

# BiomathHW07

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## Problem #1

The following is a generic code to classify the behavior of a linear system based on it's trace and eigenvalues. I will use it in each part of the problem

```
classify <- function(mat)
{
  trace <- sum(diag(mat)); deter <- det(mat)
  delta <- deter - trace^2/4
  if(deter < 0){return("saddle")}
  if(deter ==0)
  {
    if(trace > 0){return("unstable line")}
    if(trace < 0){return("stable line")}
    if(trace ==0){return("uniform motion")}
  }
  if(trace > 0)
  {
    if(delta < 0){return("source")}
    if(delta ==0){return("degenerate source")}
    if(delta > 0){return("spiral source")}
  }
  if(trace ==0){return("center")}
  if(trace < 0)
  {
    if(delta < 0){return("sink")}
    if(delta ==0){return("degenerate sink")}
    if(delta > 0){return("spiral sink")}
  }
}
}

plotSystem <- function(mat)
{
  simple <- function(t,state,parameters){
    with(as.list(c(state,parameters)),{
      dx <- a*state[1] + b*state[2]
      dy <- c*state[1] + d*state[2]
      list(c(dx,dy))
    })}
  ff <- flowField(simple,
                  xlim = c(-2, 2), ylim = c(-2, 2),
                  parameters = c(a=mat[1],b=mat[3],c=mat[2],d=mat[4]),
                  points = 19,add = FALSE)
  state <- matrix(c(1,1,1,-1,-1,1,-1,-1,0,2,0,-2,-1,0,1,0),
                  8, 2, byrow = TRUE)
  trajs <- trajectory(simple,y0 = state, tlim = c(0, 10),
                     parameters = c(a=mat[1],b=mat[3],c=mat[2],d=mat[4]),add=TRUE)
}
```

(a)

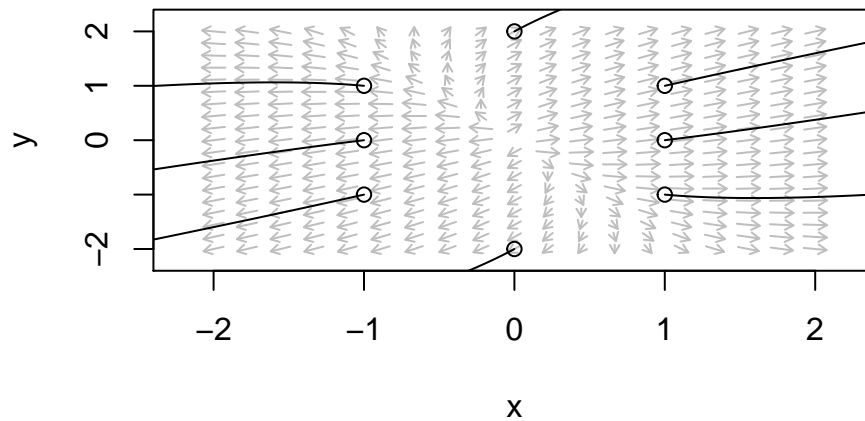
$$\dot{x} = 6x + 2y$$

$$\dot{y} = 2x + 3y$$

```
## [1] "Phase portrait type:  source"
```

Therefore we expect to have two distinct real eigenvalues:

```
## [1] "Eigenvalues:  7" "Eigenvalues:  2"
```



(b)

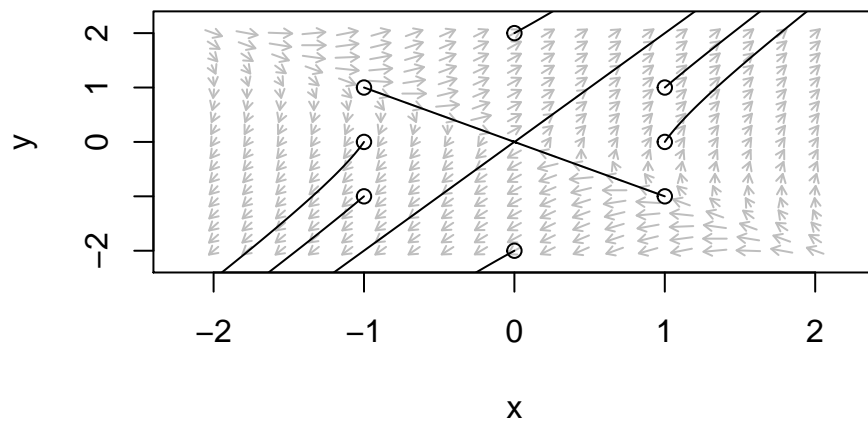
$$\dot{x} = x + 2y$$

$$\dot{y} = 4x + 3y$$

```
## [1] "Phase portrait type:  saddle"
```

Therefore we expect two distinct real eigenvalues:

```
## [1] "Eigenvalues:  5" "Eigenvalues: -1"
```



(c)

$$\dot{x} = -2x + 4y$$

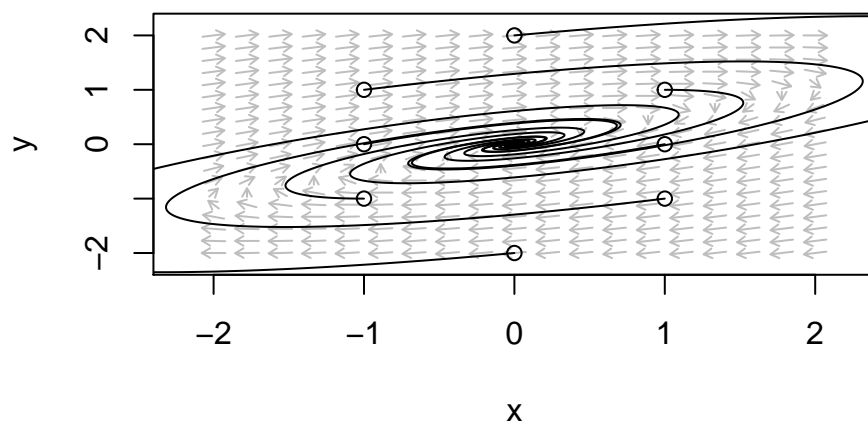
$$\dot{y} = -x + y$$

## [1] "Phase portrait type: spiral sink"

Therefore we expect a complex conjugate pair of eigenvalues:

## [1] "Eigenvalues: -0.5+1.3228756555323i"

## [2] "Eigenvalues: -0.5-1.3228756555323i"



(d)

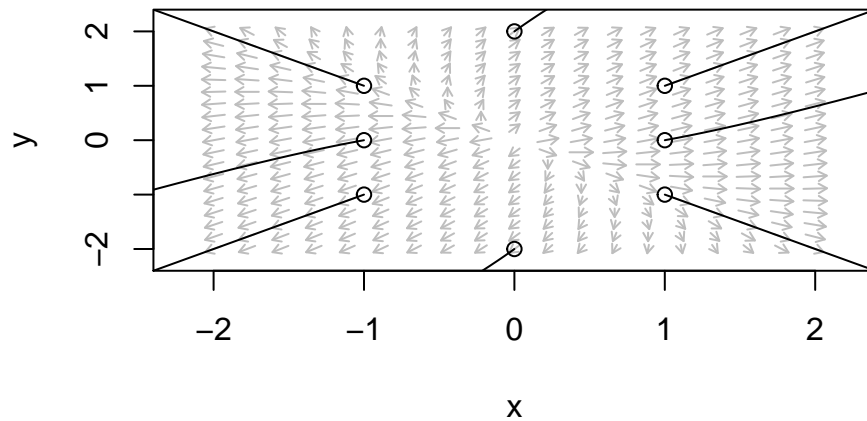
$$\dot{x} = 2x + y$$

$$\dot{y} = x + 2y$$

```
## [1] "Phase portrait type:  source"
```

Therefore we expect two distinct real eigenvalues:

```
## [1] "Eigenvalues:  3" "Eigenvalues:  1"
```



(e)

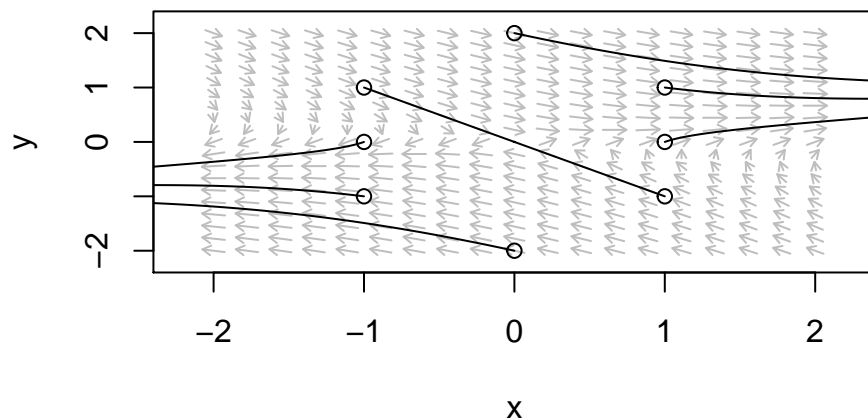
$$\dot{x} = x + 5y$$

$$\dot{y} = x - 3y$$

```
## [1] "Phase portrait type:  saddle"
```

Therefore we expect two distinct real eigenvalues:

```
## [1] "Eigenvalues:  -4" "Eigenvalues:  2"
```



(f)

$$\dot{x} = -1x + ay$$

$$\dot{y} = 0x + ay$$

for  $a \neq 0$

This function changes behavior, dependent on the value of  $a$ , when it crosses  $\det(f) = 0, \operatorname{tr}(f) = 0, \det(f) = \frac{\operatorname{tr}(f)^2}{4}$  lines on the trace-determinant plane. So I will find these critical values algebraically.

$$\operatorname{tr}(f) = -1 + a = 0 \rightarrow \boxed{a = 1}$$

$a = 1$  is not critical value since  $\det(f) < 0$

$$\det(f) = -a = 0 \rightarrow \boxed{a = 0}$$

$$\begin{aligned} \delta &= \det(f) - \frac{\operatorname{tr}(f)^2}{4} = -a - \frac{(a-1)^2}{4} \\ &= a^2 + 2a + 1 \rightarrow \boxed{a = -1} \end{aligned}$$

Classification and visualization for  $a = -1.5, -1.0, -0.5, 0, 0.5$ :

```
f_n1.5 <- matrix(c(-1,0,-1.5,-1.5),nrow = 2)
f_n1.0 <- matrix(c(-1,0,-1.0,-1.0),nrow = 2)
f_n0.5 <- matrix(c(-1,0,-0.5,-0.5),nrow = 2)
f_p0.0 <- matrix(c(-1,0,+0.0,+0.0),nrow = 2)
f_p0.5 <- matrix(c(-1,0,+0.5,+0.5),nrow = 2)
```

```
classify(f_n1.5)
```

```
## [1] "sink"
```

```
classify(f_n1.0)
```

```
## [1] "degenerate sink"
```

```
classify(f_n0.5)
```

```
## [1] "sink"
```

```
classify(f_p0.0)
```

```
## [1] "stable line"
```

```
classify(f_p0.5)
```

```
## [1] "saddle"
```

Therefore we expect the following eigenvalue combinations (in order): -Distinct real

-One real

-Distinct real

-Distinct real

-Distinct real

```
## [1] "Eigenvalues: -1.5" "Eigenvalues: -1"
```

```
## [1] "Eigenvalues: -1" "Eigenvalues: -1"
```

```
## [1] "Eigenvalues: -1" "Eigenvalues: -0.5"
```

```
## [1] "Eigenvalues: 0" "Eigenvalues: -1"
```

```
## [1] "Eigenvalues: -1" "Eigenvalues: 0.5"
```

```
fs <- list(f_n1.5,f_n1.0,f_n0.5,f_p0.0,f_p0.5)
```

```
deters <- lapply(fs,det)
```

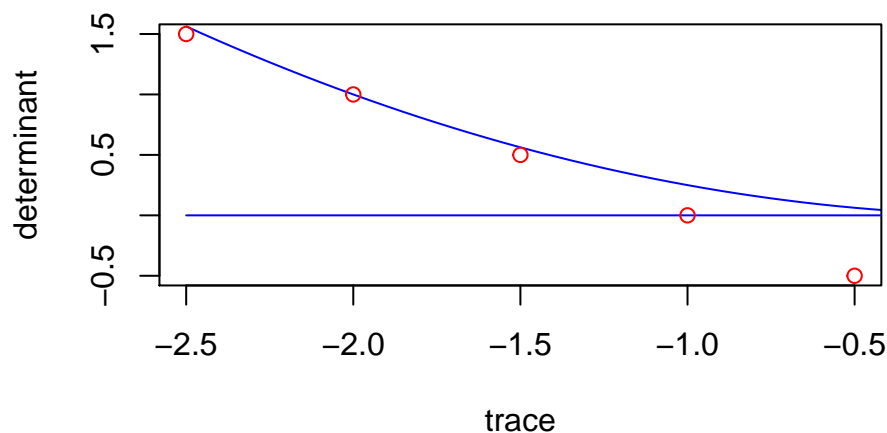
```
traces <- lapply(fs,function(A){sum(diag(A))})
```

```
vals <- seq(from = -2.5,to=2.5,length.out=101)
```

```
plot(vals,vals^2/4,col="blue",  
      xlim=c(-2.5,-.5),ylim=c(-0.5,1.5),  
      type="l",xlab="trace",ylab="determinant")
```

```
points(vals,rep(0,101),col="blue",type="l")
```

```
points(traces,deters,col="red")
```



Therefore, the system exhibits the following behavior:

Sink for  $(-\infty < a < -1) \cup (-1, 0)$

Degenerate sink for  $a = -1$

Stable line for  $a = 0$   
Saddle for  $(0 < a < \infty)$