```
In [1]: using DataFrames
In [2]: using Distributions
In [3]: using(Gadfly)
```

## 200 SNP (=10 chromosomes \* 20 SNP/chr)

```
SNP = convert(Array, readtable("/home/nicole/Jupyter/HERdata/data/Clean/newdata/ch1to10.200SNP.noID", separa
Out[4]: 1442x200 Array{Int64,2}:
                                                                      0
        corSNP = cor(SNP)
In [5]:
        nRows = size(corSNP, 1)
Out[5]: 200
In [6]: LDSNP = zeros(nRows-1,20);
```

```
In [7]: for i = 1:(nRows-20)
          LDSNP[i,:] = corSNP[i,(i+1):(i+20)].^2
      end
In [8]: y = mean(LDSNP, 1)
      sort(y,2)
Out[8]: 1x20 Array{Float64,2}:
       In [9]: plot(x=(1:20)/20*1,y=y)
Out[9]:
          0.3
          0.2
         У
          0.1
           0.0
            0.0
                                 0.5
                                                      1.0
                                  Χ
```

## 2,000 SNP (=10 chromosomes \* 200 SNP/chr)

```
In [10]: SNP = convert(Array, readtable("/home/nicole/Jupyter/HERdata/data/Clean/newdata/ch1to10.2000SNP.noID", separ
Out[10]: 1442x2000 Array{Int64,2}:
             1 0
                   1 1
                                              0
                                            1
                                               0
                                                                       0
                                            0
                                              1
                            0
                                            1
In [11]:
         corSNP = cor(SNP)
         nRows = size(corSNP,1)
Out[11]: 2000
In [12]:
         LDSNP
               = zeros(nRows-1,20);
In [13]: for i = 1:(nRows-20)
             LDSNP[i,:] = corsnP[i,(i+1):(i+20)].^2
         end
```

```
In [14]: y = mean(LDSNP,1)
    sort(y,2)
Out[14]: 1x20 Array{Float64,2}:
```

0.073607 0.077073 0.0812765 0.0817273 ... 0.171783 0.211088 0.300319

In [15]: plot(x=(1:20)/20\*1,y=y)

Out[15]:

