

Package ‘Sample.Size’

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Type Package

Title Sample size calculation

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Description Computes the required sample size using the optimal designs with multiple constraints proposed in Mayo et al.(2010). This optimal method is designed for two-arm, randomized phase II clinical trials, and the required sample size can be optimized either using fixed or flexible randomization allocation ratios.

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SampleSize_OptimalDesign-package
Sample size calculation

Description

Computes the required sample size using the optimal designs with multiple constraints proposed in Mayo et al.(2010). This optimal method is designed for two-arm, randomized phase II clinical trials, and the required sample size can be optimized either using fixed or flexible randomization allocation ratios.

Details

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 Type: Package
 Version: 1.0
 Date: 2013-12-03
 License: GPL-2

Author(s)

Wei Jiang, Jonathan Mahnken, Matthew Mayo Maintainer: Wei Jiang<wjiang@kumc.edu>

Examples

```

Sample.Size(0.3, 0.6, 0.15, 0.15, 0.15, Allratio_c = 1, Allratio_e = 3)
Sample.Size(0.3, 0.6, 0.15, 0.15, 0.15)

```

Sample.Size

Sample size calculation

Description

Computes the required sample size using the optimal designs with multiple constraints proposed in Mayo et al.(2010). This optimal method is designed for two-arm, randomized phase II clinical trials, and the required sample size can be optimized either using fixed or flexible randomization allocation ratios.

Usage

```
Sample.Size(pi_c, pi_e, gamma_c, gamma_e, gamma_delta, Allratio_c = NA, Allratio_e = NA)
```

Arguments

pi_c	Response rate for control
pi_e	Response rate for experiment
gamma_c	Upper bound for gammaC
gamma_e	Upper bound for gammaE
gamma_delta	Upper bound for gammaDelta
Allratio_c	Allocation Design
Allratio_e	Allocation Design

Value

Prints required sample sizes, 1 to 1 allocation design and 1 to 3 allocation design

Author(s)

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Examples

```

Sample.Size(0.3, 0.6, 0.15, 0.15, 0.15, Allratio_c = 1, Allratio_e = 3)
Sample.Size(0.3, 0.6, 0.15, 0.15, 0.15)
## The function is currently defined as
function(pi_c, pi_e, gamma_c, gamma_e, gamma_delta, Allratio_c = NA,
        Allratio_e = NA)
{
  c1 <- 1
  e1 <- 1
  dmax1 <- ceiling(max(((pi_c * (1 - pi_c))/(c1 * gamma_c^2)),
    ((pi_e * (1 - pi_e))/(e1 * gamma_e^2)), ((pi_c * (1 -
      pi_c) + (c1/e1) * pi_e * (1 - pi_e))/(c1 * gamma_delta^2))))
  nc_fixed1 <- c1 * dmax1
  ne_fixed1 <- e1 * dmax1
  n_fixed1 <- nc_fixed1 + ne_fixed1
  nc_ast <- (pi_c * (1 - pi_c) + sqrt(pi_c * (1 - pi_c) * pi_e *
    (1 - pi_e)))/gamma_delta^2
  ne_ast <- pi_e * (1 - pi_e) * (gamma_delta^2 - pi_c * (1 -
    pi_c) * nc_ast^(-1))^(-1)
  nc_prime <- pi_c * (1 - pi_c)/gamma_c^2
  fnc_prime <- pi_e * (1 - pi_e) * (gamma_delta^2 - pi_c *
    (1 - pi_c) * nc_prime^(-1))^(-1)
  ne_prime <- pi_e * (1 - pi_e)/gamma_e^2
  fne_prime <- pi_c * (1 - pi_c) * (gamma_delta^2 - pi_e *
    (1 - pi_e) * ne_prime^(-1))^(-1)
  if (nc_ast > nc_prime & ne_ast > ne_prime) {
    c1 <- ceiling(nc_ast)
    e1 <- floor(ne_ast)
    c2 <- floor(nc_ast)
    e2 <- ceiling(ne_ast)
    dmax1 <- max(((pi_c * (1 - pi_c))/(c1 * gamma_c^2)),
      ((pi_e * (1 - pi_e))/(e1 * gamma_e^2)), ((pi_c *
        (1 - pi_c) + (c1/e1) * pi_e * (1 - pi_e))/(c1 *
        gamma_delta^2)))
    dmax2 <- max(((pi_c * (1 - pi_c))/(c2 * gamma_c^2)),
      ((pi_e * (1 - pi_e))/(e2 * gamma_e^2)), ((pi_c *
        (1 - pi_c) + (c2/e2) * pi_e * (1 - pi_e))/(c2 *
        gamma_delta^2)))
    if (dmax1 <= 1) {
      nc_flex <- ceiling(nc_ast)
      ne_flex <- floor(ne_ast)
      n_flex <- nc_flex + ne_flex
      sample_flex <- data.frame(nc_flex, ne_flex, n_flex)
      sample_flex <- data.frame(nc_flex, ne_flex, n_flex)
    }
  }
}

```

```

}
else if (dmax2 <= 1) {
  nc_flex <- floor(nc_ast)
  ne_flex <- ceiling(ne_ast)
  n_flex <- nc_flex + ne_flex
  sample_flex <- data.frame(nc_flex, ne_flex, n_flex)
  sample_flex <- data.frame(nc_flex, ne_flex, n_flex)
}
else {
  nc_flex <- ceiling(nc_ast)
  ne_flex <- ceiling(ne_ast)
  n_flex <- nc_flex + ne_flex
  sample_flex <- data.frame(nc_flex, ne_flex, n_flex)
}
}
else if (nc_ast <= nc_prime & ne_ast > ne_prime) {
  c1 <- ceiling(nc_prime)
  e1 <- floor(max(fnc_prime, ne_prime))
  dmax1 <- max(((pi_c * (1 - pi_c))/(c1 * gamma_c^2)),
              ((pi_e * (1 - pi_e))/(e1 * gamma_e^2)), ((pi_c *
              (1 - pi_c) + (c1/e1) * pi_e * (1 - pi_e))/(c1 *
              gamma_delta^2)))
  if (dmax1 <= 1) {
    nc_flex <- ceiling(nc_prime)
    ne_flex <- floor(max(fnc_prime, ne_prime))
    n_flex <- nc_flex + ne_flex
    sample_flex <- data.frame(nc_flex, ne_flex, n_flex)
  }
  else {
    nc_flex <- ceiling(nc_prime)
    ne_flex <- ceiling(max(fnc_prime, ne_prime))
    n_flex <- nc_flex + ne_flex
    sample_flex <- data.frame(nc_flex, ne_flex, n_flex)
  }
}
else if (nc_ast > nc_prime & ne_ast <= ne_prime) {
  c1 <- floor(max(fne_prime, nc_prime))
  e1 <- ceiling(ne_prime)
  dmax1 <- max(((pi_c * (1 - pi_c))/(c1 * gamma_c^2)),
              ((pi_e * (1 - pi_e))/(e1 * gamma_e^2)), ((pi_c *
              (1 - pi_c) + (c1/e1) * pi_e * (1 - pi_e))/(c1 *
              gamma_delta^2)))
  if (dmax1 <= 1) {
    nc_flex <- floor(max(fne_prime, nc_prime))
    ne_flex <- ceiling(ne_prime)
    n_flex <- nc_flex + ne_flex
    sample_flex <- data.frame(nc_flex, ne_flex, n_flex)
  }
  else {
    nc_flex <- ceiling(max(fne_prime, nc_prime))
    ne_flex <- ceiling(ne_prime)
    n_flex <- nc_flex + ne_flex
    sample_flex <- data.frame(nc_flex, ne_flex, n_flex)
  }
}

```

```

    }
  }
  else if (nc_ast <= nc_prime & ne_ast <= ne_prime) {
    nc_flex <- ceiling(nc_prime)
    ne_flex <- ceiling(ne_prime)
    n_flex <- nc_flex + ne_flex
    sample_flex <- data.frame(nc_flex, ne_flex, n_flex)
  }
  cat("Specified values for parameters:", "\n")
  cat("Response rates:", "\n", paste("control =", pi_c, "experiment =",
    pi_e), "\n")
  cat("Upper bounds for constraints:", "\n", "gammaC =", gamma_c,
    "gammaE =", gamma_e, "gammaDelta =", gamma_delta, "\n",
    "\n", "\n")
  cat("Required sample sizes:", "\n")
  cat("[1] Optimal Design:", "\n", paste("nc =", nc_flex, "ne =",
    ne_flex, "n =", n_flex), "\n")
  cat("[2] 1 to 1 Allocation Design:", "\n", paste("nc =",
    nc_fixed1, "ne =", ne_fixed1, "n =", n_fixed1), "\n")
  if (!is.na(Allratio_c) | !is.na(Allratio_e)) {
    c <- Allratio_c
    e <- Allratio_e
    dmax <- ceiling(max(((pi_c * (1 - pi_c))/(c * gamma_c^2)),
      ((pi_e * (1 - pi_e))/(e * gamma_e^2)), ((pi_c * (1 -
        pi_c) + (c/e) * pi_e * (1 - pi_e))/(c * gamma_delta^2))))
    nc_fixed <- c * dmax
    ne_fixed <- e * dmax
    n_fixed <- nc_fixed + ne_fixed
    sample_fixed2 <- data.frame(nc_fixed, ne_fixed, n_fixed)
    cat(paste("[3]", c, "to", e, "Allocation Design:"), "\n",
      paste("nc =", nc_fixed, "ne =", ne_fixed, "n =",
        n_fixed), "\n")
  }
}

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