Stat 230: Linear Models Homework 4 Professor Ding Lev Golod

QUESTION 5, PART (1)

What happens if you don't use a log-transformation? Certain terms like n^n will go to infinity and mess up the calculation.

```
## A function to calculate log(f(k,n,p,phi)) for one value of k
logf \leftarrow function(k, n = 100, p = 0.3, phi = 0.5){
  terms = rep(NA, 7)
  terms[1] = lgamma(n+1)
                                                    # does not depend on k
  terms[2] = -lgamma(k+1)
  terms[3] = -lgamma(n-k+1)
  \# terms[4] = ((-phi+1) * k) * log(k)
  \# terms[5] = ((-phi+1) * (n-k)) * log(n-k)
  \# terms[6] = ((-phi+1) * n) * log(n)
                                                      # does not depend on k
  terms[4] = ((-phi+1) * k)
                                * log(k/n)
  terms[5] = ((-phi+1) * (n-k)) * log((n-k)/n)
  terms[6] = (phi * k)
                                  * log(p)
  terms[7] = (phi * (n-k))
                                  * log(1-p)
  terms <- ifelse(is.nan(terms),0, terms)</pre>
  sum_terms <- sum(terms)</pre>
 return(sum_terms)
}
## Find the denominator for n = 1000
denom <- sum(sapply(seq.int(0,1000), function(k) exp(logf(k,n=1000))))
# denom
## Find the denominator for n = 100, 500, 1000, 2000
ns \leftarrow c(100, 500, 1000, 2000)
names(ns) <- paste0('n=',ns)</pre>
denoms <- sapply(ns, function(n) sum(sapply(seq.int(0,n),</pre>
                                              function(k) exp(logf(k,n))))
print(denoms)
## n=100 n=500 n=1000 n=2000
## 1.4189 1.4151 1.4147 1.4144
## determine average runtime for each n value
reps=100
times <- sapply(ns, function(n)</pre>
  system.time( replicate(reps, sum(sapply(seq.int(0,n),
```

```
function(k) exp(logf(k,n))))
  )[3]
)
print(times)
## n=100.elapsed n=500.elapsed n=1000.elapsed n=2000.elapsed
##
            0.269
                           1.174
                                          2.442
QUESTION 5, PART (2)
## vectorized fn to find the denominator
vecdenom <- function(n, p = 0.3, phi = 0.5){
  ks <- seq.int(0,n)
  terms \leftarrow rep(lgamma(n+1), n+1) +
   -lgamma(ks+1) +
   -lgamma(n-ks+1) +
   ifelse(ks == 0 | ks == 100, 0, ((-phi+1) * ks ) * log(ks/n) ) +
   ifelse(ks == 0 | ks == 100, 0, ((-phi+1) * (n-ks)) * log((n-ks)/n)) +
   (phi * ks) * log(p) +
   (phi * (n-ks)) * log(1-p)
    # + phi * (k * log(p/(1-p)) + n * log(1-p))
  sum(exp(terms))
}
denoms2 <- sapply(ns, vecdenom)</pre>
print(denoms)
## n=100 n=500 n=1000 n=2000
## 1.4189 1.4151 1.4147 1.4144
times2 <- sapply(ns, function(n) system.time(replicate(reps, vecdenom(n)))[3])
print(times2)
## n=100.elapsed n=500.elapsed n=1000.elapsed n=2000.elapsed
##
            0.010
                           0.020
                                          0.033
                                                         0.063
QUESTION 5, PART (3)
cat(sprintf('How much faster is the vectorized version when n = 2000?
%s times faster', floor(times[4]/times2[4])))
## How much faster is the vectorized version when n = 2000?
## 75 times faster
```