

# Extract a thermodynamically consistent subnetwork from a given model

**Author(s): Ronan M.T. Fleming, School of Medicine, University of Galway**

**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','ctrsRemoveMethod','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		

```

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

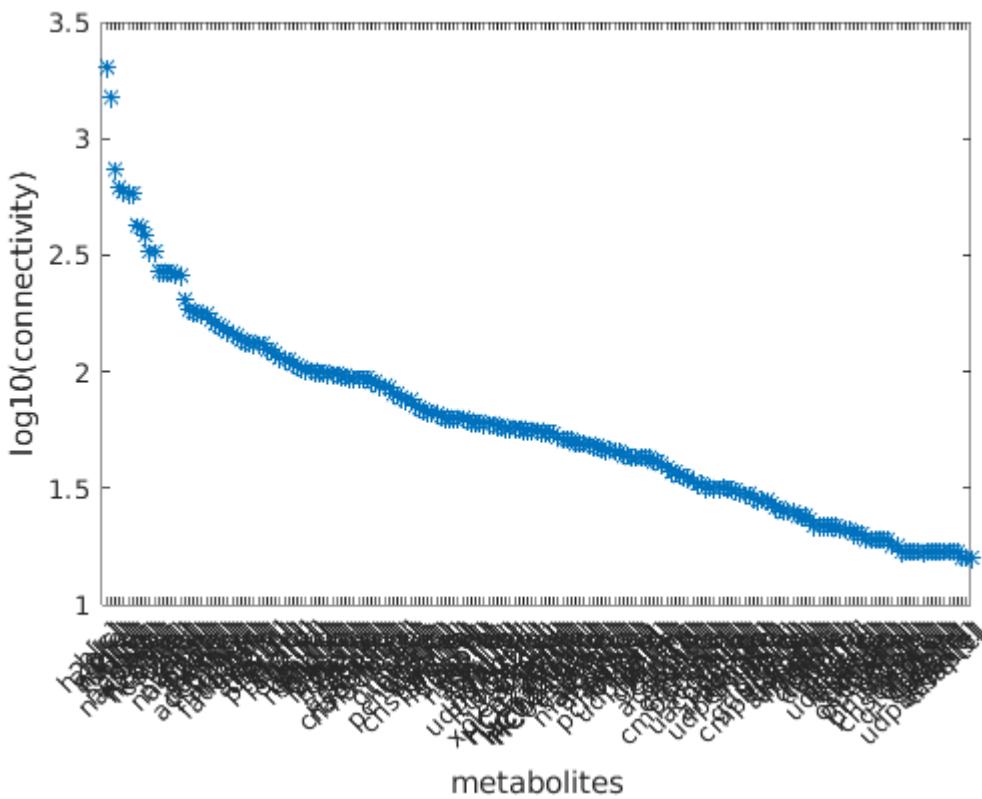
## Data to define a thermodynamically consistent subnetwork

Setup random data to select a random subset

```

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

```



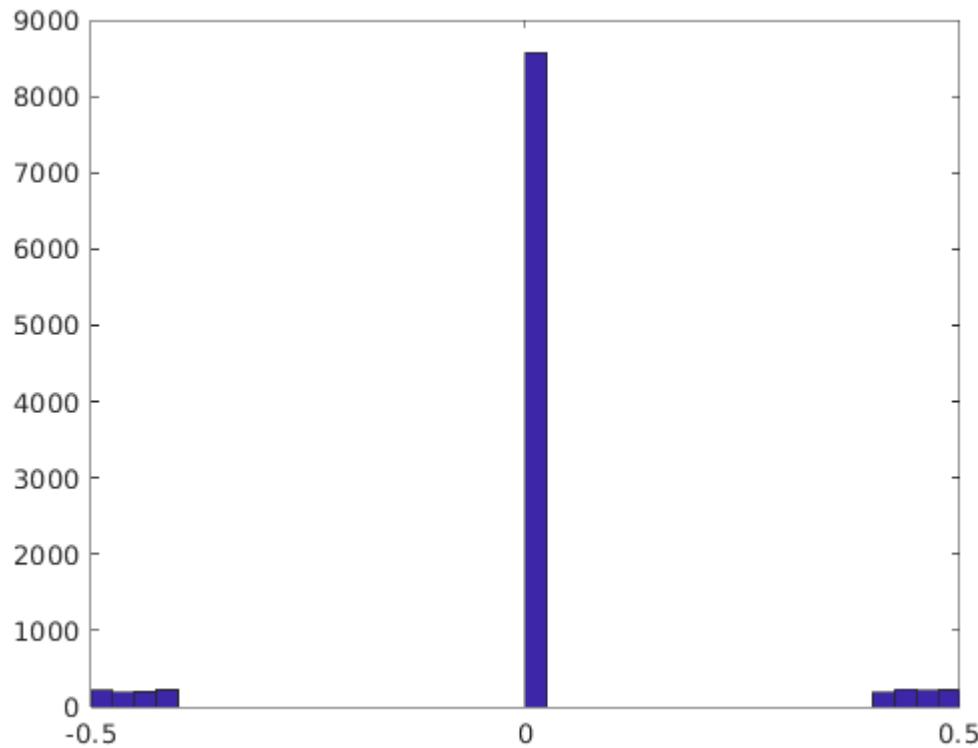
```
[nMet,nRxn]=size(stoichFluxThermoConsistModel.S);
rxnWeights=rand(nRxn,1)-0.5;
rxnWeights(~stoichFluxThermoConsistModel.SConsistentRxnBool)=0;

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)

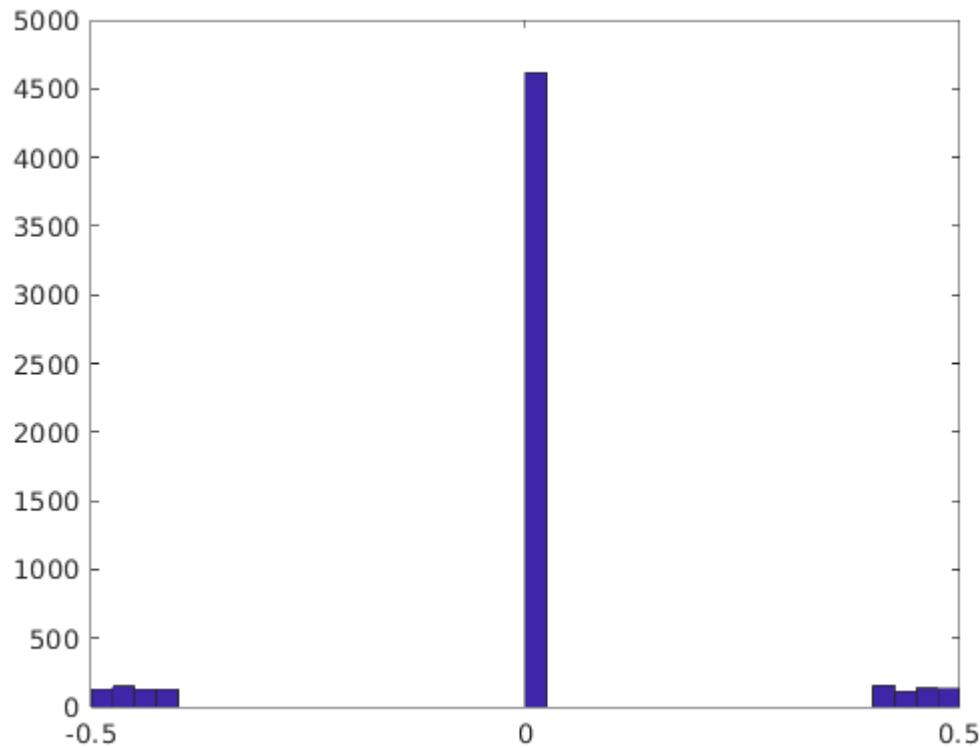
```



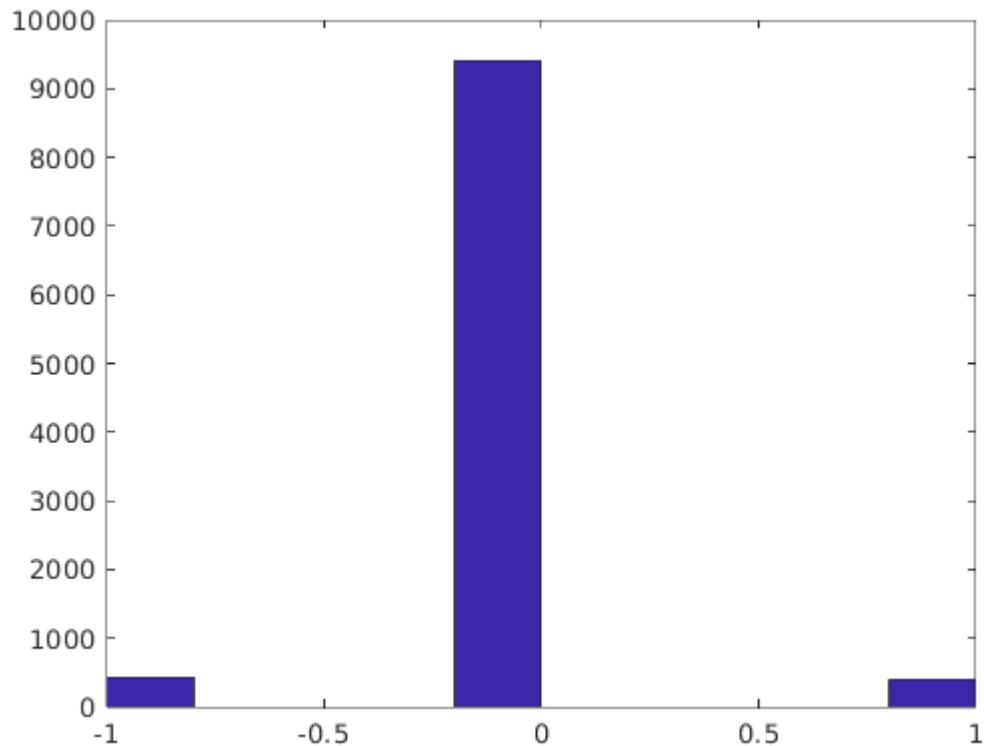
```

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)

```

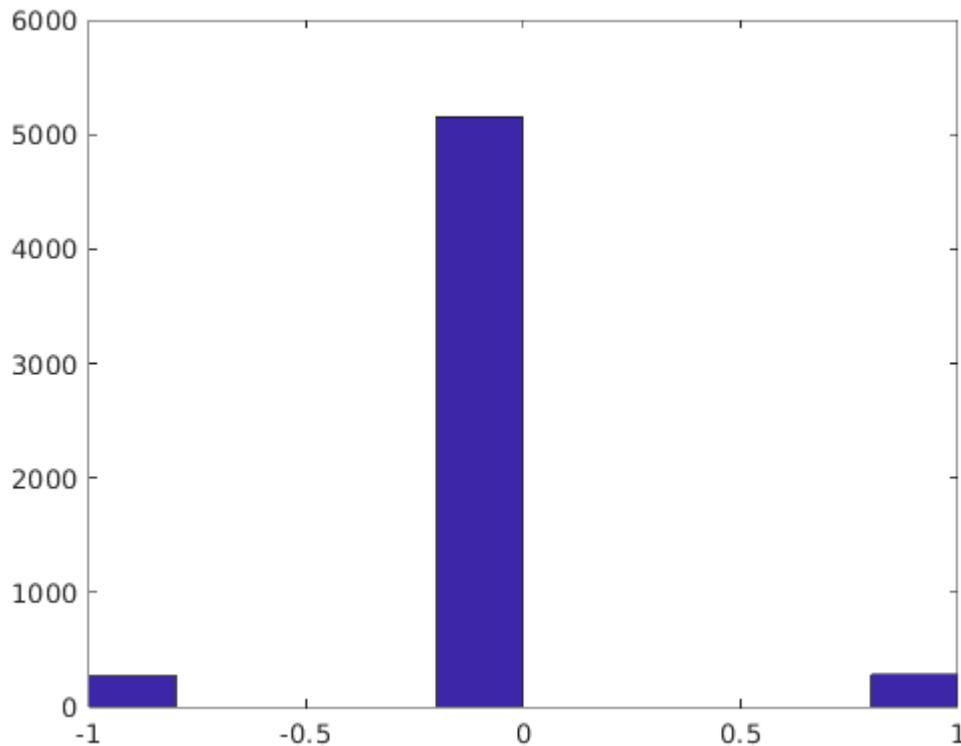


```
nlt=length(coreRxnBool);
activeInactiveRxn=zeros(nlt,1);
activeInactiveRxn(coreRxnBool)=1;
activeInactiveRxn(removeRxnBool)=-1;
hist(activeInactiveRxn)
```



```
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 alphal                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: 'pqzwr's'
    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 alphal	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

Optimise cardinality reached the stopping criterion. Finished.

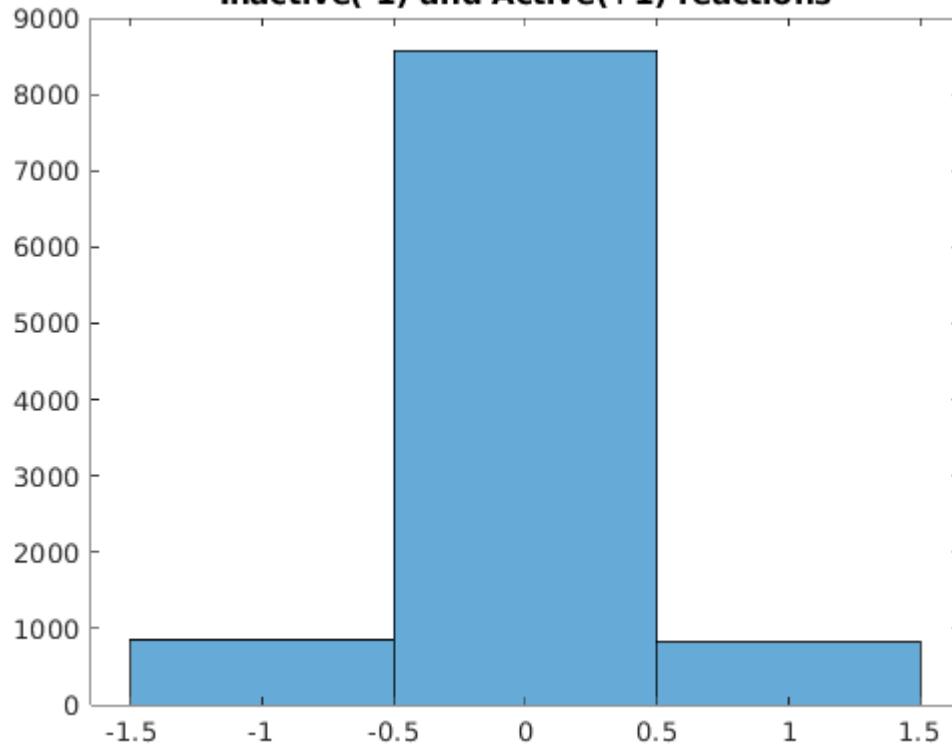
100.00% thermodynamically feasible internal fluxes (checked by cycleFreeFlux method).

iter. nz.flux.%it.feas.int.flux. %feas.inc.flux. nz.prod. %it.feas.nz.prod. %feas.inc.prod.

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

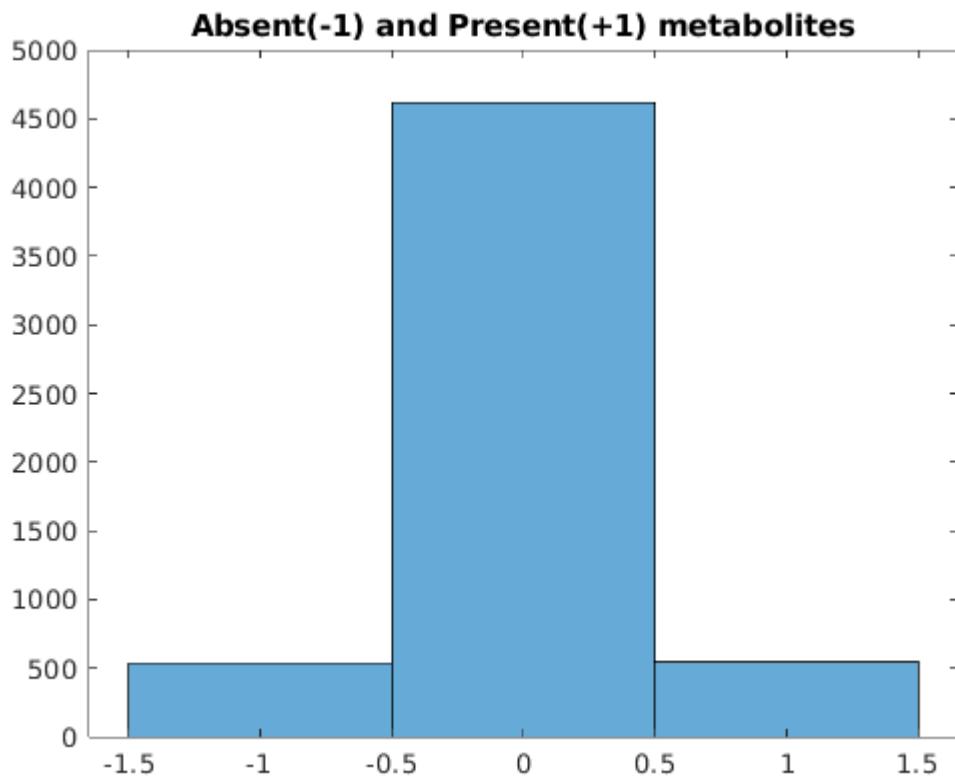
iter. nz.flux.%it.feas.int.flux. %feas.inc.flux. nz.prod. %it.feas.nz.prod. %feas.inc.prod.  
thermoKernel terminating early: n = nMax = 20

### Inactive(-1) and Active(+1) reactions



**Reaction confusion matrix, accuracy = 0.84148**





**Metabolite confusion matrix, accuracy = 0.80425**

		Target Class	
		Active	Inactive
Predicted Class	Active	74.1% 403	13.2% 71
	Inactive	25.9% 141	86.8% 468

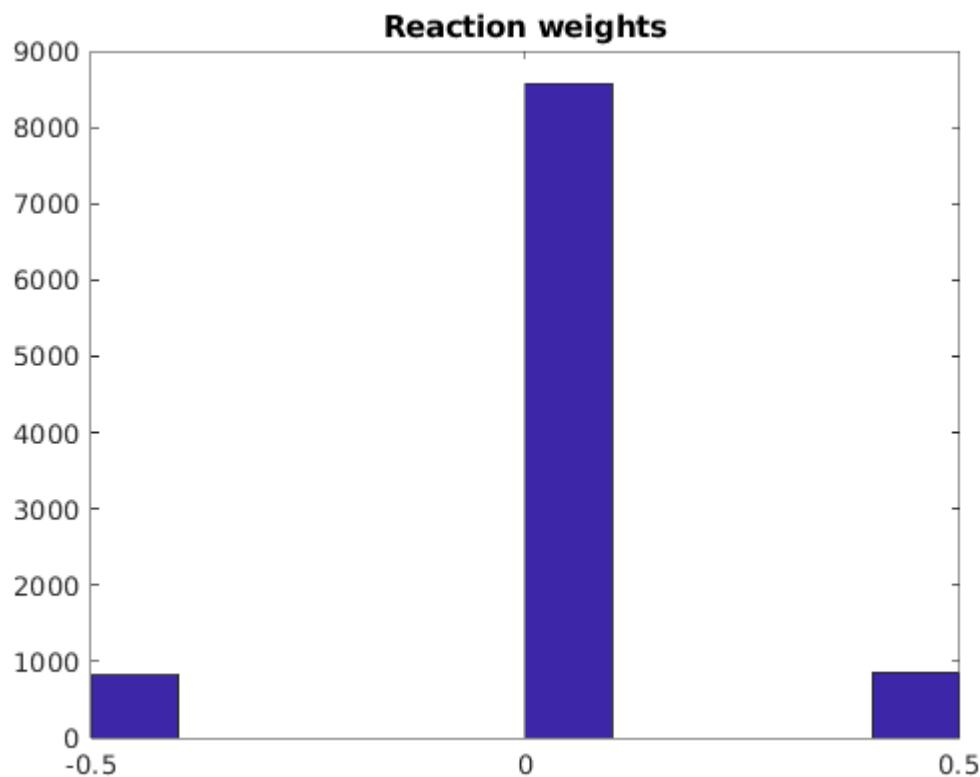
--- thermoKernel END ----

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

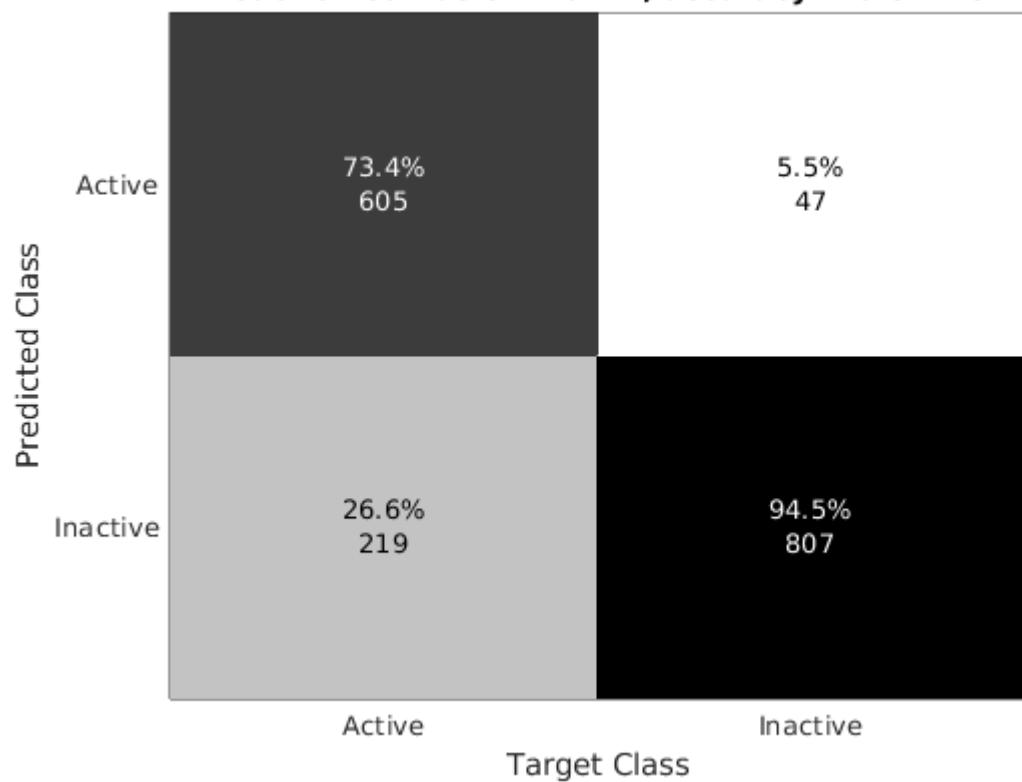
```
nMet = 2203  
nRxn = 2986
```

Compare the target versus predicted model

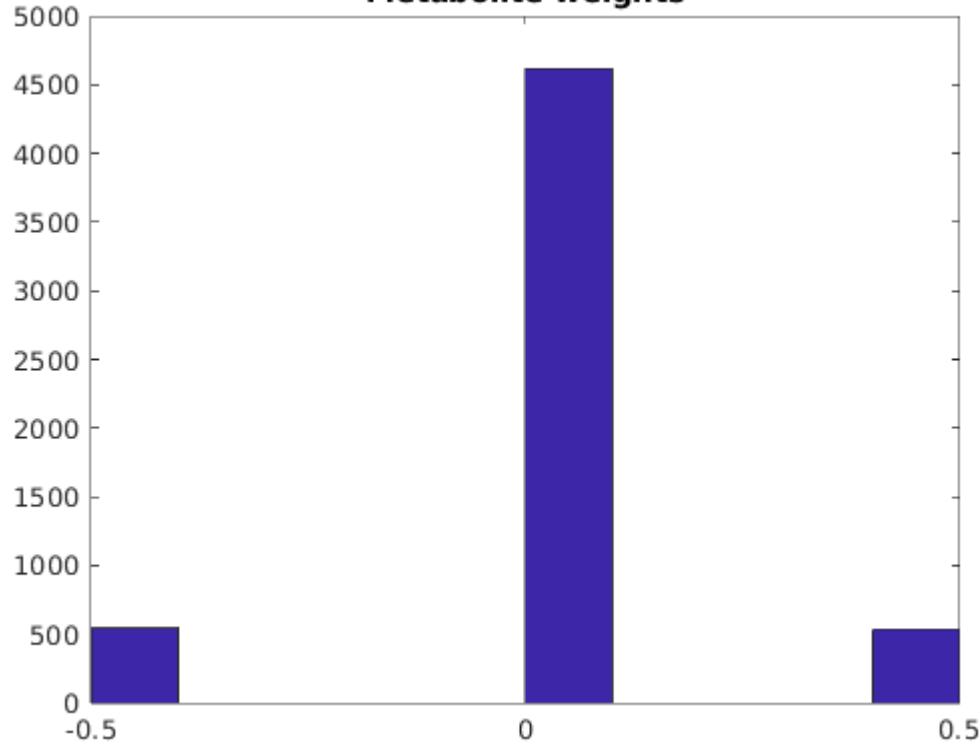
```
%plotThermoCoreStats(activeInactiveRxn, presentAbsentMet,  
thermoModelMetBool, thermoModelRxnBool);  
plotThermoKernelExtractStats(stoichFluxThermoConsistModel,  
activeInactiveRxn, rxnWeights, presentAbsentMet, metWeights,  
thermoModelMetBool, thermoModelRxnBool)
```



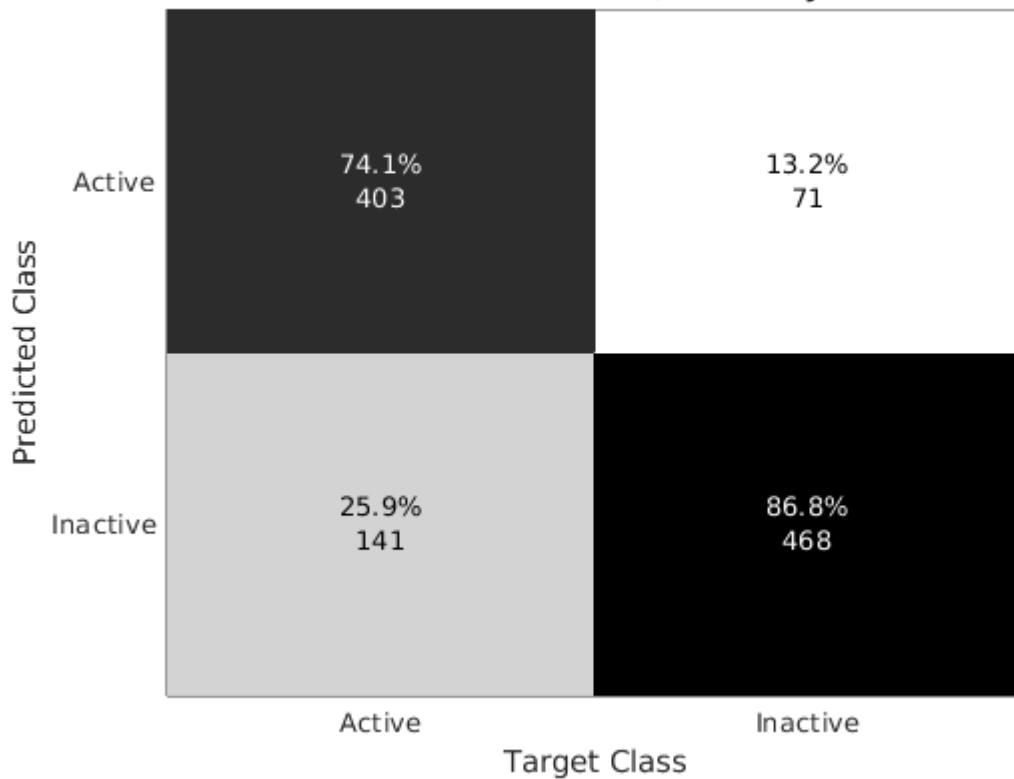
### Reaction confusion matrix, accuracy = 0.84148



### Metabolite weights



### 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: 'pqzwr'
    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.
    0.5 = 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 lambdal              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 deltal               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 alphal                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      -10.000
7       5.70      1.3982      -1.2      0.95       10     1.23553     0.51       0      -10.000
8       8.54      0.78275     -1.5      -0.51       9.8     0.832228     0.5       0      -12.000
9      12.81      0.52353     -1.8      -2.3       9.5     0.832228     0.55       0      -12.000
10      19.22      0.37265     -0.29      -2.6       9.1     0.832228     0.62       0      -12.000
11      28.83      0.33297     -0.11      -2.7       8.9     0.832228     0.73       0      -12.000
12      43.25      0.19468     -0.5      -3.2       8.7     0.832228     0.83       0      -13.210
13      64.87      0.099656     -0.91      -4.1       8.7     0.832228     0.83       0      -15.030
14      97.31      0.11652      -1.5      -5.6       8.6     0.832228     0.83       0      -15.030
15     145.96      0.12539     -0.03      -5.6       8.6     0.832228     0.83       0      -15.030
16     218.95      0.067216     -0.018     -5.6       8.6     0.832228     0.83       0      -15.030
17     328.42      0.13964     -0.012     -5.6       8.6     0.832228     0.83       0      -15.030

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.
    0.5 = 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))

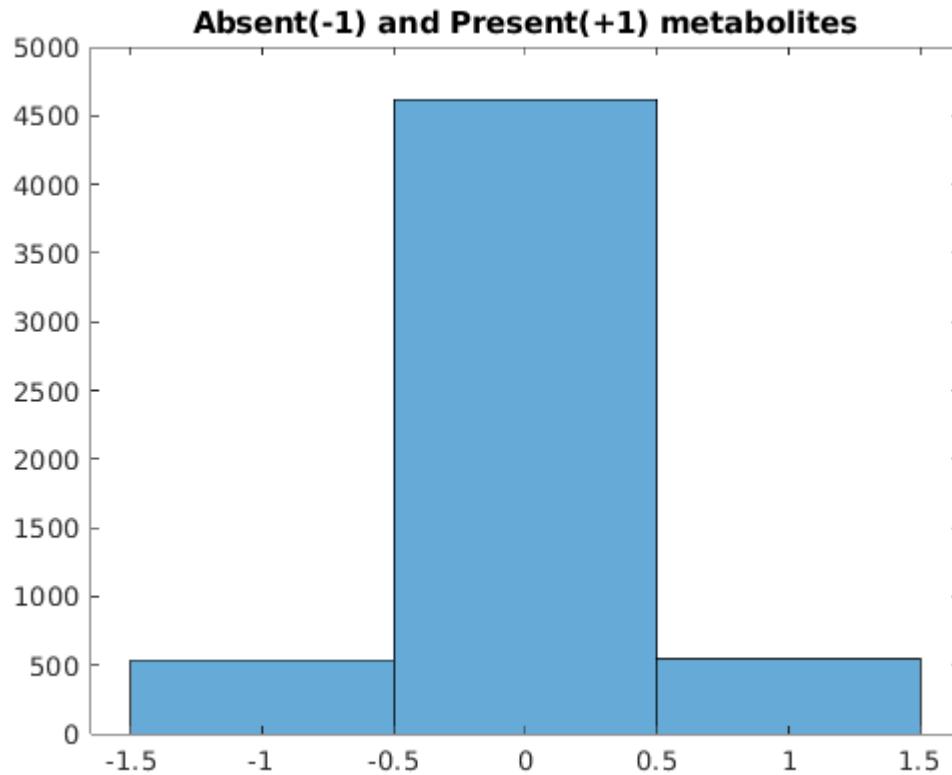
36920 cardinality free variables:
    0.46 mean(c(r))        -0 min(c(r))        1 max(c(r))
    0 alphal               0 min(o(r))         0 max(o(r))

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

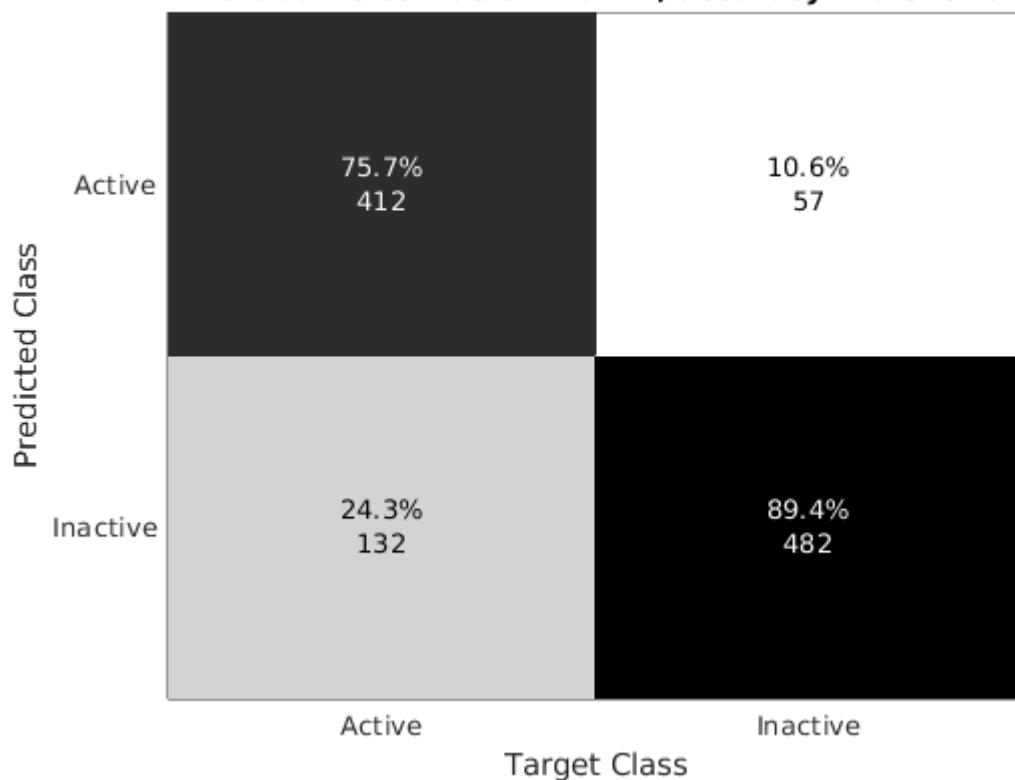
```

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.  
thermoKernel terminating early: n = nMax = 20



### Metabolite confusion matrix, accuracy = 0.82548



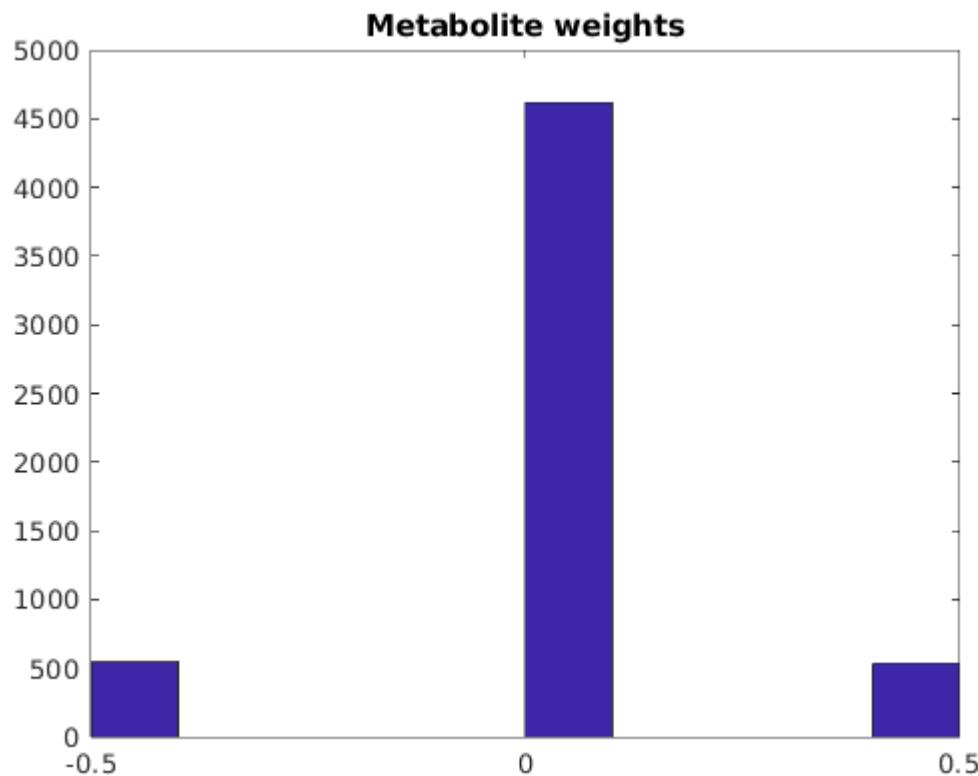
```
--- thermoKernel END ---
```

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

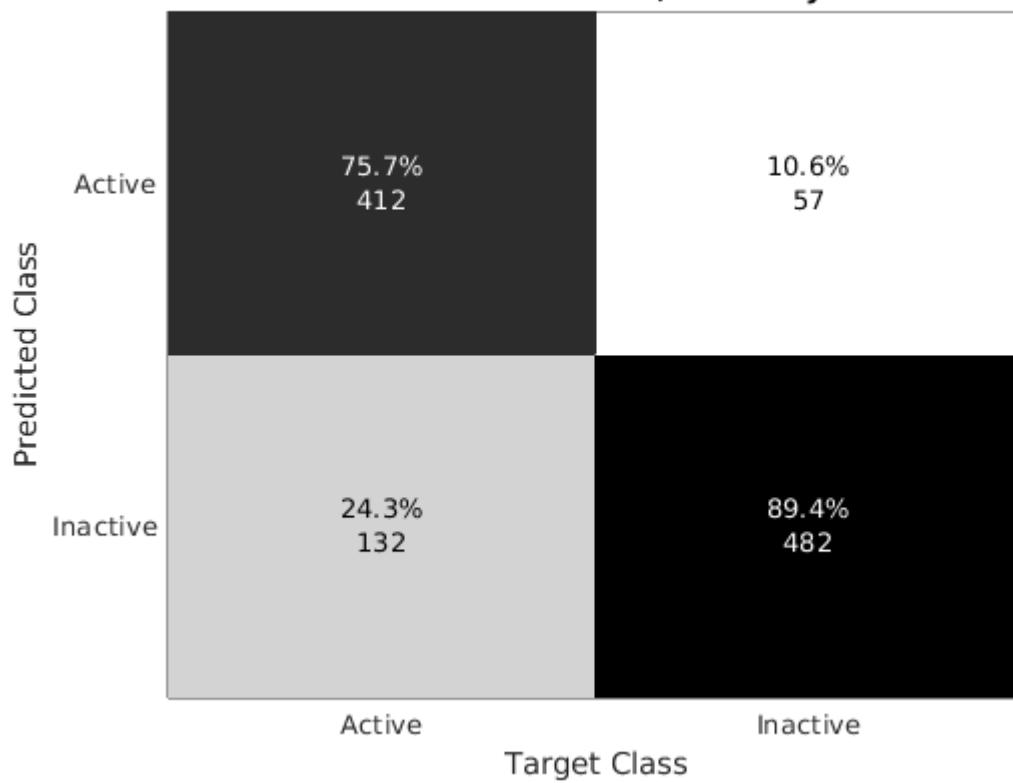
```
nMet = 1870  
nRxn = 2635
```

Compare the target versus predicted model

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



**Metabolite confusion matrix, accuracy = 0.82548**



### 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: '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 = 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 alphal                 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: '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 = 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	9.6798e+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

Optimise cardinality reached the stopping criterion. Finished.

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

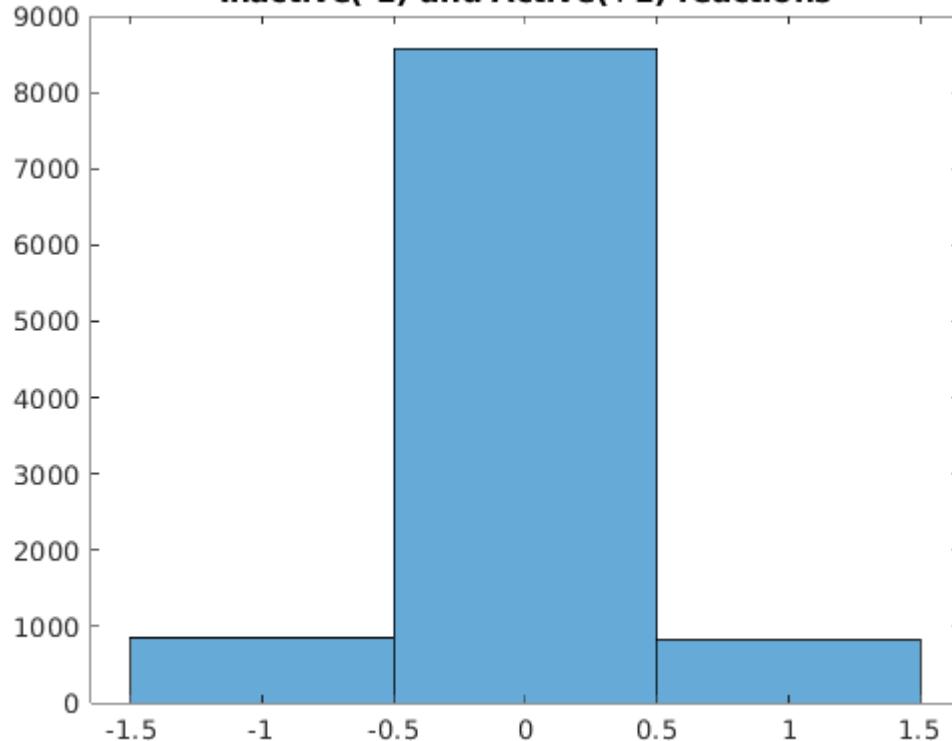
iter.	nz.flux.%it.feas.int.flux.	%feas.inc.flux.	nz.prod.	%it.feas.nz.prod.	%feas.inc.prod.	fo
1	100	1.00	0.01	72	1.15	NaN
2	99	1.00	0.01	70	1.21	NaN
3	113	1.00	0.02	79	1.15	NaN
4	113	1.00	0.03	83	1.12	NaN
5	181	1.00	0.07	134	1.14	NaN
6	240	1.00	0.14	183	1.14	NaN
7	192	1.00	0.20	128	1.38	NaN
8	184	1.00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

```

18      116          1.00          0.67          87          1.20          NaN
19      147          1.00          0.68          101         1.40          NaN
20      199          1.00          0.70          139         1.42          NaN
iter. nz.flux.%it.feas.int.flux. %feas.inc.flux. nz.prod. %it.feas.nz.prod. %feas.inc.prod.
thermoKernel terminating early: n = nMax = 20

```

### Inactive(-1) and Active(+1) reactions



### Reaction confusion matrix, accuracy = 0.83313



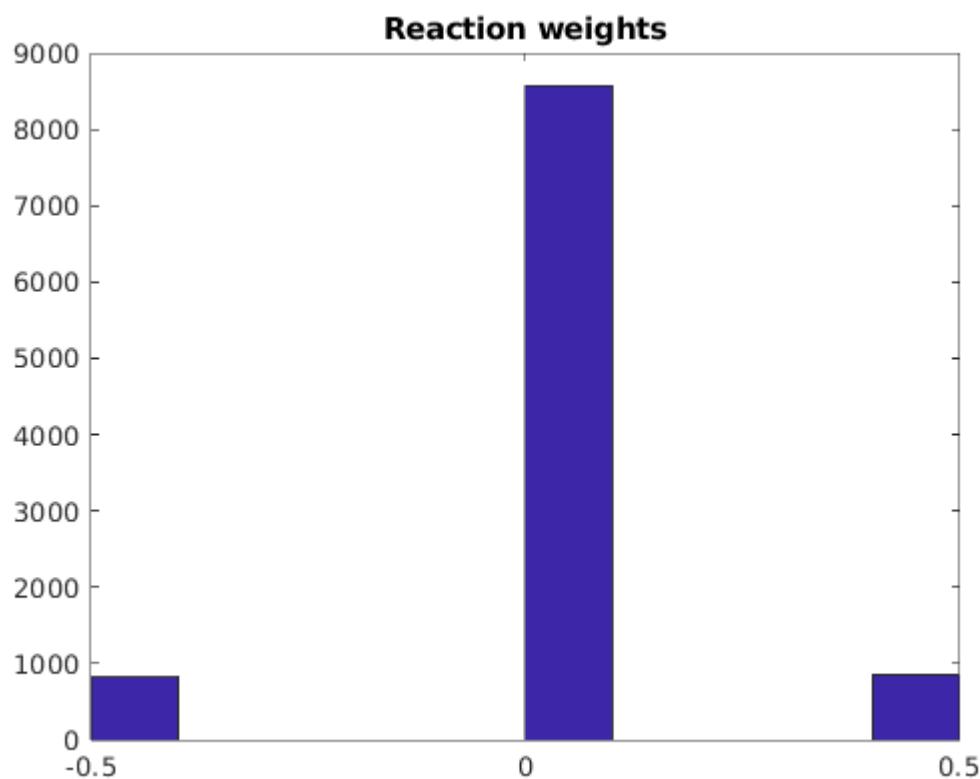
```
--- thermoKernel END ----
```

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

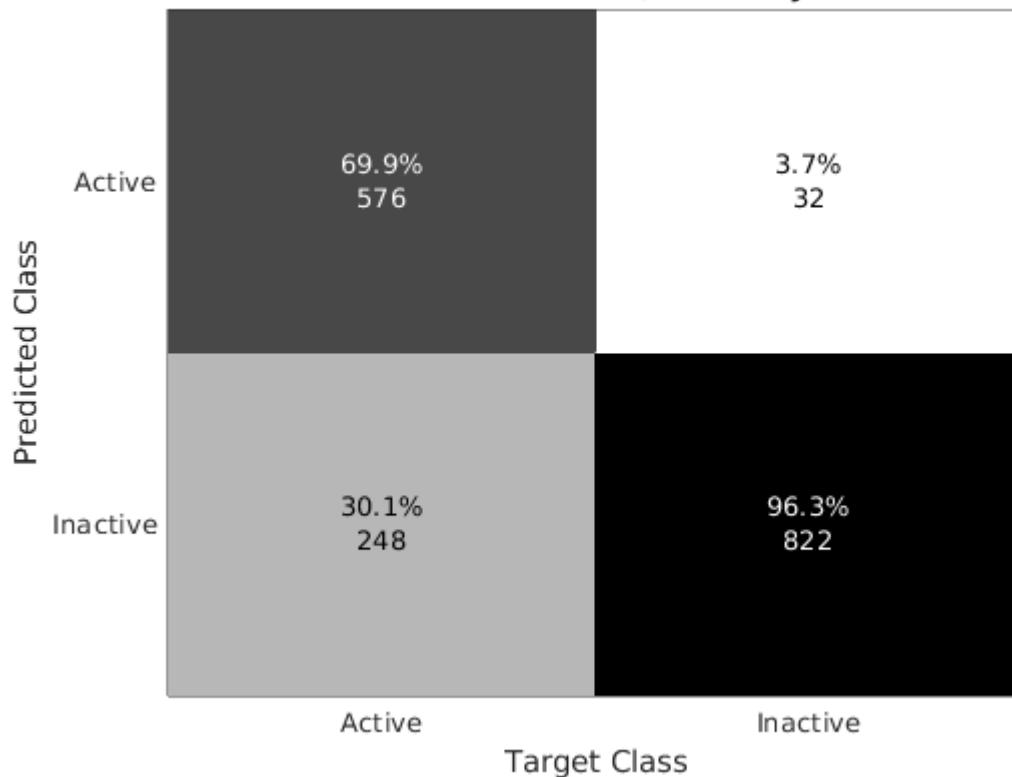
```
nMet = 1441  
nRxn = 1924
```

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 deltal              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 alphal              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: 'pqzwr'
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 alphal                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. fo
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

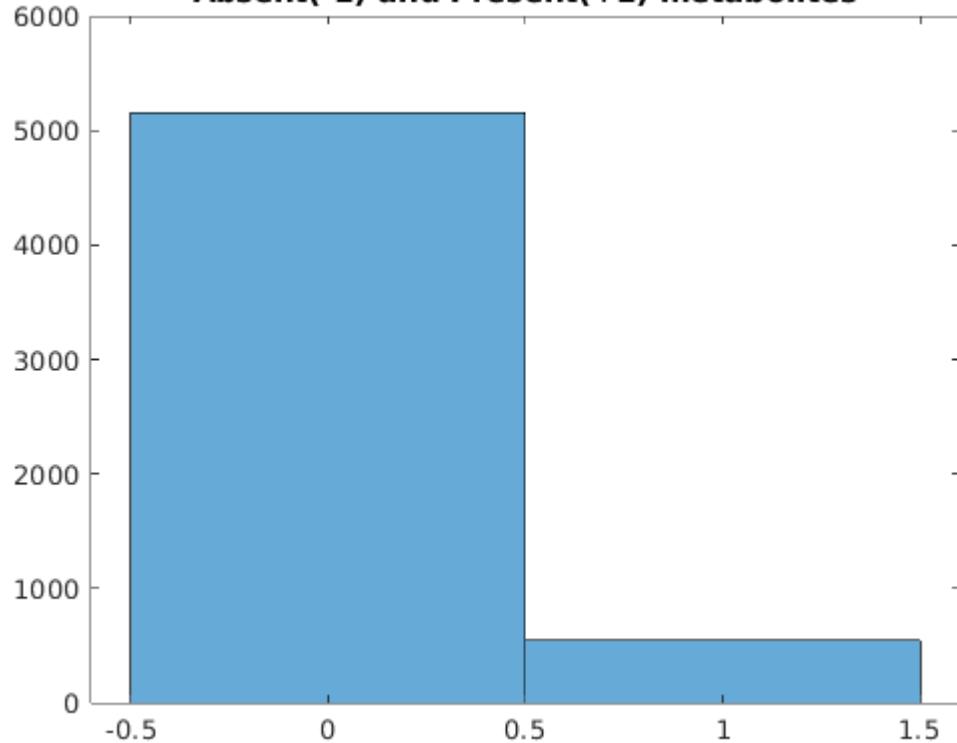
```

```

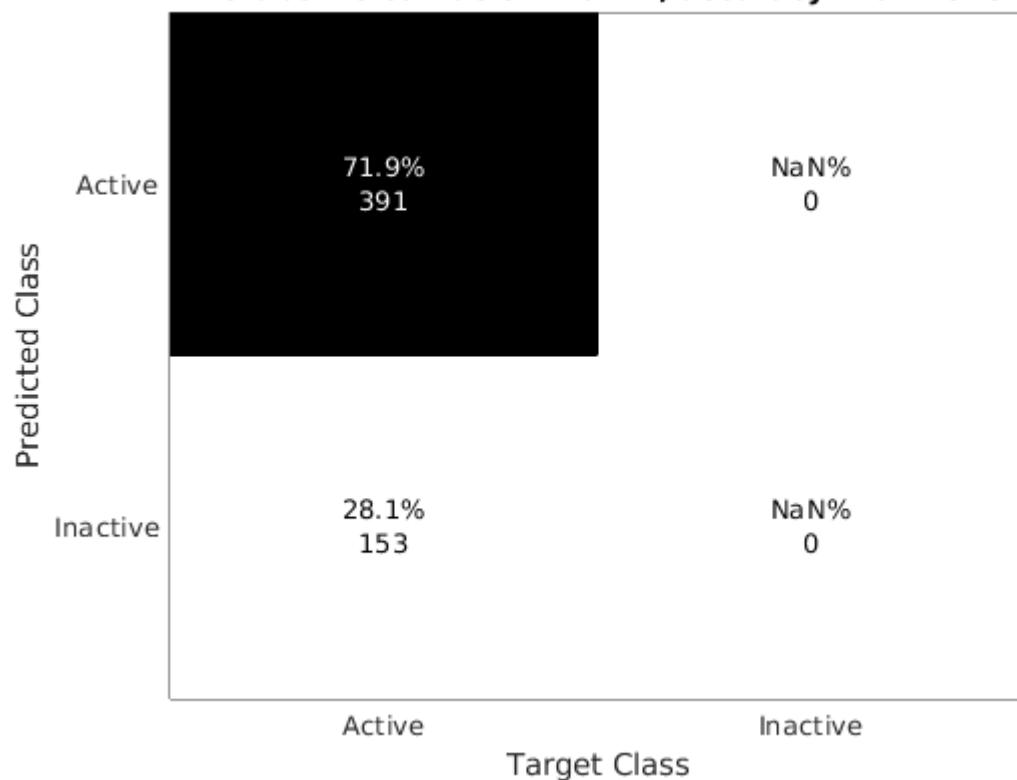
16      129      1.00      NaN      92      1.30      0.69      gre
17      164      1.00      NaN     116      1.30      0.69      gre
18      166      1.00      NaN     123      1.26      0.70      gre
19      165      1.00      NaN     117      1.35      0.71      gre
20      141      1.00      NaN      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.
thermoKernel terminating early: n = nMax = 20

```

**Absent(-1) and Present(+1) metabolites**



### Metabolite confusion matrix, accuracy = 0.71875



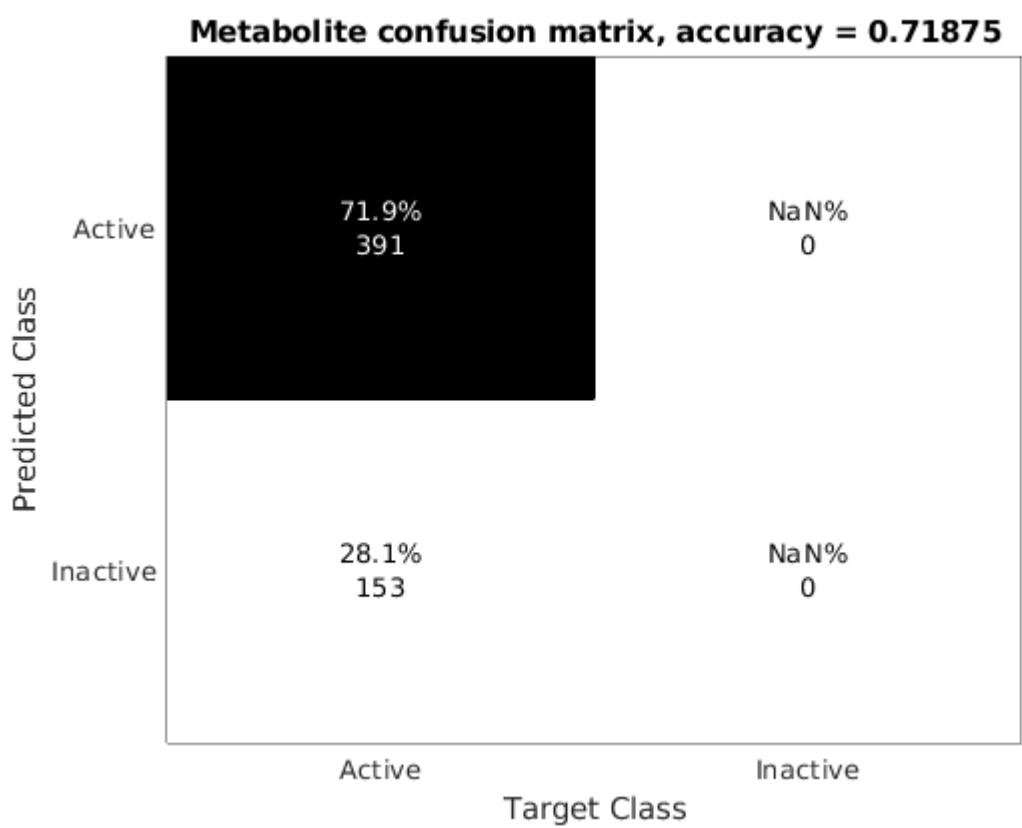
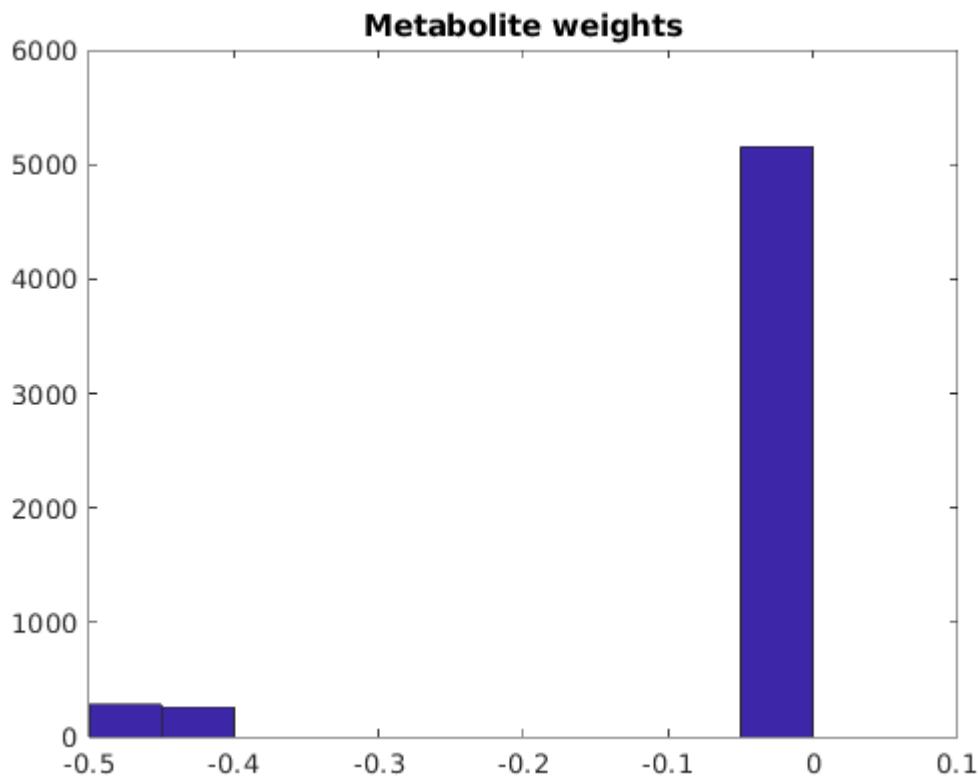
```
--- thermoKernel END ---
```

```
[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 alphal	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	9.6384e+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

itn	theta	$\ dx\ $	del_obj	obj	linear	$\ x\ _0$	$a(x)$	$\ x\ _1$	$\ y\ $
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

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 = 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	deltal	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

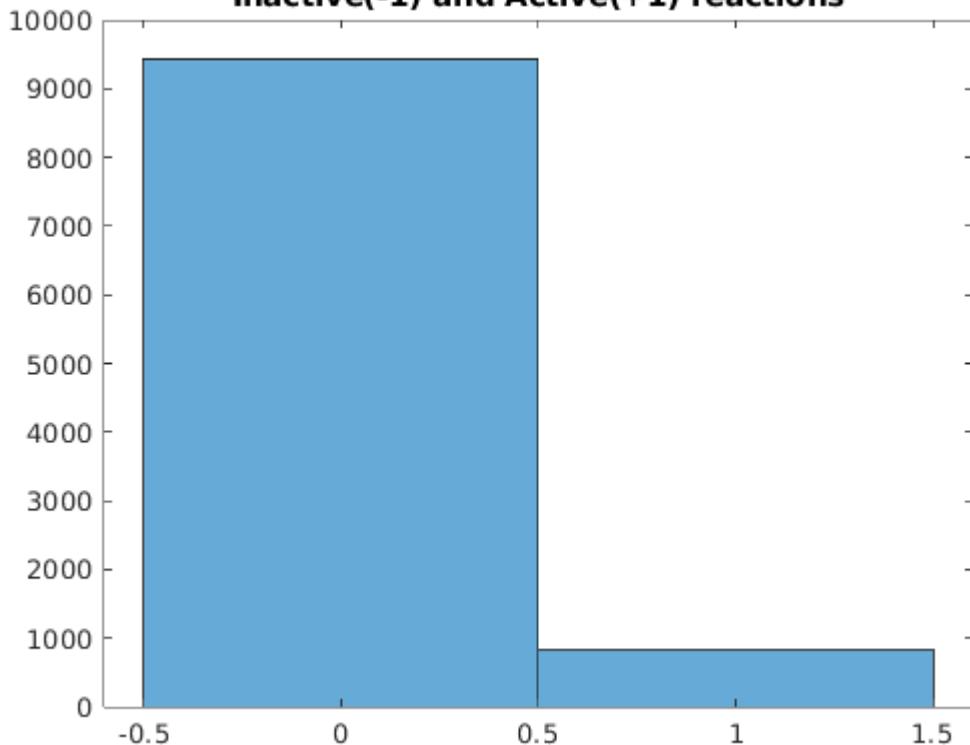
12	43.25	0.45629	-0.35	3	8.9	0	0	0	-5.832
13	64.87	0.57354	-0.22	2.8	8.7	0	0	0	-5.832
14	97.31	0.4286	-0.13	2.7	8.5	0	0	0	-5.832
15	145.96	0.145	-0.036	2.7	8.5	0	0	0	-5.832
16	218.95	0.11648	-0.019	2.6	8.5	0	0	0	-5.832
17	328.42	0.083385	-0.012	2.6	8.5	0	0	0	-5.832
itn	theta	dx	del_obj	obj	linear	x  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.	fc
1	111	1.00	0.02	74	1.24	NaN
2	110	1.00	0.02	76	1.24	NaN
3	132	1.00	0.03	93	1.22	NaN
4	127	1.00	0.04	94	1.18	NaN
5	215	1.00	0.10	152	1.24	NaN
6	190	1.00	0.15	138	1.23	NaN
7	247	1.00	0.21	175	1.29	NaN
8	268	1.00	0.28	193	1.34	NaN
9	300	1.00	0.36	217	1.27	NaN
10	284	1.00	0.43	201	1.42	NaN
11	234	1.00	0.47	162	1.37	NaN
12	325	1.00	0.53	235	1.39	NaN
13	217	1.00	0.56	155	1.39	NaN
14	223	1.00	0.59	158	1.29	NaN
15	197	1.00	0.60	137	1.37	NaN
16	136	1.00	0.61	94	1.29	NaN
17	189	1.00	0.63	135	1.39	NaN
18	222	1.00	0.65	148	1.47	NaN
19	316	1.00	0.68	219	1.42	NaN
20	155	1.00	0.69	114	1.34	NaN

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

### Inactive(-1) and Active(+1) reactions



### Reaction confusion matrix, accuracy = 0.68811

		Target Class	
		Active	Inactive
Predicted Class	Active	68.8% 567	NaN% 0
	Inactive	31.2% 257	NaN% 0

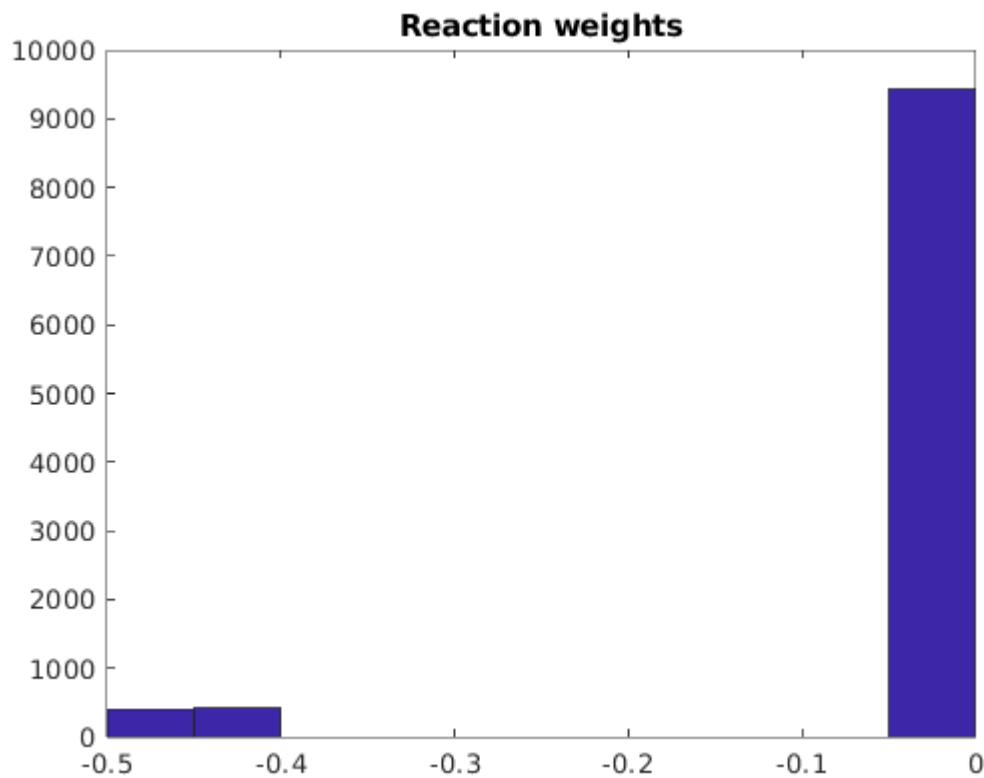
--- thermoKernel END ---

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

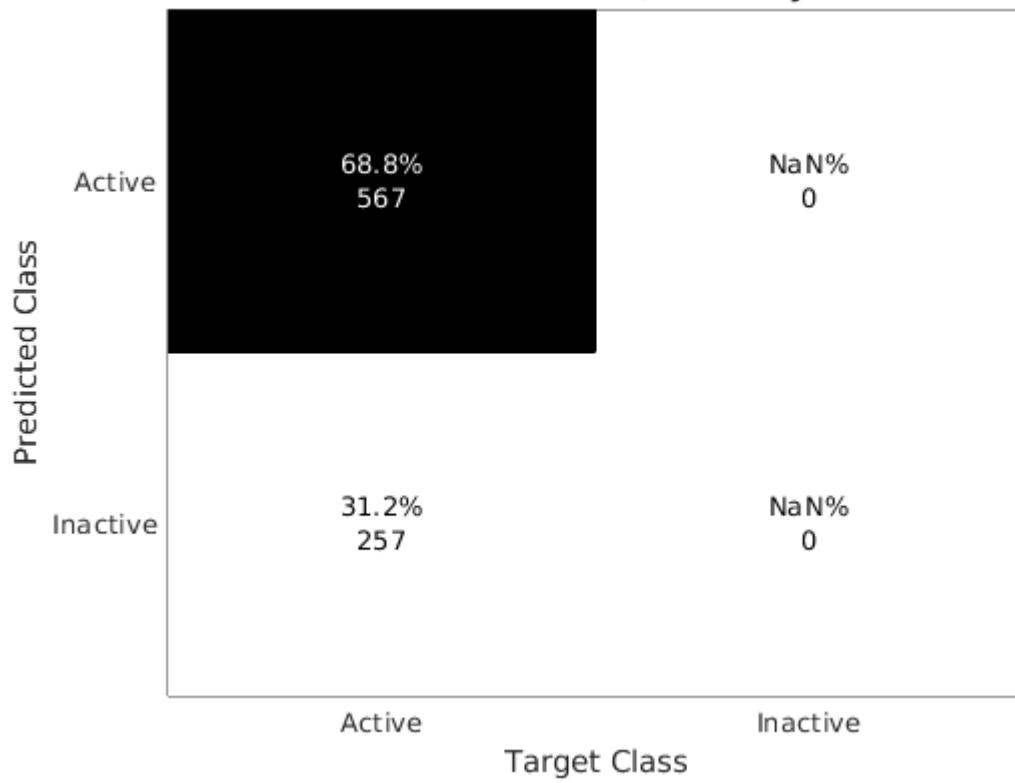
```
nMet = 1449  
nRxn = 1915
```

Compare the target versus predicted model

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



**Reaction confusion matrix, accuracy = 0.68811**



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## REFERENCES

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- [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>