

Quasi-potentials: A framework for analyzing stochastic dynamical systems in ecology

Quasi-potential Tutorial

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QPot: Quasi-Potential Analysis for Stochastic Differential Equations

A R package that introduces a suite of functions to assist in analysis of SDEs

Key functions

Function	Main arguments	Description
TSTraj()	Deterministic skeleton, σ , T , Δt	Creates a realization (time series) of the stochastic differential equations.
TSPlot()	TSTraj() output	Plots a realization of the stochastic differential equations, with an optional histogram side-plot. Plots can additionally be two-dimensional, which show realizations in (X, Y) -space.
TSDensity()	TSTraj() output	Creates a density plot of a trajectory in (X, Y) -space in one or two dimensions.
QPotential()	Deterministic skeleton, stable equilibria, bounds, mesh (number of divisions along each axis)	Creates a matrix corresponding to a discretized version of the local quasi-potential function for each equilibrium.
QPGlobal()	Local quasi-potential matrices, unstable equilibria	Creates a global quasi-potential surface.
QPIinterp()	Global quasi-potential, (x, y) -coordinates	Evaluates the global quasi-potential at (x, y) .
QPContour()	Global quasi-potential	Creates a contour plot of the quasi-potential.
VecDecomAll()	Global quasi-potential, deterministic skeleton, bounds	Creates three vector fields: the deterministic skeleton, the negative gradient of the quasi-potential, and the remainder vector field. To find each field individually, the functions VecDecomVec(), VecDecomGrad(), or VecDecomRem() can be used.
VecDecomPlot()	Deterministic skeleton, gradient, or remainder field	Creates a vector field plot for the vector, gradient, or remainder field.

Steps for the analysis of Stochastic Differential Equations

- 1) Analyze the deterministic model
Find equilibria, determine stability
- 2) Simulate the equations, include stochasticity
- 3) Calculate the local quasi-potentials
For each stable equilibrium
- 4) Calculate the global quasi-potential
(if only one stable equilibrium, then local = global)
- 5) Visualize the global quasi-potential
- 6) Visualize the vector field decompositions

Our Equations: Lotka-Volterra Competition

Species X

$$\frac{dx}{dt} = r_x x \frac{K_x - x - \alpha_{xy}y}{K_x}$$

Species Y

$$\frac{dy}{dt} = r_y y \frac{K_y - y - \alpha_{yx}x}{K_y}$$

x = state variable, density of species x

y = state variable, density of species y

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α_{xy} = competition coefficient,
converts individuals of species y into species x equivalents

α_{yx} = competition coefficient,
converts individuals of species x into species y equivalents

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Four different outcomes based on parameter values

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Four different outcomes based on parameter values

X wins $K_x > \frac{K_y}{\alpha_{yx}}$ $K_y < \frac{K_x}{\alpha_{xy}}$

Our Equations: Lotka-Volterra Competition

Species X

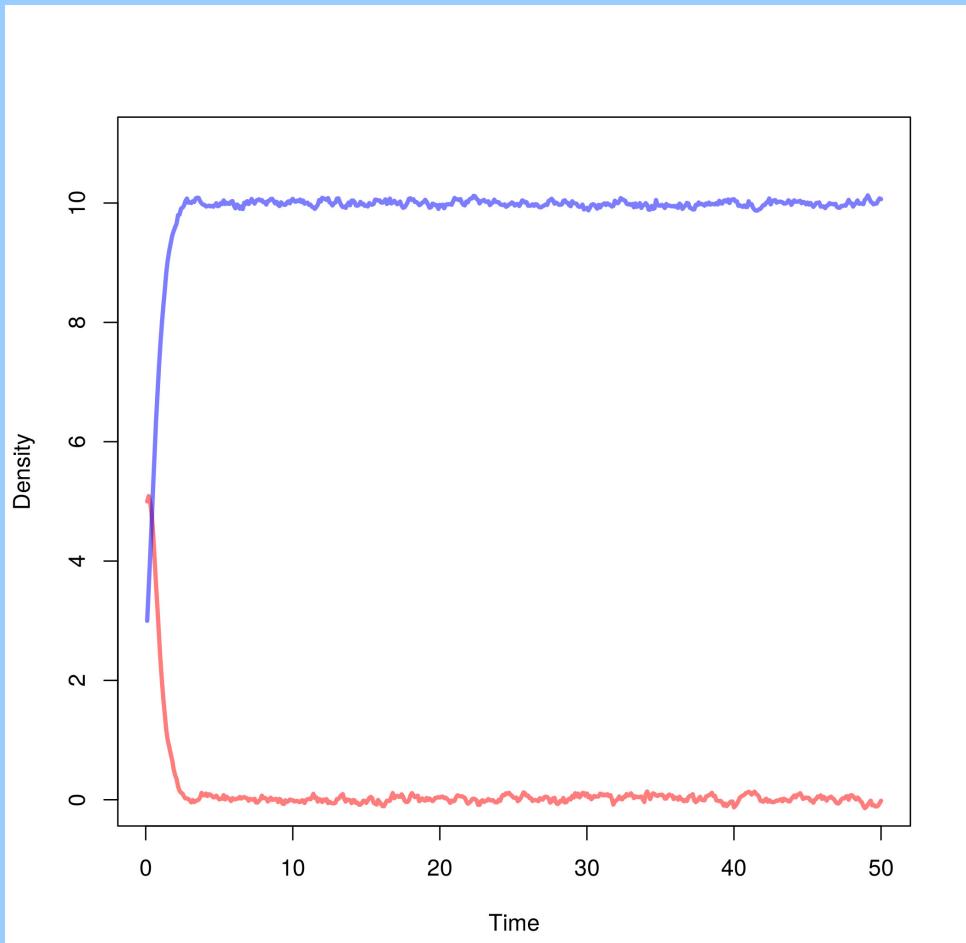
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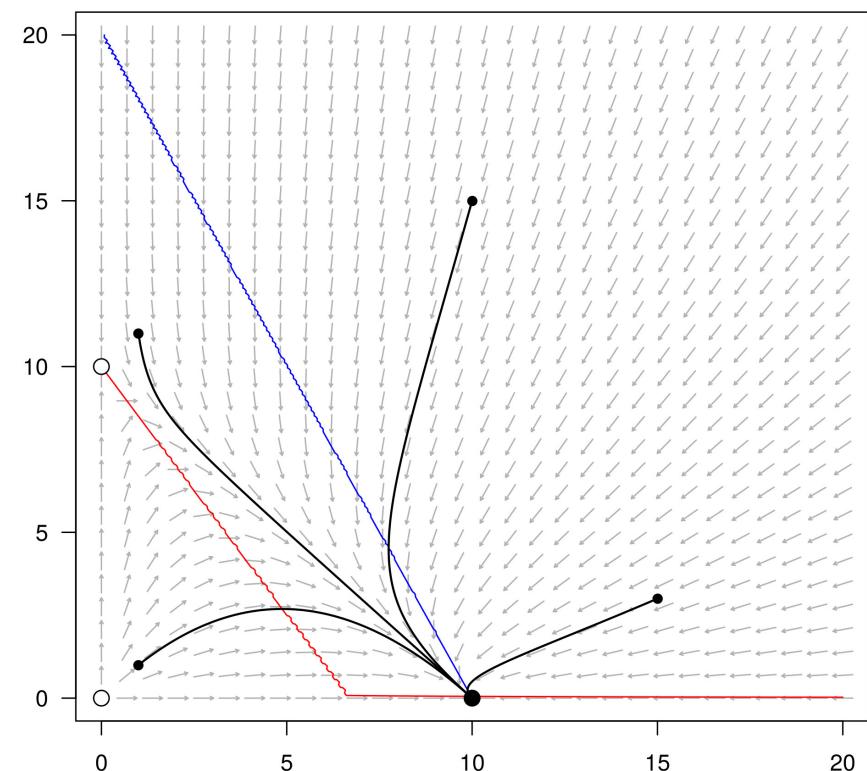
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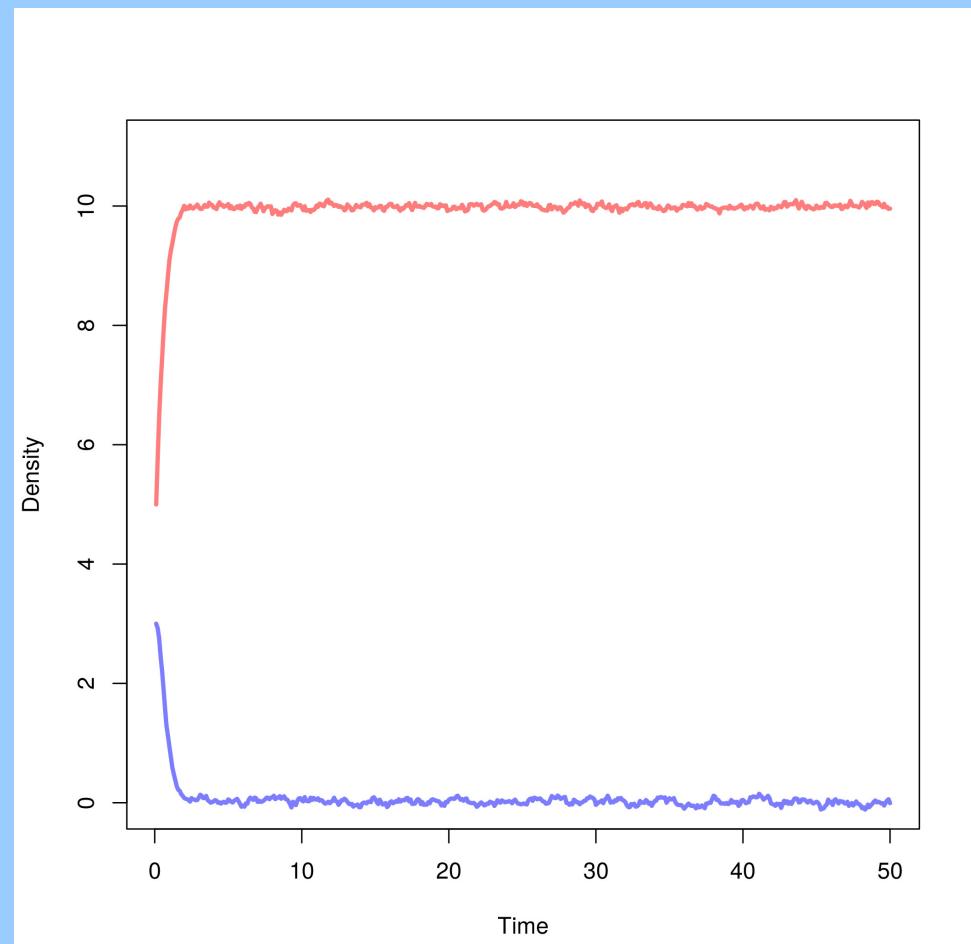
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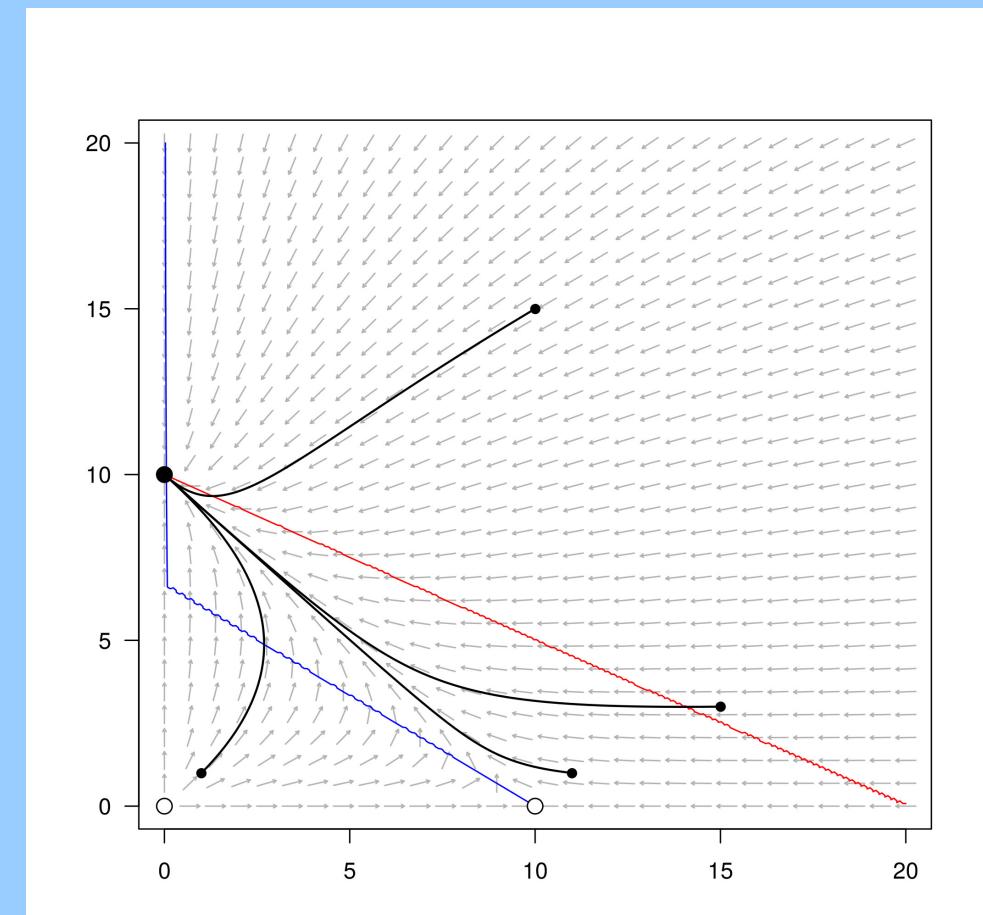
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condition $K_x > \frac{K_y}{\alpha_{yx}}$ $K_y > \frac{K_x}{\alpha_{xy}}$
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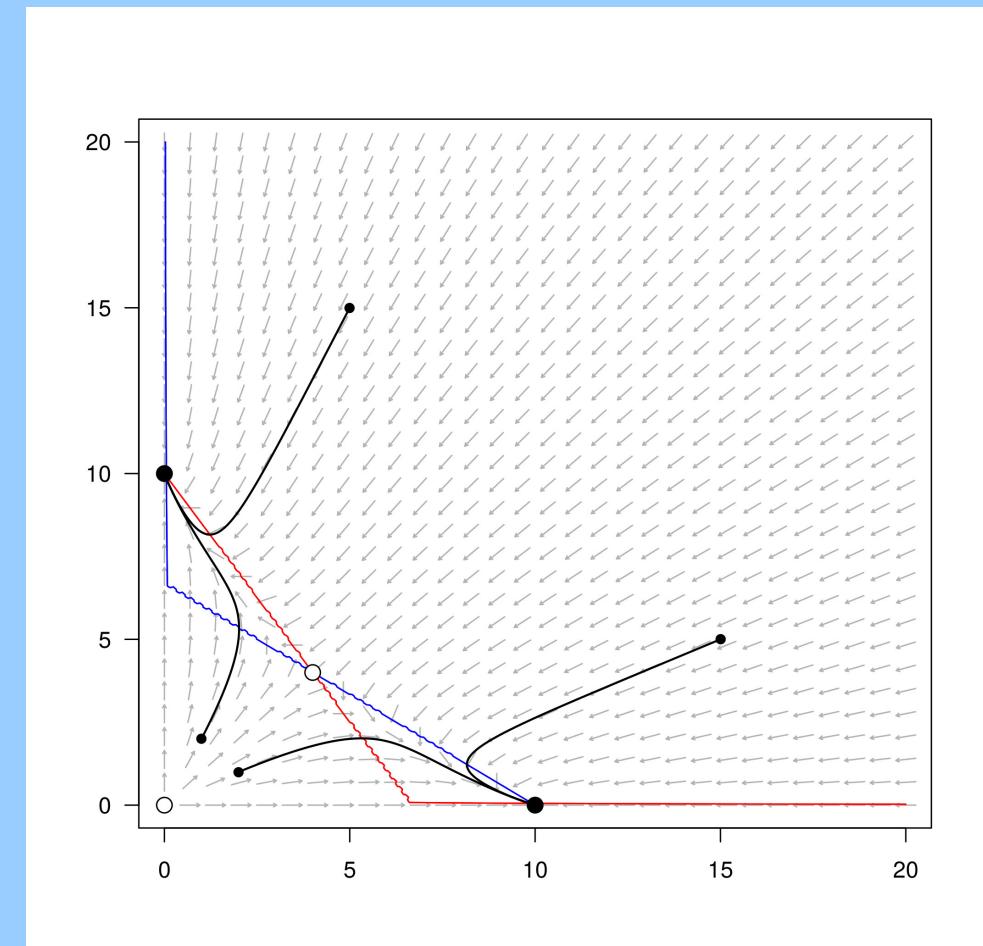
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Initial condition dependence $K_x > \frac{K_y}{\alpha_{yx}}$ $K_y > \frac{K_x}{\alpha_{xy}}$



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Coexist $K_x < \frac{K_y}{\alpha_{yx}}$ $K_y < \frac{K_x}{\alpha_{xy}}$

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Four different outcomes based on parameter values

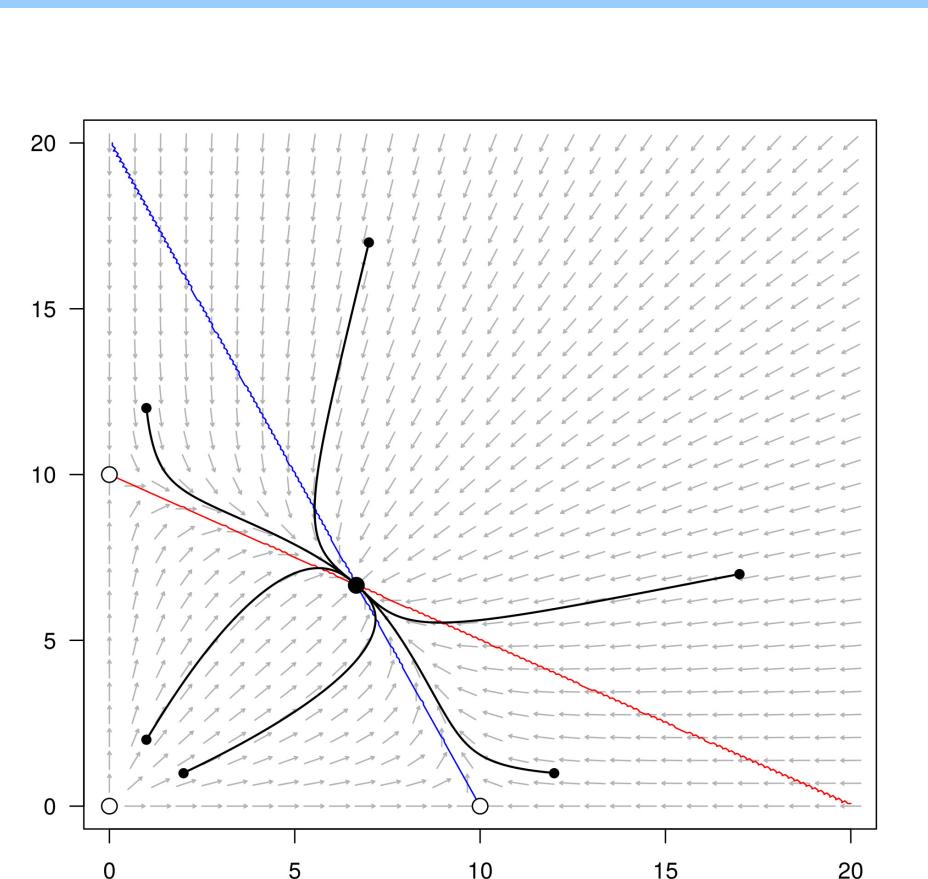
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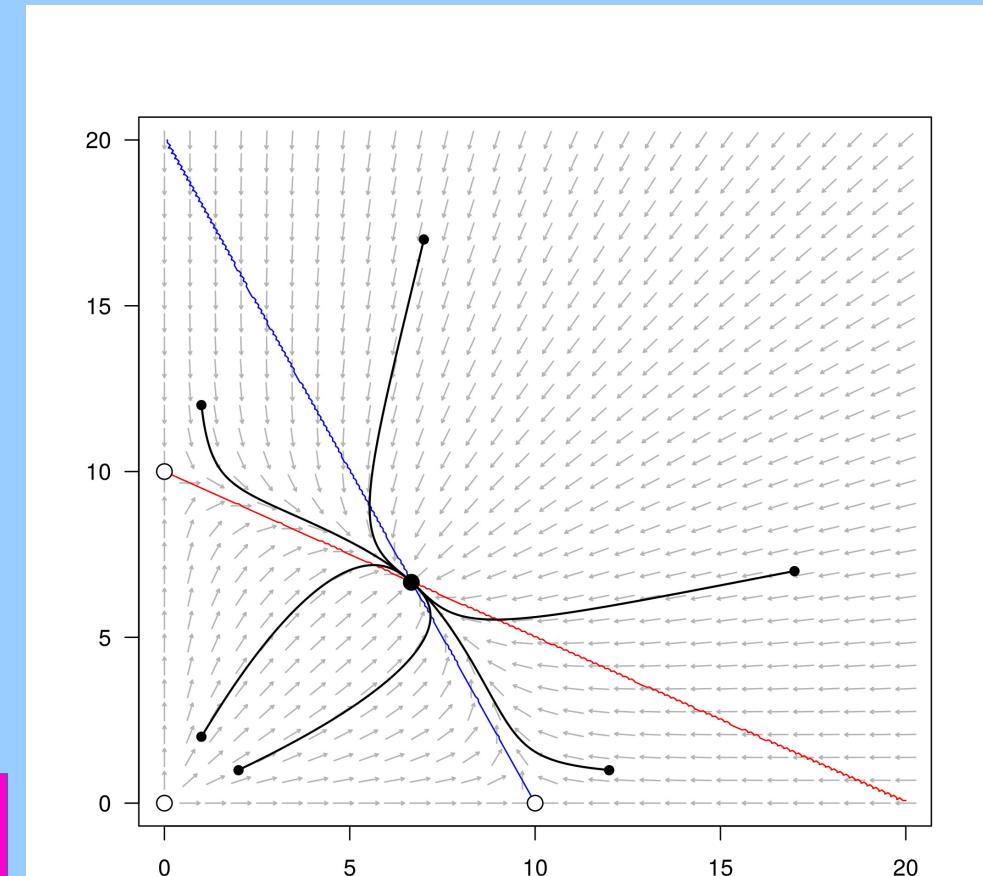
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Coexist $K_x < \frac{K_y}{\alpha_{yx}}$ $K_y < \frac{K_x}{\alpha_{xy}}$



Lotka-Volterra Competition: Coexistence

Coexistence $K_x < \frac{K_y}{\alpha_{yx}}$ $K_y < \frac{K_x}{\alpha_{xy}}$

- 1) Analyze the deterministic model
Find equilibria, determine stability

$$0 = r_x x \frac{K_x - x - \alpha_{xy} y}{K_x}$$

$$0 = r_y y \frac{K_y - y - \alpha_{yx} x}{K_y}$$

(0,0) unstable equilibrium

$$\left(\frac{K_x - \alpha_{xy} K_y}{1 - \alpha_{xy} \alpha_{yx}}, \frac{K_y - \alpha_{yx} K_x}{1 - \alpha_{xy} \alpha_{yx}} \right)$$

(K_x ,0) unstable equilibrium

stable equilibrium

(0, K_y) unstable equilibrium

Lotka-Volterra Competition: Coexistence

Install these packages: (order is wrong in handout)

```
install.packages('Qpot')
install.packages('plot3D')
install.packages('deSolve')
install.packages('phaseR')
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in R, anything after the # is ignored

Lotka-Volterra Competition: Coexistence

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```
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library(phaseR)
```

in R, anything after the # is ignored

Lotka-Volterra Competition: Coexistence

- 1) Analyze the deterministic model
Find equilibria, determine stability

```
# parameters that lead to coexistence
model.state.coexist <- c(rx = 4, ry = 4, kx = 10, ky = 10, alphaxy = 0.5, alphayx = 0.5)

require(phaseR)
competition.model <- function(t, y, parameters){
  x <- y[1]                                # initial state of species x
  y <- y[2]                                # initial state of species y
  rx <- parameters["rx"]                  # growth rate of species x
  ry <- parameters["ry"]                  # growth rate of species y
  alphaxy <- parameters["alphaxy"]        # competition coefficient
  alphayx <- parameters["alphayx"]        # competition coefficient
  kx <- parameters["kx"]                  # carrying capacity species x
  ky <- parameters["ky"]                  # carrying capacity species y
  dx <- rx*x*(kx - x - alphaxy*y)*(1/kx) # equation for species x
  dy <- ry*y*(ky - y - alphayx*x)*(1/ky) # equation for species y
  list(c(dx,dy))
}
```

Lotka-Volterra Competition: Coexistence

1) Analyze the deterministic model
Find equilibria, determine stability

```
# this function will draw the vector field that describes how the densities will change
flowField(deriv = competition.model, x.lim = c(0, 20), y.lim = c(0, 20),
parameters = model.state.coexist, add = F, points = 30)

# Draw the null clines, where dx/dt = 0 or dy/dt = 0
supp.print <- nullclines(deriv = competition.model, x.lim = c(0, 20),
y.lim = c(0, 20), parameters = model.state.coexist,
col = c("blue", "red"), points = 250)

#plot points on graph for stable equilibrium (black) and unstable (white)
points(x = 0, y = 10 , pch = 21 , bg = "white" , cex = 1.5)
points(x = 10, y = 0 , pch = 21 , bg = "white" , cex = 1.5)
points(x = 6.667, y = 6.667 , pch = 21 , bg = "black" , cex = 1.5)
points(x = 0, y = 0 , pch = 21 , bg = "white" , cex = 1.5)

xstart = runif(1,0,20)  #1 random point between 0 and 20 from uniform dist
ystart = runif(1,0,20)  # change this to a value to set start point
traj <- trajectory(competition.model,y0=c(xstart,ystart),
parameters = model.state.coexist , t.end = 250)
```

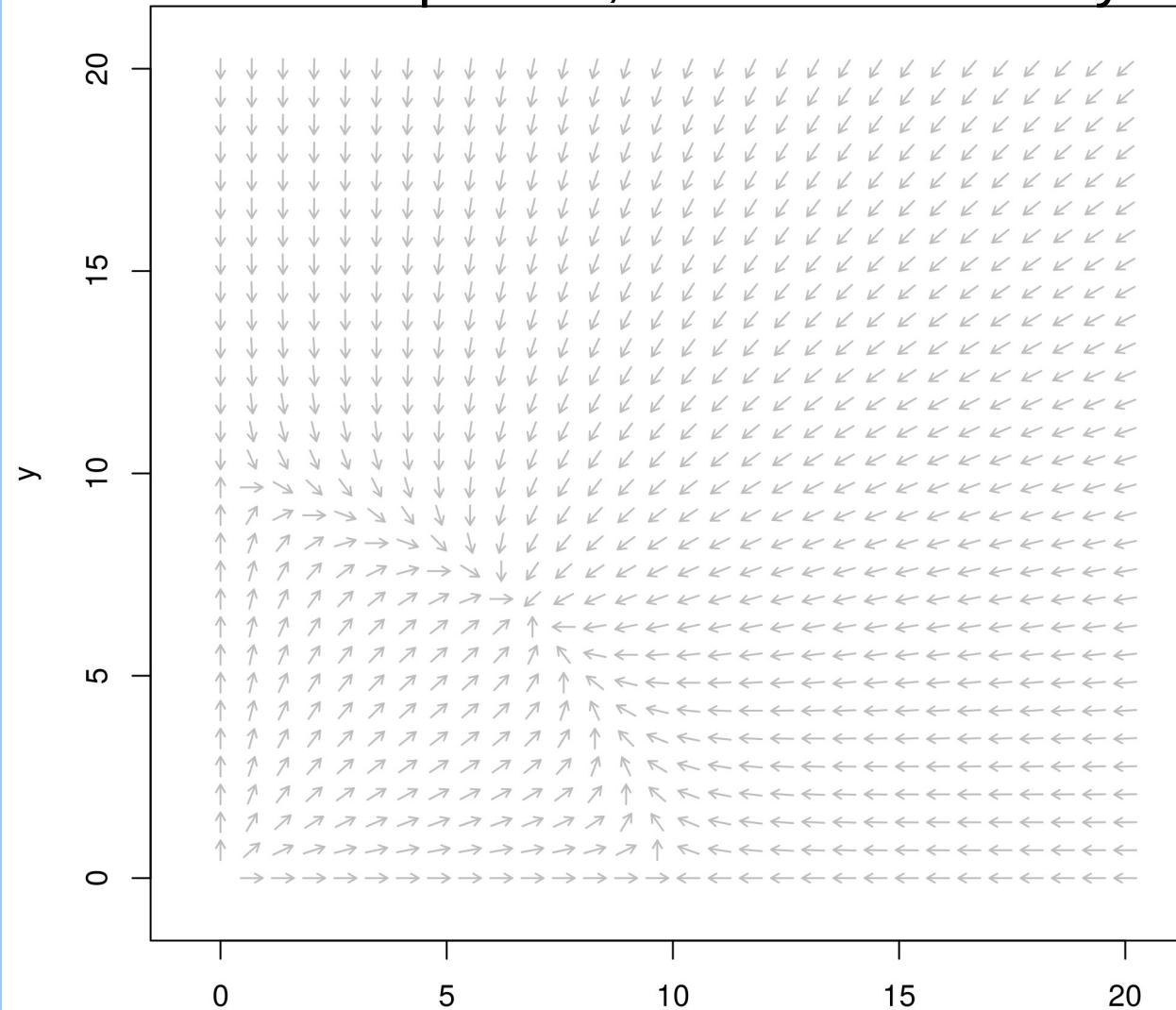
Lotka-Volterra Competition: Coexistence

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Lotka-Volterra Competition: Coexistence

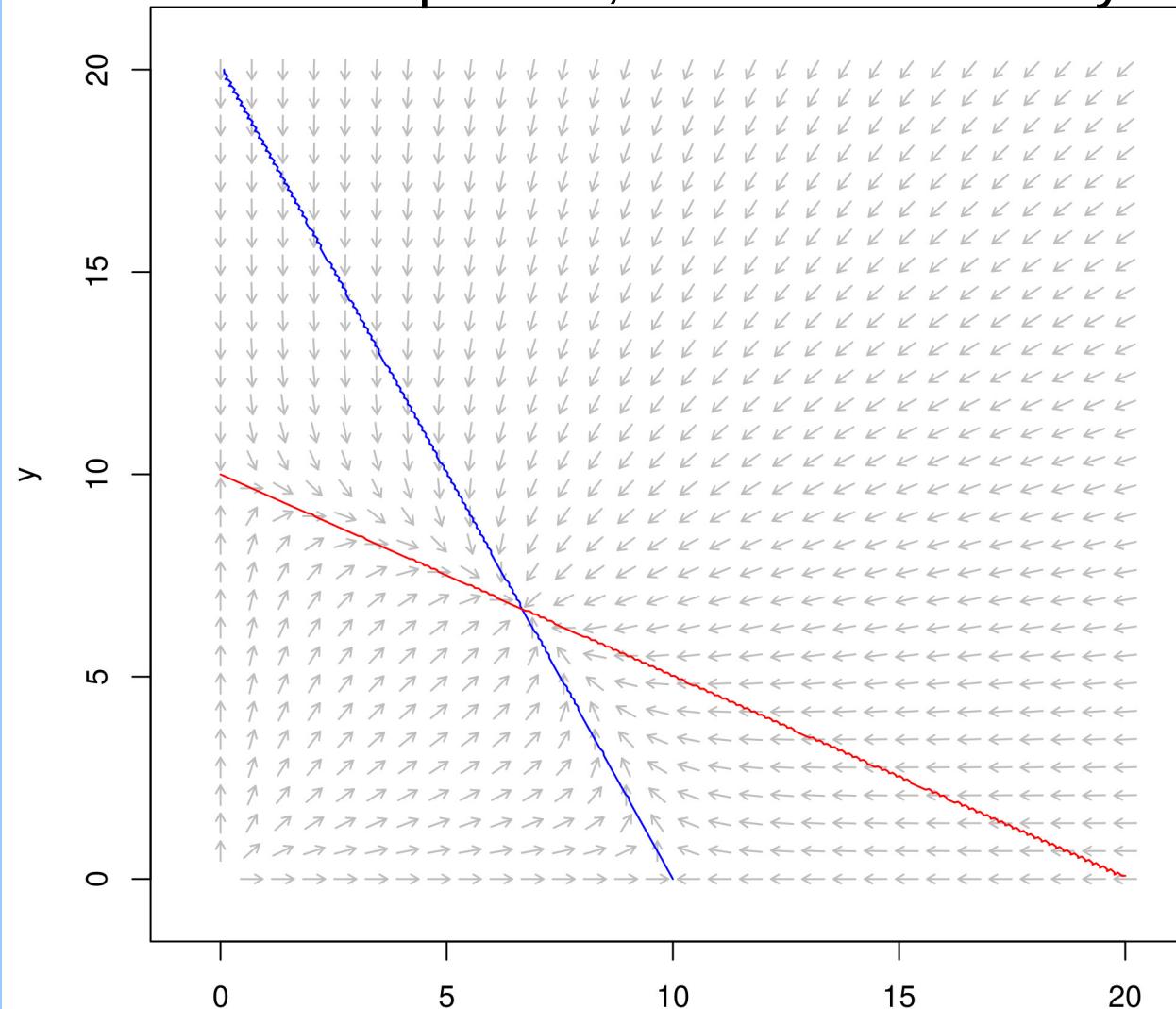
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Lotka-Volterra Competition: Coexistence

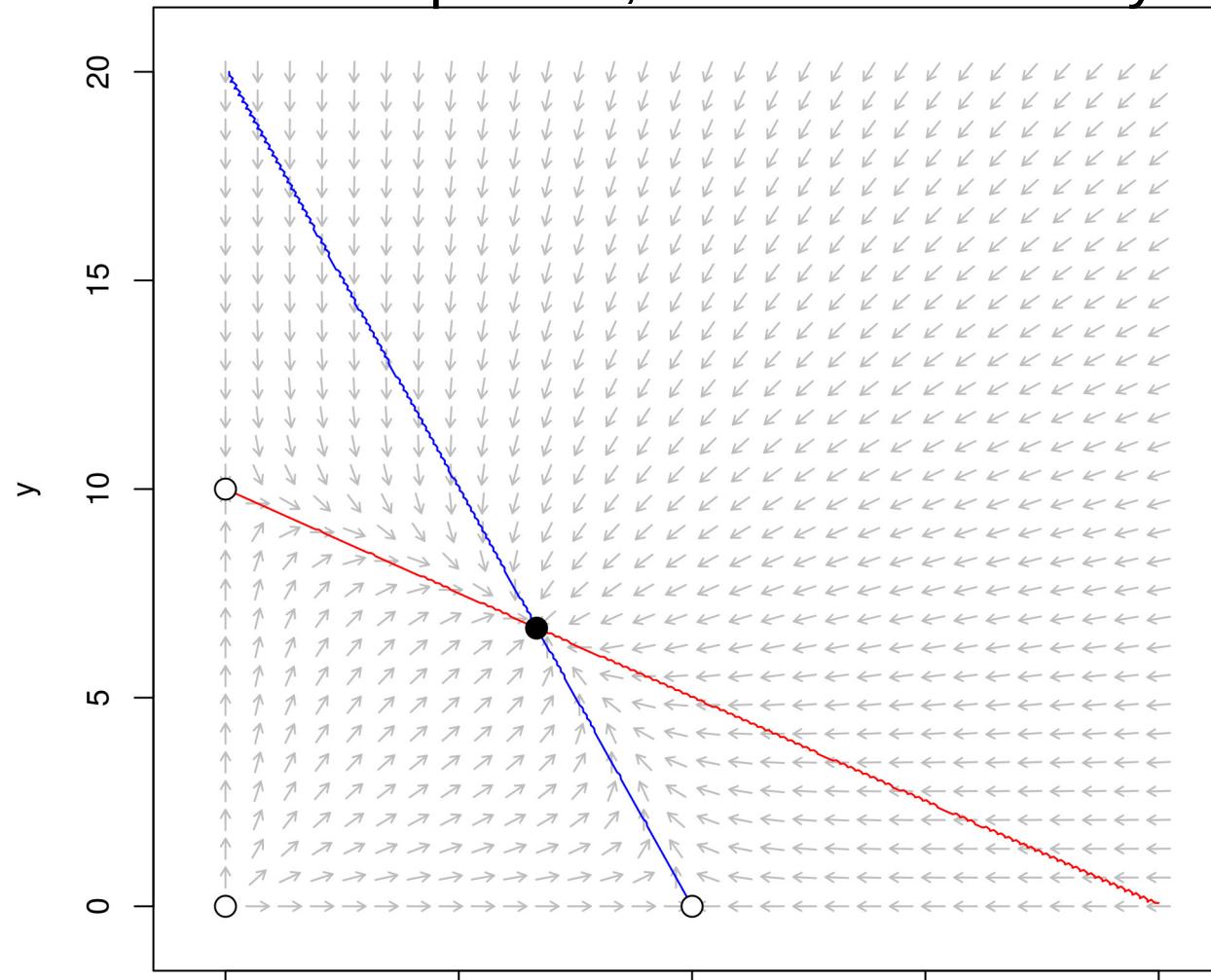
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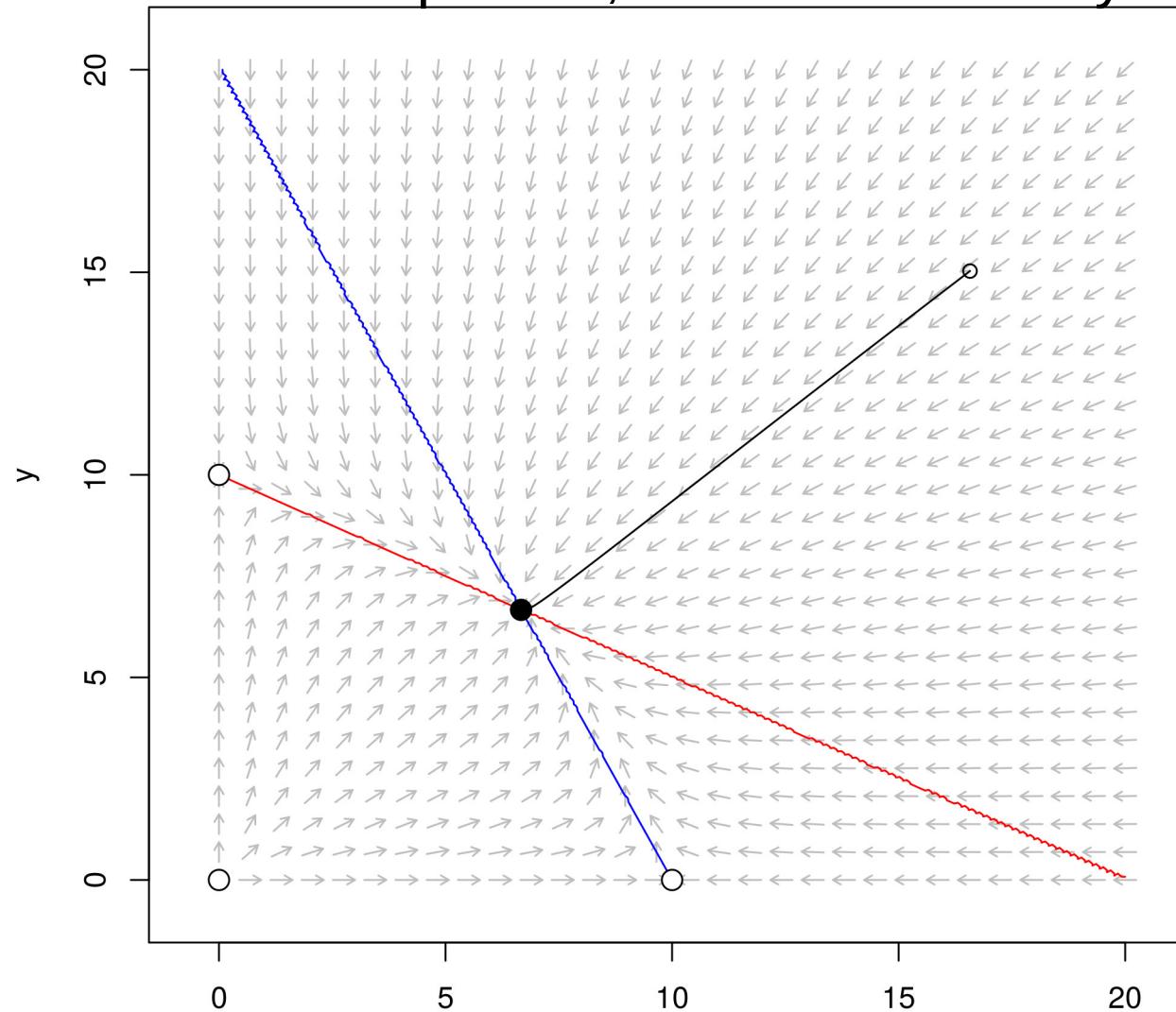
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points(0,10 ,pch = 21 , col = "black", bg = "white" , cex = 1.5)
points(10,0 , pch = 21 , col = "black" , bg = "white" , cex = 1.5)
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Lotka-Volterra Competition: Coexistence

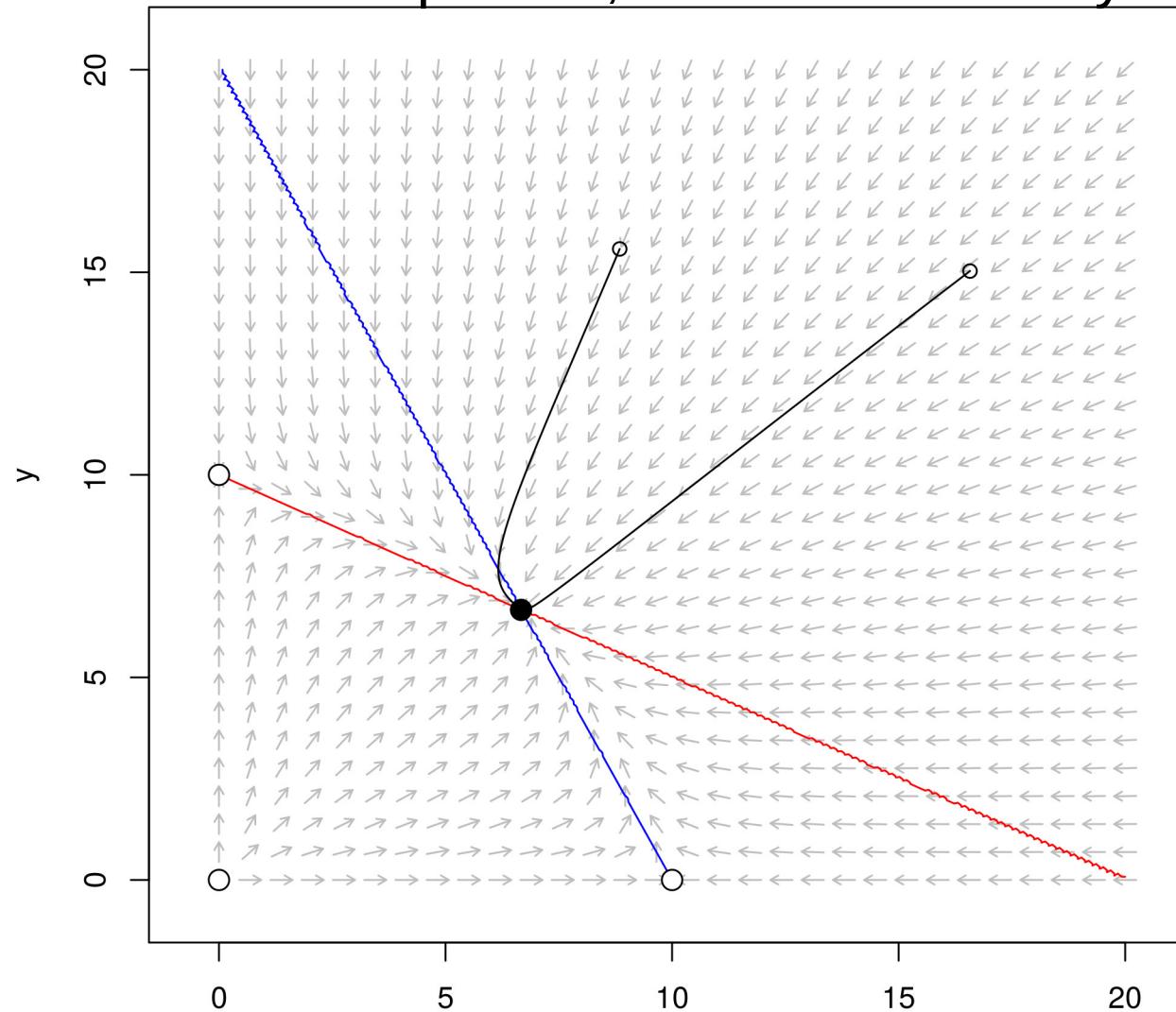
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Lotka-Volterra Competition: Coexistence

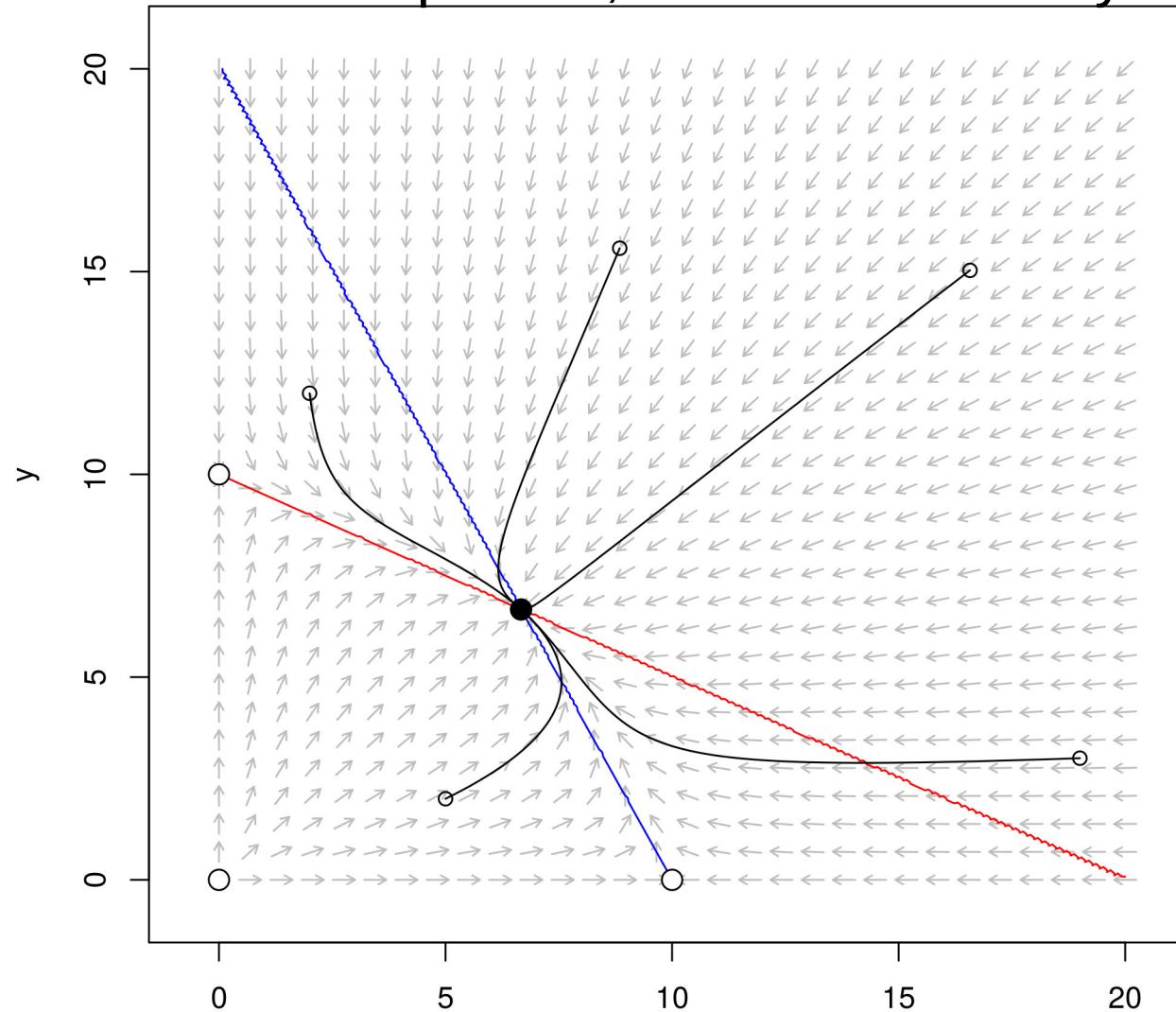
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Lotka-Volterra Competition: Coexistence

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Lotka-Volterra Competition: Coexistence

Coexistence $K_x < \frac{K_y}{\alpha_{yx}}$ $K_y < \frac{K_x}{\alpha_{xy}}$

2) Simulate the equations, include stochasticity

Lotka-Volterra Competition: Coexistence

Coexistence $K_x < \frac{K_y}{\alpha_{yx}}$ $K_y < \frac{K_x}{\alpha_{xy}}$

2) Simulate the equations, include stochasticity

```
dxstring <- 'rx*x*(kx - x - alphaxy*y)*(1/kx)' # L-V competition
```

```
dystring <- 'ry*y*(ky - y - alphayx*x)*(1/ky)' # from earlier
```

```
# Parameters included in step one, but in case you are skipping around
```

```
model.state.coexist <- c(rx = 4, ry = 4, kx = 10, ky = 10, alphaxy = 0.5, alphayx = 0.5)
```

```
dx <- Model2String(model = dxstring, parms = model.state.coexist)
```

```
dy <- Model2String(model = dystring, parms = model.state.coexist)
```

Lotka-Volterra Competition: Coexistence

Coexistence $K_x < \frac{K_y}{\alpha_{yx}}$ $K_y < \frac{K_x}{\alpha_{xy}}$

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```

```
dx <- Model2String(model = dxstring, parms = model.state.coexist)
```

```
dy <- Model2String(model = dystring, parms = model.state.coexist)
```

```
# We need to define some values for our simulation
```

```
init.cond <- c(x = 3, y = 10) # initial state of simulation
```

```
sigma = 0.1 # level of stochasticity, sigma = 0 is deterministic model
```

```
timesteps = 1000 # number of time steps
```

```
deltat = 0.1 # size of time step
```

```
# Run a stochastic simulation, but TSTraj does not return a plot
```

```
timesimulation <- TSTraj(y0 = init.cond, time = timesteps,  
deltat = deltat, x.rhs = dx, y.rhs = dy, sigma = sigma)
```

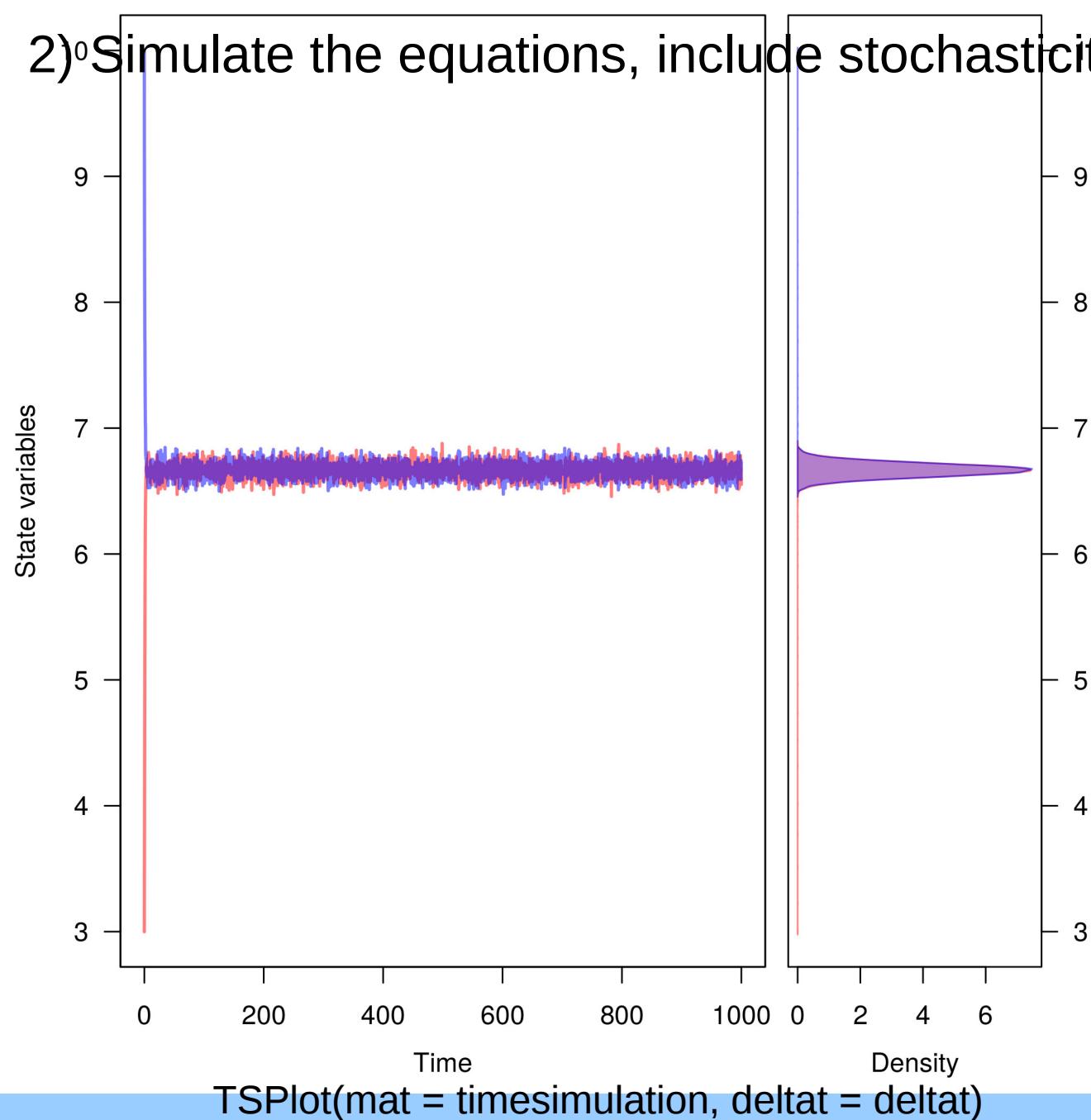
Lotka-Volterra Competition: Coexistence

2) Simulate the equations, include stochasticity

```
TSPlot(mat = timesimulation, deltat = deltat)
```

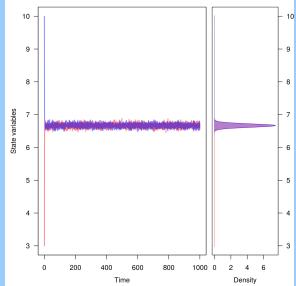
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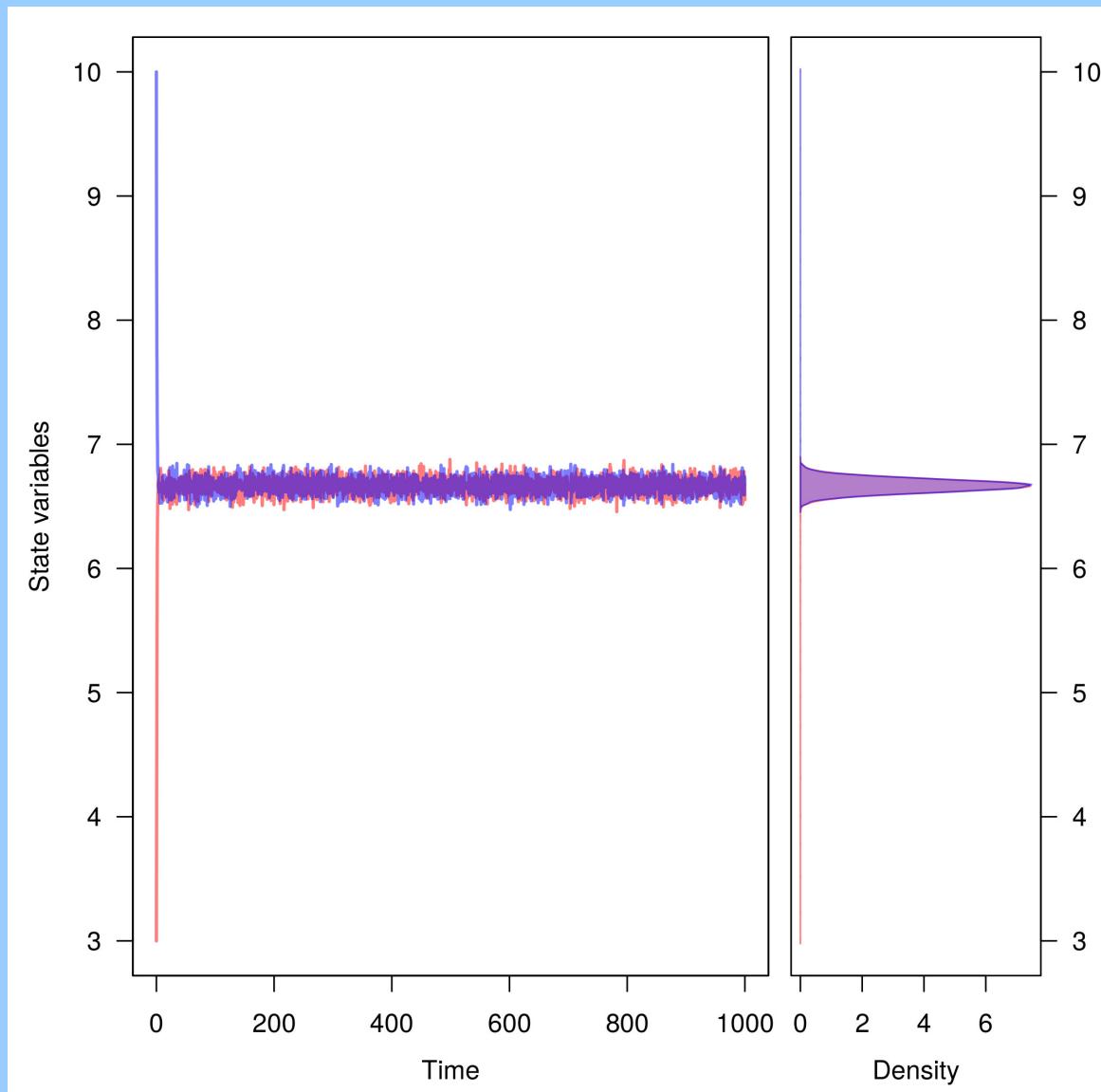


Lotka-Volterra Competition: Coexistence

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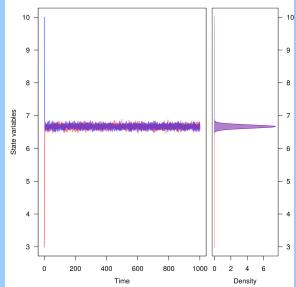


TSPlot(timesimulation,
deltat = deltat)

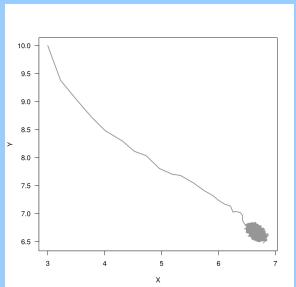


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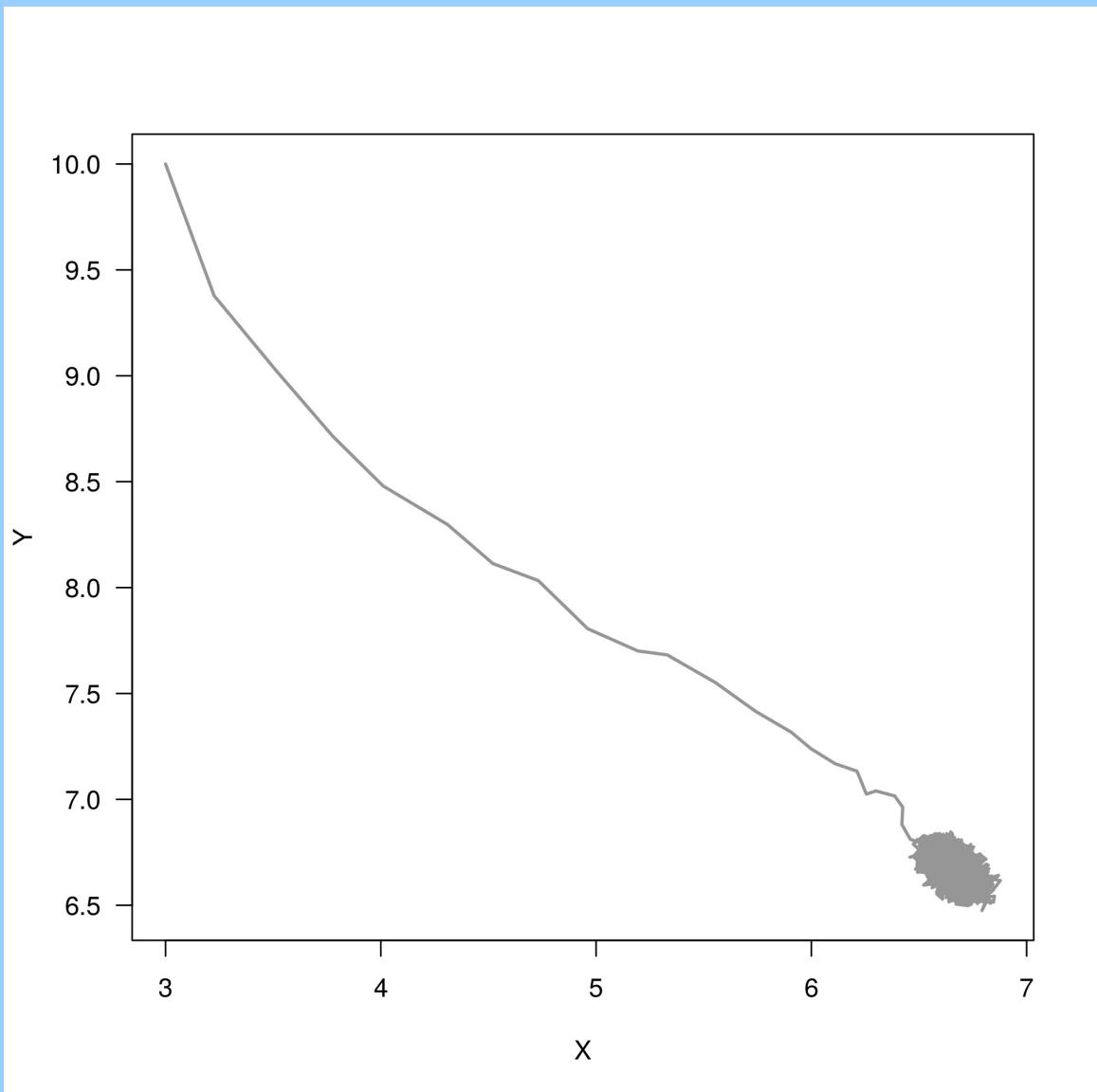
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`TSPlot(timesimulation,
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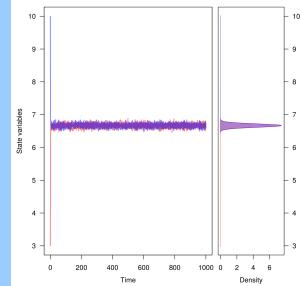


`TSPlot(timesimulation,
deltat = model.deltat,
dim = 2)`

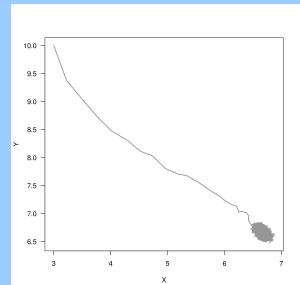


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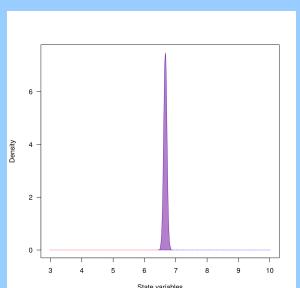
2) Simulate the equations, include stochasticity



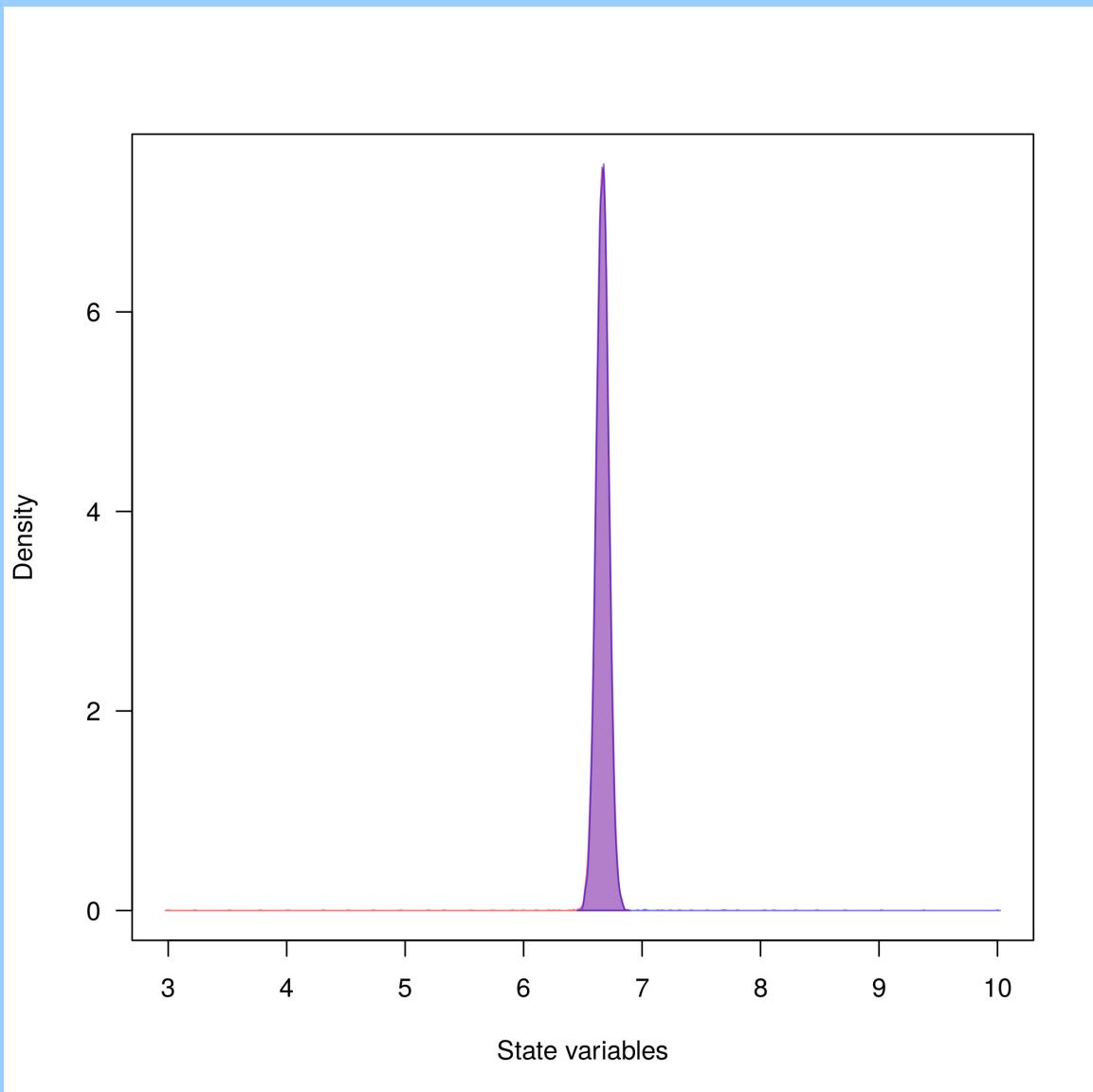
`TSPlot(timesimulation,
deltat = deltat)`



`TSPlot(timesimulation,
deltat = model.deltat,
dim = 2)`

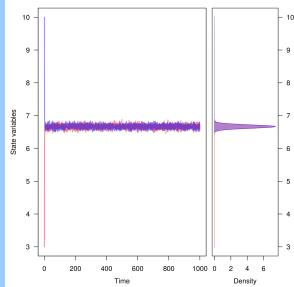


`TSDensity(timesimulation,
dim = 1)`

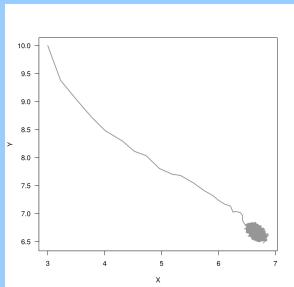


Lotka-Volterra Competition: Coexistence

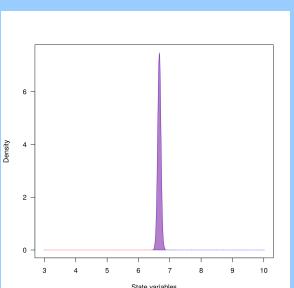
2) Simulate the equations, include stochasticity



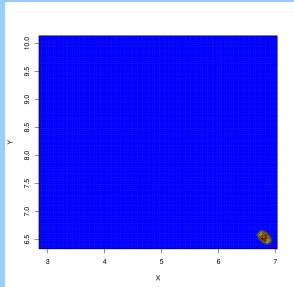
`TSPlot(timesimulation,
deltat = deltat)`



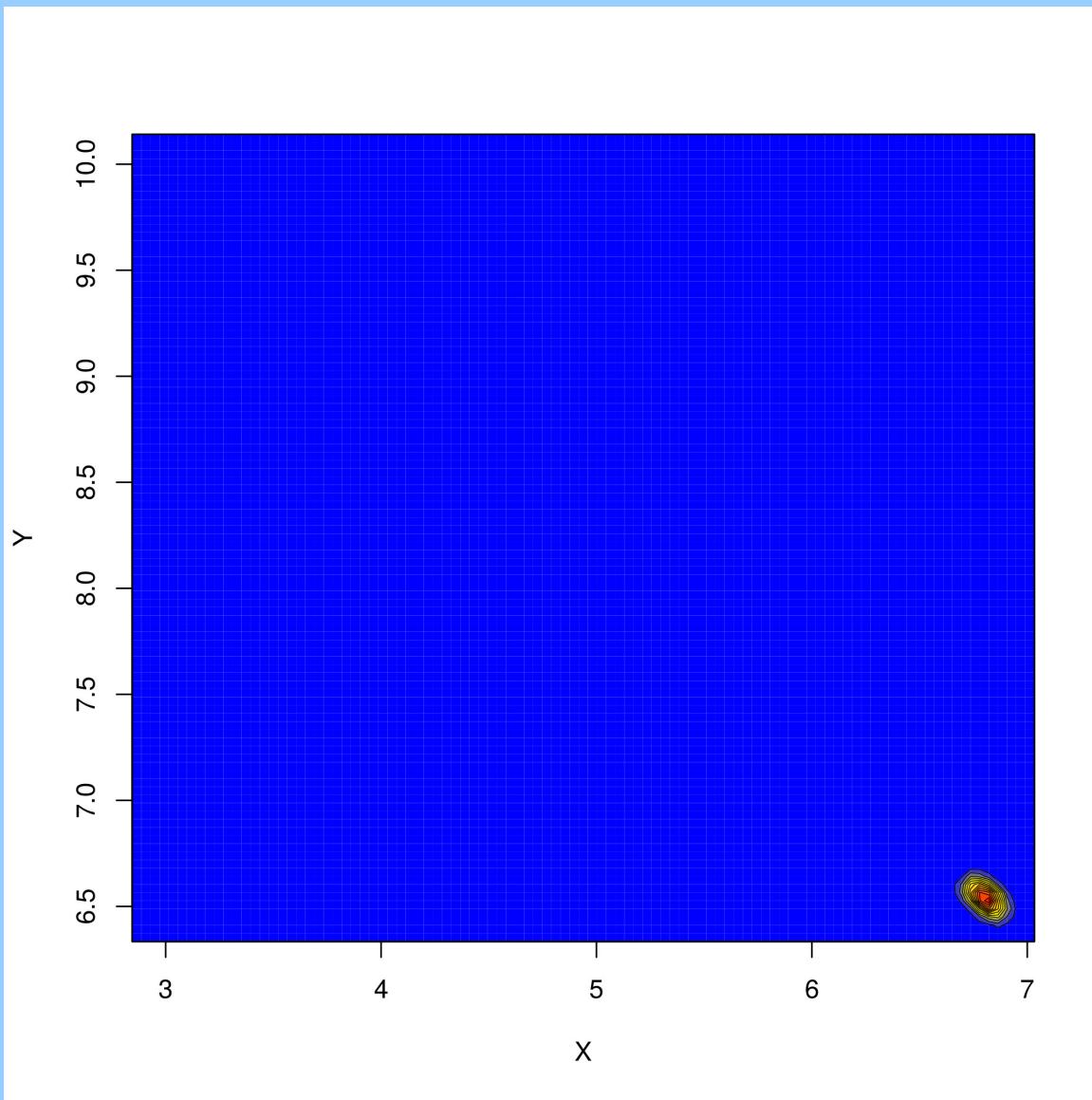
`TSPlot(timesimulation,
deltat = model.deltat,
dim = 2)`



`TSDensity(timesimulation,
dim = 1)`



`TSDensity(timesimulation,
dim = 2)`



Lotka-Volterra Competition: Coexistence

3) Calculate the local quasi-potentials

```
bounds.x = c(-5,15)    # upper and lower limit of x  
bounds.y = c(-5,15)    # upper and lower limit of y
```

```
step.number.x = 2000    # number of divisions in bounds.x  
step.number.y = 2000    # number of divisions in bounds.y
```

```
xinit = 6.667          # equilibrium x value  
yinit = 6.667          # equilibrium y value
```

```
# Compute the quasi-potential using the ordered upwind method  
coexist <- QPotential(x.rhs = dx, x.start = xinit, x.bound = bounds.x,  
                      x.num.steps = step.number.x, y.rhs = dy, y.start = yinit,  
                      y.bound = bounds.y, y.num.steps = step.number.y)
```

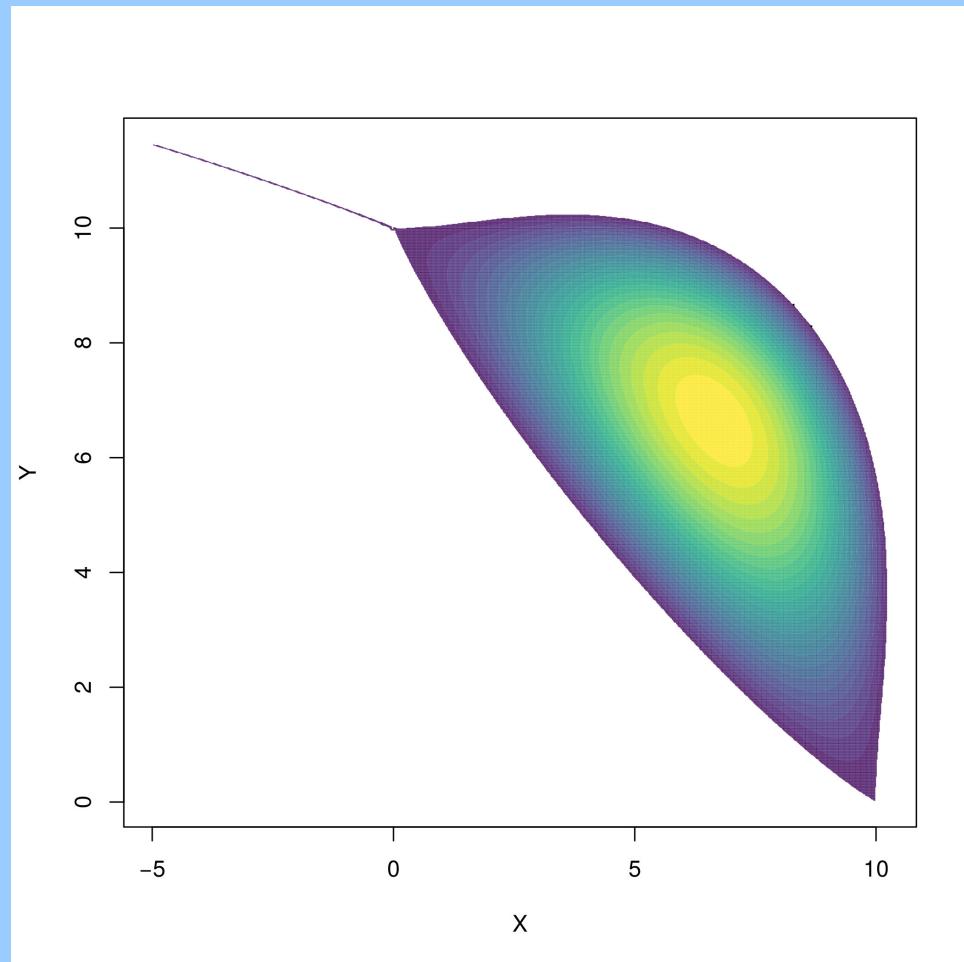
Lotka-Volterra Competition: Coexistence

3) Calculate the local quasi-potentials

```
QPContour(surface = coexist, dens = c(1000, 1000), x.bound = bounds.x,  
y.bound = bounds.y, c.parm = 0)
```

Lotka-Volterra Competition: Coexistence

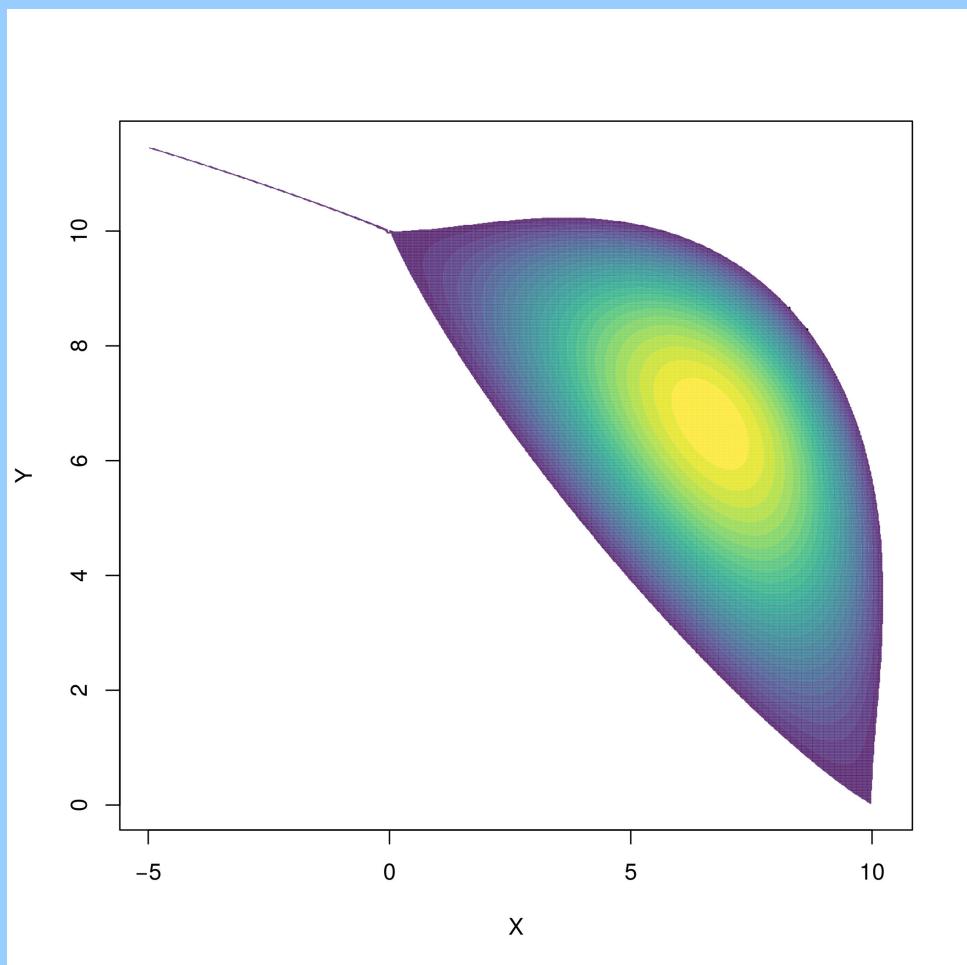
3) Calculate the local quasi-potentials



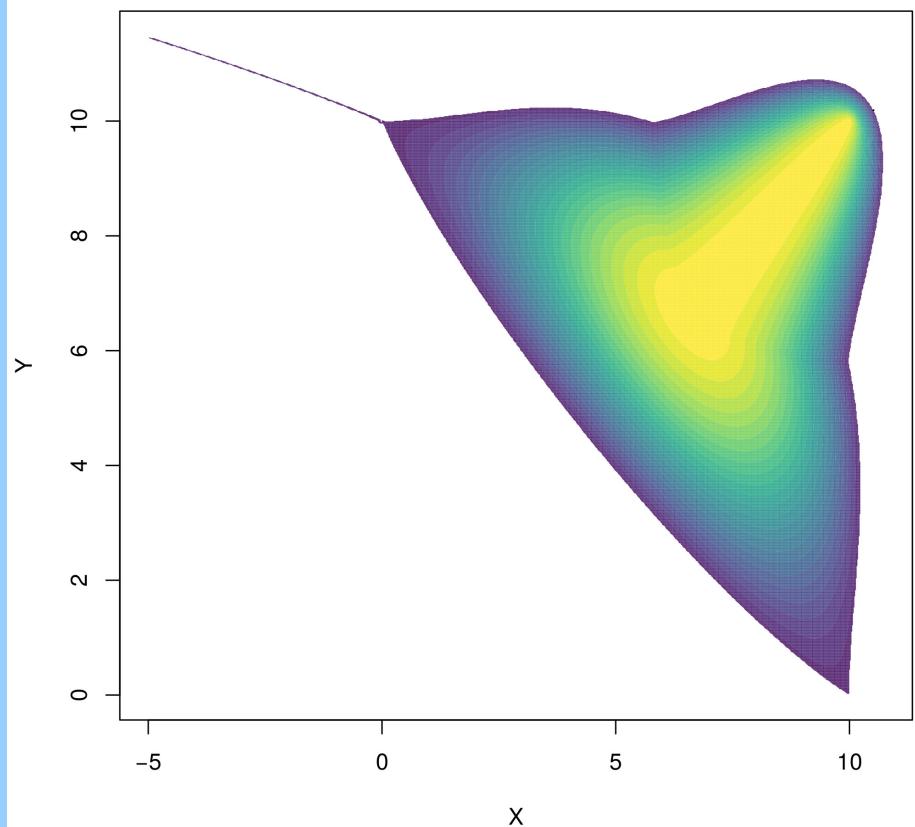
```
QPContour(surface = coexist, dens = c(1000, 1000), x.bound = bounds.x,  
y.bound = bounds.y, c.parm = 0)
```

Lotka-Volterra Competition: Coexistence

3) Calculate the local quasi-potentials



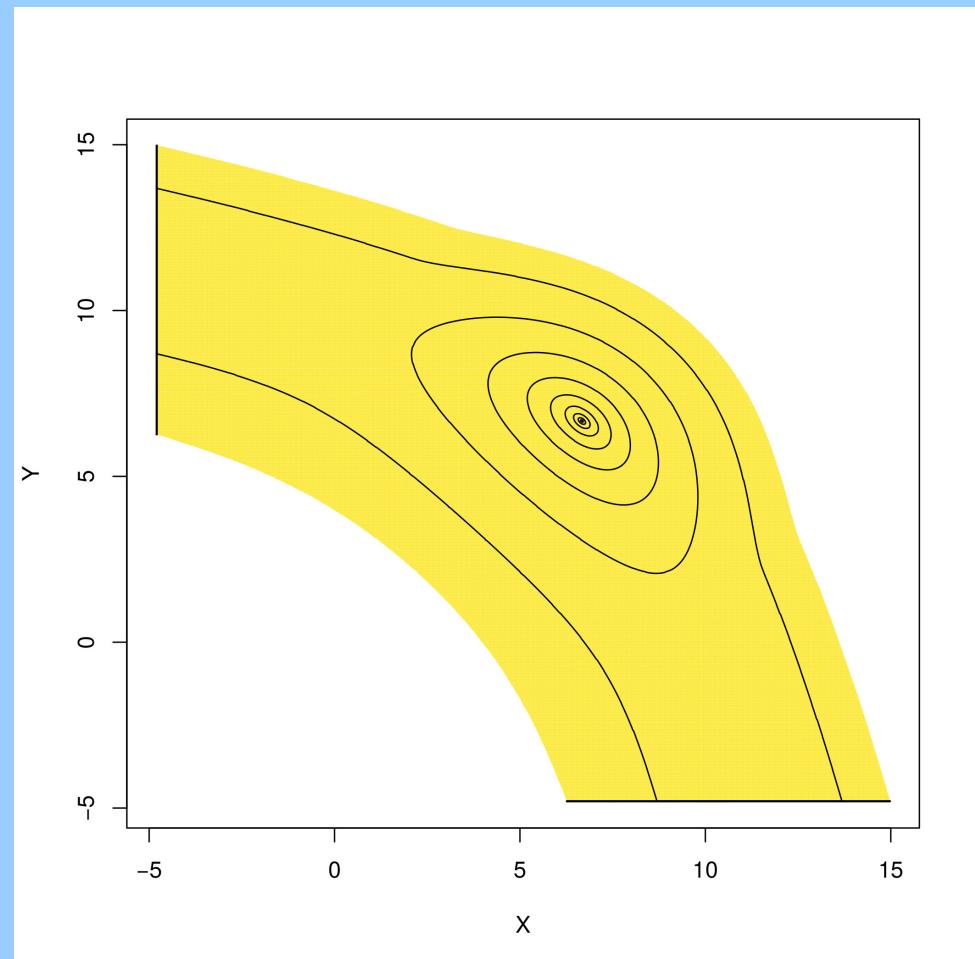
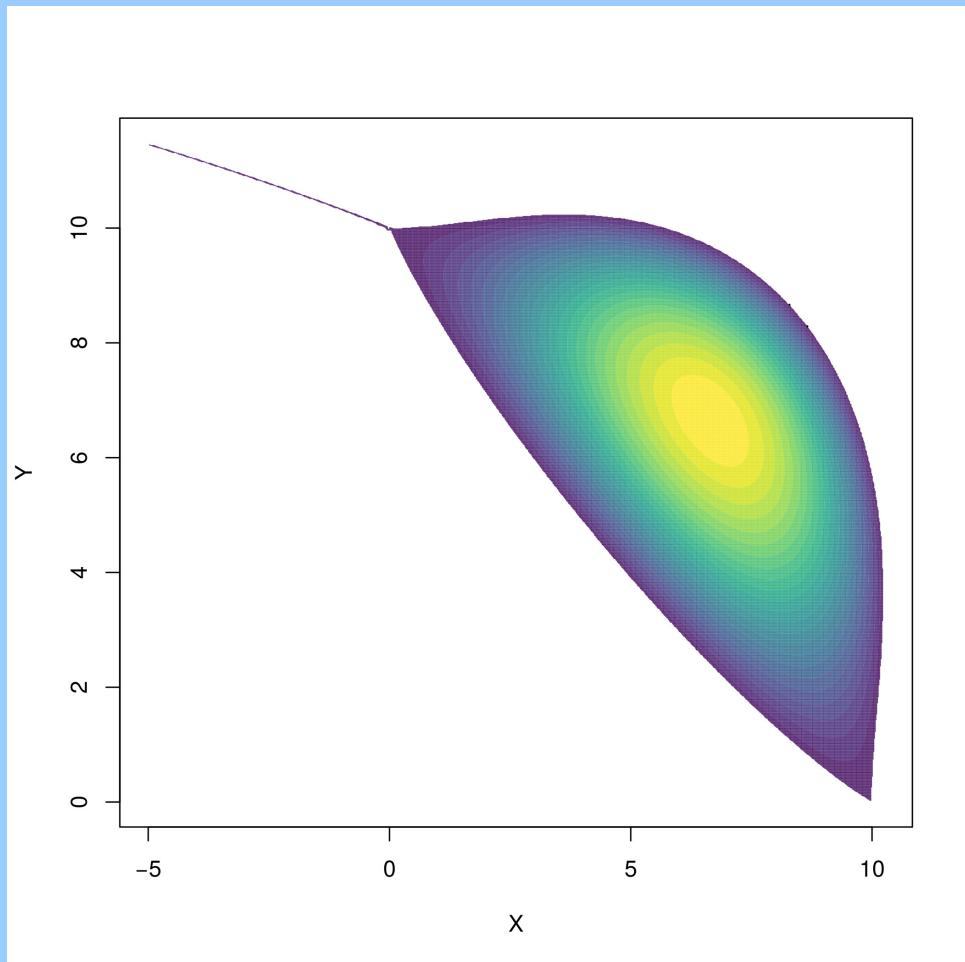
What happens
if you don't start at equilibrium



```
QPContour(surface = coexist, dens = c(1000, 1000), x.bound = bounds.x,  
y.bound = bounds.y, c.parm = 0)
```

Lotka-Volterra Competition: Coexistence

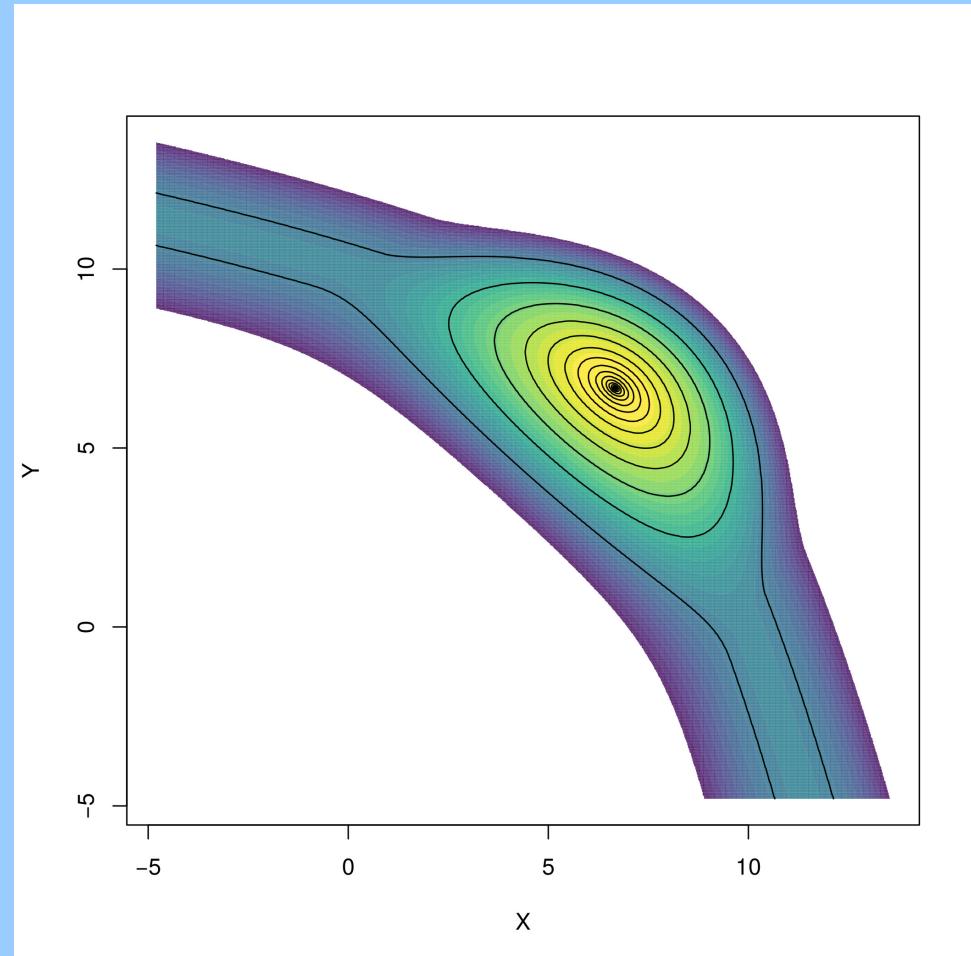
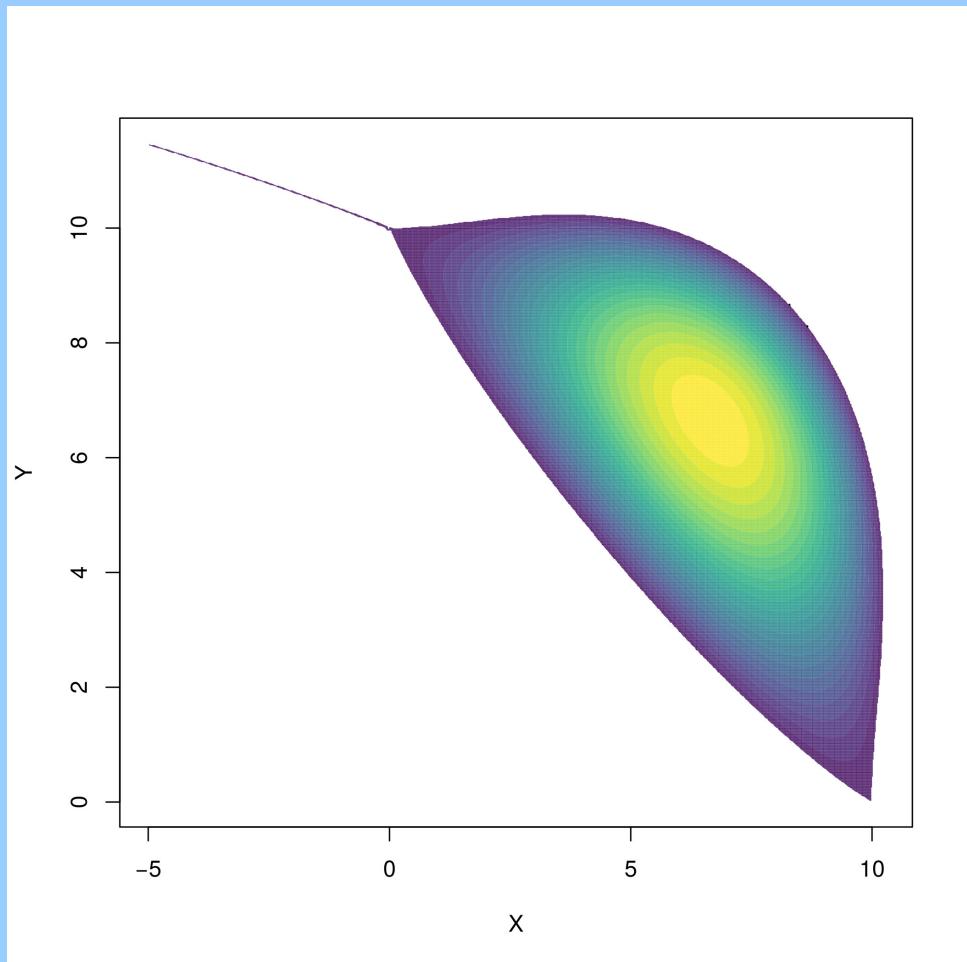
3) Calculate the local quasi-potentials



```
QPContour(surface = coexist.BOUNCE, dens = c(1000, 1000), x.bound = bounds.x,  
y.bound = bounds.y, c.parm = 10)
```

Lotka-Volterra Competition: Coexistence

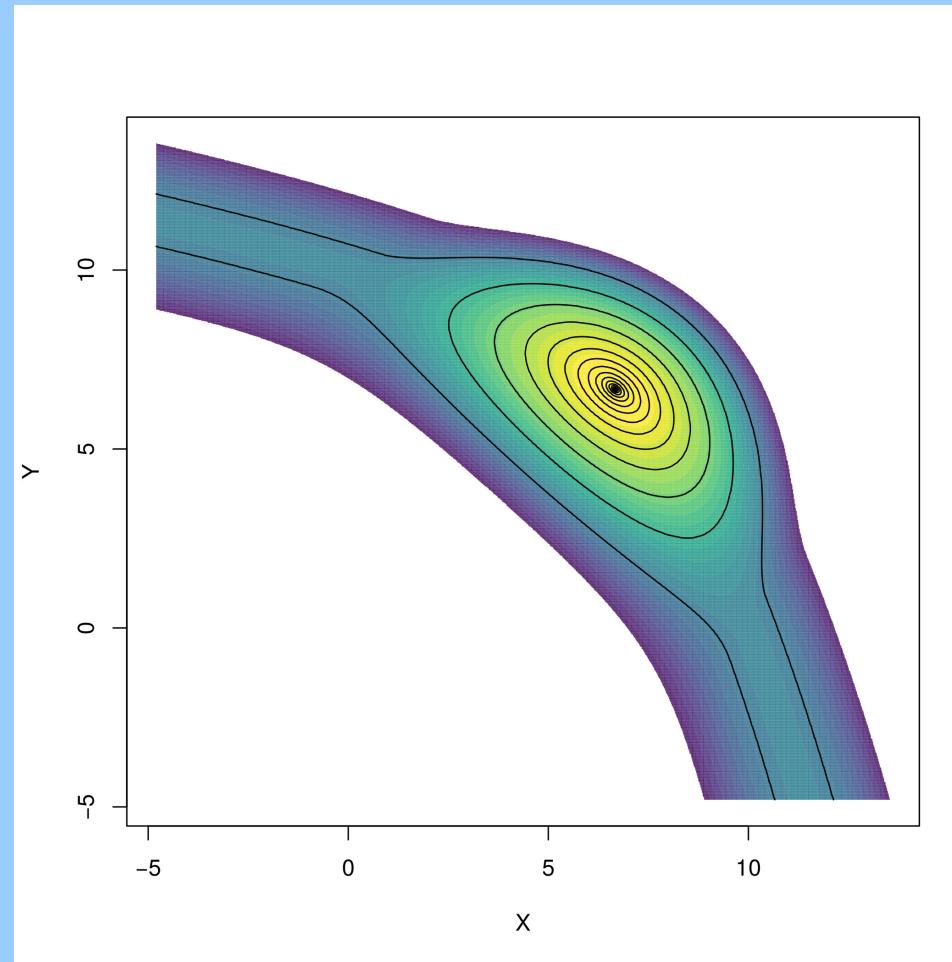
3) Calculate the local quasi-potentials



```
QPContour(surface = coexist.BOUNCE.REMOVED, dens = c(1000, 1000), x.bound = bounds.x,  
y.bound = bounds.y, c.parm = 10)
```

Lotka-Volterra Competition: Coexistence

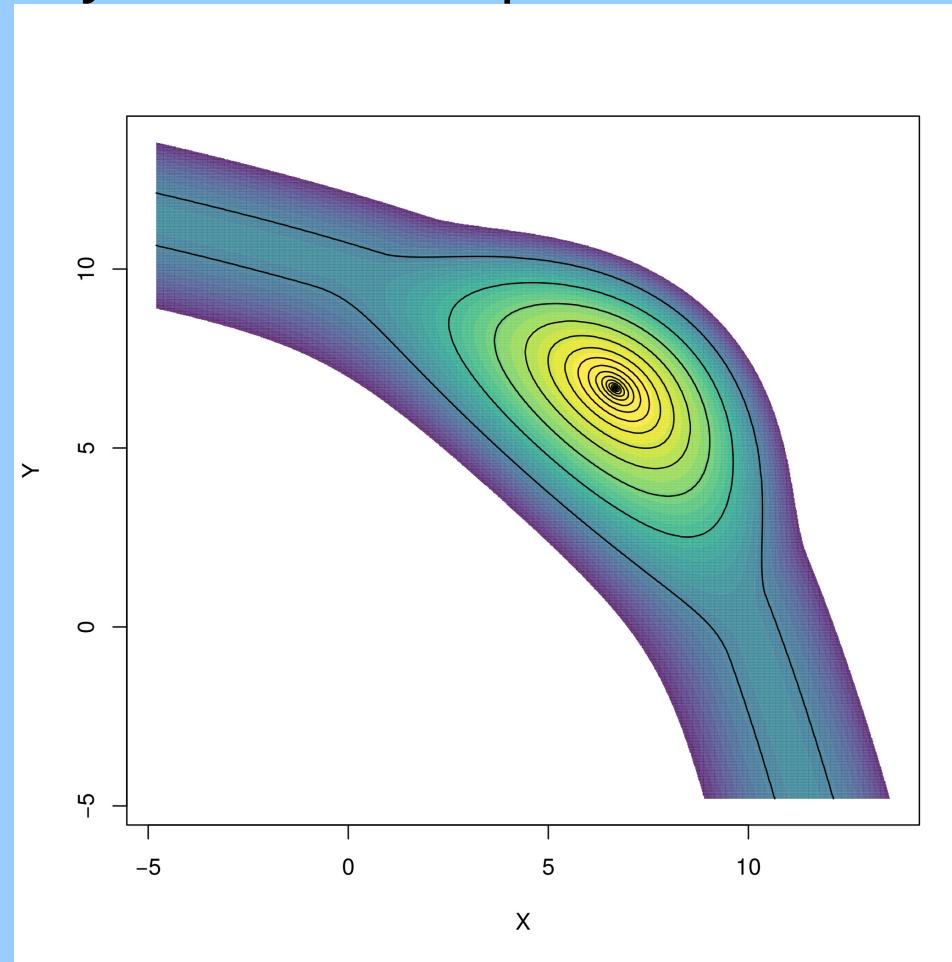
3) Calculate the local quasi-potentials



```
QPContour(surface = coexist.BOUNCE.REMOVED, dens = c(1000, 1000), x.bound = bounds.x,  
y.bound = bounds.y, c.parm = 10)
```

Lotka-Volterra Competition: Coexistence

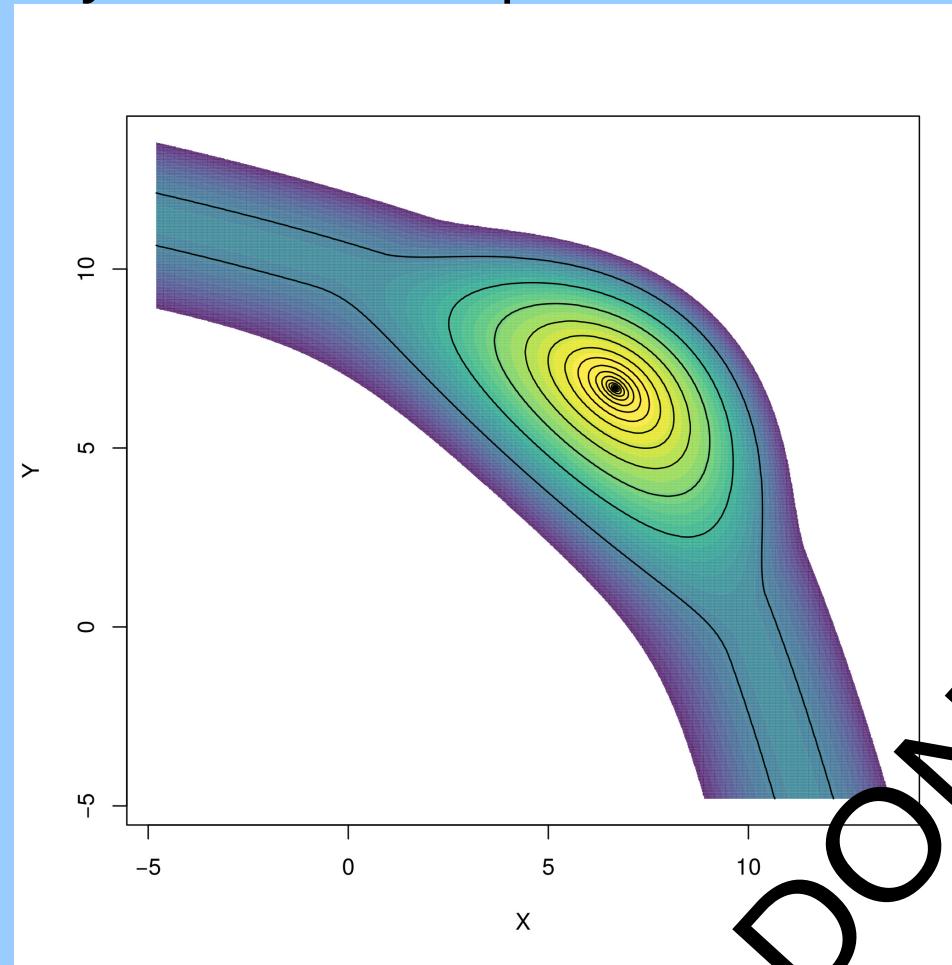
4) Calculate the global quasi-potential
(if only one stable equilibrium, then local = global)



```
QPContour(surface = coexist.BOUNCE.REMOVED, dens = c(1000, 1000), x.bound = bounds.x,  
y.bound = bounds.y, c.parm = 10)
```

Lotka-Volterra Competition: Coexistence

4) Calculate the global quasi-potential
(if only one stable equilibrium, then local = global)



DONE!

In later examples,
we will have to combine
local quasi-potentials
into a global
quasi-potential

```
QPCContour(surface = coexist.BOUNCE.REMOVED, dens = c(1000, 1000), x.bound = bounds.x,  
y.bound = bounds.y, c.parm = 10)
```

Lotka-Volterra Competition: Coexistence

5) Visualize the global quasi-potential

```
require(plot3D)
subsample.x = 200 # 3D plotting all the points takes forever
subsample.y = 200 # so we subsample and only plot 200*200 points

my.theta = 110          # rotation on 3D graph
my.phi = 0              # altitude of 3D graph
my.zlim = c(-0.001, 25) # range of z axis on plot, by making many
                        # graphs, we decided 25 looks good
                        # also notice that 25 removed the funk
my.zlab = intToUtf8(0x03A6) # this allows us to plot the letter phi
```

Lotka-Volterra Competition: Coexistence

5) Visualize the global quasi-potential

```
require(plot3D)
subsample.x = 200 # 3D plotting all the points takes forever
subsample.y = 200 # so we subsample and only plot 200*200 points

my.theta = 110          # rotation on 3D graph
my.phi = 0              # altitude of 3D graph
my.zlim = c(-0.001, 25) # range of z axis on plot, by making many
                        # graphs, we decided 25 looks good
                        # also notice that 25 removed the funk
my.zlab = intToUtf8(0x03A6) # this allows us to plot the letter phi

# again, need to subsample from the bigger matrix to make graphing faster
coexist.BOUNCE.SUBSAMPLE <- coexist.BOUNCE.REMOVED[
  seq(1,step.number.x,step.number.x/subsample.x),
  seq(1,step.number.y,step.number.y/subsample.y) ]
```

Lotka-Volterra Competition: Coexistence

5) Visualize the global quasi-potential

```
require(plot3D)
subsample.x = 200 # 3D plotting all the points takes forever
subsample.y = 200 # so we subsample and only plot 200*200 points

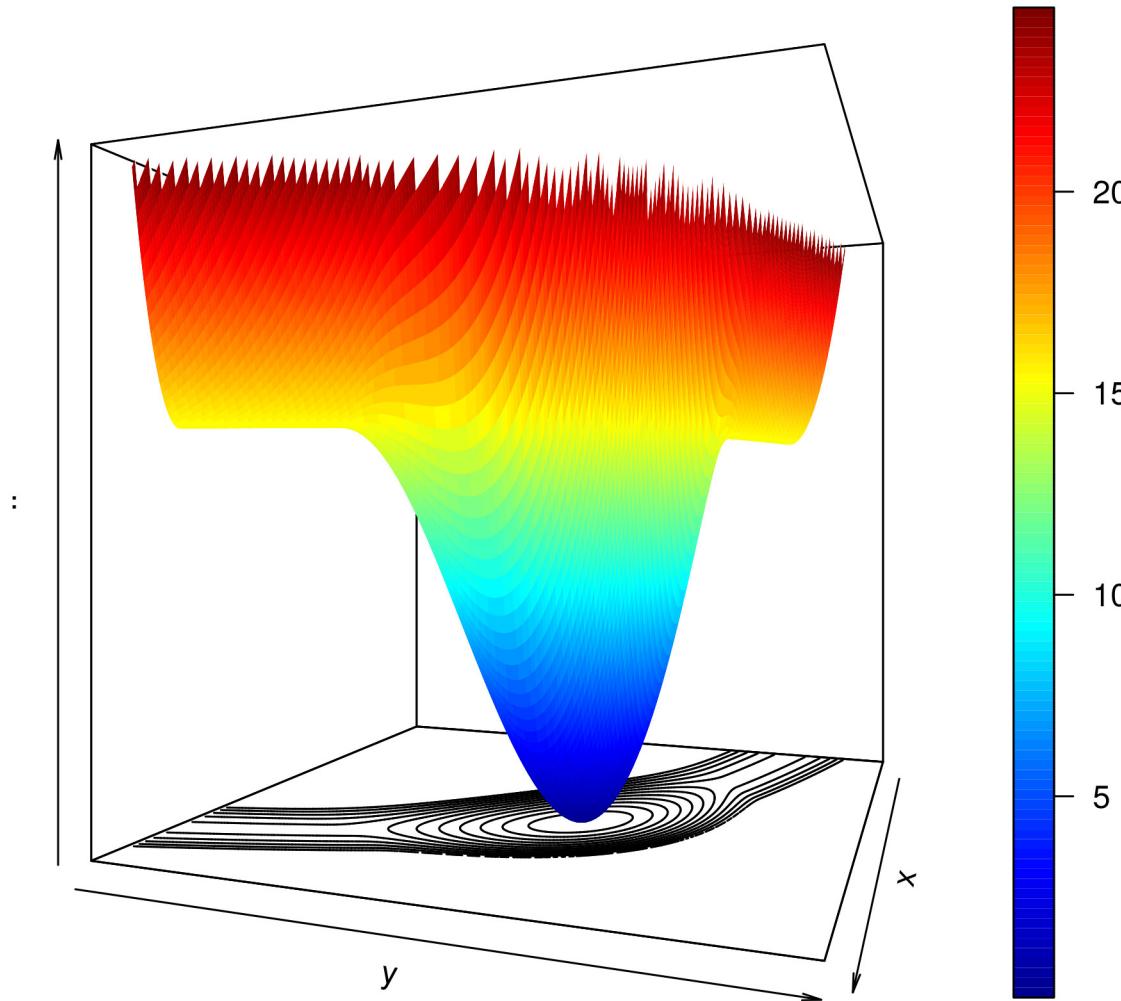
my.theta = 110          # rotation on 3D graph
my.phi = 0              # altitude of 3D graph
my.zlim = c(-0.001, 25) # range of z axis on plot, by making many
                        # graphs, we decided 25 looks good
                        # also notice that 25 removed the funk
my.zlab = intToUtf8(0x03A6) # this allows us to plot the letter phi

# again, need to subsample from the bigger matrix to make graphing faster
coexist.BOUNCE.SUBSAMPLE <- coexist.BOUNCE.REMOVED[
  seq(1,step.number.x,step.number.x/subsample.x),
  seq(1,step.number.y,step.number.y/subsample.y) ]

# plot the quasipotential when we used bounce = 'b'
persp3D(z = coexist.BOUNCE.SUBSAMPLE, theta = my.theta, phi = my.phi,
        contour = T, zlim = my.zlim, zlab = my.zlab)
```

Lotka-Volterra Competition: Coexistence

5) Visualize the global quasi-potential



```
persp3D(z = coexist.BOUNCE.SUBSAMPLE, theta = my.theta, phi = my.phi,  
contour = T, zlim = my.zlim, zlab = my.zlab)
```

Lotka-Volterra Competition: Coexistence

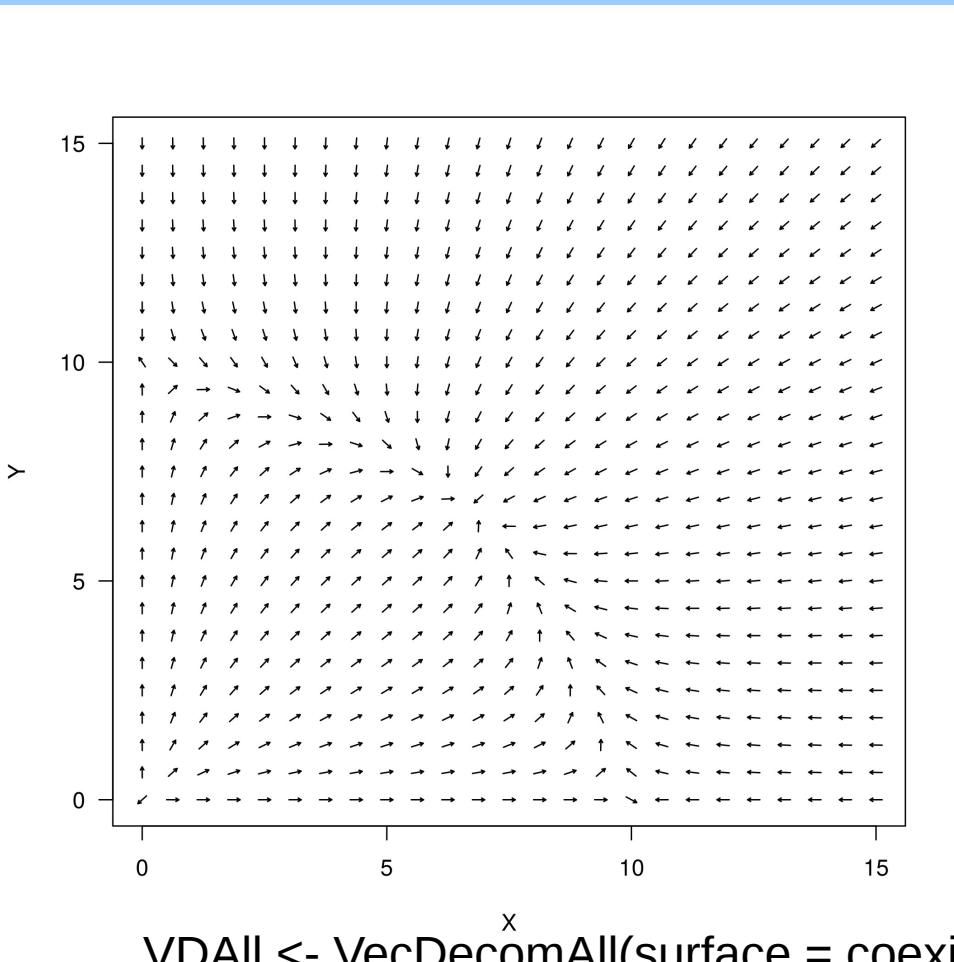
- 6) Visualize the vector field decompositions
 - a) deterministic skeleton

```
VDAll <- VecDecomAll(surface = coexist.BOUNCE, x.rhs = dx,
                      y.rhs = dy, x.bound = bounds.x, y.bound = bounds.y)
x.skel = VDAll[,1]; y.skel = VDAll[,2]
VecDecomPlot(x.field = x.skel, y.field = y.skel, dens = c(25, 25), x.bound = bounds.x,
              y.bound = bounds.y, arrow.type = "equal", xlim = c(0, 15), ylim = c(0, 15),
              tail.length = 0.25, head.length = 0.025)
```

Lotka-Volterra Competition: Coexistence

6) Visualize the vector field decompositions

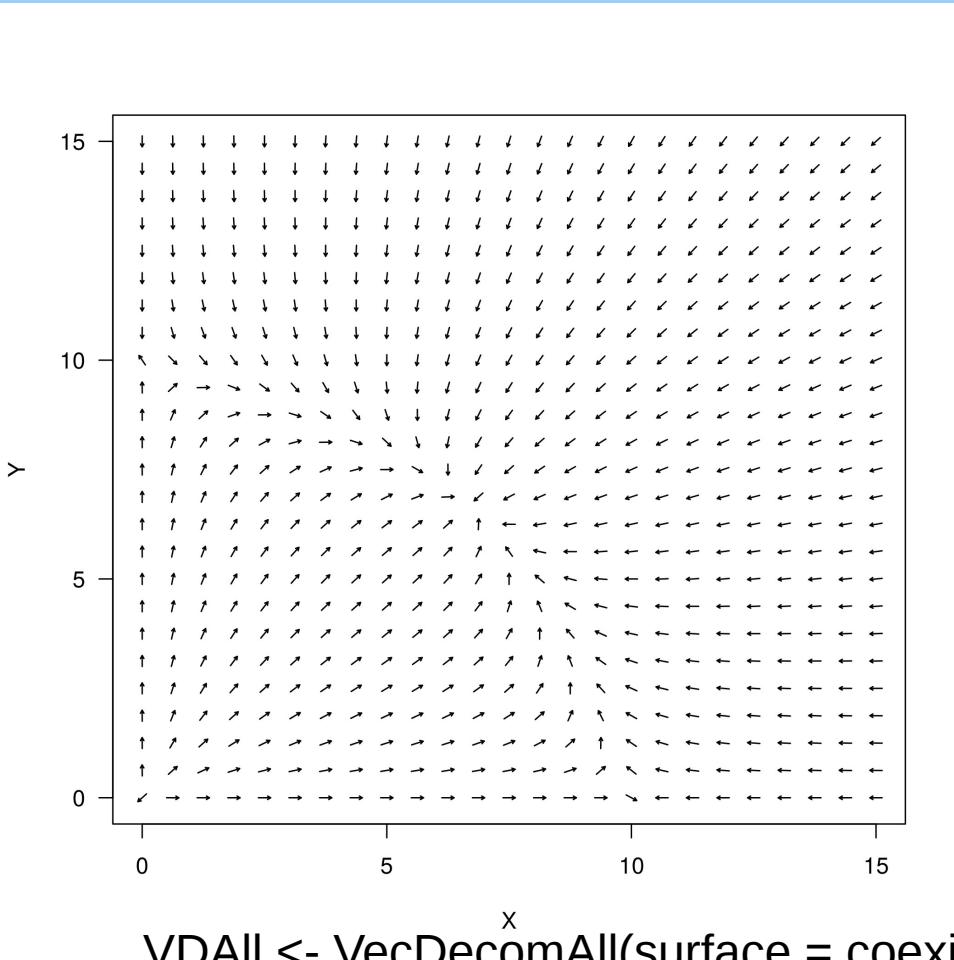
a) deterministic skeleton



```
VDAll <- VecDecomAll(surface = coexist.BOUNCE, x.rhs = dx,
                      y.rhs = dy, x.bound = bounds.x, y.bound = bounds.y)
x.skel = VDAll[,1]; y.skel = VDAll[,2]
VecDecomPlot(x.field = x.skel, y.field = y.skel, dens = c(25, 25), x.bound = bounds.x,
             y.bound = bounds.y, arrow.type = "equal", xlim = c(0, 15), ylim = c(0, 15),
             tail.length = 0.25, head.length = 0.025)
```

Lotka-Volterra Competition: Coexistence

- 6) Visualize the vector field decompositions
 - a) deterministic skeleton



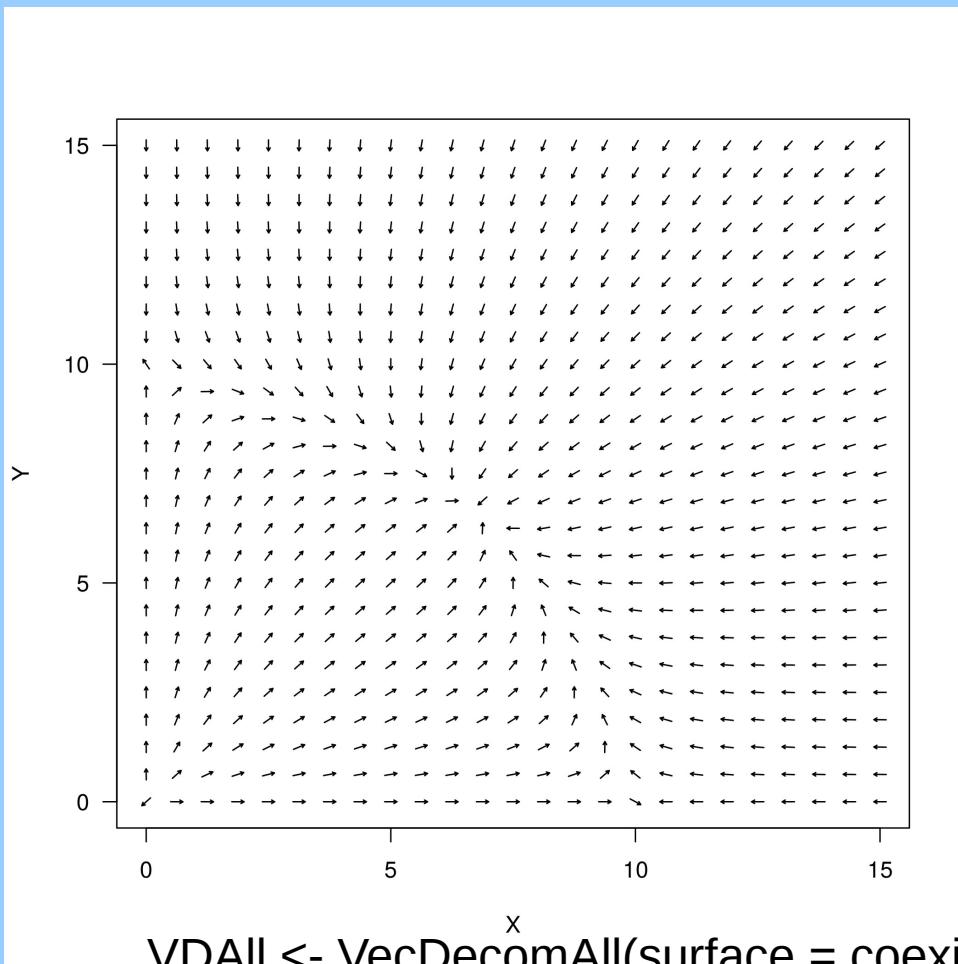
Should look familiar

```
VDAll <- VecDecomAll(surface = coexist.BOUNCE, x.rhs = dx,  
                      y.rhs = dy, x.bound = bounds.x, y.bound = bounds.y)  
x.skel = VDAll[,1]; y.skel = VDAll[,2]  
VecDecomPlot(x.field = x.skel, y.field = y.skel, dens = c(25, 25), x.bound = bounds.x,  
             y.bound = bounds.y, arrow.type = "equal", xlim = c(0, 15), ylim = c(0, 15),  
             tail.length = 0.25, head.length = 0.025)
```

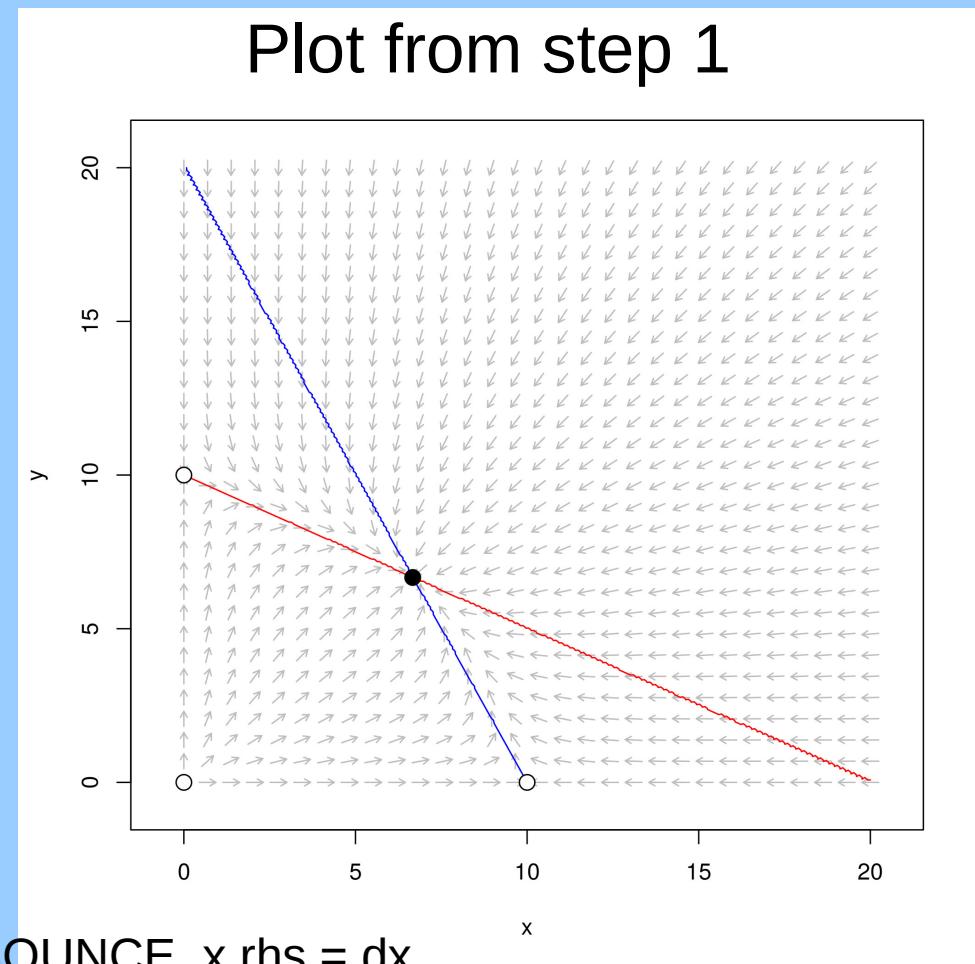
Lotka-Volterra Competition: Coexistence

6) Visualize the vector field decompositions

a) deterministic skeleton



```
X  
VDAll <- VecDecomAll(surface = coexist.BOUNCE, x.rhs = dx,  
y.rhs = dy, x.bound = bounds.x, y.bound = bounds.y)  
x.skel = VDAll[,1]; y.skel = VDAll[,2]  
VecDecomPlot(x.field = x.skel, y.field = y.skel, dens = c(25, 25), x.bound = bounds.x,  
y.bound = bounds.y, arrow.type = "equal", xlim = c(0, 15), ylim = c(0, 15),  
tail.length = 0.25, head.length = 0.025)
```

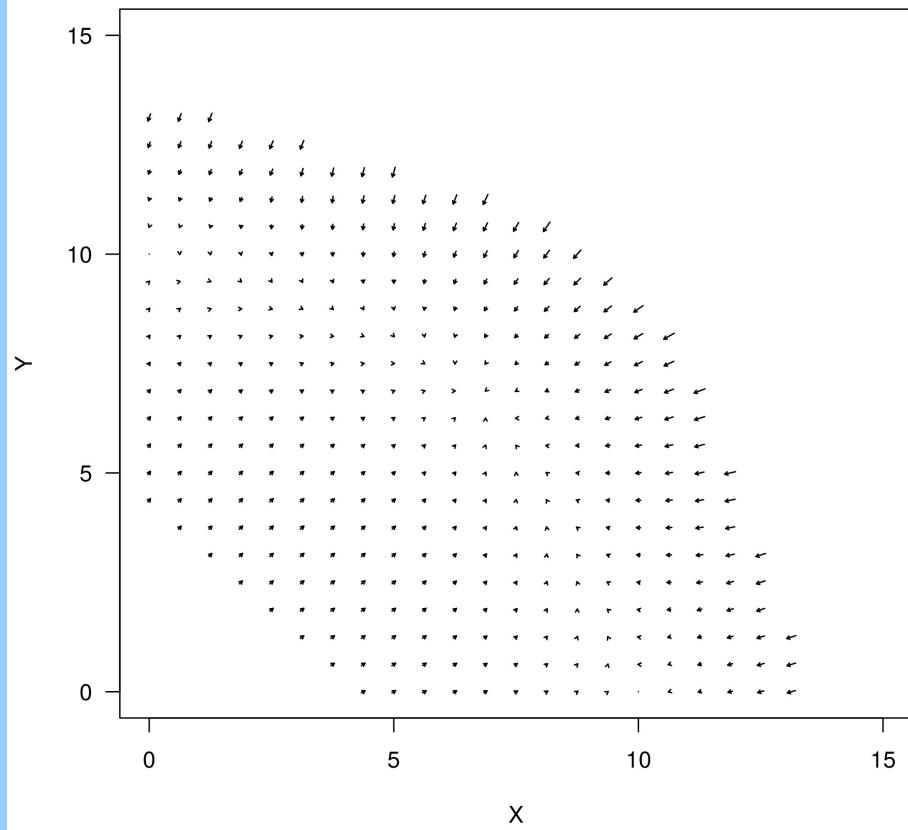


Plot from step 1

Lotka-Volterra Competition: Coexistence

- 6) Visualize the vector field decompositions
 - b) visualize the gradient

arrow.type = 'proportional'



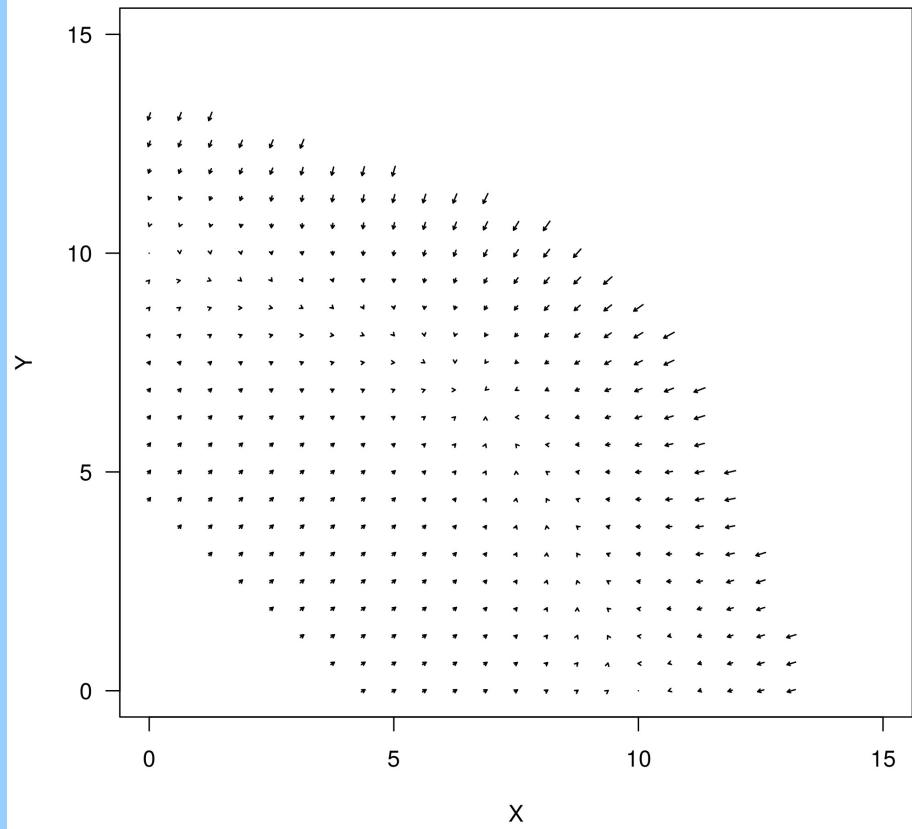
```
VecDecomPlot(x.field = VDAII[,3], y.field = VDAII[,4], dens = c(25, 25),  
x.bound = bounds.x, y.bound = bounds.y, arrow.type = "XXXXXXXXXX",  
tail.length = 0.25, head.length = 0.025, xlim = c(0,15), ylim = c(0,15))
```

Lotka-Volterra Competition: Coexistence

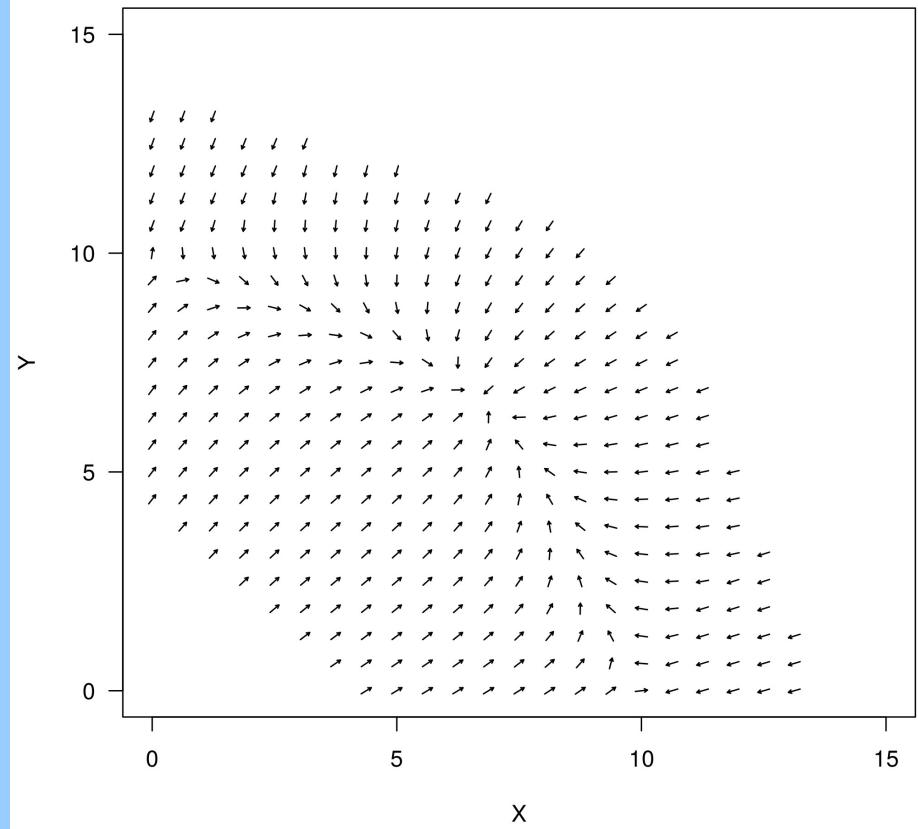
6) Visualize the vector field decompositions

b) visualize the gradient

arrow.type = 'proportional'



arrow.type = 'equal'

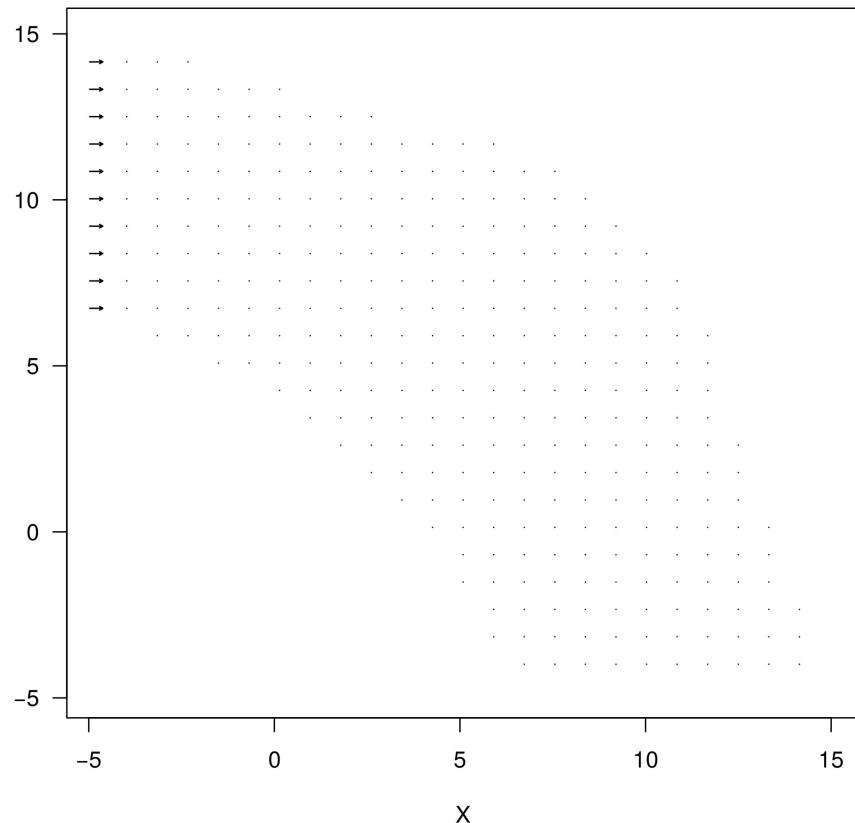


```
VecDecomPlot(x.field = VDAII[,3], y.field = VDAII[,4], dens = c(25, 25),  
x.bound = bounds.x, y.bound = bounds.y, arrow.type = "XXXXXXXXXX",  
tail.length = 0.25, head.length = 0.025, xlim = c(0,15), ylim = c(0,15))
```

Lotka-Volterra Competition: Coexistence

- 6) Visualize the vector field decompositions
 - b) visualize the remainder

```
arrow.type = 'proportional'
```



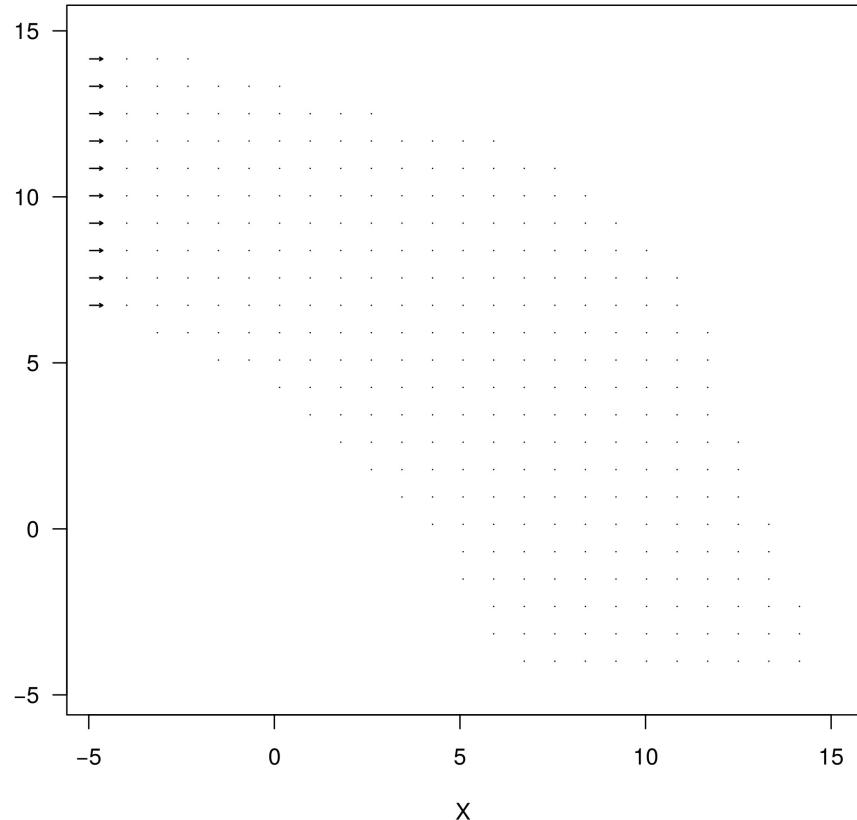
```
VecDecomPlot(x.field = VDAII[,5], y.field = VDAII[,6], dens = c(25, 25), x.bound = bounds.x,  
y.bound = bounds.y, arrow.type = "XXXXXXXXX",  
tail.length = 0.35, head.length = 0.025)
```

Lotka-Volterra Competition: Coexistence

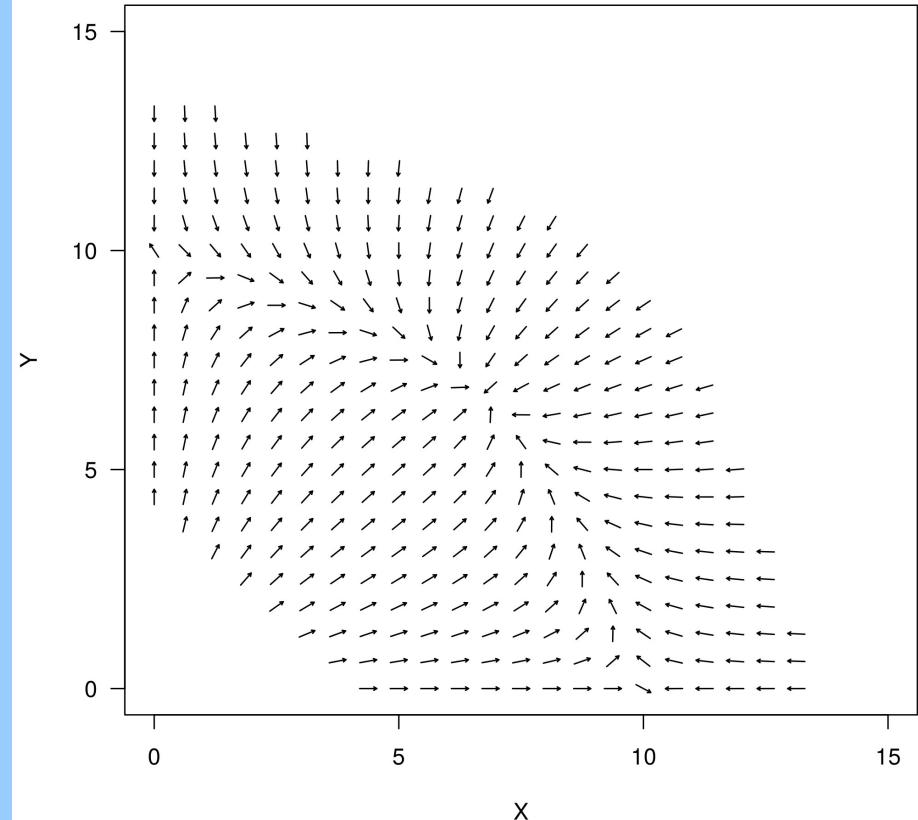
6) Visualize the vector field decompositions

b) visualize the remainder

arrow.type = 'proportional'



arrow.type = 'equal'

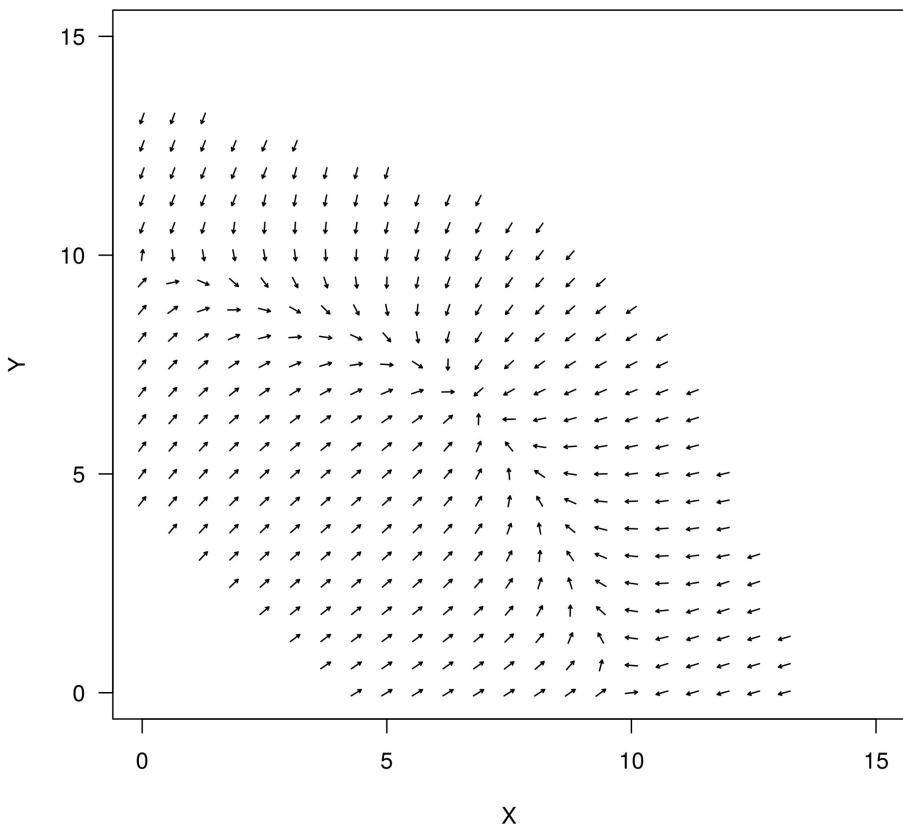


```
VecDecomPlot(x.field = VDAII[,5], y.field = VDAII[,6], dens = c(25, 25), x.bound = bounds.x,  
y.bound = bounds.y, arrow.type = "XXXXXXXXX",  
tail.length = 0.35, head.length = 0.025)
```

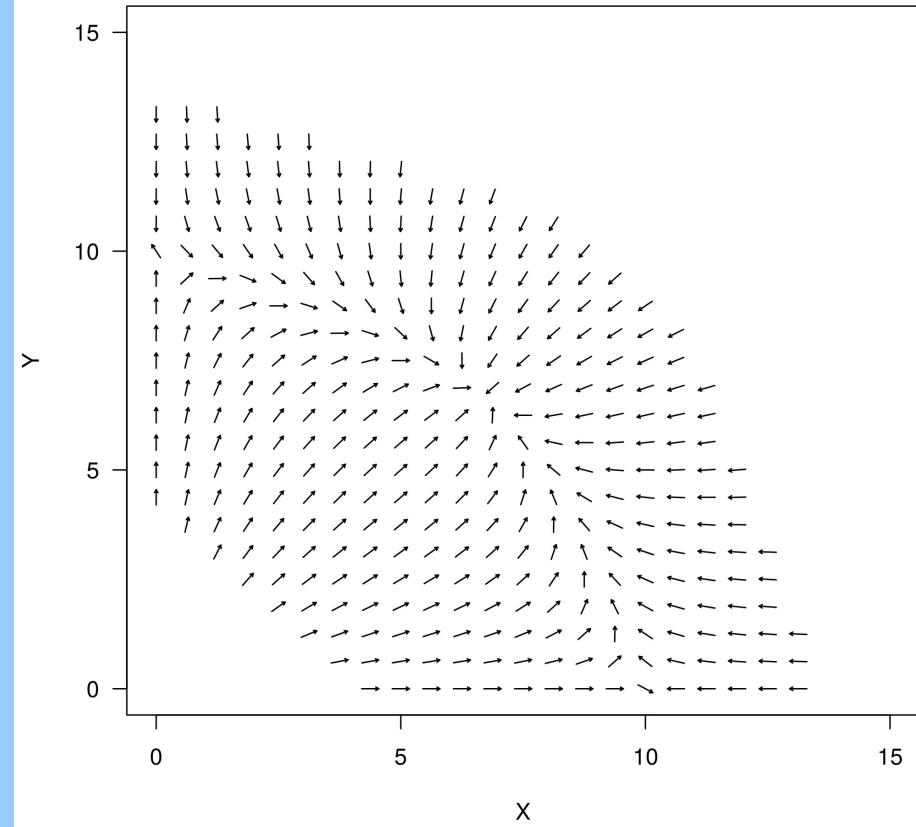
Lotka-Volterra Competition: Coexistence

6) Visualize the vector field decompositions

Gradient



Remainder

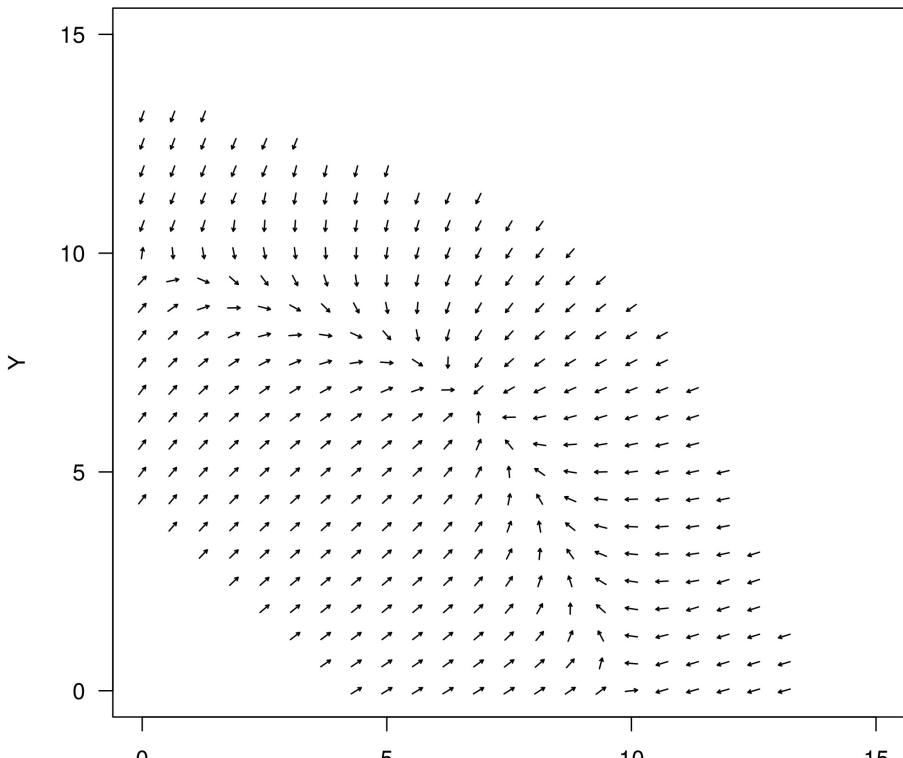


arrow.type = "equal"

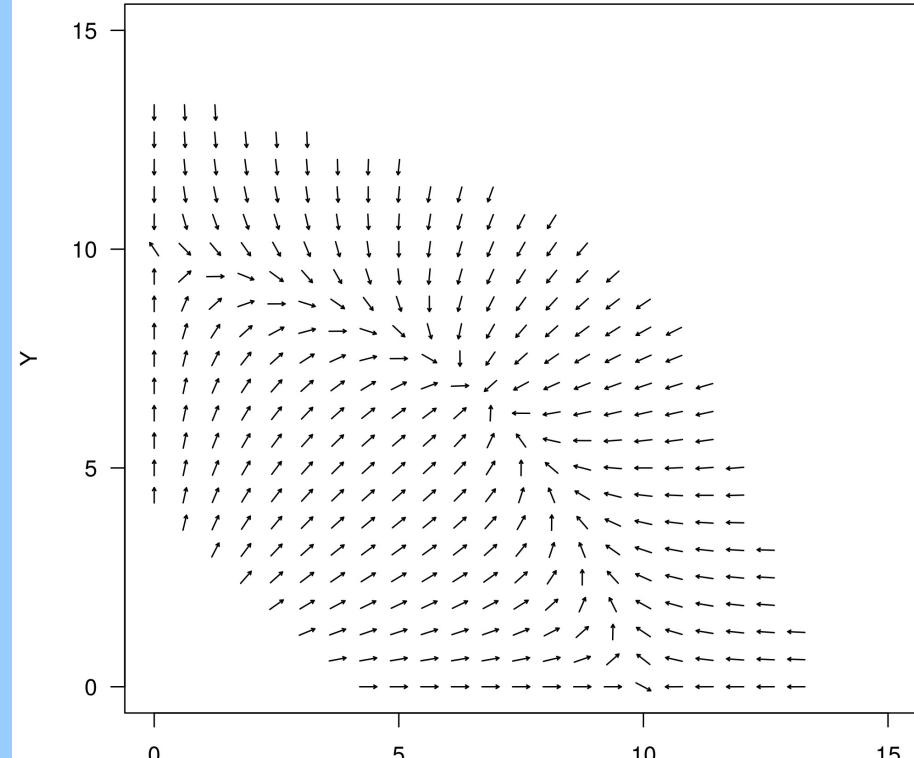
Lotka-Volterra Competition: Coexistence

6) Visualize the vector field decompositions

Gradient



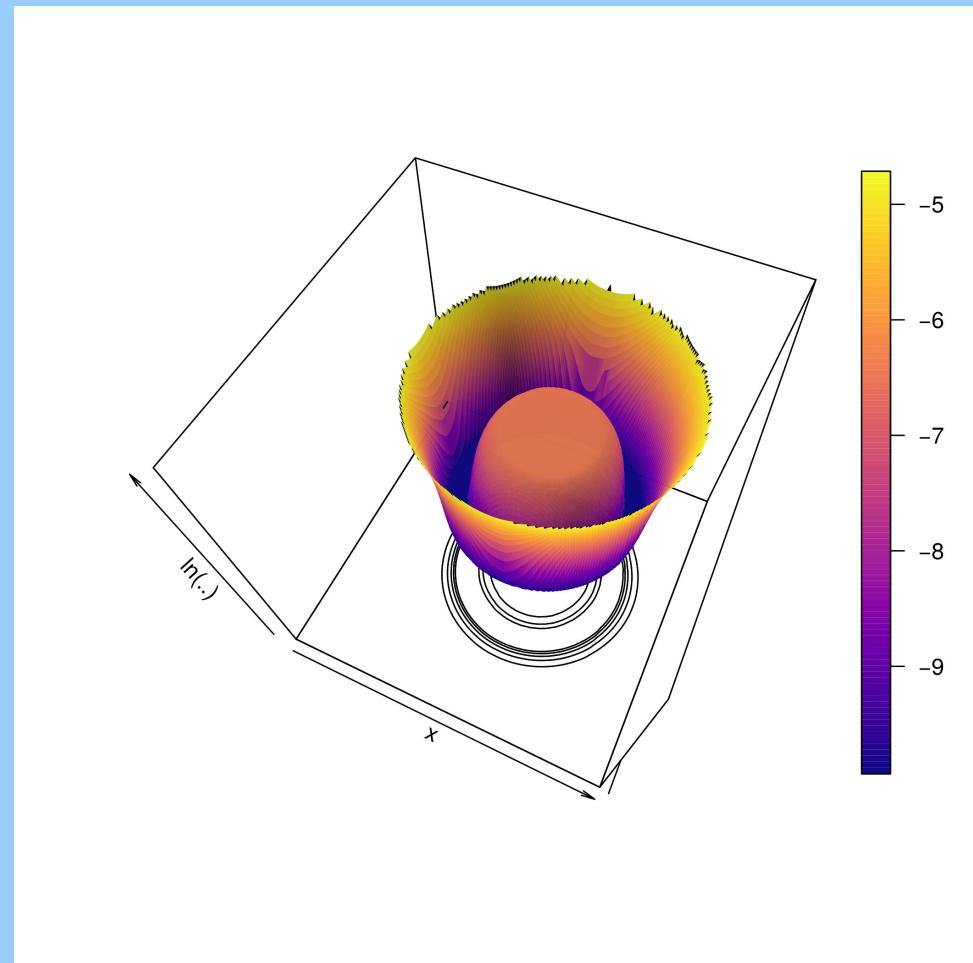
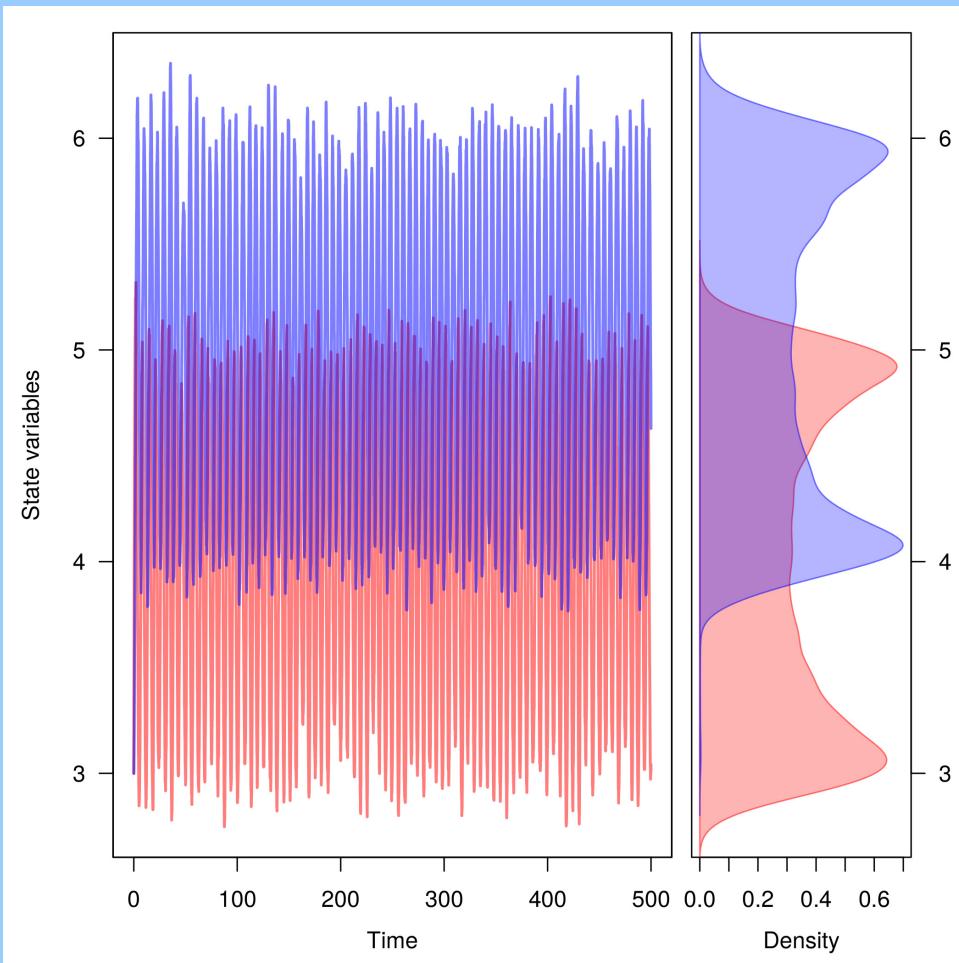
Remainder



In this case, the two look similar,
but could look different.

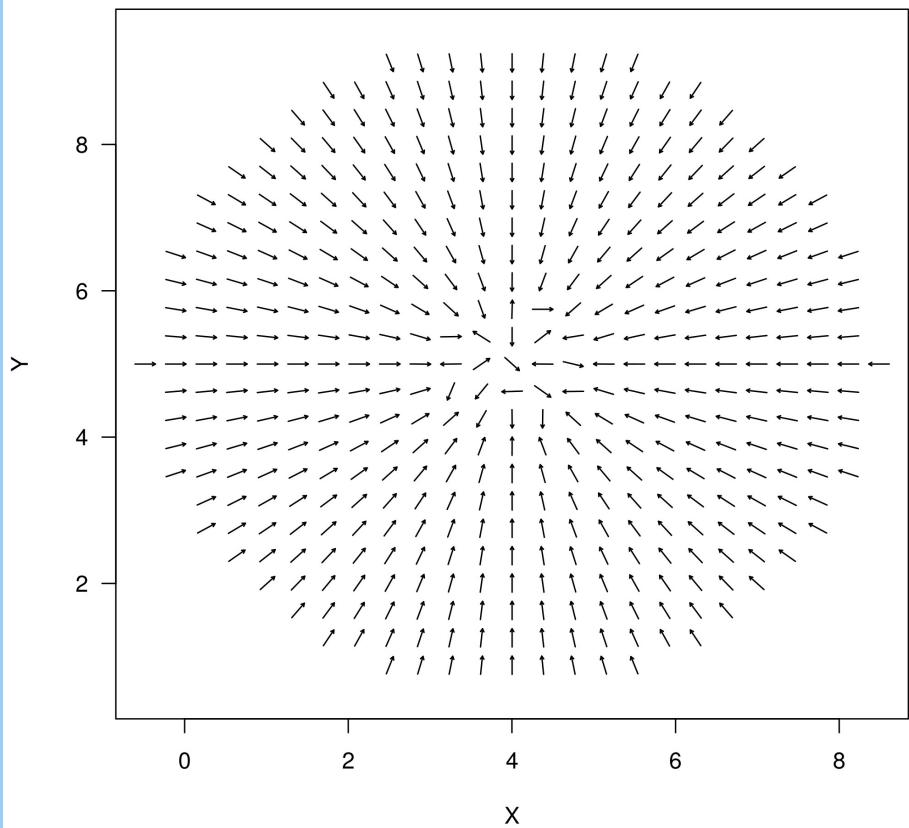
arrow.type = "equal"

Limit-cycle

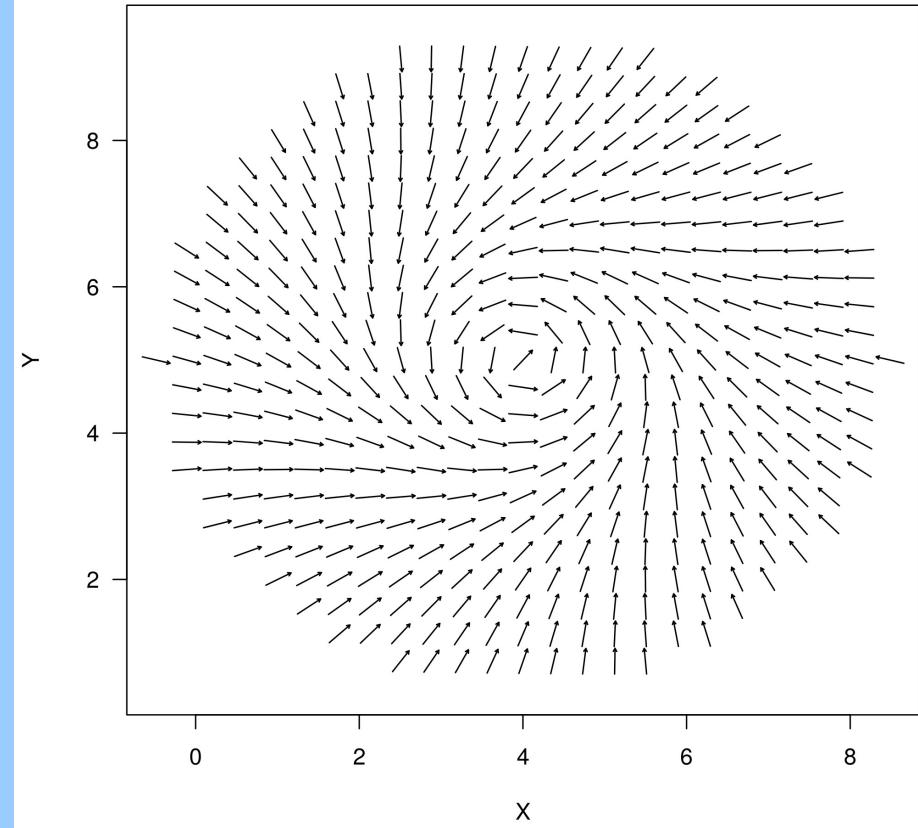


Limit-cycle

Gradient



Remainder



remainder vector field, which is the “force that causes trajectories to circulate around level sets of the quasi-potential.”

Other Examples

At stieha.com

stieha@hotmail.com
stieha@case.edu

1) Competition with initial condition dependence

This shows an example of having multiple stable equilibria,

Where you need to use QPGlobal() to merge

Depending on the parameter values,

may require changing boundary conditions to get good results

2) Consumer-resource model with two stable equilibria

Appeared in talk, also appears in Moore et al. 2016 as Example 1

Have multiple stable equilibria, so need to combine local quasi-potentials

into a global quasi-potential

3) Consumer-resource model with limit cycle (no code yet)

Appeared in talk,

Same model as #2, but different parameter values

4) System of equations with a limit cycle

Example 2 in Moore et al. 2016

5) Combining local quasi-potentials into a global quasi-potential

In a difficult system

Example 3 in Moore et al. 2016.

Download article for code