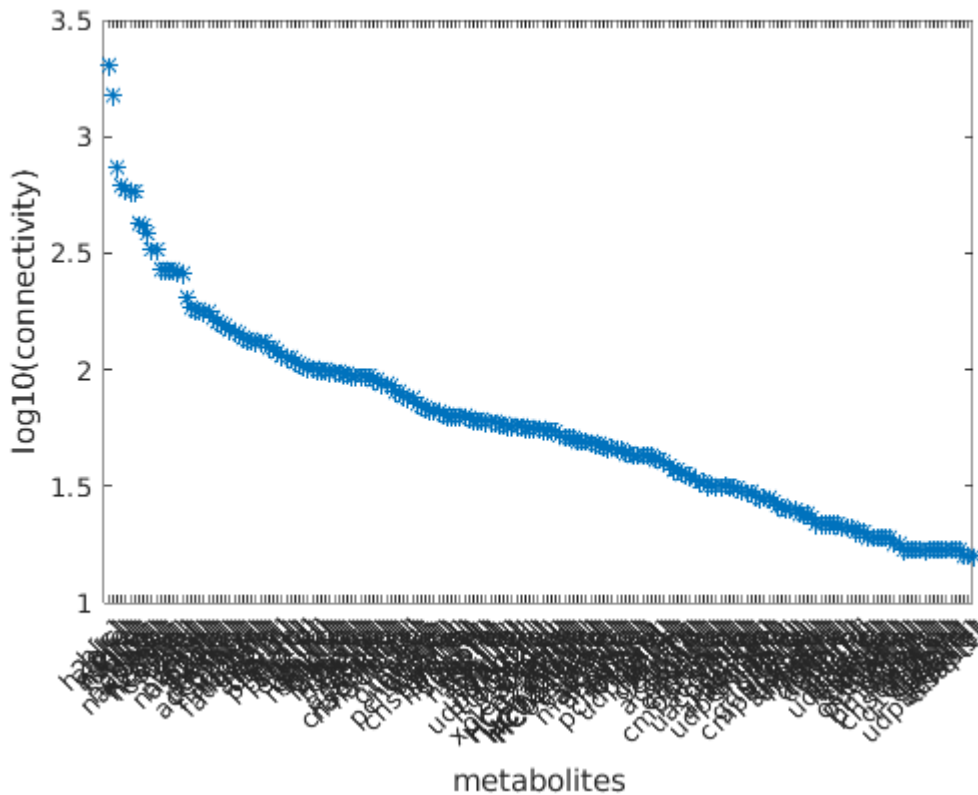
```
[stoichFluxThermoConsistModel,rankK,nnzK,timeTaken] =  
internalNullspace(stoichFluxThermoConsistModel);  
rankK
```

```
rankK = 5485
```

## Data to define a thermodynamically consistent subnetwork

Setup random data to select a random subset

```
param.n=200;  
[rankMetConnectivity,rankMetInd,rankConnectivity] =  
rankMetabolicConnectivity(stoichFluxThermoConsistModel,param);
```



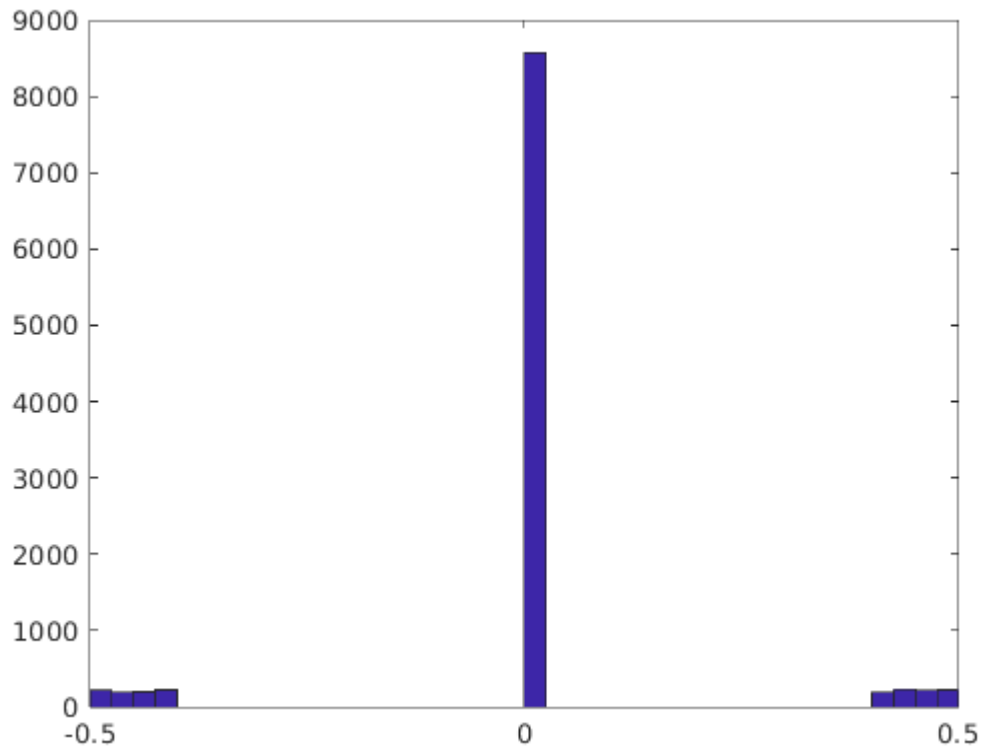
```
[nMet,nRxn]=size(stoichFluxThermoConsistModel.S);  
rxnWeights=rand(nRxn,1)-0.5;  
rxnWeights(~stoichFluxThermoConsistModel.SConsistentRxnBool)=0;  
  
coreRxnBool=rxnWeights<-0.45;  
removeRxnBool=rxnWeights>0.45;  
if 0
```



```

rxnWeights(rxnWeights>0.4)=1;
rxnWeights(rxnWeights<-0.4)=-1;
end
rxnWeights(rxnWeights>=-0.4 & rxnWeights<=0.4)=0;
hist(rxnWeights,40)

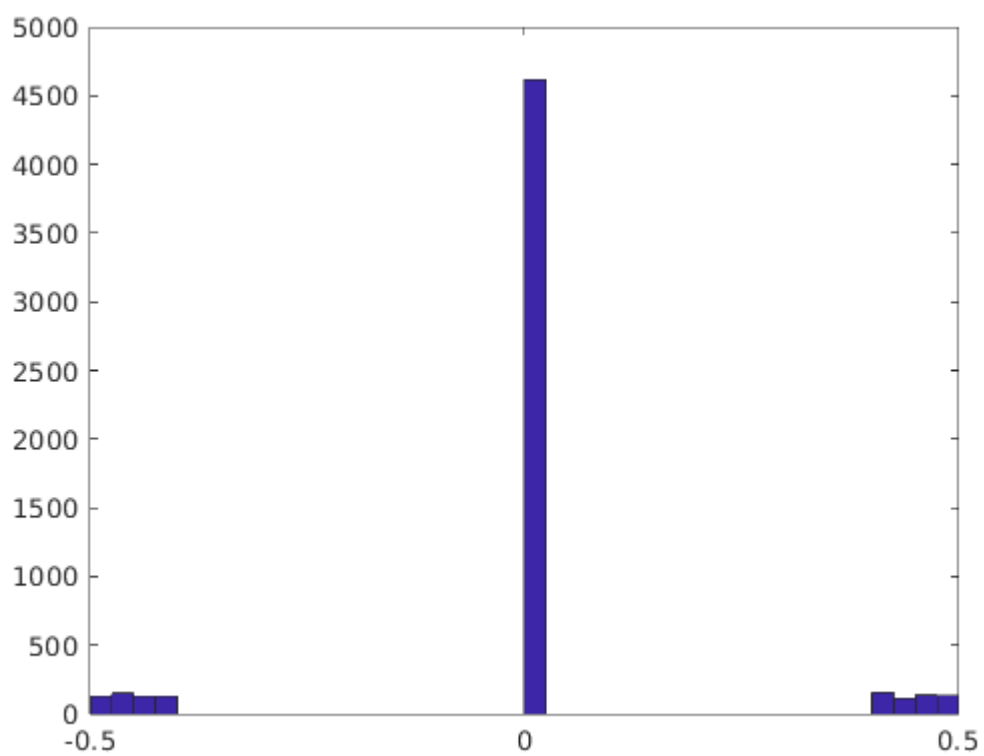
```



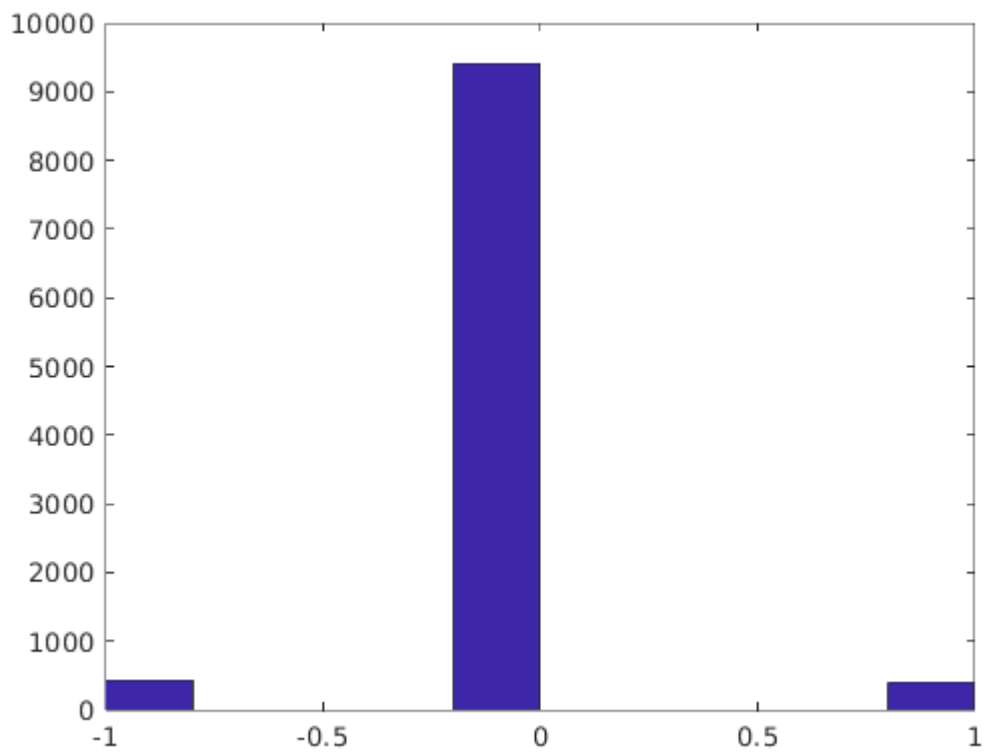
```

metWeights=rand(nMet,1)-0.5;
metWeights(rankMetInd(1:200))=0;
coreMetBool=metWeights<-0.45;
removeMetBool=metWeights>0.45;
if 0
    metWeights(metWeights>0.4)=1;
    metWeights(metWeights<-0.4)=-1;
end
metWeights(metWeights>=-0.4 & metWeights<=0.4)=0;
hist(metWeights,40)

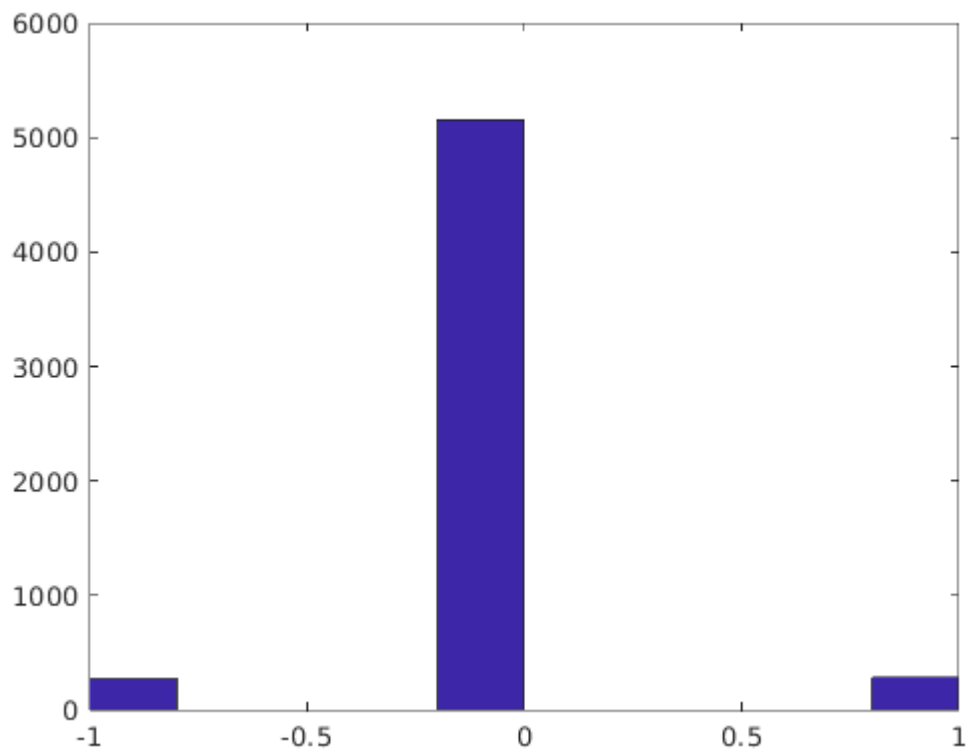
```



```
nlt=length(coreRxnBool);  
activeInactiveRxn=zeros(nlt,1);  
activeInactiveRxn(coreRxnBool)=1;  
activeInactiveRxn(removeRxnBool)=-1;  
hist(activeInactiveRxn)
```



```
mlt=length(coreMetBool);  
presentAbsentMet=zeros(mlt,1);  
presentAbsentMet(coreMetBool)=1;  
presentAbsentMet(removeMetBool)=-1;  
if 0  
    activeInactiveRxn(:)=0;  
    presentAbsentMet(:)=0;  
end  
param.normalizeZeroNormWeights=0;  
  
hist(presentAbsentMet)
```



**Compute the smallest thermodynamically consistent subnetwork given a list of present/absent metabolites and active/inactive reactions**

```
[tissueModel, thermoModelMetBool, thermoModelRxnBool] =
thermoKernel(stoichFluxThermoConsistModel, activeInactiveRxn, rxnWeights,
presentAbsentMet, metWeights,param);
```

```
--- thermoKernel START ---
thermoKernel parameters:
    printLevel: 1
    relaxBounds: 0
    acceptRepairedFlux: 1
        n: 200
    normalizeZeroNormWeights: 0
        formulation: 'pqzwrS'
        epsilon: 1.0000e-06
    removeOrphanGenes: 1
        nbMaxIteration: 30
        nMax: 20
    iterationMethod: 'greedyRandom'

    warmStartMethod: 'random'
        formulation: 'pqzwrS'
    thetaMultiplier: 1.5000
        theta: 0.5000
    regularizeOuter: 1
        epsilon: 1.0000e-06
    printLevel: 1
    relaxBounds: 0
    acceptRepairedFlux: 1
```

```

thermoConsistencyMethod: 'cycleFreeFlux'
    bigNum: 10000
    debug: 0

optCardThermo objective data:
    1 = beta, the global weight on one-norm of internal reaction rate.
    -0.5 = min(g0), the local weight on zero-norm of internal reaction rate.
    0.5 = max(g0), the local weight on zero-norm of internal reaction rate.
    -0.5 = min(h0), the local weight on zero-norm of metabolite production rate.
    0.5 = max(h0), the local weight on zero-norm of metabolite production rate.

optimizeCardinality objective data:

1393 min cardinality variables:
    0 mean(c(p))          -0 min(c(p))          -0 max(c(p))
    1 lambda0             0.4 min(k)             0.5 max(k)
    0 lambda1             0 min(o(p))             0 max(o(p))

1912 max cardinality variables:
    0 mean(c(q))          -0 min(c(q))          -0 max(c(q))
    1 delta0             0.4 min(d)             0.5 max(d)
    0 delta1             0 min(o(q))             0 max(o(q))

35242 cardinality free variables:
    0.48 mean(c(r))       -0 min(c(r))          1 max(c(r))
    0 alpha1             0 min(o(r))             0 max(o(r))

itn    theta    ||dx||    del_obj    obj    linear    ||x||0    a(x)    ||x||1    ||y||
  1    0.50    9.6992e+05    -8.5e+07    7.9        11    1.64389    0.11        0    -14.90
  2    0.75     4.5121     -1.7        6.2        12    2.0597    0.16        0    -15.84
  3    1.12     2.8094     -1.8        4.4        12    1.65769    0.25        0    -17.51
  4    1.69     3.0116      -2         2.4        12    1.23553    0.43        0    -18.81
  5    2.53     3.0252     -2.3        0.041      13    1.23553    0.45        0    -19.68
  6    3.80     2.7007     -3.3       -3.3       13    1.23553    0.47        0    -24.78
  7    5.70     2.0612     -4.9       -8.2       14    1.23553    0.87        0    -25.69
  8    8.54     1.9653     -2.8       -11       13    1.23553    0.9         0    -27.03
  9   12.81     1.3137     -3.8       -15       12    1.23553    0.95        0    -30.49
 10   19.22     1.2354      -4        -19       11    1.23553    1           0    -32.32
 11   28.83     0.809      -2.2       -21       10    1.23553    1.1         0    -32.32
 12   43.25     0.71837    -0.57      -21       9.6    1.23553    1.2         0    -32.32
 13   64.87     0.46434    -0.55      -22       9.2    1.23553    1.2         0    -33.26
 14   97.31     0.33824    -1.1       -23       8.9    1.23553    1.2         0    -33.26
 15  145.96     0.21621    -0.14      -23       8.8    1.23553    1.2         0    -33.26
 16  218.95     0.2235     -0.088     -23       8.7    1.23553    1.2         0    -33.26
 17  328.42     0.10113    -0.058     -23       8.6    1.23553    1.2         0    -33.26
itn    theta    ||dx||    del_obj    obj    linear    ||x||0    a(x)    ||x||1    ||y||
Optimise cardinality reached the stopping criterion. Finished.
100.00% thermodynamically feasible internal fluxes (checked by cycleFreeFlux method).
    warmStartMethod: 'random'
    formulation: 'pqzwrS'
    thetaMultiplier: 1.5000
    theta: 0.5000
    regularizeOuter: 1
    epsilon: 1.0000e-06
    printLevel: 1
    relaxBounds: 0
    acceptRepairedFlux: 1
thermoConsistencyMethod: 'cycleFreeFlux'
    bigNum: 10000
    debug: 0

optCardThermo objective data:
    1 = beta, the global weight on one-norm of internal reaction rate.
    -0.5 = min(g0), the local weight on zero-norm of internal reaction rate.

```

0.5 = max(g0), the local weight on zero-norm of internal reaction rate.  
-0.5 = min(h0), the local weight on zero-norm of metabolite production rate.  
0.5 = max(h0), the local weight on zero-norm of metabolite production rate.

optimizeCardinality objective data:

1393 min cardinality variables:

0 mean(c(p))	-0 min(c(p))	-0 max(c(p))
1 lambda0	0.4 min(k)	0.5 max(k)
0 lambda1	0 min(o(p))	0 max(o(p))

1912 max cardinality variables:

0 mean(c(q))	-0 min(c(q))	-0 max(c(q))
1 delta0	0.4 min(d)	0.5 max(d)
0 delta1	0 min(o(q))	0 max(o(q))

35242 cardinality free variables:

0.48 mean(c(r))	-0 min(c(r))	1 max(c(r))
0 alpha1	0 min(o(r))	0 max(o(r))

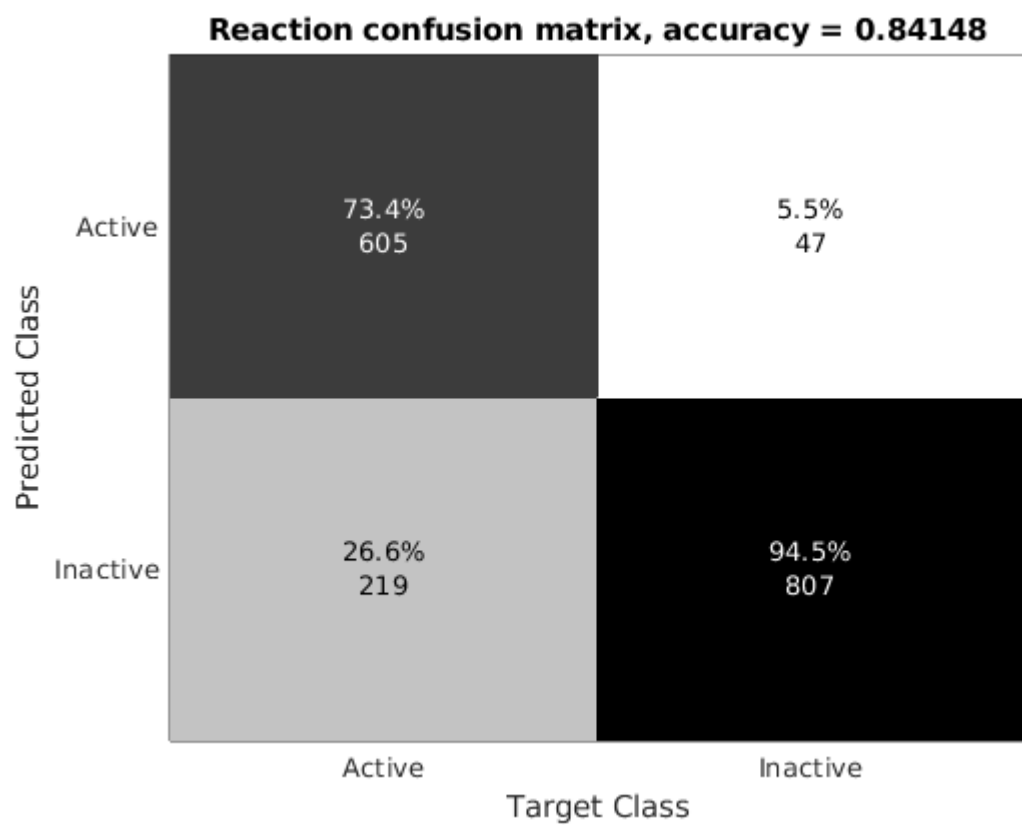
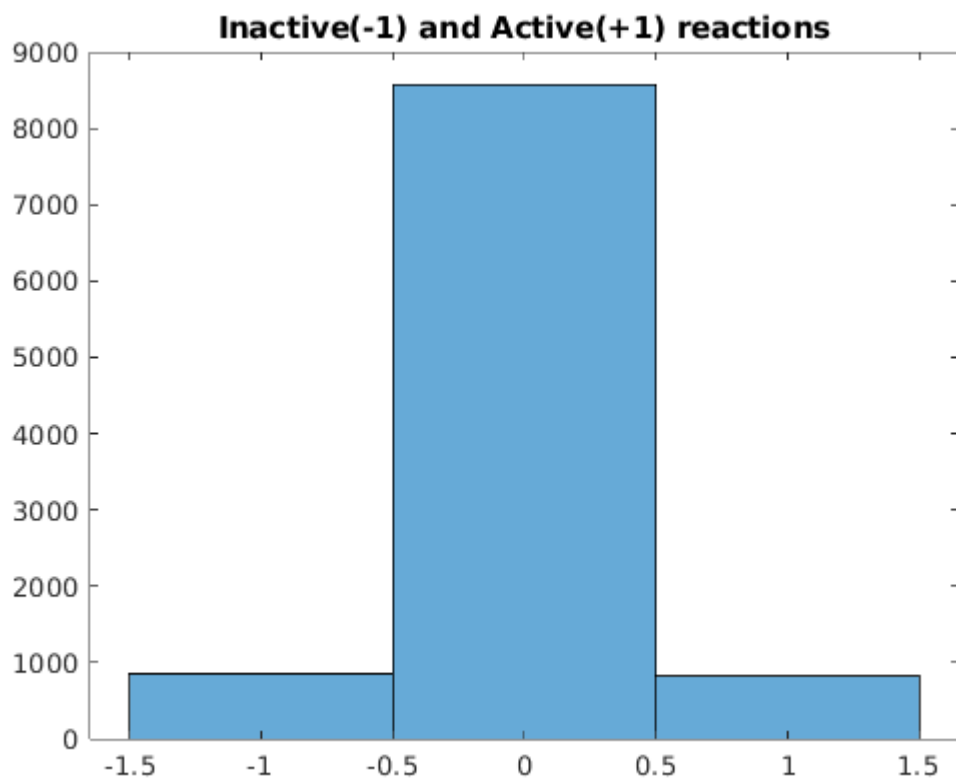
itn	theta	dx	del_obj	obj	linear	x  0	a(x)	x  1	y
1	0.50	9.6862e+05	-8.4e+07	8.1	11	3.74292	0.11	0	-13.29
2	0.75	2.3033	-1	7.1	10	2.06605	0.16	0	-12.36
3	1.12	2.5538	-1.1	6	10	2.06605	0.26	0	-13.24
4	1.69	2.8214	-1.1	4.8	11	1.23553	0.43	0	-14.56
5	2.53	3.0167	-2.9	2	12	1.23553	0.45	0	-17.11
6	3.80	2.0716	-2.1	-0.084	12	1.23553	0.47	0	-18.83
7	5.70	2.5196	-3.8	-3.9	13	0.832228	0.47	0	-20.67
8	8.54	1.9952	-3.6	-7.5	13	0.832228	0.5	0	-22.59
9	12.81	1.589	-3	-10	12	0.832228	0.55	0	-23.07
10	19.22	1.0613	-1.5	-12	10	0.832228	0.62	0	-23.07
11	28.83	0.69458	-0.65	-13	9.7	0.832228	0.73	0	-23.07
12	43.25	0.49103	-0.32	-13	9.3	0.832228	0.83	0	-23.07
13	64.87	0.24192	-0.24	-13	9	0.832228	0.83	0	-23.07
14	97.31	0.21816	-0.55	-14	8.9	0.832228	0.83	0	-23.47
15	145.96	0.17636	-0.075	-14	8.8	0.832228	0.83	0	-23.47
16	218.95	0.097141	-0.048	-14	8.7	0.832228	0.83	0	-23.47
17	328.42	0.16623	-0.031	-14	8.7	0.832228	0.83	0	-23.47
itn	theta	dx	del_obj	obj	linear	x  0	a(x)	x  1	y

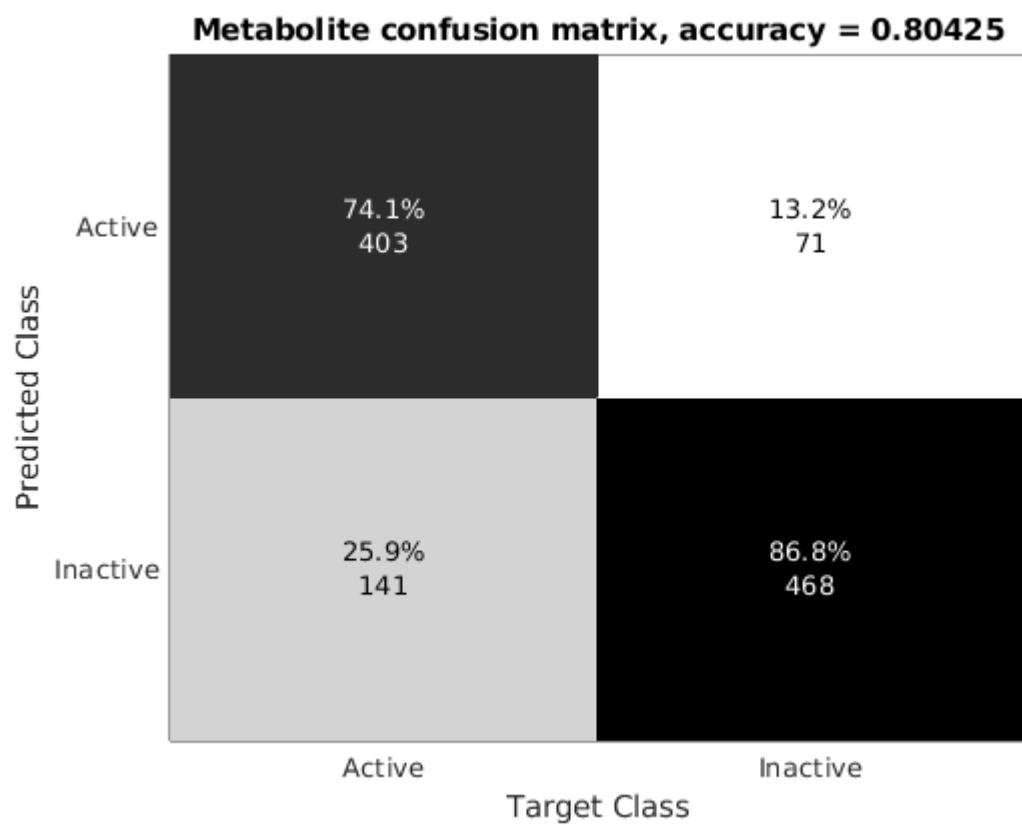
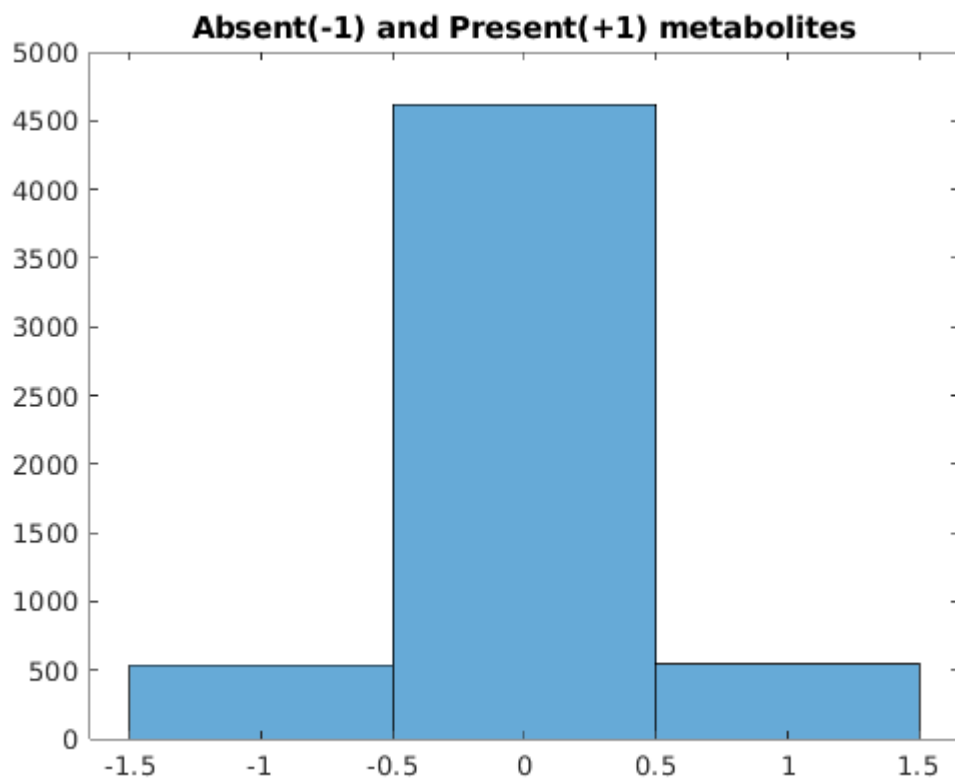
Optimise cardinality reached the stopping criterion. Finished.

100.00% thermodynamically feasible internal fluxes (checked by cycleFreeFlux method).

iter.	nz.flux.%it.feas.int.flux.	%feas.inc.flux.	nz.prod.	%it.feas.nz.prod.	%feas.inc.prod.	fo
1	178	1.00	0.02	121	1.29	0.03
2	208	1.00	0.04	155	1.19	0.10
3	335	1.00	0.07	249	1.20	0.22
4	424	1.00	0.11	301	1.28	0.35
5	393	1.00	0.20	272	1.30	0.42
6	505	1.00	0.29	354	1.29	0.51
7	280	1.00	0.34	208	1.28	0.54
8	378	1.00	0.41	281	1.30	0.58
9	395	1.00	0.45	291	1.31	0.63
10	272	1.00	0.50	211	1.28	0.65
11	281	1.00	0.54	203	1.36	0.67
12	315	1.00	0.59	228	1.36	0.68
13	189	1.00	0.61	134	1.29	0.69
14	358	1.00	0.65	258	1.40	0.70
15	299	1.00	0.68	218	1.28	0.72
16	166	1.00	0.69	116	1.30	0.72
17	275	1.00	0.70	191	1.45	0.73
18	249	1.00	0.71	175	1.45	0.73
19	150	1.00	0.72	107	1.38	0.74
20	235	1.00	0.73	176	1.36	0.74
iter.	nz.flux.%it.feas.int.flux.	%feas.inc.flux.	nz.prod.	%it.feas.nz.prod.	%feas.inc.prod.	fo

thermoKernel terminating early: n = nMax = 20





--- thermoKernel END ----

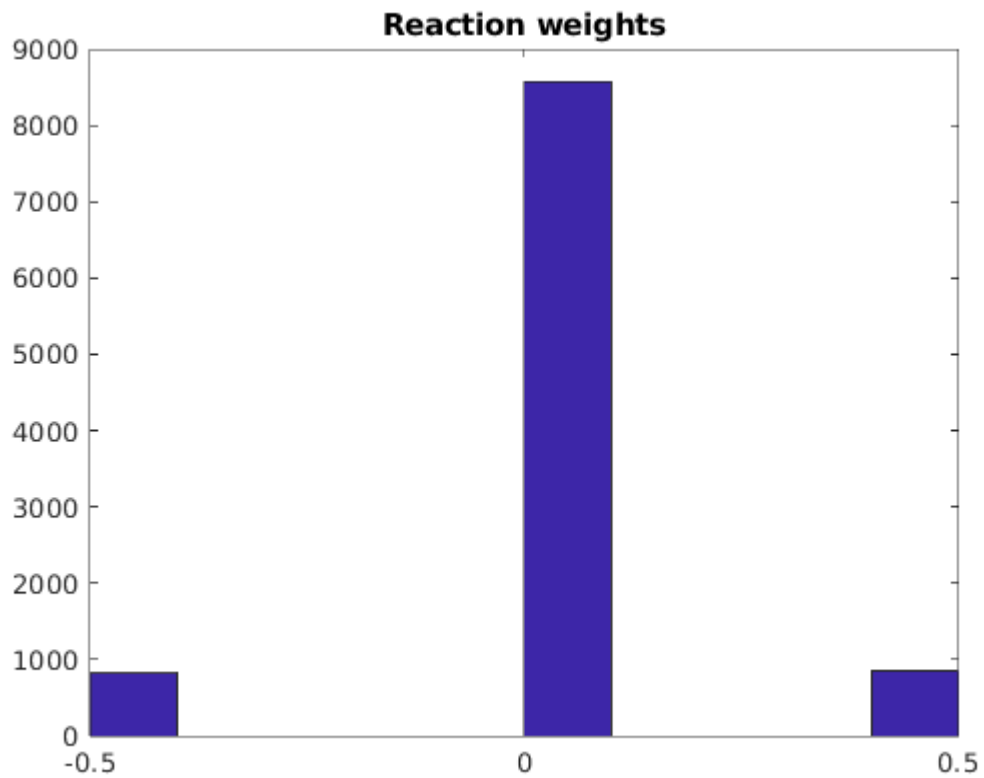
```
[nMet,nRxn]=size(tissueModel.S)
```

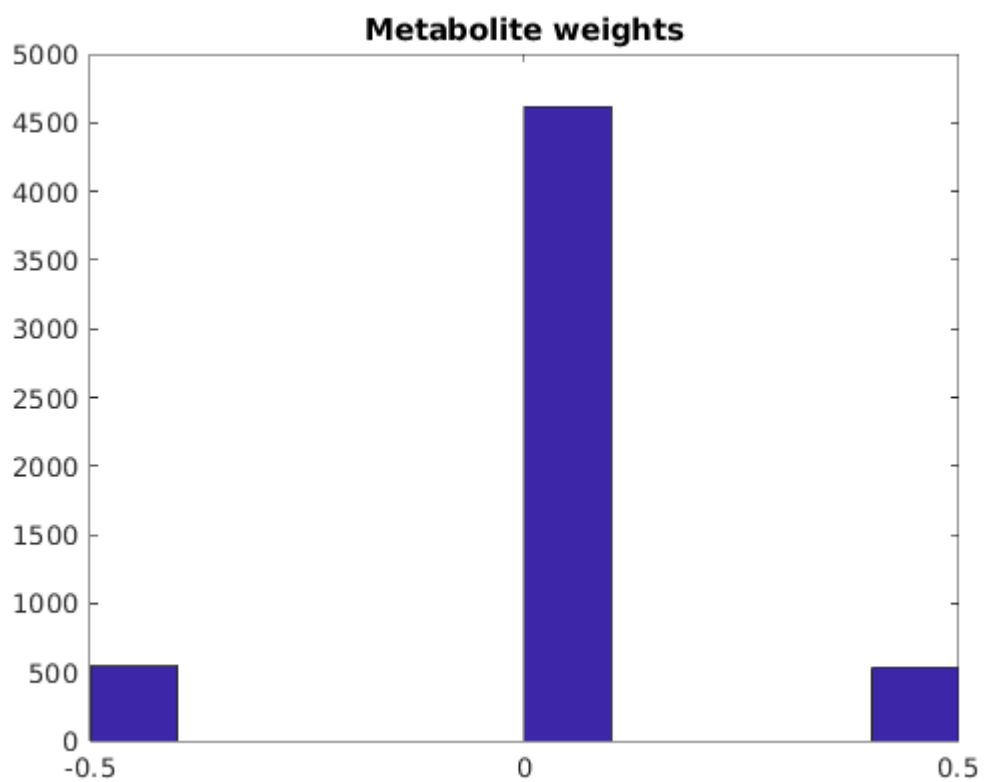
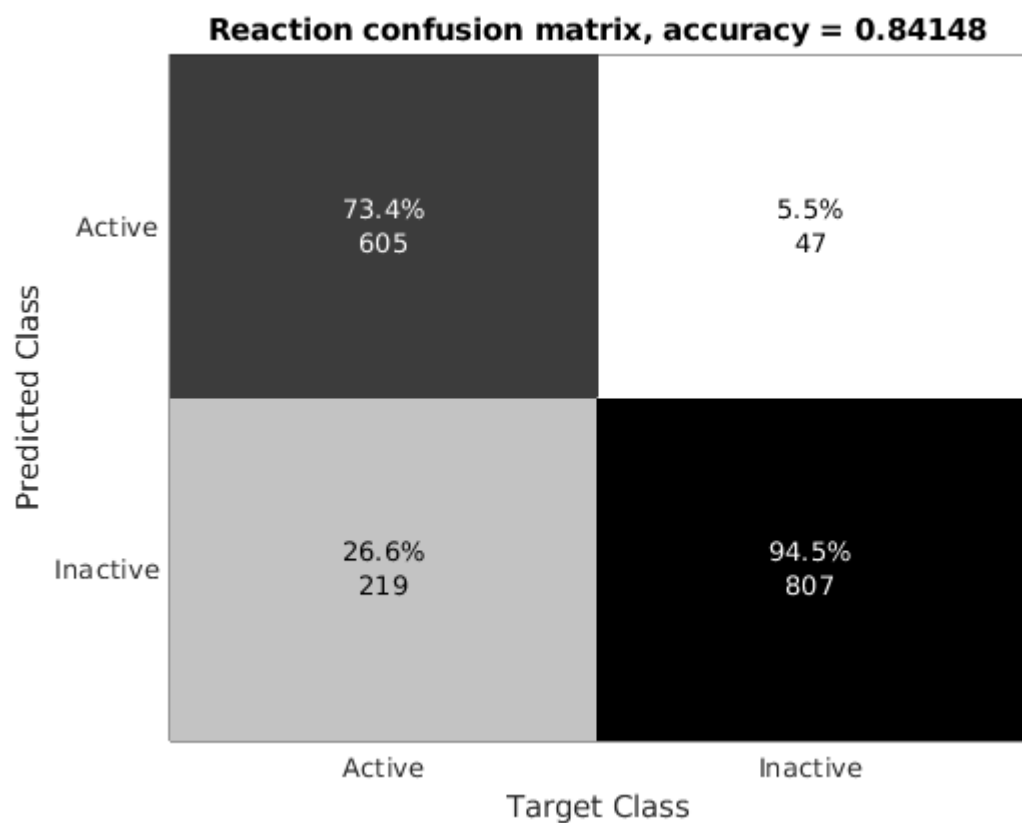


nMet = 2203  
nRxn = 2986

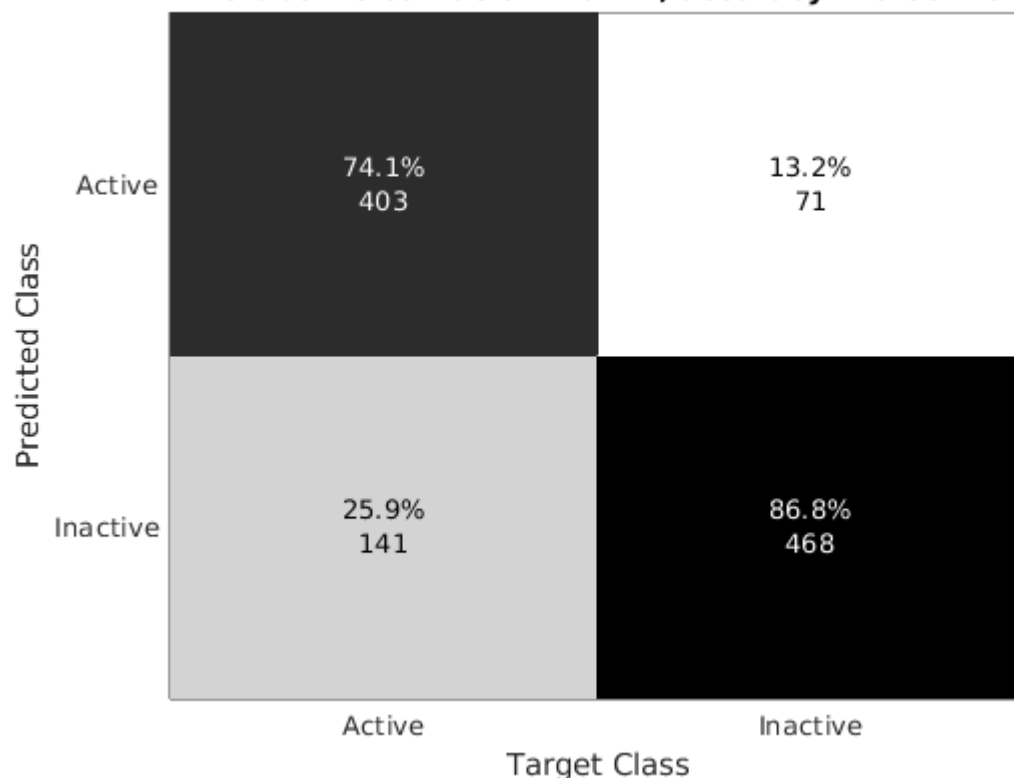
Compare the target versus predicted model

```
%plotThermoCoreStats(activeInactiveRxn, presentAbsentMet,  
thermoModelMetBool, thermoModelRxnBool);  
plotThermoKernelExtractStats(stoichFluxThermoConsistModel,  
activeInactiveRxn, rxnWeights, presentAbsentMet, metWeights,  
thermoModelMetBool, thermoModelRxnBool)
```





**Metabolite confusion matrix, accuracy = 0.80425**



### Save weights

```
rxnWeightsTmp=rxnWeights;
metWeightsTmp=metWeights;
```

### Submodel with just metabolites specified

```
metWeights=metWeightsTmp;
rxnWeights(:)=0;
[tissueModel, thermoModelMetBool, thermoModelRxnBool] =
thermoKernel(stoichFluxThermoConsistModel, activeInactiveRxn, rxnWeights,
presentAbsentMet, metWeights,param);
```

```
--- thermoKernel START ---
thermoKernel parameters:
    printLevel: 1
    relaxBounds: 0
    acceptRepairedFlux: 1
                    n: 200
    normalizeZeroNormWeights: 0
    formulation: 'pqzwrs'
    epsilon: 1.0000e-06
    removeOrphanGenes: 1
    nbMaxIteration: 30
    nMax: 20
    iterationMethod: 'greedyRandom'

    warmStartMethod: 'random'
```

```

        formulation: 'pqzwrns'
        thetaMultiplier: 1.5000
            theta: 0.5000
        regularizeOuter: 1
            epsilon: 1.0000e-06
            printLevel: 1
            relaxBounds: 0
        acceptRepairedFlux: 1
        thermoConsistencyMethod: 'cycleFreeFlux'
            bigNum: 10000
            debug: 0

optCardThermo objective data:
    1 = beta, the global weight on one-norm of internal reaction rate.
    0 = min(g0), the local weight on zero-norm of internal reaction rate.
    0 = max(g0), the local weight on zero-norm of internal reaction rate.
    -0.5 = min(h0), the local weight on zero-norm of metabolite production rate.
    0.5 = max(h0), the local weight on zero-norm of metabolite production rate.

optimizeCardinality objective data:

539 min cardinality variables:
    0 mean(c(p))                0 min(c(p))                0 max(c(p))
    1 lambda0                    0.4 min(k)                0.5 max(k)
    0 lambda1                    0 min(o(p))                0 max(o(p))

1088 max cardinality variables:
    0 mean(c(q))                0 min(c(q))                0 max(c(q))
    1 delta0                    0.4 min(d)                0.5 max(d)
    0 delta1                    0 min(o(q))                0 max(o(q))

36920 cardinality free variables:
    0.46 mean(c(r))             -0 min(c(r))             1 max(c(r))
    0 alpha1                    0 min(o(r))             0 max(o(r))

itn      theta      ||dx||      del_obj      obj      linear      ||x||0      a(x)      ||x||1      ||y||
1        0.50      9.6648e+05      -8.4e+07      7.9        10        2.08529      0.087      0        -7.858
2        0.75        2.4394        -0.69        7.2         9.4        1.67053      0.13      0        -8.287
3        1.12        1.334         -0.47        6.7         9.3        1.23553      0.24      0        -8.287
4        1.69        3.7656         -1          5.7         12         1.23553      0.36      0        -8.287
5        2.53        3.2162        -1.7         4          11         1.23553      0.45      0        -8.737
6        3.80        2.1001        -1.8         2.1         11         1.23553      0.47      0        -10.00
7        5.70        1.3982        -1.2         0.95        10         1.23553      0.51      0        -10.00
8        8.54        0.78275       -1.5        -0.51        9.8        0.832228      0.5       0        -12.
9       12.81        0.52353       -1.8        -2.3         9.5        0.832228      0.55      0        -12.
10      19.22        0.37265       -0.29       -2.6         9.1        0.832228      0.62      0        -12.
11      28.83        0.33297       -0.11       -2.7         8.9        0.832228      0.73      0        -12.
12      43.25        0.19468       -0.5        -3.2         8.7        0.832228      0.83      0        -13.21
13      64.87        0.099656      -0.91       -4.1         8.7        0.832228      0.83      0        -15.03
14      97.31        0.11652       -1.5        -5.6         8.6        0.832228      0.83      0        -15.03
15     145.96        0.12539       -0.03       -5.6         8.6        0.832228      0.83      0        -15.03
16     218.95        0.067216      -0.018      -5.6         8.6        0.832228      0.83      0        -15.03
17     328.42        0.13964      -0.012      -5.6         8.6        0.832228      0.83      0        -15.03
itn      theta      ||dx||      del_obj      obj      linear      ||x||0      a(x)      ||x||1      ||y||
Optimise cardinality reached the stopping criterion. Finished.
100.00% thermodynamically feasible internal fluxes (checked by cycleFreeFlux method).
        warmStartMethod: 'random'
            formulation: 'pqzwrns'
            thetaMultiplier: 1.5000
                theta: 0.5000
            regularizeOuter: 1
                epsilon: 1.0000e-06
                printLevel: 1
                relaxBounds: 0

```

```

acceptRepairedFlux: 1
thermoConsistencyMethod: 'cycleFreeFlux'
    bigNum: 10000
    debug: 0

```

optCardThermo objective data:

```

1 = beta, the global weight on one-norm of internal reaction rate.
0 = min(g0), the local weight on zero-norm of internal reaction rate.
0 = max(g0), the local weight on zero-norm of internal reaction rate.
-0.5 = min(h0), the local weight on zero-norm of metabolite production rate.
0.5 = max(h0), the local weight on zero-norm of metabolite production rate.

```

optimizeCardinality objective data:

539 min cardinality variables:

```

0 mean(c(p))          0 min(c(p))          0 max(c(p))
1 lambda0             0.4 min(k)            0.5 max(k)
0 lambda1             0 min(o(p))          0 max(o(p))

```

1088 max cardinality variables:

```

0 mean(c(q))          0 min(c(q))          0 max(c(q))
1 delta0             0.4 min(d)            0.5 max(d)
0 delta1             0 min(o(q))          0 max(o(q))

```

36920 cardinality free variables:

```

0.46 mean(c(r))      -0 min(c(r))          1 max(c(r))
0 alpha1             0 min(o(r))          0 max(o(r))

```

itn	theta	dx	del_obj	obj	linear	x  0	a(x)	x  1	y
1	0.50	9.7116e+05	-8.5e+07	7.9	10	2.08529	0.079	0	-7.121
2	0.75	2.4188	-0.68	7.2	9.5	1.23553	0.11	0	-7.551
3	1.12	2.6941	-0.72	6.5	10	1.23553	0.24	0	-8.353
4	1.69	4.9777	-0.99	5.5	11	1.23553	0.36	0	-8.353
5	2.53	4.4276	-1.7	3.8	11	1.23553	0.45	0	-8.802
6	3.80	2.3513	-2.2	1.6	11	1.23553	0.47	0	-10.52
7	5.70	1.6188	-1.2	0.4	10	1.23553	0.51	0	-10.52
8	8.54	0.63996	-0.59	-0.19	9.6	1.23553	0.56	0	-10.52
9	12.81	0.55001	-0.9	-1.1	9.4	0.832228	0.55	0	-12.35
10	19.22	0.4508	-1.5	-2.6	9.1	0.832228	0.62	0	-12.35
11	28.83	0.42169	-0.11	-2.7	8.9	0.832228	0.73	0	-12.35
12	43.25	0.24485	-0.032	-2.8	8.7	0.832228	0.83	0	-12.35
13	64.87	0.19244	-0.075	-2.9	8.7	0.832228	0.83	0	-12.35
14	97.31	0.10717	-0.042	-2.9	8.6	0.832228	0.83	0	-12.35
15	145.96	0.083129	-0.023	-2.9	8.6	0.832228	0.83	0	-12.35
16	218.95	0.070749	-0.015	-2.9	8.6	0.832228	0.83	0	-12.35
17	328.42	0.14835	-0.0098	-2.9	8.6	0.832228	0.83	0	-12.35
itn	theta	dx	del_obj	obj	linear	x  0	a(x)	x  1	y

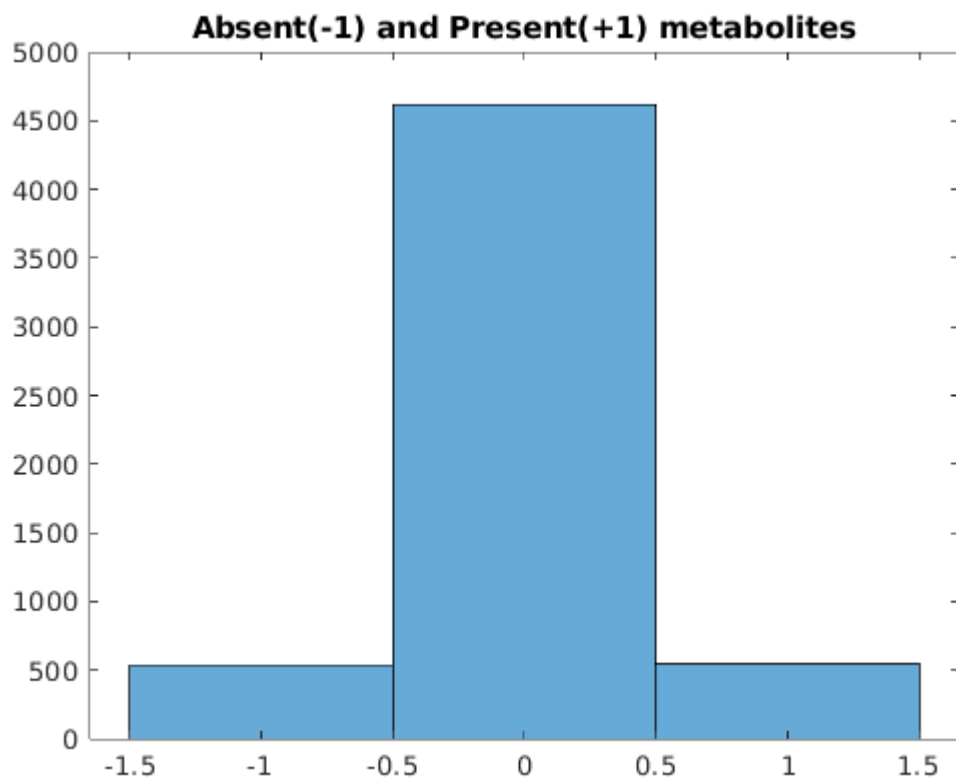
Optimise cardinality reached the stopping criterion. Finished.

100.00% thermodynamically feasible internal fluxes (checked by cycleFreeFlux method).

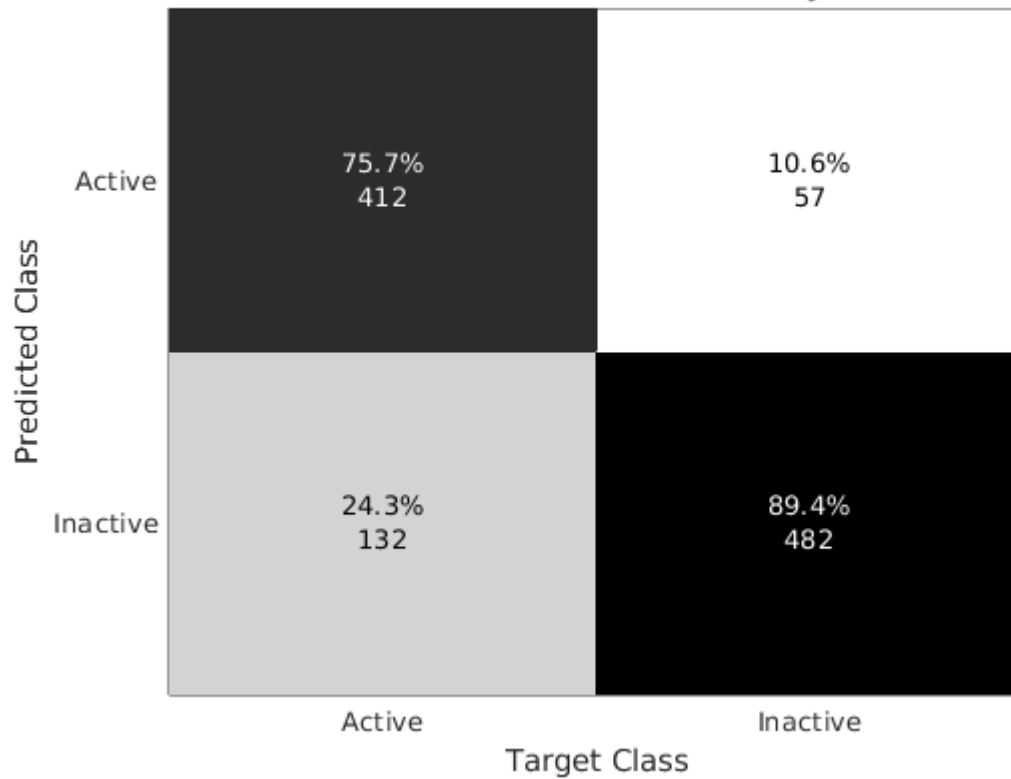
iter.	nz.flux.	%it.feas.int.flux.	%feas.inc.flux.	nz.prod.	%it.feas.nz.prod.	%feas.inc.prod.	fo
1	137	1.00	NaN	95	1.21	0.03	gre
2	186	1.00	NaN	139	1.17	0.08	gre
3	297	1.00	NaN	226	1.19	0.20	gre
4	467	1.00	NaN	349	1.30	0.37	gre
5	203	1.00	NaN	149	1.23	0.43	gre
6	282	1.00	NaN	213	1.24	0.51	gre
7	285	1.00	NaN	201	1.32	0.56	gre
8	213	1.00	NaN	156	1.30	0.60	gre
9	168	1.00	NaN	124	1.27	0.62	gre
10	132	1.00	NaN	97	1.20	0.63	gre
11	248	1.00	NaN	184	1.33	0.66	gre
12	111	1.00	NaN	78	1.22	0.66	gre
13	200	1.00	NaN	148	1.36	0.68	gre
14	112	1.00	NaN	77	1.25	0.68	gre

15	984	1.00	NaN	709	1.26	0.70
16	1076	1.00	NaN	789	1.24	0.74
17	1024	1.00	NaN	744	1.27	0.74
18	1000	1.00	NaN	726	1.26	0.75
19	974	1.00	NaN	732	1.22	0.75
20	965	1.00	NaN	718	1.24	0.76

iter.    nz.flux.%it.feas.int.flux.    %feas.inc.flux.    nz.prod.    %it.feas.nz.prod.    %feas.inc.prod.    fo  
 thermoKernel terminating early: n = nMax = 20



**Metabolite confusion matrix, accuracy = 0.82548**



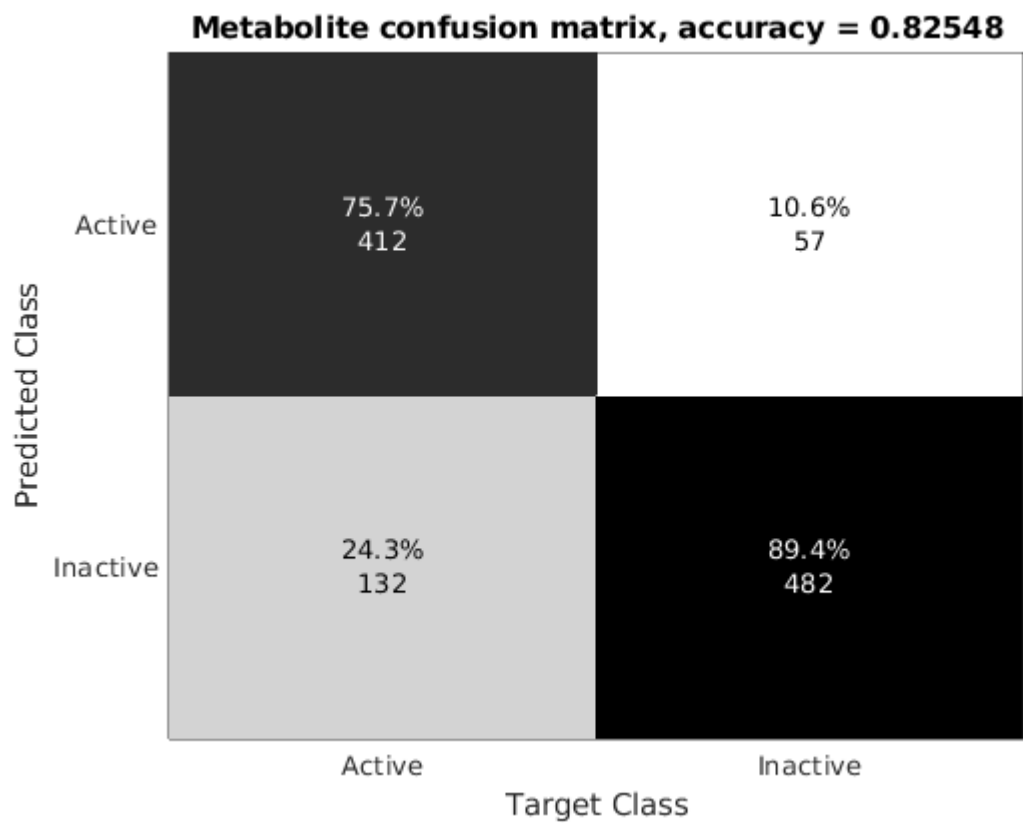
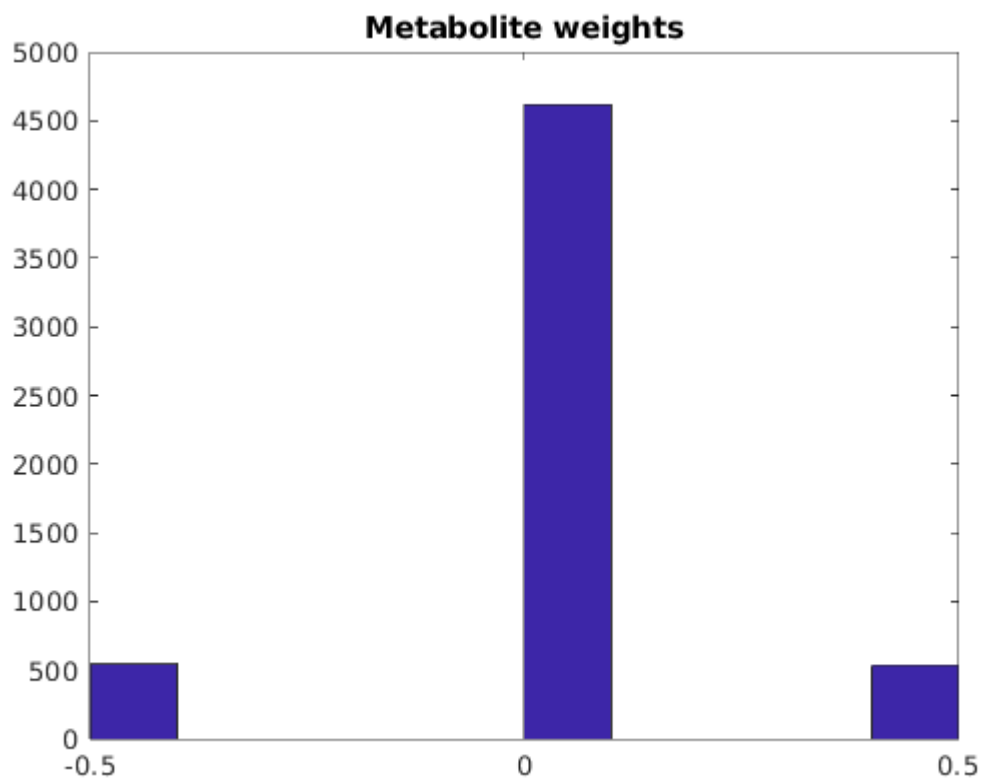
--- thermoKernel END ----

```
[nMet,nRxn]=size(tissueModel.S)
```

```
nMet = 1870
nRxn = 2635
```

Compare the target versus predicted model

```
plotThermoKernelExtractStats(stoichFluxThermoConsistModel,
activeInactiveRxn, rxnWeights, presentAbsentMet, metWeights,
thermoModelMetBool, thermoModelRxnBool)
```



### Submodel with just reactions specified

```
rxnWeights=rxnWeightsTmp;
```



```
metWeights(:)=0;
[tissueModel, thermoModelMetBool, thermoModelRxnBool] =
thermoKernel(stoichFluxThermoConsistModel, activeInactiveRxn, rxnWeights,
presentAbsentMet, metWeights,param);
```

```
--- thermoKernel START ----
thermoKernel parameters:
    printLevel: 1
    relaxBounds: 0
    acceptRepairedFlux: 1
        n: 200
    normalizeZeroNormWeights: 0
        formulation: 'pqzwrns'
        epsilon: 1.0000e-06
    removeOrphanGenes: 1
        nbMaxIteration: 30
        nMax: 20
        iterationMethod: 'greedyRandom'

    warmStartMethod: 'random'
        formulation: 'pqzwrns'
    thetaMultiplier: 1.5000
        theta: 0.5000
    regularizeOuter: 1
        epsilon: 1.0000e-06
        printLevel: 1
        relaxBounds: 0
    acceptRepairedFlux: 1
    thermoConsistencyMethod: 'cycleFreeFlux'
        bigNum: 10000
        debug: 0
```

```
optCardThermo objective data:
    1 = beta, the global weight on one-norm of internal reaction rate.
    -0.5 = min(g0), the local weight on zero-norm of internal reaction rate.
    0.5 = max(g0), the local weight on zero-norm of internal reaction rate.
    0 = min(h0), the local weight on zero-norm of metabolite production rate.
    0 = max(h0), the local weight on zero-norm of metabolite production rate.
```

```
optimizeCardinality objective data:
```

```
854 min cardinality variables:
    -0 mean(c(p))          -0 min(c(p))          -0 max(c(p))
    1 lambda0              0.4 min(k)              0.5 max(k)
    0 lambda1              0 min(o(p))              0 max(o(p))
```

```
824 max cardinality variables:
    -0 mean(c(q))          -0 min(c(q))          -0 max(c(q))
    1 delta0              0.4 min(d)              0.5 max(d)
    0 delta1              0 min(o(q))              0 max(o(q))
```

```
36869 cardinality free variables:
    0.46 mean(c(r))        -0 min(c(r))          1 max(c(r))
    0 alpha1              0 min(o(r))          0 max(o(r))
```

itn	theta	dx	del_obj	obj	linear	x  0	a(x)	x  1	y
1	0.50	9.6945e+05	-8.5e+07	8.4	8.4	0.408362	0.02	0	-3.084
2	0.75	0.082294	-0.032	8.4	8.4	0.408362	0.028	0	-3.506
3	1.12	0.25722	-0.046	8.3	8.5	0.408362	0.03	0	-3.506
4	1.69	0.41187	-0.093	8.2	8.5	0.422161	0.0076	0	-3.506
5	2.53	5.4552e-14	-0.15	8.1	8.5	0.422161	0.011	0	-3.506
itn	theta	dx	del_obj	obj	linear	x  0	a(x)	x  1	y

Optimise cardinality reached the stopping criterion. Finished.

100.00% thermodynamically feasible internal fluxes (checked by cycleFreeFlux method).

```
warmStartMethod: 'random'
formulation: 'pqzwrns'
thetaMultiplier: 1.5000
theta: 0.5000
regularizeOuter: 1
epsilon: 1.0000e-06
printLevel: 1
relaxBounds: 0
acceptRepairedFlux: 1
thermoConsistencyMethod: 'cycleFreeFlux'
bigNum: 10000
debug: 0
```

optCardThermo objective data:

```
1 = beta, the global weight on one-norm of internal reaction rate.
-0.5 = min(g0), the local weight on zero-norm of internal reaction rate.
0.5 = max(g0), the local weight on zero-norm of internal reaction rate.
0 = min(h0), the local weight on zero-norm of metabolite production rate.
0 = max(h0), the local weight on zero-norm of metabolite production rate.
```

optimizeCardinality objective data:

854 min cardinality variables:

```
-0 mean(c(p))          -0 min(c(p))          -0 max(c(p))
1 lambda0              0.4 min(k)              0.5 max(k)
0 lambda1              0 min(o(p))              0 max(o(p))
```

824 max cardinality variables:

```
-0 mean(c(q))          -0 min(c(q))          -0 max(c(q))
1 delta0              0.4 min(d)              0.5 max(d)
0 delta1              0 min(o(q))              0 max(o(q))
```

36869 cardinality free variables:

```
0.46 mean(c(r))        -0 min(c(r))          1 max(c(r))
0 alpha1              0 min(o(r))          0 max(o(r))
```

itn	theta	dx	del_obj	obj	linear	x  0	a(x)	x  1	y  1
1	0.50	9.6798e+05	-8.5e+07	8.4	8.4	0.830523	0.02	0	-3.506
2	0.75	0.11192	-0.032	8.4	8.4	0.408362	0.028	0	-3.506
3	1.12	0.25722	-0.046	8.3	8.5	0.408362	0.03	0	-3.506
4	1.69	0.41187	-0.093	8.2	8.5	0.422161	0.0076	0	-3.506
5	2.53	5.4552e-14	-0.15	8.1	8.5	0.422161	0.011	0	-3.506
itn	theta	dx	del_obj	obj	linear	x  0	a(x)	x  1	y  1

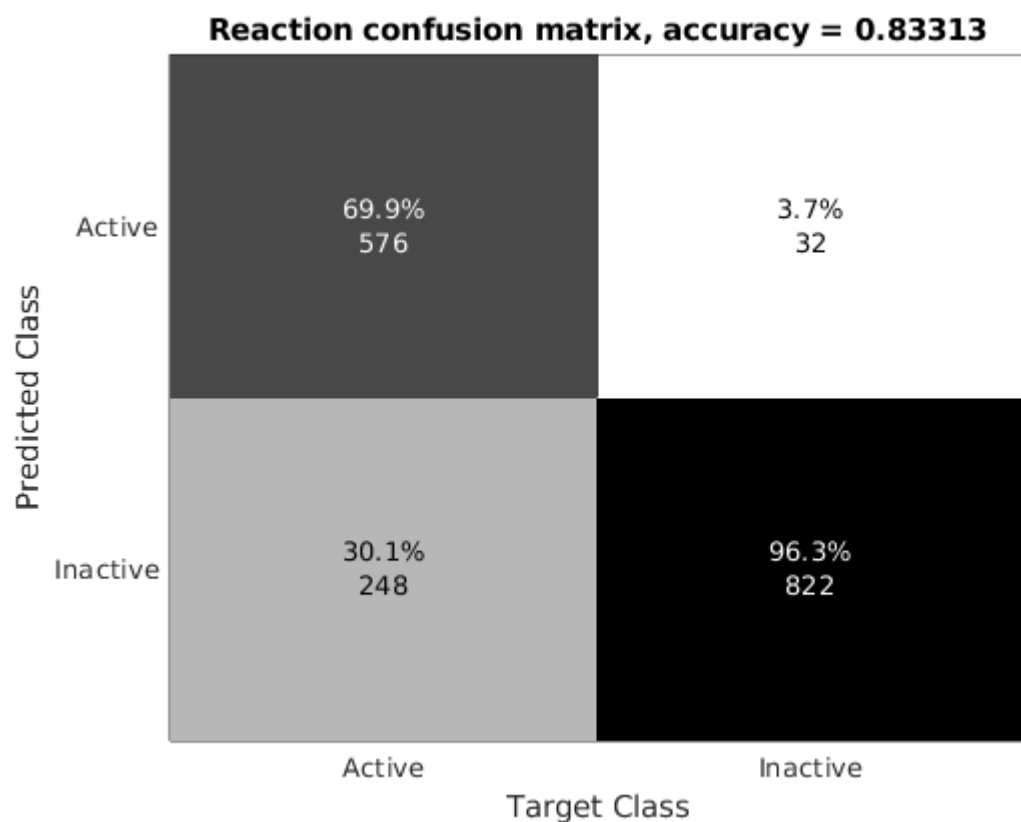
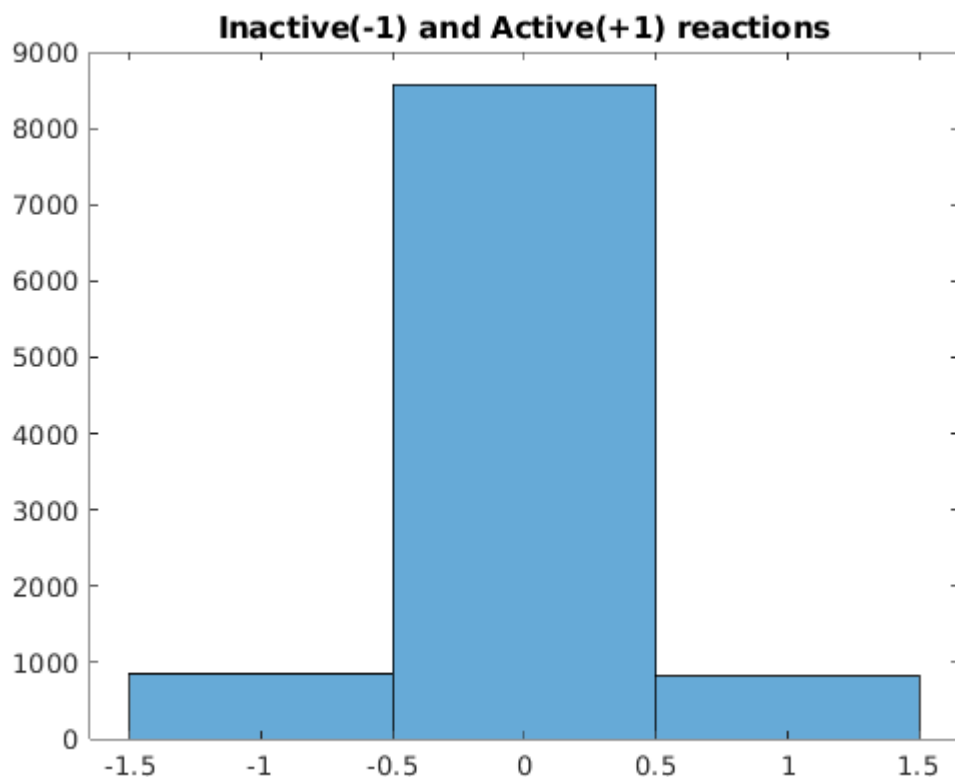
Optimise cardinality reached the stopping criterion. Finished.

100.00% thermodynamically feasible internal fluxes (checked by cycleFreeFlux method).

iter.	nz.flux.	%it.feas.int.flux.	%feas.inc.flux.	nz.prod.	%it.feas.nz.prod.	%feas.inc.prod.	fo
1	100	1.00	0.01	72	1.15	NaN	gre
2	99	1.00	0.01	70	1.21	NaN	gre
3	113	1.00	0.02	79	1.15	NaN	gre
4	113	1.00	0.03	83	1.12	NaN	gre
5	181	1.00	0.07	134	1.14	NaN	gre
6	240	1.00	0.14	183	1.14	NaN	gre
7	192	1.00	0.20	128	1.38	NaN	gre
8	184	1.00	0.24	130	1.31	NaN	gre
9	276	1.00	0.30	192	1.38	NaN	gre
10	267	1.00	0.35	193	1.28	NaN	gre
11	309	1.00	0.42	227	1.29	NaN	gre
12	293	1.00	0.49	209	1.44	NaN	gre
13	295	1.00	0.54	205	1.39	NaN	gre
14	214	1.00	0.57	173	1.25	NaN	gre
15	340	1.00	0.62	249	1.37	NaN	gre
16	284	1.00	0.66	203	1.37	NaN	gre
17	177	1.00	0.67	122	1.42	NaN	gre

18	116	1.00	0.67	87	1.20	NaN	gre
19	147	1.00	0.68	101	1.40	NaN	gre
20	199	1.00	0.70	139	1.42	NaN	gre

iter. nz.flux.%it.feas.int.flux. %feas.inc.flux. nz.prod. %it.feas.nz.prod. %feas.inc.prod. fo  
 thermoKernel terminating early: n = nMax = 20



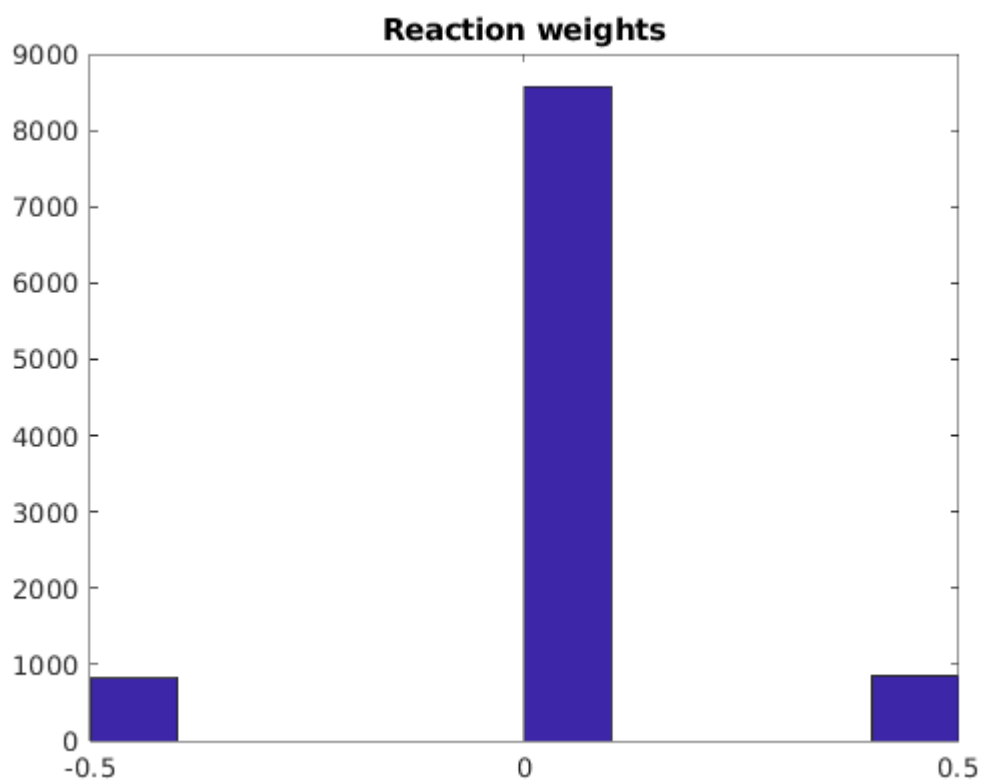
```
--- thermoKernel END ----
```

```
[nMet,nRxn]=size(tissueModel.S)
```

```
nMet = 1441  
nRxn = 1924
```

Compare the target versus predicted model

```
plotThermoKernelExtractStats(stoichFluxThermoConsistModel,  
activeInactiveRxn, rxnWeights, presentAbsentMet, metWeights,  
thermoModelMetBool, thermoModelRxnBool)
```



**Reaction confusion matrix, accuracy = 0.83313**



## Submodel with just active metabolites specified

```
metWeightsRed=metWeightsTmp;  
rxnWeightsRed=rxnWeightsTmp*0;  
metWeightsRed(metWeightsRed>=0)=0;
```

```
[tissueModel, thermoModelMetBool, thermoModelRxnBool] =  
thermoKernel(stoichFluxThermoConsistModel, activeInactiveRxn, rxnWeightsRed,  
presentAbsentMet, metWeightsRed,param);
```

```
--- thermoKernel START ----  
thermoKernel parameters:  
    printLevel: 1  
    relaxBounds: 0  
    acceptRepairedFlux: 1  
        n: 200  
    normalizeZeroNormWeights: 0  
        formulation: 'pqzwrS'  
        epsilon: 1.0000e-06  
    removeOrphanGenes: 1  
        nbMaxIteration: 30  
        nMax: 20  
    iterationMethod: 'greedyRandom'  
  
    warmStartMethod: 'random'  
        formulation: 'pqzwrS'
```

```

thetaMultiplier: 1.5000
    theta: 0.5000
regularizeOuter: 1
    epsilon: 1.0000e-06
    printLevel: 1
    relaxBounds: 0
acceptRepairedFlux: 1
thermoConsistencyMethod: 'cycleFreeFlux'
    bigNum: 10000
    debug: 0

optCardThermo objective data:
    1 = beta, the global weight on one-norm of internal reaction rate.
    0 = min(g0), the local weight on zero-norm of internal reaction rate.
    0 = max(g0), the local weight on zero-norm of internal reaction rate.
    -0.5 = min(h0), the local weight on zero-norm of metabolite production rate.
    0 = max(h0), the local weight on zero-norm of metabolite production rate.

optimizeCardinality objective data:

0 min cardinality variables:
    NaN mean(c(p))          NaN min(c(p))          NaN max(c(p))
    1 lambda0              NaN min(k)              NaN max(k)
    0 lambda1              NaN min(o(p))            NaN max(o(p))

1088 max cardinality variables:
    0 mean(c(q))            0 min(c(q))            0 max(c(q))
    1 delta0                0.4 min(d)              0.5 max(d)
    0 delta1                0 min(o(q))            0 max(o(q))

37459 cardinality free variables:
    0.45 mean(c(r))         -0 min(c(r))         1 max(c(r))
    0 alfa1                 0 min(o(r))         0 max(o(r))

itn   theta   ||dx||   del_obj   obj   linear   ||x||0   a(x)   ||x||1   ||y||
1     0.50   9.6587e+05 -8.5e+07   7.4     13        0        0        0   -14.77
2     0.75   5.8717    -2.6     4.8     14        0        0        0   -14.32
3     1.12   3.0027    -2.1     2.7     13        0        0        0   -15.21
4     1.69   5.6907    -2.2     0.55    13        0        0        0   -17.00
5     2.53   5.2054    -4.5    -3.9     14        0        0        0   -20.17
6     3.80   2.462     -3.4    -7.3     13        0        0        0   -21.59
7     5.70   1.606     -2.7    -10     12        0        0        0   -22.5
8     8.54   1.0881    -1.7    -12     11        0        0        0   -22.5
9    12.81   0.80985   -0.89   -13     10        0        0        0   -22.5
10   19.22   0.49523   -0.58   -13     9.4        0        0        0   -22.5
11   28.83   0.38668   -0.36   -13        9        0        0        0   -22.5
12   43.25   0.28276   -0.38   -14     8.8        0        0        0   -23.47
13   64.87   0.21412   -0.94   -15     8.7        0        0        0   -23.47
14   97.31   0.16629   -0.083  -15     8.6        0        0        0   -23.47
15  145.96   0.084169  -0.26   -15     8.5        0        0        0   -25.28
16  218.95   0.12117   -1.6    -17     8.5        0        0        0   -25.28
17  328.42   0.16767   -0.14   -17     8.5        0        0        0   -25.74
itn   theta   ||dx||   del_obj   obj   linear   ||x||0   a(x)   ||x||1   ||y||
Optimise cardinality reached the stopping criterion. Finished.
100.00% thermodynamically feasible internal fluxes (checked by cycleFreeFlux method).
    warmStartMethod: 'random'
    formulation: 'pqzwrS'
    thetaMultiplier: 1.5000
    theta: 0.5000
regularizeOuter: 1
    epsilon: 1.0000e-06
    printLevel: 1
    relaxBounds: 0
acceptRepairedFlux: 1

```

```

thermoConsistencyMethod: 'cycleFreeFlux'
    bigNum: 10000
    debug: 0

optCardThermo objective data:
    1 = beta, the global weight on one-norm of internal reaction rate.
    0 = min(g0), the local weight on zero-norm of internal reaction rate.
    0 = max(g0), the local weight on zero-norm of internal reaction rate.
    -0.5 = min(h0), the local weight on zero-norm of metabolite production rate.
    0 = max(h0), the local weight on zero-norm of metabolite production rate.

optimizeCardinality objective data:

0 min cardinality variables:
    NaN mean(c(p))          NaN min(c(p))          NaN max(c(p))
    1 lambda0              NaN min(k)              NaN max(k)
    0 lambda1              NaN min(o(p))            NaN max(o(p))

1088 max cardinality variables:
    0 mean(c(q))            0 min(c(q))            0 max(c(q))
    1 delta0                0.4 min(d)              0.5 max(d)
    0 delta1                0 min(o(q))            0 max(o(q))

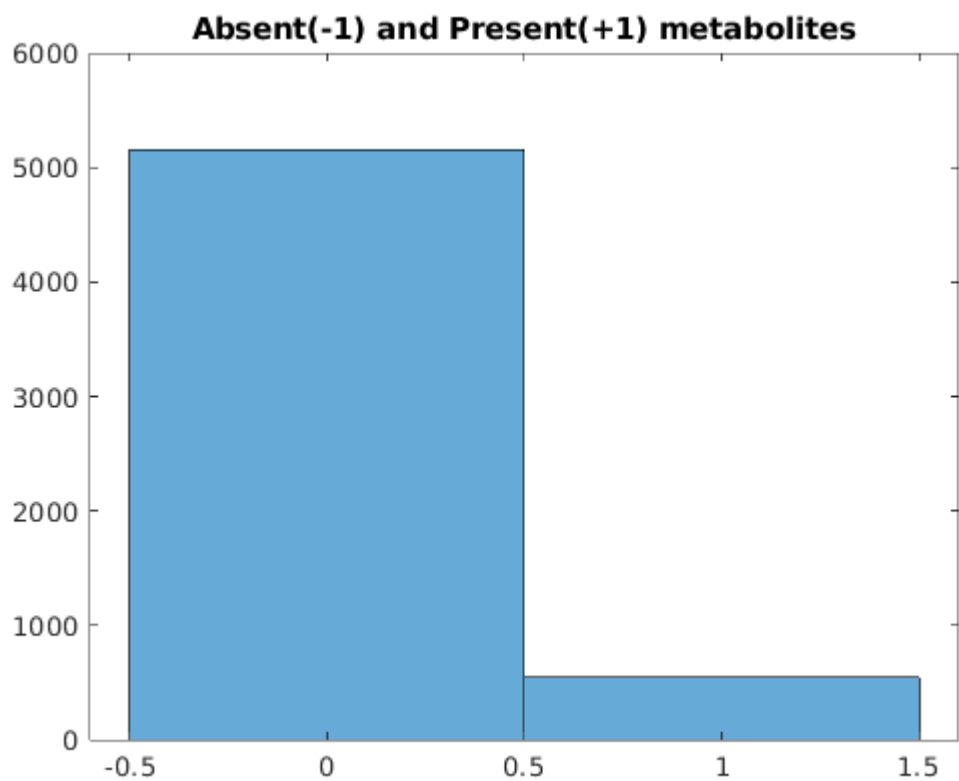
37459 cardinality free variables:
    0.45 mean(c(r))         -0 min(c(r))          1 max(c(r))
    0 alpha1                0 min(o(r))          0 max(o(r))

itn      theta      ||dx||      del_obj      obj      linear      ||x||0      a(x)      ||x||1      ||y||
1        0.50      9.6895e+05      -8.5e+07      8.1        9          0          0          0          -6.131
2        0.75        1.691        -0.42        7.7        8.8        0          0          0          -6.561
3        1.12        2.8196       -0.42        7.3        10        0          0          0          -6.561
4        1.69        3.3125       -0.91        6.4        11        0          0          0          -6.561
5        2.53        3.3113       -2.3         4.1        11        0          0          0          -7.845
6        3.80        1.408        -1           3          11        0          0          0          -8.276
7        5.70        1.4232       -1.5         1.5        10        0          0          0          -8.728
8        8.54        0.75486      -0.82        0.71       9.6        0          0          0          -9.635
9       12.81        0.5787       -1.1        -0.43      9.2        0          0          0          -9.635
10      19.22        0.45877      -0.29       -0.72      8.9        0          0          0          -9.635
11      28.83        0.19759      -0.19       -0.91      8.7        0          0          0          -9.635
12      43.25        0.17445      -0.12        -1         8.6        0          0          0          -9.635
13      64.87        0.1615       -0.44       -1.5       8.5        0          0          0          -11.45
14      97.31        0.14097      -1.5        -2.9       8.5        0          0          0          -11.45
15     145.96        0.12588      -0.026       -3         8.5        0          0          0          -11.45
16     218.95        0.086138     -0.015       -3         8.5        0          0          0          -11.45
17     328.42        0.074785     -0.01        -3         8.5        0          0          0          -11.45
itn      theta      ||dx||      del_obj      obj      linear      ||x||0      a(x)      ||x||1      ||y||
Optimise cardinality reached the stopping criterion. Finished.
100.00% thermodynamically feasible internal fluxes (checked by cycleFreeFlux method).
iter.  nz.flux.%it.feas.int.flux.  %feas.inc.flux.  nz.prod.%it.feas.nz.prod.  %feas.inc.prod.  fc
1      122          1.00          NaN          86          1.16          0.02          gre
2      230          1.00          NaN          172         1.16          0.10          gre
3      231          1.00          NaN          183         1.18          0.18          gre
4      352          1.00          NaN          263         1.22          0.31          gre
5      283          1.00          NaN          206         1.32          0.40          gre
6      274          1.00          NaN          199         1.37          0.47          gre
7      275          1.00          NaN          201         1.34          0.54          gre
8      175          1.00          NaN          139         1.19          0.56          gre
9      130          1.00          NaN          97          1.20          0.58          gre
10     307          1.00          NaN          235         1.34          0.62          gre
11     243          1.00          NaN          183         1.31          0.65          gre
12     120          1.00          NaN          84          1.29          0.65          gre
13     218          1.00          NaN          158         1.32          0.67          gre
14     113          1.00          NaN          80          1.24          0.67          gre
15     131          1.00          NaN          99          1.30          0.68          gre

```

16	129	1.00	NaN	92	1.30	0.69	gre
17	164	1.00	NaN	116	1.30	0.69	gre
18	166	1.00	NaN	123	1.26	0.70	gre
19	165	1.00	NaN	117	1.35	0.71	gre
20	141	1.00	NaN	95	1.36	0.72	gre

iter.    nz.flux.%it.feas.int.flux.    %feas.inc.flux.    nz.prod.    %it.feas.nz.prod.    %feas.inc.prod.    fo  
 thermoKernel terminating early: n = nMax = 20





**Metabolite confusion matrix, accuracy = 0.71875**



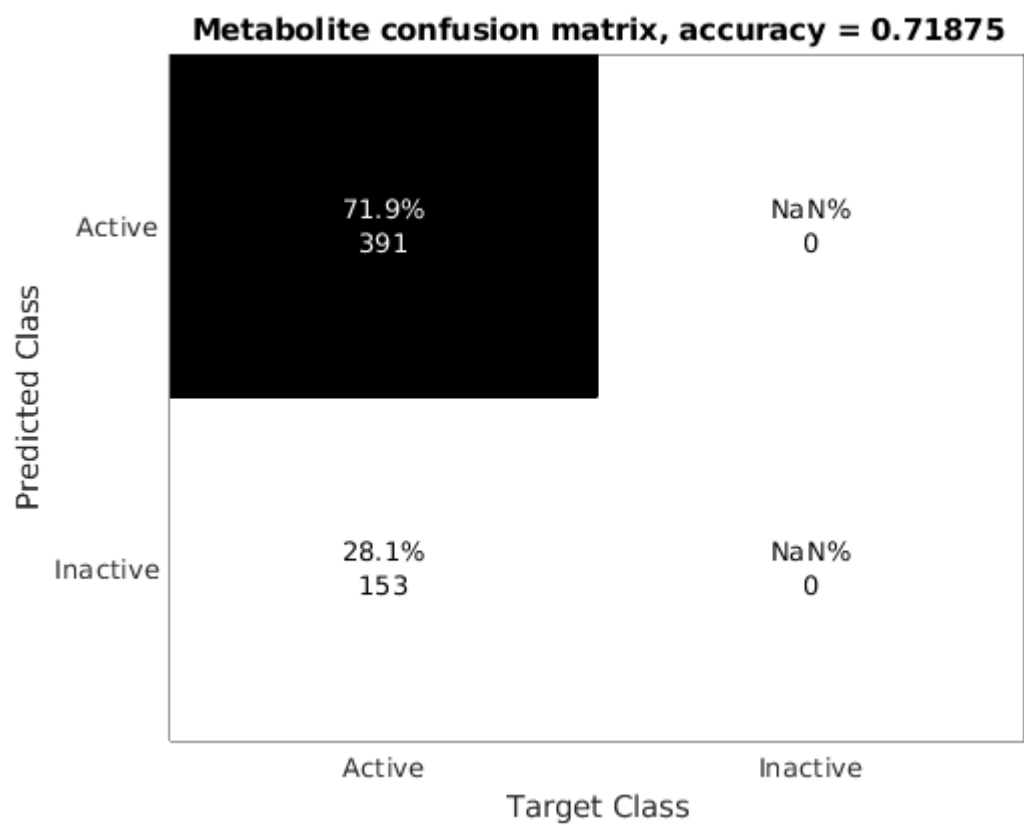
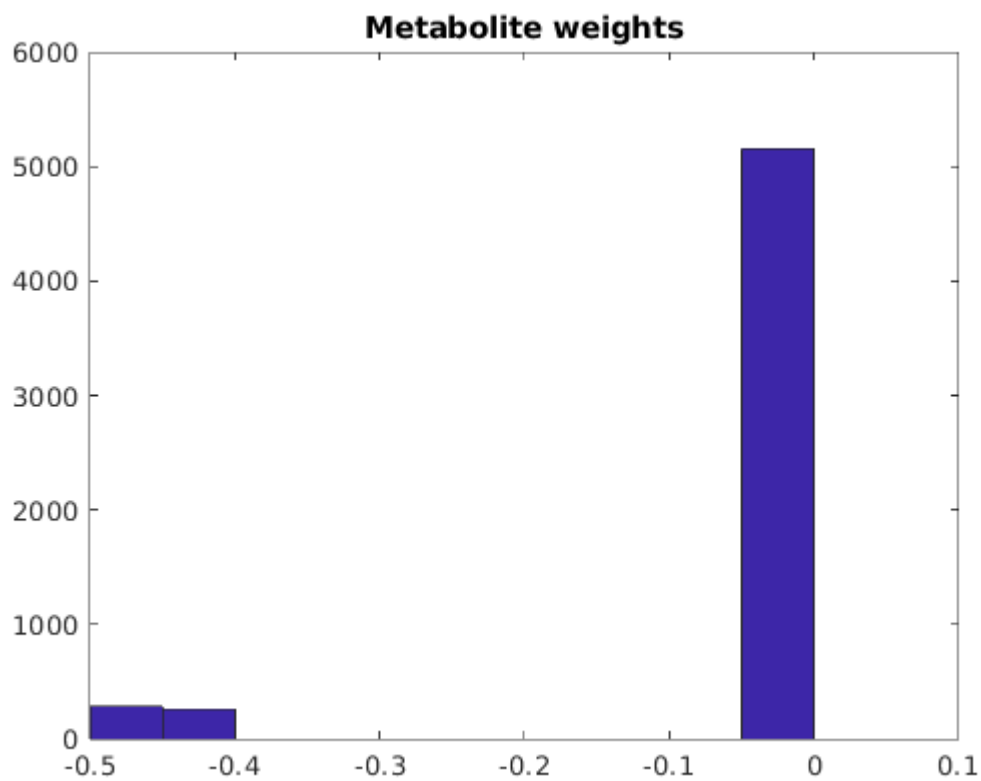
--- thermoKernel END ----

```
[nMet,nRxn]=size(tissueModel.S)
```

```
nMet = 1448
nRxn = 1753
```

Compare the target versus predicted model

```
plotThermoKernelExtractStats(stoichFluxThermoConsistModel,
activeInactiveRxn, rxnWeightsRed, presentAbsentMet, metWeightsRed,
thermoModelMetBool, thermoModelRxnBool)
```



**Submodel with just active reactions specified**

```

rxnWeightsRed=rxnWeightsTmp;
metWeightsRed=metWeightsTmp*0;
rxnWeightsRed(rxnWeightsRed>=0)=0;
[tissueModel, thermoModelMetBool, thermoModelRxnBool] =
thermoKernel(stoichFluxThermoConsistModel, activeInactiveRxn, rxnWeightsRed,
presentAbsentMet, metWeightsRed,param);

```

```

--- thermoKernel START ----
thermoKernel parameters:
    printLevel: 1
    relaxBounds: 0
    acceptRepairedFlux: 1
        n: 200
normalizeZeroNormWeights: 0
    formulation: 'pqzwrS'
    epsilon: 1.0000e-06
removeOrphanGenes: 1
    nbMaxIteration: 30
        nMax: 20
    iterationMethod: 'greedyRandom'

    warmStartMethod: 'random'
        formulation: 'pqzwrS'
    thetaMultiplier: 1.5000
        theta: 0.5000
regularizeOuter: 1
    epsilon: 1.0000e-06
    printLevel: 1
    relaxBounds: 0
    acceptRepairedFlux: 1
thermoConsistencyMethod: 'cycleFreeFlux'
    bigNum: 10000
    debug: 0

```

optCardThermo objective data:

```

1 = beta, the global weight on one-norm of internal reaction rate.
-0.5 = min(g0), the local weight on zero-norm of internal reaction rate.
0 = max(g0), the local weight on zero-norm of internal reaction rate.
-0 = min(h0), the local weight on zero-norm of metabolite production rate.
-0 = max(h0), the local weight on zero-norm of metabolite production rate.

```

optimizeCardinality objective data:

0 min cardinality variables:

NaN mean(c(p))	NaN min(c(p))	NaN max(c(p))
1 lambda0	NaN min(k)	NaN max(k)
0 lambda1	NaN min(o(p))	NaN max(o(p))

824 max cardinality variables:

-0 mean(c(q))	-0 min(c(q))	-0 max(c(q))
1 delta0	0.4 min(d)	0.5 max(d)
0 delta1	0 min(o(q))	0 max(o(q))

37723 cardinality free variables:

0.45 mean(c(r))	-0 min(c(r))	1 max(c(r))
0 alpha1	0 min(o(r))	0 max(o(r))

itn	theta	dx	del_obj	obj	linear	x  0	a(x)	x  1	y
1	0.50	9.6384e+05	-8.4e+07	8.4	8.5	0	0	0	-3.118
2	0.75	1.1692	-0.091	8.3	8.4	0	0	0	-3.548
3	1.12	0.15183	-0.062	8.3	8.4	0	0	0	-3.506
4	1.69	0.092443	-0.084	8.2	8.4	0	0	0	-3.506

5	2.53	0.44167	-0.14	8.1	8.5	0	0	0	-3.506
6	3.80	0.31473	-0.25	7.8	8.5	0	0	0	-3.956
7	5.70	1.8987	-0.76	7	9.6	0	0	0	-4.903
8	8.54	1.0204	-1.2	5.8	9.4	0	0	0	-5.394
9	12.81	1.4536	-1.1	4.8	10	0	0	0	-5.832
10	19.22	0.67982	-0.85	3.9	9.8	0	0	0	-5.832
11	28.83	0.60377	-0.54	3.4	9.2	0	0	0	-5.832
12	43.25	0.45629	-0.35	3	8.9	0	0	0	-5.832
13	64.87	0.57354	-0.22	2.8	8.7	0	0	0	-5.832
14	97.31	0.4286	-0.13	2.7	8.5	0	0	0	-5.832
15	145.96	0.145	-0.036	2.7	8.5	0	0	0	-5.832
16	218.95	0.11648	-0.019	2.6	8.5	0	0	0	-5.832
17	328.42	0.083385	-0.012	2.6	8.5	0	0	0	-5.832

```

itn      theta      ||dx||      del_obj      obj      linear      ||x||0      a(x)      ||x||1      ||y||
Optimise cardinality reached the stopping criterion. Finished.
100.00% thermodynamically feasible internal fluxes (checked by cycleFreeFlux method).
  warmStartMethod: 'random'
  formulation: 'pqzwrns'
  thetaMultiplier: 1.5000
  theta: 0.5000
  regularizeOuter: 1
  epsilon: 1.0000e-06
  printLevel: 1
  relaxBounds: 0
  acceptRepairedFlux: 1
  thermoConsistencyMethod: 'cycleFreeFlux'
  bigNum: 10000
  debug: 0

optCardThermo objective data:
  1 = beta, the global weight on one-norm of internal reaction rate.
  -0.5 = min(g0), the local weight on zero-norm of internal reaction rate.
  0 = max(g0), the local weight on zero-norm of internal reaction rate.
  -0 = min(h0), the local weight on zero-norm of metabolite production rate.
  -0 = max(h0), the local weight on zero-norm of metabolite production rate.

optimizeCardinality objective data:

0 min cardinality variables:
  NaN mean(c(p))      NaN min(c(p))      NaN max(c(p))
  1 lambda0      NaN min(k)      NaN max(k)
  0 lambda1      NaN min(o(p))      NaN max(o(p))

824 max cardinality variables:
  -0 mean(c(q))      -0 min(c(q))      -0 max(c(q))
  1 delta0      0.4 min(d)      0.5 max(d)
  0 delta1      0 min(o(q))      0 max(o(q))

37723 cardinality free variables:
  0.45 mean(c(r))      -0 min(c(r))      1 max(c(r))
  0 alpha1      0 min(o(r))      0 max(o(r))

itn      theta      ||dx||      del_obj      obj      linear      ||x||0      a(x)      ||x||1      ||y||
1      0.50      9.6834e+05      -8.4e+07      8.4      8.5      0      0      0      -3.540
2      0.75      1.251      -0.098      8.3      8.4      0      0      0      -3.970
3      1.12      0.091425      -0.056      8.3      8.4      0      0      0      -3.506
4      1.69      0.092443      -0.084      8.2      8.4      0      0      0      -3.506
5      2.53      0.44167      -0.14      8.1      8.5      0      0      0      -3.506
6      3.80      0.31473      -0.25      7.8      8.5      0      0      0      -3.956
7      5.70      1.8987      -0.76      7      9.6      0      0      0      -4.903
8      8.54      1.0204      -1.2      5.8      9.4      0      0      0      -5.394
9      12.81      1.4536      -1.1      4.8      10      0      0      0      -5.832
10     19.22      0.67982      -0.85      3.9      9.8      0      0      0      -5.832
11     28.83      0.60377      -0.54      3.4      9.2      0      0      0      -5.832

```

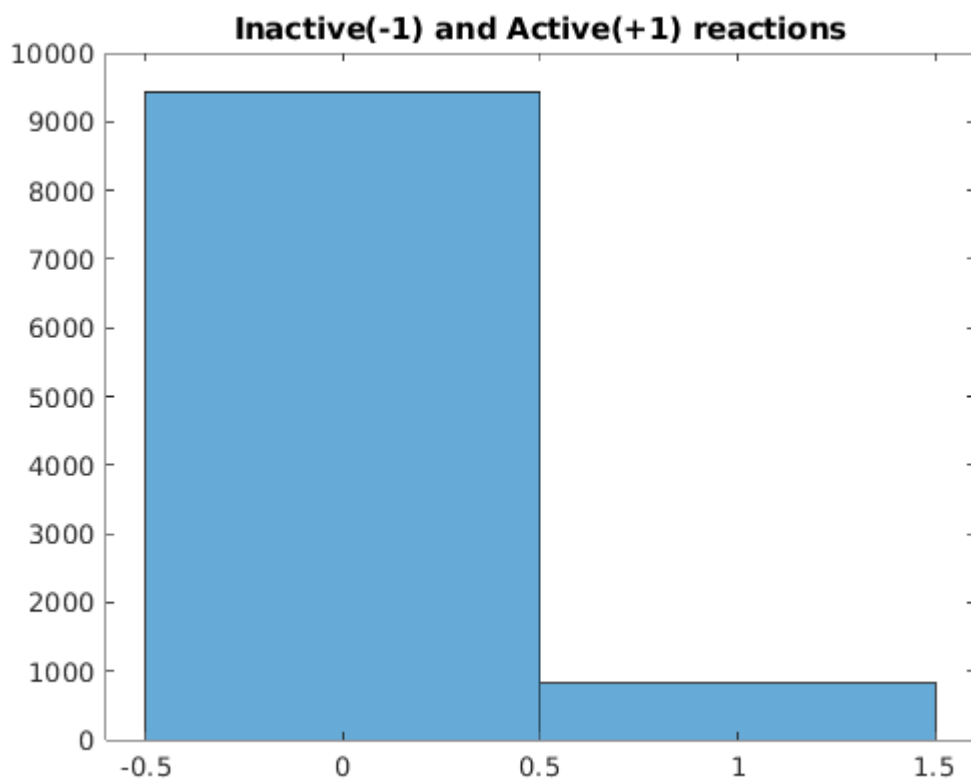
12	43.25	0.45629	-0.35	3	8.9	0	0	0	-5.832
13	64.87	0.57354	-0.22	2.8	8.7	0	0	0	-5.832
14	97.31	0.4286	-0.13	2.7	8.5	0	0	0	-5.832
15	145.96	0.145	-0.036	2.7	8.5	0	0	0	-5.832
16	218.95	0.11648	-0.019	2.6	8.5	0	0	0	-5.832
17	328.42	0.083385	-0.012	2.6	8.5	0	0	0	-5.832
itn	theta	dx	del_obj	obj	linear	x  0	a(x)	x  1	y

Optimise cardinality reached the stopping criterion. Finished.

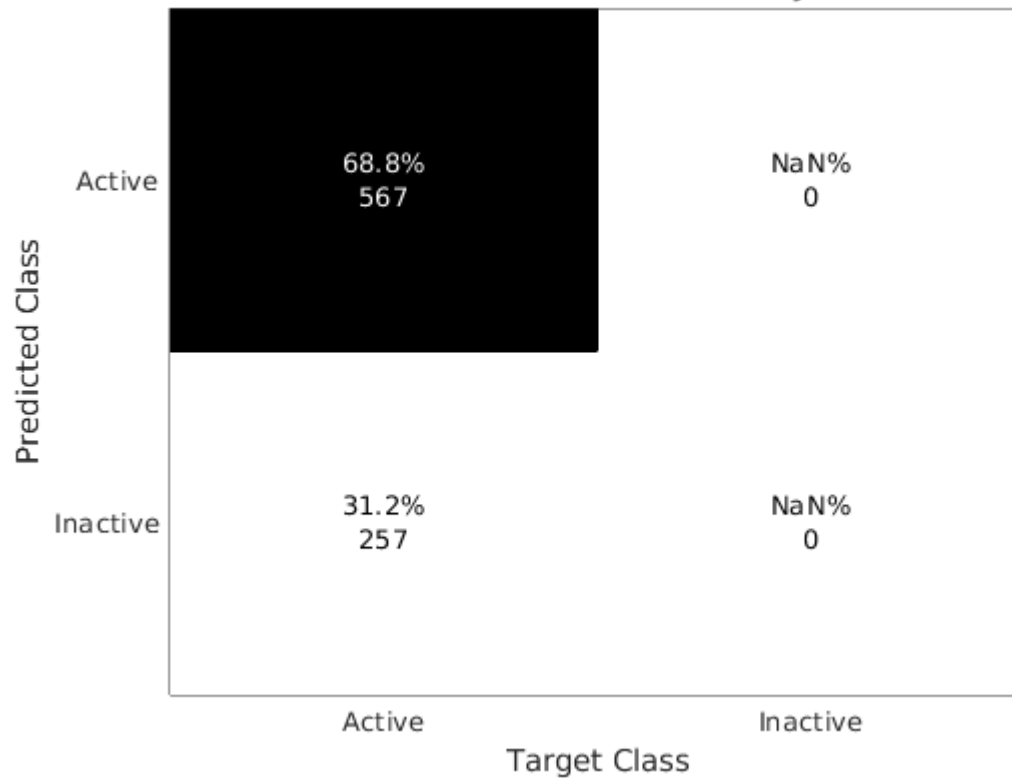
100.00% thermodynamically feasible internal fluxes (checked by cycleFreeFlux method).

iter.	nz.flux.	%it.feas.int.flux.	%feas.inc.flux.	nz.prod.	%it.feas.nz.prod.	%feas.inc.prod.	fo
1	111	1.00	0.02	74	1.24	NaN	gre
2	110	1.00	0.02	76	1.24	NaN	gre
3	132	1.00	0.03	93	1.22	NaN	gre
4	127	1.00	0.04	94	1.18	NaN	gre
5	215	1.00	0.10	152	1.24	NaN	gre
6	190	1.00	0.15	138	1.23	NaN	gre
7	247	1.00	0.21	175	1.29	NaN	gre
8	268	1.00	0.28	193	1.34	NaN	gre
9	300	1.00	0.36	217	1.27	NaN	gre
10	284	1.00	0.43	201	1.42	NaN	gre
11	234	1.00	0.47	162	1.37	NaN	gre
12	325	1.00	0.53	235	1.39	NaN	gre
13	217	1.00	0.56	155	1.39	NaN	gre
14	223	1.00	0.59	158	1.29	NaN	gre
15	197	1.00	0.60	137	1.37	NaN	gre
16	136	1.00	0.61	94	1.29	NaN	gre
17	189	1.00	0.63	135	1.39	NaN	gre
18	222	1.00	0.65	148	1.47	NaN	gre
19	316	1.00	0.68	219	1.42	NaN	gre
20	155	1.00	0.69	114	1.34	NaN	gre

iter.	nz.flux.	%it.feas.int.flux.	%feas.inc.flux.	nz.prod.	%it.feas.nz.prod.	%feas.inc.prod.	fo
thermoKernel terminating early: n = nMax = 20							



**Reaction confusion matrix, accuracy = 0.68811**



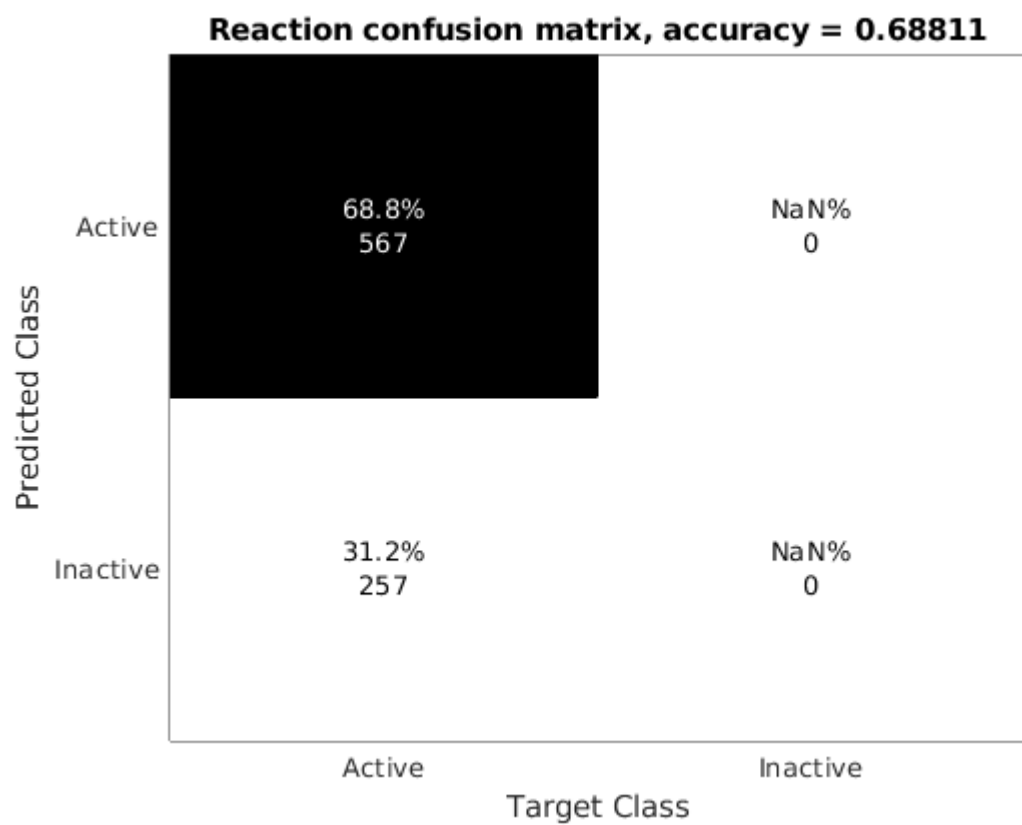
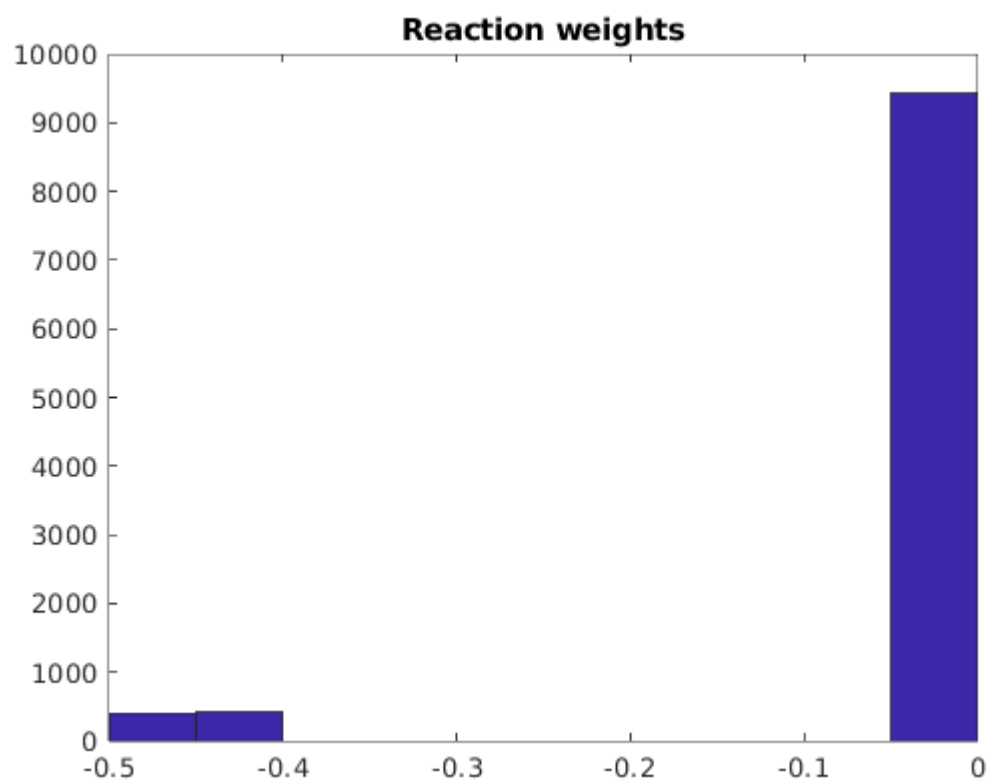
--- thermoKernel END ----

```
[nMet,nRxn]=size(tissueModel.S)
```

```
nMet = 1449
nRxn = 1915
```

Compare the target versus predicted model

```
plotThermoKernelExtractStats(stoichFluxThermoConsistModel,
activeInactiveRxn, rxnWeightsRed, presentAbsentMet, metWeightsRed,
thermoModelMetBool, thermoModelRxnBool)
```



## Acknowledgments

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## REFERENCES

- [1] Ronan M T Fleming, Hulda S Haraldsdottir, Le Hoai Minh, Phan Tu Vuong, Thomas Hankemeier, Ines Thiele, Cardinality optimization in constraint-based modelling: application to human metabolism, *Bioinformatics*, Volume 39, Issue 9, September 2023, btad450, <https://doi.org/10.1093/bioinformatics/btad450>
- [2] Preciat, G., Wegrzyn, A. B., Luo, X., Thiele, I., Hankemeier, T., & Fleming, R. M. T. (2025). XomicsToModel: omics data integration and generation of thermodynamically consistent metabolic models. *Nature Protocols*. <https://doi.org/10.1038/s41596-025-01288-9>