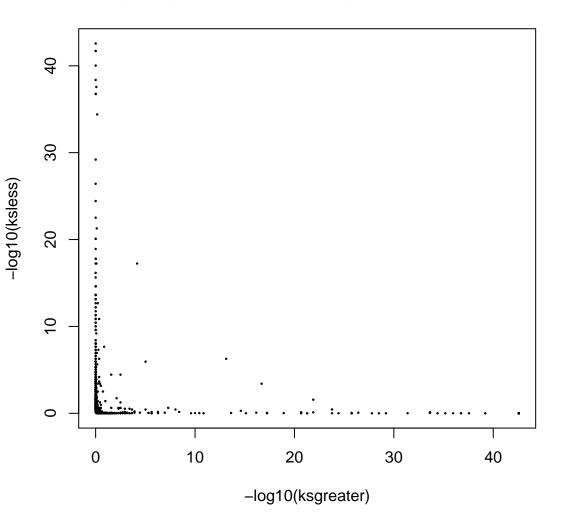
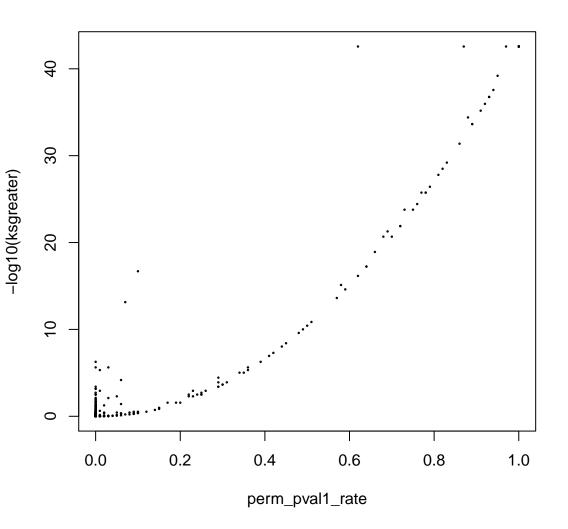
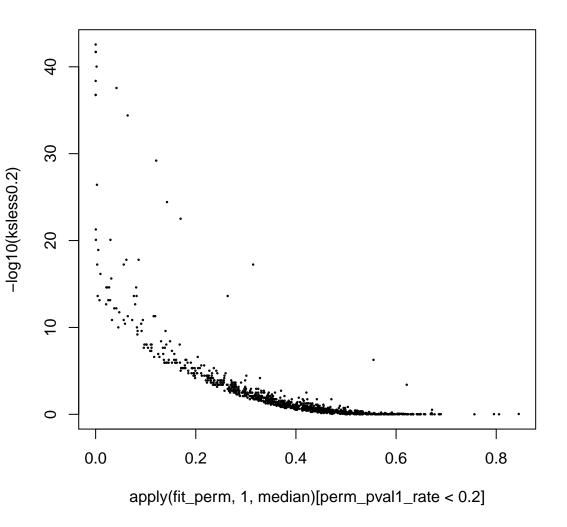


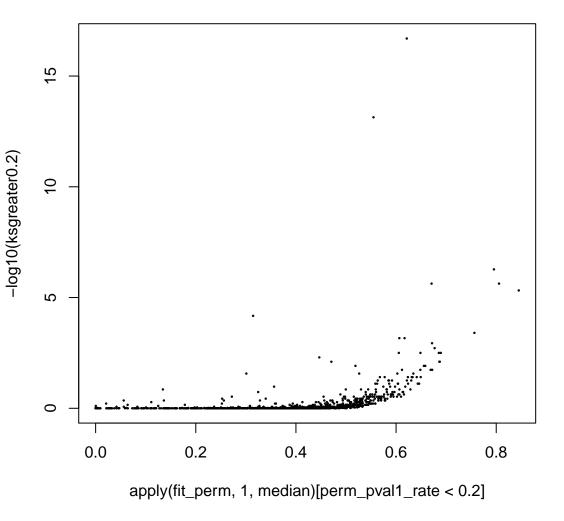
sig\_KSgreater: 66.167%, sig\_KSless: 10.467%



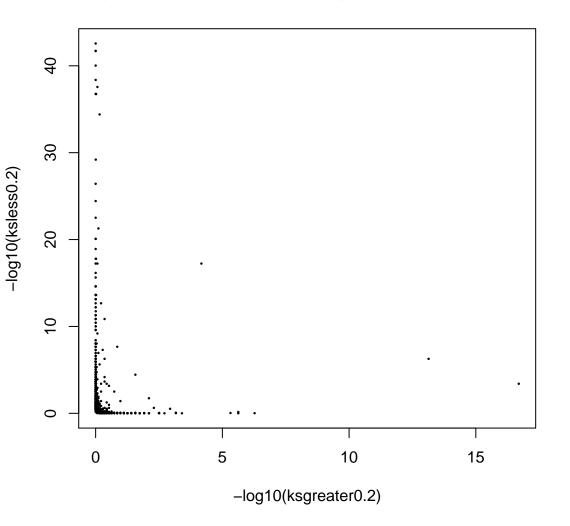
cor: 0.998



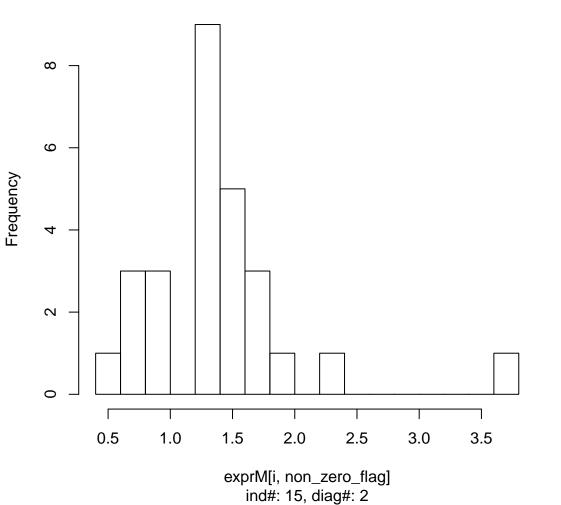




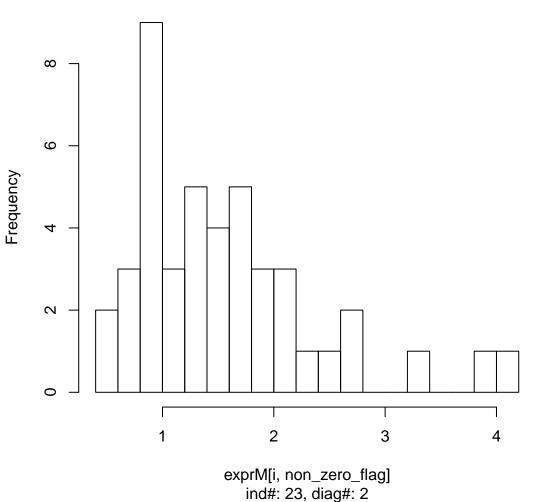
sig\_KSgreater0.2: 1.934%, sig\_KSless0.2: 30.174%



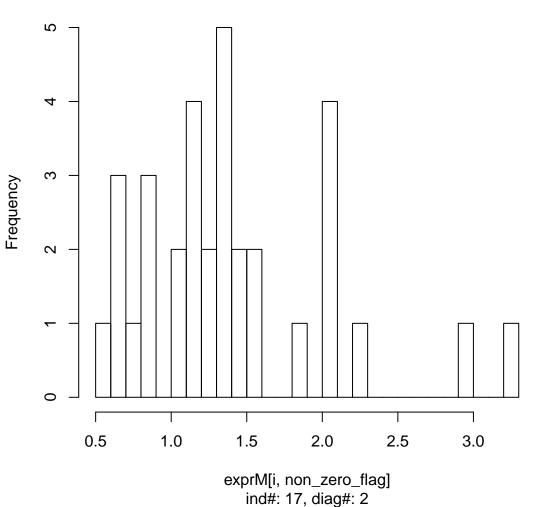
# log expression of gene#2013, pval ob=0.1826, non-zero num=2



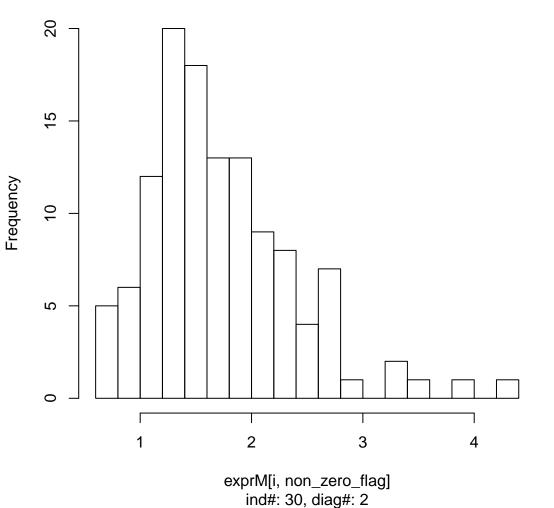
## log expression of gene#680, pval ob=0.0759, non-zero num=4



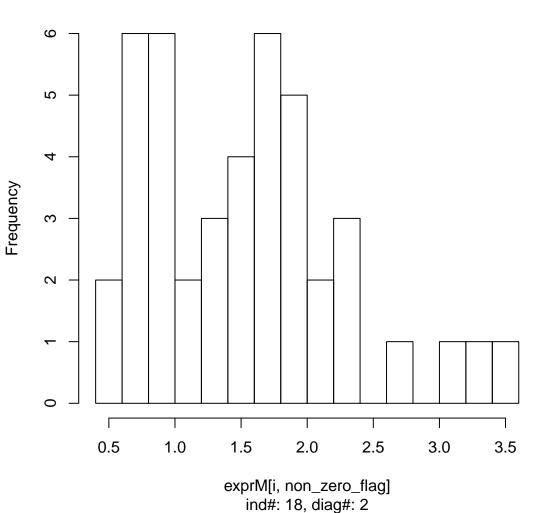
# log expression of gene#2273, pval ob=0.0342, non-zero num=3



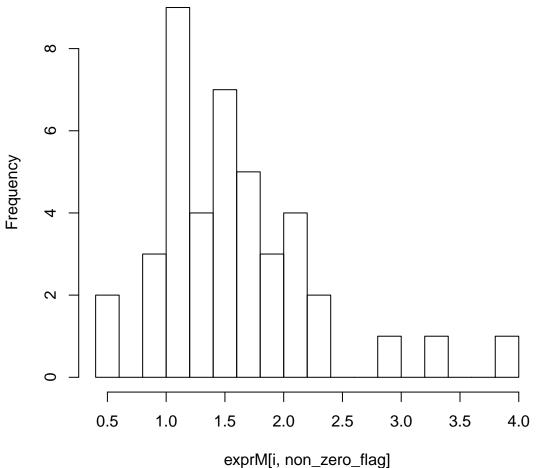
# log expression of gene#730, pval ob=0.5338, non-zero num=12



# log expression of gene#212, pval ob=0.9372, non-zero num=4

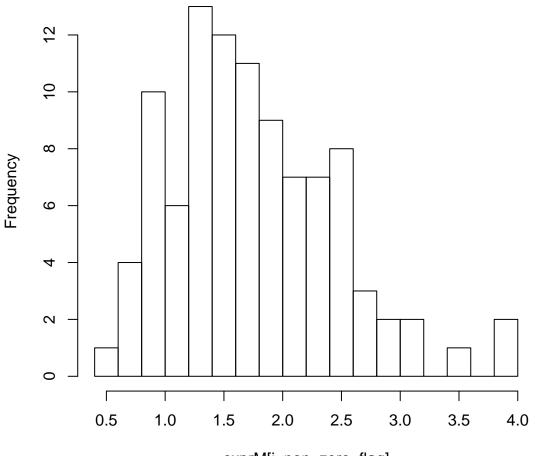


# log expression of gene#1047, pval ob=0.4215, non-zero num=4



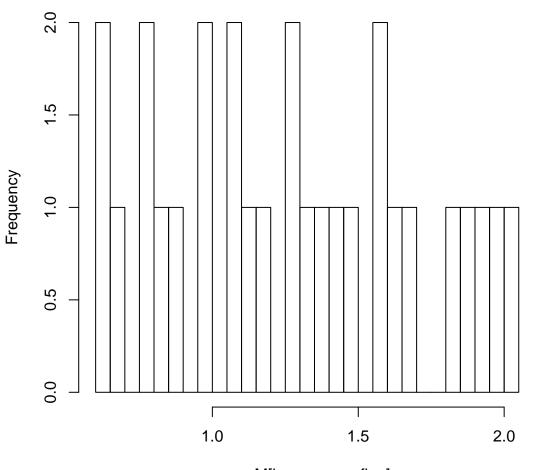
ภาพ[เ, non\_zero\_แล ind#: 20, diag#: 2

# log expression of gene#1558, pval ob=0.9793, non-zero num=9



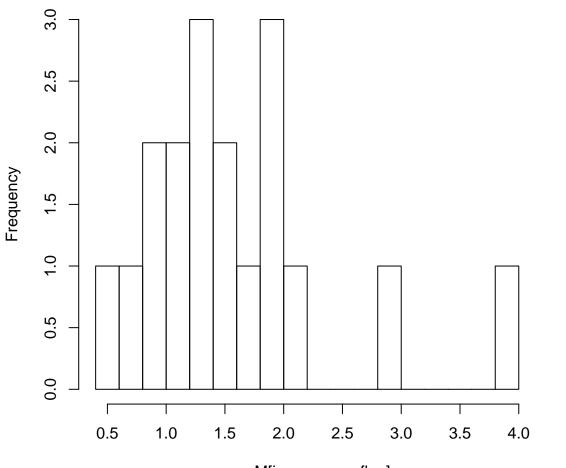
exprM[i, non\_zero\_flag] ind#: 25, diag#: 2

# log expression of gene#1, pval ob=0.2426, non-zero num=28



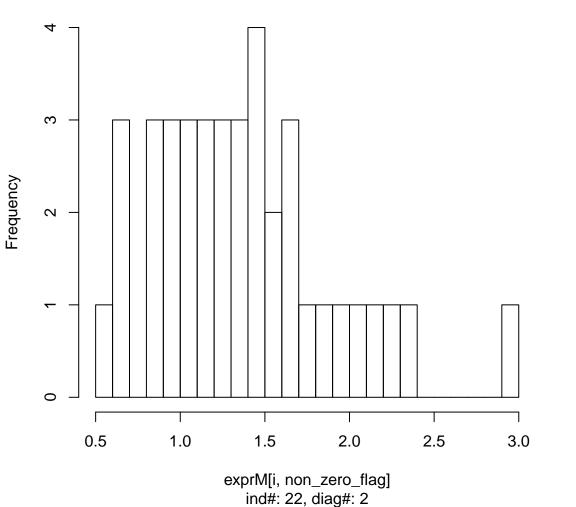
exprM[i, non\_zero\_flag] ind#: 17, diag#: 2

#### log expression of gene#559, pval ob=0, non-zero num=18

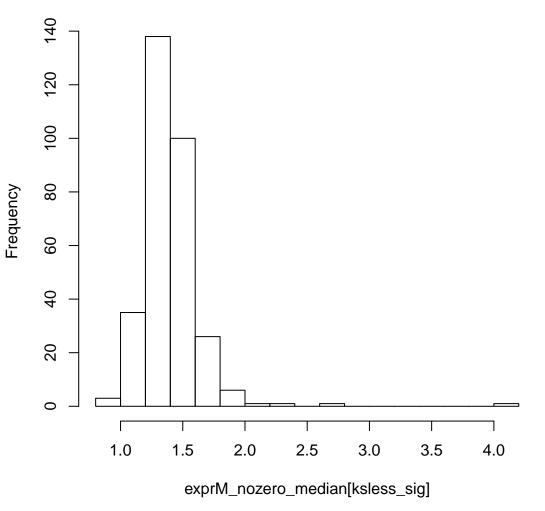


exprM[i, non\_zero\_flag] ind#: 12, diag#: 2

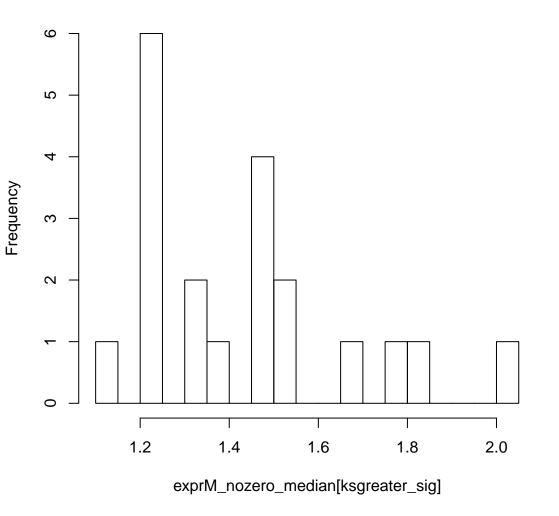
# log expression of gene#2587, pval ob=0.9105, non-zero num=3



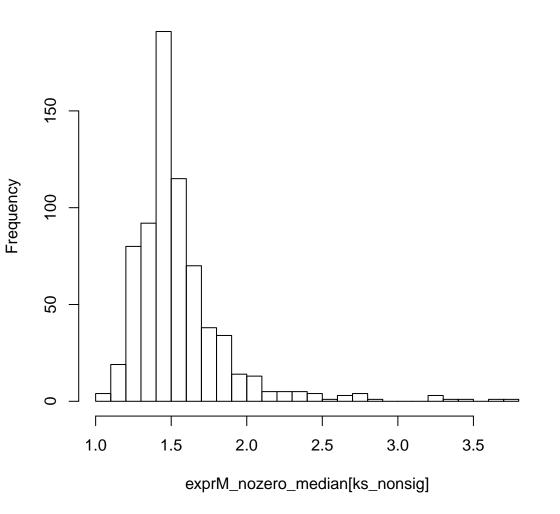
median of nozero log-expres of genes, pval1\_rate<0.2,ksless s



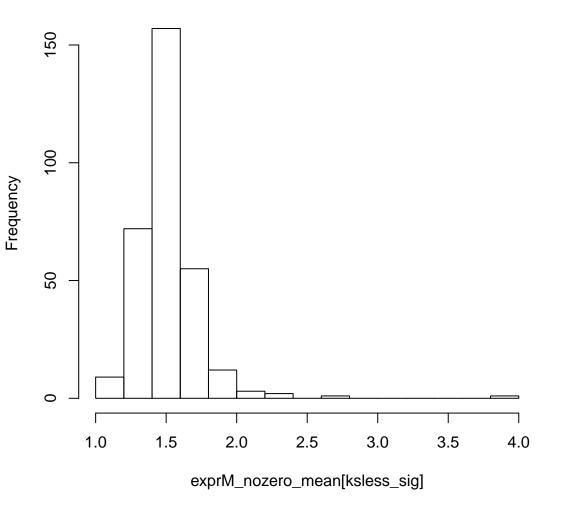
### median of nozero log-expres of genes,pval1\_rate<0.2,ksgreater



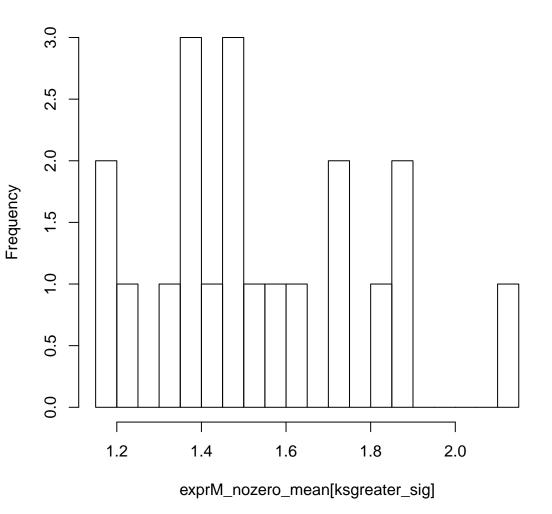
### median of nozero log-expres of genes,pval1\_rate<0.2,ks no si



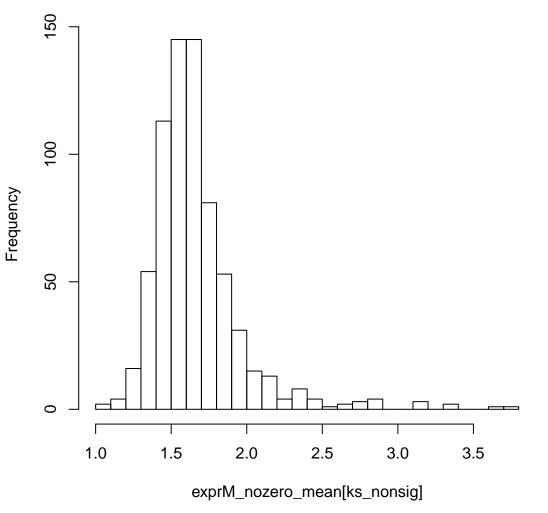
mean of nozero log-expres of genes, pval1\_rate<0.2,ksless si



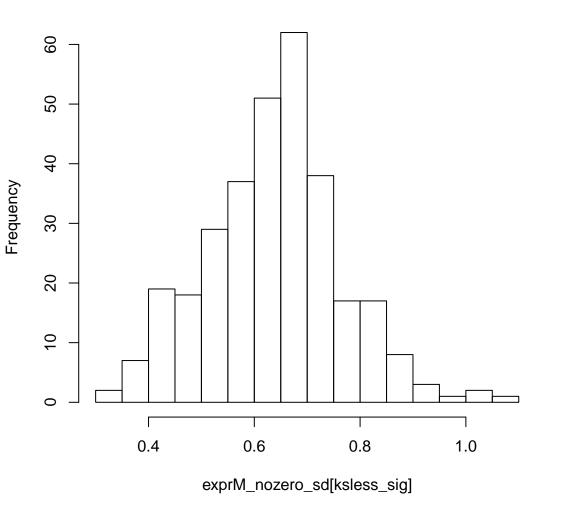
## mean of nozero log-expres of genes,pval1\_rate<0.2,ksgreaters



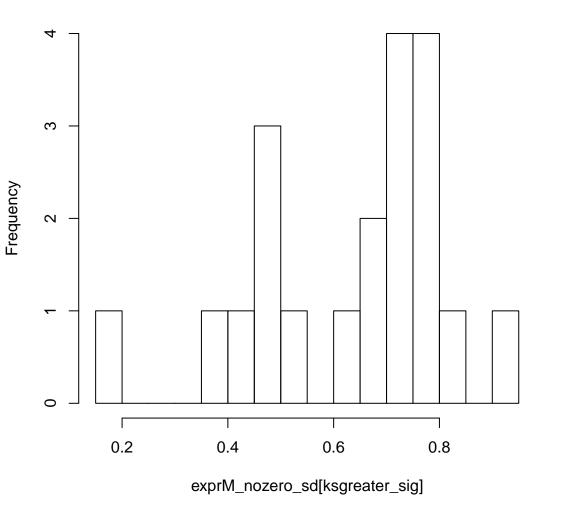
mean of nozero log-expres of genes,pval1\_rate<0.2,ks no sig



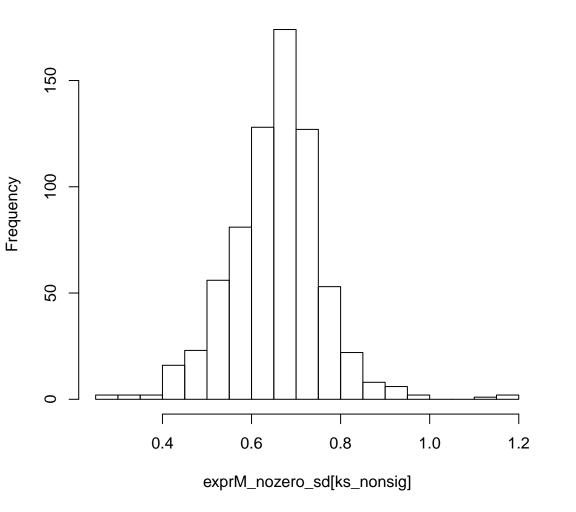
### sd of nozero log-expres of genes, pval1\_rate<0.2,ksless sig



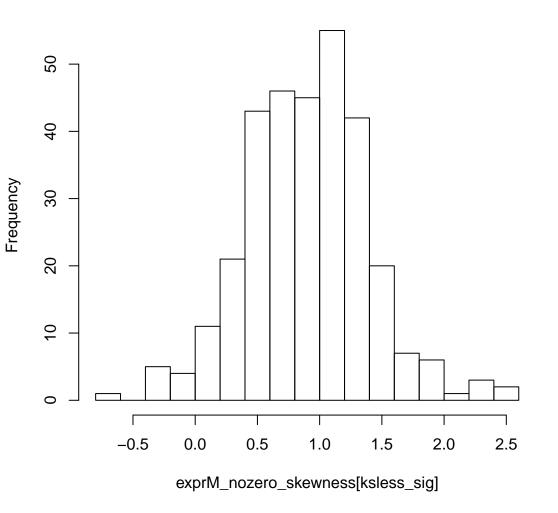
### sd of nozero log-expres of genes,pval1\_rate<0.2,ksgreater signal



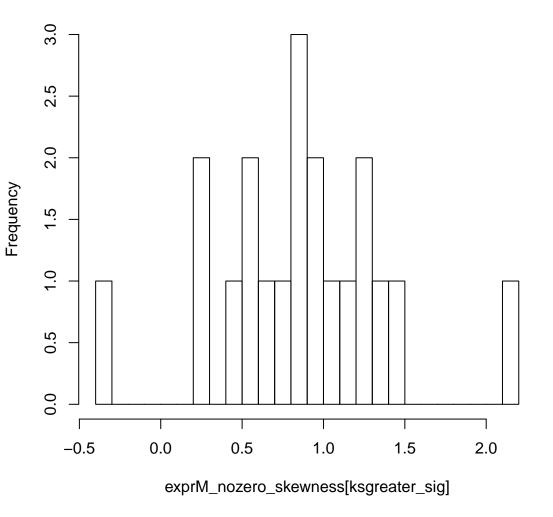
#### sd of nozero log-expres of genes,pval1\_rate<0.2,ks no sig



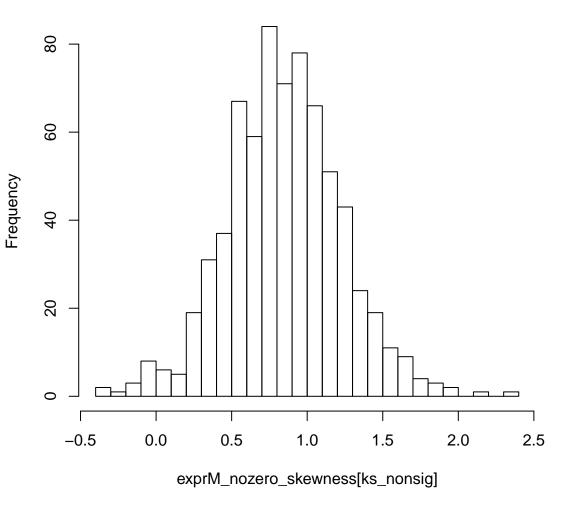
#### skewness of nozero log-expres of genes, pval1\_rate<0.2,ksless



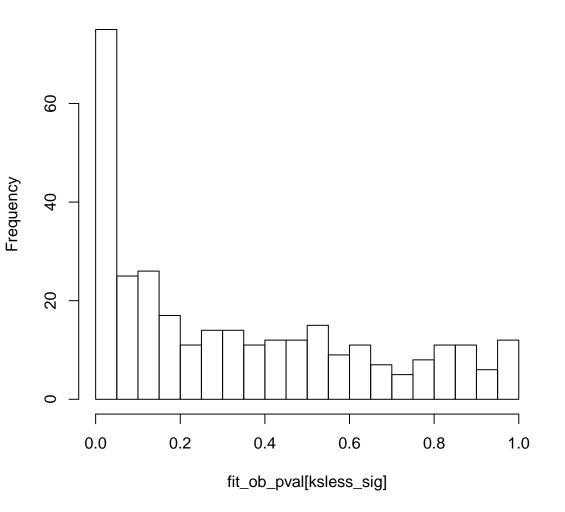
## skewness of nozero log-expres of genes,pval1\_rate<0.2,ksgreate



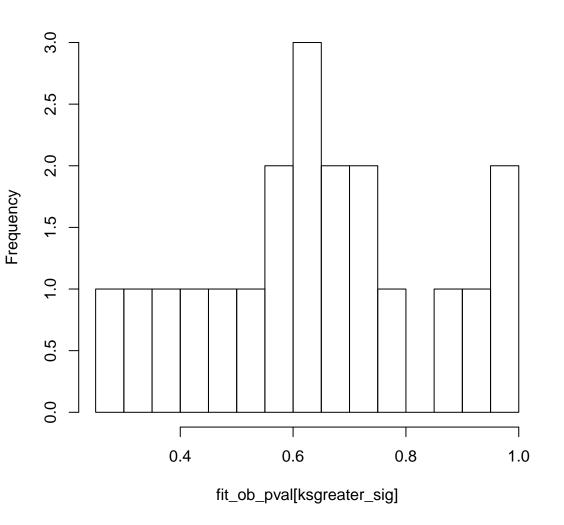
## skewness of nozero log-expres of genes,pval1\_rate<0.2,ks no



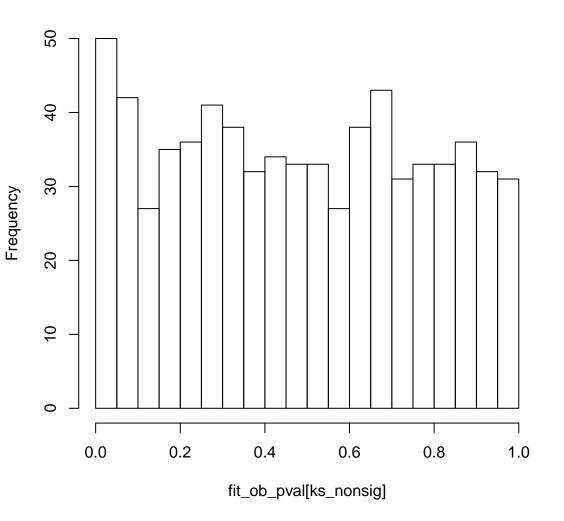
#### observed pvalues with pval1\_rate<0.2,ksless sig



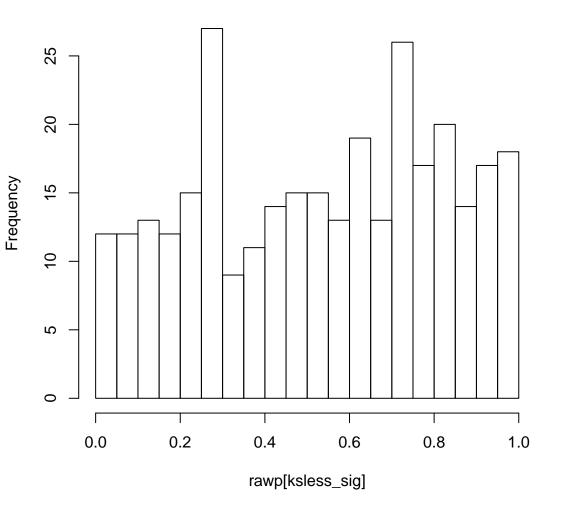
#### observed pvalues with pval1\_rate<0.2,ksgreater sig



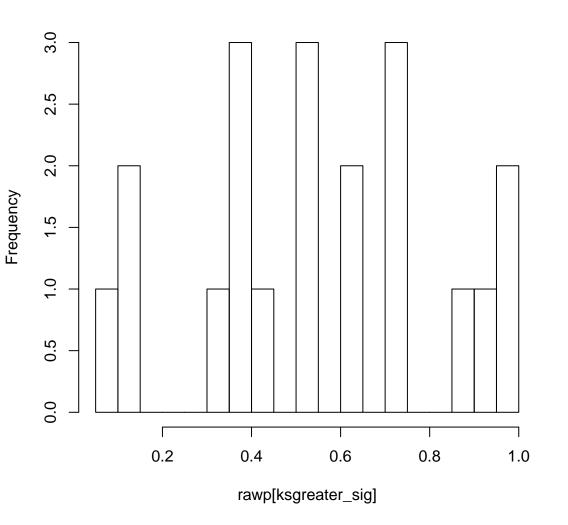
#### observed pvalues with pval1\_rate<0.2,ks no sig



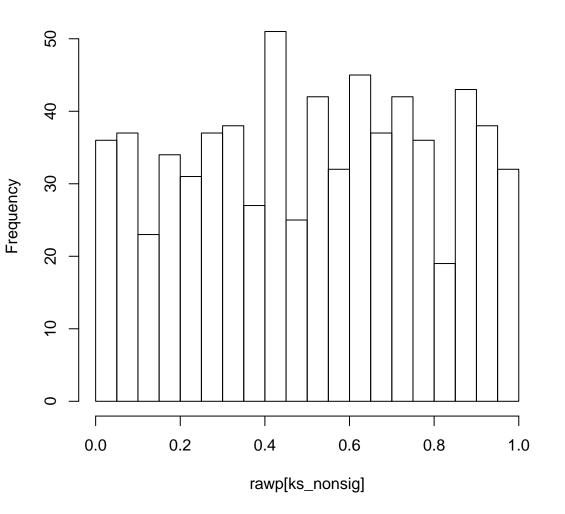
#### permutation pvalues with pval1\_rate<0.2,ksless sig



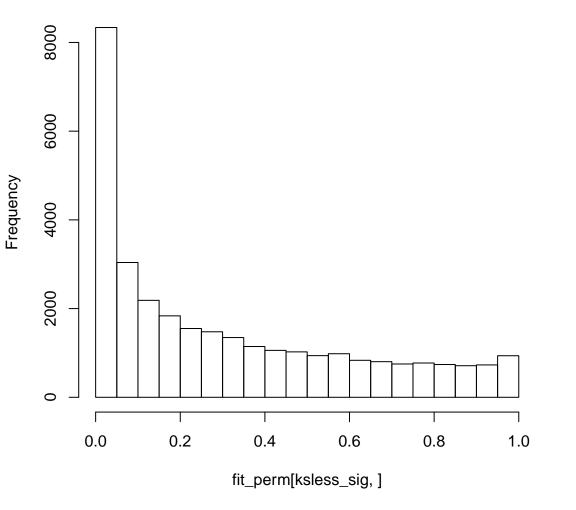
#### permutation pvalues with pval1\_rate<0.2,ksgreater sig



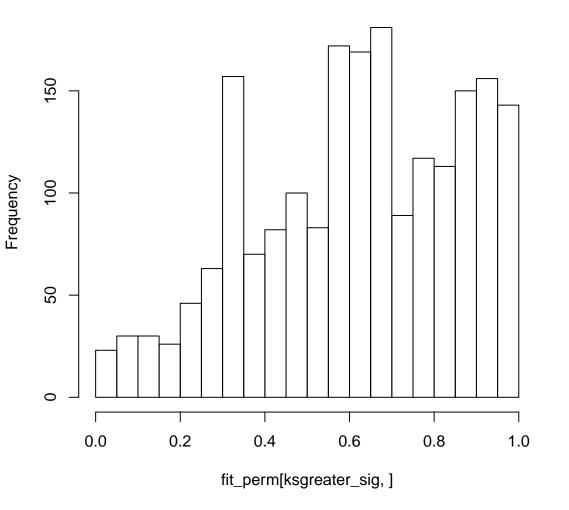
#### permutation pvalues with pval1\_rate<0.2,ks no sig



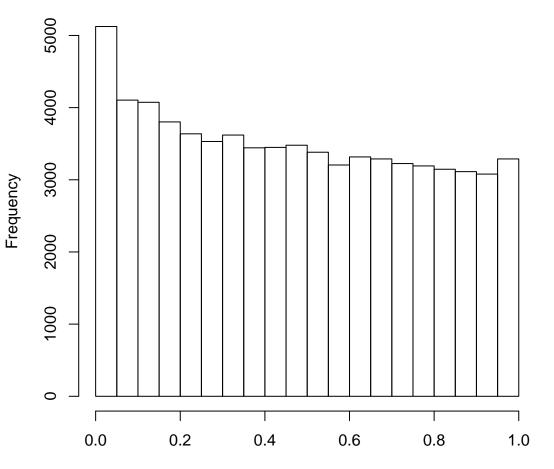
## pvalues from permutation data with pval1\_rate<0.2,ksless sig



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



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



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