Example 1

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```
library(knitr)
setwd("/Users/KevQuant/Desktop/Depaul/csc495/wk1/ex1")
read_chunk("example01_2.R")
knitr::opts_chunk$set(echo = TRUE)
```

Load packages

```
# Load packages
library("ggplot2")
# Must load other packages first
library("sand")
```

Load the data

```
# Load the data.
setwd("/Users/KevQuant/Desktop/Depaul/csc495/wk1/ex1")
dolphin <- read.graph("dolphin2.graphml", format="graphml")</pre>
```

Graph information

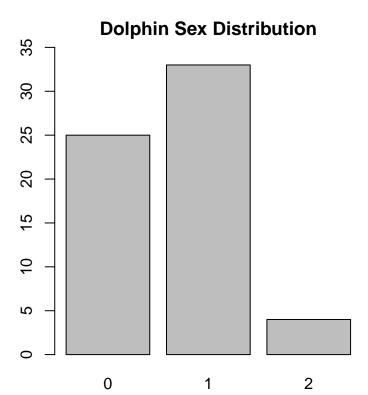
```
# Display basic information about the graph
# Summary
summary(dolphin)
## IGRAPH U--- 62 159 --
## + attr: label (v/c), Sex (v/n), id (v/c)
# Is it directed? (use functions with _ rather than .)
is_directed(dolphin)
## [1] FALSE
# Is it simple?
is_simple(dolphin)
## [1] TRUE
# Count nodes
vcount(dolphin)
## [1] 62
# Count edges
ecount(dolphin)
## [1] 159
```

Graph elements

```
# List graph elements
# Nodes
V(dolphin)
## + 62/62 vertices:
## [1] 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23
## [24] 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43 44 45 46
## [47] 47 48 49 50 51 52 53 54 55 56 57 58 59 60 61 62
# Edges
E(dolphin)
## + 159/159 edges:
    [1] 4-- 9 6--10 7--10 1--11 3--11 6--14 7--14 10--14 1--15 4--15
   [11] 1--16 15--17 2--18 7--18 10--18 14--18 16--19 2--20 8--20 9--21
##
   [21] 17--21 19--21 19--22 18--23 15--25 16--25 19--25 18--26 2--27 26--27
## [31] 2--28 8--28 18--28 26--28 27--28 2--29 9--29 21--29 11--30 19--30
## [41] 22--30 25--30 8--31 20--31 29--31 18--32 10--33 14--33 13--34 15--34
   [51] 17--34 22--34 15--35 34--35 30--36 2--37 21--37 24--37 15--38 17--38
##
   [61] 21--38 34--38 37--39 1--40 8--40 15--40 16--40 34--40 37--40 40--62
## [71] 2-41 10-41 14-41 1-42 3-42 11-42 31-42 15-43 30-43 34-43
## [81] 38--43 43--62 3--44 21--44 35--44 38--44 9--45 16--45 19--45 22--45
## [91] 24--45 25--45 30--45 45--62 43--46 1--47 11--47 21--47 29--47 31--47
## + ... omitted several edges
```

Histogram

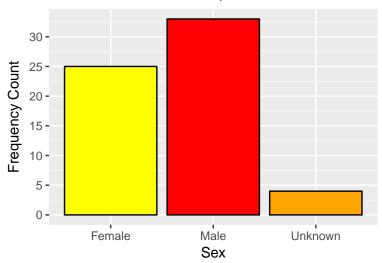
```
# Histogram of sex distribution
par(mar=c(2,2,2,2))
barplot(table(V(dolphin)$Sex), ylim = c(0,35),main = "Dolphin Sex Distribution")
```



GGPlot version

```
# GGPlot version
# Note that node attributes cannot be factors
sex<-factor(V(dolphin)$Sex,c(0,1,2),labels = c("Female","Male","Unknown"))
g <- ggplot(data.frame(sex),aes(x=sex))
g<-g+geom_histogram(binwidth=1,stat="count",colour="black",fill=c("yellow","red","orange"))
## Warning: Ignoring unknown parameters: binwidth, bins, pad
g<-g+xlab("Sex")
g<-g+ylab("Frequency Count")
g<-g+ggtitle("Sex Distribution in Dolphin Dataset")
g<-g+scale_y_continuous(breaks = seq(0,40,5))
print(g)</pre>
```

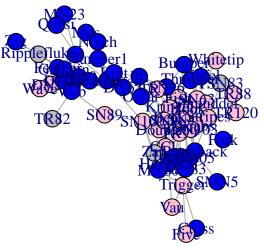
Sex Distribution in Dolphin Dataset



```
##Altervative plot option
#g <- g + geom_bar()
#print(g)</pre>
```

Network visualization

```
# Create a network visualization. Note that the layout has a random element, so your visualization will
# Make the male nodes blue, female nodes pink, and unknown grey
dolphin.col<-factor(V(dolphin)$Sex,c(0,1,2),labels = c("pink","blue","grey"))
plot(dolphin,vertex.color=as.character(dolphin.col))</pre>
```



```
#altnative
#dolphin.col<-rep("pink",each=vcount(dolphin))
#dolphin.col<-seq(1:vcount(dolphin))
#dolphin.col[V(dolphin)$Sex==1]<-"lightblue"
#dolphin.col[V(dolphin)$Sex==2]<-"grey"
#dolphin.col[V(dolphin)$Sex==0]<-"red"
#plot(dolphin,vertex.color=as.character(f))</pre>
```

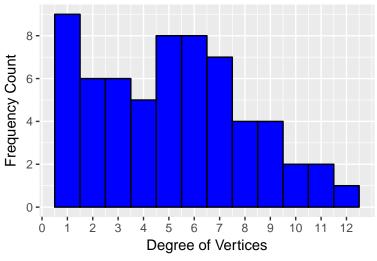
Degree distribution

```
# Degree distribution
# Compute degrees of each node (degree sequence)
deg<-degree(dolphin)</pre>
deg
##
                  3
                                                8 12
   [1]
                                 6
                                    7
                                       5
## [24]
                 3 5
                       5
                          9
                             5
                                1
                                    3 10
                                         5
                                            1
                                               7
                                                   8
                                                              6 7 4 11
                                2
                                    2
              2 7 10 4 2 7
                                      9
                                         1
                                            5
## [47]
```

Histogram

```
# Histogram with ggplot (boundary, binwidth)
g<-ggplot(data.frame(Degree=deg),aes(x=Degree))
g<-g+geom_histogram(binwidth=1,stat="bin",col="black",fill="blue")
g<-g+ylab("Frequency Count")
g<-g+xlab("Degree of Vertices")
g<-g+gtitle("Histogram of Degree of Dolphin Vertice")
g<-g+scale_x_continuous(breaks = 0:12)
g<-g+scale_y_continuous(breaks = seq(0,10,2))
print(g)</pre>
```

Histogram of Degree of Dolphin Vertice



```
#Altnative option
#g<-g+geom_bar()
#print(g)</pre>
```

Degree distribution by sex

Setting up the data

```
# Degree distribution of male and female dolphins.

# Set up the data
```

```
# First make a data frame with the degree and sex. Then remove the "Unknown" dolphins.
dol.df<-data.frame(Degree=deg, Sex=V(dolphin)$Sex)
dol.df<-dol.df[dol.df$Sex!=2,]

# Then remake the sex factor so that the "Unknown" option is omitted.
dol.df$Sex<-factor(dol.df$Sex,c(0,1),labels=c("Female","Male"))</pre>
```

Plotting

```
# GGPlot boxplot
g<-ggplot(dol.df,aes(x=Sex,y=Degree,color=Sex))
g<-g+geom_boxplot(outlier.colour="red", outlier.shape=16,outlier.size=3)
g<-g+ggtitle("Degree distribution of male and female dolphins")
g<-g+scale_y_continuous(breaks = seq(0,13,2))
print(g)</pre>
```

Degree distribution of male and female dolp

