

dalGen: borrow both arms

Yi Li

08/12/2022

```
setwd("/Users/yili/Desktop/EPIB 635/Final Project")
library(VGAM)
```

```
## Warning: package 'VGAM' was built under R version 4.1.2
```

```
## Loading required package: stats4
```

```
## Loading required package: splines
```

Modify the code a bit to borrow treatment arm as well.

```
# posterior weight: corresponds to  $w_R$  in Eq. (7) in Schmidli et al. (2014)
w_bar = function(w, y, n, a, b, a0 = 1, b0 = 1) {
  f = beta(a + y, b + n - y)/beta(a, b)
  f0 = beta(a0 + y, b0 + n - y)/beta(a0, b0)
  w_bar = w*f/(w*f + (1 - w)*f0)
  return(w_bar)
}

options(warn = -1)
trial2 = function(TE, w,
                  ni = c(132, 264),
                  # keep the same total no. of cases at interim and final analyses
                  a0 = 1, b0 = 1) {
  n = c(ni[1], diff(ni, 1))
  theta = (1 - TE)/(2 - TE)
  p_eff = NULL
  p_success = NULL
  y_i = 0
  i = 0

  repeat {
    i = i + 1
    y_i = y_i + rbinom(1, n[i], theta) #  $y_i \sim \text{Binom}(n_i, \theta)$ 
    ai = 1 + y_i
    bi = 1 + ni[i] - y_i

    ### Add robust MAP prior
    # posterior weight:
    wb = w_bar(w, y_i, n[i], ai, bi)
```

```

# P(TE > 0|data):
# In power_and_mixture_priors.html: y1 is # of cases in historical trial (ctrl arm) , and y is # of cases in trt arm

peff_new = wb*pbeta(0.5,
                    ai + 226 + 180, # add 180 cases in trt to y1 in MAP code
                    bi + (226 + 180) - 226) +
(1-wb)*pbeta(0.5,
              ai,
              bi)

p_eff = c(p_eff, peff_new)

p_success = c(p_success,
              wb*pbetabinom.ab(q = 117,
                                size = 264,
                                shape1 = ai + 226 + 180,
                                # add 180 cases in trt to y1
                                shape2 = bi + (226 + 180) - 226,
                                log = FALSE) +
              (1-wb)*pbetabinom.ab(q = 117,
                                    size = 264,
                                    shape1 = ai,
                                    shape2 = bi,
                                    log = FALSE)
              )

if (i <= 1) {
  # P(TE > 0|data) > 0.995;
  if (peff_new > 0.995 | p_success[length(p_success)] < 0.05) break
} else break
}
out = list(round(p_eff, 2), round(p_success, 8))
return(out)
}

trial2(TE = 0.7, w = 0.7)

```

```

## [[1]]
## [1] 0.06 1.00
##
## [[2]]
## [1] 0.05668654 0.99995963

```

```

trial2(TE = 0.7, w = 0.5) # decrease w because trt arm in historical trial may not be reliable

```

```

## [[1]]
## [1] 0.12 1.00
##
## [[2]]
## [1] 0.1190636 0.9999741

```

3.1 FPR

FPR looks fine if we keep $w = 0.7$

```
options(warn = -1)
fp = NULL
p_eff <- p_eff_unlist <- check <- list()
for (m in 1:10000) {
  p_eff = trial2(TE = 0, w = 0.7)
  p_eff_unlist = unlist(p_eff[[1]])
  check = ifelse(length(p_eff_unlist) < 2,
                 p_eff_unlist[length(p_eff_unlist)] > 0.995,
                 p_eff_unlist[length(p_eff_unlist)] > 0.986)
  fp = c(fp, check)
}
head(fp)
```

```
## [1] FALSE FALSE FALSE FALSE FALSE FALSE
```

```
mean(fp, na.rm = T)
```

```
## [1] 0.0028
```

```
sum(is.na(fp))
```

```
## [1] 0
```

3.2 TPR

$w = 0.7$: power is not high enough

```
options(warn = -1)
tp = NULL
for (m in 1:10000) {
  p_eff = trial2(TE = 0.7, w = 0.7)
  p_eff_unlist = unlist(p_eff[[1]])
  check = ifelse(length(p_eff_unlist) < 2,
                 p_eff_unlist[length(p_eff_unlist)] > 0.995,
                 p_eff_unlist[length(p_eff_unlist)] > 0.986)
  tp = c(tp, check)
}
mean(tp, na.rm = T)
```

```
## [1] 0.7931
```

```
sum(is.na(tp))
```

```
## [1] 0
```

$w = 0.5$: power is 1

```

options(warn = -1)
tp = NULL
for (m in 1:10000) {
  p_eff = trial2(TE = 0.7, w = 0.5)
  p_eff_unlist = unlist(p_eff[[1]])
  check = ifelse(length(p_eff_unlist) < 2,
                  p_eff_unlist[length(p_eff_unlist)] > 0.995,
                  p_eff_unlist[length(p_eff_unlist)] > 0.986)
  tp = c(tp, check)
}
mean(tp, na.rm = T)

```

```
## [1] 0.9976
```

```
sum(is.na(tp))
```

```
## [1] 0
```