library(powerly)

# Run the method.

results <- powerly(

range\_lower = 150,

range\_upper = 800,

samples = 30,

replications = 50,

measure = "sen",

measure\_value = 0.6,

statistic = "power",

statistic\_value = 0.8,

model = "ggm",

nodes = 10,

density = 0.4)

summary(results)

plot(results, step = 1)

plot(results, step = 2)

plot(results, step = 3)