

Group Randomized Trial Design For Targeted Agent – A Simulation Study

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```
library(powerSurvEpi)
library(stats)
library(base)
library(survival)
library(PwrGSD)
library(ggplot2)
```

The simulation

```
simulation <- function(sims,countries,positive_only,assumption,test,T0,T1){
  # sims: How many simulations
  # countries: How many countries
  # positive_only: Include only positive patients or not
  # T0: End of accrual (in year)
  # T1: End of study (in year)
  #
  # test: Type of hypothesis testing
  # # "log rank": Perform a log rank test
  # # "t": Perform a one-tailed t test on the five-year survival rate
  # assumption: ssumption of efficacy of the drug
  # # "best": 45% for +, 40% for -
  # # "unif": unif(0.35-0.45) for +, unif(0.3-0.4) for -
  T0 <- 365*T0
  T1 <- 365*T1
  Total_N <- 0
  f <- c()
  significance <- c()
  p_list <- c()
  N <- 100

  for (round in 1:sims){
    if (round%%50 == 0){
      cat("Round ", round, "\n")
    }
    big_dat <- data.frame()
    cur_id <- 1
    cur_time <- 0
    accrual_end <- 0
    i <- 1
    while(accrual_end==0){
      set.seed(round*1001+i)
      x <- rbinom(N,1,0.25) #Biomarker status
      g <- rbinom(N,1,0.5) #Group assignment
    }
  }
}
```

```

sp_list <- c()
s <- c()
c <- c()
delta <- c() #Censoring status
follow_up <- c() #Actual follow-up days since intake (considering censoring)
five_yr_live <- c() #Alive at five year of follow-up or not
accrued <- N
for (j in 1:N){
  #Simulate entry time in days
  gap <- rexp(1,(72*countries)/365) #Si-S{i-1}, assuming 3 countries: 72*3 = 216
  cur_time <- cur_time+gap
  if (cur_time > T0){ #Stop Accrual
    accrued <- j-1
    accrual_end <- 1
    break
  }
  s <- c(s,cur_time)
  #Simulate survival probability of 5 years
  sp <- 0
  if (g[j] == 1){
    if (assumption=="unif"){
      #Uniform Scenario
      if (x[j] == 1) sp <- runif(1,0.35,0.45)
      else sp <- runif(1,0.3,0.4)
    }
    else if (assumption=="best"){
      #Best case scenario
      if (x[j] == 1) sp <- 0.45
      else sp <- 0.4
    }
  }else{
    sp <- 0.3
  }
  sp_list <- c(sp_list,sp)
  #Simulate censoring time since intake (in days)
  c_day <- rexp(1,0.05/365)
  c <- c(c,c_day)
}
if (accrued == 0){
  break
}
lambda <- -log(sp_list)/5
t <- c() #Survival time since intake (in days)
for (j in 1:accrued){
  st <- rexp(1,lambda[j])*365
  t <- c(t,st)
  delta <- c(delta,ifelse(c[j]<=t[j],1,0)) #right censored
  follow_up <- c(follow_up,min(t[j],c[j]))
  if (s[j]+follow_up[j]>T1){
    follow_up[j] <- T1-s[j]
    delta[j] <- 1 #right censored
  }
  five_yr_live <- c(five_yr_live,ifelse(t[j]>=1825,1,0))
}

```

```

    }
    id <- seq(cur_id,cur_id+accrued-1)
    cur_id <- cur_id+accrued
    if (accrual_end){
      x <- x[1:accrued]
      g <- g[1:accrued]
    }
    dat <- data.frame(id=id,x=x,g=g,s=s,t=t,c=c,delta=delta,follow_up=follow_up,five_yr_live=five_yr_live)
    big_dat <- rbind(big_dat,dat)
    i <- i + 1
  }
  if(positive_only){
    sub <- subset(big_dat,x==1)
  }else{
    sub <- big_dat
  }
  Total_N <- Total_N + length(sub$id)
  f <- c(f,sum(sub$follow_up)/length(sub$follow_up))
  if (test=="log rank"){
    s <- survdiff(Surv(follow_up,delta==0) ~ g,data=sub)
    p <- 1 - pchisq(s$chisq, 1)
    p_list <- c(p_list,p)
    #print(p)
    significance <- c(significance,ifelse(p<=0.05,1,0))
  }else if (test == "t"){
    sub <- subset(sub,c>t) #delete dropouts
    trt <- subset(sub,g==1)
    ctr <- subset(sub,g==0)
    #print(length(ctr$five_yr_live))
    s <- t.test(ctr$five_yr_live,trt$five_yr_live,alternative = "less", var.equal = TRUE)
    p_list <- c(p_list,s$p.value)
    significance <- c(significance,ifelse(s$p.value<=0.05,1,0))
  }else{
    print("Please enter correct test type!")
    quit()
  }
}

cat("T0 = ", T0, "\n")
cat("T1 = ", T1, "\n")
cat("test = ", test, "\n")
cat("assumption = ", assumption, "\n")
cat("positive_only = ", positive_only, "\n")
cat("countries = ", countries, "\n")
cat("power = ",mean(significance), "\n")
cat("Mean total sample size = ",Total_N/sims, "\n")
cat("Mean follow_up time per subject= ",mean(f), "\n\n\n")

sim <- data.frame(T0=T0,T1=T1,assumption=assumption,positive_only=positive_only,countries=countries,sims=sims)
return (sim)
}

# direct output to a file

```

```
#sink("/Users/river/Documents/699/Project_4/output.txt", append=TRUE, split=FALSE)
```

```
sim_results <- data.frame()

for (countries in 3:4){
  for (positive_only in c(TRUE,FALSE)){
    for (assumption in c("best","unif")){
      for (T0 in 3:5){
        for (T1 in 5:10){
          test <- "log rank"
          sims <- 0 #Set to 0 for output purpose only; should be 1000
          sim <- simulation(sims,countries,positive_only,assumption,test,T0,T1)
          sim_results<-rbind(sim_results,sim)
        }
      }
    }
  }
}

#sink()
sim_results$T0 <- sim_results$T0/365
sim_results$T1 <- sim_results$T1/365
```

Simulation Results

```
#write.table(sim_results, file = "sim_results.csv", sep = ",", col.names = NA, qmethod = "double")
#For output purpose only, loading saved results from simulation
sim_results <- read.table("/Users/river/Documents/699/Project_4/sim_results.csv",sep=",",header = TRUE)
#Listing results that have power >= 0.7
subset(sim_results,power>=0.7)[,2:9]
```

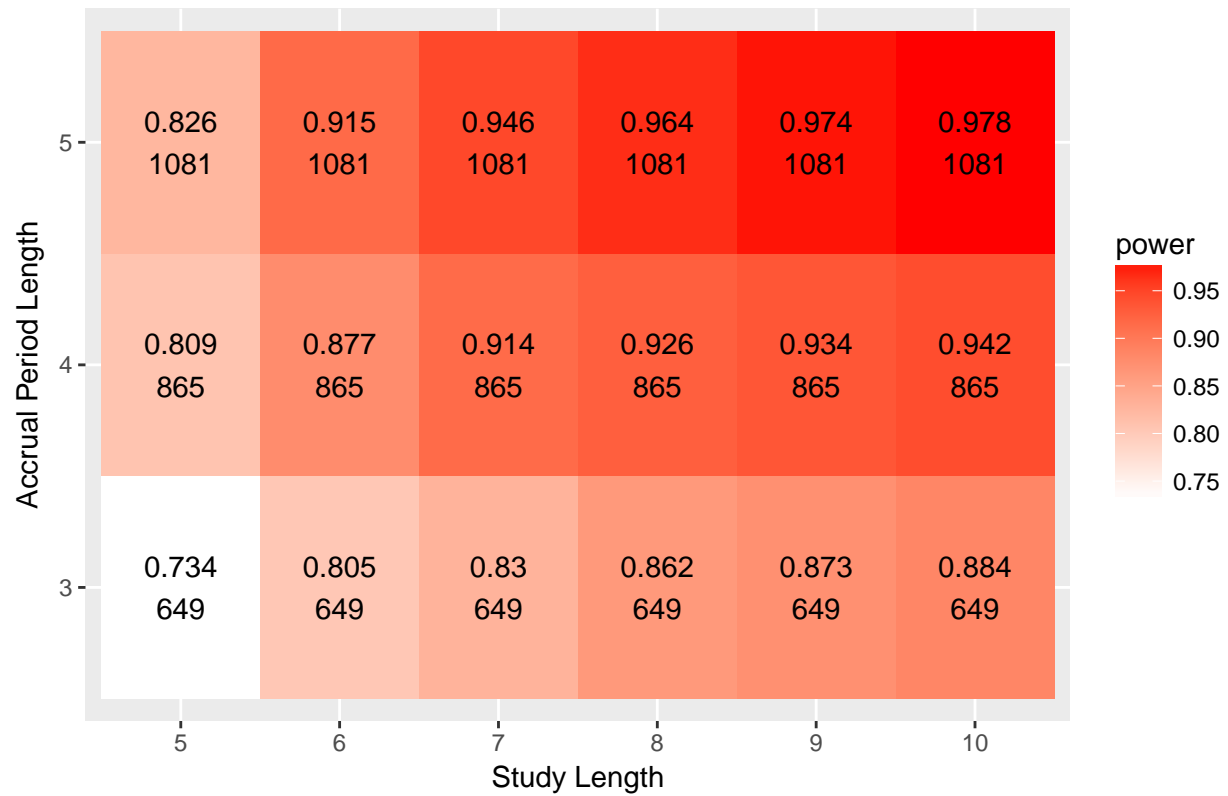
##	T0	T1	assumption	positive_only	countries	sample_size	power	follow_up
## 12	4	10	best	TRUE	3	216.560	0.713	1262.2956
## 16	5	8	best	TRUE	3	270.299	0.720	1070.3796
## 17	5	9	best	TRUE	3	270.299	0.753	1159.2970
## 18	5	10	best	TRUE	3	270.299	0.787	1229.1874
## 37	3	5	best	FALSE	3	648.561	0.734	826.5350
## 38	3	6	best	FALSE	3	648.561	0.805	960.5547
## 39	3	7	best	FALSE	3	648.561	0.830	1064.6032
## 40	3	8	best	FALSE	3	648.561	0.862	1145.2311
## 41	3	9	best	FALSE	3	648.561	0.873	1207.7733
## 42	3	10	best	FALSE	3	648.561	0.884	1256.2934
## 43	4	5	best	FALSE	3	865.106	0.809	732.7468
## 44	4	6	best	FALSE	3	865.106	0.877	888.0599
## 45	4	7	best	FALSE	3	865.106	0.914	1008.5421
## 46	4	8	best	FALSE	3	865.106	0.926	1102.0410
## 47	4	9	best	FALSE	3	865.106	0.934	1174.6468
## 48	4	10	best	FALSE	3	865.106	0.942	1230.9584
## 49	5	5	best	FALSE	3	1080.642	0.826	620.0889
## 50	5	6	best	FALSE	3	1080.642	0.915	800.9620
## 51	5	7	best	FALSE	3	1080.642	0.946	941.0860

##	52	5	8	best	FALSE	3	1080.642	0.964	1049.7834
##	53	5	9	best	FALSE	3	1080.642	0.974	1134.1100
##	54	5	10	best	FALSE	3	1080.642	0.978	1199.5141
##	77	3	9	best	TRUE	4	216.560	0.700	1238.0129
##	78	3	10	best	TRUE	4	216.560	0.713	1290.9061
##	81	4	7	best	TRUE	4	288.530	0.745	1026.9596
##	82	4	8	best	TRUE	4	288.530	0.772	1125.3156
##	83	4	9	best	TRUE	4	288.530	0.814	1202.5775
##	84	4	10	best	TRUE	4	288.530	0.822	1263.2088
##	86	5	6	best	TRUE	4	360.406	0.729	812.2796
##	87	5	7	best	TRUE	4	360.406	0.796	957.3533
##	88	5	8	best	TRUE	4	360.406	0.842	1070.9712
##	89	5	9	best	TRUE	4	360.406	0.866	1160.1464
##	90	5	10	best	TRUE	4	360.406	0.886	1230.1165
##	109	3	5	best	FALSE	4	865.106	0.850	826.7241
##	110	3	6	best	FALSE	4	865.106	0.898	960.9169
##	111	3	7	best	FALSE	4	865.106	0.918	1065.0849
##	112	3	8	best	FALSE	4	865.106	0.933	1145.9542
##	113	3	9	best	FALSE	4	865.106	0.938	1208.7147
##	114	3	10	best	FALSE	4	865.106	0.947	1257.4264
##	115	4	5	best	FALSE	4	1152.824	0.899	732.9600
##	116	4	6	best	FALSE	4	1152.824	0.949	888.3083
##	117	4	7	best	FALSE	4	1152.824	0.964	1008.8364
##	118	4	8	best	FALSE	4	1152.824	0.978	1102.3353
##	119	4	9	best	FALSE	4	1152.824	0.984	1174.8590
##	120	4	10	best	FALSE	4	1152.824	0.989	1231.0889
##	121	5	5	best	FALSE	4	1441.158	0.928	619.9900
##	122	5	6	best	FALSE	4	1441.158	0.974	800.9211
##	123	5	7	best	FALSE	4	1441.158	0.983	941.1725
##	124	5	8	best	FALSE	4	1441.158	0.991	1049.8935
##	125	5	9	best	FALSE	4	1441.158	0.997	1134.2076
##	126	5	10	best	FALSE	4	1441.158	0.998	1199.5456
##	138	4	10	unif	FALSE	4	1150.819	0.706	1183.4834
##	142	5	8	unif	FALSE	4	1438.970	0.718	1017.7893
##	143	5	9	unif	FALSE	4	1438.970	0.764	1095.4418
##	144	5	10	unif	FALSE	4	1438.970	0.794	1154.7509

```
best_3_f <- subset(sim_results,assumption=="best" & countries==3 & positive_only==FALSE)
unif_4_f <- subset(sim_results,assumption=="unif" & countries==4 & positive_only==FALSE)
best_4_f <- subset(sim_results,assumption=="best" & countries==4 & positive_only==FALSE)
best_4_t <- subset(sim_results,assumption=="best" & countries==4 & positive_only==TRUE)
```

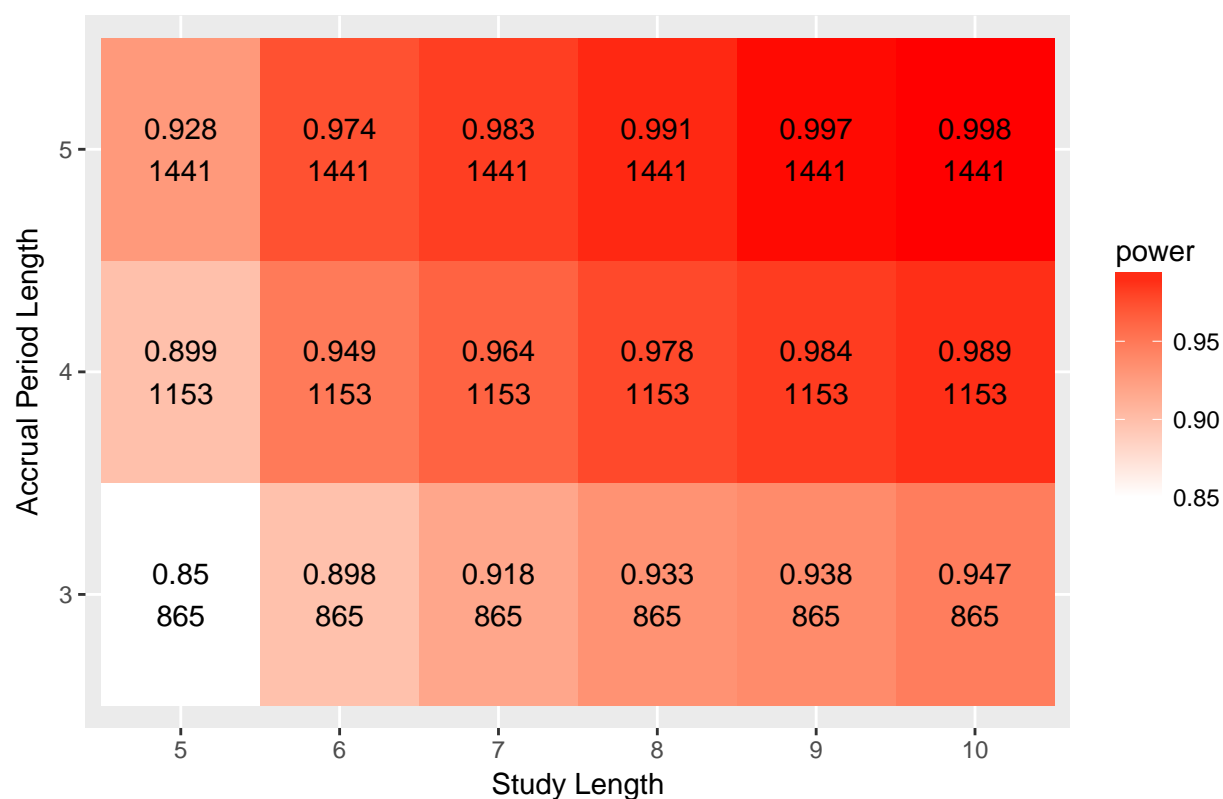
```
ggplot(best_3_f, aes(as.factor(T1), as.factor(T0))) +
  geom_tile(aes(fill = power)) +
  geom_text(aes(label = paste(power,round(sample_size), sep="\n"))) +
  scale_fill_gradient(low = "white", high = "red") +
  ggtitle("Best case scenario, 3 countries, including all") +
  labs(x="Study Length",y="Accrual Period Length")
```

Best case scenario, 3 countries, including all



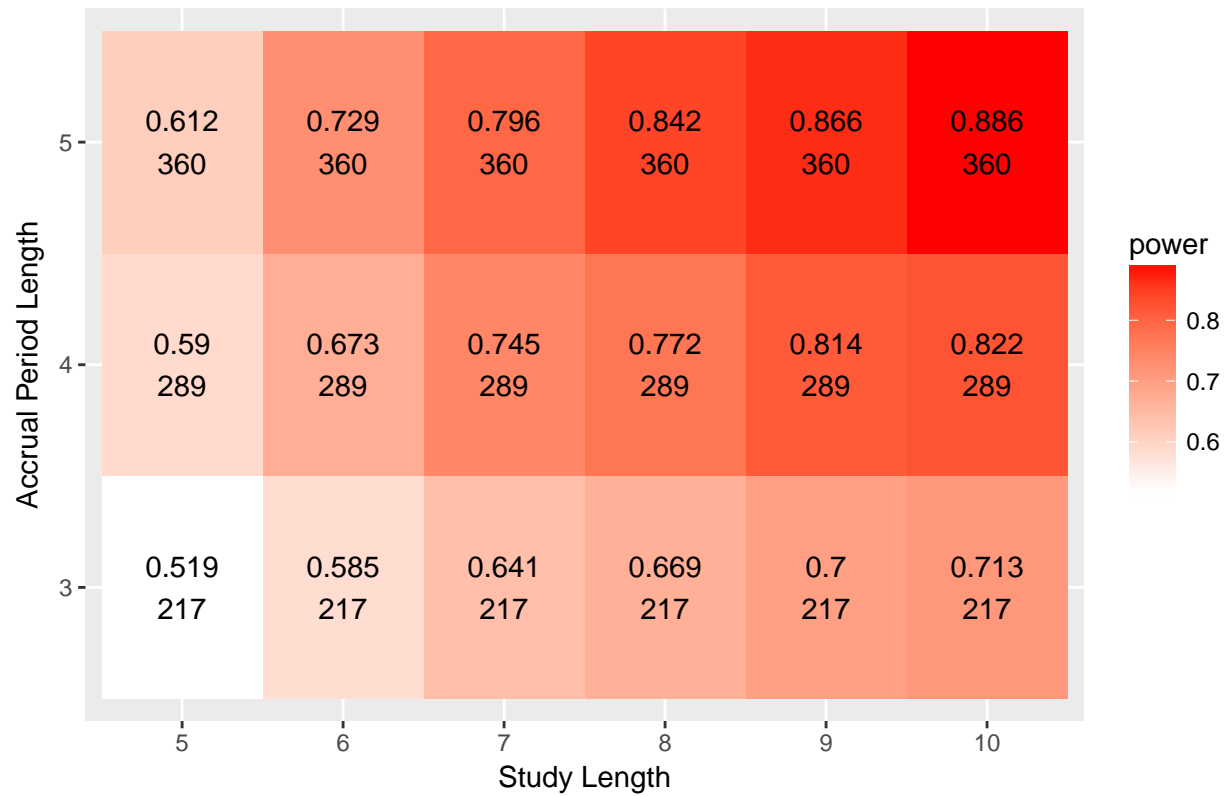
```
ggplot(best_4_f, aes(as.factor(T1), as.factor(T0))) +
  geom_tile(aes(fill = power)) +
  geom_text(aes(label = paste(power, round(sample_size), sep="\n"))) +
  scale_fill_gradient(low = "white", high = "red") +
  ggtitle("Best case scenario, 4 countries, including all") +
  labs(x="Study Length", y="Accrual Period Length")
```

Best case scenario, 4 countries, including all



```
ggplot(best_4_t, aes(as.factor(T1), as.factor(T0))) +
  geom_tile(aes(fill = power)) +
  geom_text(aes(label = paste(power, round(sample_size), sep="\n"))) +
  scale_fill_gradient(low = "white", high = "red") +
  ggtitle("Best case scenario, 4 countries, positive only") +
  labs(x="Study Length", y="Accrual Period Length")
```

Best case scenario, 4 countries, positive only



```
ggplot(unif_4_f, aes(as.factor(T1), as.factor(T0))) +
  geom_tile(aes(fill = power)) +
  geom_text(aes(label = paste(power, round(sample_size), sep="\n"))) +
  scale_fill_gradient(low = "white", high = "red") +
  ggtitle("Uniform scenario, 4 countries, including all") +
  labs(x="Study Length", y="Accrual Period Length")
```


Uniform scenario, 4 countries, including all

