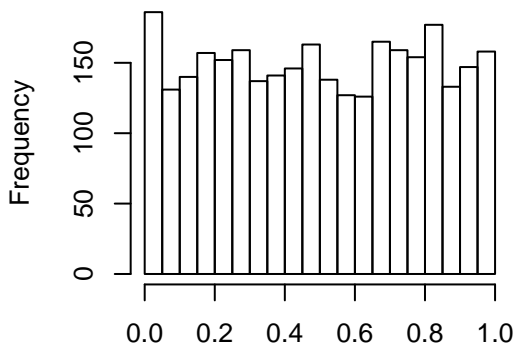
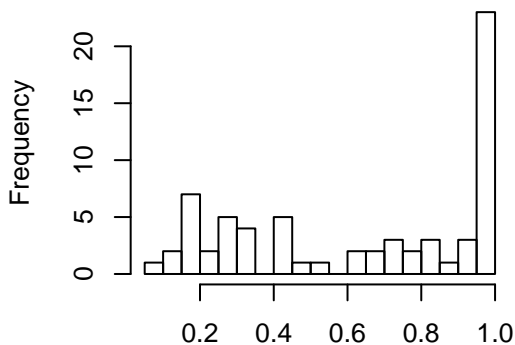


perm pvalues



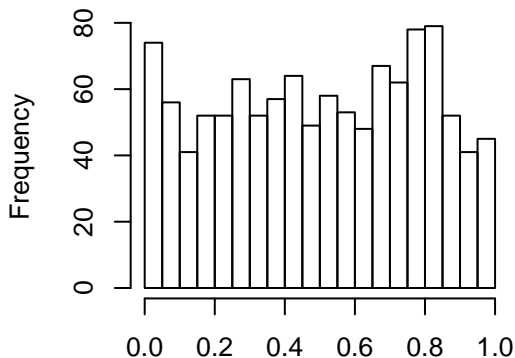
rawp  
n=2996

s,rawM\_0zero\_num<max(20,min(rawM\_0zero\_num,100))



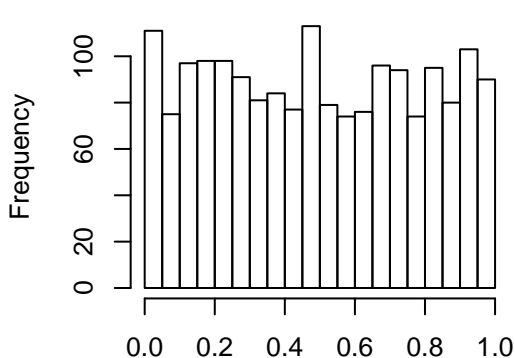
rawp[rawM\_0zero\_num < max(20, min(rawM\_0zero\_num, 100))]  
n=67, ks\_pval=0.00145793618262746

perm pvalues,rawM\_0zero\_num <= 20-



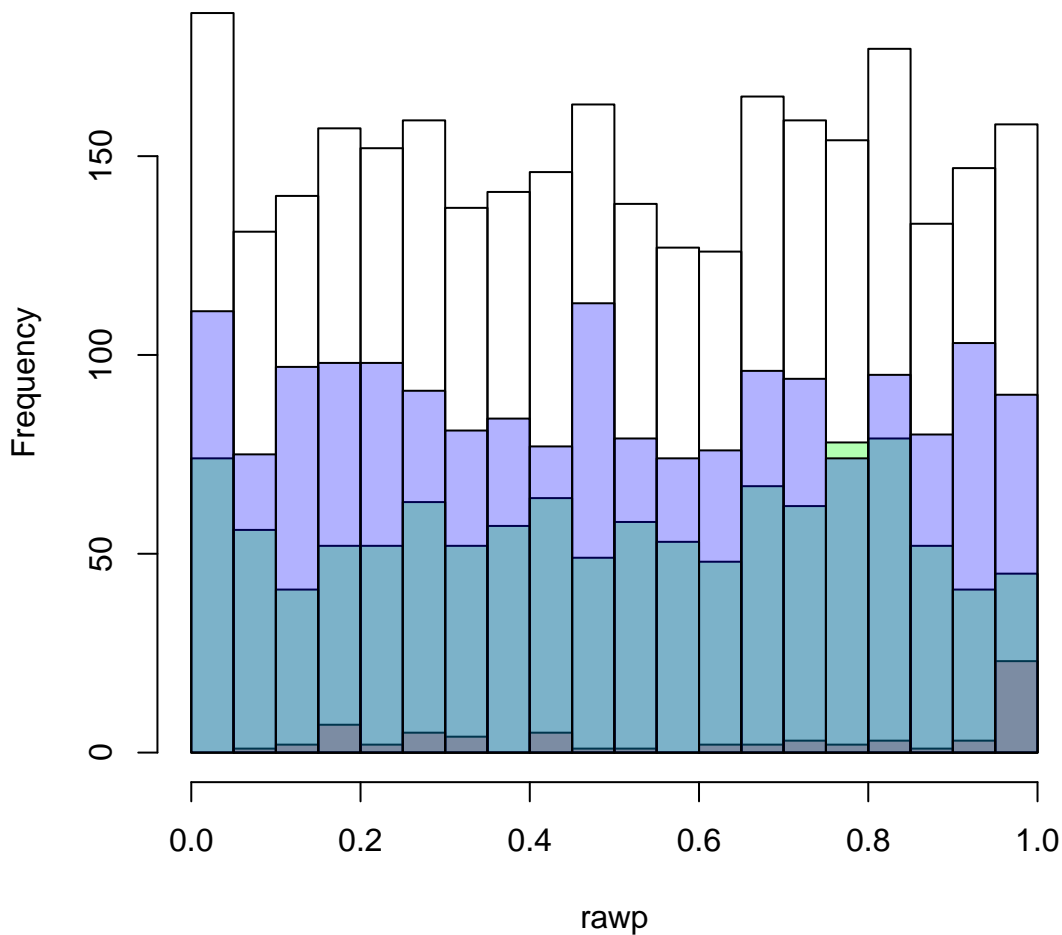
rawp[rawM\_0zero\_num <= 20 & rawM\_0zero\_num <= 100]  
n=1143, ks\_pval=0.552662806545381

perm pvalues,rawM\_0zero\_num >= 100-

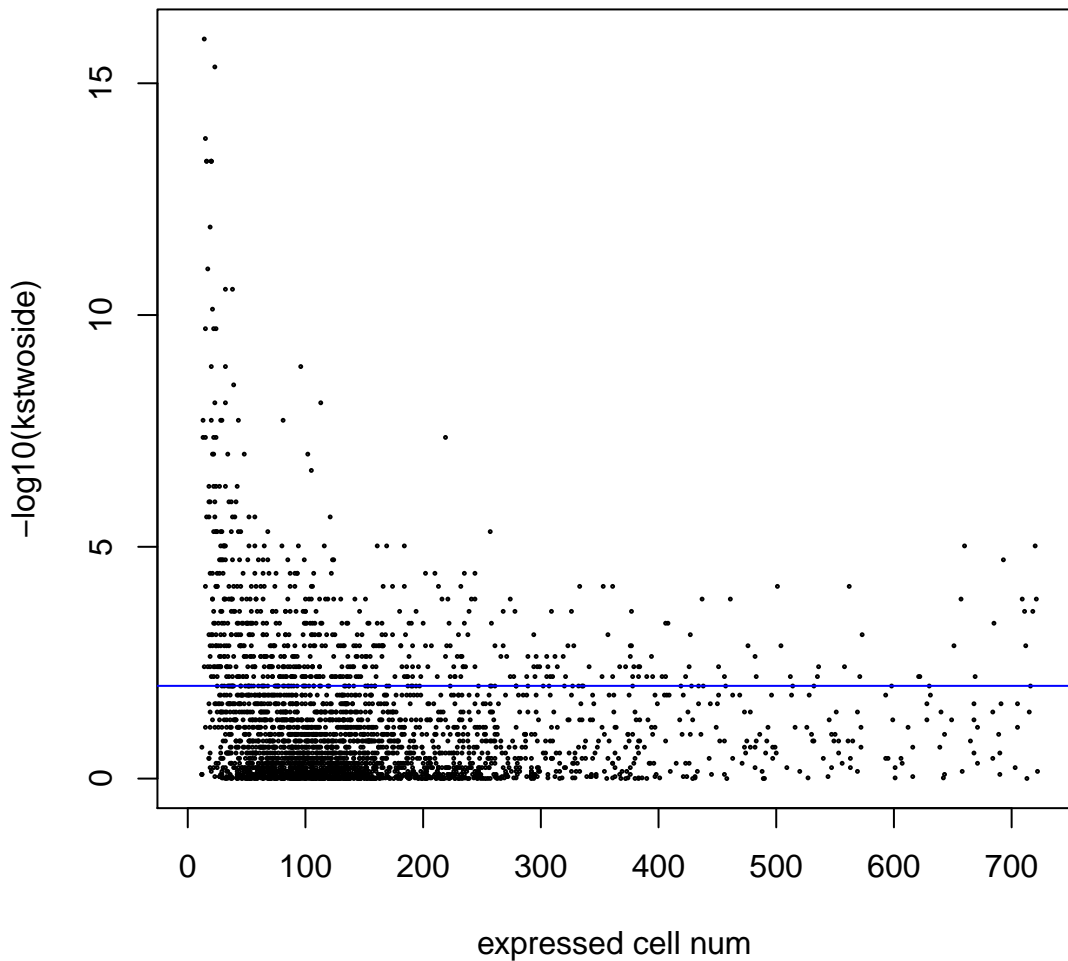


rawp[rawM\_0zero\_num >= 100]  
n=1786, ks\_pval=0.838057682170524

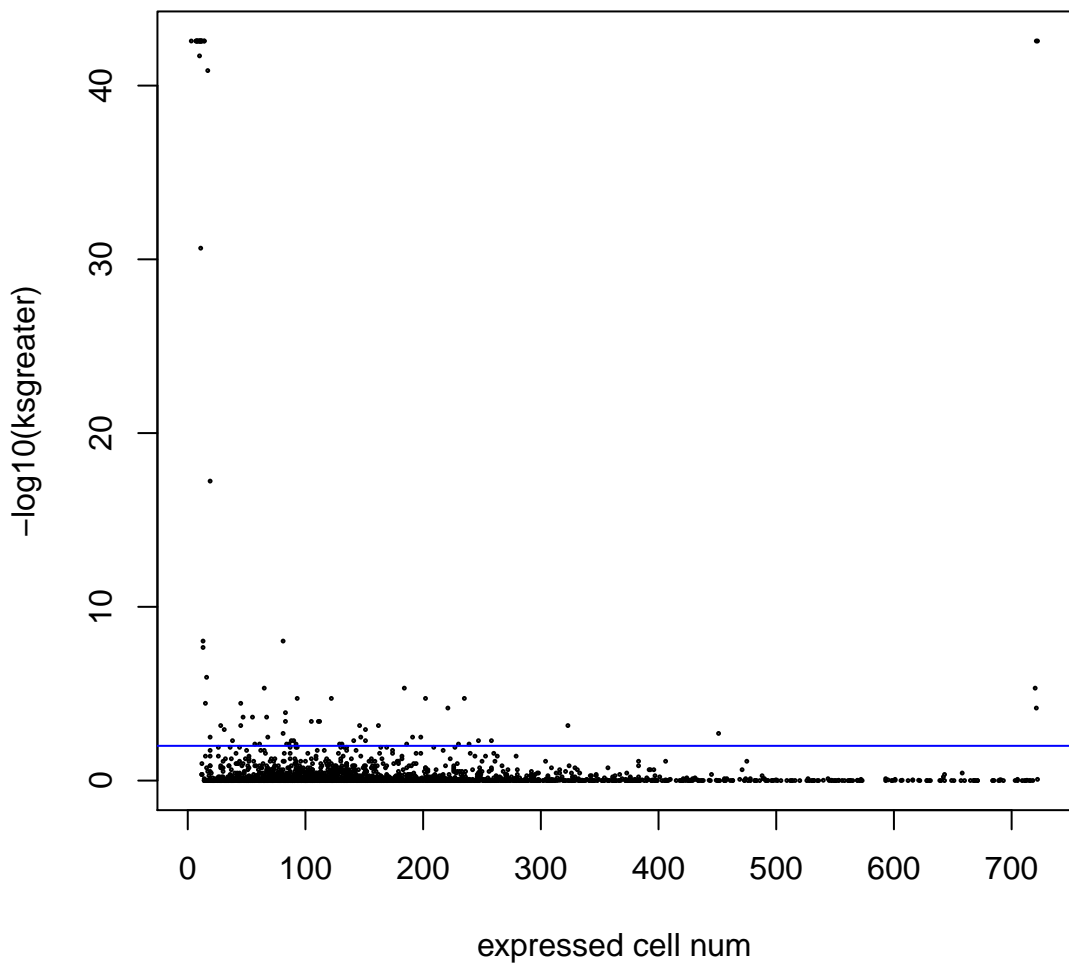
# perm pvalues



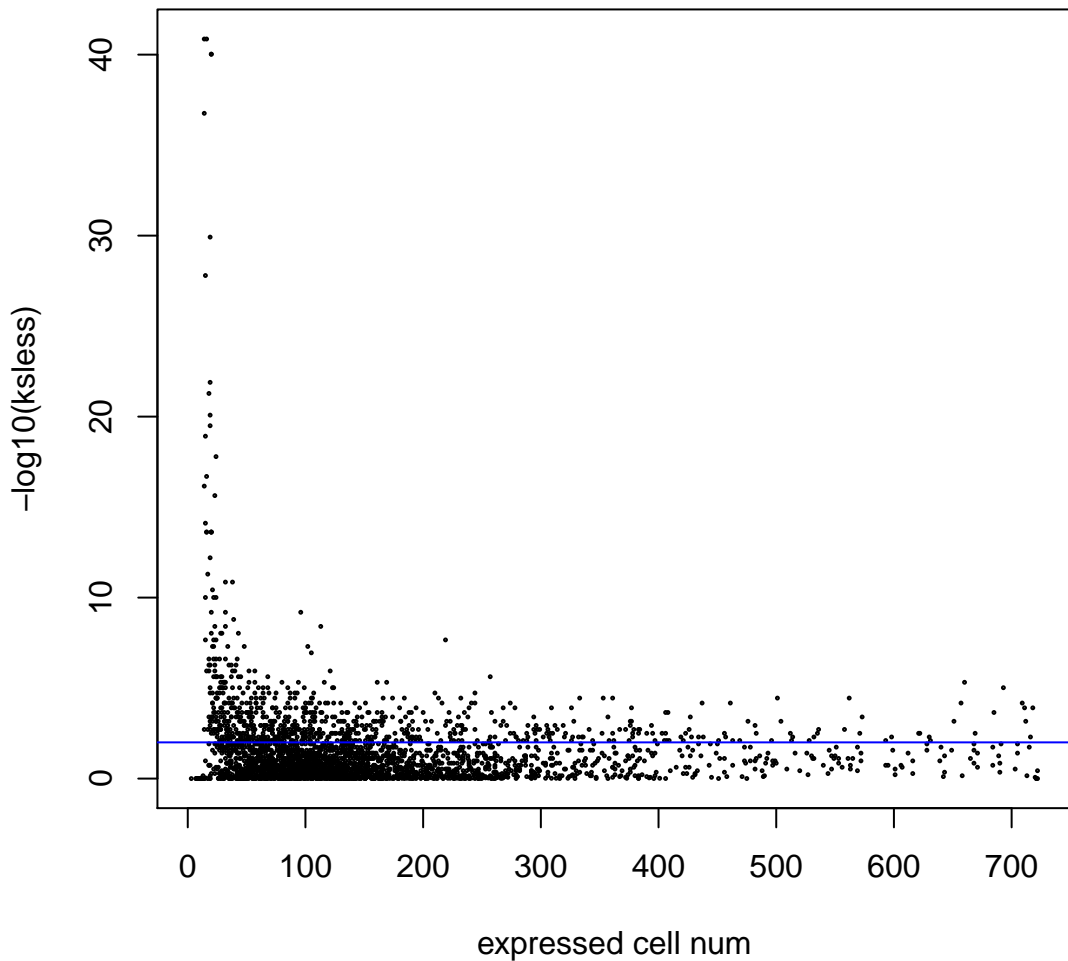
**sig\_KStwoside: 20.995%**



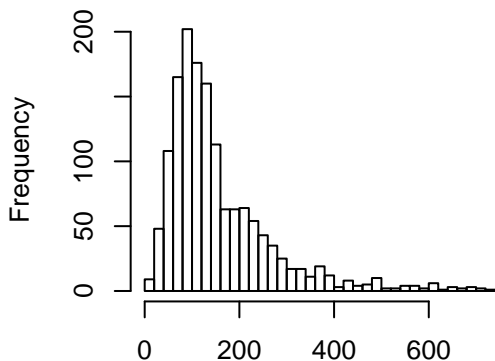
**sig\_KSgreater: 2.57%**



**sig\_KSless: 25.033%**

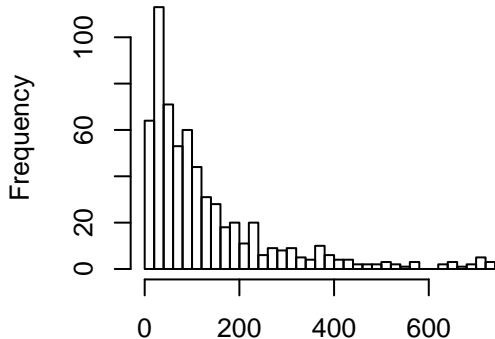


**expression cell num,kstwoside>0.2**



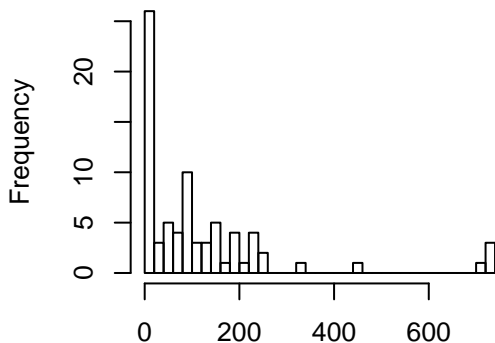
rawM\_0zero\_num[kstwoside > 0.2]

**expression cell num,kstwoside<0.01**



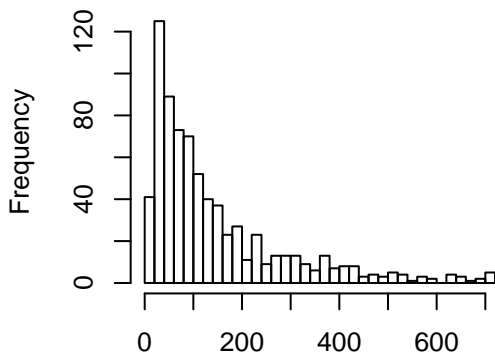
rawM\_0zero\_num[kstwoside < 0.01]

**expression cell num,ksgreater<0.01**



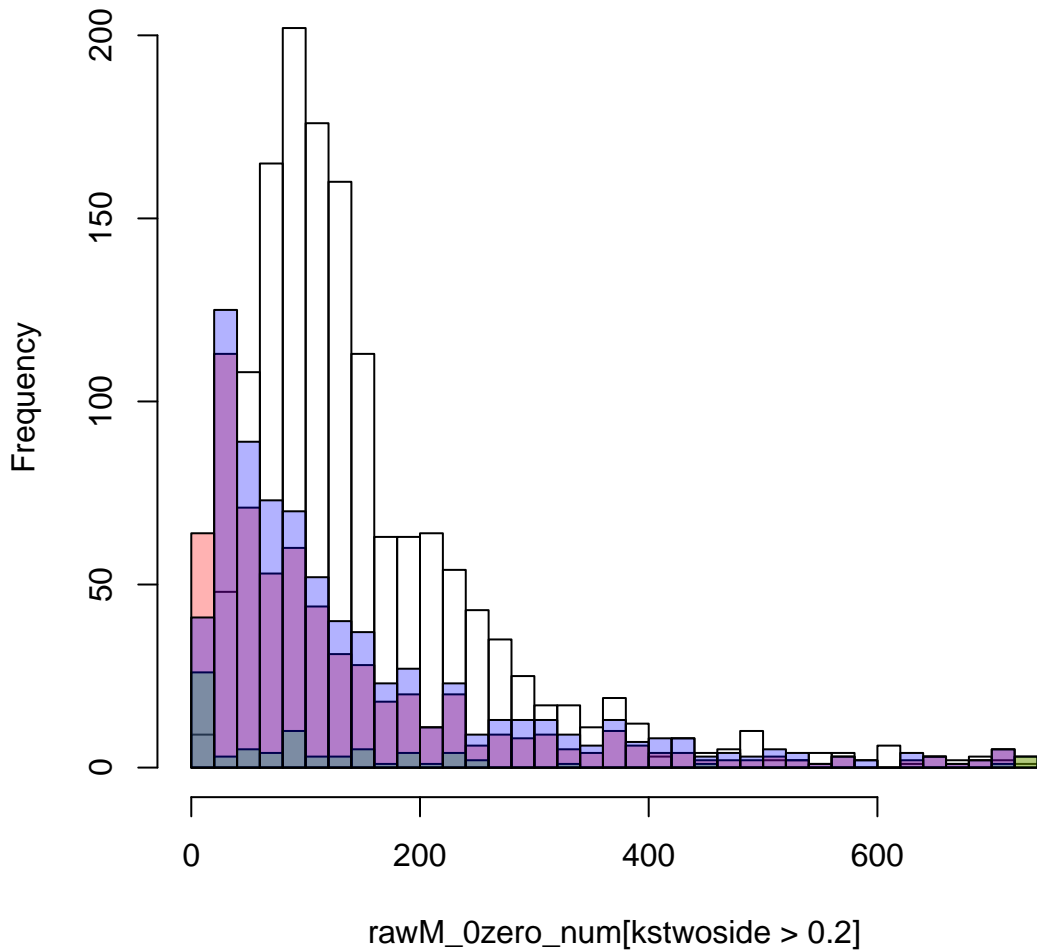
rawM\_0zero\_num[ksgreater < 0.01]

**expression cell num,ksless<0.01**

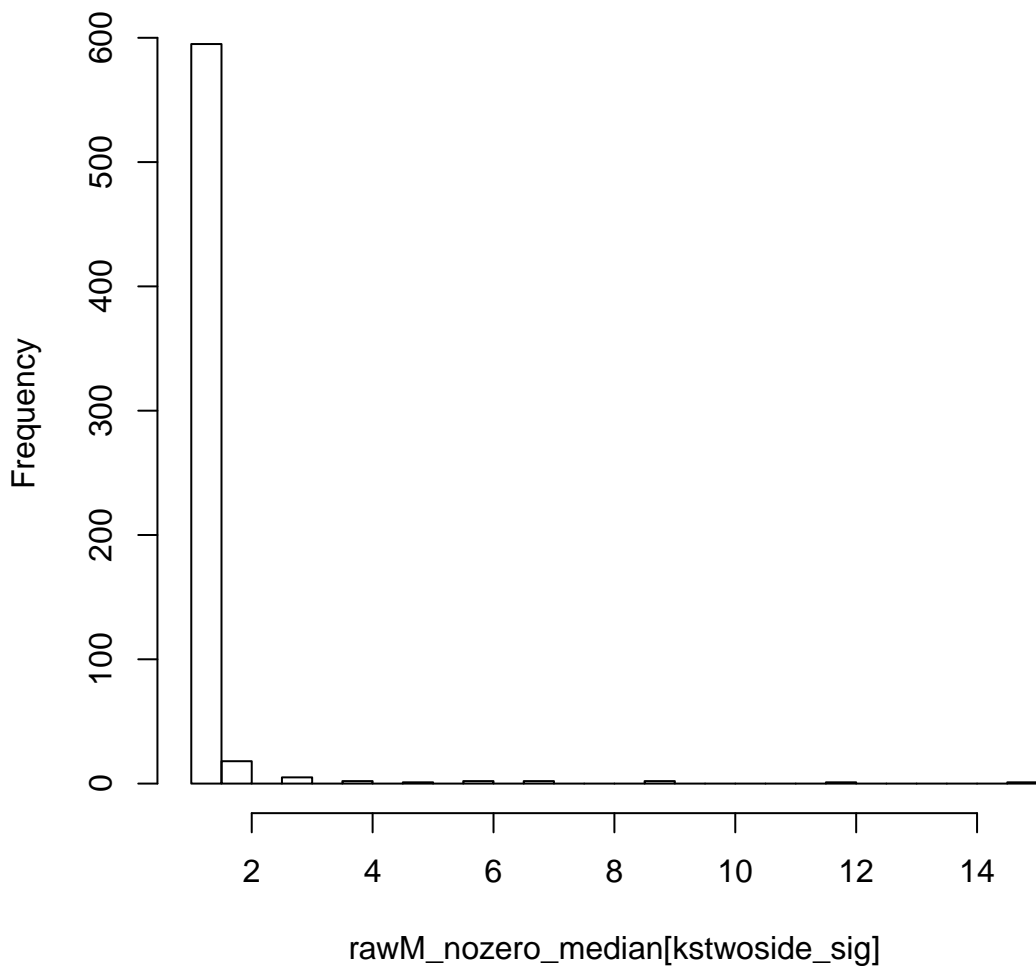


rawM\_0zero\_num[ksless < 0.01]

## expression cell num

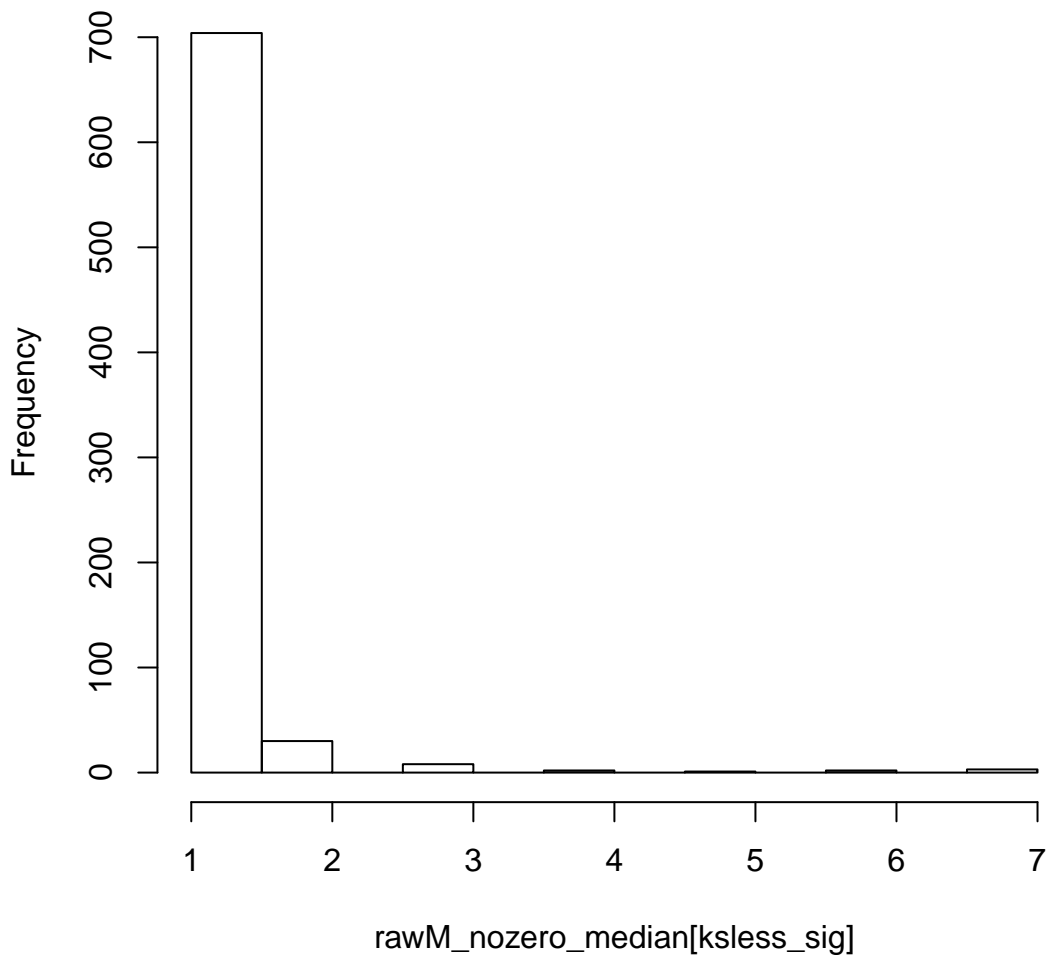


# median of nozero log-express of genes, kstwoside sig

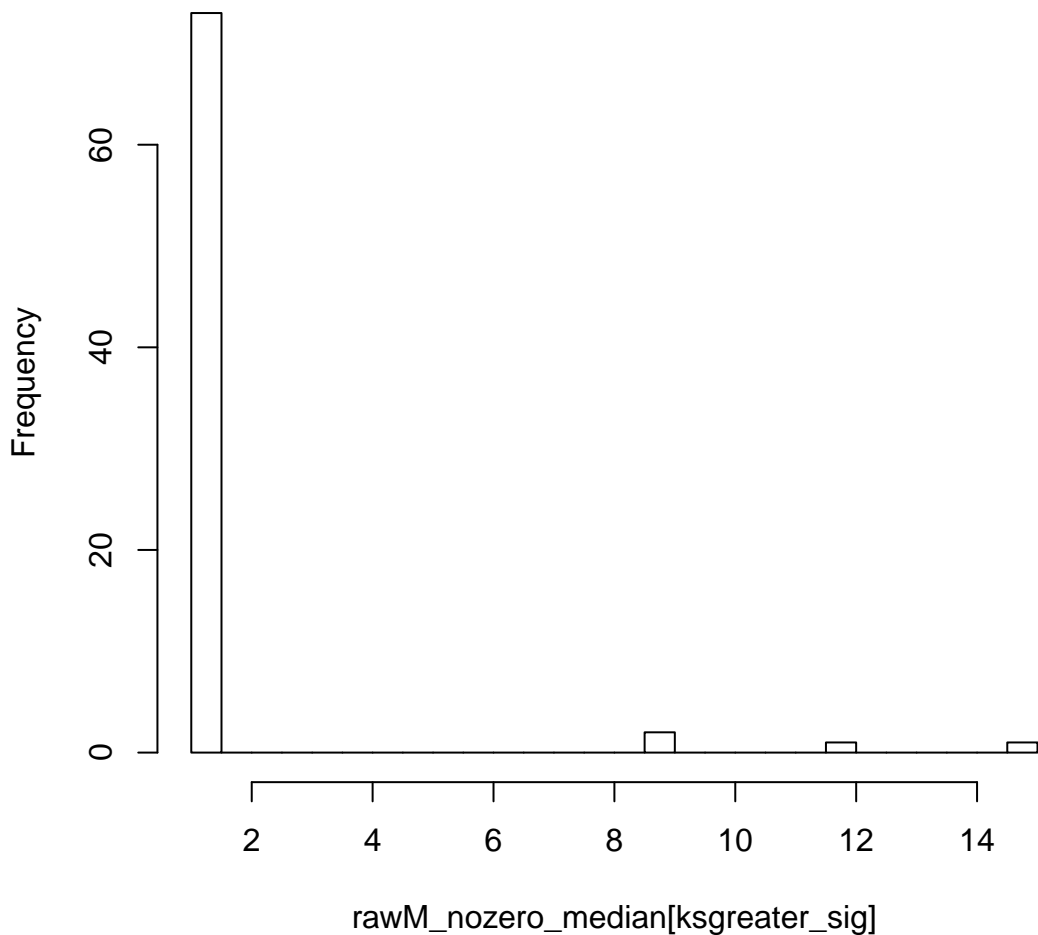




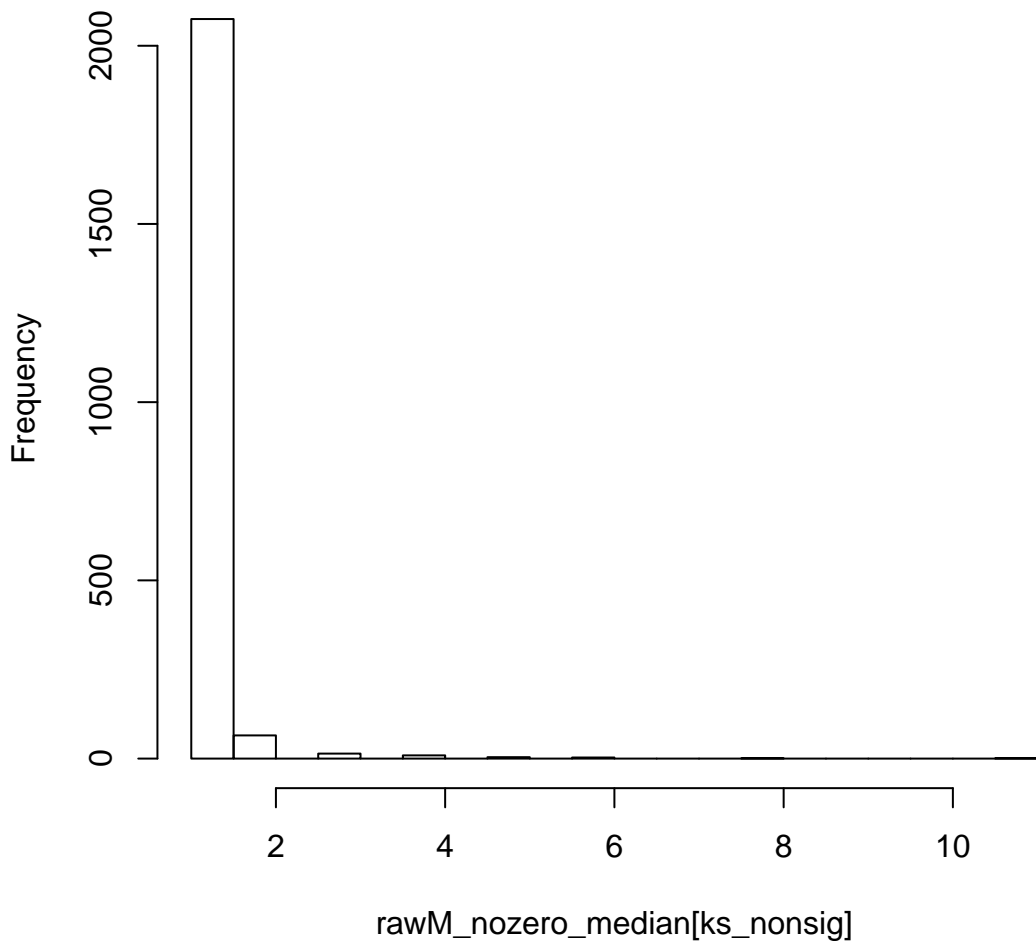
# median of nozero log-express of genes, ksless sig



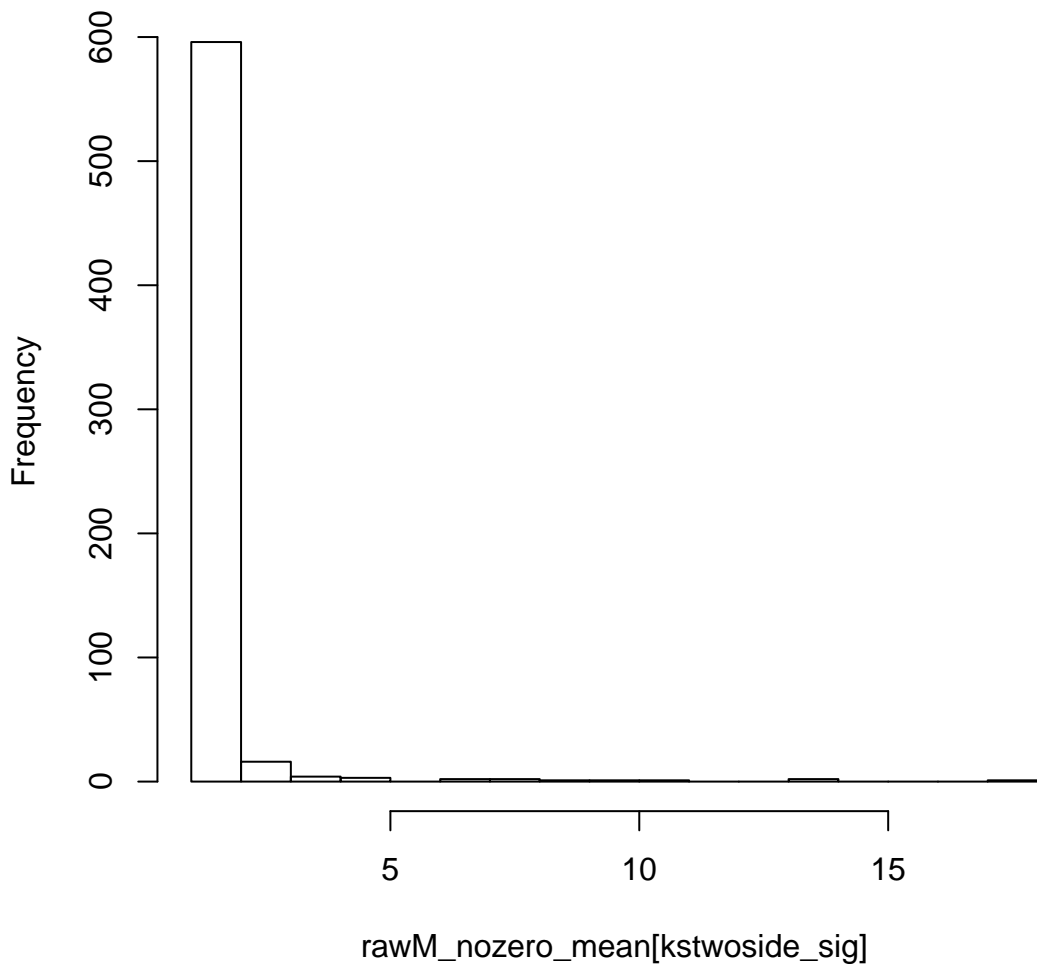
# median of nozero log-express of genes,ksgreater sig



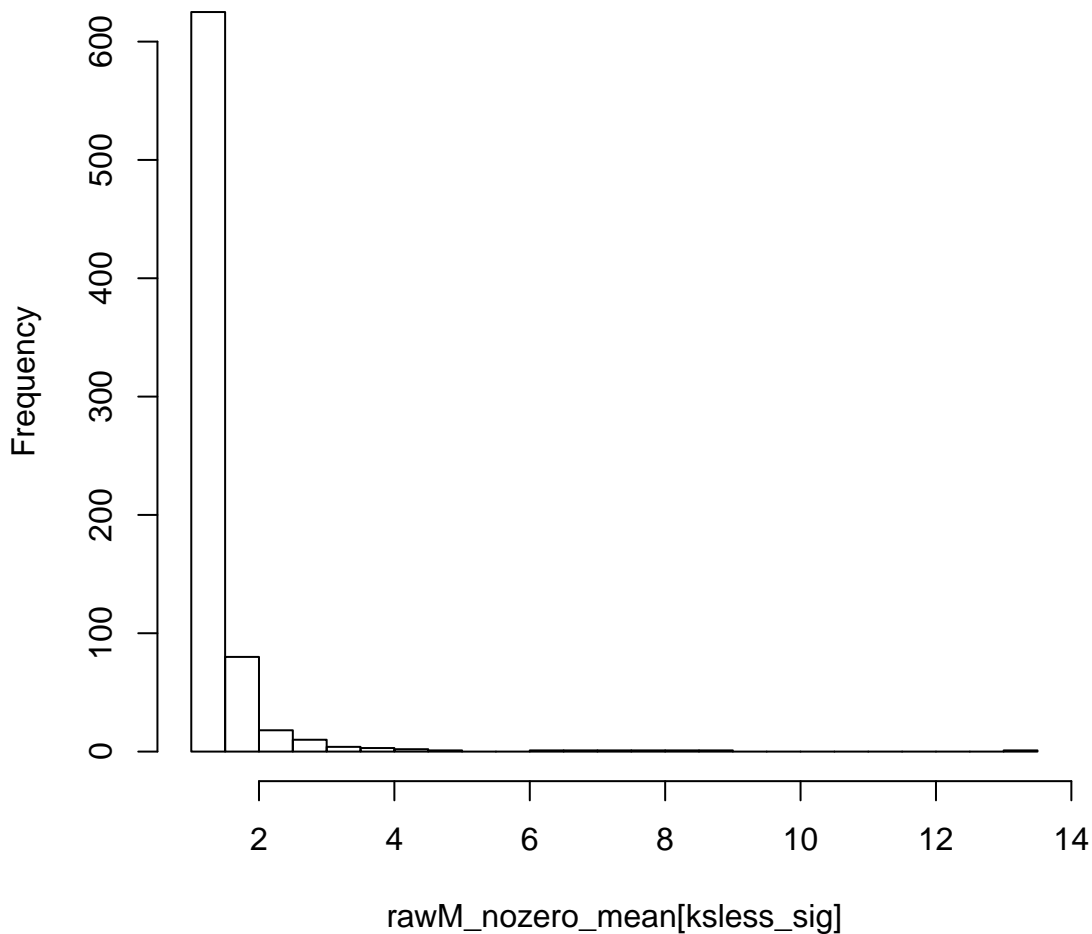
# median of nozero log-expres of genes,ks no sig



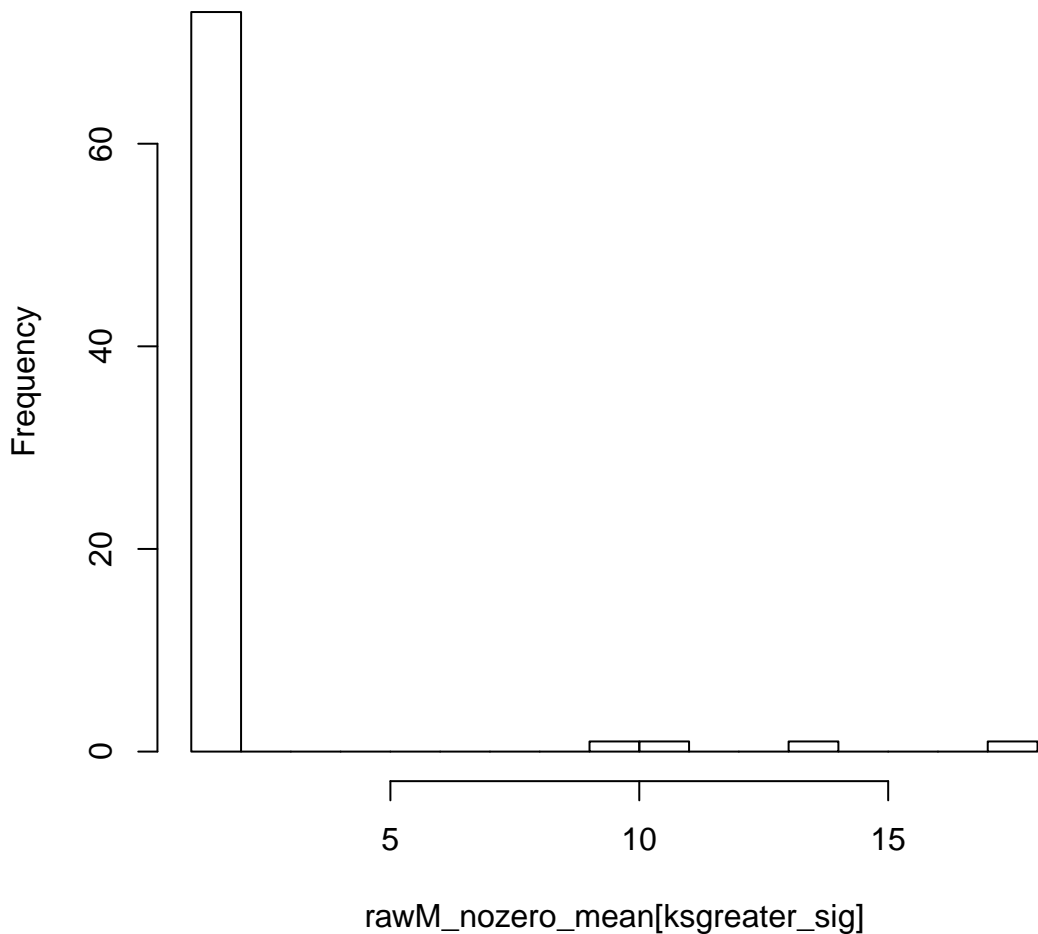
# mean of nozero log-express of genes, kstwoside sig



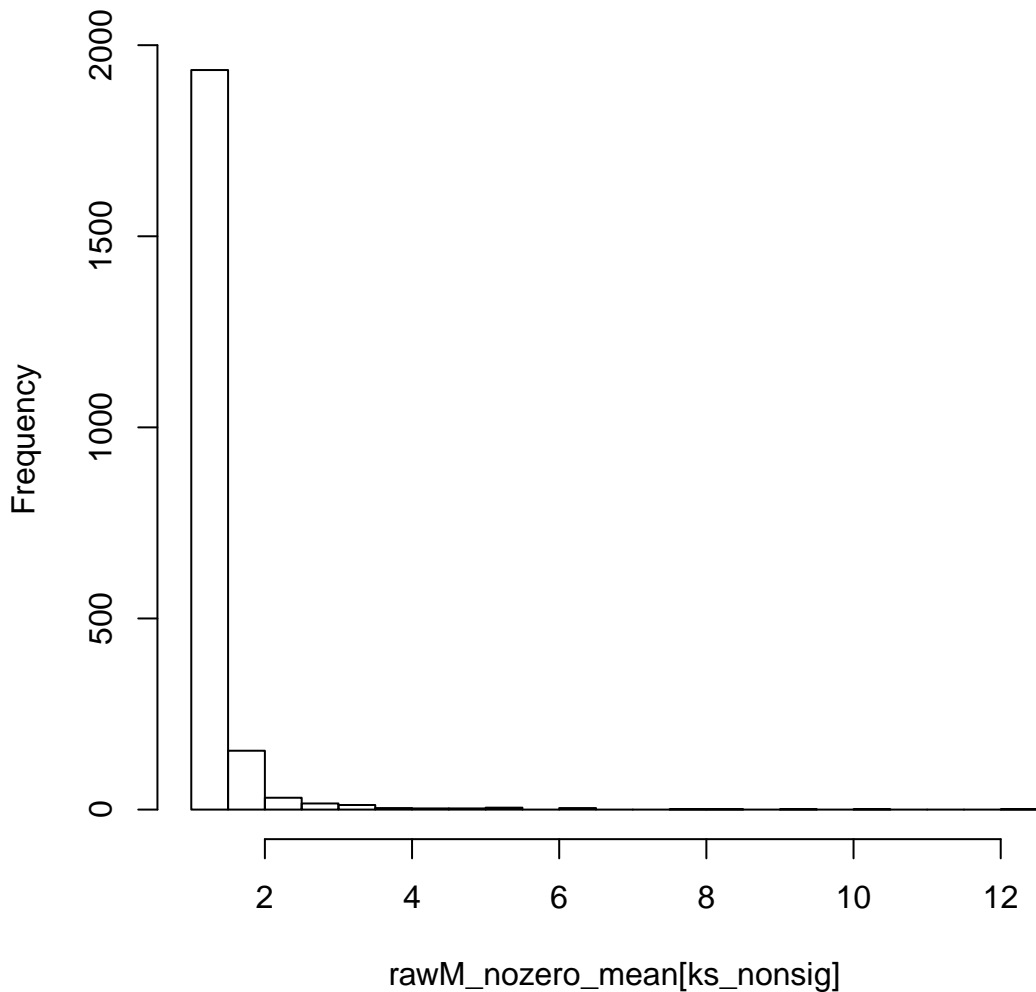
# mean of nozero log-express of genes, ksless sig



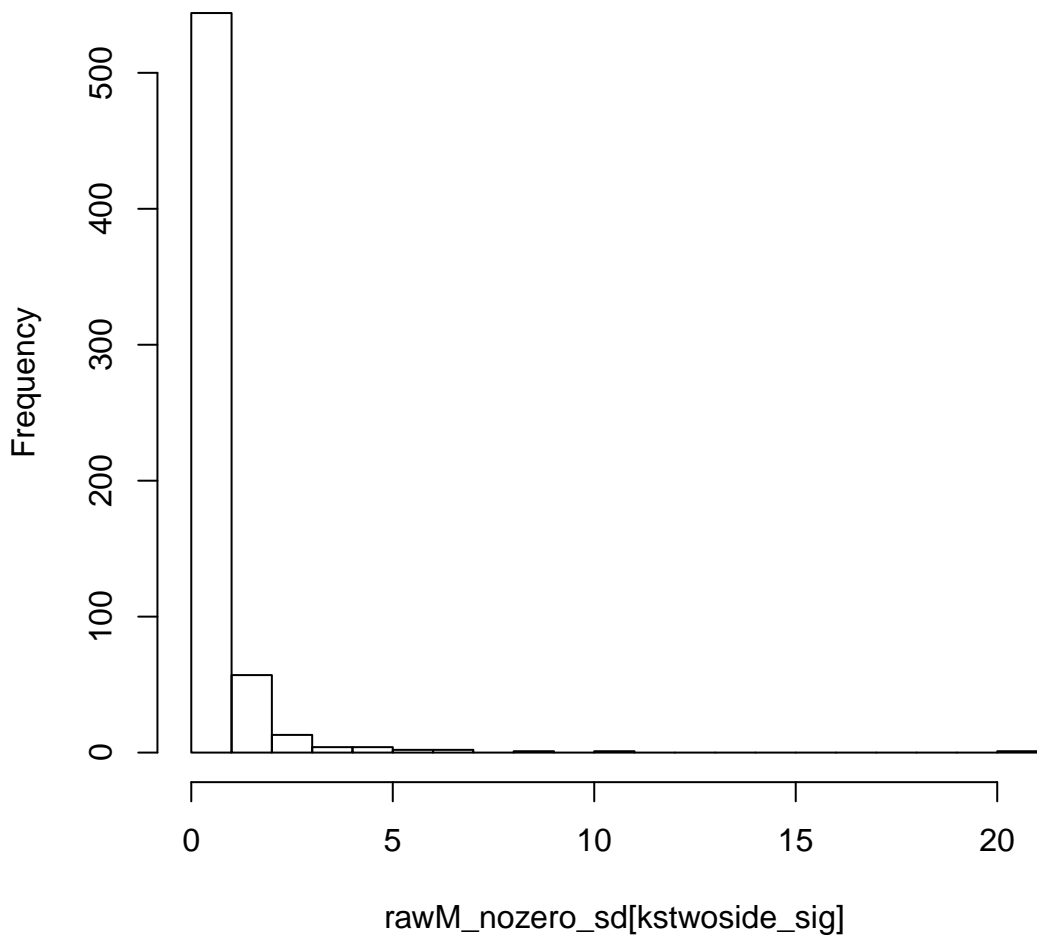
# mean of nozero log-expres of genes,ksgreater sig



# mean of nozero log-expres of genes,ks no sig

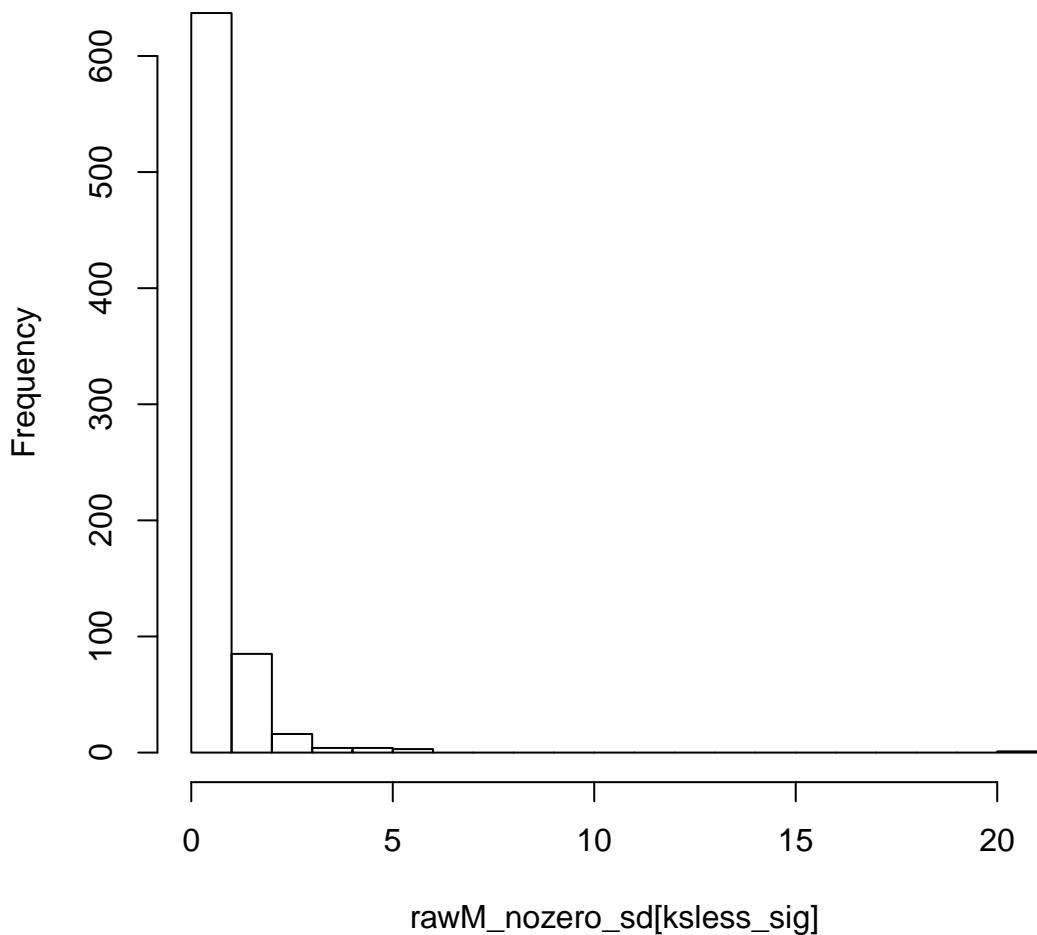


# sd of nozero log-expres of genes, kstwoside sig

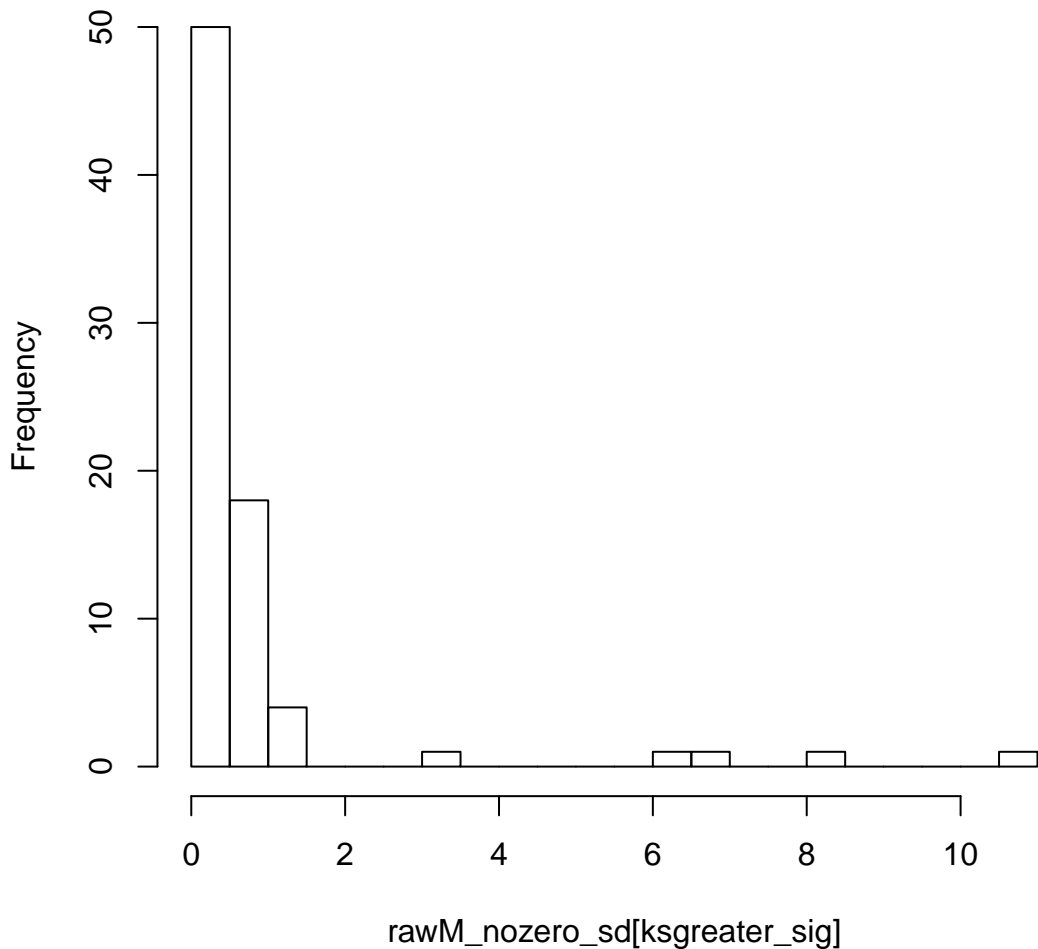




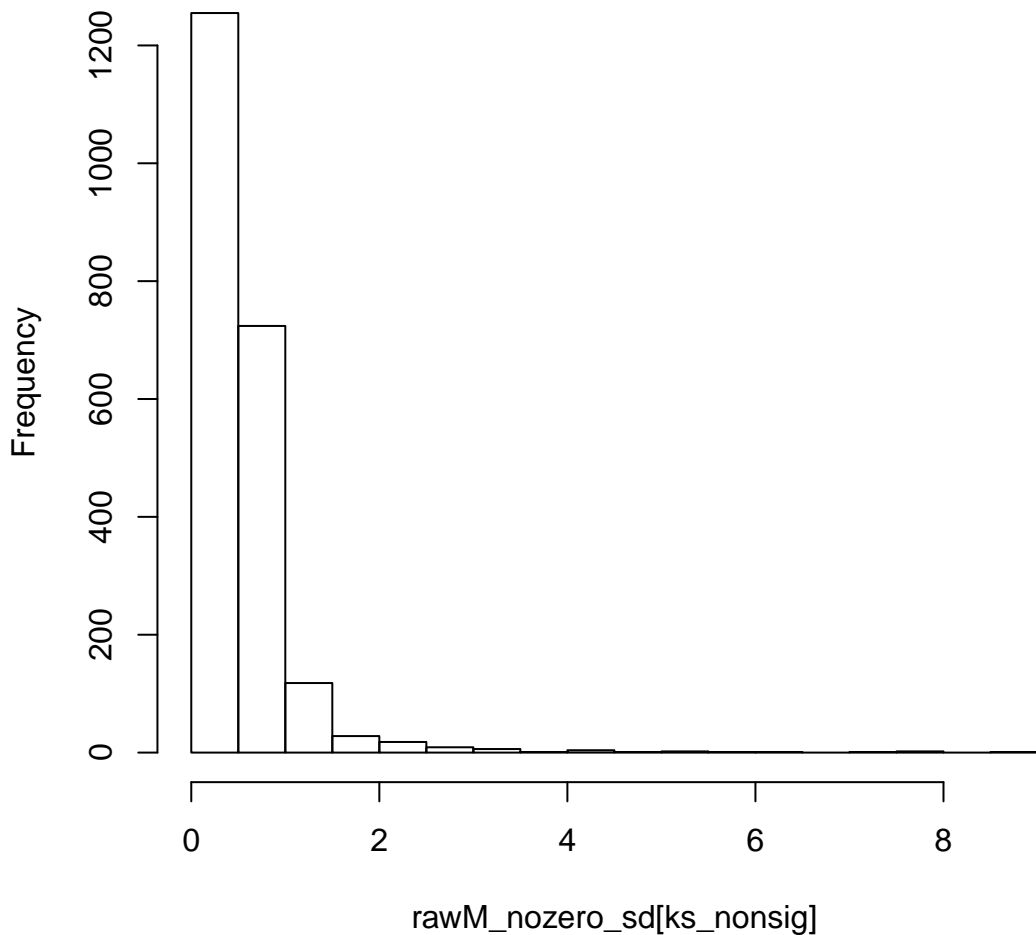
# sd of nozero log-express of genes, ksless sig



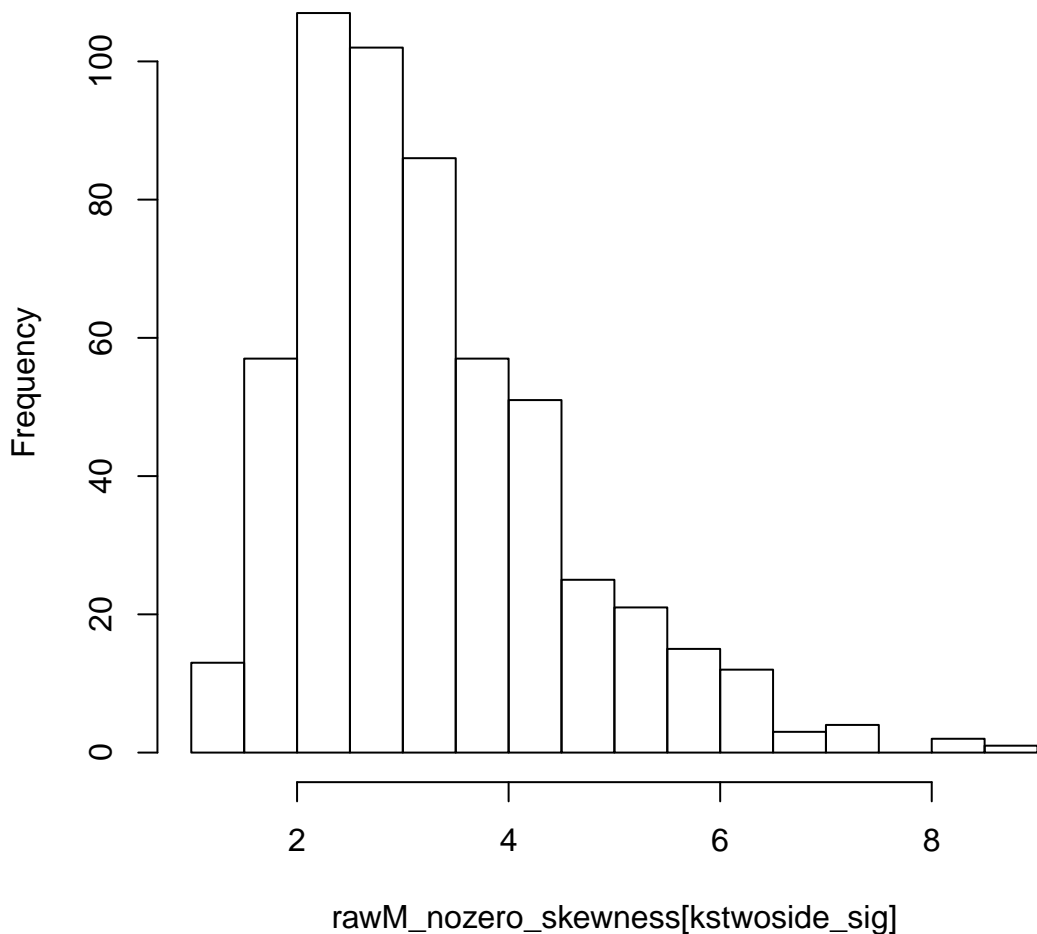
# sd of nozero log-expres of genes,ksgreater sig



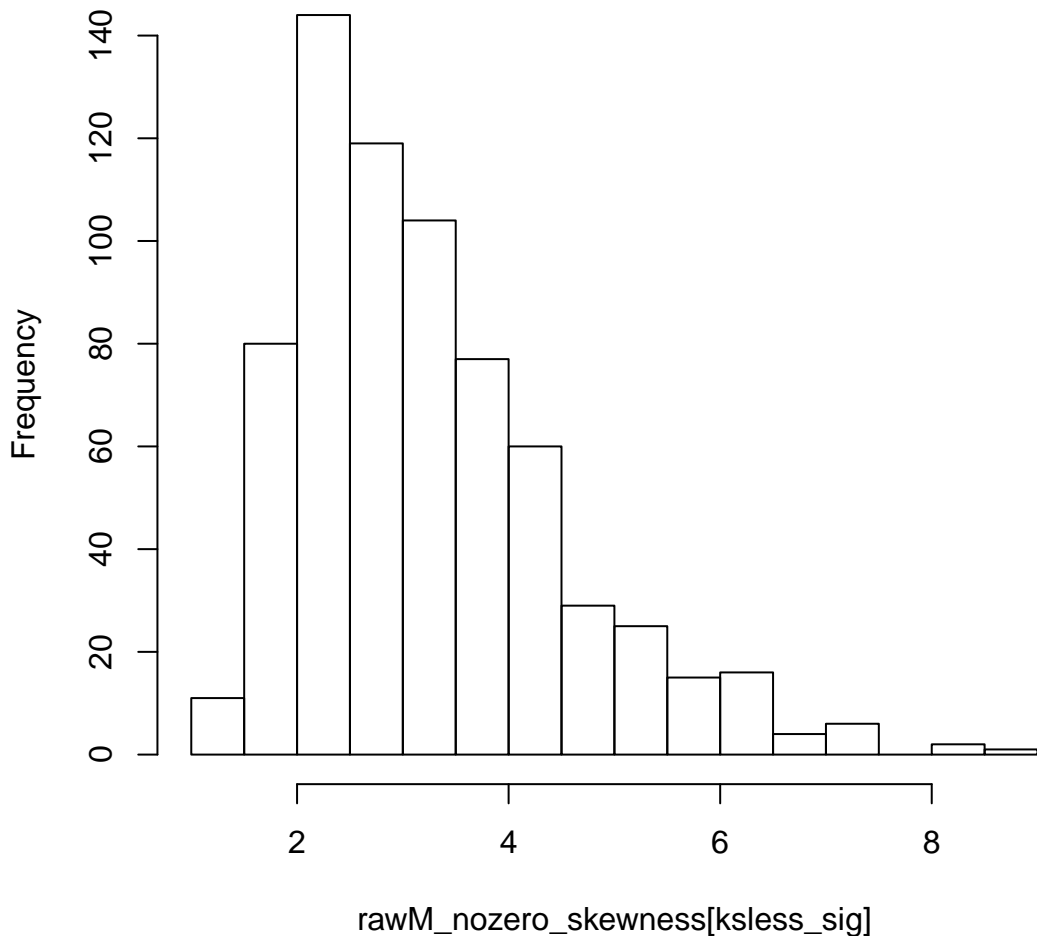
# sd of nozero log-express of genes,ks no sig



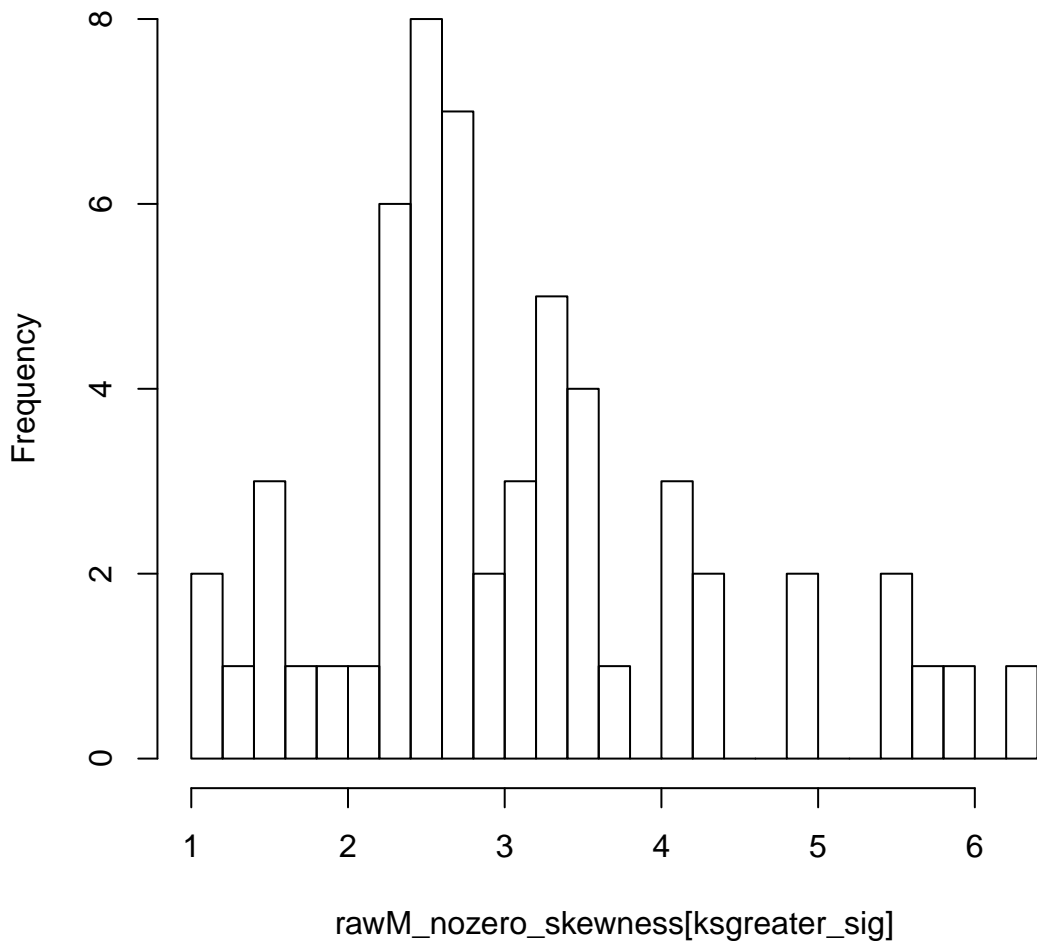
# skewness of nozero log-express of genes, kstwo side sig



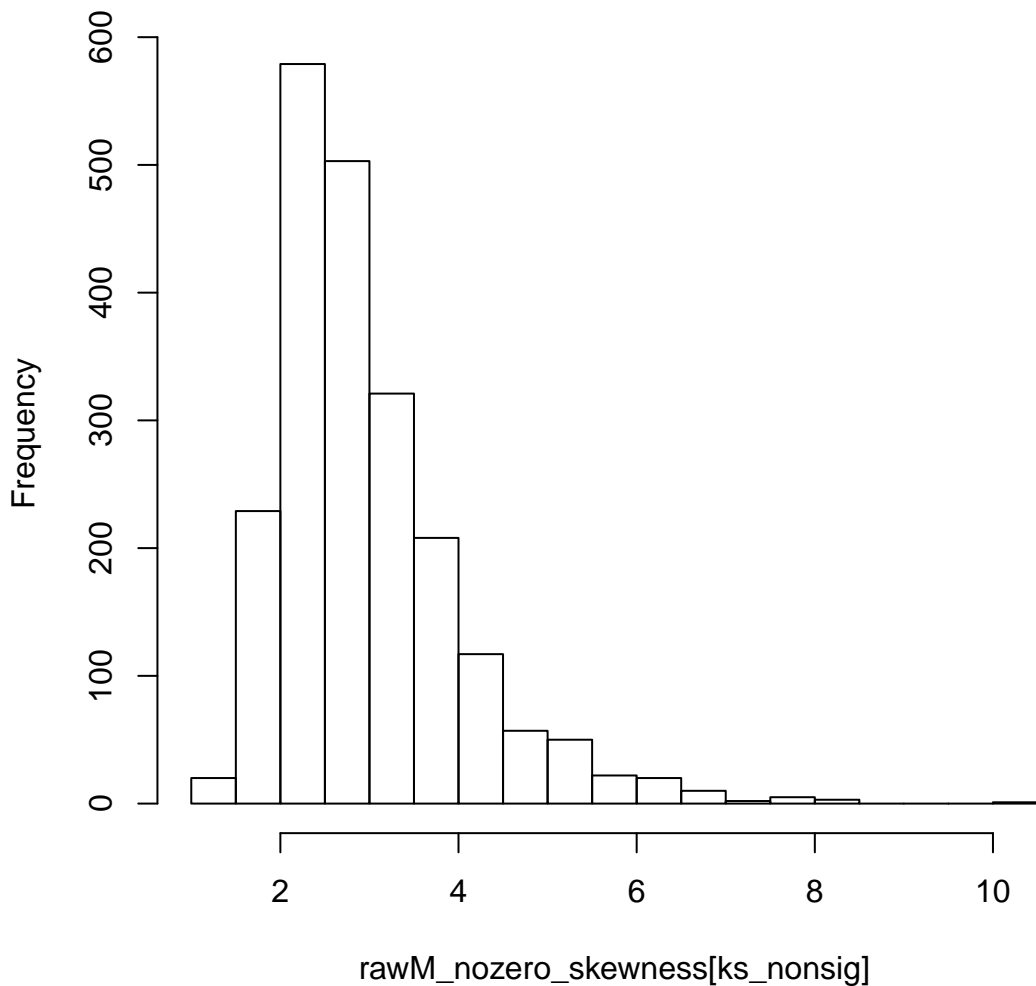
# skewness of nozero log-express of genes, ksless sig



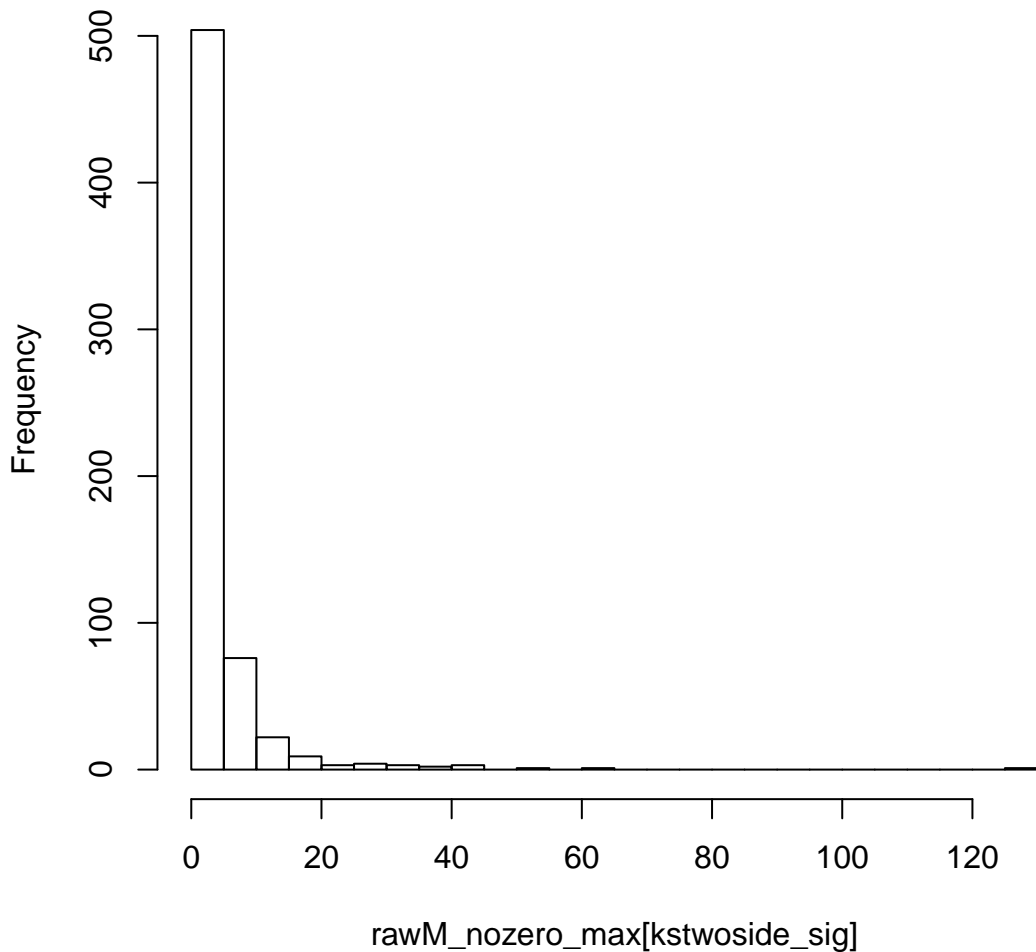
# skewness of nozero log-express of genes,ksgreater sig



# skewness of nozero log-express of genes,ks no sig

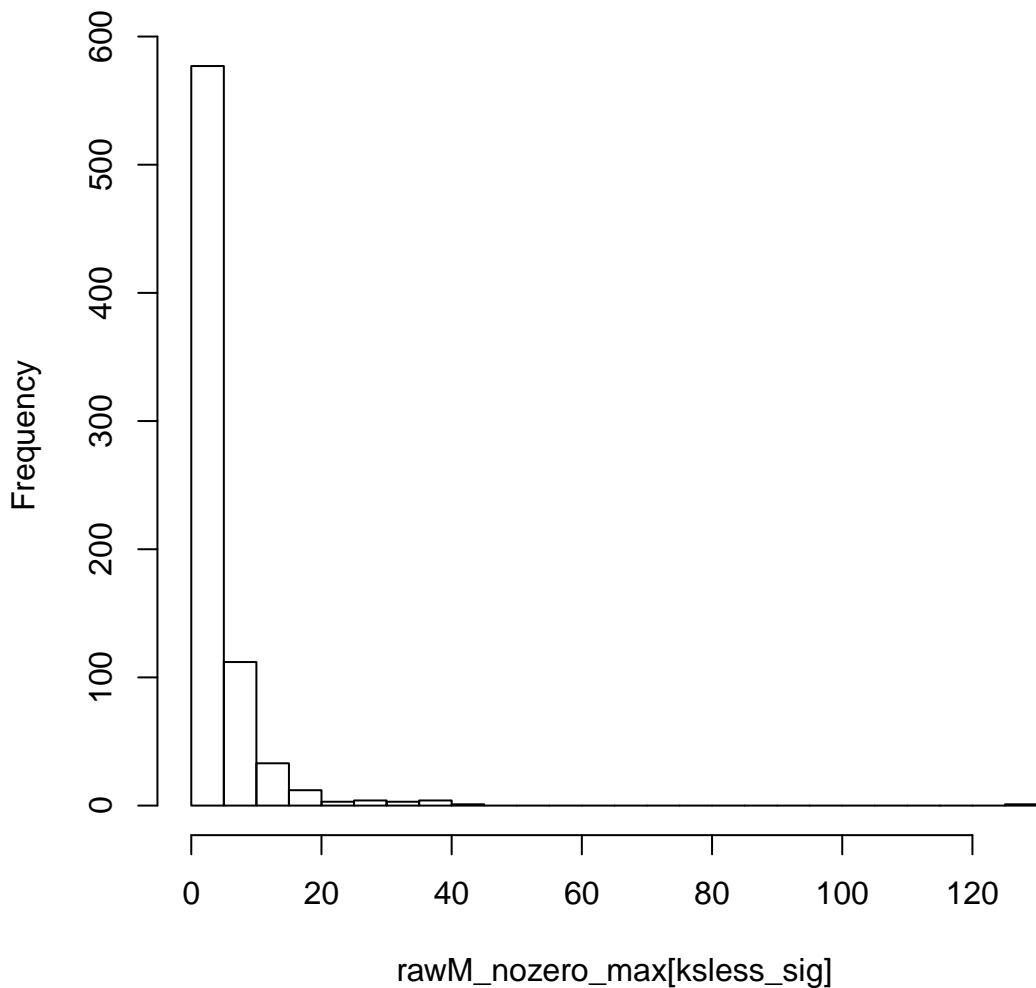


# max of nozero log-express of genes, kstvoside sig

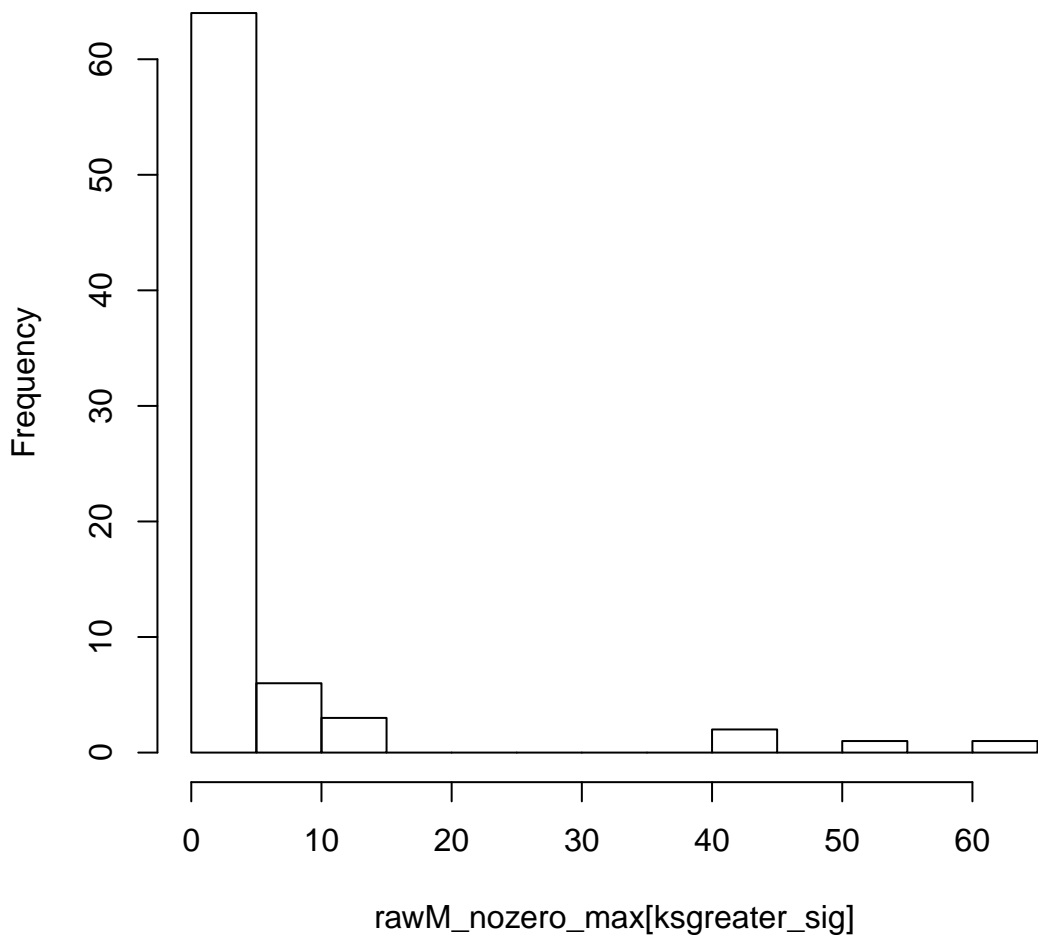




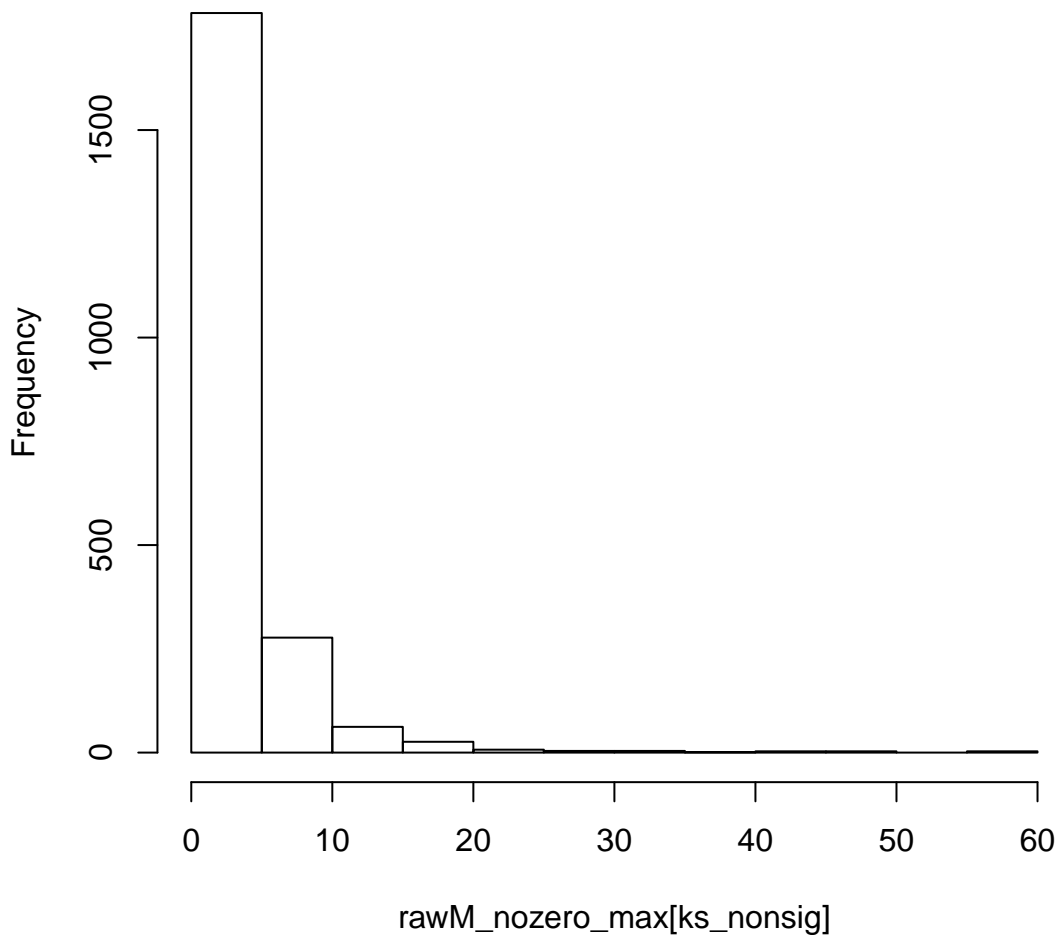
# max of nozero log-express of genes, ksless sig



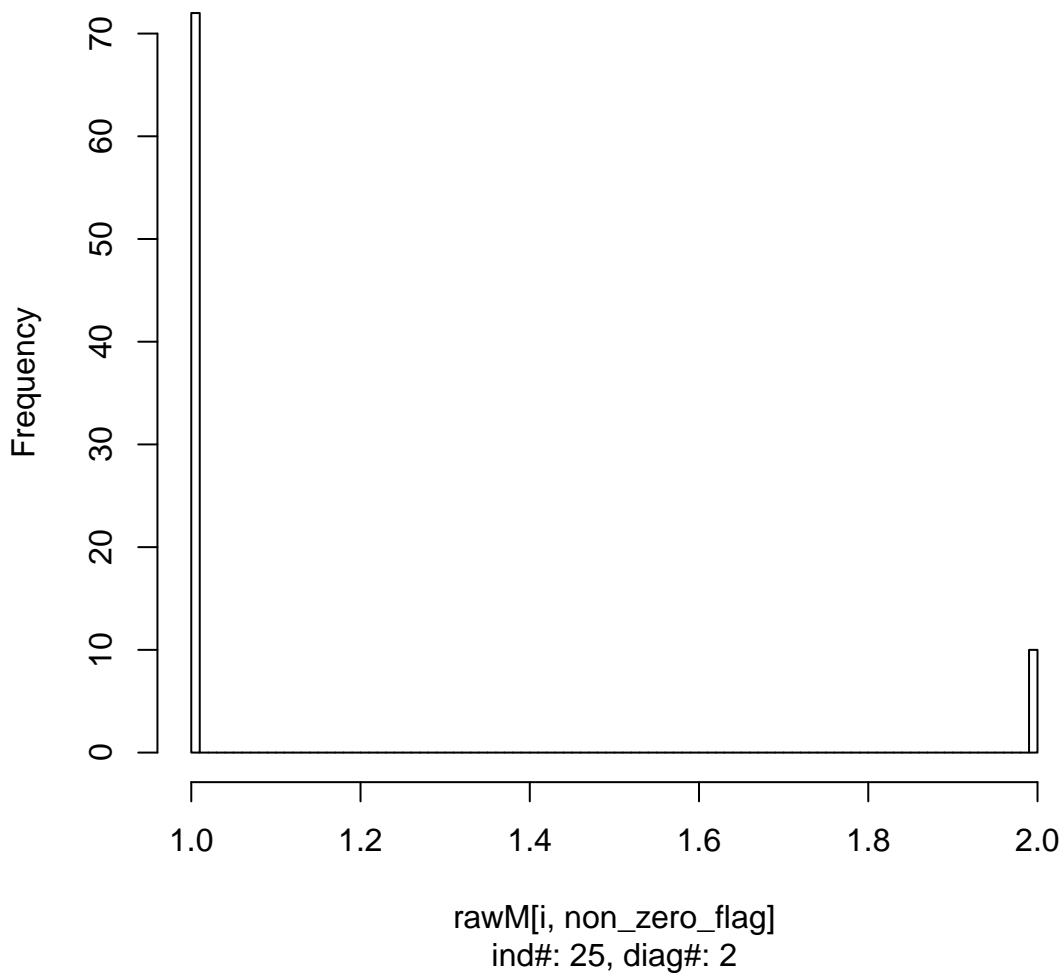
# max of nozero log-express of genes,ksgreater sig



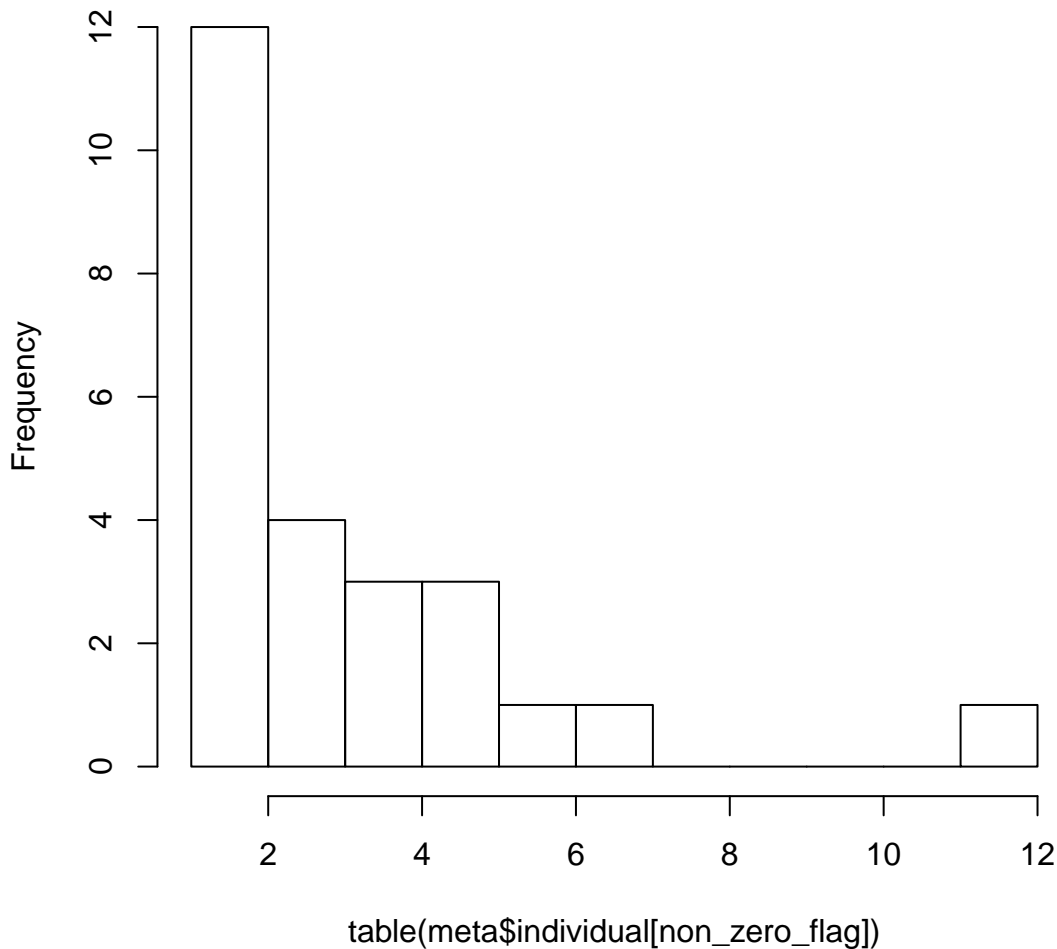
# max of nozero log-expres of genes,ks no sig



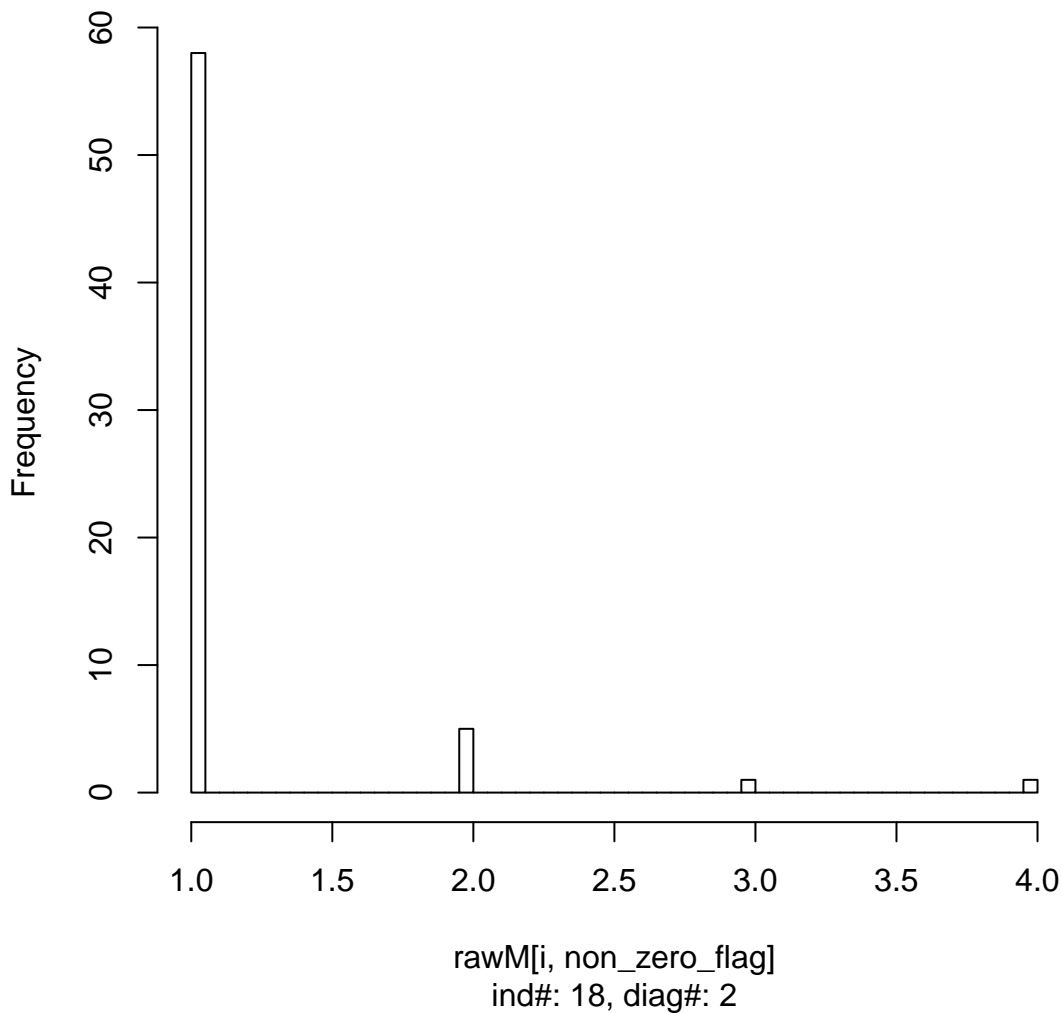
less sig: log expression of gene#3, pval ob=0.3323, non-zero nu



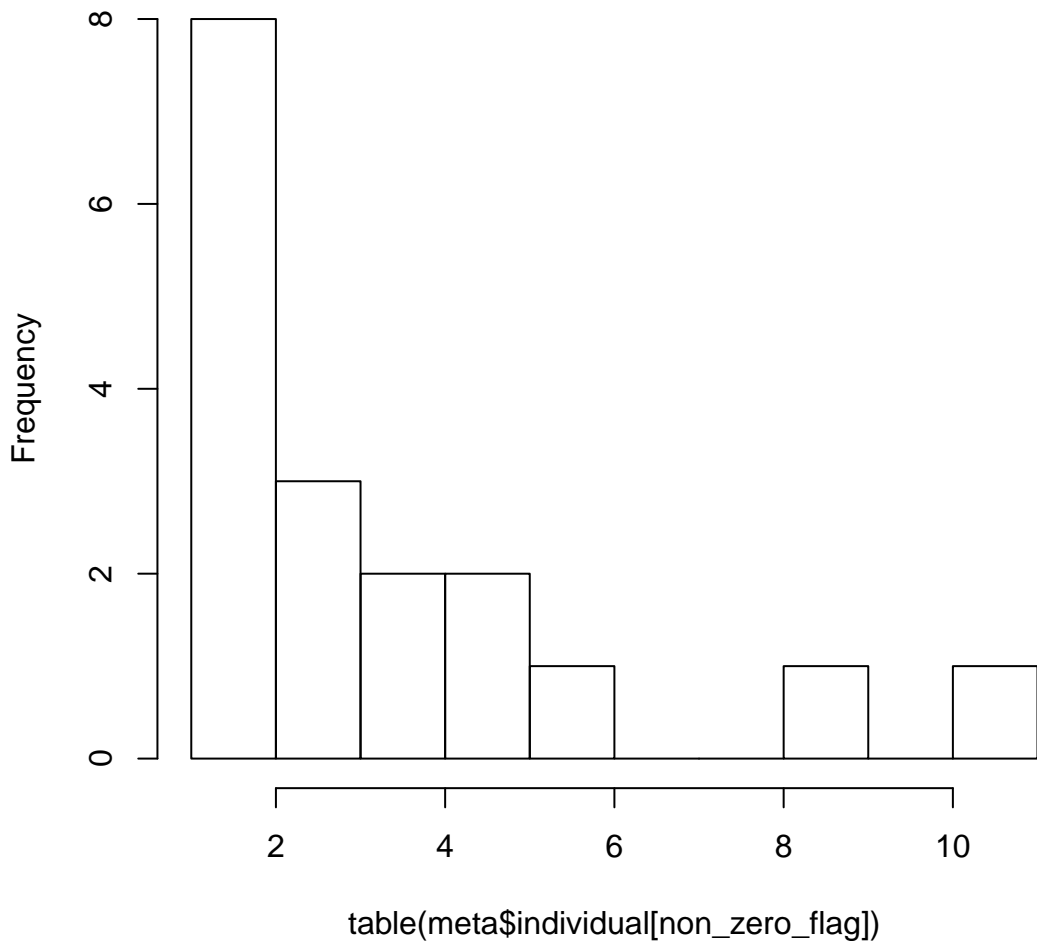
# KSless sig: individual expression cell count of gene#3



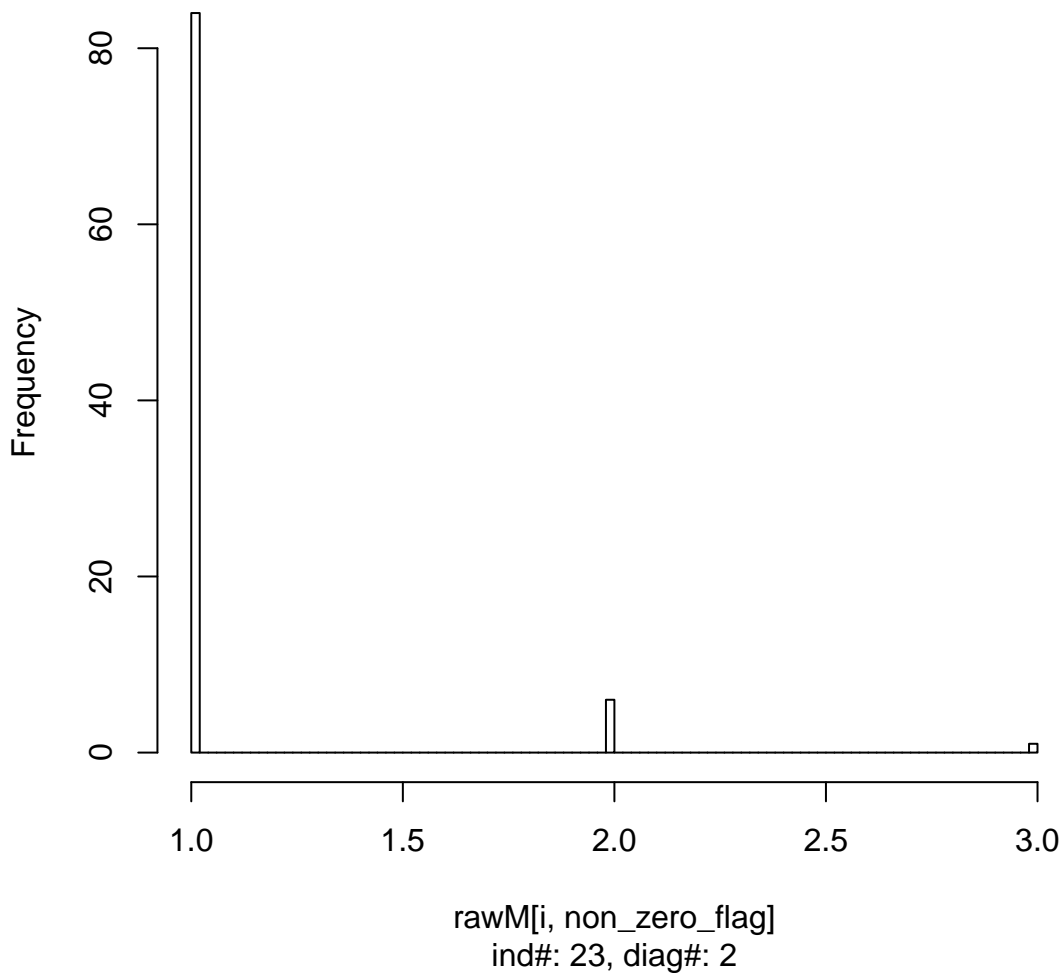
Sless sig: log expression of gene#5, pval ob=0.9099, non-zero nu



# KSless sig: individual expression cell count of gene#5

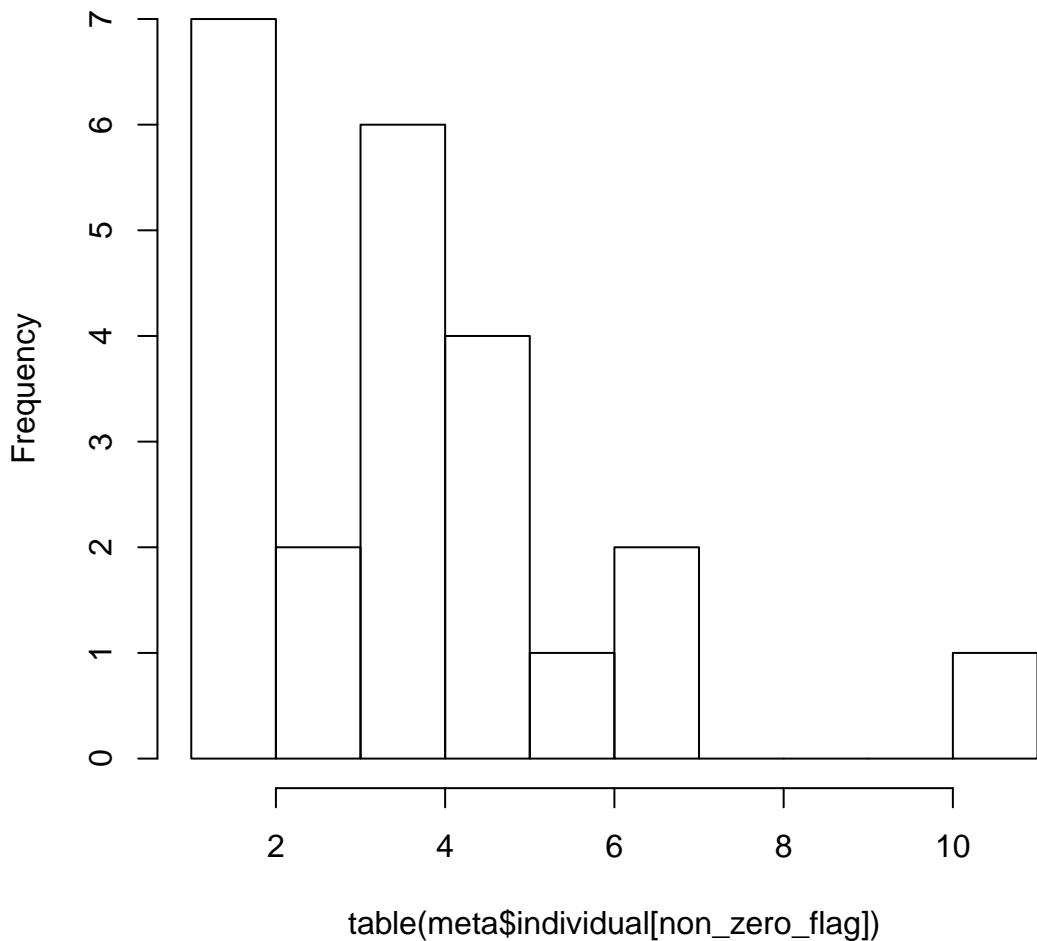


Less sig: log expression of gene#6, pval ob=0.6589, non-zero nu

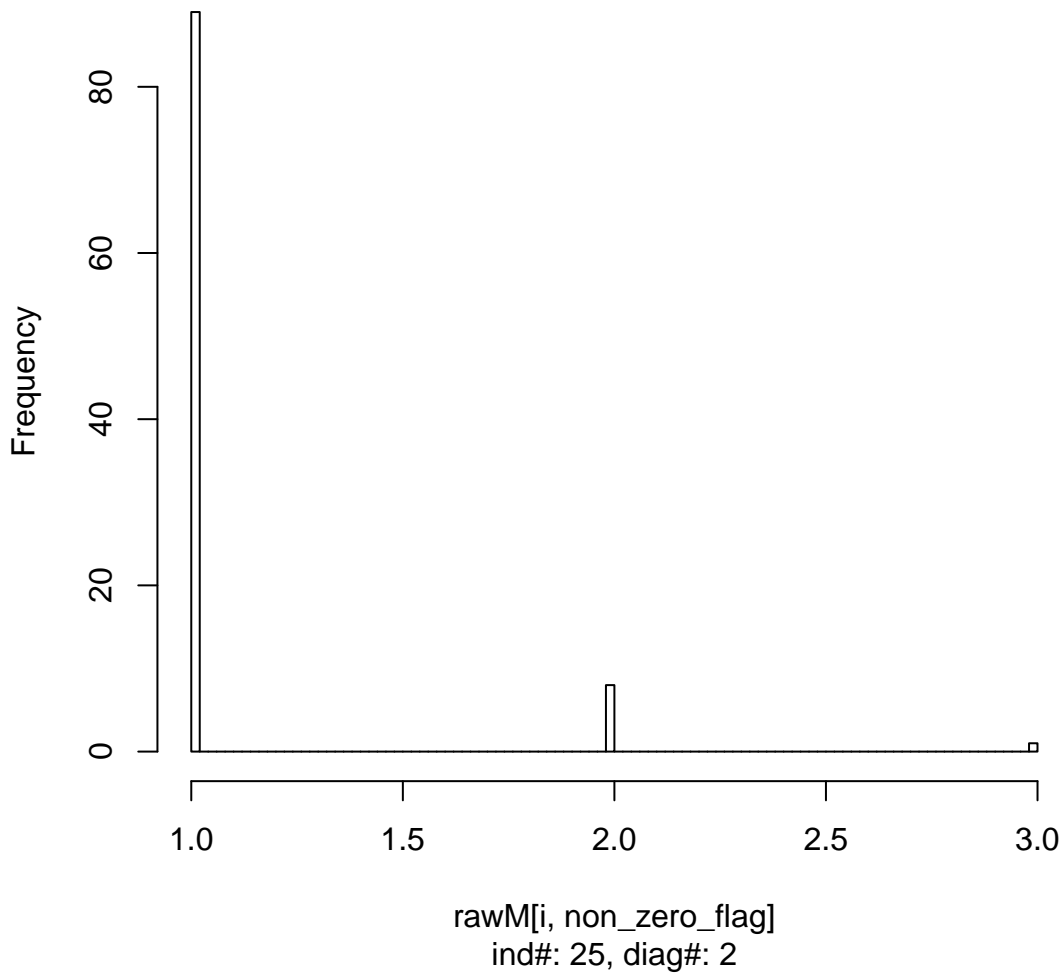




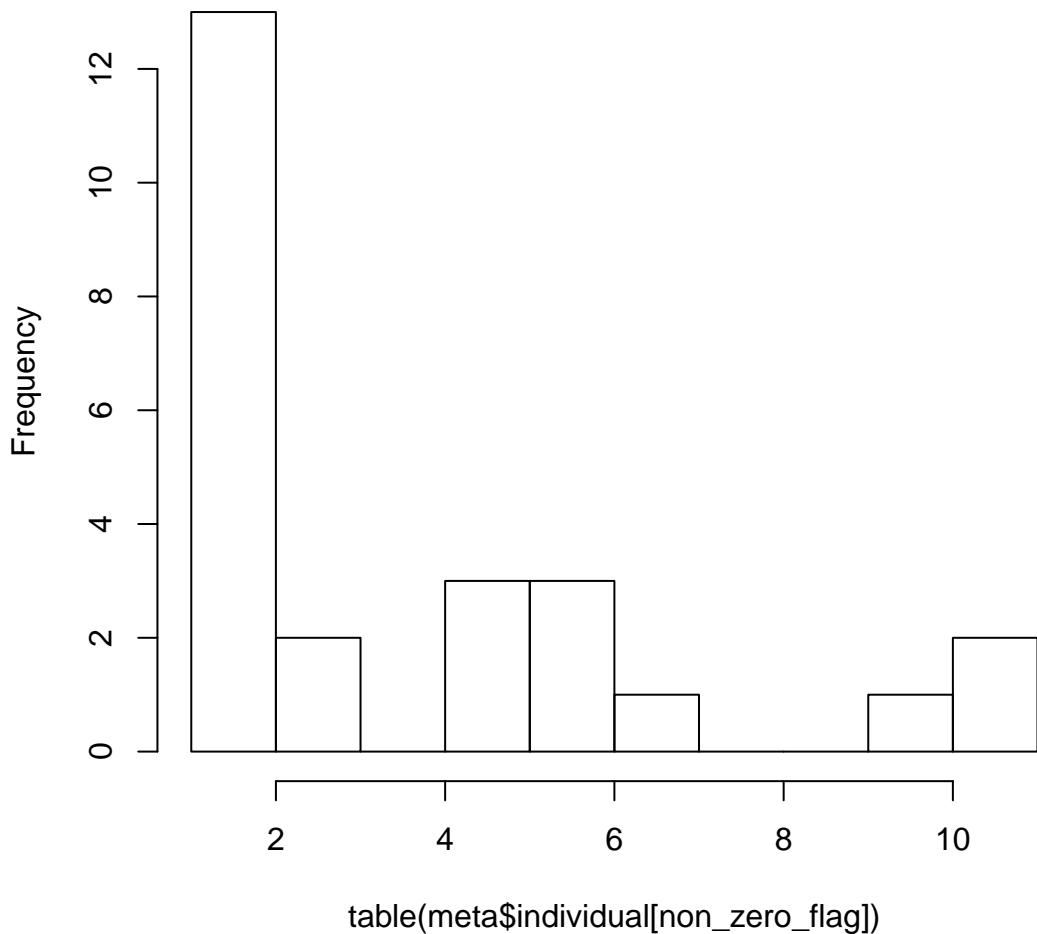
# KSless sig: individual expression cell count of gene#6



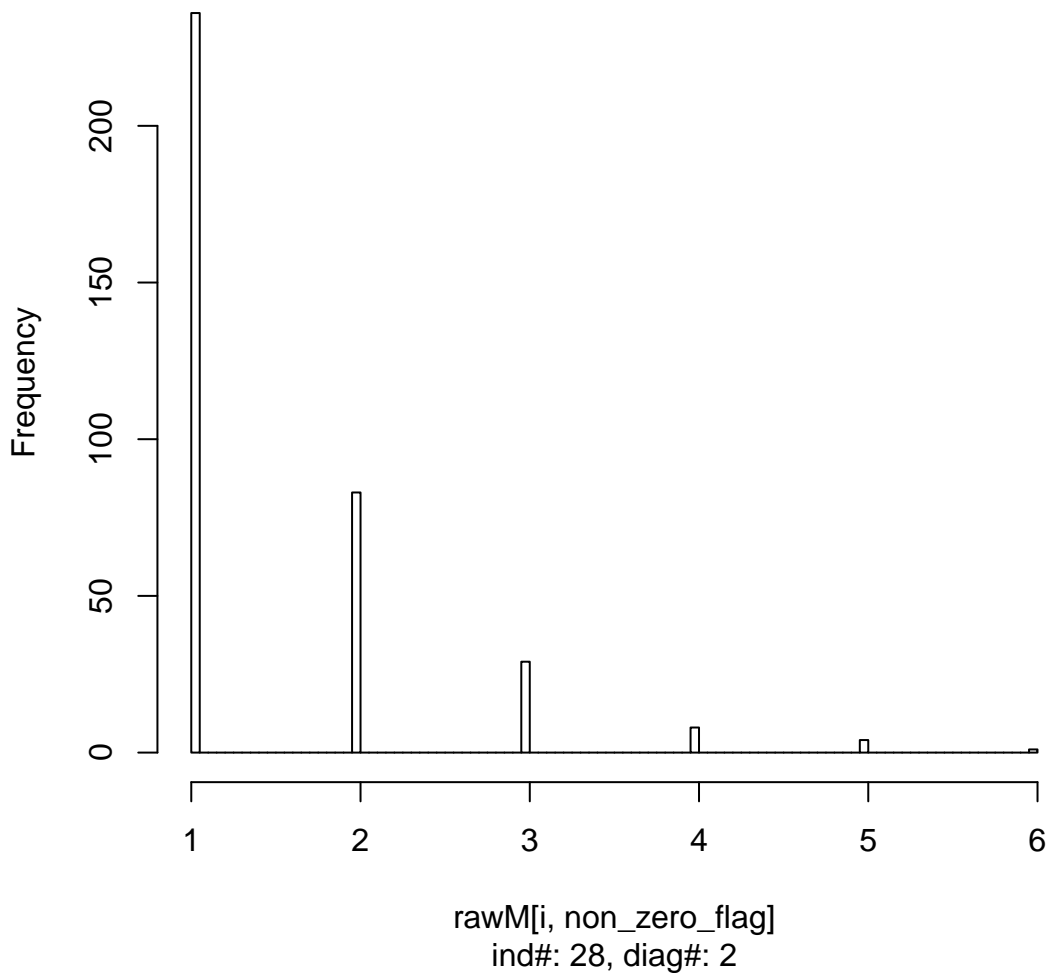
less sig: log expression of gene#15, pval ob=0.5012, non-zero n



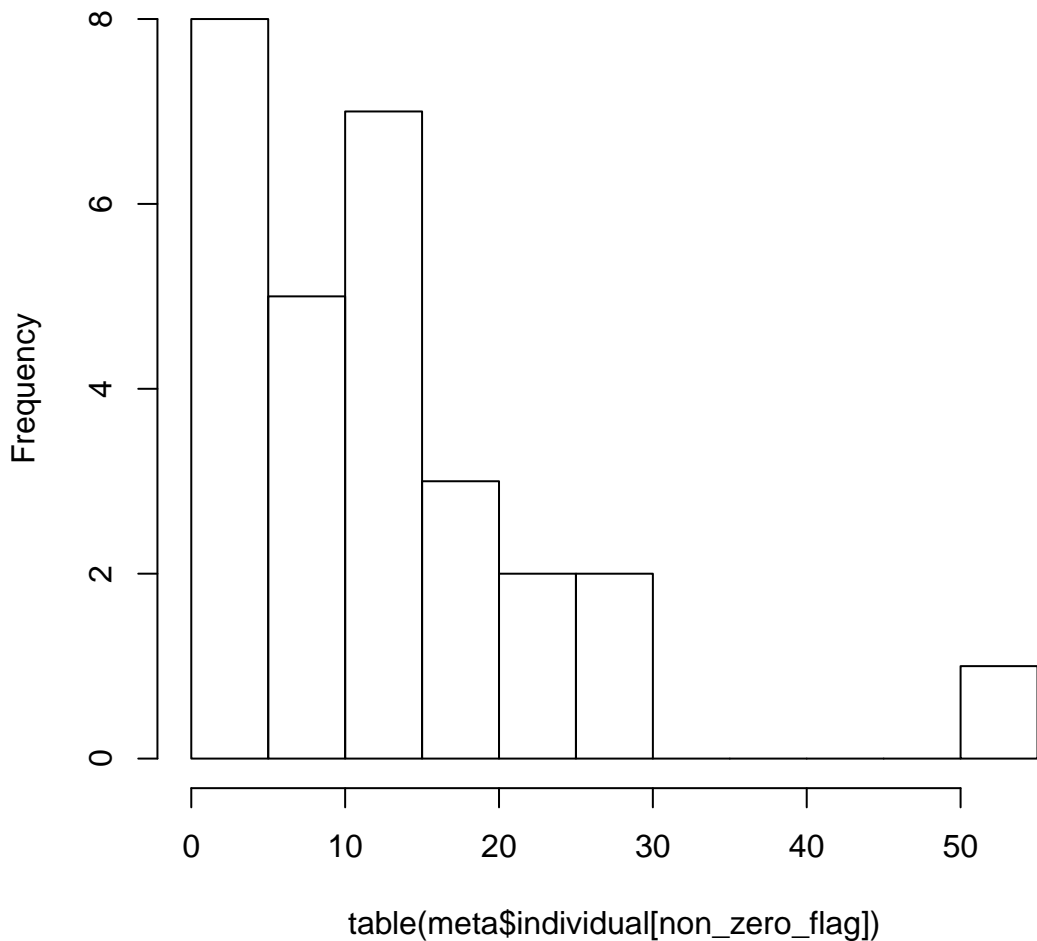
# KSless sig: individual expression cell count of gene#15



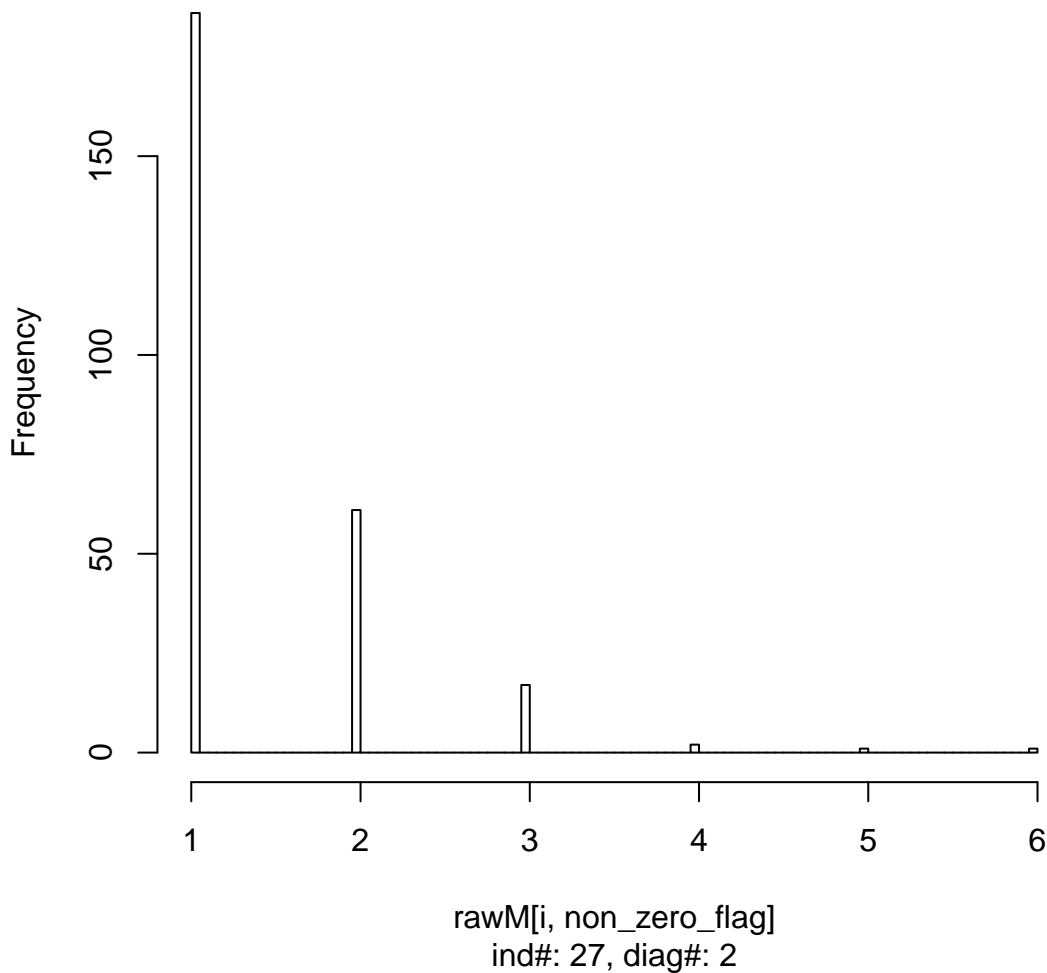
less sig: log expression of gene#30, pval ob=0.4617, non-zero nu



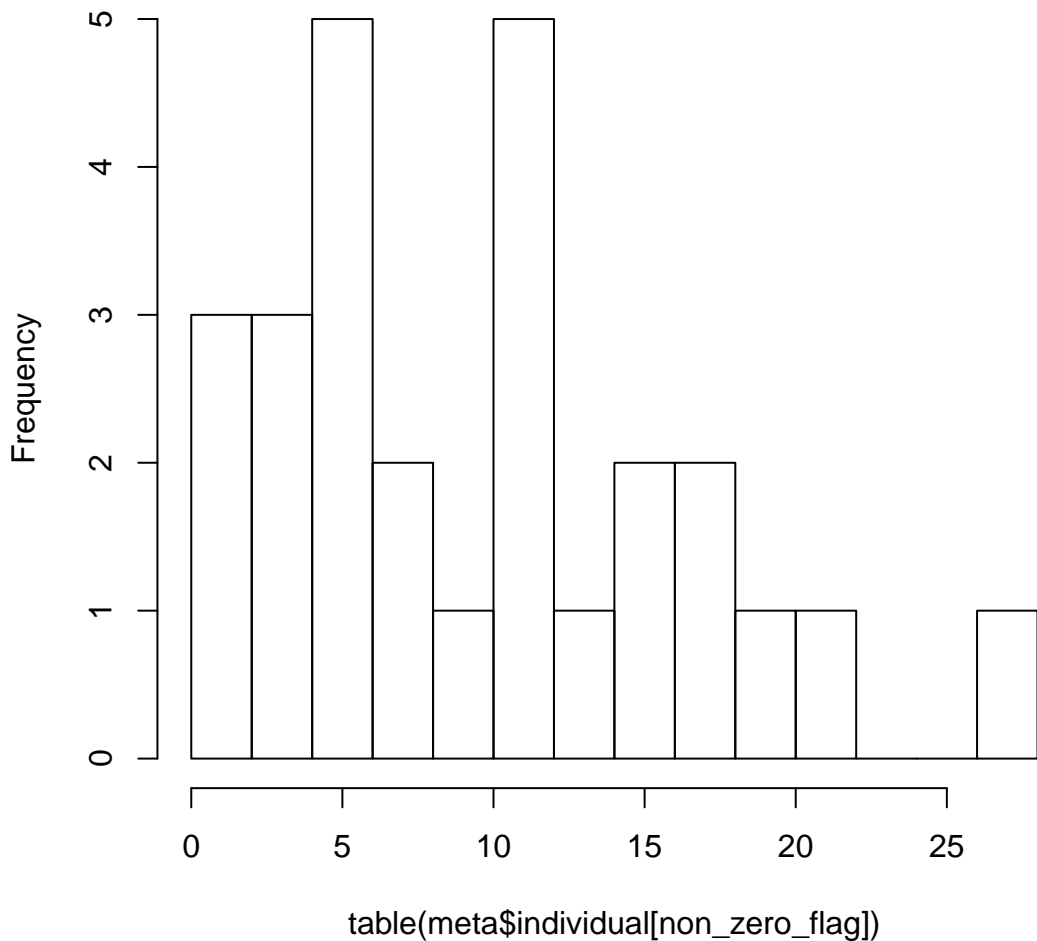
# KSless sig: individual expression cell count of gene#30



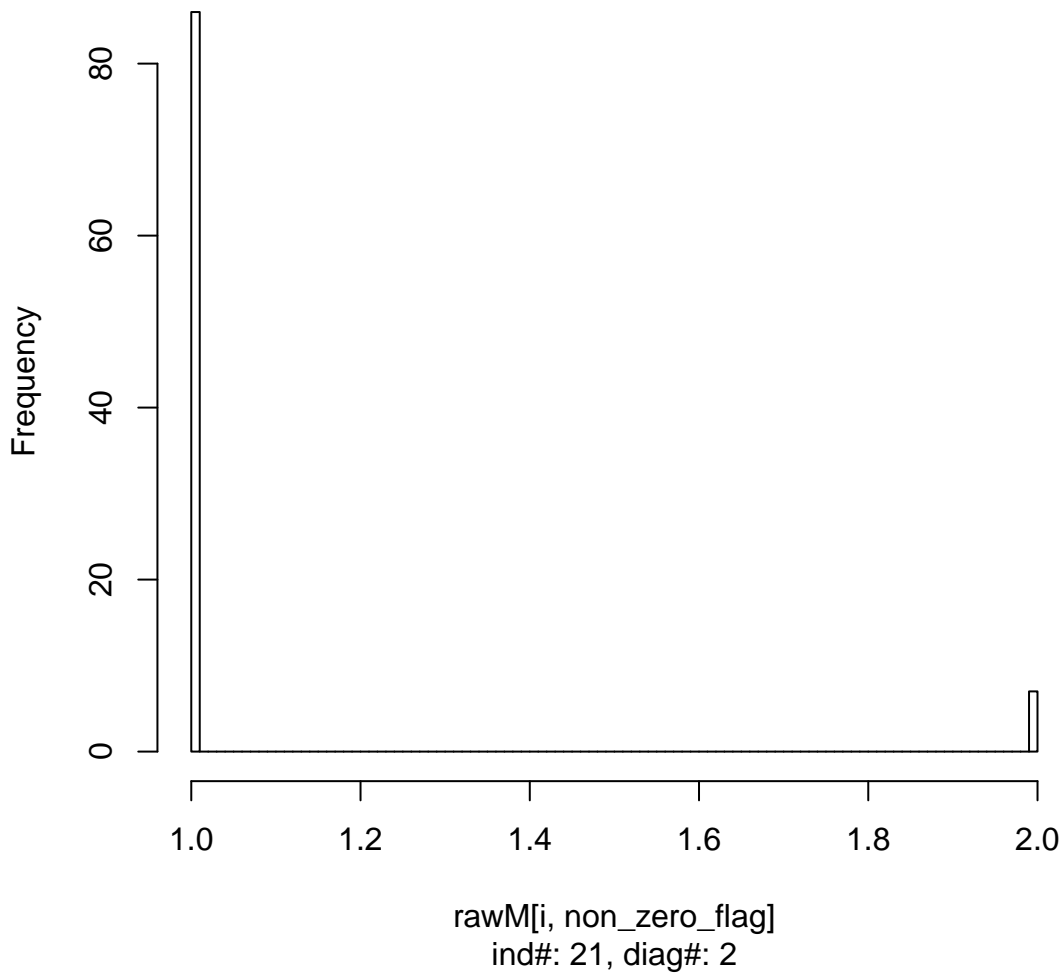
less sig: log expression of gene#31, pval ob=0.1334, non-zero nu



# KSless sig: individual expression cell count of gene#31

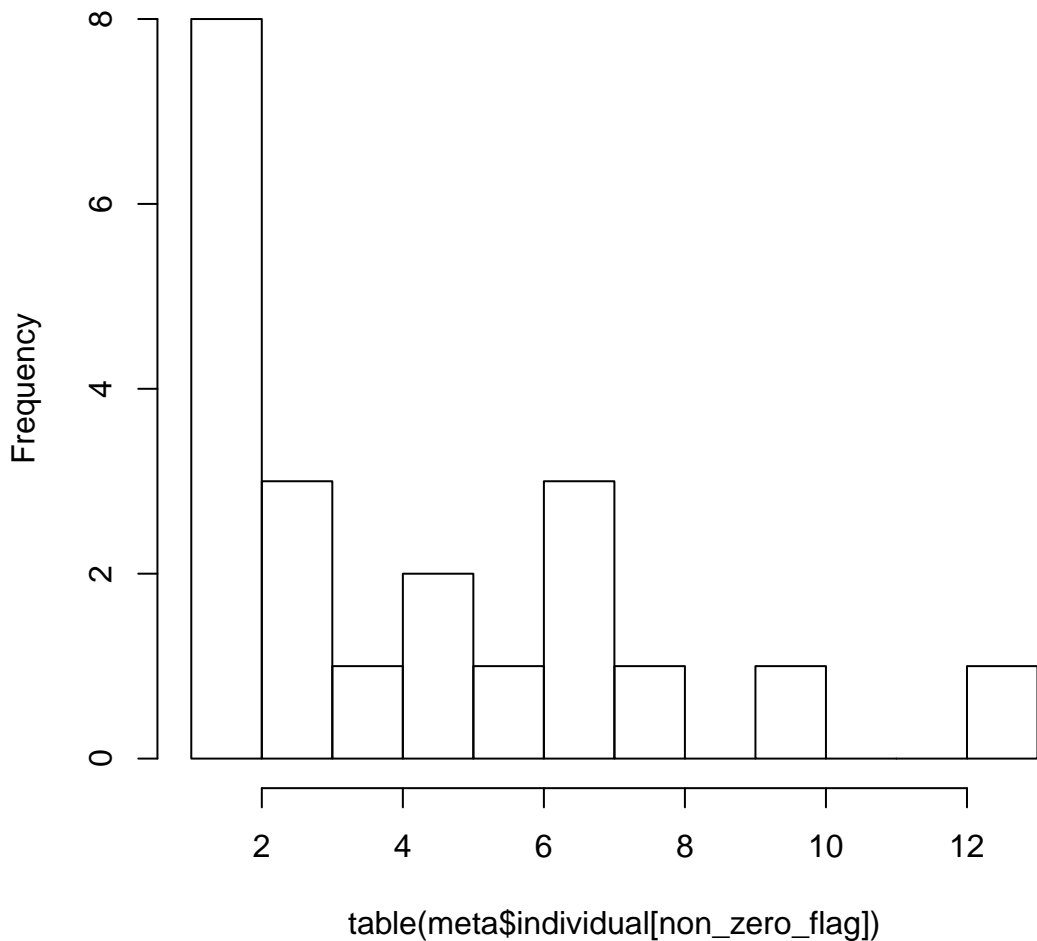


less sig: log expression of gene#35, pval ob=0.0013, non-zero n

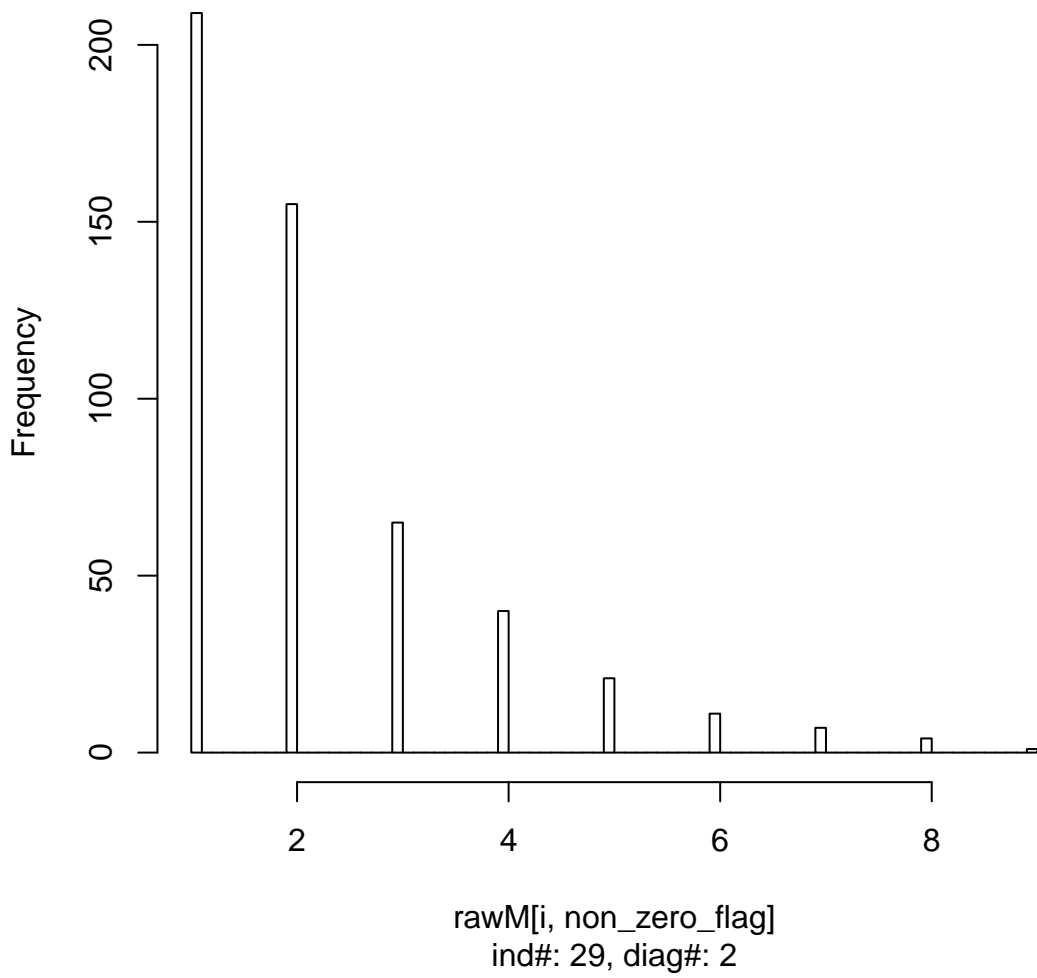




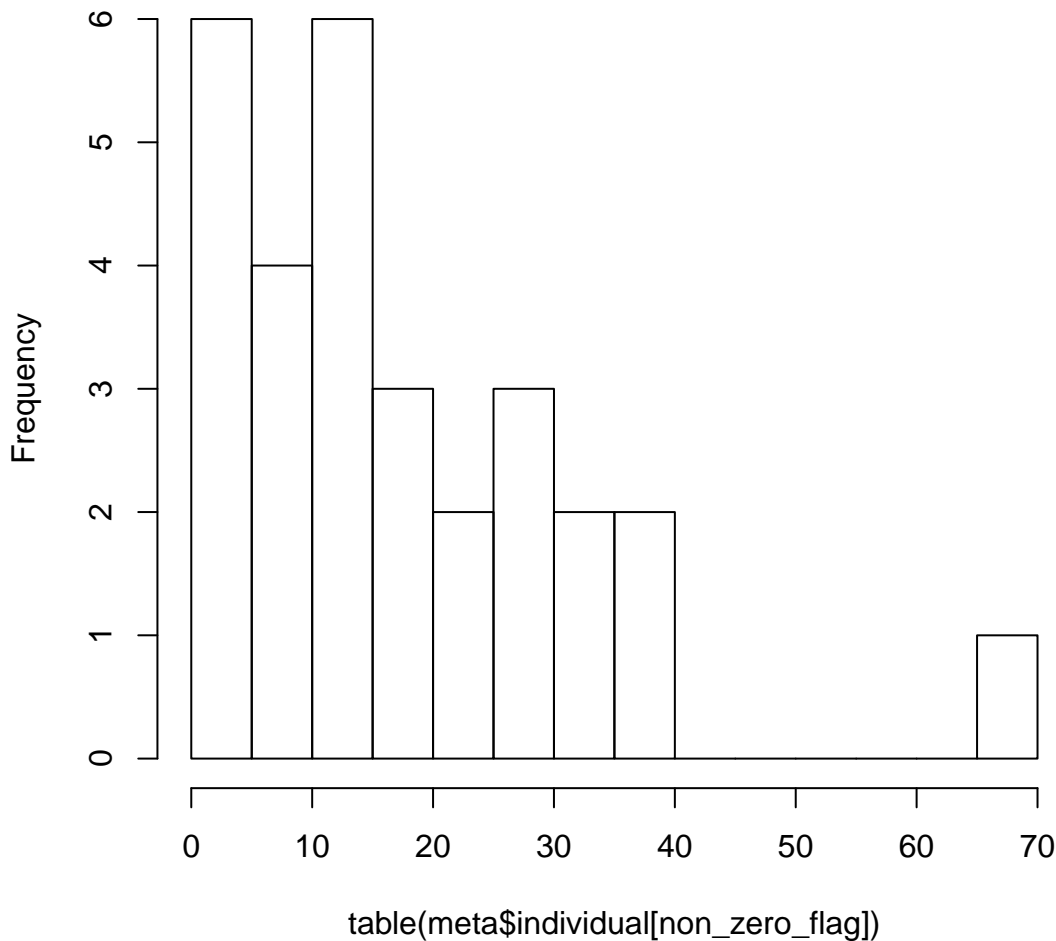
# KSless sig: individual expression cell count of gene#35



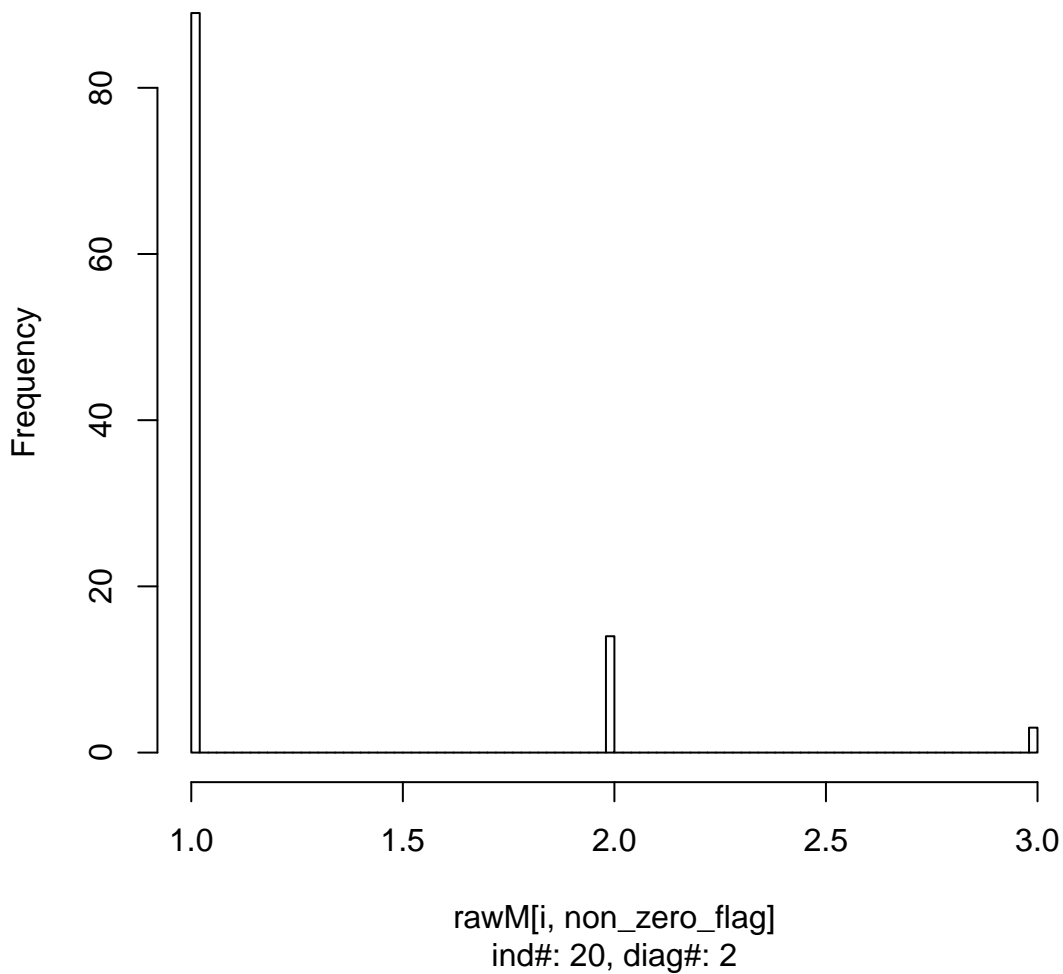
less sig: log expression of gene#48, pval ob=0.9311, non-zero nu



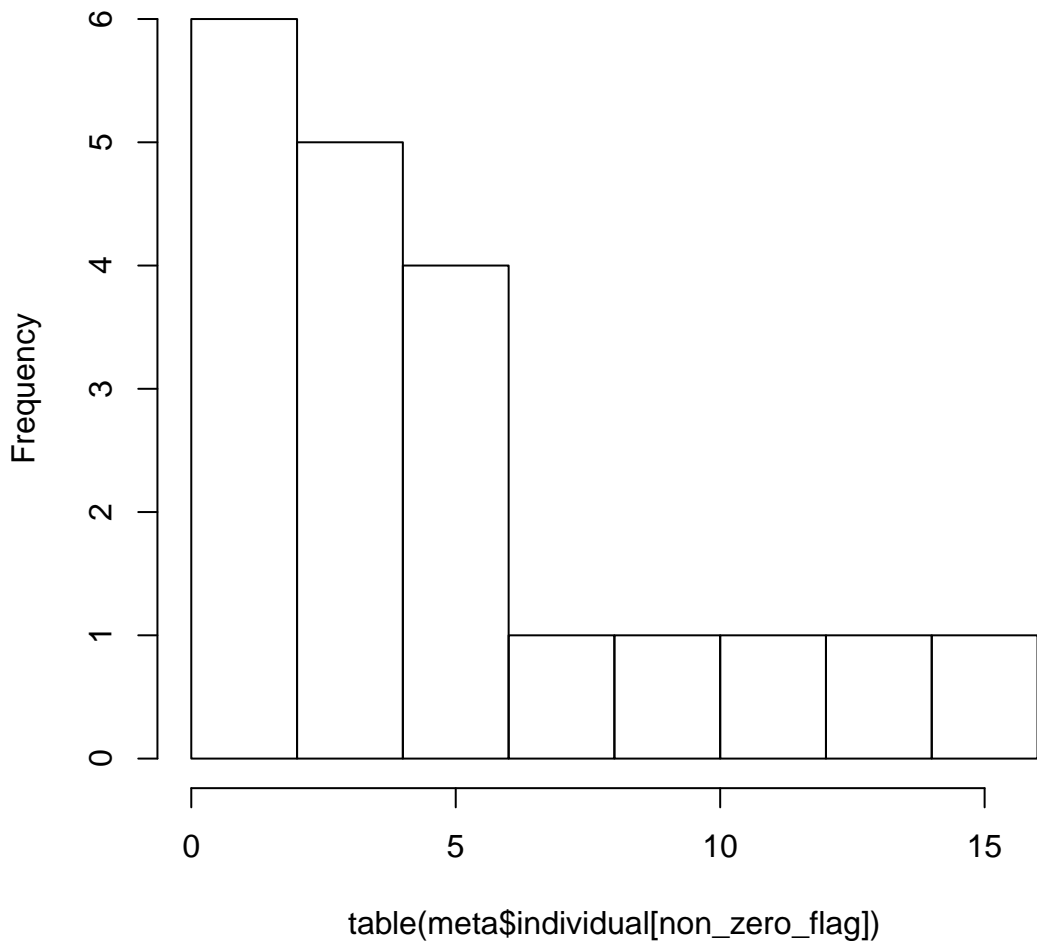
# KSless sig: individual expression cell count of gene#48



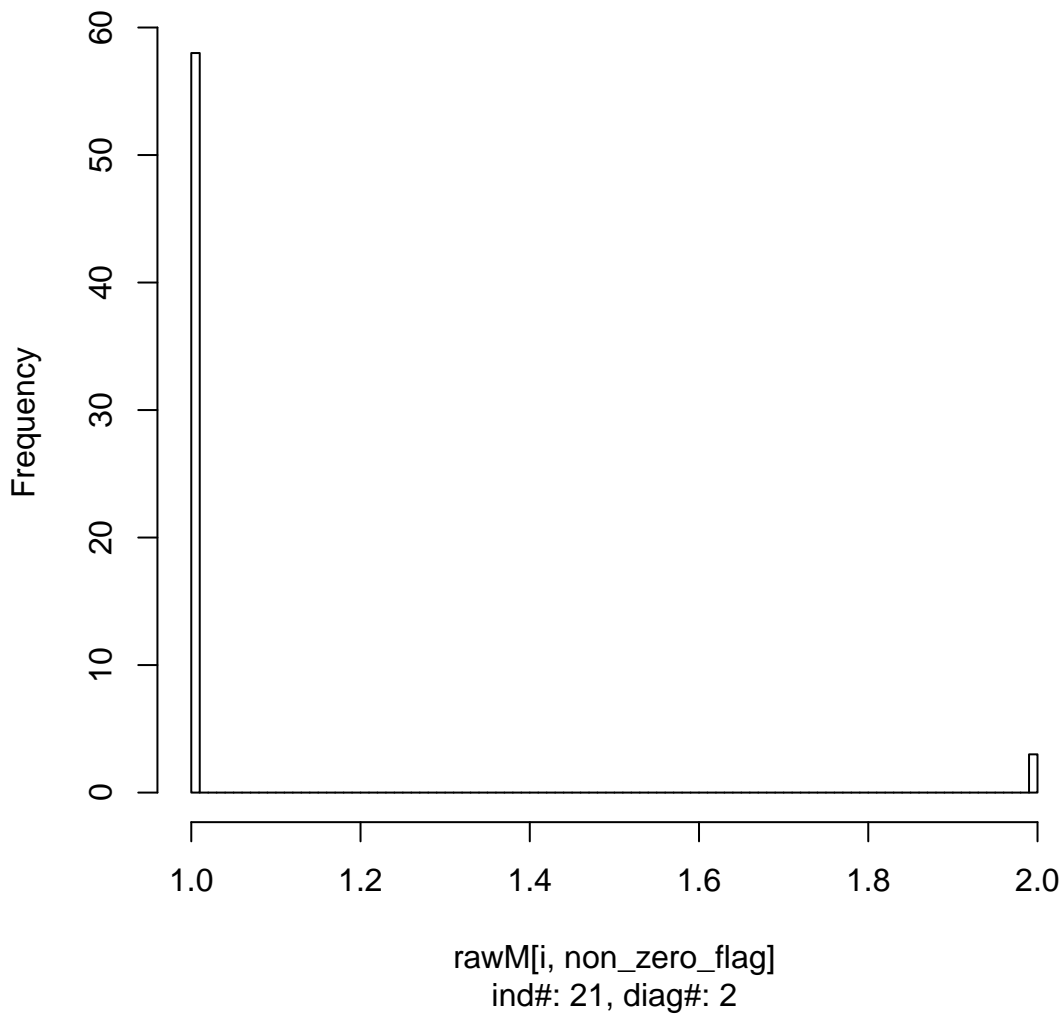
less sig: log expression of gene#49, pval ob=0.0201, non-zero nu



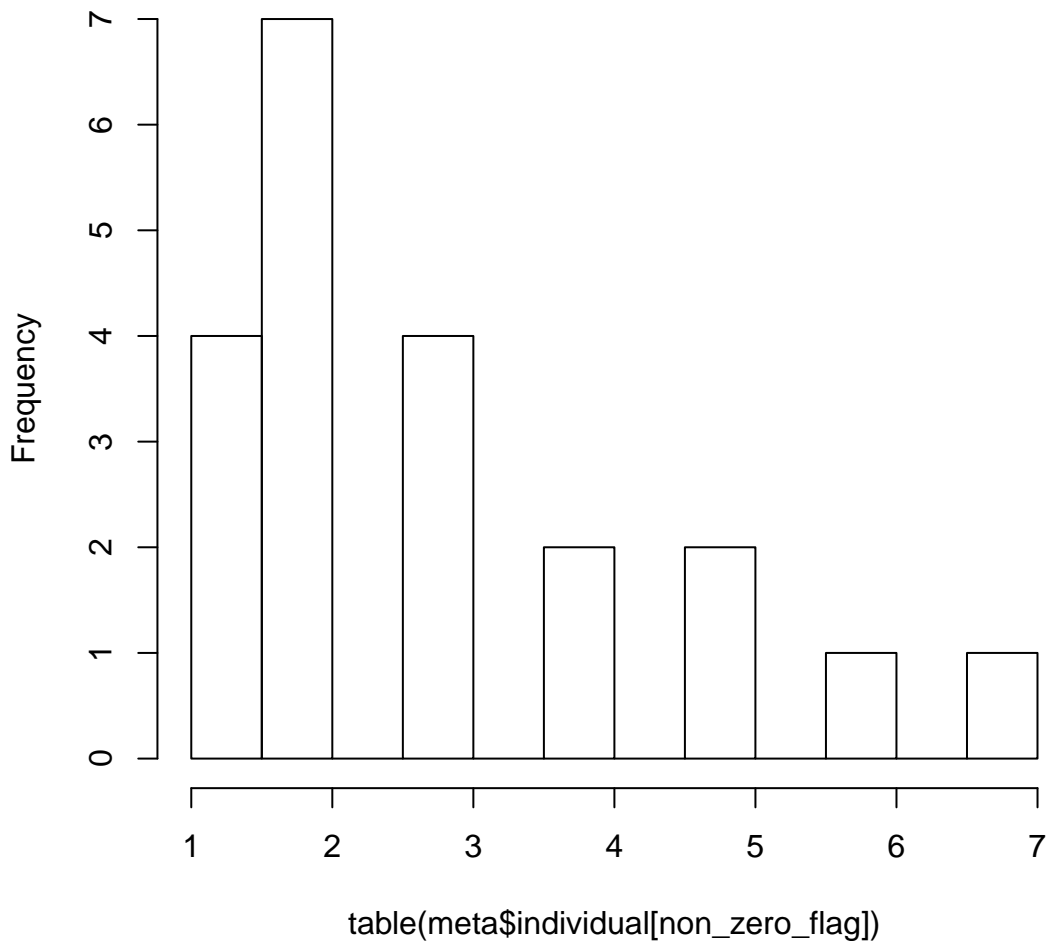
# KSless sig: individual expression cell count of gene#49



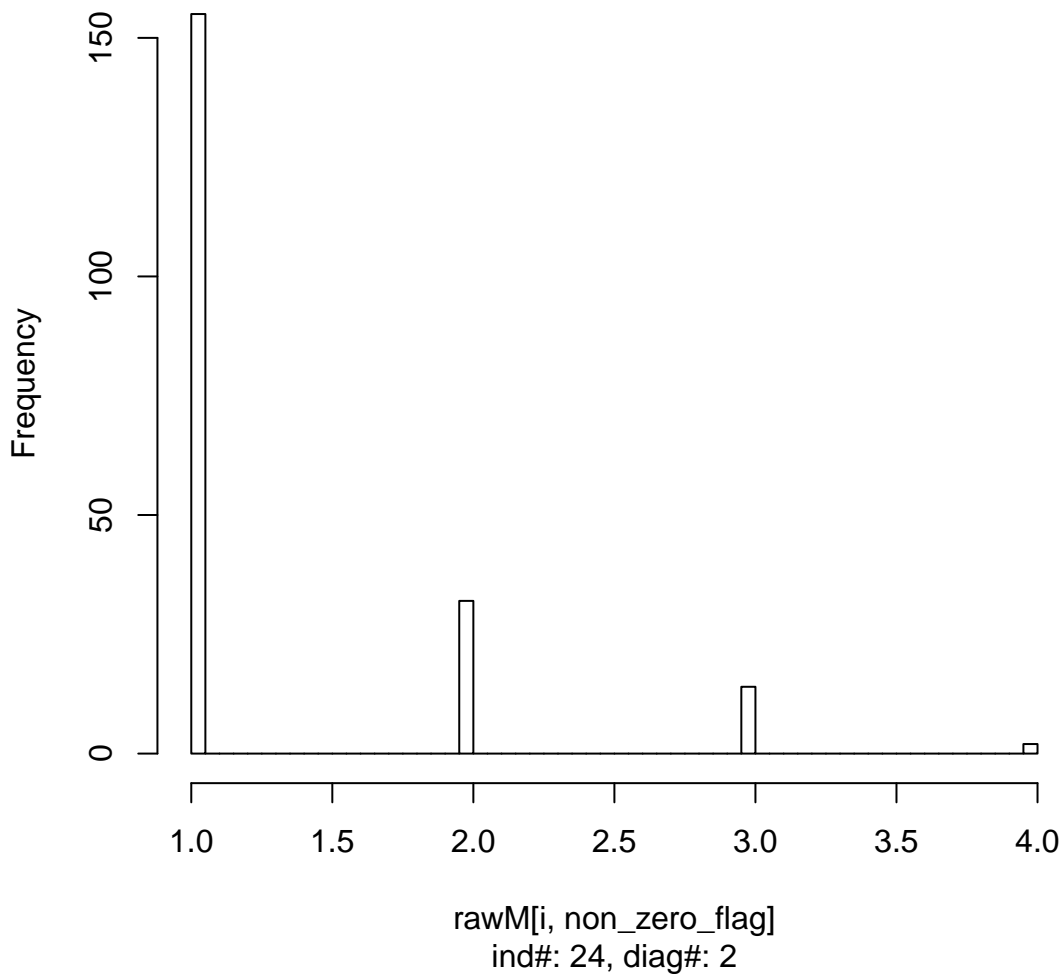
less sig: log expression of gene#56, pval ob=0.5181, non-zero n



# KSless sig: individual expression cell count of gene#56

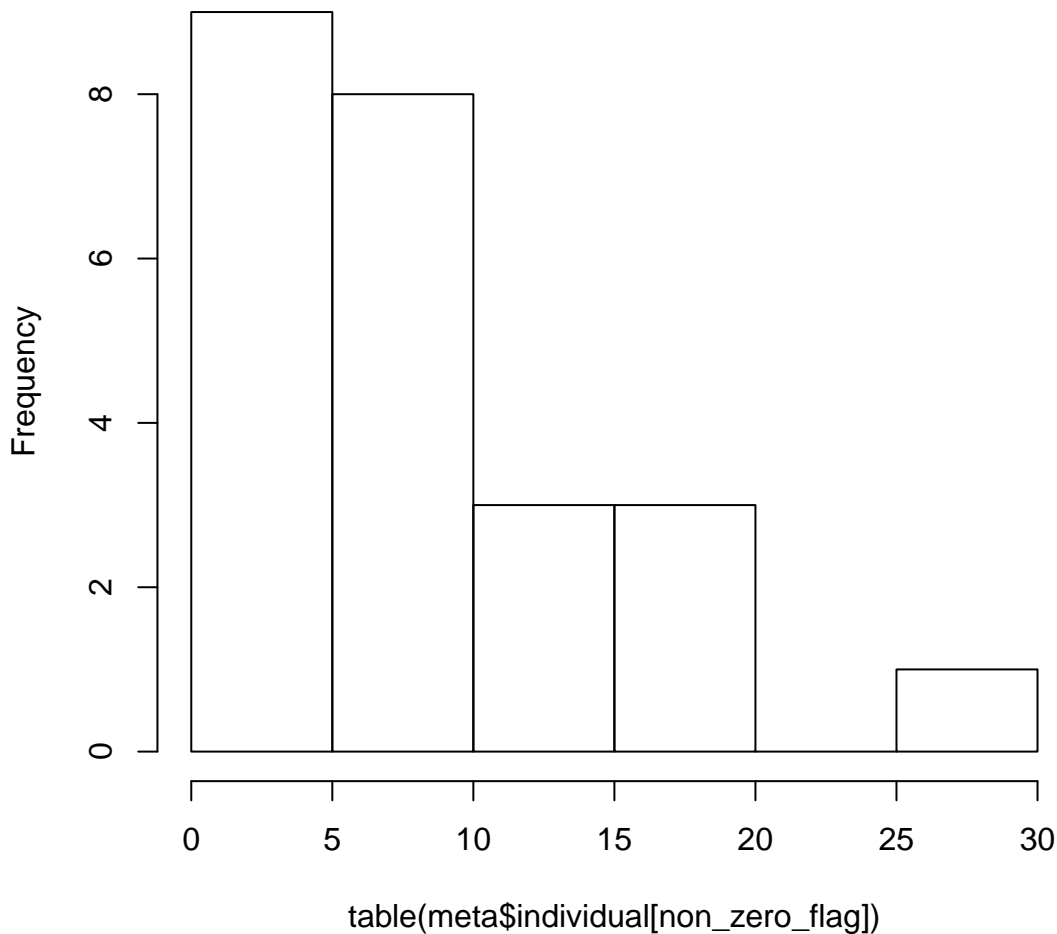


**S nonsig: log expression of gene#1, pval ob=0.8202, non-zero nu**

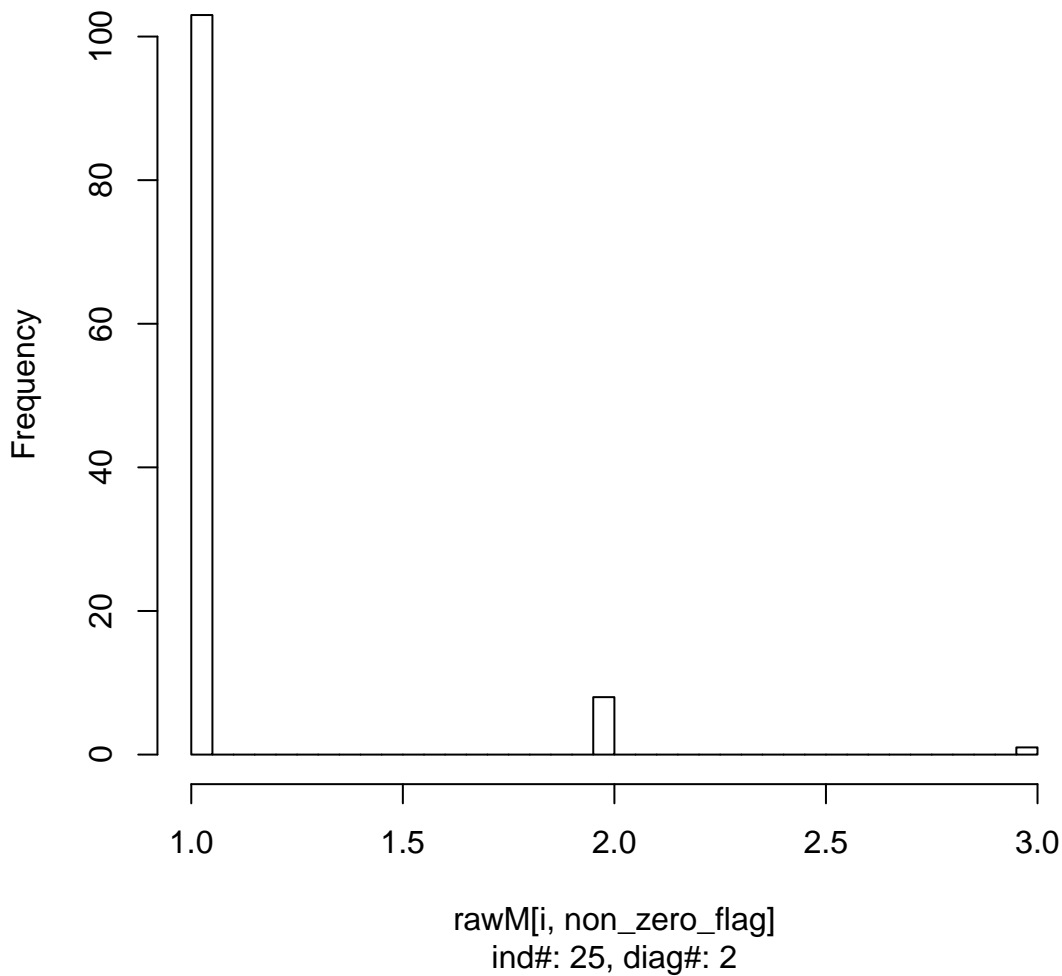




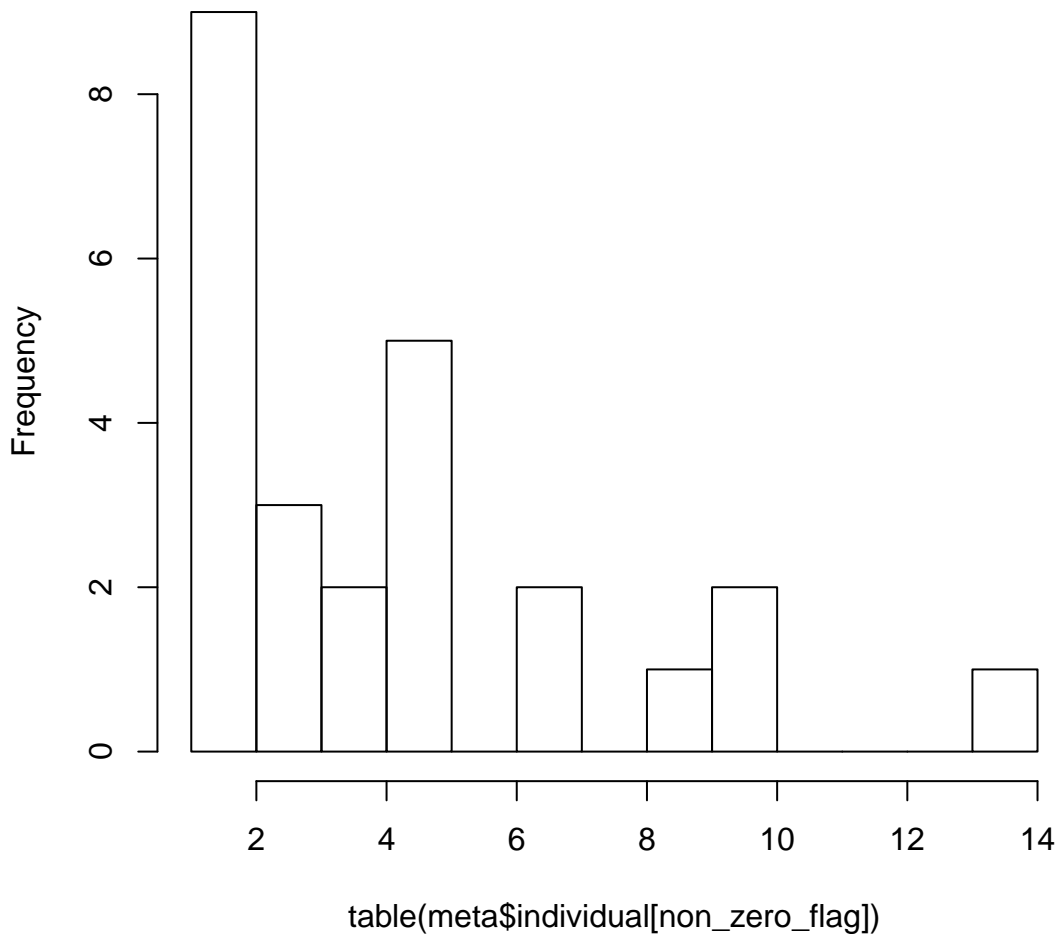
# KSless nonsig: individual expression cell count of gene#1



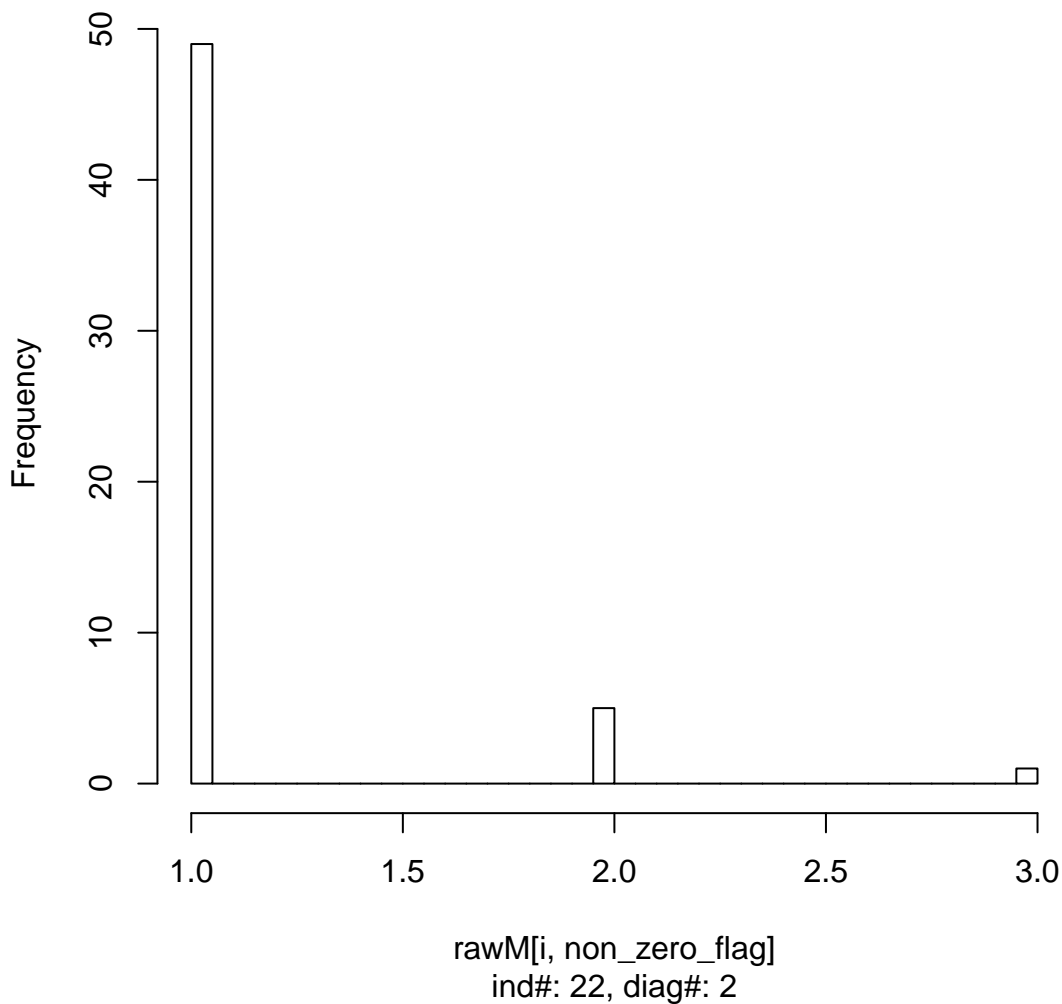
**S nonsig: log expression of gene#4, pval ob=0.4129, non-zero nu**



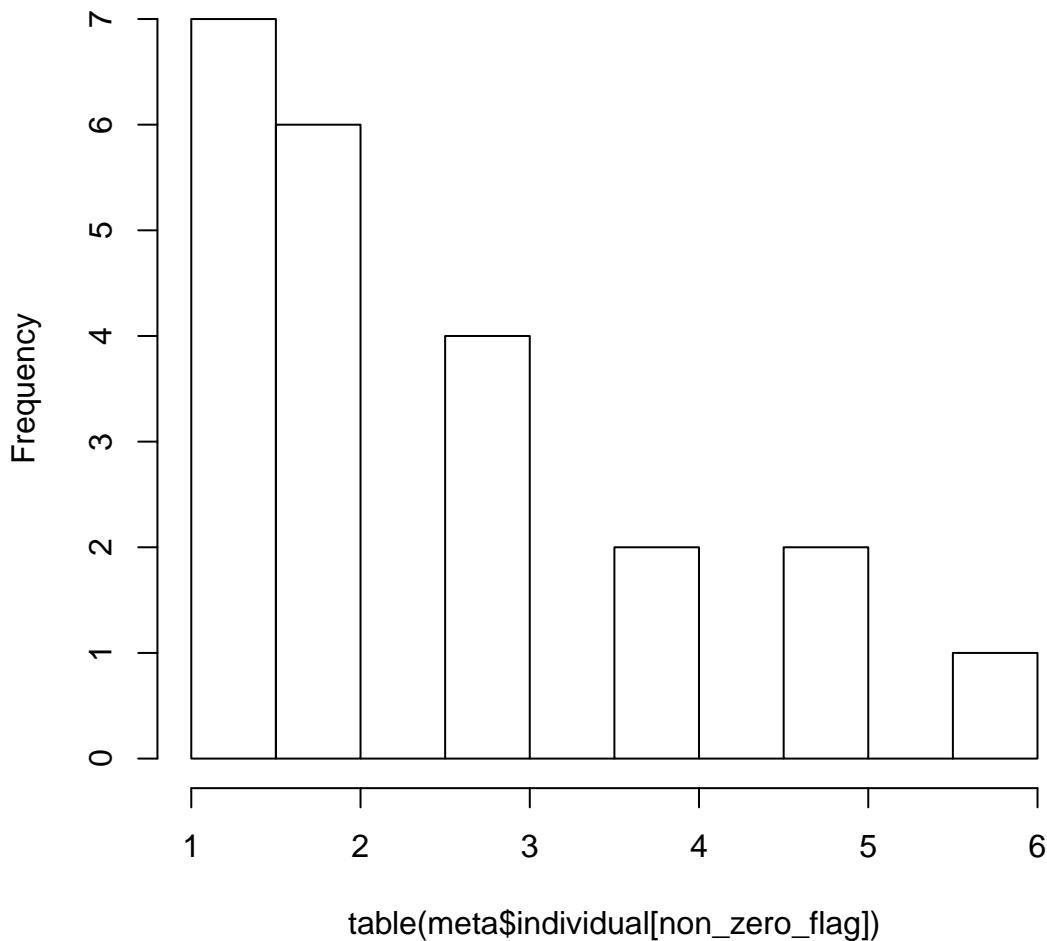
# KSless nonsig: individual expression cell count of gene#4



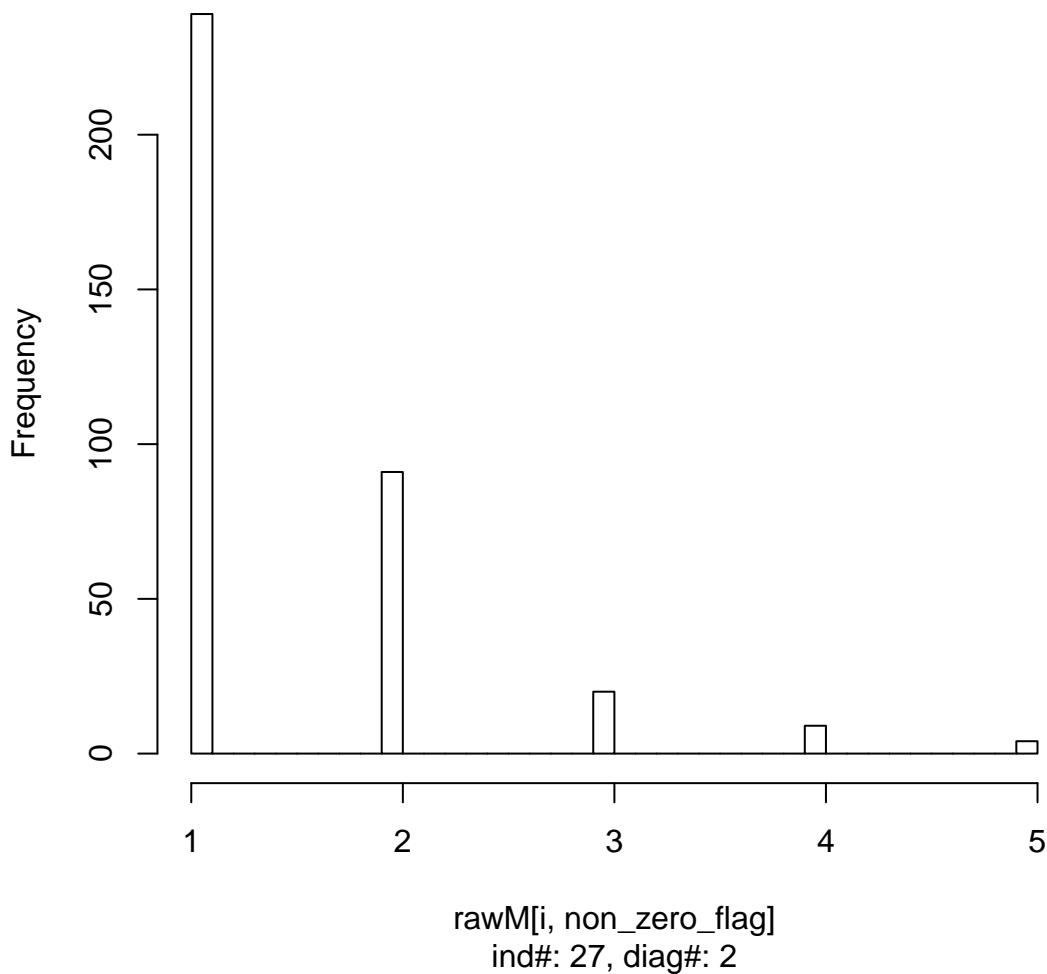
**S nonsig: log expression of gene#9, pval ob=0.7142, non-zero nu**



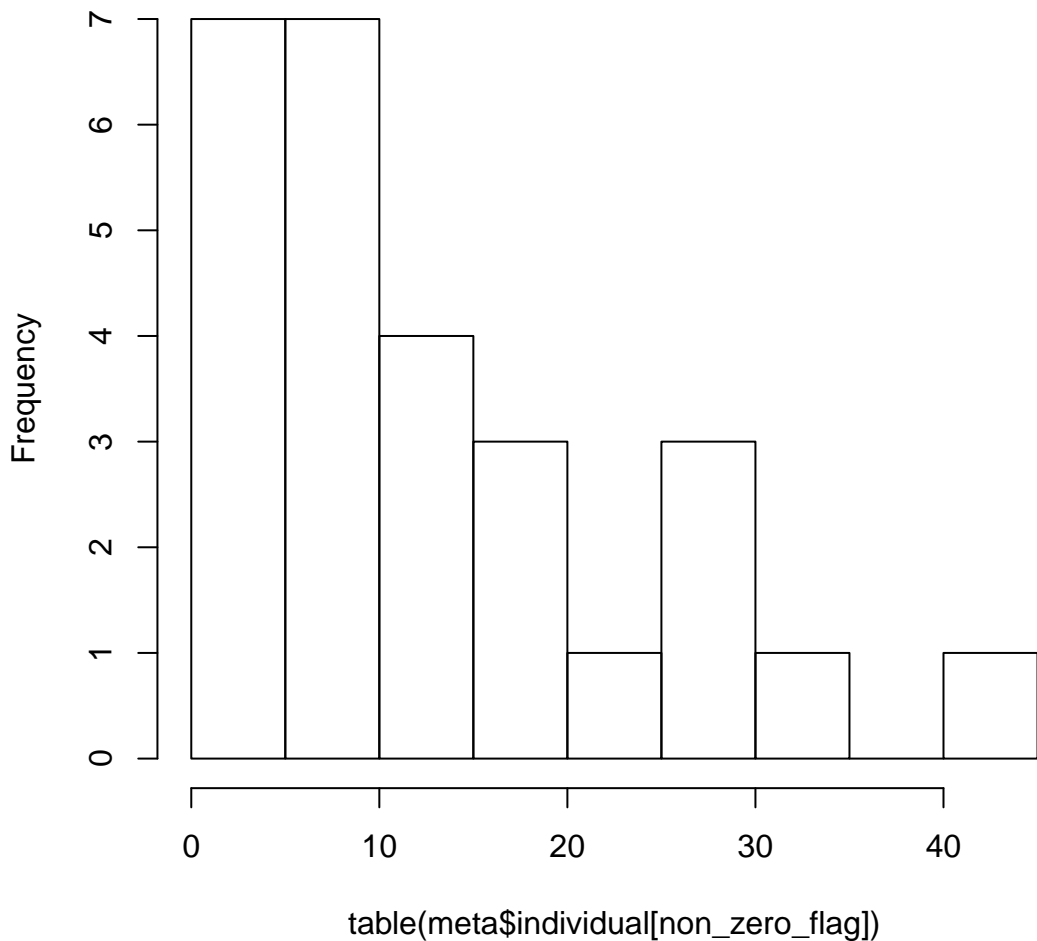
# KSless nonsig: individual expression cell count of gene#9



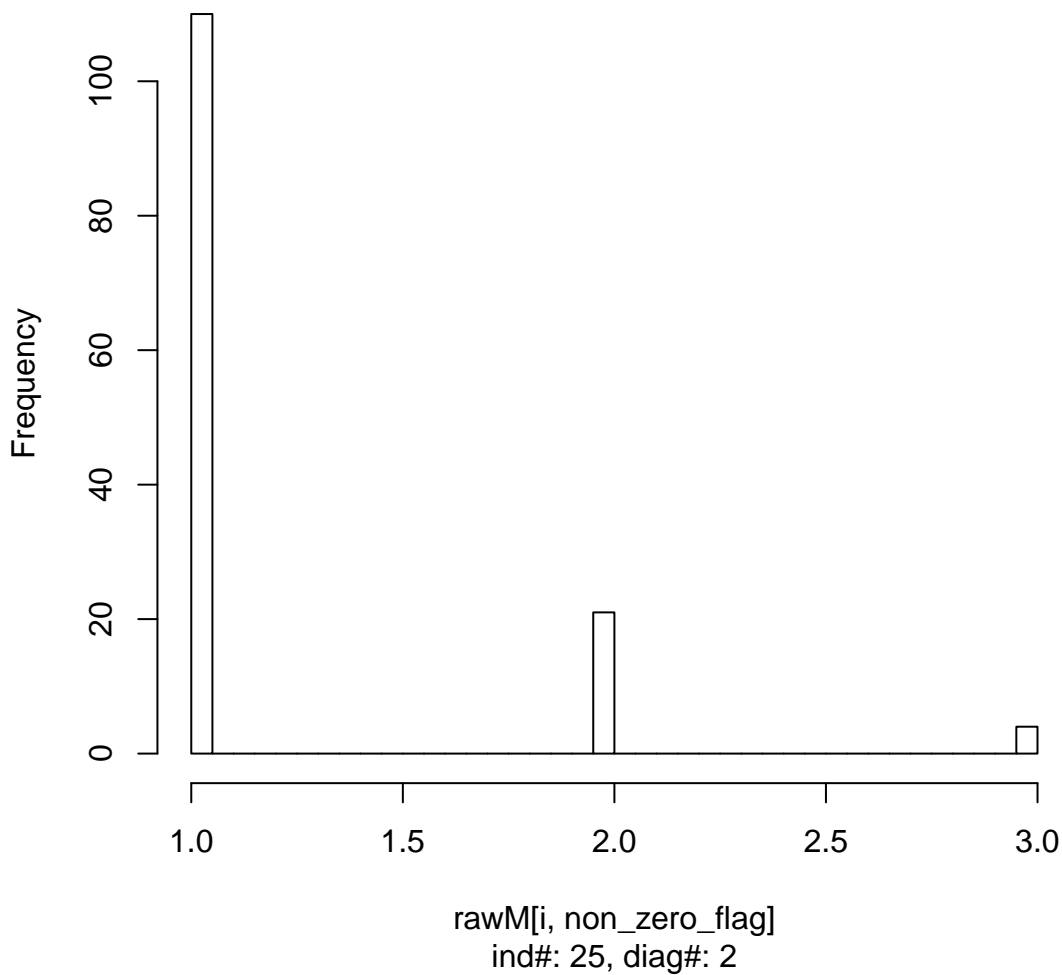
**nsig: log expression of gene#12, pval ob=0.3004, non-zero nu**



# KSless nonsig: individual expression cell count of gene#12

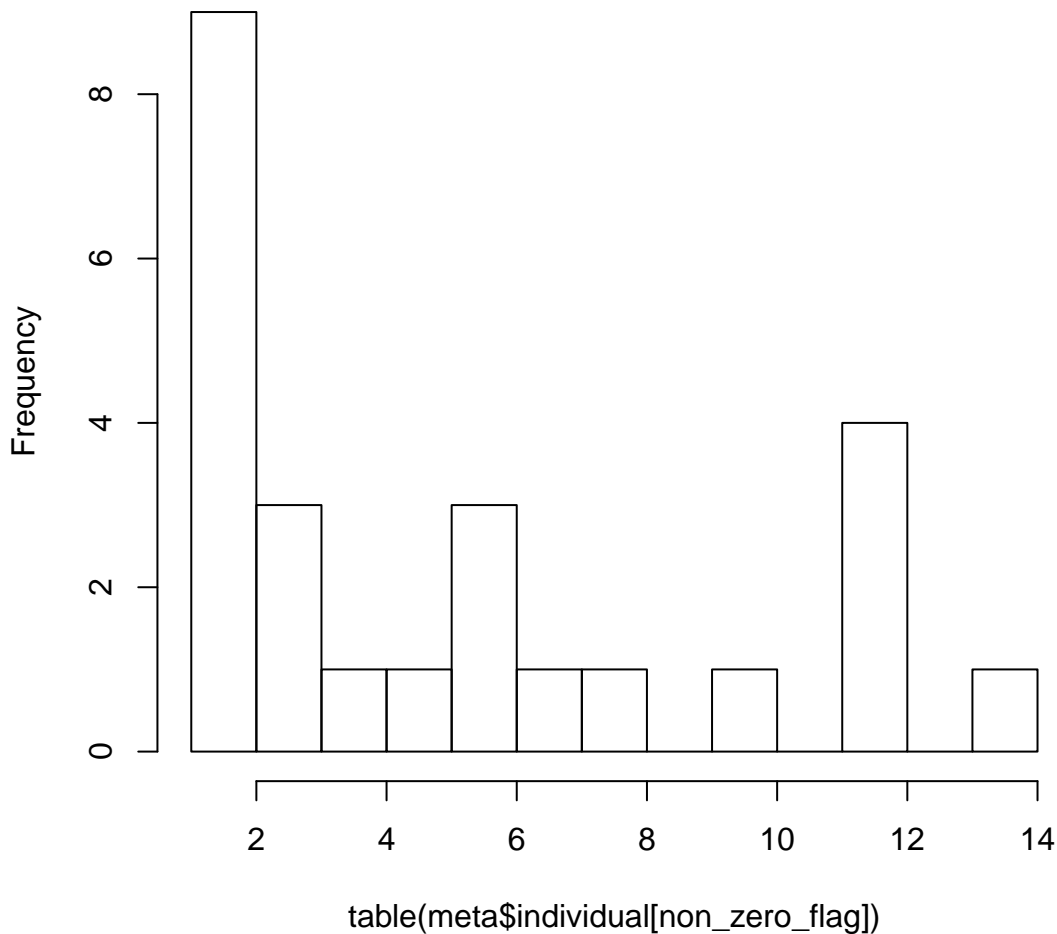


**ns nonsig: log expression of gene#14, pval ob=0.1191, non-zero nu**

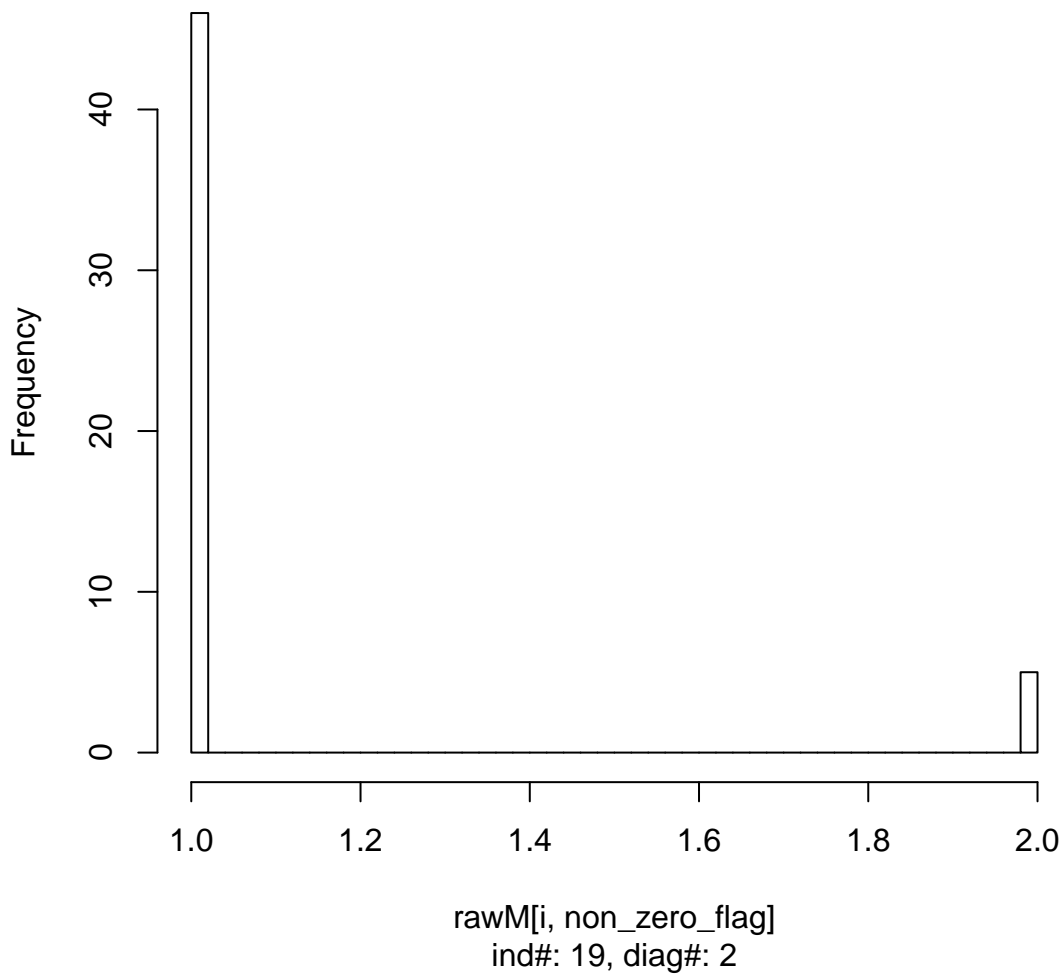




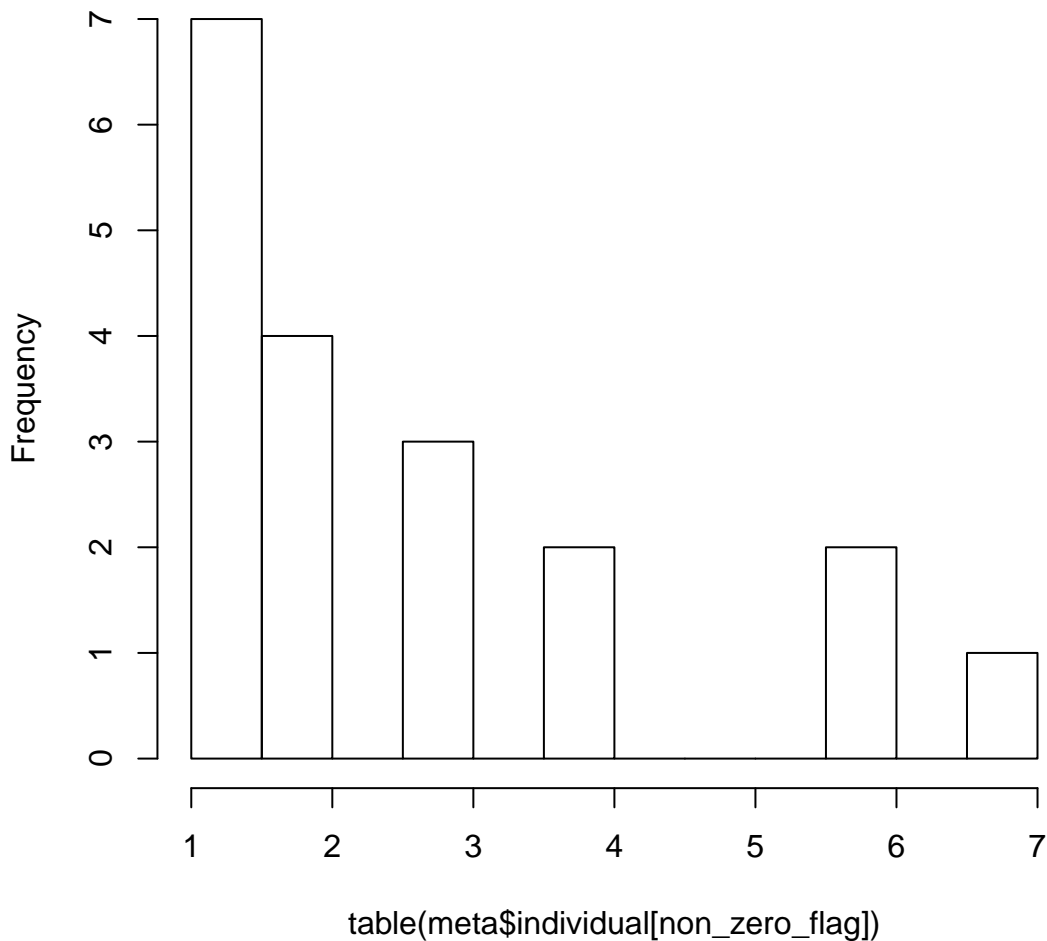
# KSless nonsig: individual expression cell count of gene#14



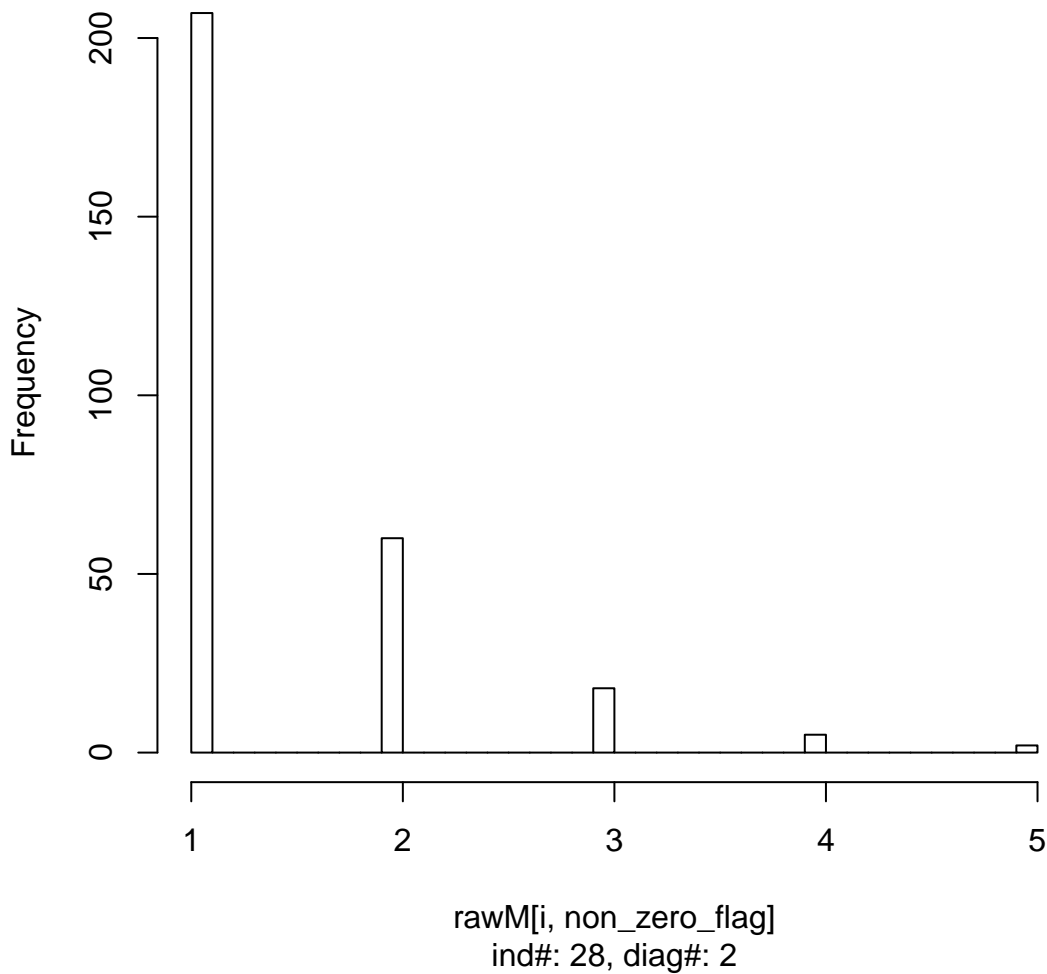
**S nonsig: log expression of gene#17, pval ob=0.9952, non-zero n**



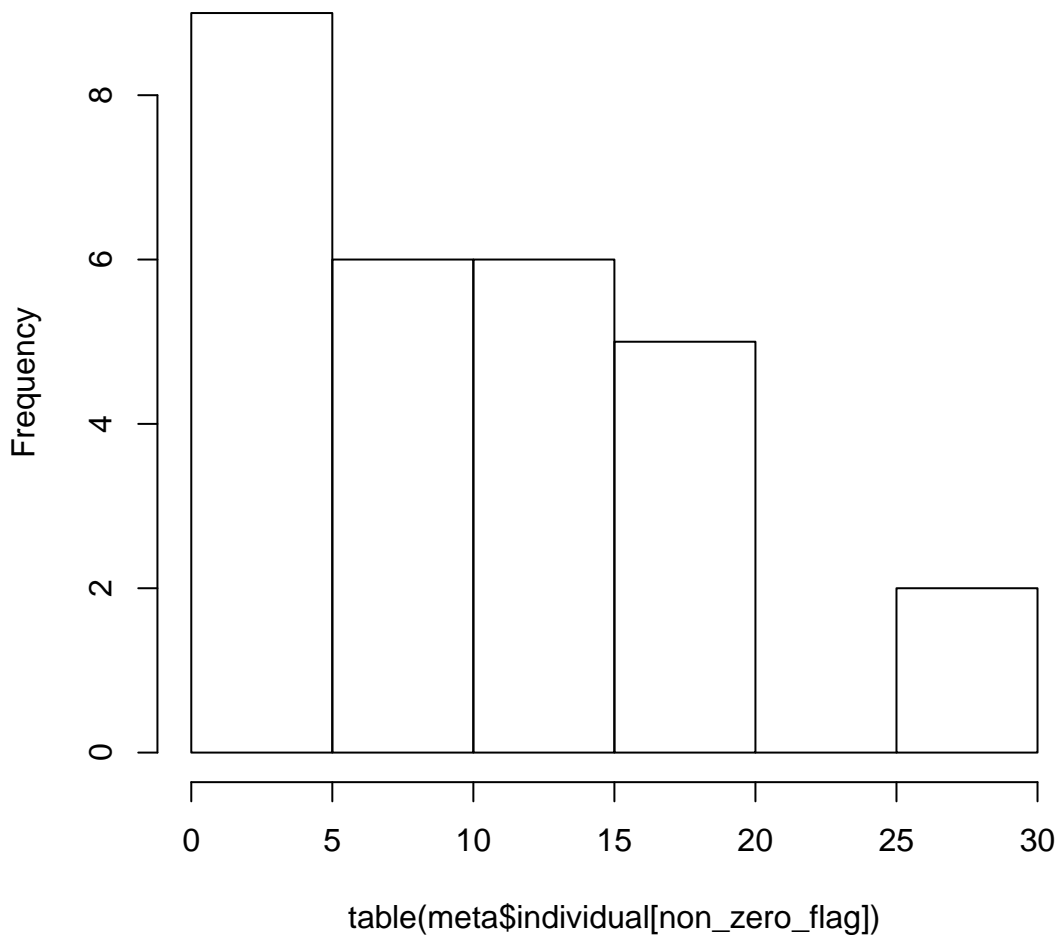
# KSless nonsig: individual expression cell count of gene#17



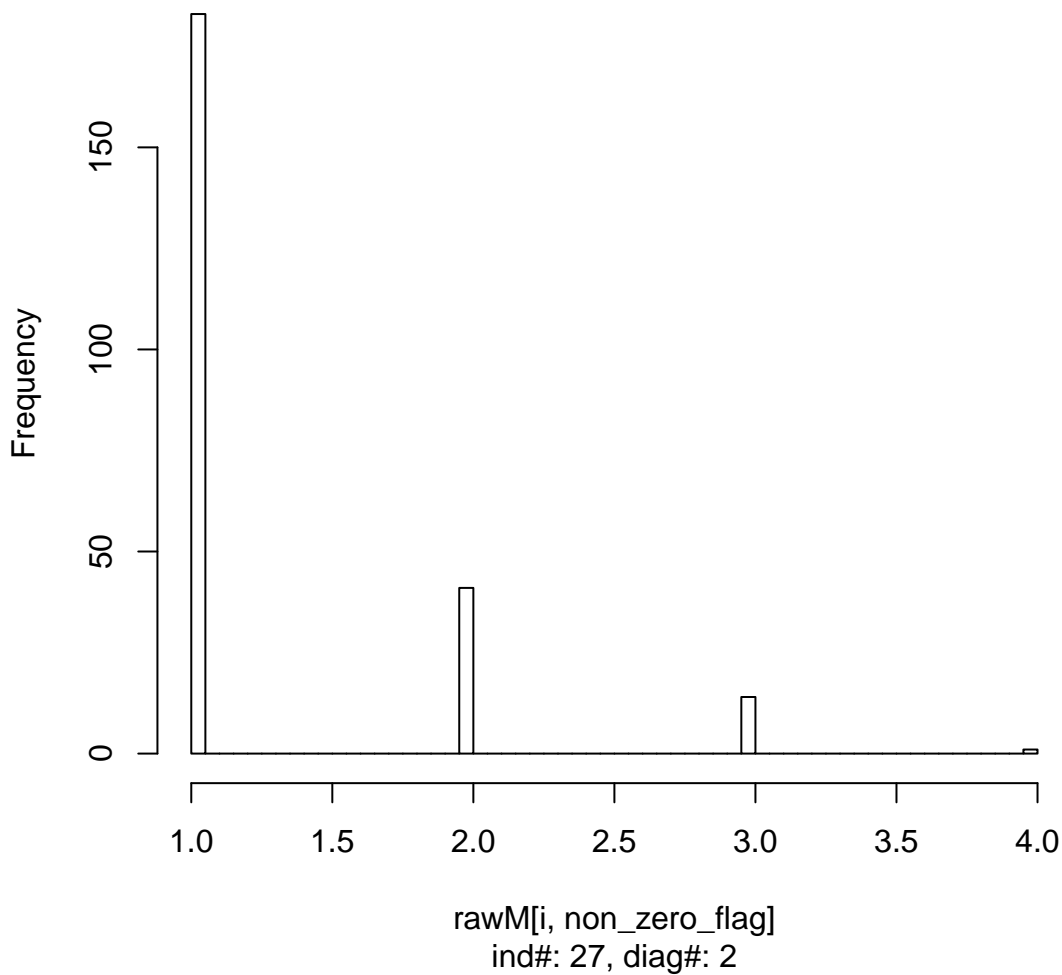
**S nonsig: log expression of gene#19, pval ob=0.456, non-zero nu**



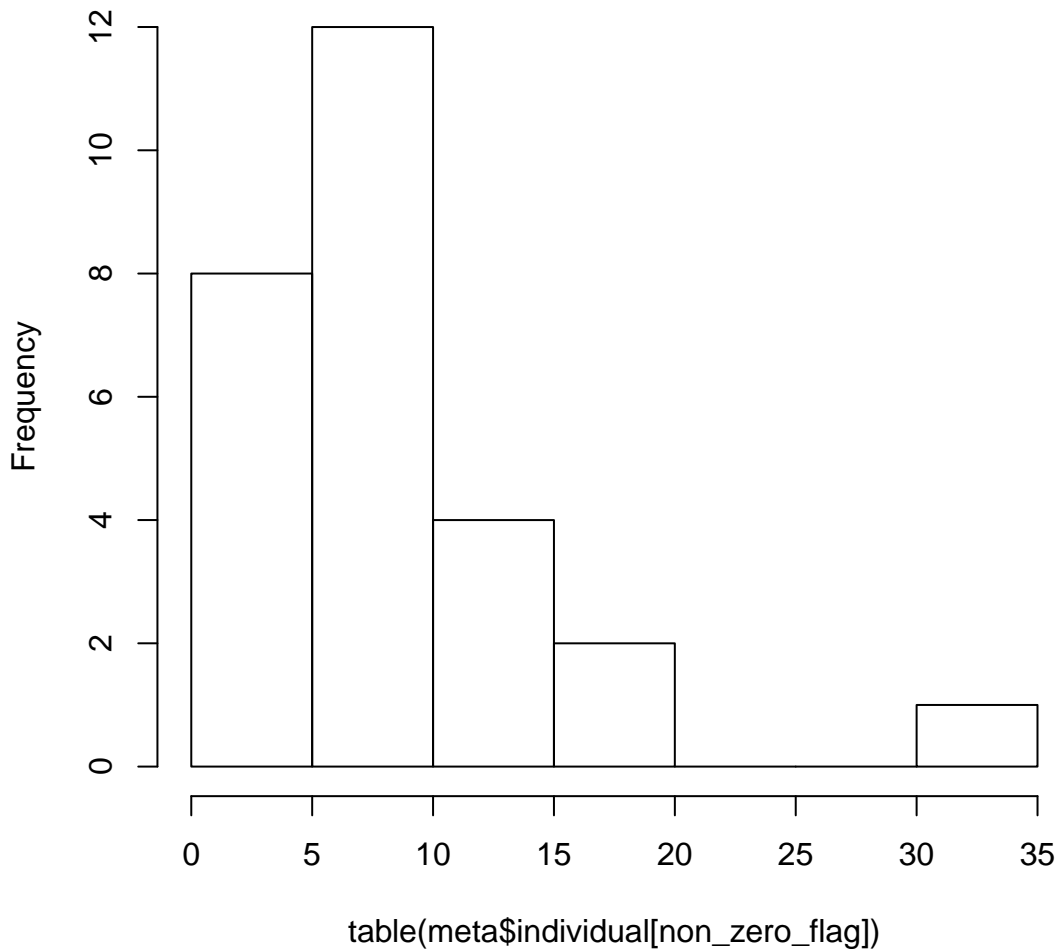
# KSless nonsig: individual expression cell count of gene#19



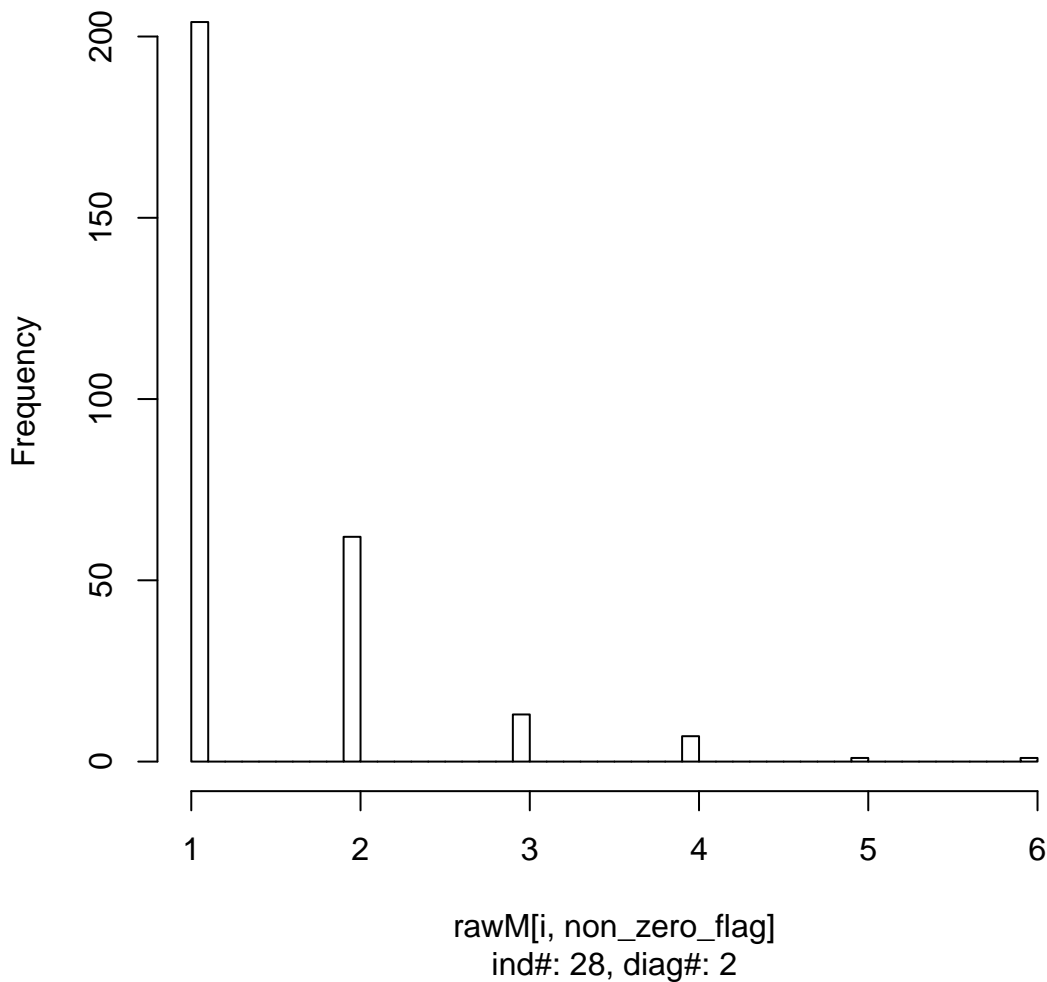
**nonsig: log expression of gene#20, pval ob=0.1412, non-zero nu**



# KSless nonsig: individual expression cell count of gene#20

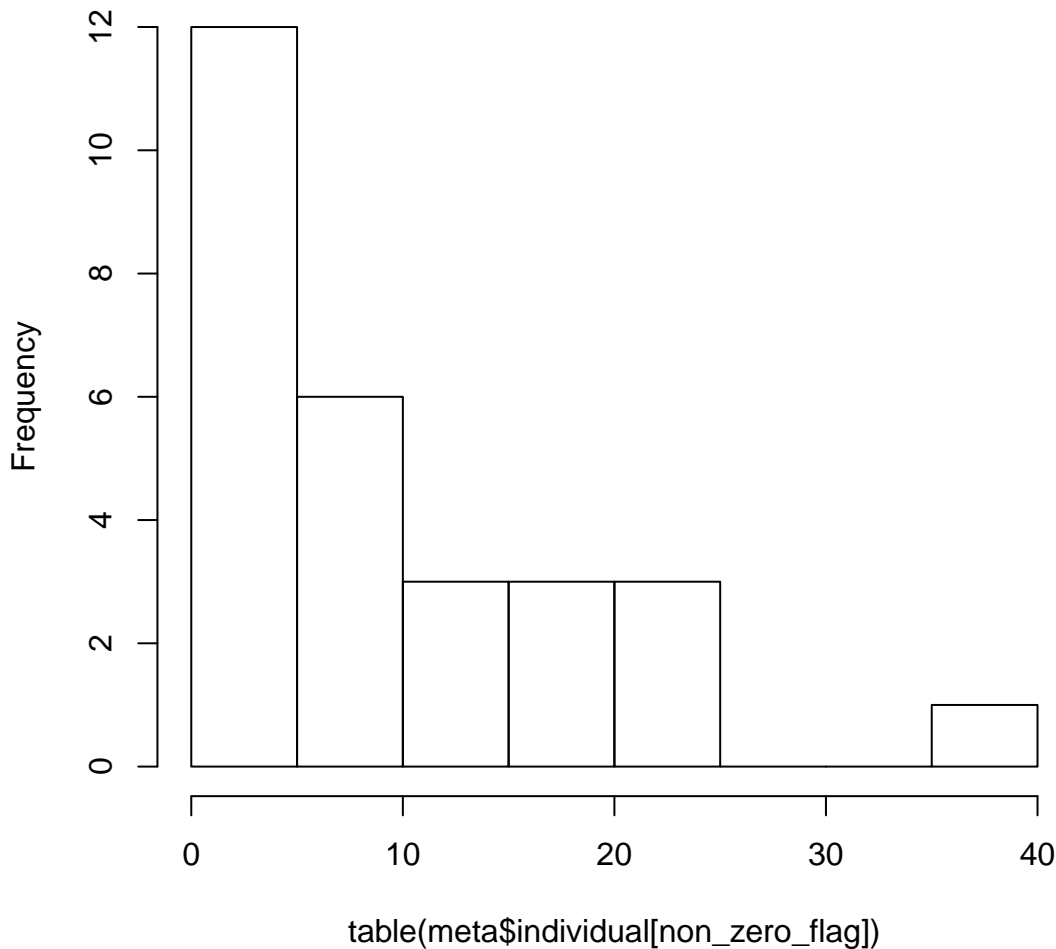


**S nonsig: log expression of gene#21, pval ob=0.746, non-zero nu**

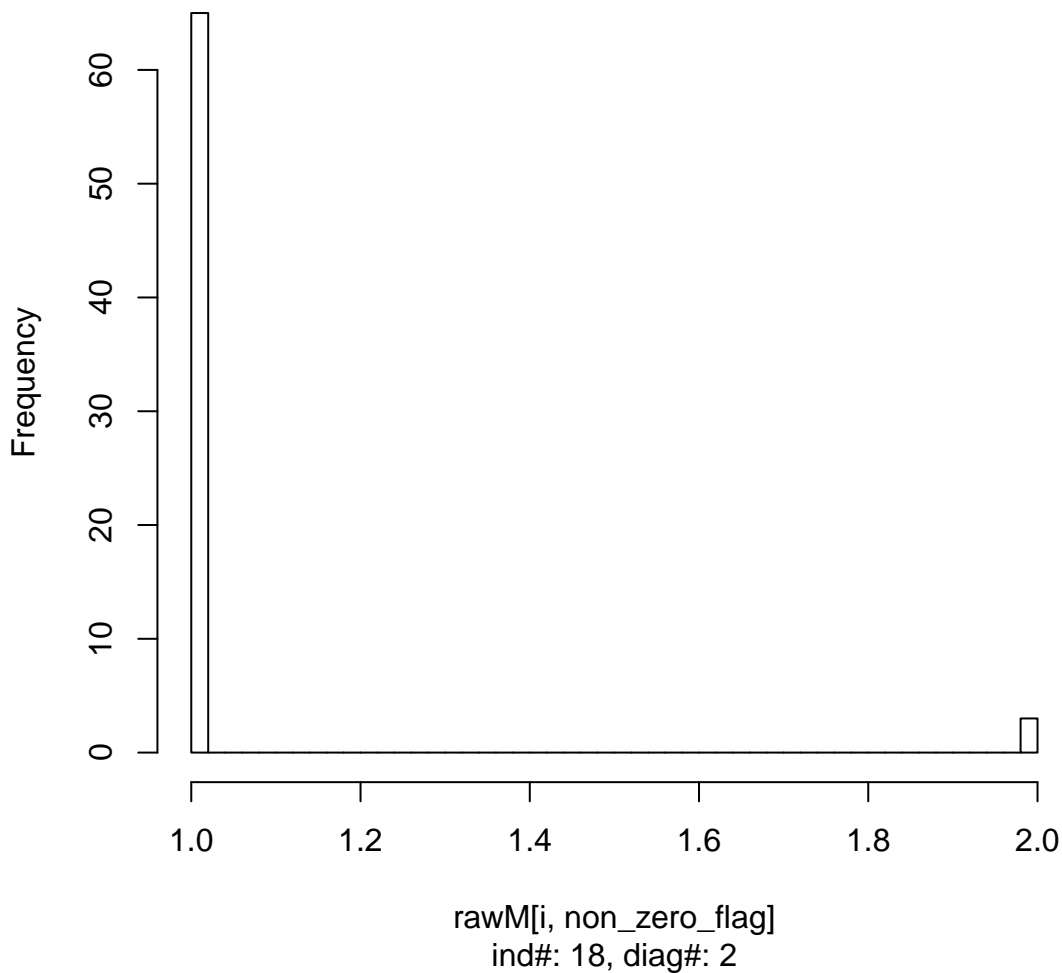




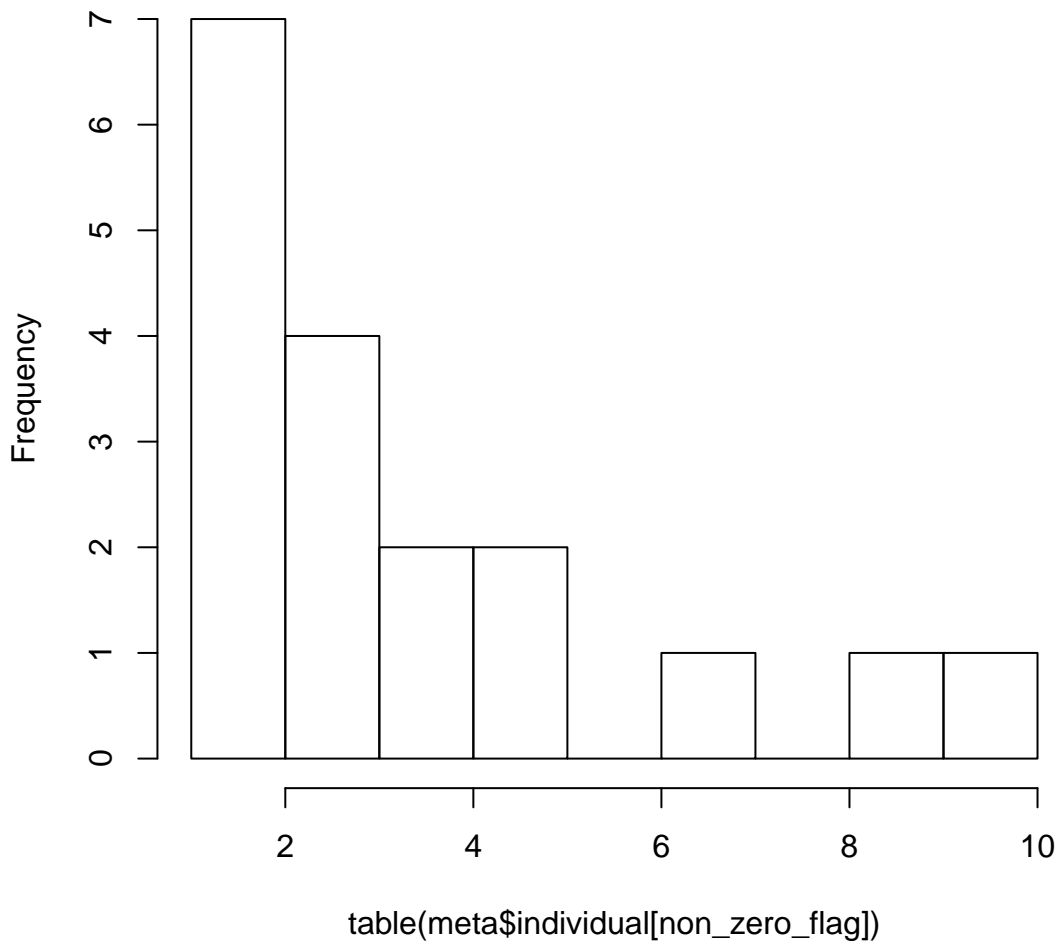
# KSless nonsig: individual expression cell count of gene#21



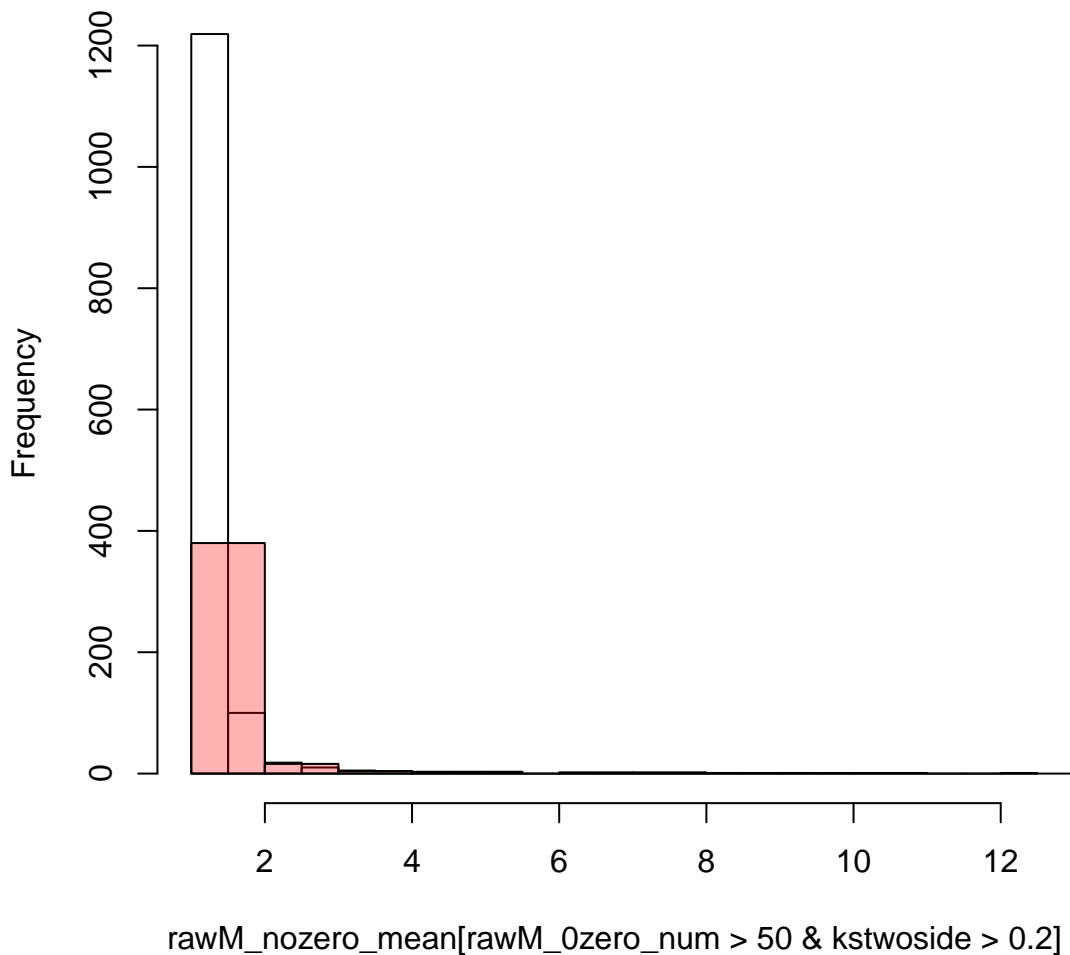
**S nonsig: log expression of gene#22, pval ob=0.4597, non-zero n**



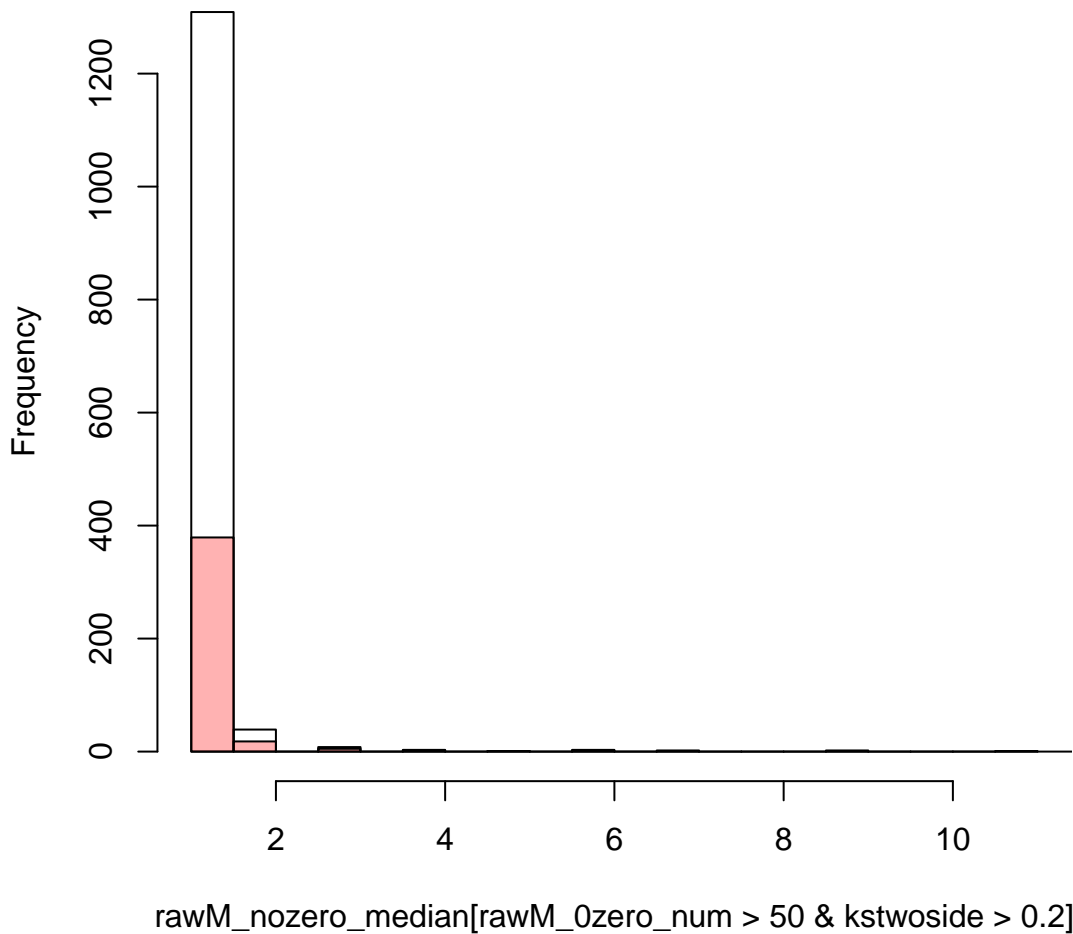
# KSless nonsig: individual expression cell count of gene#22



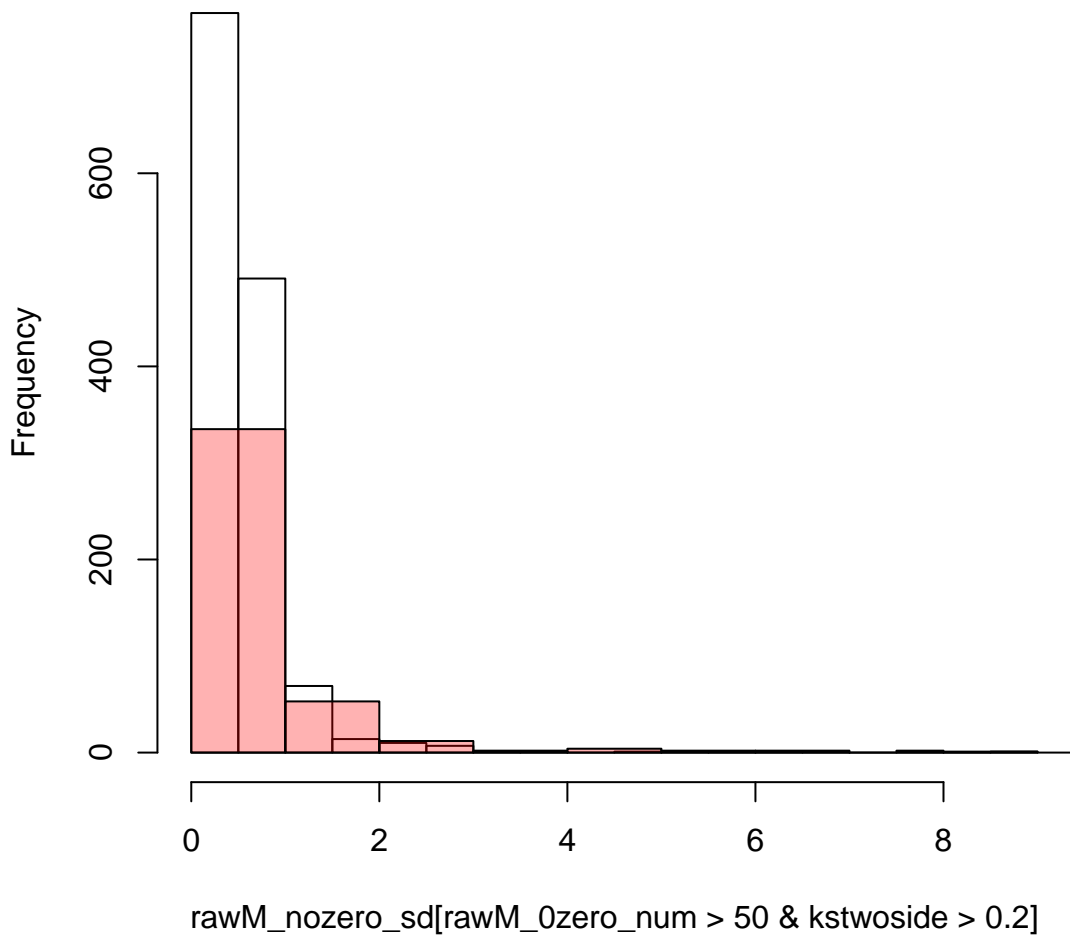
Histogram of rawM\_nozero\_mean[rawM\_0zero\_num > 50 & kstwsid



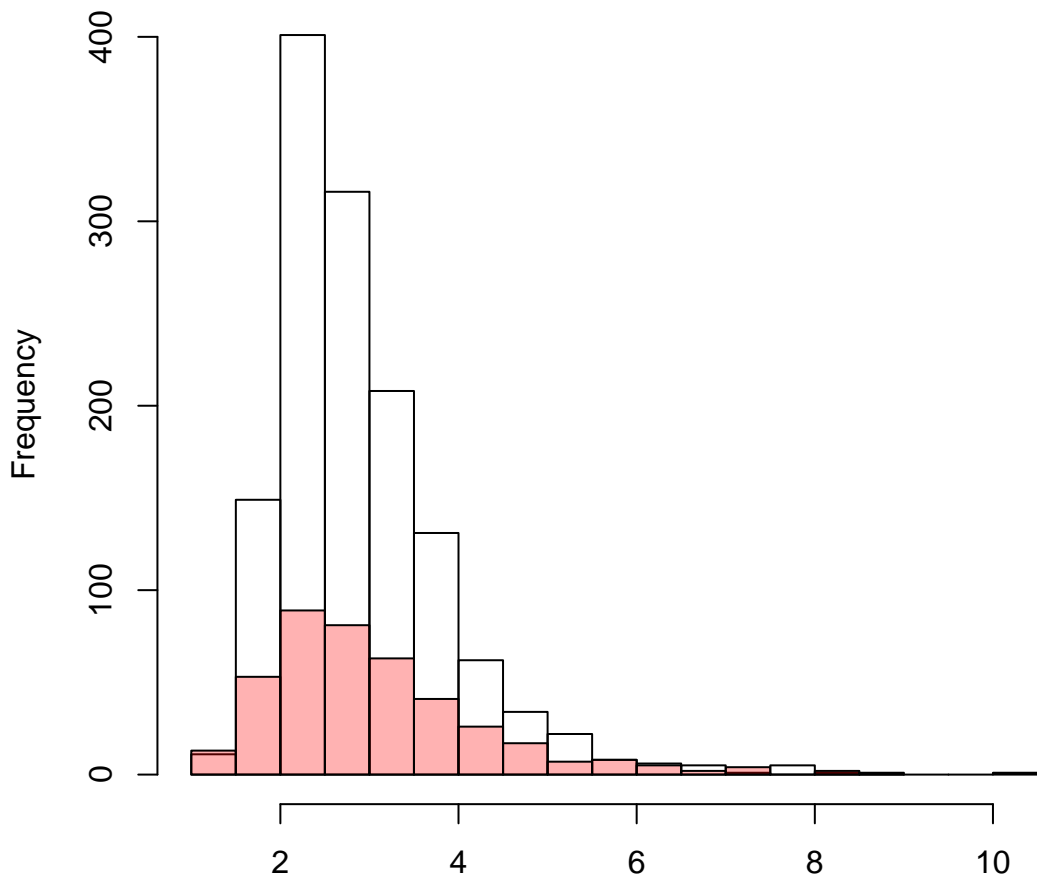
ogram of rawM\_nozero\_median[rawM\_0zero\_num > 50 & kstwosi



histogram of rawM\_nozero\_sd[rawM\_0zero\_num > 50 & kstwoside

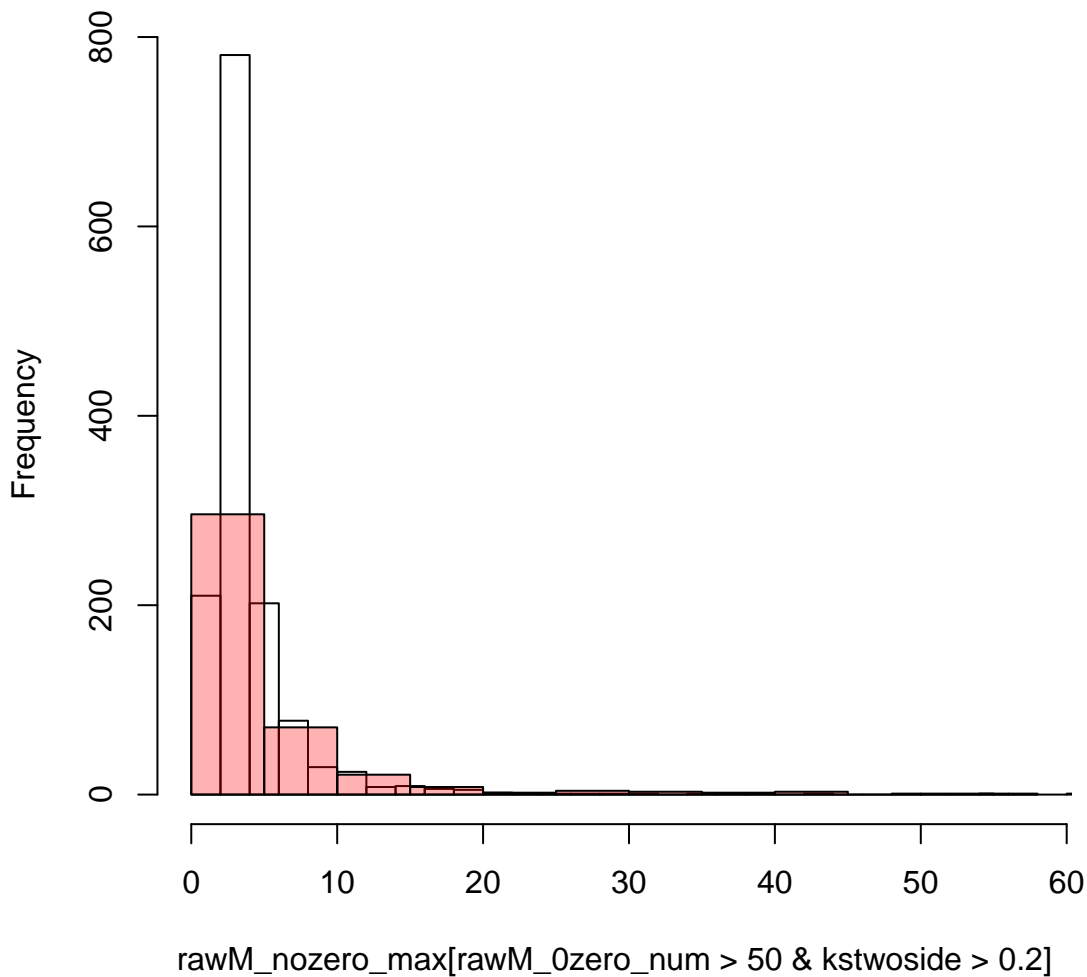


gram of rawM\_nozero\_skewness[rawM\_0zero\_num > 50 & kstws



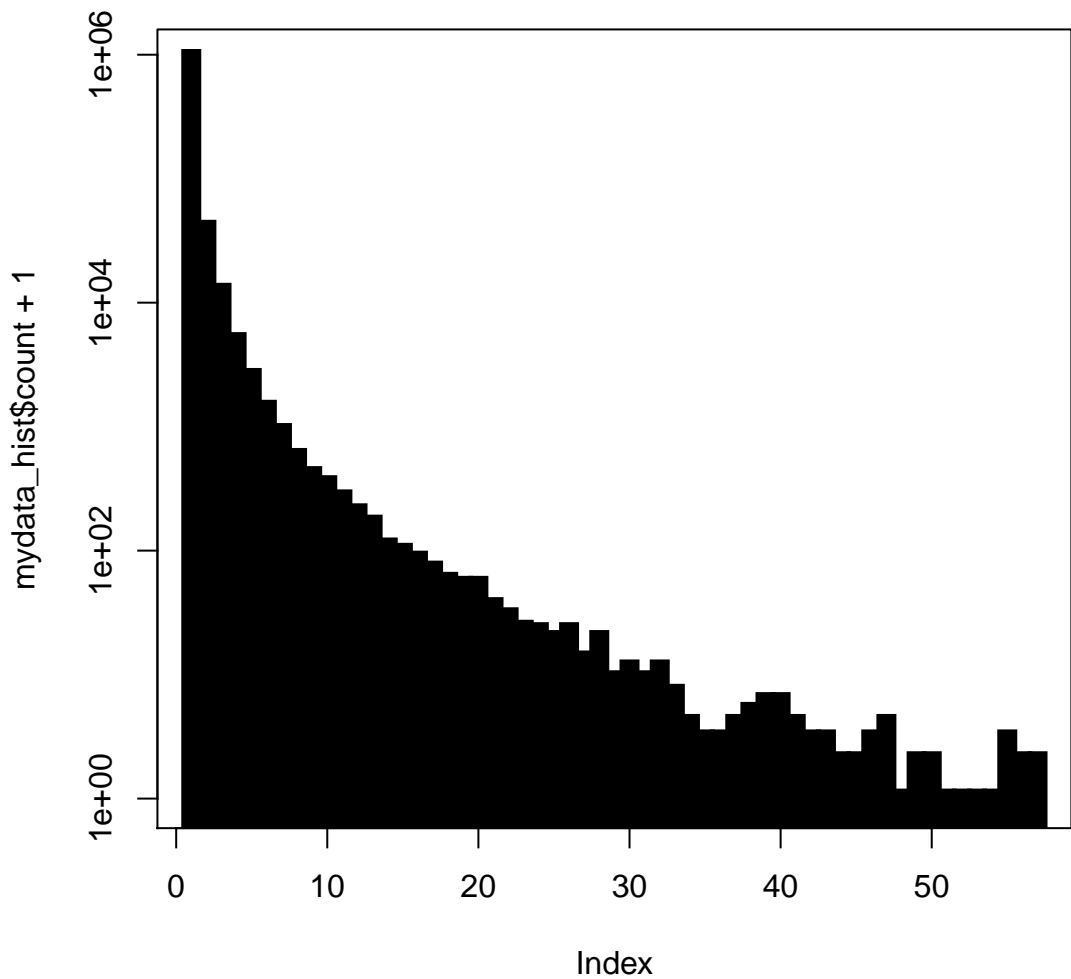
rawM\_nozero\_skewness[rawM\_0zero\_num > 50 & kstwside > 0.2]

Histogram of rawM\_nozero\_max[rawM\_0zero\_num > 50 & kstwoside

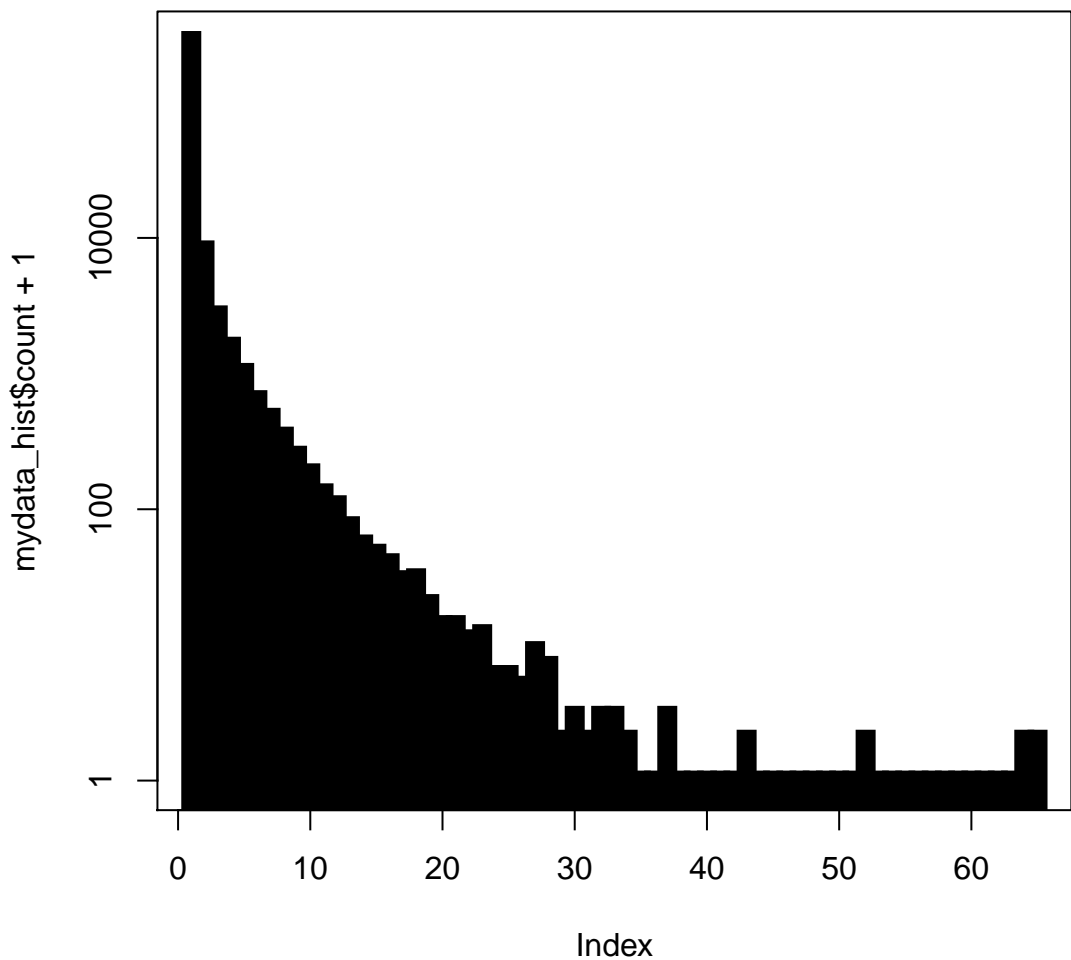




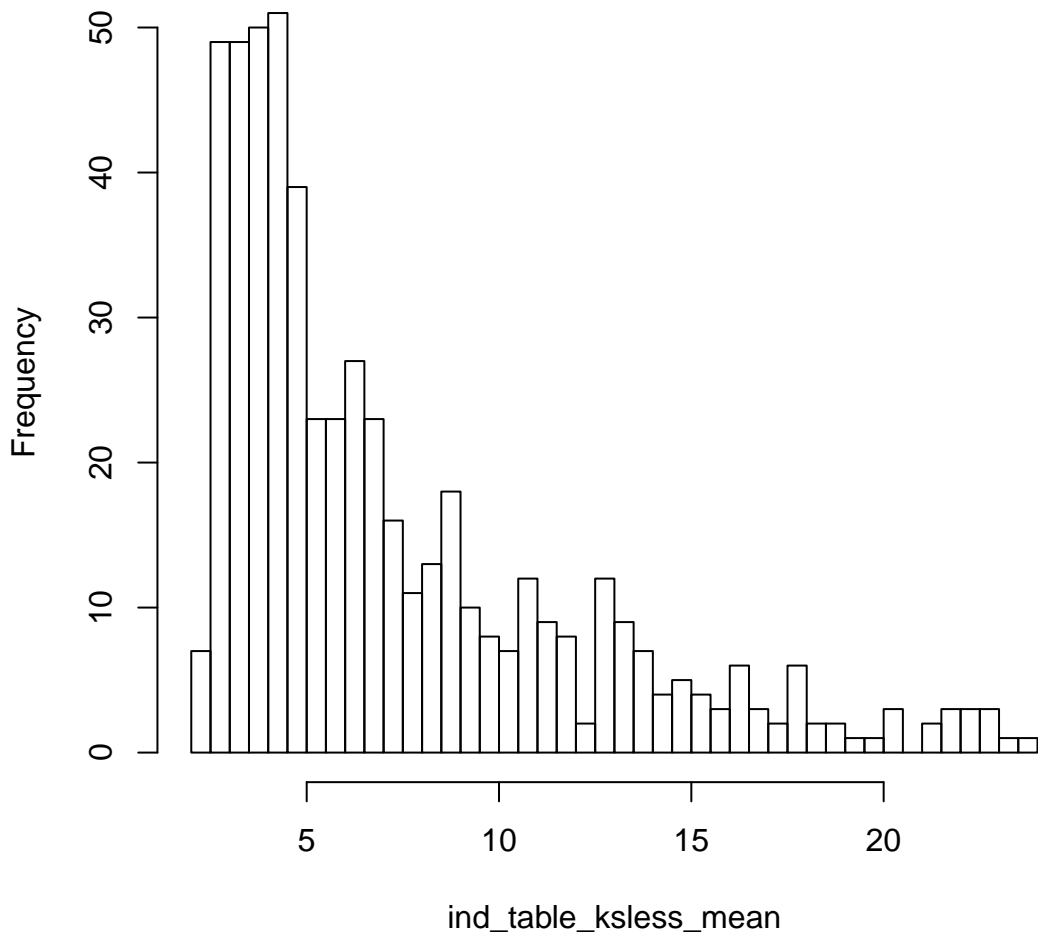
genes  $\log(\text{expression} + 1)$  with least 50 cell expression and kstwosi



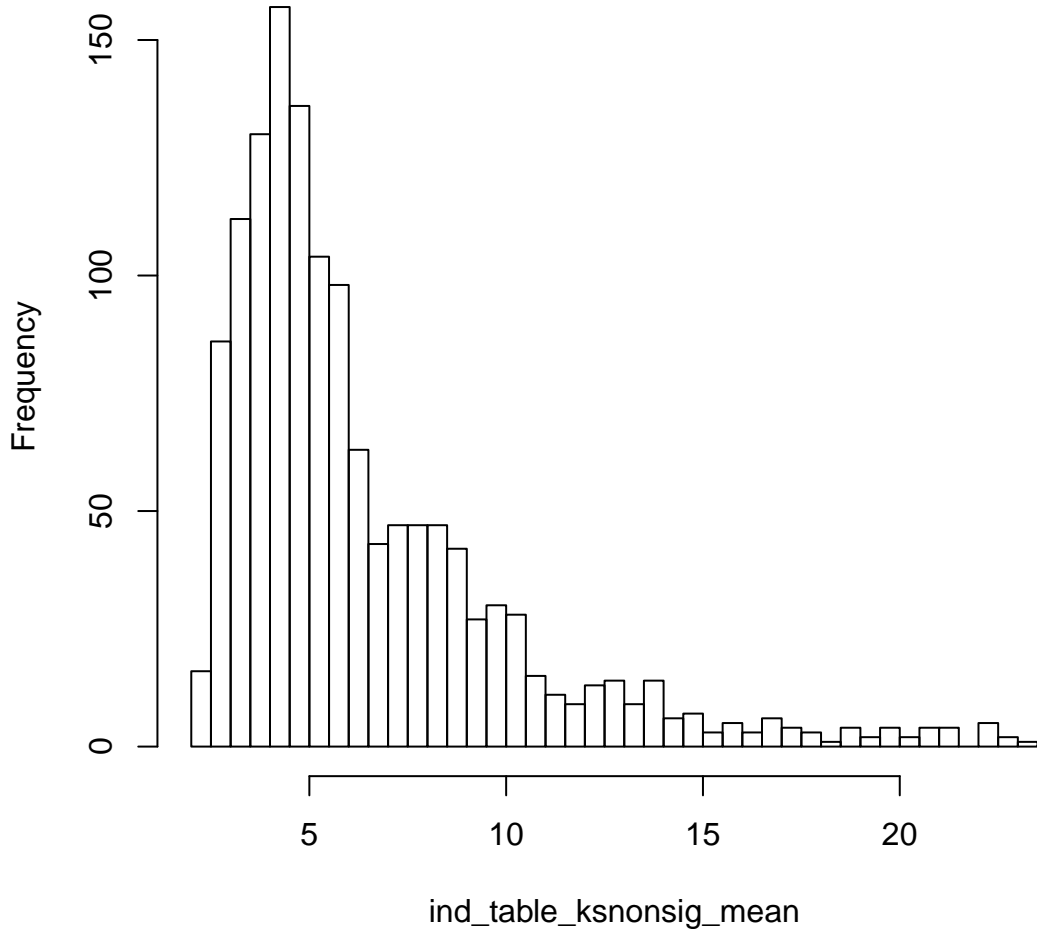
nes  $\log(\text{expression} + 1)$  with least 50 cell expression and kstwo



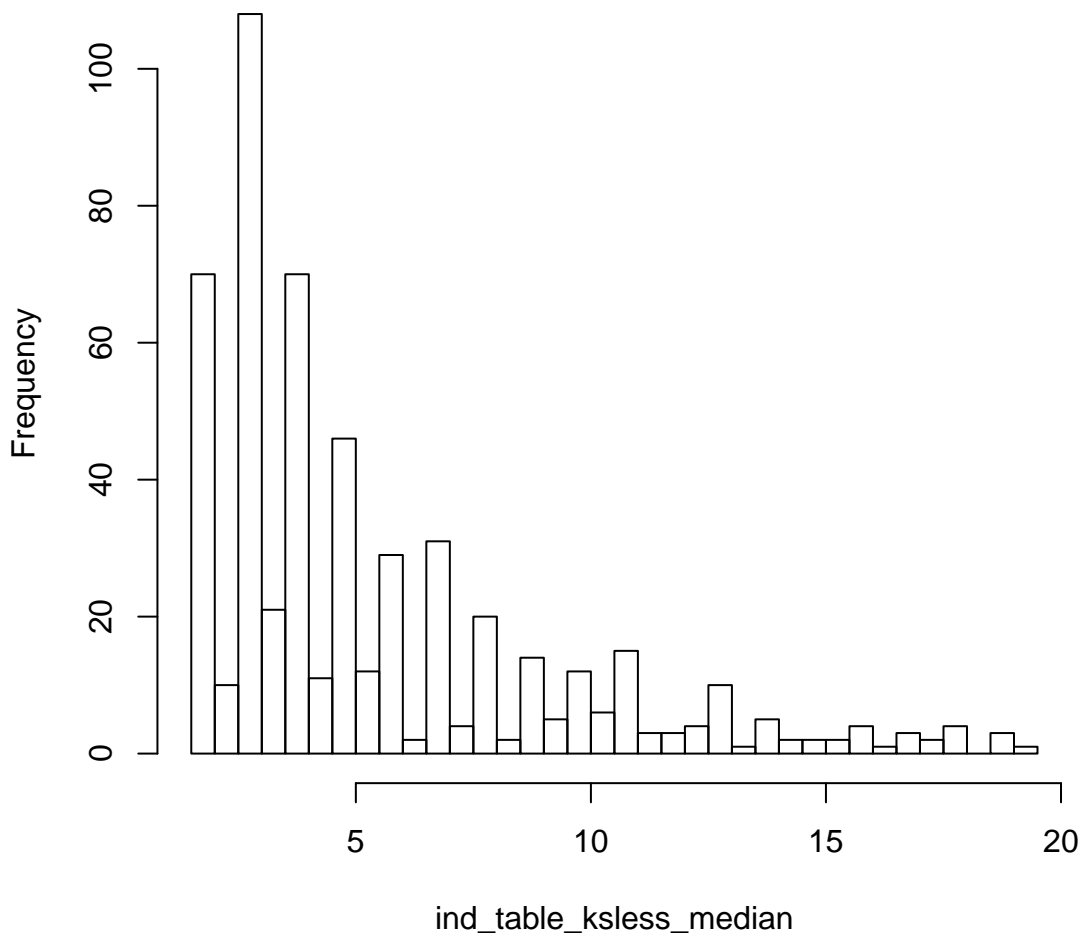
**Histogram of ind\_table\_ksless\_mean**



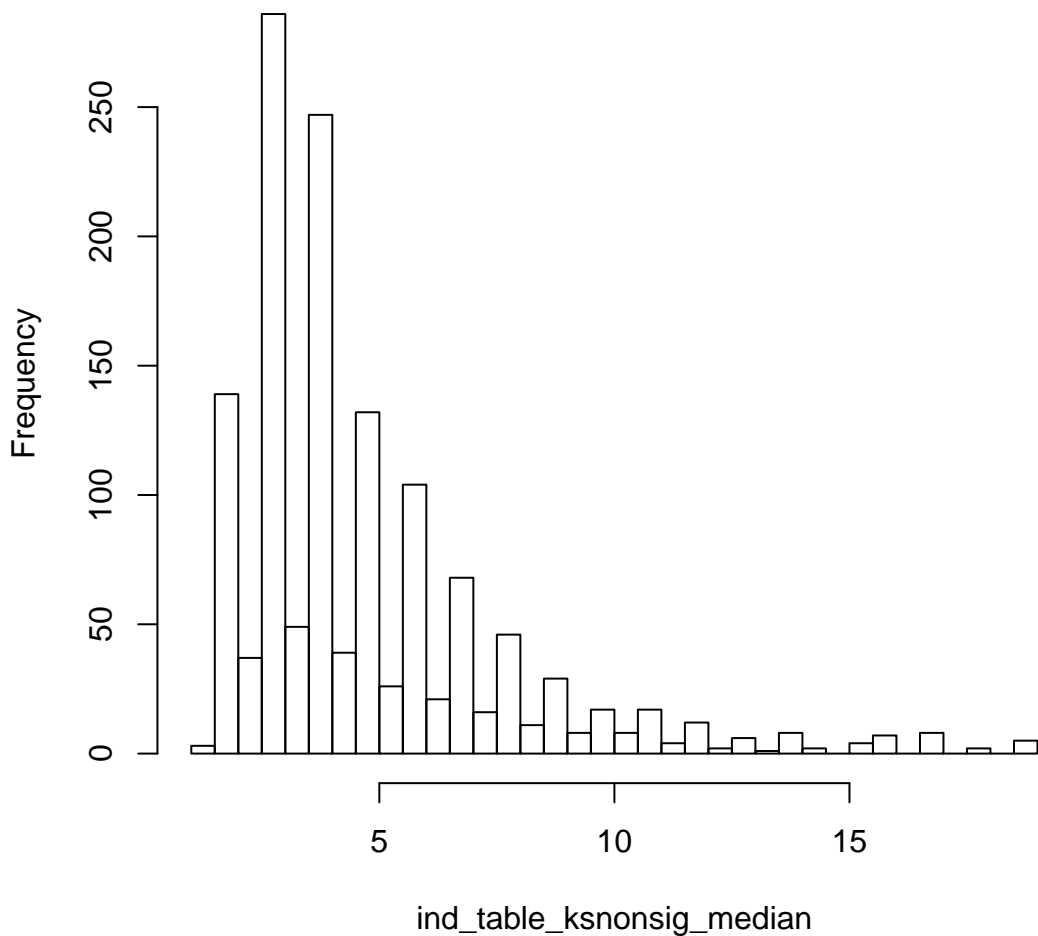
**Histogram of ind\_table\_ksnonsig\_mean**



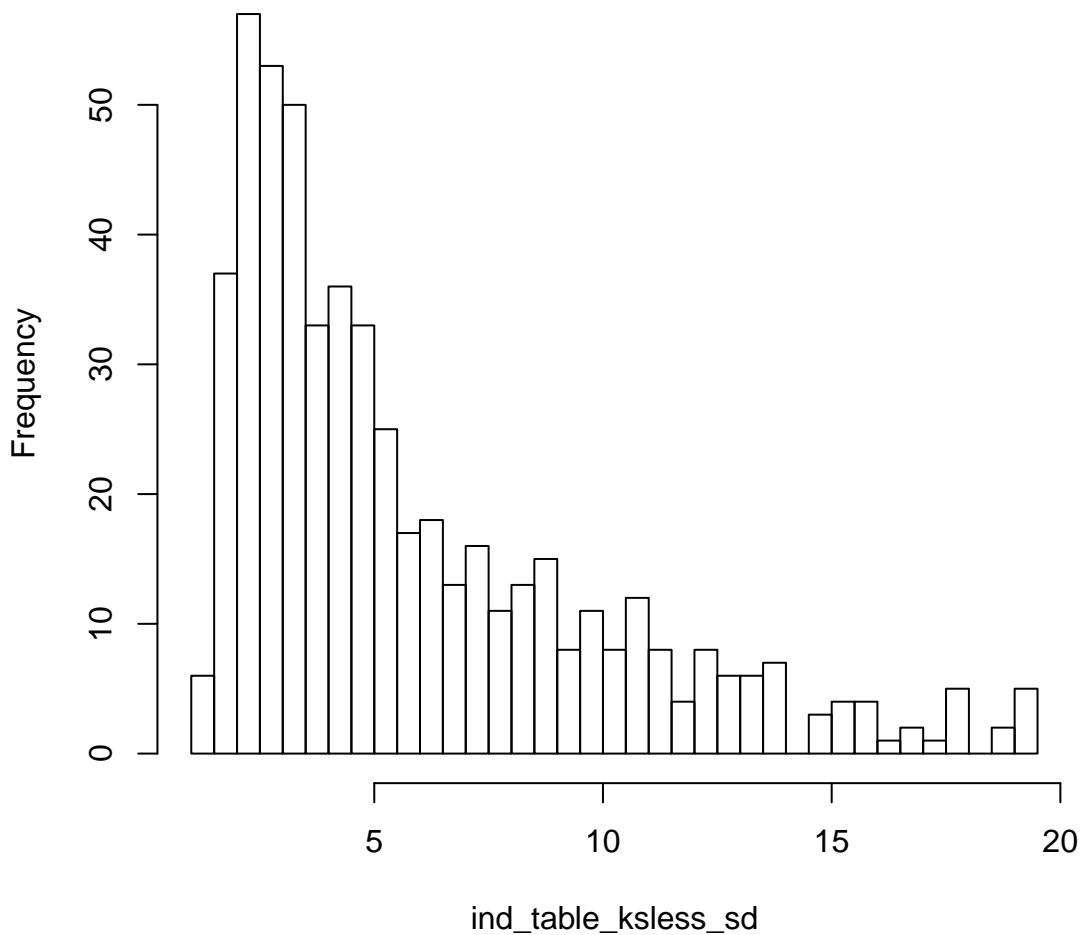
**Histogram of ind\_table\_ksless\_median**



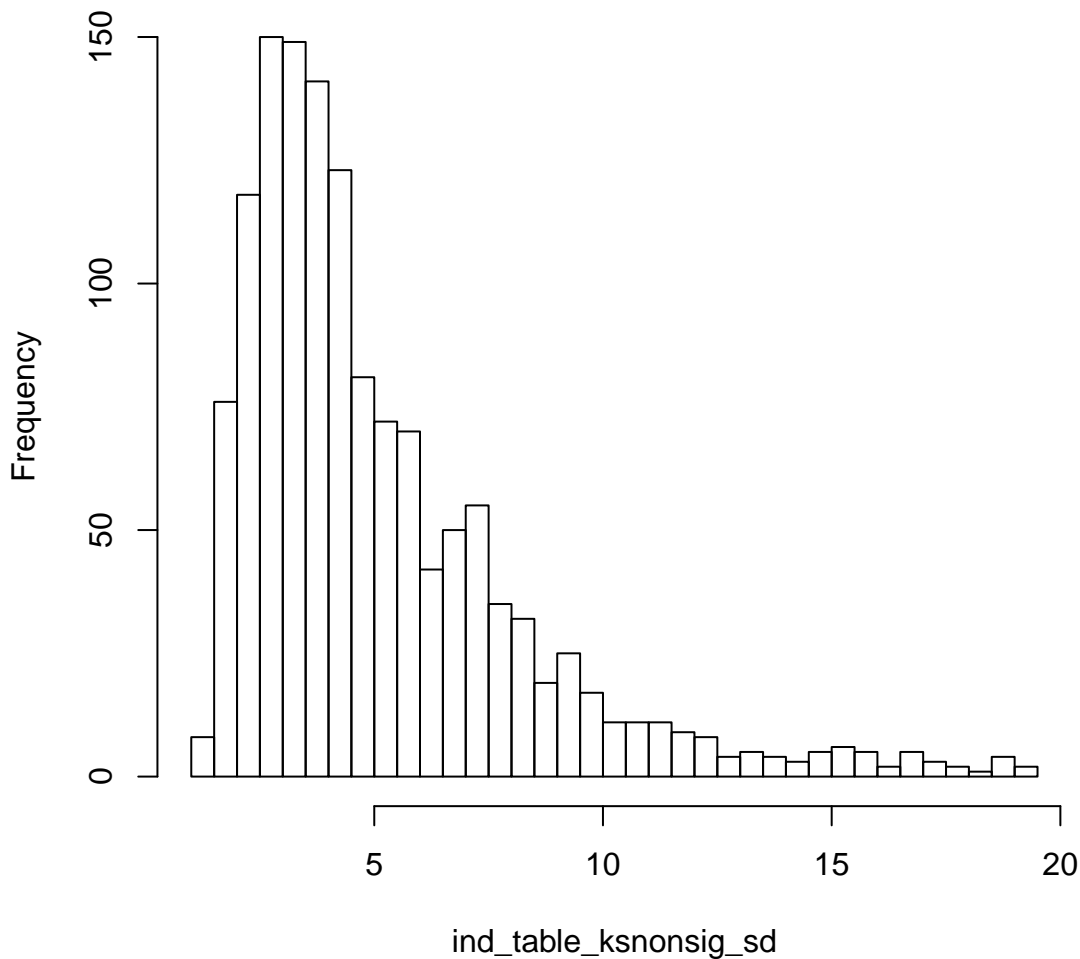
**Histogram of ind\_table\_ksnonsig\_median**



**Histogram of ind\_table\_ksless\_sd**

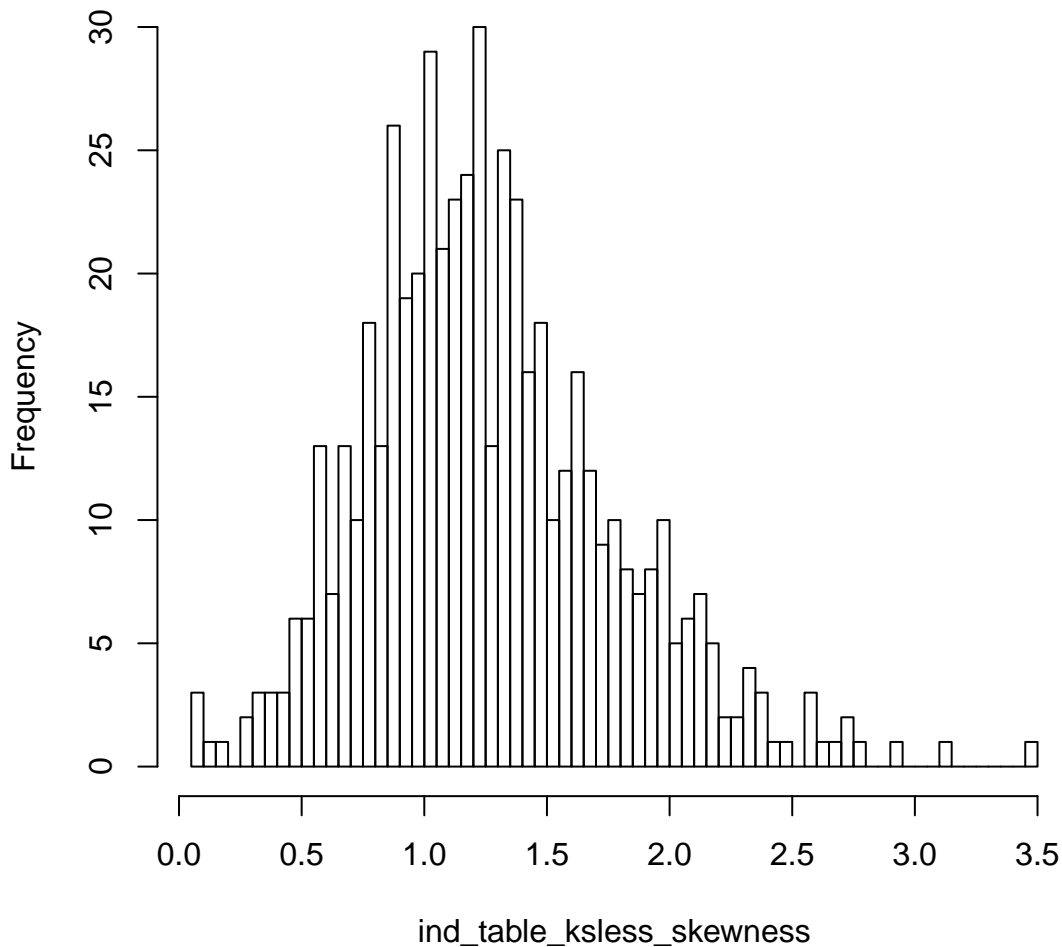


**Histogram of ind\_table\_ksnonsig\_sd**

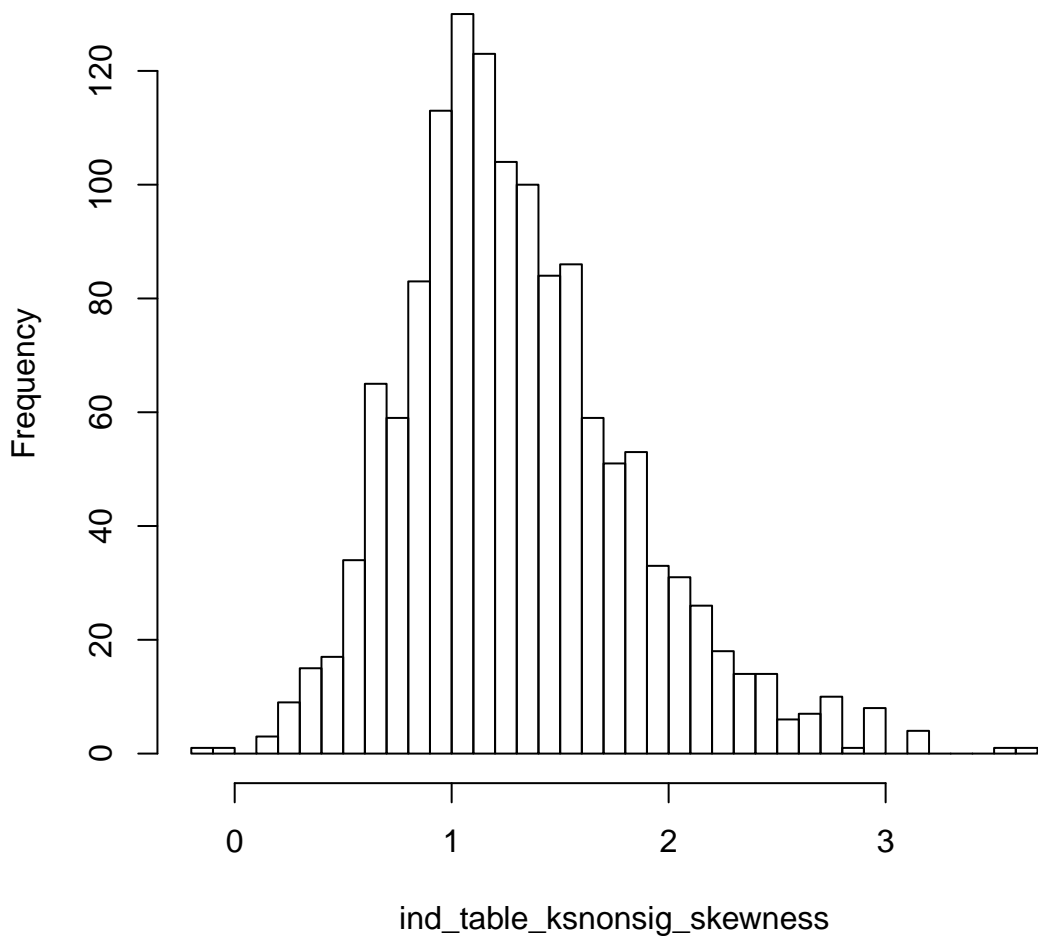




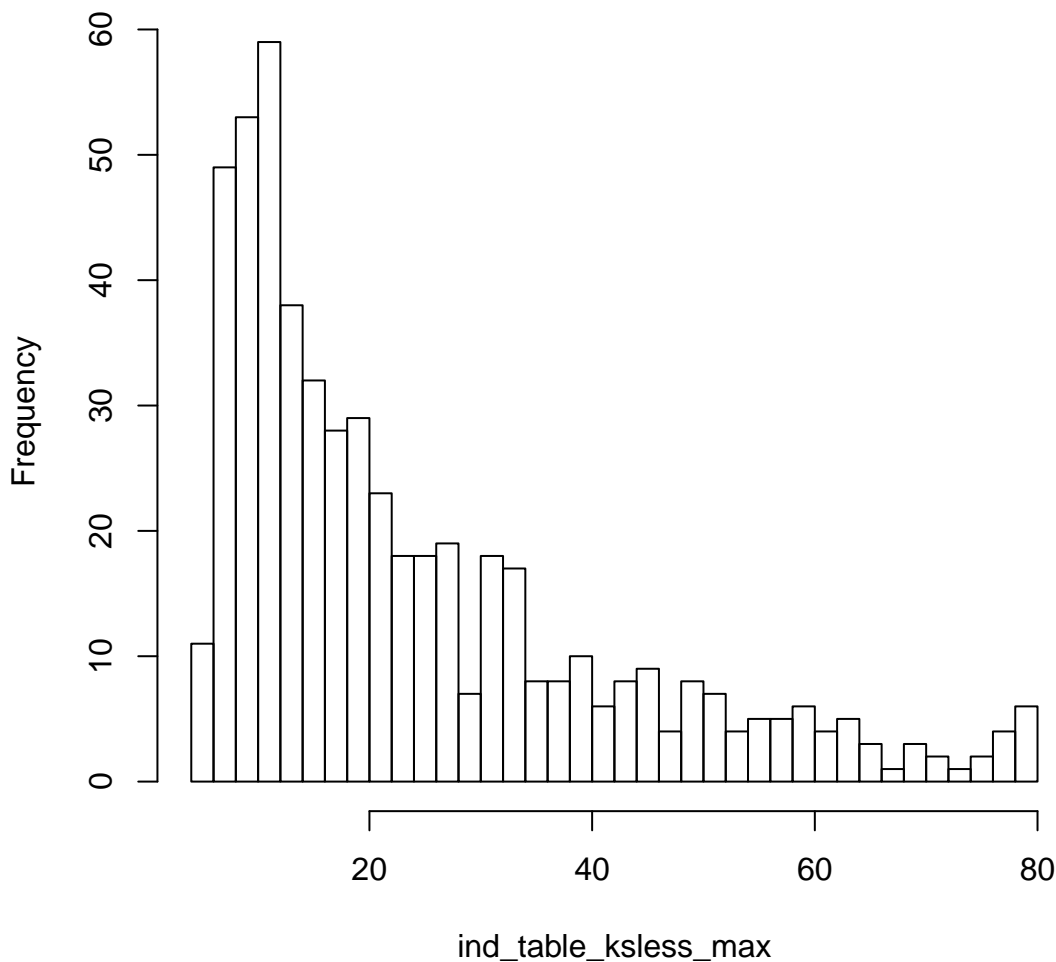
**Histogram of ind\_table\_ksless\_skewness**



**Histogram of ind\_table\_ksnonsig\_skewness**



**Histogram of ind\_table\_ksless\_max**



**Histogram of ind\_table\_ksnonsig\_max**

