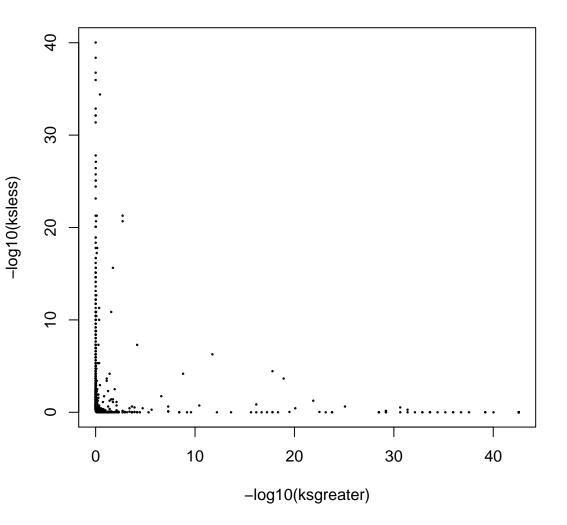
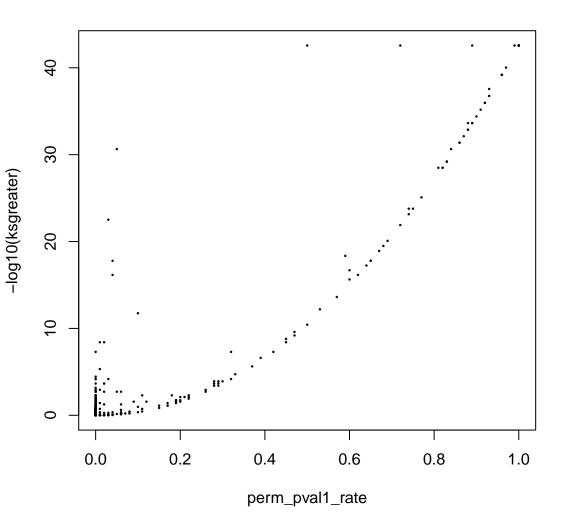
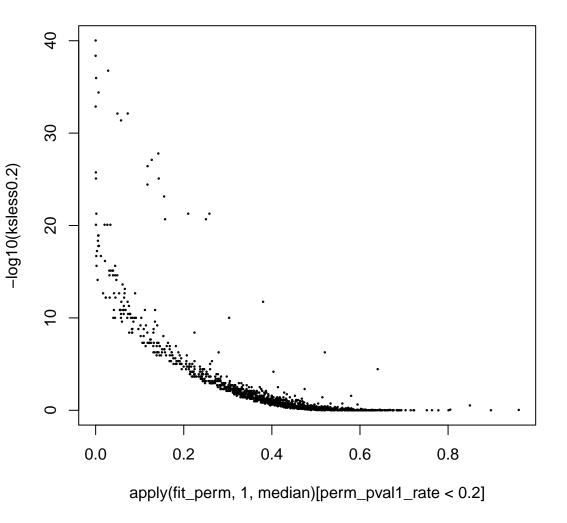


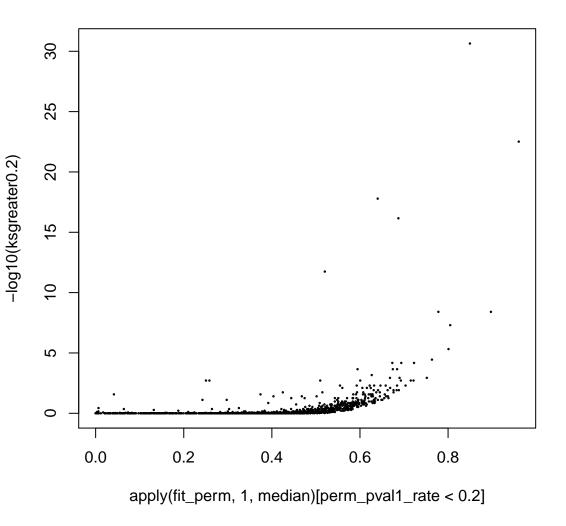
sig\_KSgreater: 57.933%, sig\_KSless: 12.833%



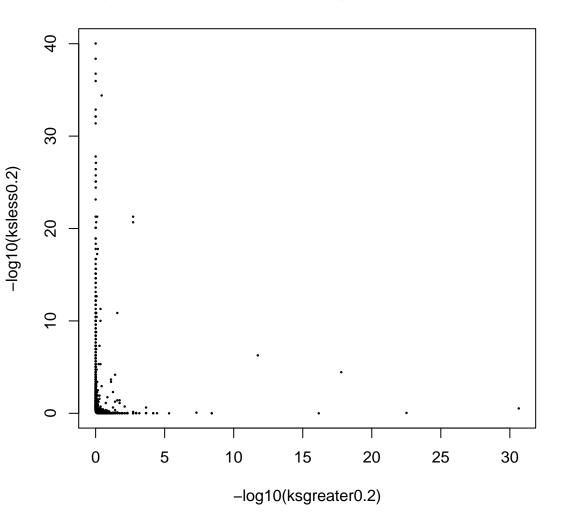
cor: 0.997



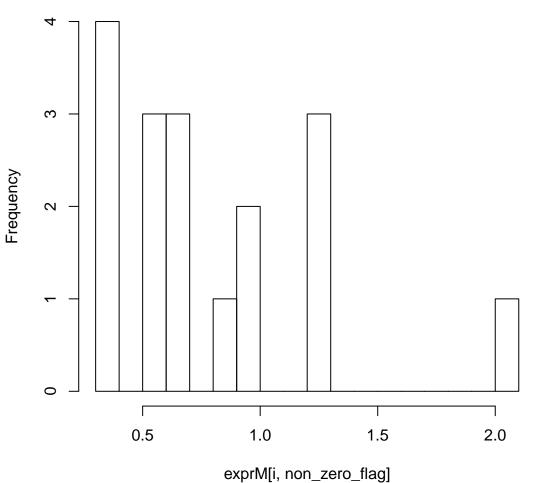




sig\_KSgreater0.2: 3.079%, sig\_KSless0.2: 29.253%

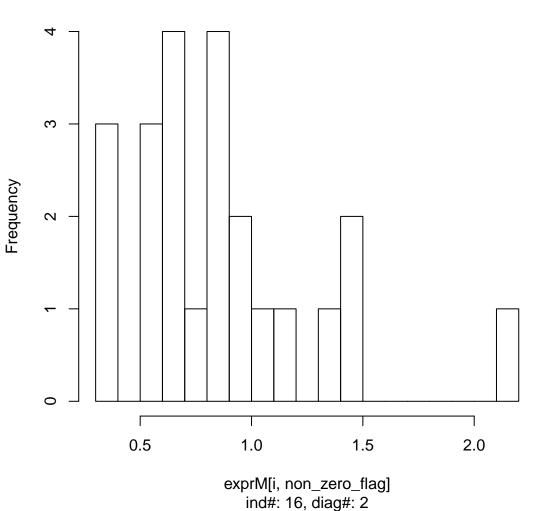


# log expression of gene#1454, pval ob=0.9007, non-zero num=1

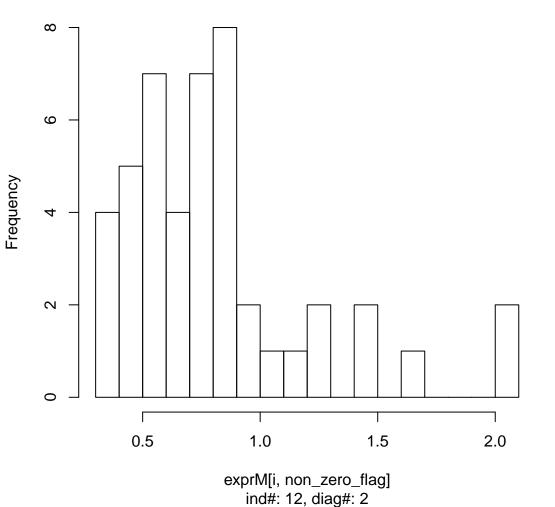


ind#: 12, diag#: 2

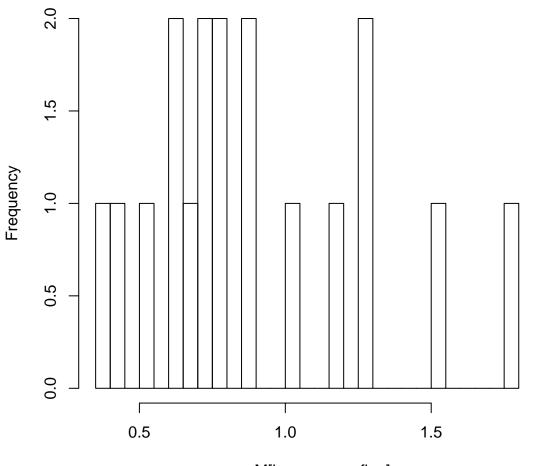
## log expression of gene#2709, pval ob=0.1548, non-zero num=2



### log expression of gene#561, pval ob=0.0824, non-zero num=4

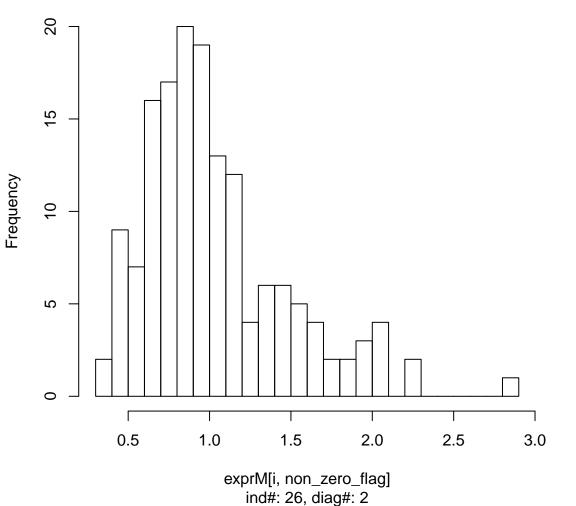


#### log expression of gene#2370, pval ob=0.042, non-zero num=1

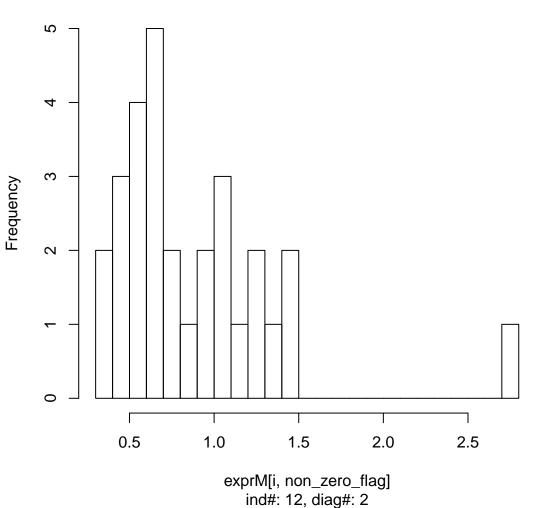


exprM[i, non\_zero\_flag] ind#: 12, diag#: 2

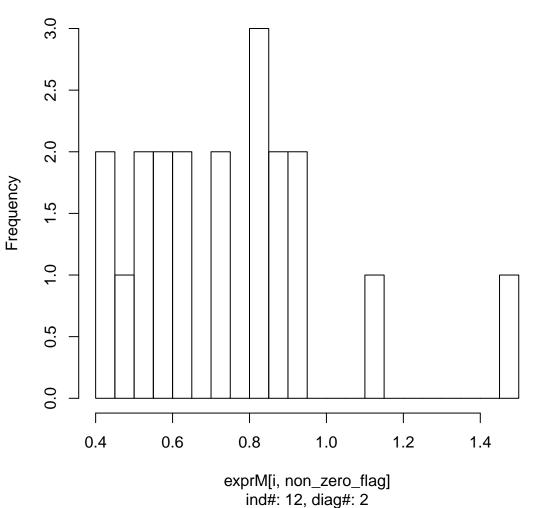
## log expression of gene#861, pval ob=0.2108, non-zero num=15



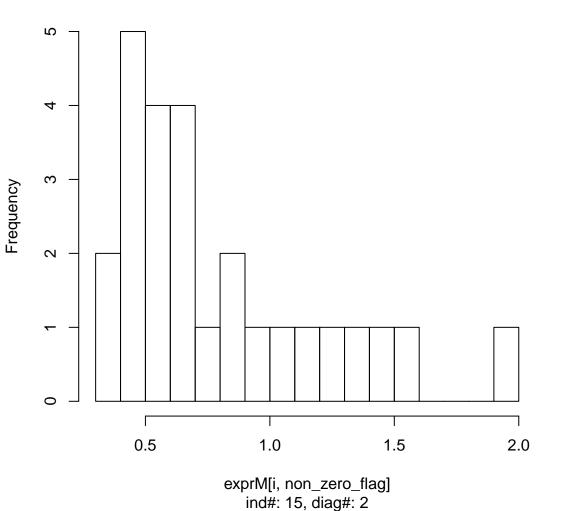
## log expression of gene#1347, pval ob=0.8418, non-zero num=2



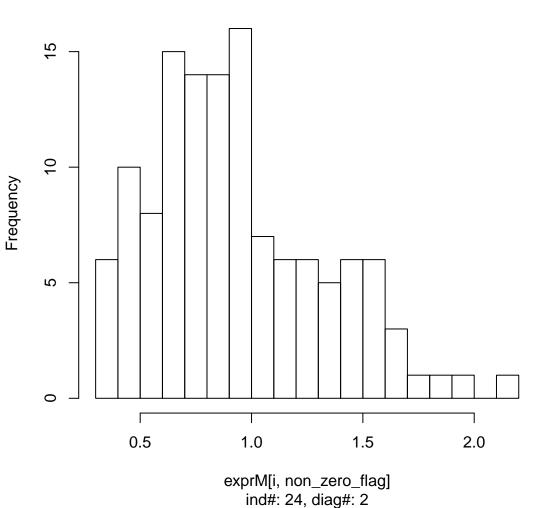
## log expression of gene#2565, pval ob=0.6857, non-zero num=2



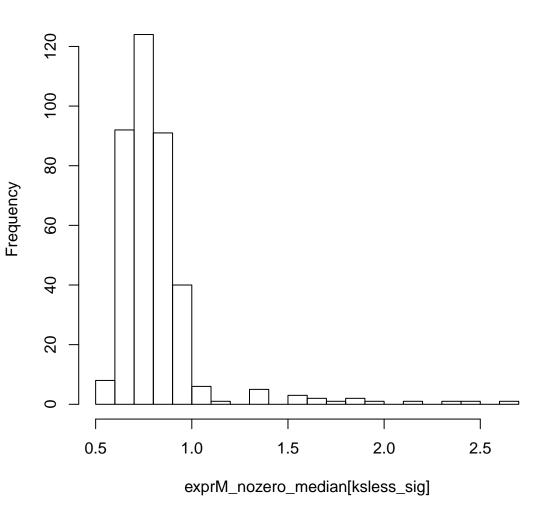
### log expression of gene#559, pval ob=0.4357, non-zero num=2



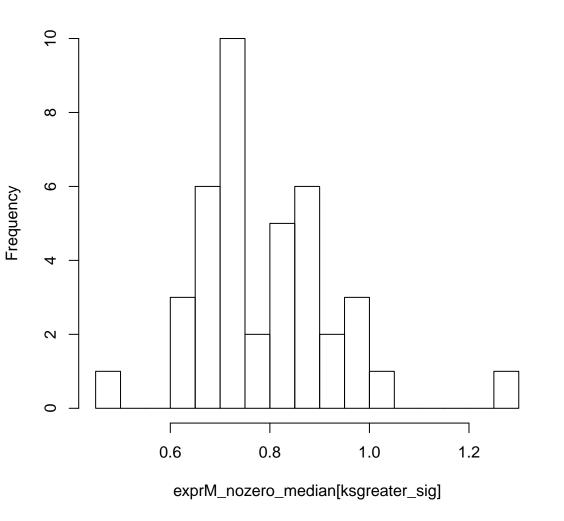
## log expression of gene#193, pval ob=0.2757, non-zero num=12



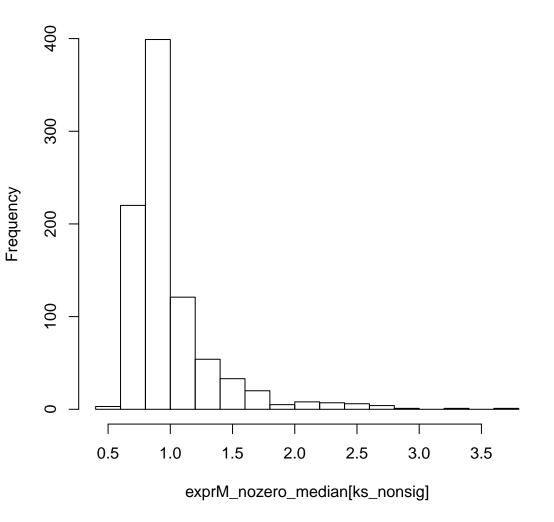
### median of nozero log-expres of genes, pval1\_rate<0.2,ksless s



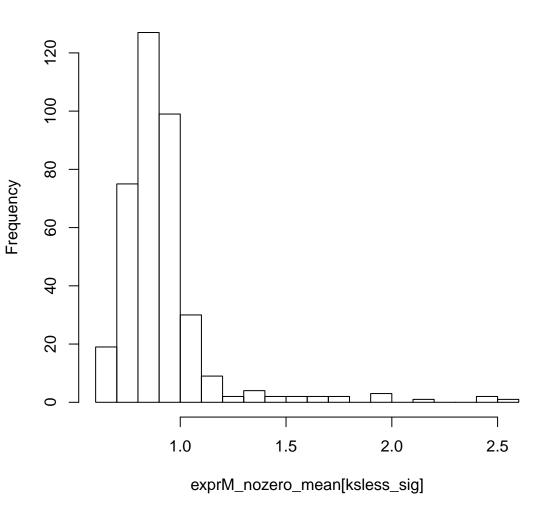
#### median of nozero log-expres of genes,pval1\_rate<0.2,ksgreater



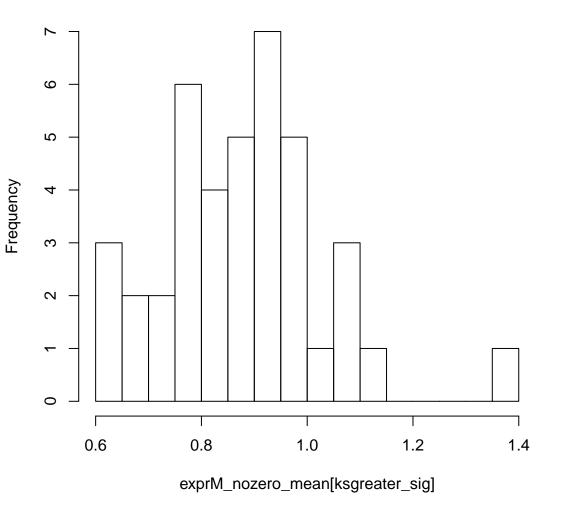
median of nozero log-expres of genes,pval1\_rate<0.2,ks no si



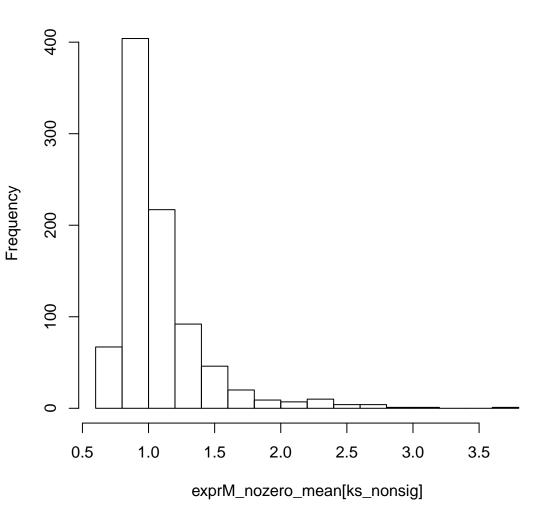
mean of nozero log-expres of genes, pval1\_rate<0.2,ksless si



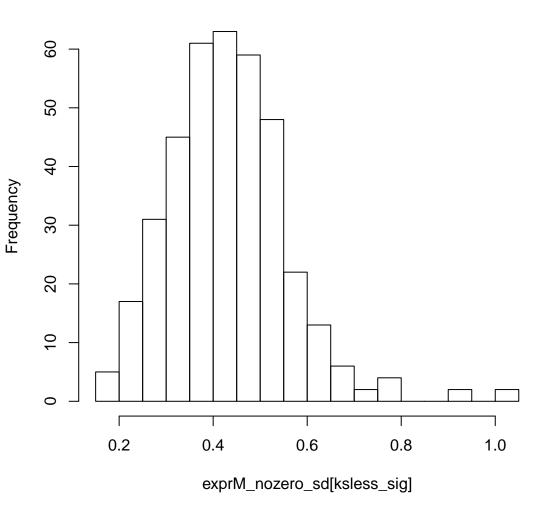
#### mean of nozero log-expres of genes,pval1\_rate<0.2,ksgreaters



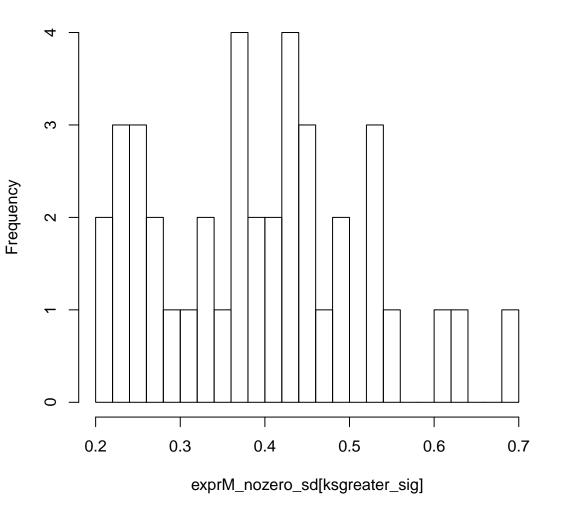
mean of nozero log-expres of genes,pval1\_rate<0.2,ks no sig



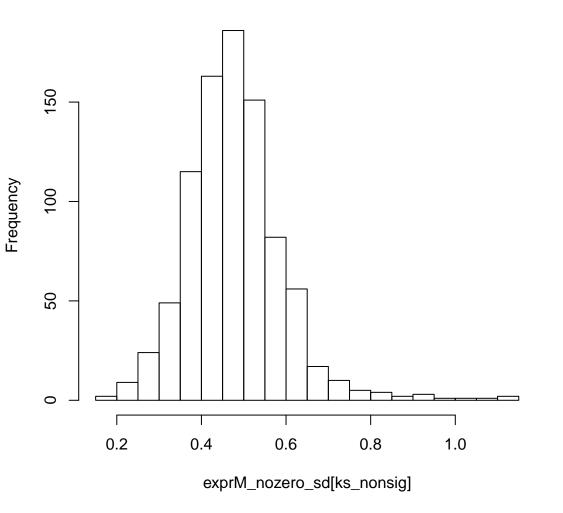
#### sd of nozero log-expres of genes, pval1\_rate<0.2,ksless sig



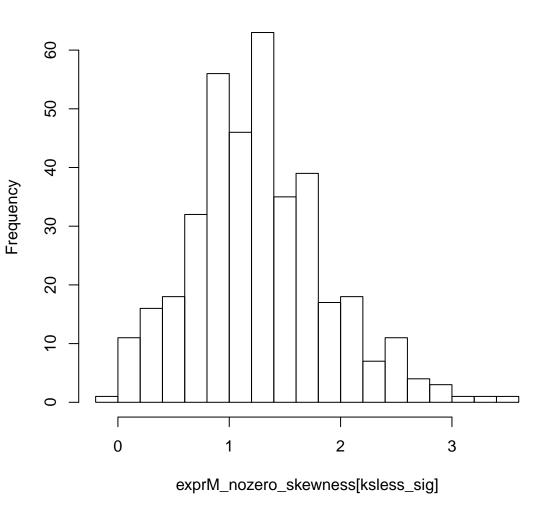
#### sd of nozero log-expres of genes,pval1\_rate<0.2,ksgreater signal



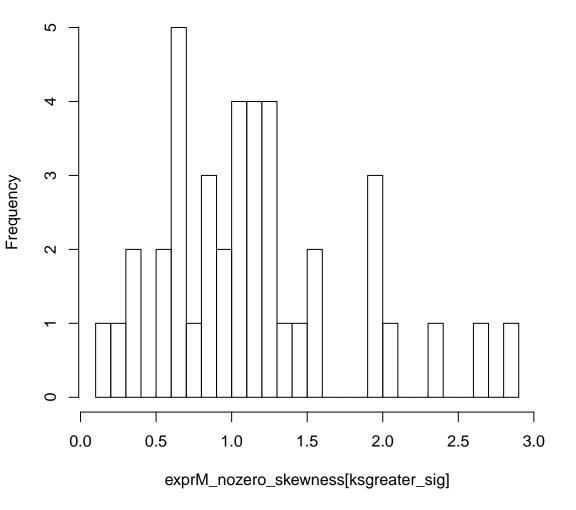
#### sd of nozero log-expres of genes,pval1\_rate<0.2,ks no sig



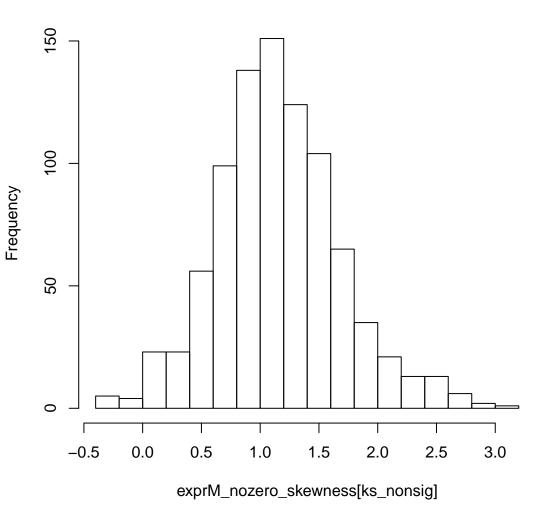
#### skewness of nozero log-expres of genes, pval1\_rate<0.2,ksless



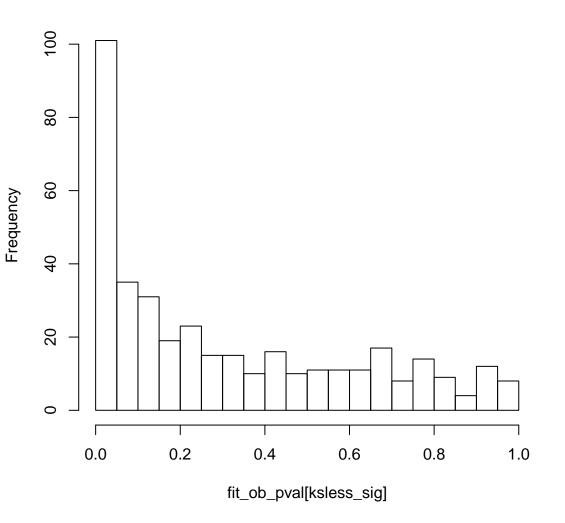
### skewness of nozero log-expres of genes,pval1\_rate<0.2,ksgreate



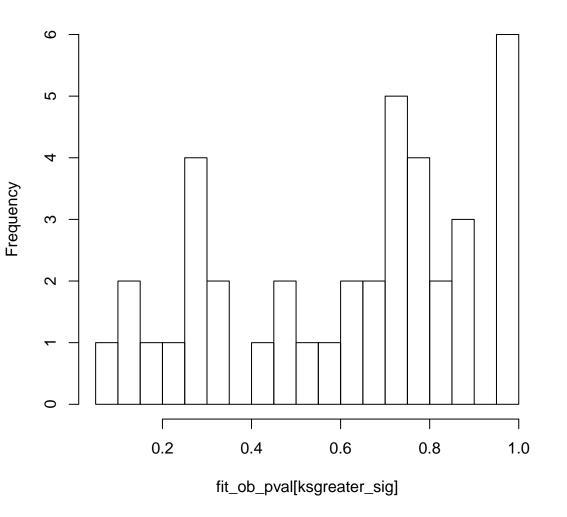
#### skewness of nozero log-expres of genes,pval1\_rate<0.2,ks no



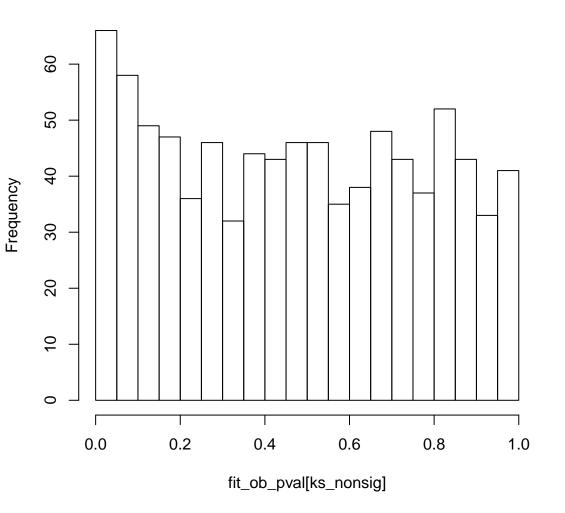
#### observed pvalues with pval1\_rate<0.2,ksless sig



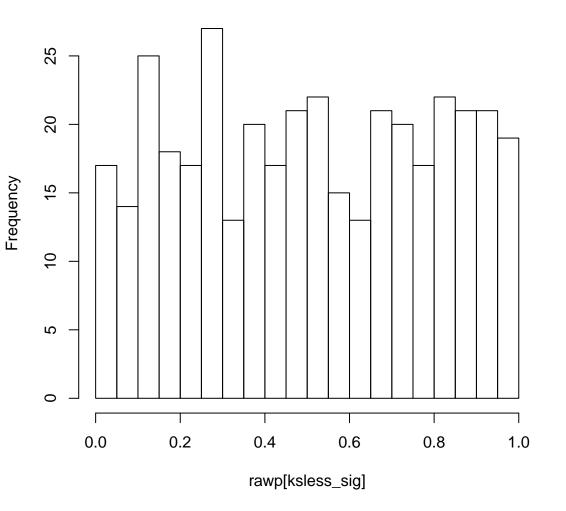
#### observed pvalues with pval1\_rate<0.2,ksgreater sig



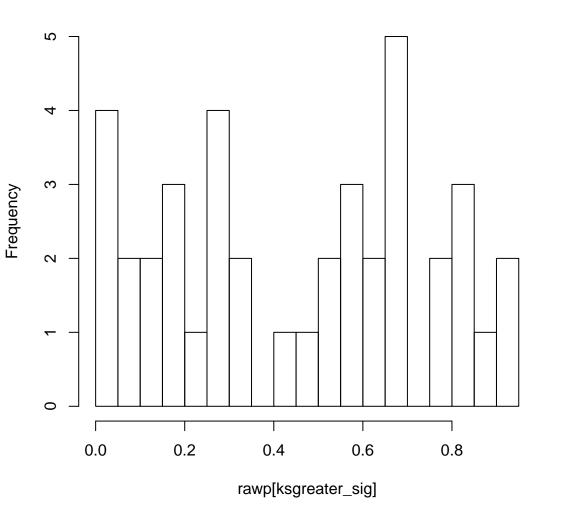
#### observed pvalues with pval1\_rate<0.2,ks no sig



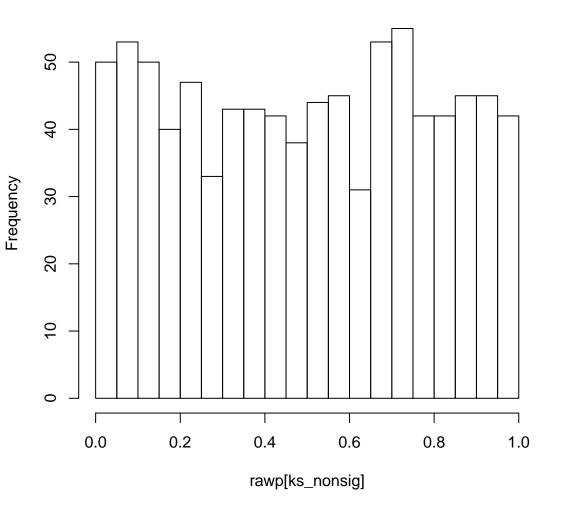
#### permutation pvalues with pval1\_rate<0.2,ksless sig



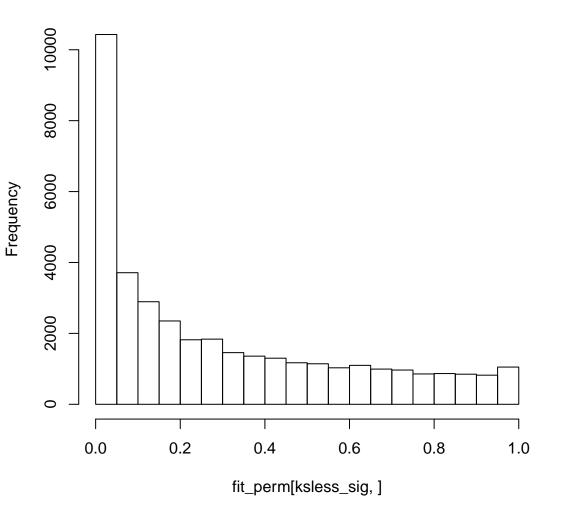
#### permutation pvalues with pval1\_rate<0.2,ksgreater sig



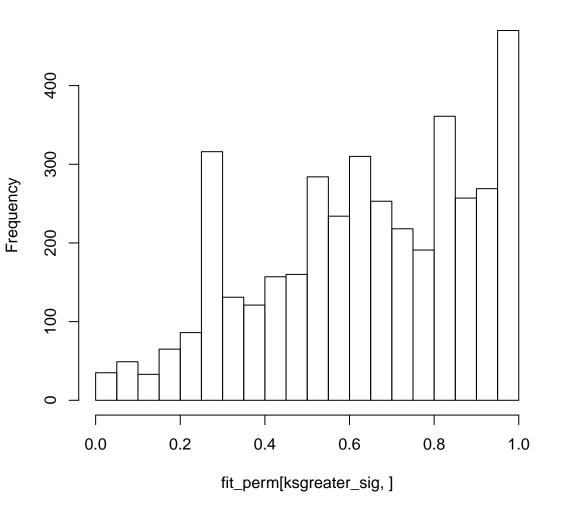
#### permutation pvalues with pval1\_rate<0.2,ks no sig



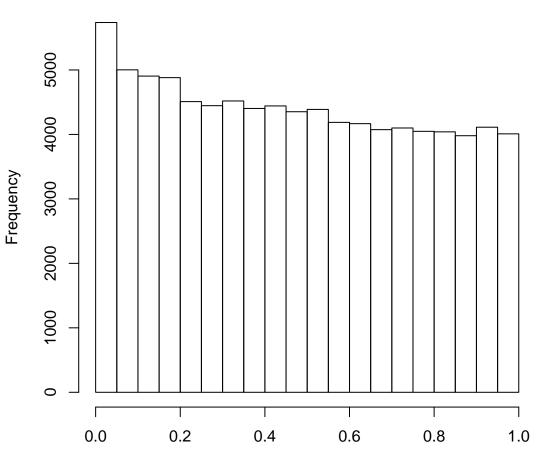
#### pvalues from permutation data with pval1\_rate<0.2,ksless sig



#### pvalues from permutation data with pval1\_rate<0.2,ksgreater s



#### pvalues from permutation data with pval1\_rate<0.2,ks no sig



 $fit\_perm[perm\_pval1\_rate < 0.2 \& ksgreater >= 0.01 \& ksless > 0.01, ]$