BiomathHW07

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Problem #1

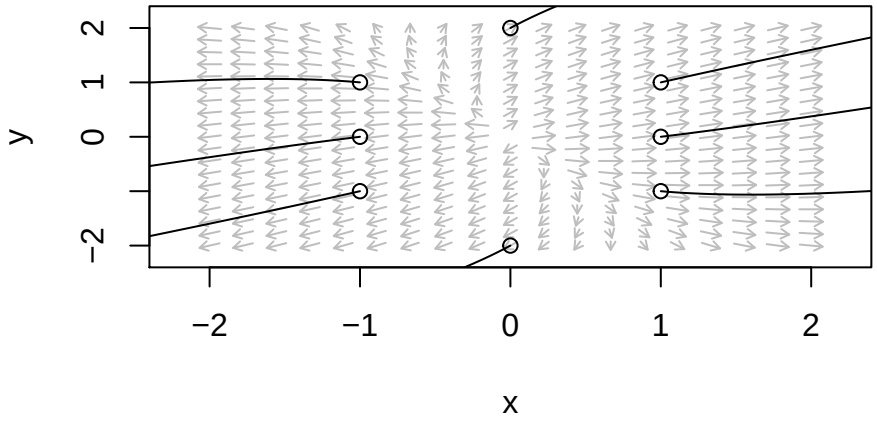
The following is a generic code to classify the behavior of a linear system based on it's trace and eigenvalues. I will use it in each part of the problem

```
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)

$$\dot{x} = 6x + 2y$$
$$\dot{y} = 2x + 3y$$

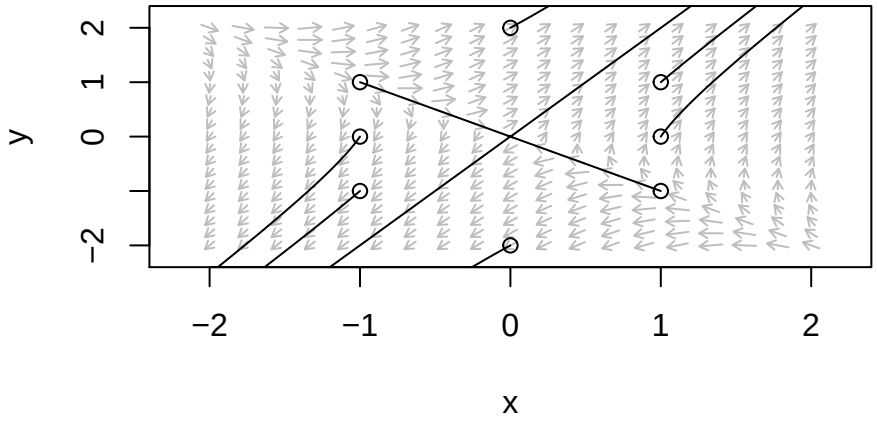
```
## [1] "Phase portrait type:  source"
## [1] "Eigenvalues:  7" "Eigenvalues:  2"
```



(b)

$$\dot{x} = x + 2y$$
$$\dot{y} = 4x + 3y$$

```
## [1] "Phase portrait type:  saddle"
## [1] "Eigenvalues:  5"  "Eigenvalues:  -1"
```

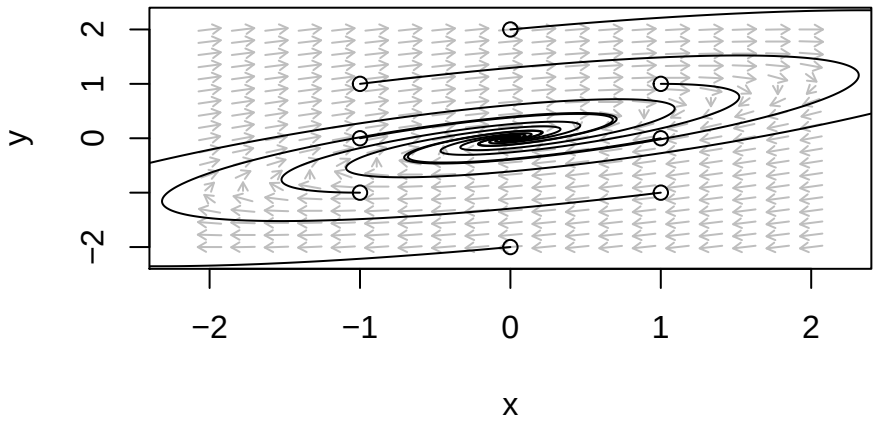


(c)

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

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

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

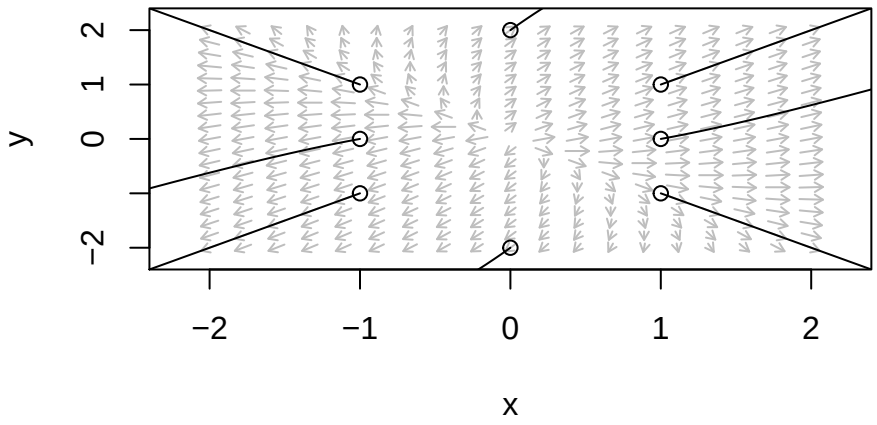


(d)

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

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

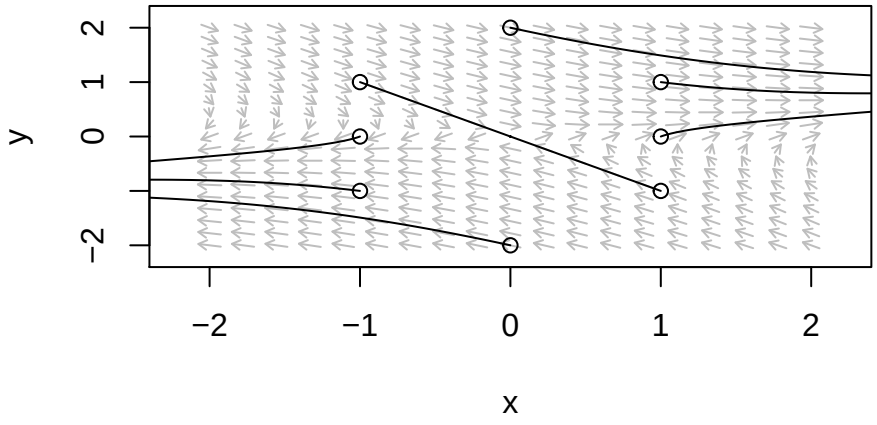
```
## [1] "Eigenvalues:  3" "Eigenvalues:  1"
```



(e)

$$\dot{x} = x + 5y$$
$$\dot{y} = x - 3y$$

```
## [1] "Phase portrait type:  saddle"
## [1] "Eigenvalues:  -4" "Eigenvalues:  2"
```



(f)

$$\dot{x} = -1x + ay$$
$$\dot{y} = 0x + ay$$
$$\text{for } a \neq 0$$

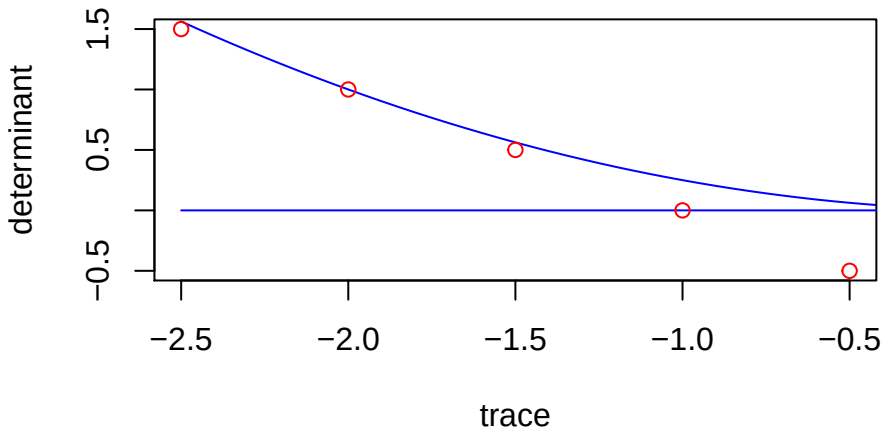
This function changes behavior, dependent on the value of a, when it crosses $det(f) = 0, tr(f) = 0, det(f) = \frac{tr(f)^2}{4}$ lines on the trace-determinant plane. So I will find these critical values alegbraically.

$$
\begin{aligned}
tr(f) = -1 + a = 0 \quad &\rightarrow \quad \boxed{a = 1} \\
a = 1 \text{ is not critical value since } &det(d) < 0 \\
det(f) = -a = 0 \quad &\rightarrow \quad \boxed{a = 0} \\
delta = det(f) - \frac{tr(f)^2}{4} &= -a - \frac{(a-1)^2}{4} \\
= a^2 + 2a + 1 \quad &\rightarrow \quad \boxed{a = -1}
\end{aligned}
$$

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"

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 $(0 < a < \infty)$