

SNP-wise General Weights2.0

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We first redefine the log10BF function (note that we rename log10BF to emphasize that it is indeed in base 10).

```
log10BF = function(g,y,sigmaa) {
  p=dim(g)[2]
  n=dim(g)[1]
  if (is.null(dim(g)[2])){
    g = g - mean(g)
    y = y - mean(y)
    n=length(g)
    X = g
    invnu = 1/sigmaa^2
    invOmega = invnu + t(X) %*% X
    B = (t(X) %*% cbind(y))/invOmega
    invOmega0 = n
    return(-0.5*log10(det(invOmega)) + 0.5*log10(invOmega0) - log10(sigmaa) - (n/2)*(log10
(t(y- X %*% B) %*% y) - log10(t(y) %*% y)))
  }
  else {
    g = scale(g, scale = FALSE)
    y = scale(y, scale = FALSE)
    X = g
    invnu = diag(rep(1/sigmaa^2, p))
    invOmega = invnu + t(X) %*% X
    B = solve(invOmega, t(X) %*% cbind(y))
    invOmega0 = n
    return(-0.5*log10(det(invOmega)) + 0.5*log10(invOmega0) - p*log10(sigmaa) - (n/2)*(l
og10(t(y- X %*% B) %*% y) - log10(t(y) %*% y - n*mean(y)^2)))
  }
}
```

We then redefine the snpwise_weights function to do several things: add a π parameter that allows us to make all priors smaller or larger based on previous knowledge (e.g., $\pi = 10^{-3}$ if we generally expect 1 in 1000 SNPs to be significant), redefine the "post" variable as "log10post", and include a normalizing constant in the output weights:

```

snpwise_weights = function(X,y,priorpi,sigmaa) {
  models <- c()
  log10post <- c()
  log10bf <- c()

  #get number of parameters
  if (is.null(dim(X)[2])) {
    par <- 1
  }
  else {
    par <- dim(X)[2]
  }

  if (missing(priorpi)){
    p <- par
  }
  else {
    p <- 1/priorpi
  }

  #get all possible parameter combinations
  l <- rep(list(0:1),par)
  combs <- expand.grid(l)
  m <- dim(combs)[1]

  for (i in c(1:m)){
    numlist <- c()
    incl <- c()
    numincl <- 0

    for (j in c(1:par)){
      if (combs[i,j]==1){
        incl <- cbind(incl, X[,j])
        numincl <- numincl + 1
      }
    }
    numlist <- c(numlist, numincl)

    #if only one variable in model
    if (numincl==1) {
      for (k in c(1:par)){
        if (combs[i,k]==1){
          varin <- k
        }
      }
      newmod <- paste0("X", varin)

      #get weight
      prior <- (1/p)*(1-1/p)^(par-1)
      lbf <- log10BF(X[,varin], y, sigmaa)
      log10bf <- c(log10bf, lbf)
      post <- log10(prior) + lbf
    }
  }
}

```

```

#if multiple variables in model
else if (numincl != 0){
  newmod <- ""
  for (j in c(1:par)){

    if (combs[i,j]==1){
      var <- paste0("X", j)
      newmod <- paste(newmod, sep =",", var)
    }
  }
  newmod <- sub('.', '', newmod)
  #get weight
  prior <- (1/p)^numincl * (1-1/p)^(numincl-1)
  lbf <- log10BF(incl, y, sigmaa)
  log10bf <- c(log10bf, lbf)
  post <- log10(prior) + lbf
}

#if null model
else {
  log10bf <- c(log10bf, 0)
  post <- log10((1-1/p)^(p))
  newmod <- "NULL"
}

models <- c(models, newmod)
#add weight to list
log10post <- c(log10post, post)
}

#find log10 normalizing constant
maxlog10post <- max(log10post)
log10norm <- maxlog10post + log10(sum(10^(log10post-maxlog10post)))
wts <- 10^((log10post)-log10norm)
log10weights <- log10(wts)

#return dataframe of each model and its posterior weight
toreturn <- data.frame(models, log10post, wts, log10norm, log10bf)
names(toreturn)[1] <- "models"
names(toreturn)[2] <- "log10postscores"
names(toreturn)[3] <- "weights"
names(toreturn)[4] <- "log10norm"
names(toreturn)[5] <- "log10bayesfactor"

return(toreturn)
}

```

We first test this function without inputting any prior π , and thus defaulting to $1/p$. We can also do a sanity to confirm that the weights sum to 1:

```
x1 <- rnorm(100)
x3 <- rnorm(100)
x2 <- x1 + x3 + rnorm(100, sd = 0.5)
y <- x1 + x3 + rnorm(100)
X <- cbind(x1, x2, x3)

test1 <- snpwise_weights(X, y, sigmaa = 0.5)
test1
```

##	models	log10postscores	weights	log10norm	log10bayesfactor
## 1	NULL	-0.5282738	6.049175e-26	24.69003	0.000000
## 2	X1	12.1398224	2.817036e-13	24.69003	12.969126
## 3	X2	20.7256458	1.085465e-04	24.69003	21.554950
## 4	X1,X2	22.0699362	2.398315e-03	24.69003	23.200270
## 5	X3	5.5119809	6.636679e-20	24.69003	6.341285
## 6	X1,X3	24.6366134	8.842668e-01	24.69003	25.766947
## 7	X2,X3	19.8770641	1.538275e-05	24.69003	21.007398
## 8	X1,X2,X3	23.7439183	1.132109e-01	24.69003	25.527465

```
sum(test1$weights)
```

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

We can next test the function with the same data as above, but setting $\pi = 10^{-3}$:

```
test2 <- snpwise_weights(X, y, priorpi = 10^(-3), sigmaa = 0.5)
test2
```

##	models	log10postscores	weights	log10norm	log10bayesfactor
## 1	NULL	-0.4345118	5.912632e-21	19.79371	0.000000
## 2	X1	9.9682571	1.494685e-10	19.79371	12.969126
## 3	X2	18.5540806	5.759346e-02	19.79371	21.554950
## 4	X1,X2	17.1998355	2.547582e-03	19.79371	23.200270
## 5	X3	3.3404156	3.521342e-17	19.79371	6.341285
## 6	X1,X3	19.7665126	9.393020e-01	19.79371	25.766947
## 7	X2,X3	15.0069633	1.634015e-05	19.79371	21.007398
## 8	X1,X2,X3	16.5265956	5.406152e-04	19.79371	25.527465

```
sum(test2$weights)
```

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

With this parameter set, we can see that the models with multiple variables are penalized much more than before. Where in the first example the largest weighted model was X1,X3, followed by X2, they are now reversed.