# Marketing Models - Assignment 1

E.C. van Groningen

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## Load libraries

#### Load the data

```
data <- load("Assignment_1.RData")
```

# Assignment 1

Inspect the data

```
head(Conjoint_data)
```

```
price choice
##
     respondent_id profile
                                   form
                                          noapply disinfect bio
                         1 Concentrate 200 times
## 1
                 1
                                                        Yes No 35 cents
## 2
                 1
                         2
                                Powder 200 times
                                                        Yes No 35 cents
                                                                               0
## 3
                 1
                         3
                                Premix 100 times
                                                        Yes Yes 49 cents
                                                                               1
## 4
                         4
                                Powder 200 times
                                                        Yes Yes 49 cents
                                                                               0
                 1
                         5
## 5
                 1
                                Powder 50 times
                                                        Yes No 79 cents
                                                                               0
## 6
                         6 Concentrate 200 times
                                                         No Yes 79 cents
                 1
                                                                               0
```

#### Estimate the model

```
# Set baseline for variables in regression
Conjoint_data$form <- relevel(Conjoint_data$form, ref = "Powder")
Conjoint_data$noapply <- relevel(Conjoint_data$noapply, ref = "200 times")
Conjoint_data$disinfect <- relevel(Conjoint_data$disinfect, ref = "No")
Conjoint_data$price <- relevel(Conjoint_data$price, ref = "35 cents")

# Estimate model
model <-
    glm(choice ~ form + noapply + disinfect + bio + price,
        family = "binomial",
        data = Conjoint_data)

# Show results
results <- summary(model)
results

## ## Call:
## glm(formula = choice ~ form + noapply + disinfect + bio + price,</pre>
```

```
## glm(formula = choice ~ form + noapply + disinfect + bio + price,
      family = "binomial", data = Conjoint_data)
##
## Deviance Residuals:
##
      Min
                1Q
                     Median
                                  3Q
                                          Max
## -2.1680 -0.7913
                     0.3276
                                       2.1374
                              0.8173
##
## Coefficients:
                   Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                     1.1357
                                0.4662 2.436 0.014859 *
## formConcentrate
                     0.6482
                                0.3293 1.968 0.049035 *
## formPremix
                    -0.2519
                                0.3347 -0.753 0.451638
## noapply100 times -0.2335
                                0.3457 -0.675 0.499416
## noapply50 times
                    -0.5904
                                0.3601 -1.639 0.101111
## disinfectYes
                                0.2998 3.717 0.000202 ***
                    1.1143
## bioYes
                    0.2762
                                0.2862
                                        0.965 0.334477
## price49 cents
                    -1.3828
                                0.3609 -3.831 0.000128 ***
## price79 cents
                    -3.0606
                                0.3810 -8.034 9.46e-16 ***
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
```

```
##
## Logistic Regression Model
Dependent variable:
##
               -----
##
                       choice
## Constant
                      1.136**
                      (0.466)
## formConcentrate
                     0.648**
                       (0.329)
## formPremix
                      -0.252
                      (0.335)
                       -0.234
## noapply100 times
                      (0.346)
## noapply50 times
                      -0.590
                      (0.360)
## disinfectYes
                     1.114***
                      (0.300)
## bioYes
                       0.276
                      (0.286)
                     -1.383***
## price49 cents
                      (0.361)
                     -3.061***
## price79 cents
                     (0.381)
## Observations
                        330
## Log Likelihood
                     -174.477
## Akaike Inf. Crit.
                     366.955
## Note:
               *p<0.1; **p<0.05; ***p<0.01
```

#### Create empty part-worth

```
# Set levels part-worth
names(partworth$form) <- levels(Conjoint_data$form)</pre>
names(partworth$noapply) <- levels(Conjoint_data$noapply)</pre>
names(partworth$disinfect) <- levels(Conjoint data$disinfect)</pre>
names(partworth$bio) <- levels(Conjoint_data$bio)</pre>
names(partworth$price) <- levels(Conjoint_data$price)</pre>
# Set insignificant coefficients to zero
coeffs <- results$coefficients[,1] * (results$coefficients[,4] < .05)</pre>
coeffs[1]
## (Intercept)
##
      1.135682
# Fill the part-worth table
partworth$form[2:3] <- coeffs[2:3]</pre>
partworth$noapply[2:3] <- coeffs[4:5]</pre>
partworth$disinfect[2] <- coeffs[6]</pre>
partworth$bio[2] <- coeffs[7]</pre>
partworth$price[2:3] <- coeffs[8:9]</pre>
# Show results
partworth
## $form
##
        Powder Concentrate
                                  Premix
                               0.0000000
##
     0.0000000 0.6481925
##
## $noapply
## 200 times 100 times 50 times
##
           0
                      0
                                 Λ
##
## $disinfect
        No
                  Yes
## 0.000000 1.114257
##
## $bio
## No Yes
##
    0
##
## $price
## 35 cents 49 cents 79 cents
## 0.000000 -1.382753 -3.060619
Compute probability
```

```
max(partworth$disinfect) + max(partworth$bio) +
    max(partworth$price)))
prob
```

## [1] 0.947754

## Question 2

#### Question 2a

```
lik_bgnbd <- function(para) {</pre>
  # This is the likelihood function with CLV_data
  \# Change CLV_data to the names your datasets; the variables should be the same.
  # unpack 4 parameters:
  # r and alpha are the parameters of the gamma distribution of lambda
  # a and b are the parameters of the beta distribution ofr the "death" rate
  r <- para[1]
  alpha <- para[2]</pre>
  a <- para[3]
  b <- para[4]
  # unpack three variables:
  # x - # of transactions (frequency)
  \# t_x - the most recent time of transaction (recency)
  \# T - the end time for the Poisson process
  x <- CLV_data$x
  t_x <- CLV_data$t_x</pre>
  T <- CLV_data$T
  # A1 - A4 corresponds to the log of 4 terms in the likelihood function
  # See p.280 of Fader et al. (2005)
  # They are the log of A1-A4 for the log-likelihood
  A1 \leftarrow lgamma(r+x) + r*log(alpha) - lgamma(r)
  A2 \leftarrow lgamma(a+b) + lgamma(b+x) - lgamma(b) - lgamma(a+b+x)
  A3 <- -(r+x)*log(alpha+T)
  A4 \leftarrow rep(0,length(x))
  idx <- x>0
  if (sum(idx)>0) {
    A4[idx] \leftarrow \log(a/(b+x[idx]-1)) - (r+x[idx])*\log(alpha+t_x[idx])
  }
  lh \leftarrow sum(A1+A2+log(exp(A3)+idx*exp(A4)))
  # To return the minus of likelihood for minimization
  # maximize likelihood = minimize negative likelihood
  return(-lh)
}
```

```
pred_Y <- function(para, H, t) {</pre>
  # Given estimated parameters, for a customer with history \{x,t_{\perp}x,T\}, to
  # predict her # of transactions in t time.
  # Please see Equation (10) of Fader et al. (2005), p.279;
  # This function uses a package "hypergeo" for function evaluation;
  # As evaluation of 2F1(.) is somewhat costly, this function is to accommodate
  # only 1 customer;
  # You can extend the function to calculate for many customers by iterations or
  # use lapply(.);
# The function takes three input:
  # para - a vector of the 4 parameters;
  # H - a data.frame with only 1 row (for 1 customer) and three variables
  \# \{x, t_x, T\};
  # t - how long into the future?
  # unpack 4 parameters:
  \# r and alpha are the parameters of the gamma distribution of lambda
  # a and b are the parameters of the beta distribution of the "death" rate
  r <- para[1]
  alpha <- para[2]</pre>
  a <- para[3]
  b <- para[4]
  # unpack three variables:
  \# x - \# of transactions (frequency)
  # t \times x - the most recent time of transaction (recency)
  \# T - the end time for the Poisson process
  x <- H$x
  t_x <- H$t_x
  T <- H$T
  # the term of the Gaussian hypergeometric function 2F1(.)
  hg <- hypergeo::hypergeo(r+x,b+x,a+b+x-1,t/(alpha+T+t))
  hg <- Re(hg) # hg is a real number in the complex format Re(.) makes it real.
  # the numerator term, given hg
  Y1 \leftarrow (a+b+x-1)/(a-1)*(1-(((alpha+T)/(alpha+T+t))^(r+x))*hg)
  Y2 <- 1
  if (x>0) {
    Y2 \leftarrow Y2 + a/(b+x-1)*(((alpha+T)/(alpha+t_x))^(r+x))
 return(Y1/Y2)
}
```

#### Estimating the Parameters of the BG/NBD model

```
# to specify the constraint matrix
ui <- diag(4)
ci <- rep(0,4)</pre>
```

```
# para: the starting values; a vector with 4 elements corresponding to
# (r, alpha, a, b).
# you can change this to other values or test different values.
para <- rep(.01,4)
# to run the constrained optimization
results <- constrOptim(para, lik_bgnbd, NULL, ui, ci)
# to get the estimated parameters in the order of (r,alpha,a,b)
para <- results$par</pre>
names(para) <- c("r", "alpha", "a", "b")</pre>
stargazer(para, summary=FALSE, title="Parameters GB/NBD model", type = "text")
##
## Parameters GB/NBD model
## =========
       alpha a b
## -----
## 0.232 8.957 1.194 3.191
## -----
```

#### Predict the No. of Transactions

## 4 4 0.753 ## 5 5 0.089 ## -----