R Notebook

```
if (require("pacman")) {
    library(pacman)
  }else{
    install.packages("pacman")
    library(pacman)
  }
```

```
## Loading required package: pacman
```

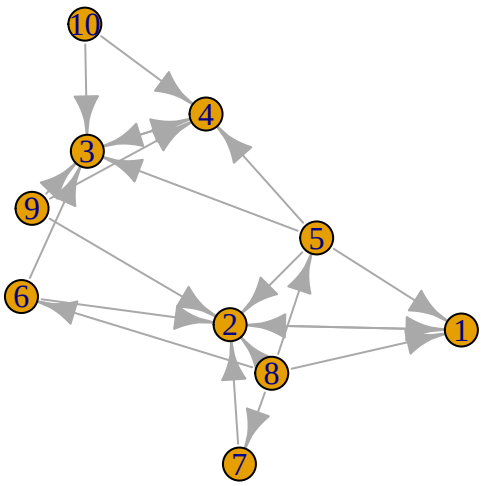
```
  pacman::p_load(tidyverse)
```

Implementing Page Rank Methods

Example Graph

Consider the following Graph taken from the paper:

```
g1 <- igraph::graph.formula(1++2, 1+-8, 1+-5, 2+-5, 2+-7, 2+-8, 2+-6, 2+-9, 3++4, 3+-5, 3+-6, 3+-9, 3+-
plot(g1)
```



Adjacency Matrix

The adjacency Matrix is given by:

```
A <- igraph::get.adjacency(g1, names = TRUE, sparse = FALSE) %>%
  as.matrix()

## Adjust the Order
(A <- A[order(as.integer(row.names(A))), order(as.integer(colnames(A)))])
```

```
##    1 2 3 4 5 6 7 8 9 10
## 1  0 1 0 0 0 0 0 0 0  0
```

```
## 2  1 0 0 0 0 0 0 0 0  0
## 3  0 0 0 1 0 0 0 0 0  0
## 4  0 0 1 0 0 0 0 0 0  0
## 5  1 1 1 1 0 0 0 0 0  0
## 6  0 1 1 0 0 0 0 0 0  0
## 7  0 1 0 0 0 0 0 0 0  0
## 8  1 1 0 0 1 1 1 0 0  0
## 9  0 1 1 1 0 0 0 0 0  0
## 10 0 0 1 1 0 0 0 0 0  0
```

State Distribution

The state distribution is the transpose of the adjacency matrix:

```
(p0 <- t(A))
```

```
##    1 2 3 4 5 6 7 8 9 10
## 1  0 1 0 0 1 0 0 1 0  0
## 2  1 0 0 0 1 1 1 1 1  0
## 3  0 0 0 1 1 1 0 0 1  1
## 4  0 0 1 0 1 0 0 0 1  1
## 5  0 0 0 0 0 0 0 1 0  0
## 6  0 0 0 0 0 0 0 1 0  0
## 7  0 0 0 0 0 0 0 1 0  0
## 8  0 0 0 0 0 0 0 0 0  0
## 9  0 0 0 0 0 0 0 0 0  0
## 10 0 0 0 0 0 0 0 0 0  0
```

Probability Transition Matrix

The probability transition matrix is such that each column of the initial state distribution (i.e. the transposed adjacency matrix) is scaled to 1.

```
p0 %*% diag(1/colSums(p0))
```

```
##    [,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8]      [,9] [,10]
## 1     0    1    0    0 0.25  0.0    0  0.2 0.0000000   0.0
## 2     1    0    0    0 0.25  0.5    1  0.2 0.3333333   0.0
## 3     0    0    0    1 0.25  0.5    0  0.0 0.3333333   0.5
## 4     0    0    1    0 0.25  0.0    0  0.0 0.3333333   0.5
## 5     0    0    0    0 0.00  0.0    0  0.2 0.0000000   0.0
## 6     0    0    0    0 0.00  0.0    0  0.2 0.0000000   0.0
## 7     0    0    0    0 0.00  0.0    0  0.2 0.0000000   0.0
## 8     0    0    0    0 0.00  0.0    0  0.0 0.0000000   0.0
## 9     0    0    0    0 0.00  0.0    0  0.0 0.0000000   0.0
## 10    0    0    0    0 0.00  0.0    0  0.0 0.0000000   0.0
```

```
adj_to_probTrans <- function(adjMat) {
  t(adjMat) %*% diag(1/colSums(t(adjMat)))
}

(T <- adj_to_probTrans(A)) %>% round(2)
```

Create a Function

```
##    [,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8] [,9] [,10]
```

```
## 1     0    1    0    0 0.25  0.0    0  0.2 0.00   0.0
## 2     1    0    0    0 0.25  0.5    1  0.2 0.33   0.0
## 3     0    0    0    1 0.25  0.5    0  0.0 0.33   0.5
## 4     0    0    1    0 0.25  0.0    0  0.0 0.33   0.5
## 5     0    0    0    0 0.00  0.0    0  0.2 0.00   0.0
## 6     0    0    0    0 0.00  0.0    0  0.2 0.00   0.0
## 7     0    0    0    0 0.00  0.0    0  0.2 0.00   0.0
## 8     0    0    0    0 0.00  0.0    0  0.0 0.00   0.0
## 9     0    0    0    0 0.00  0.0    0  0.0 0.00   0.0
## 10    0    0    0    0 0.00  0.0    0  0.0 0.00   0.0
```

Page Rank Random Surfer

The random surfer page rank method modifies the probability transition matrix T so that the method works also for non-ergodic graphs by introducing the possibility of a random jump, we'll call the surfer transition matrix S:

$$S = \lambda T + (1 - \lambda)\, B : \tag{1}$$

$$\tag{2}$$

$$B = \begin{bmatrix} \frac{1}{N} & \frac{1}{N} & \cdots & \frac{1}{N} \\ \frac{1}{N} & \frac{1}{N} & \cdots & \frac{1}{N} \\ \vdots & \vdots & \ddots & \vdots \\ \frac{1}{N} & \frac{1}{N} & \cdots & \frac{1}{N} \end{bmatrix} \tag{3}$$

$$N = ||V|| \tag{4}$$

$$\lambda \in [0, 1] \tag{5}$$

```
B <- matrix(rep(1/nrow(T), length.out = nrow(T)**2), nrow = nrow(T))
l <- 0.8

S <- l*T+(1-l)*B
```

Eigen Value Method

The eigenvector corresponding to the the eigenvalue of 1 will be the stationary point:

```
eigen(S, symmetric = FALSE)
```

```
## eigen() decomposition
## $values
##  [1]  1.000000e+00  8.000000e-01 -8.000000e-01 -8.000000e-01  3.117428e-09
##  [6] -3.117428e-09  1.233252e-17 -6.831762e-18  1.351981e-18  1.827902e-34
##
## $vectors
##              [,1]          [,2]          [,3]          [,4]          [,5]
##  [1,] -0.48531271  5.000000e-01  1.074642e-03  7.070996e-01  6.735753e-01
##  [2,] -0.52732002  5.000000e-01 -1.074642e-03 -7.070996e-01  9.622505e-02
##  [3,] -0.49152601 -5.000000e-01  7.071060e-01  3.197134e-03  9.622505e-02
##  [4,] -0.47977477 -5.000000e-01 -7.071060e-01 -3.197134e-03  2.886751e-01
##  [5,] -0.05288058  7.620569e-17  1.212133e-16 -3.419297e-18 -3.849002e-01
##  [6,] -0.05288058  7.620569e-17 -7.099634e-17 -3.419297e-18 -3.849002e-01
##  [7,] -0.05288058  7.620569e-17  2.943750e-17 -3.419297e-18 -3.849002e-01
##  [8,] -0.04558671  6.926804e-17 -1.948070e-18 -4.183296e-17 -7.499367e-09
```

```
##  [9,] -0.04558671  6.926804e-17  7.359376e-18 -4.183296e-17 -7.499367e-09
## [10,] -0.04558671  6.926804e-17  7.359376e-18 -4.183296e-17 -7.499367e-09
##                [,6]          [,7]          [,8]          [,9]         [,10]
##  [1,] -6.735753e-01 -2.658112e-01  5.357798e-01 -4.171123e-01  2.122431e-01
##  [2,] -9.622504e-02  1.896313e-01 -1.330801e-01 -7.334472e-02  1.241240e-01
##  [3,] -9.622504e-02  1.990137e-01 -1.665251e-01  2.591495e-01 -1.598069e-04
##  [4,] -2.886751e-01 -3.972519e-02  1.099005e-02 -1.886035e-02  6.690506e-02
##  [5,]  3.849002e-01 -7.585250e-01  5.323206e-01  2.933789e-01 -4.964959e-01
##  [6,]  3.849002e-01  4.774778e-01 -3.550303e-01  5.560197e-01 -1.341297e-01
##  [7,]  3.849002e-01  2.204565e-01 -5.047228e-01  1.987554e-01  1.561493e-01
##  [8,] -7.499367e-09 -1.208463e-16  1.668414e-16 -2.685569e-16  3.416862e-17
##  [9,] -7.499367e-09 -1.125898e-02  4.013399e-02 -3.989931e-01 -5.316106e-01
## [10,] -7.499367e-09 -1.125898e-02  4.013399e-02 -3.989931e-01  6.029746e-01
```

So in this case ~~the~~ a stationary point is $\langle -0.49, -0.53, -0.49, -0.48, -0.05, -0.05, -0.05, -0.04, -0.04, -0.04 \rangle$ which can be verified:

$$1\vec{p} = S\vec{p}$$

```
(p     <- eigen(S)$values[1] * eigen(S)$vectors[,1])
```

```
##  [1] -0.48531271 -0.52732002 -0.49152601 -0.47977477 -0.05288058 -0.05288058
##  [7] -0.05288058 -0.04558671 -0.04558671 -0.04558671
```

```
(p_new <- S %*% p)
```

```
##           [,1]
## 1  -0.48531271
## 2  -0.52732002
## 3  -0.49152601
## 4  -0.47977477
## 5  -0.05288058
## 6  -0.05288058
## 7  -0.05288058
## 8  -0.04558671
## 9  -0.04558671
## 10 -0.04558671
```

Power Value Method

Using the power method should give the same result, which it indeed does if it starts at the solution:

```
# p     <- c(1, 0, 0, 0, 0, 0, 0, 0, 0, 0)
# p_new
# p_new <- S %*% p

while (sum(round(p, 9) != round(p_new, 9))) {
    (p     <- p_new)
    (p_new <- S %*% p)
}

p_new
```

```
##           [,1]
## 1  -0.48531271
## 2  -0.52732002
```

```
## 3  -0.49152601
## 4  -0.47977477
## 5  -0.05288058
## 6  -0.05288058
## 7  -0.05288058
## 8  -0.04558671
## 9  -0.04558671
## 10 -0.04558671
```

```
p
```

```
##  [1] -0.48531271 -0.52732002 -0.49152601 -0.47977477 -0.05288058 -0.05288058
##  [7] -0.05288058 -0.04558671 -0.04558671 -0.04558671
```

Comparison

However it seems to converge to a different solution if the IC is otherwise:

```
p     <- c(1, 0, 0, 0, 0, 0, 0, 0, 0, 0)
p_new <- S %*% p

while (sum(round(p, 9) != round(p_new, 9))) {
    (p     <- p_new)
    (p_new <- S %*% p)
}

p_new
```

```
##         [,1]
## 1  0.2129185
## 2  0.2313481
## 3  0.2156444
## 4  0.2104889
## 5  0.0232000
## 6  0.0232000
## 7  0.0232000
## 8  0.0200000
## 9  0.0200000
## 10 0.0200000
```

```
p
```

```
##         [,1]
## 1  0.2129185
## 2  0.2313481
## 3  0.2156444
## 4  0.2104889
## 5  0.0232000
## 6  0.0232000
## 7  0.0232000
## 8  0.0200000
## 9  0.0200000
## 10 0.0200000
```