Specifying a probability cutoff (the minimal probability for which we can say that two genes are linked), we can plot the network :

```
showNetwork (result,    0.3 )

## Loading required package:   sna
## sna:   Tools for Social Network Analysis
## Version 2.3-2 created on 2014-01-13.
## copyright (c) 2005, Carter T. Butts, University of California-Irvine
##    For citation information, type citation("sna").
##    Type help(package="sna") to get started.
##
```