portal sim

```
library(tidyverse)
## -- Attaching packages --
## v ggplot2 3.0.0
                      v purrr
                                  0.2.5
## v tibble 1.4.2
                       v dplyr
                                  0.7.6
## v tidyr
             0.8.1
                       v stringr 1.3.1
## v readr
             1.1.1
                       v forcats 0.3.0
## -- Conflicts -----
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                     masks stats::lag()
p2 <- read.csv("~/../Dropbox/negative_population_trends/portal2.csv")
p3 <- read.csv("~/../Dropbox/negative_population_trends/portal3.csv")
p4 <- read.csv("~/../Dropbox/negative_population_trends/portal4.csv")</pre>
p2 %>% group_by(bar, model) %>%
  summarize(samplesize=n(),
            negativebeta = sum(beta<0),</pre>
            percent = negativebeta/samplesize*100)
## # A tibble: 4 x 5
## # Groups:
               bar [?]
##
     bar
                                         model samplesize negativebeta percent
##
     <fct>
                                                     <int>
                                                                  <int>
                                                                           <dbl>
## 1 True Population Size, 2 Highest P~ line~
                                                     20000
                                                                  13063
                                                                           65.3
## 2 True Population Size, 2 Highest P~ log
                                                     20000
                                                                  12970
                                                                           64.8
## 3 True Population Size, 2 Random Po~ line~
                                                     20000
                                                                   9986
                                                                           49.9
## 4 True Population Size, 2 Random Po~ log
                                                     20000
                                                                  10013
                                                                           50.1
p3 %>% group_by(bar, model) %>%
  summarize(samplesize=n(),
            negativebeta = sum(beta<0),</pre>
            percent = negativebeta/samplesize*100)
## # A tibble: 4 x 5
## # Groups:
               bar [?]
##
     har
                                         model samplesize negativebeta percent
##
     <fct>
                                         <fct>
                                                    <int>
                                                                  <int>
                                                                           <dbl>
## 1 True Population Size, 2 Highest P~ line~
                                                     20000
                                                                  13239
                                                                           66.2
## 2 True Population Size, 2 Highest P~ log
                                                     20000
                                                                  13129
                                                                           65.6
## 3 True Population Size, 2 Random Po~ line~
                                                     20000
                                                                  10067
                                                                           50.3
## 4 True Population Size, 2 Random Po~ log
                                                     20000
                                                                  10039
                                                                           50.2
p4 %>% group_by(bar, model) %>%
  summarize(samplesize=n(),
            negativebeta = sum(beta<0),</pre>
            percent = negativebeta/samplesize*100)
## # A tibble: 4 x 5
## # Groups:
               bar [?]
##
     bar
                                         model samplesize negativebeta percent
##
     <fct>
                                         <fct>
                                                     <int>
                                                                  <int>
                                                                           <dbl>
```

##	1	True	Population	Size,	2 Highest P~	line~	20000	13167	65.8
##	2	True	Population	Size,	2 Highest P~	log	20000	13043	65.2
##	3	True	Population	Size,	2 Random Po~	line~	20000	9900	49.5
##	4	True	Population	Size,	2 Random Po~	log	20000	9877	49.4