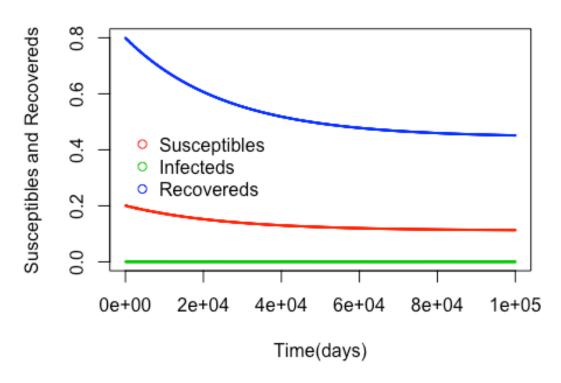
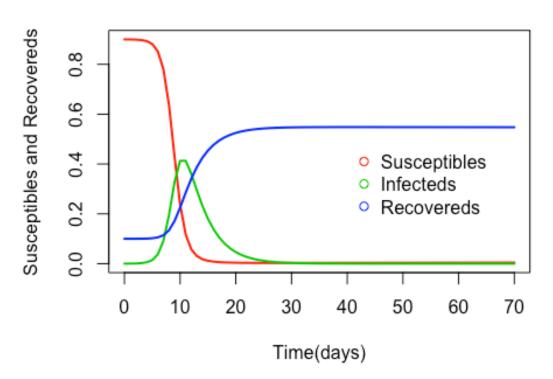
## SIR model with mortality(frequency-dependent)

```
## It is the SIR model with a probability of mortality, and uneugal births and
## This code assumes Frequency-Dependent Transmission
## Parameters:
# rho: The mortality probability;
# nu: The population level birth rate;
# miu: Per captia death rate, and the population level birth rate;
# beta: Product of contact rates and transmission probability;
# gamma: Recovery Rate;
# X: The number or density of susceptible individuals;
# Y: The number or density of infectious individuals;
# Z: The number or density of recovered individuals.
library(deSolve)
SIR.freq = function(time, state, pars) {
  with(as.list(c(state,pars)), {
    dX = nu - beta * X0 * Y0 / (X0 + Y0 + Z0) - miu * X0
    dY = beta * X0 * Y0 / (X0 + Y0 + Z0) - (gamma+miu)/(1-rho) * Y0
    dZ = gamma * Y0 - miu * Z0
    return(list(c(dX,dY,dZ)))
 })
# condition 1
yini = c(X0 = 0.2, Y0 = 1e-4, Z0 = 1-0.2-1e-4)
pars = c(\text{rho} = 0.5, \text{nu} = 1/(70*365), \text{miu} = 1/(70*365), \text{beta} = 520/365, \text{gamma} =
times = seq(0,1e5,by = 1)
out = ode(func = SIR.freq, y = yini, parms = pars, times = times)
out = as.data.frame(out)
out$time = NULL
matplot(times,out,type = "1", xlab = "Time(days)", ylab = "Susceptibles and
Recovereds", main = "SIR Model with a probability of mortality",
        lwd = 2, lty = 1, col = 2:4)
legend(40,0.5,c("Susceptibles","Infecteds","Recovereds"), pch = 1, col =
2:4,bty = "n")
```

## SIR Model with a probability of mortality



## SIR Model with a probability of mortality



## SIR Model with a probability of mortality

