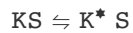


Bistable motif: detailed balancing and its consequences

Finding the condition of multistationarity

We consider the following reactions:



The species of the system are:



In total, there are 11 reactions and 6 species.

We firstly construct the ordinary differential equations based on mass-action kinetics. Then compute the determinant of Jacobian, using the solution at critical point (steady state) to calculate the determinant. The (necessary) condition for multistationarity is to make determinant equal to zero (non-zero determinant implies injectivity).

```

ClearAll["Global`*"];
A = Table[0, {11}, {6}];
A[[1]][[1]] = -1;
A[[1]][[3]] = -1;
A[[1]][[5]] = 1;
A[[2]] = -A[[1]];
A[[3]][[3]] = 1; A[[3]][[2]] = 1; A[[3]][[5]] = -1;
A[[4]][[1]] = -1;
A[[4]][[4]] = -1;
A[[4]][[6]] = 1;
A[[5]] = -A[[4]];
A[[6]][[4]] = 1;
A[[6]][[2]] = 1;
A[[6]][[6]] = -1;
A[[7]][[2]] = -1;
A[[7]][[1]] = 1;
A[[8]][[3]] = -1; A[[8]][[4]] = 1; A[[9]] = -A[[8]];
A[[10]][[5]] = -1; A[[10]][[6]] = 1; A[[11]] = -A[[10]];
stoiM = Transpose[A];
(* Now we construct the rate vector *)
ks = {k1 × x3 × x1, k2 × x5, k3 × x5, k4 × x4 × x1,
      k5 × x6, k6 × x6, k7 × x2, k8 × x3, k9 × x4, k10 × x5, k11 × x6};
ssEqns = stoiM.ks;
mC = RowReduce[NullSpace[A]];
subsEqns = {ssEqns[[2]], ssEqns[[4]],
            ssEqns[[5]], ssEqns[[6]], x1 + x2 + x5 + x6 - T1, x3 + x4 + x5 + x6 - T2};
jacobian = D[subsEqns, {{x1, x2, x3, x4, x5, x6}}];
detJ = Collect[Distribute[Det[jacobian]], {x1, x2, x3, x4, x5, x6}}];
solution =
  Solve[{subsEqns[[1]], subsEqns[[2]], subsEqns[[3]], subsEqns[[4]] == 0},
        {x2, x4, x5, x6}}];
detSubs = Replace[detJ, solution[[1]], {0, Infinity}];
(* Equivilant to detSubs=detJ/.solution[[1]]; *)
polSubs = Numerator[Together[detSubs]];
finalSubs = Collect[Distribute[polSubs], x_, FactorTerms]

factor = k12 k3 k4 k5 k9 k10 + k12 k3 k4 k6 k9 k10 - k12 k4 k5 k6 k9 k10 -
  k12 k4 k62 k9 k10 - k12 k4 k5 k6 k102 - k12 k4 k62 k102 - k12 k4 k5 k7 k102 - k12 k4 k6 k7 k102 -
  k1 k2 k3 k42 k8 k11 - k1 k32 k42 k8 k11 + k1 k2 k42 k6 k8 k11 + k1 k3 k42 k6 k8 k11 -
  k12 k3 k4 k5 k10 k11 - k12 k3 k4 k6 k10 k11 - k1 k2 k42 k6 k10 k11 - k1 k3 k42 k6 k10 k11 -
  k1 k2 k42 k7 k10 k11 - k1 k3 k42 k7 k10 k11 - k12 k4 k5 k7 k10 k11 - k12 k4 k6 k7 k10 k11 -
  k1 k2 k3 k42 k112 - k1 k32 k42 k112 - k1 k2 k42 k7 k112 - k1 k3 k42 k7 k112;

Factor[factor]

term = k1 k3 k5 k9 k10 + k1 k3 k6 k9 k10 - k1 k5 k6 k9 k10 - k1 k62 k9 k10 -
  k1 k5 k6 k102 - k1 k62 k102 - k1 k5 k7 k102 - k1 k6 k7 k102 - k2 k3 k4 k8 k11 - k32 k4 k8 k11 +
  k2 k4 k6 k8 k11 + k3 k4 k6 k8 k11 - k1 k3 k5 k10 k11 - k1 k3 k6 k10 k11 - k2 k4 k6 k10 k11 -
  k3 k4 k6 k10 k11 - k2 k4 k7 k10 k11 - k3 k4 k7 k10 k11 - k1 k5 k7 k10 k11 -
  k1 k6 k7 k10 k11 - k2 k3 k4 k112 - k32 k4 k112 - k2 k4 k7 k112 - k3 k4 k7 k112;

simpTerm = FullSimplify[term]

simplerTerm = Distribute[simpTerm / (k1 * k4)] /. {(k2 + k3) / k1 → M1, (k5 + k6) / k4 → M2}

```

This above term larger than 0 should be the necessary condition.

```
condition = simplerTerm > 0
```

By mannual simplifying the term, we can have:

```

simpleCond = (k3 - k6) * (M2 * k9 * k10 - M1 * k8 * k11) >
  (k11 * M1 + k10 * M2) * ((k6 * k10 + k3 * k11) + k7 * (k10 + k11))

left = (k3 - k6) * (M2 * k9 * k10 - M1 * k8 * k11) /. {M1 → (k2 + k3) / k1, M2 → (k5 + k6) / k4}

right = (k11 * M1 + k10 * M2) * ((k6 * k10 + k3 * k11) + k7 * (k10 + k11)) /.
  {M1 → (k2 + k3) / k1, M2 → (k5 + k6) / k4}

```

To fulfill the assumption of thermodynamic conditions for the reversible reactions, we have the constraint:

$$\frac{k_1 k_{10}}{k_2 k_{11}} = \frac{k_4 k_8}{k_5 k_9}.$$

This will give us a even simple condition. Then we will example how will this condition result in the parameter space for multistationarity.

```

oriCond = simpleCond /. {M1 → (k2 + k3) / k1, M2 → (k5 + k6) / k4}

```

$$(k_3 - k_6) \left(\frac{(k_5 + k_6) k_9 k_{10}}{k_4} - \frac{(k_2 + k_3) k_8 k_{11}}{k_1} \right) > \left(\frac{(k_5 + k_6) k_{10}}{k_4} + \frac{(k_2 + k_3) k_{11}}{k_1} \right) (k_6 k_{10} + k_3 k_{11} + k_7 (k_{10} + k_{11}))$$

```

Simplify[oriCond, Assumptions →  $\frac{k_1 k_{10}}{k_2 k_{11}} == \frac{k_4 k_8}{k_5 k_9}$ ]

```

$$\frac{(k_3 - k_6) (k_1 k_6 k_9 k_{10} - k_3 k_4 k_8 k_{11})}{k_1 k_4} > \left(\frac{(k_5 + k_6) k_{10}}{k_4} + \frac{(k_2 + k_3) k_{11}}{k_1} \right) ((k_6 + k_7) k_{10} + (k_3 + k_7) k_{11})$$

Better to do it manually, then we have the condition with thermodynamic constraint:

```

thermoCond =

```

$$(k_3 - k_6) (k_6 k_2 - k_3 k_5) > \left(\frac{k_2}{k_9} \times \frac{k_5^2 + k_6}{k_5} + \frac{k_5}{k_8} \times \frac{k_2^2 + k_3}{k_2} \right) ((k_6 + k_7) k_{10} + (k_3 + k_7) k_{11})$$

$$(k_3 - k_6) (-k_3 k_5 + k_2 k_6) > \left(\frac{(k_2^2 + k_3) k_5}{k_2 k_8} + \frac{k_2 (k_5^2 + k_6)}{k_5 k_9} \right) ((k_6 + k_7) k_{10} + (k_3 + k_7) k_{11})$$

Fromt the above condition, we can get some general idea that in order to satisfy the thermodynamic condition we should have:

Necessarily:

$$k_3 > k_6 \text{ and } k_2 > k_5$$

or

$$k_3 < k_6 \text{ and } k_5 > k_2$$

With additional:

$$k_8, k_9 \gg k_{10}, k_{11} \text{ and } k_7, k_{10}, k_{11} \approx 0$$

Sampling the parameters (without thermodynamic constraint)

Here we try to sampling the parameters by enforcing the thermodynamic constraint. The parameters are sampled in biologically meaningful ranges.

(NewKernel) In[546]:=

```

ClearAll["Global`*"];
A = Table[0, {11}, {6}];

```

```

A[[1]][[1]] = -1;
A[[1]][[3]] = -1;
A[[1]][[5]] = 1;
A[[2]] = -A[[1]];
A[[3]][[3]] = 1; A[[3]][[2]] = 1; A[[3]][[5]] = -1;
A[[4]][[1]] = -1;
A[[4]][[4]] = -1;
A[[4]][[6]] = 1;
A[[5]] = -A[[4]];
A[[6]][[4]] = 1;
A[[6]][[2]] = 1;
A[[6]][[6]] = -1;
A[[7]][[2]] = -1;
A[[7]][[1]] = 1;
A[[8]][[3]] = -1; A[[8]][[4]] = 1; A[[9]] = -A[[8]];
A[[10]][[5]] = -1; A[[10]][[6]] = 1; A[[11]] = -A[[10]];
stoiM = Transpose[A];
(* Now we construct the rate vector *)
ks = {k1 × x3 × x1, k2 × x5, k3 × x5, k4 × x4 × x1,
      k5 × x6, k6 × x6, k7 × x2, k8 × x3, k9 × x4, k10 × x5, k11 × x6};
ssEqns = stoiM.ks;
mC = RowReduce[NullSpace[A]];
subEqns = {ssEqns[[2]], ssEqns[[4]],
           ssEqns[[5]], ssEqns[[6]], x1 + x2 + x5 + x6 - T1, x3 + x4 + x5 + x6 - T2};
jacobian = D[subEqns, {{x1, x2, x3, x4, x5, x6}}];
detJ = Collect[Distribute[Det[jacobian]], {x1, x2, x3, x4, x5, x6}];
solution =
  Solve[{{subEqns[[1]], subEqns[[2]], subEqns[[3]], subEqns[[4]]} == 0},
        {x2, x4, x5, x6}];
detSubs = Replace[detJ, solution[[1]], {0, Infinity}];
(* Equivilant to detSubs=detJ/.solution[[1]]; *)
polSubs = Numerator[Together[detSubs]];
finalSubs = Collect[Distribute[polSubs], x_, FactorTerms];
(*The above code is the same as first section*)

bistableKs = {};
bistableParSets = {};
SeedRandom[];
Timing[
  Do[{
    rands = Exp[RandomReal[{0, 1}, 11] * 2 * Log[1000] - Log[1000]];
    k1 = rands[[1]];
    k2 = rands[[2]];
    k3 = rands[[3]];
    k4 = rands[[4]];
    k5 = rands[[5]];
    k6 = rands[[6]];
    k7 = rands[[7]];
    k8 = rands[[8]];
    k9 = rands[[9]];
    k10 = rands[[10]];
    k11 = rands[[11]];
    left = (k3 - k6)  $\left( \frac{(k5 + k6) k9 k10}{k4} - \frac{(k2 + k3) k8 k11}{k1} \right)$ ;
    right =  $\left( \frac{(k5 + k6) k10}{k4} + \frac{(k2 + k3) k11}{k1} \right) (k6 k10 + k3 k11 + k7 (k10 + k11))$ ;
    If[left > right, {
      AppendTo[bistableKs, {k1, k2, k3, k4, k5, k6, k7, k8, k9, k10, k11}];
    }
  }

```

```

(*counter=1;hitQ=0;
While[hitQ==0&&counter<=1000,{
  x1=RandomReal[{0,1000}];
  finalSol=NSolve[finalSubs==0/.{k1→k1,k2→k2,k3→k3,k4→k4,k5→k5,
    k6→k6,k7→k7,k8→k8,k9→k9,k10→k10,k11→k11,x1→x1},{x3}];
  x3=x3/.finalSol[[1]];
  realSol=solution/.{k1→k1,k2→k2,k3→k3,k4→k4,k5→k5,
    k6→k6,k7→k7,k8→k8,k9→k9,k10→k10,k11→k11,x1→x1,x3→x3};
  T1=(x1+x2+x5+x6)/.Flatten[Append[{x1→x1,x3→x3},realSol[[1]]]];
  T2=(x3+x4+x5+x6)/.Flatten[Append[{x1→x1,x3→x3},realSol[[1]]]];
  If[T1<=1000&&T2<=1000,{
    AppendTo[bistableParSets,
      {k1,k2,k3,k4,k5,k6,k7,k8,k9,k10,k11,T1,T2,left,right}];
    hitQ=1;
  }];
  counter++;
}];*)
}],{i,10000}];
]

```

(NewKernel) Out[568]=
{0.336731, Null}

(NewKernel) In[545]:=
Length[bistableKs]

(NewKernel) Out[545]=
14 325

(NewKernel) In[490]:=
tKs = Transpose[bistableKs];
km1 = $\frac{tKs[[2]] + tKs[[3]]}{tKs[[1]]}$;
km2 = $\frac{tKs[[5]] + tKs[[6]]}{tKs[[4]]}$;
kd1 = $\frac{tKs[[8]]}{tKs[[9]]}$;
kd2 = $\frac{tKs[[10]]}{tKs[[11]]}$;

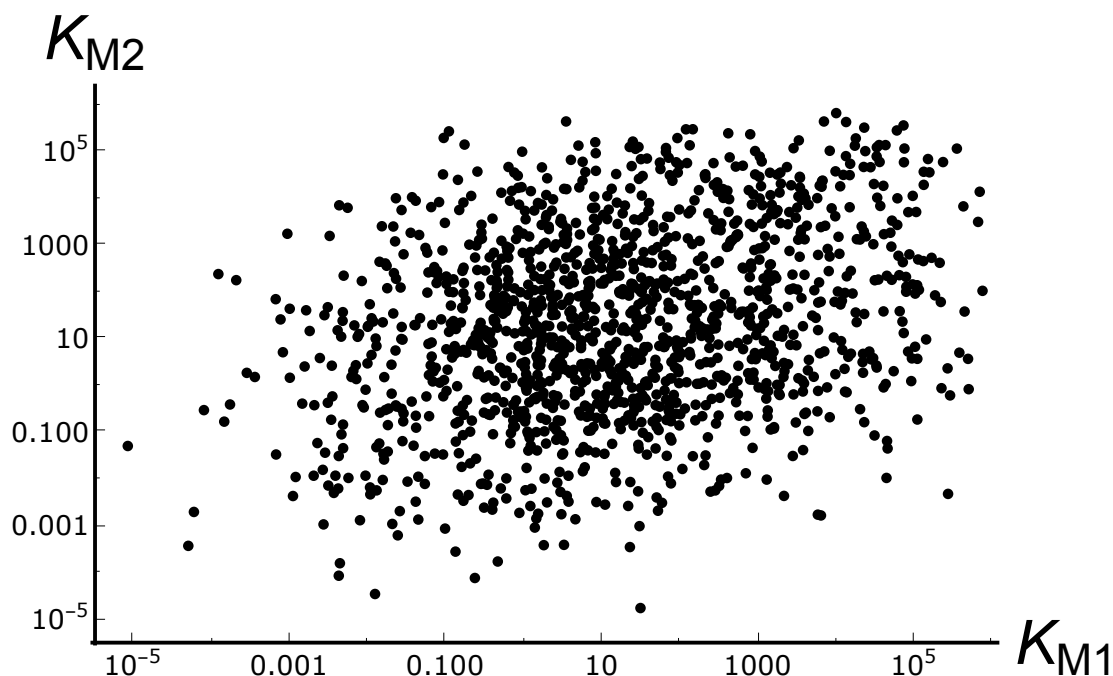
(NewKernel) In[425]:=

```

kmPlot = ListLogLogPlot[Transpose[{km1, km2}], ImageSize → Large, PlotRange → Full,
PlotLabel → None, LabelStyle → {32, GrayLevel[0]}, AxesLabel → {"KM1", "KM2"},
Ticks → Automatic, TicksStyle → Directive["Label", 14],
AxesStyle → Thick, PlotTheme → "Monochrome"]

```

(NewKernel) Out[425]=



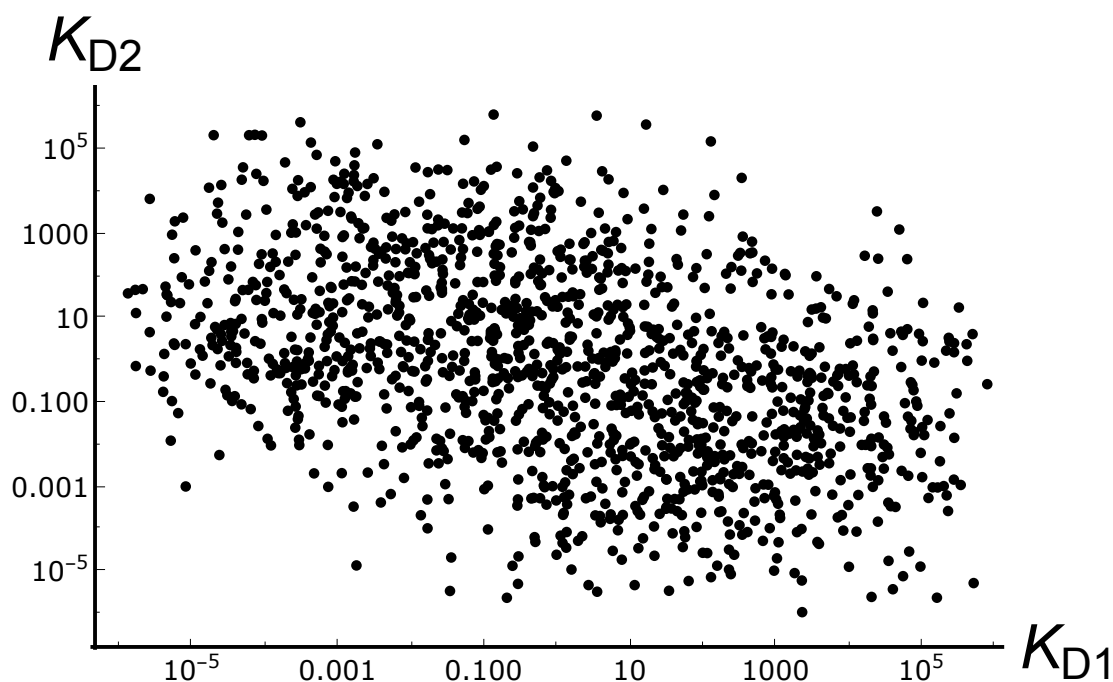
(NewKernel) In[426]:=

```

kdPlot = ListLogLogPlot[Transpose[{kd1, kd2}], ImageSize → Large, PlotRange → Full,
PlotLabel → None, LabelStyle → {32, GrayLevel[0]}, AxesLabel → {"KD1", "KD2"},
Ticks → Automatic, TicksStyle → Directive["Label", 14],
AxesStyle → Thick, PlotTheme → "Monochrome"]

```

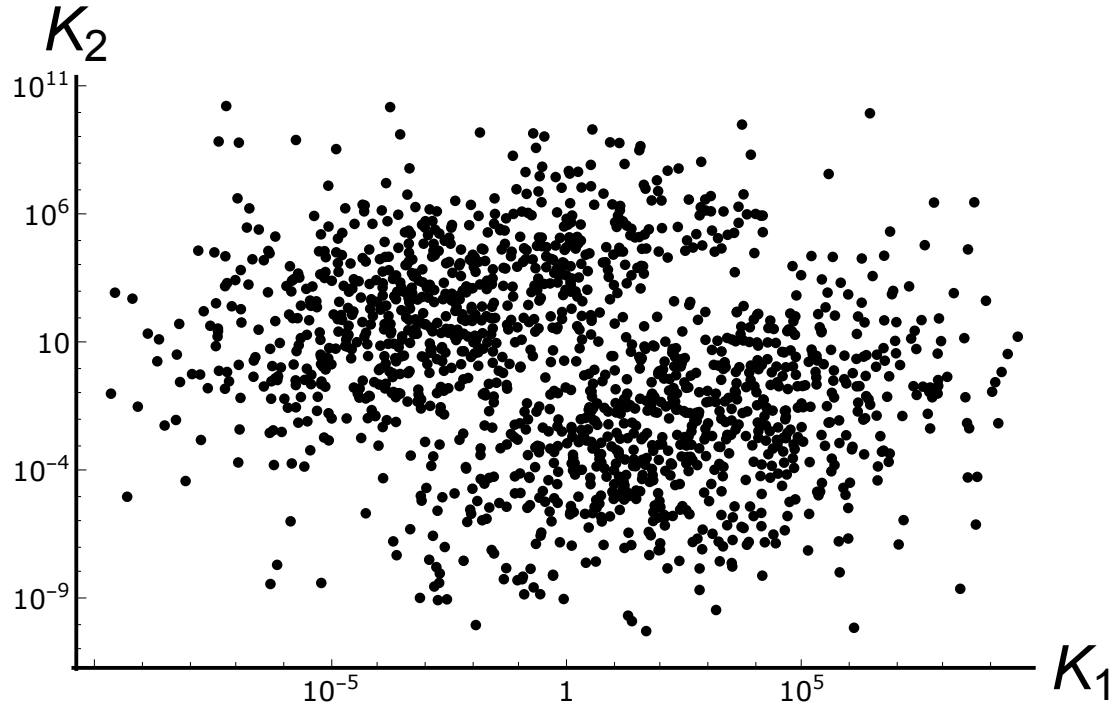
(NewKernel) Out[426]=



(NewKernel) In[497]:=

```
kdPlot = ListLogLogPlot[Transpose[{tKs[[1]] * tKs[[10]] * tKs[[5]] * tKs[[9]],
  tKs[[2]] * tKs[[11]] * tKs[[4]] * tKs[[8]]}], ImageSize → Large,
  PlotRange → Full, PlotLabel → None, LabelStyle → {32, GrayLevel[0]},
  AxesLabel → {"K1", "K2"}, Ticks → Automatic, TicksStyle → Directive["Label", 14],
  AxesStyle → Thick, PlotTheme → "Monochrome"]
```

(NewKernel) Out[497]:=



(NewKernel) In[427]:=

```
model = LinearModelFit[Transpose[{Log10[kd1], Log10[kd2]}], x, x]
```

(NewKernel) Out[427]:=

```
FittedModel[[-0.056131-0.346952x]]
```

(NewKernel) In[428]:=

```
InputForm[model["BestFit"]]
```

(NewKernel) Out[428]/InputForm=

```
-0.0561309658626419 - 0.34695151867837787*x
```

(NewKernel) In[429]:=

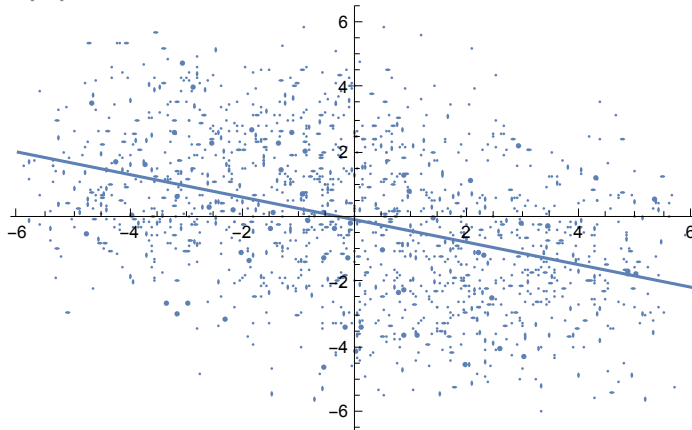
```
model["ParameterTable"]
```

(NewKernel) Out[429]:=

	Estimate	StandardError	t-Statistic	P-Value
1	-0.056131	0.05661	-0.991538	0.321594
x	-0.346952	0.0208559	-16.6356	7.66607×10 ⁻⁵⁷

```
(NewKernel) In[430]:=
Show[ListPlot[Transpose[{Log10[kd1], Log10[kd2]}]],
Plot[model["BestFit"], {x, -6, 6}]]
```

```
(NewKernel) Out[430]=
```



```
(NewKernel) In[301]:=
pc = PrincipalComponents[Log10[bistableKs]];
```

```
(NewKernel) In[300]:=
pc[[Range[1, 10]]];
```

Sampling the parameters (with thermodynamic constraint)

Here we try to sampling the parameters by enforcing the thermodynamic constraint. The parameters are sampled in biologically meaningful ranges.

Sampling the parameters related to thermodynamic constraint $k_1, k_2, k_4, k_5, k_8, k_9, k_{10}, k_{11}$. Using Gamma distribution with $\alpha = 7, \beta = 2$, and then uniformly sample 4 random numbers that sum to 1, these are for k_1, k_5, k_9, k_{10} . Sample another four uniform random number for k_2, k_4, k_8, k_{11} .

For other parameters:

```
(NewKernel) In[593]:=
ClearAll["Global`*"];
A = Table[0, {11}, {6}];
A[[1]][[1]] = -1;
A[[1]][[3]] = -1;
A[[1]][[5]] = 1;
A[[2]] = -A[[1]];
A[[3]][[3]] = 1; A[[3]][[2]] = 1; A[[3]][[5]] = -1;
A[[4]][[1]] = -1;
A[[4]][[4]] = -1;
A[[4]][[6]] = 1;
A[[5]] = -A[[4]];
A[[6]][[4]] = 1;
A[[6]][[2]] = 1;
A[[6]][[6]] = -1;
A[[7]][[2]] = -1;
A[[7]][[1]] = 1;
A[[8]][[3]] = -1; A[[8]][[4]] = 1; A[[9]] = -A[[8]];
A[[10]][[5]] = -1; A[[10]][[6]] = 1; A[[11]] = -A[[10]];
stoiM = Transpose[A];
(* Now we construct the rate vector *)
ks = {k1 × x3 × x1, k2 × x5, k3 × x5, k4 × x4 × x1,
      k5 × x6, k6 × x6, k7 × x2, k8 × x3, k9 × x4, k10 × x5, k11 × x6};
```



```

ssEqns = stoim.ks;
mC = RowReduce[NullSpace[A]];
subsEqns = {ssEqns[[2]], ssEqns[[4]],
  ssEqns[[5]], ssEqns[[6]], x1 + x2 + x5 + x6 - T1, x3 + x4 + x5 + x6 - T2};
jacobian = D[subsEqns, {{x1, x2, x3, x4, x5, x6}}];
detJ = Collect[Distribute[Det[jacobian]], {x1, x2, x3, x4, x5, x6}];
solution =
  Solve[{{subsEqns[[1]], subsEqns[[2]], subsEqns[[3]], subsEqns[[4]]} == 0},
    {x2, x4, x5, x6}];
detSubs = Replace[detJ, solution[[1]], {0, Infinity}];
(* Equivilant to detSubs=detJ/.solution[[1]]; *)
polSubs = Numerator[Together[detSubs]];
finalSubs = Collect[Distribute[polSubs], x_, FactorTerms];
(*The above code is the same as first section*)

bistableKs = {};
bistableParSets = {};
SeedRandom[];
Timing[
  Do[{
    others = Exp[RandomReal[{0, 1}, 3] * 2 * Log[1000] - Log[1000]];
    While[(rand1s = RandomReal[{0, 1}, 4];
      sum = Total@rand1s;
      psum = Switch[True, 0 ≤ # ≤ 1, 1. / 4 * sum^3, 1 < # ≤ 2, 1 / 4 * (-3 * sum^3 + 12 *
        sum^2 - 12 * sum + 4), 2 < # ≤ 3, 1 / 4 * (3 * sum^3 - 24 * sum^2 + 60 * sum -
        44), 3 < # ≤ 4, 1 / 4 * (-sum^3 + 12 * sum^2 - 48 * sum + 64)] & [sum];
      RandomReal ≤ psum)];
    rand23 = RandomVariate[DirichletDistribution[{1, 1, 1, 1}]];
    rand21 = 1 - Total@rand23;
    k1 = Exp[rand1s[[1]] * 2 * Log[1000] - Log[1000]];
    k2 = Exp[sum * rand23[[1]] * 2 * Log[1000] - Log[1000]];
    k3 = others[[1]];
    k4 = Exp[sum * rand23[[2]] * 2 * Log[1000] - Log[1000]];
    k5 = Exp[rand1s[[2]] * 2 * Log[1000] - Log[1000]];
    k6 = others[[2]];
    k7 = others[[3]];
    k8 = Exp[sum * rand23[[3]] * 2 * Log[1000] - Log[1000]];
    k9 = Exp[rand1s[[3]] * 2 * Log[1000] - Log[1000]];
    k10 = Exp[rand1s[[4]] * 2 * Log[1000] - Log[1000]];
    k11 = Exp[sum * rand21 * 2 * Log[1000] - Log[1000]];
    left = (k3 - k6)  $\left( \frac{(k5 + k6) k9 k10}{k4} - \frac{(k2 + k3) k8 k11}{k1} \right)$ ;
    right =  $\left( \frac{(k5 + k6) k10}{k4} + \frac{(k2 + k3) k11}{k1} \right) (k6 k10 + k3 k11 + k7 (k10 + k11))$ ;
    If[left > right, {
      AppendTo[bistableKs, {k1, k2, k3, k4, k5, k6, k7, k8, k9, k10, k11}];
      (*counter=1;hitQ=0;
      While[hitQ==0&&counter≤1000,{
        x1=Exp[-RandomVariate[
          ExponentialDistribution[Log[2]/(-Log[0.0001])]]*1000;
        finalSol=NSolve[finalSubs==0/.{k1→k1,k2→k2,k3→k3,k4→k4,k5→k5,
          k6→k6,k7→k7,k8→k8,k9→k9,k10→k10,k11→k11,x1→x1},{x3}];
        x3=x3/.finalSol[[1]];
        realSol=solution/.{k1→k1,k2→k2,k3→k3,k4→k4,k5→k5,
          k6→k6,k7→k7,k8→k8,k9→k9,k10→k10,k11→k11,x1→x1,x3→x3};
        T1=(x1+x2+x5+x6)/.Flatten[Append[{x1→x1,x3→x3},realSol[[1]]]];
        T2=(x3+x4+x5+x6)/.Flatten[Append[{x1→x1,x3→x3},realSol[[1]]]]];

```

```

    If[0.0001≤T1≤1000&&0.0001≤T2≤1000,{
      AppendTo[bistableParSets,
        {k1,k2,k3,k4,k5,k6,k7,k8,k9,k10,k11,T1,T2,left,right}];
      hitQ=1;
    }];
    counter++;
  }];*)
}];
}, {i, 100 000}];
]

```

```

(NewKernel) Out[615]=
{23.683, Null}

```

```

(NewKernel) In[616]:=
Length[bistableKs]

```

```

(NewKernel) Out[616]=
4378

```

```

(NewKernel) In[455]:=
tKs = Transpose[bistableKs];
km1 =  $\frac{tKs[[2]] + tKs[[3]]}{tKs[[1]]}$ ;
km2 =  $\frac{tKs[[5]] + tKs[[6]]}{tKs[[4]]}$ ;
kd1 =  $\frac{tKs[[8]]}{tKs[[9]]}$ ;
kd2 =  $\frac{tKs[[10]]}{tKs[[11]]}$ ;

```

```

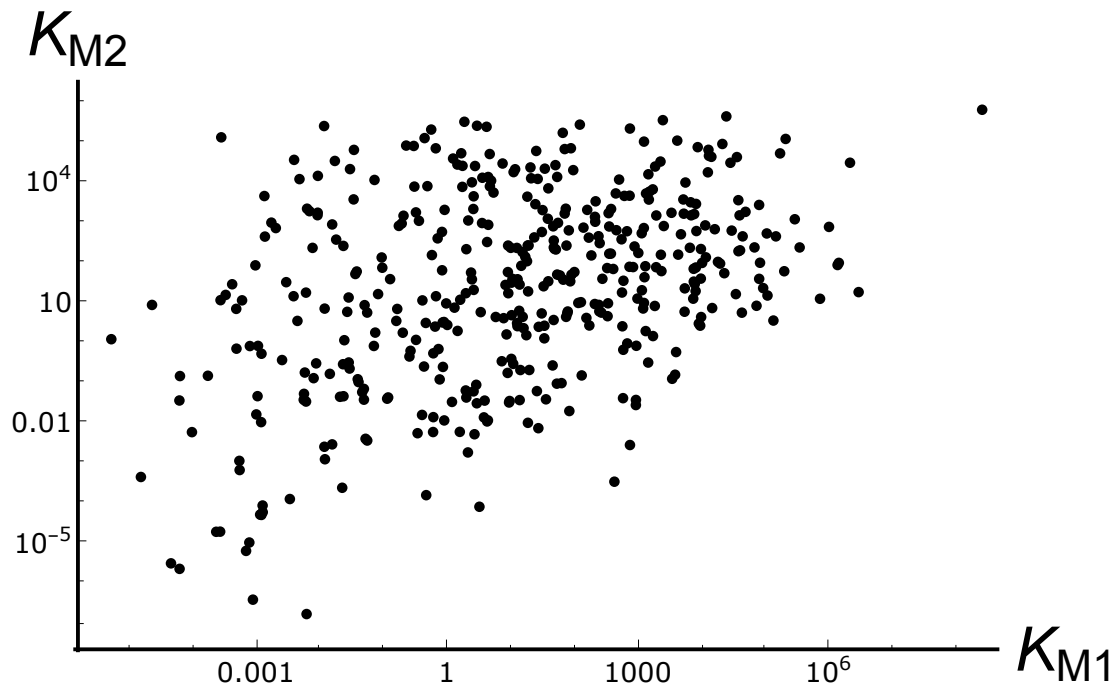
(NewKernel) In[460]:=
kmPlot = ListLogLogPlot[Transpose[{km1, km2}], ImageSize → Large, PlotRange → Full,
  PlotLabel → None, LabelStyle → {32, GrayLevel[0]}, AxesLabel → {"KM1", "KM2"},
  Ticks → Automatic, TicksStyle → Directive["Label", 14],
  AxesStyle → Thick, PlotTheme → "Monochrome"]

```

```

(NewKernel) Out[460]=

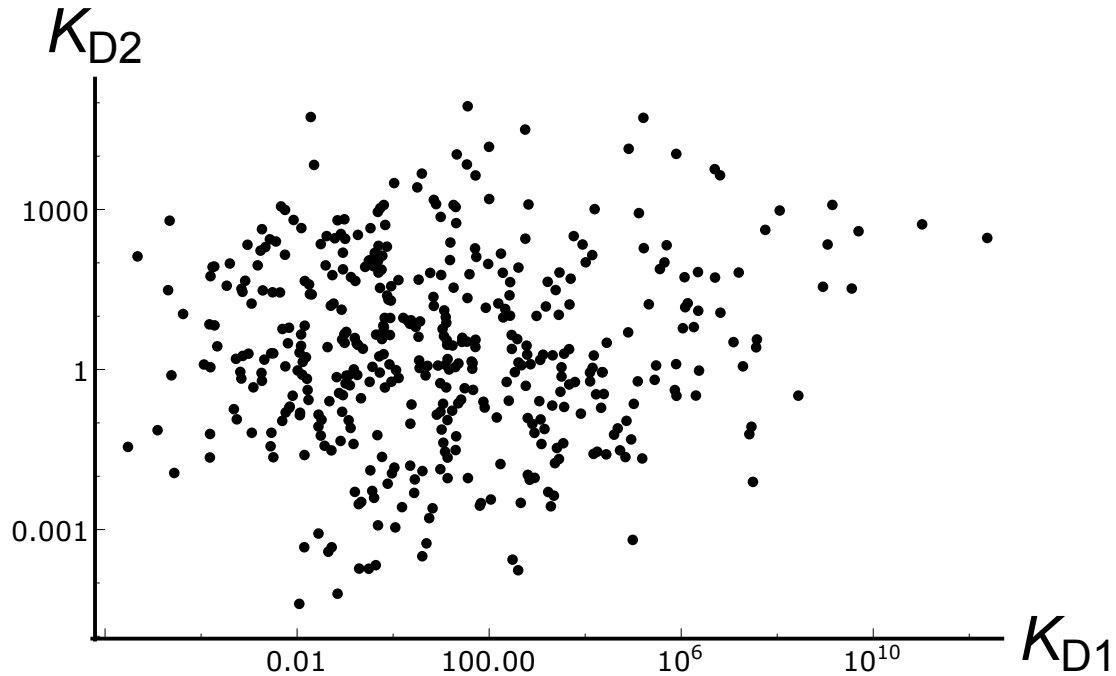
```



(NewKernel) In[461]:=

```
kdPlot = ListLogLogPlot[Transpose[{kd1, kd2}], ImageSize → Large, PlotRange → Full,
  PlotLabel → None, LabelStyle → {32, GrayLevel[0]}, AxesLabel → {"KD1", "KD2"},
  Ticks → Automatic, TicksStyle → Directive["Label", 14],
  AxesStyle → Thick, PlotTheme → "Monochrome"]
```

(NewKernel) Out[461]=



(NewKernel) In[462]:=

```
model = LinearModelFit[Transpose[{Log10[kd1], Log10[kd2]}], x, x]
```

(NewKernel) Out[462]=

```
FittedModel[0.372627+0.0354998 x]
```

(NewKernel) In[463]:=

```
InputForm[model["BestFit"]]
```

(NewKernel) Out[463]/InputForm=

```
0.37262682052688595 + 0.035499771883158025*x
```

(NewKernel) In[464]:=

```
model["ParameterTable"]
```

(NewKernel) Out[464]=

	Estimate	StandardError	t-Statistic	P-Value
1	0.372627	0.0877545	4.24624	0.0000266342
x	0.0354998	0.0274304	1.29417	0.196297

(NewKernel) In[465]:=

```
Show[ListPlot[Transpose[{Log10[kd1], Log10[kd2]}]],
  Plot[model["BestFit"], {x, -6, 6}]]
```

■ Conclusion:

These above results show that the parameter set within biological meaningful ranges can be reached by increasing the sampling size even when enforcing the thermodynamic constraint. Comparing to results from the other document (without enforcing thermodynamic constraint), the parameter space is largely reduced.

Conditions	with thermo	with thermo	without thermo	without thermo
Sampling size	only check bistability	bistability & concentrations	only check bistability	bistability & concentrations
10^4	396	N / A	1448	N / A
10^5	4378	N / A	14 325	N / A

From the sampling results, thermodynamic constraint indeed shrinks the parameter space for bistability in the motif. Although, the concentration is probably not very well sampled, the concentration is trivial for its purpose here.

Test