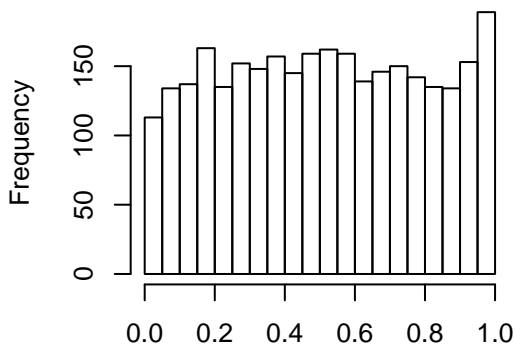
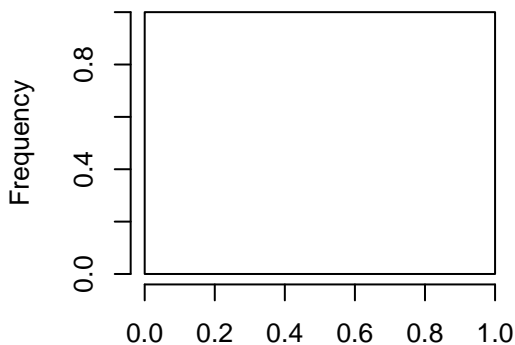


perm pvalues



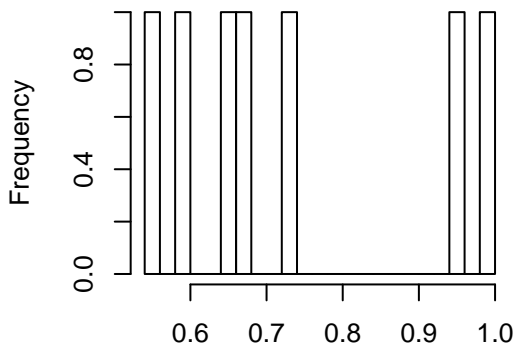
rawp  
n=2952

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



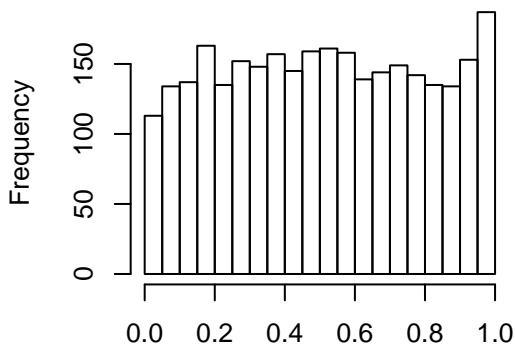
rawp[rawM\_0zero\_num < max(20, min(rawM\_0zero\_num, 100))]  
n=1, ks\_pval=1

perm pvalues,rawM\_0zero\_num >= 20



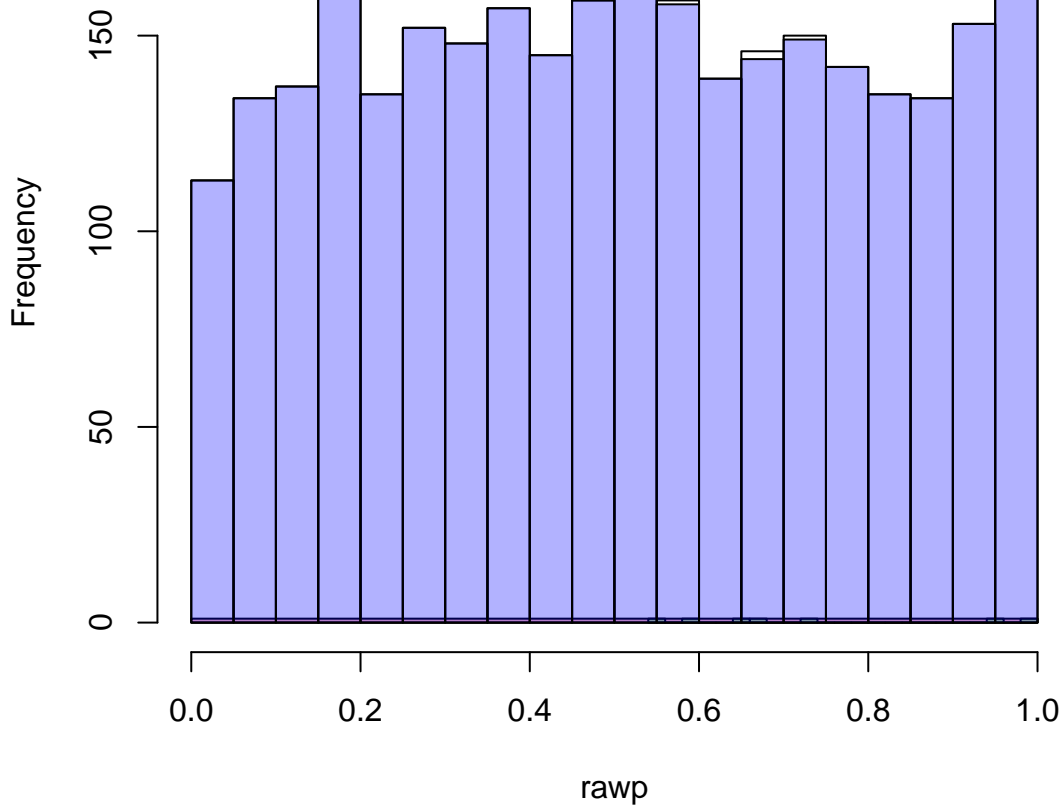
rawp[rawM\_0zero\_num >= 20 & rawM\_0zero\_num <= 100]  
n=7, ks\_pval=0.541243098374871

perm pvalues,rawM\_0zero\_num >= 100

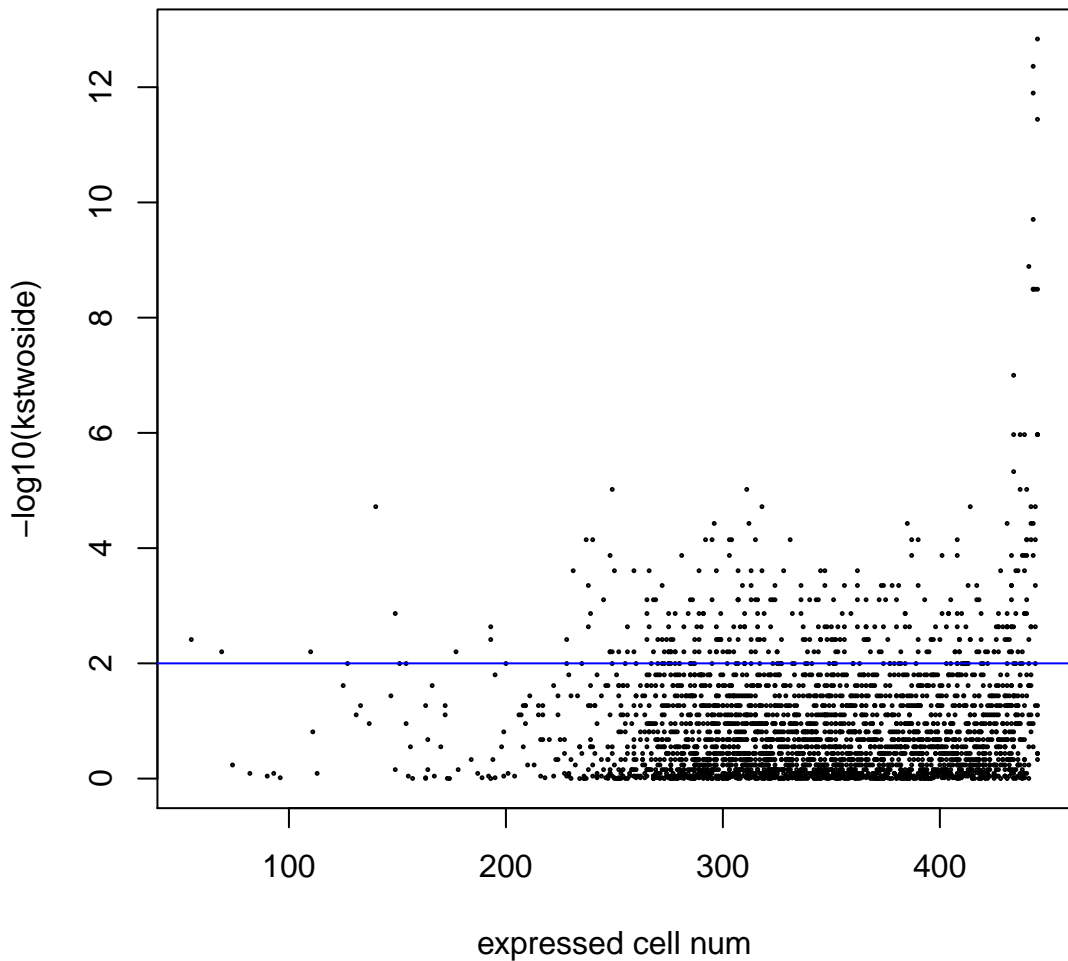


rawp[rawM\_0zero\_num >= 100]  
n=2945, ks\_pval=0.12776811909323

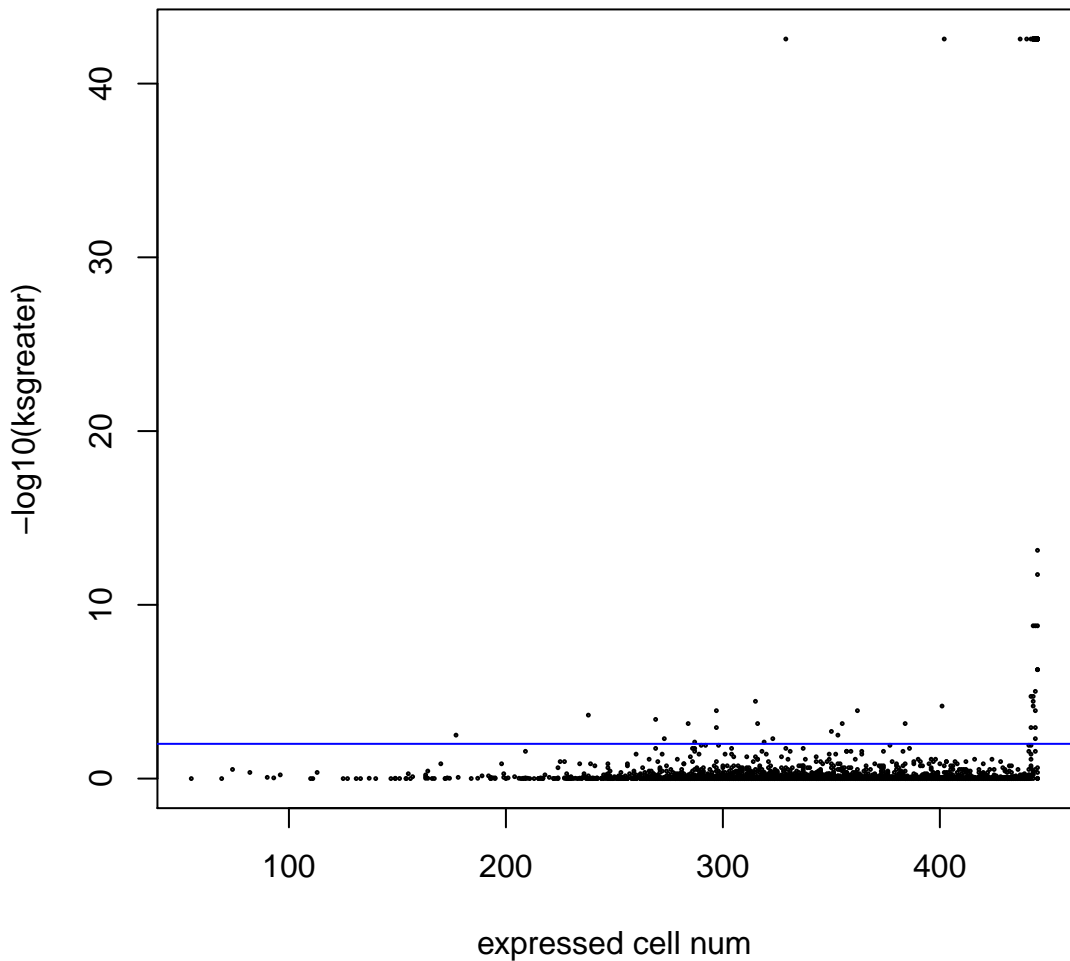
# perm pvalues



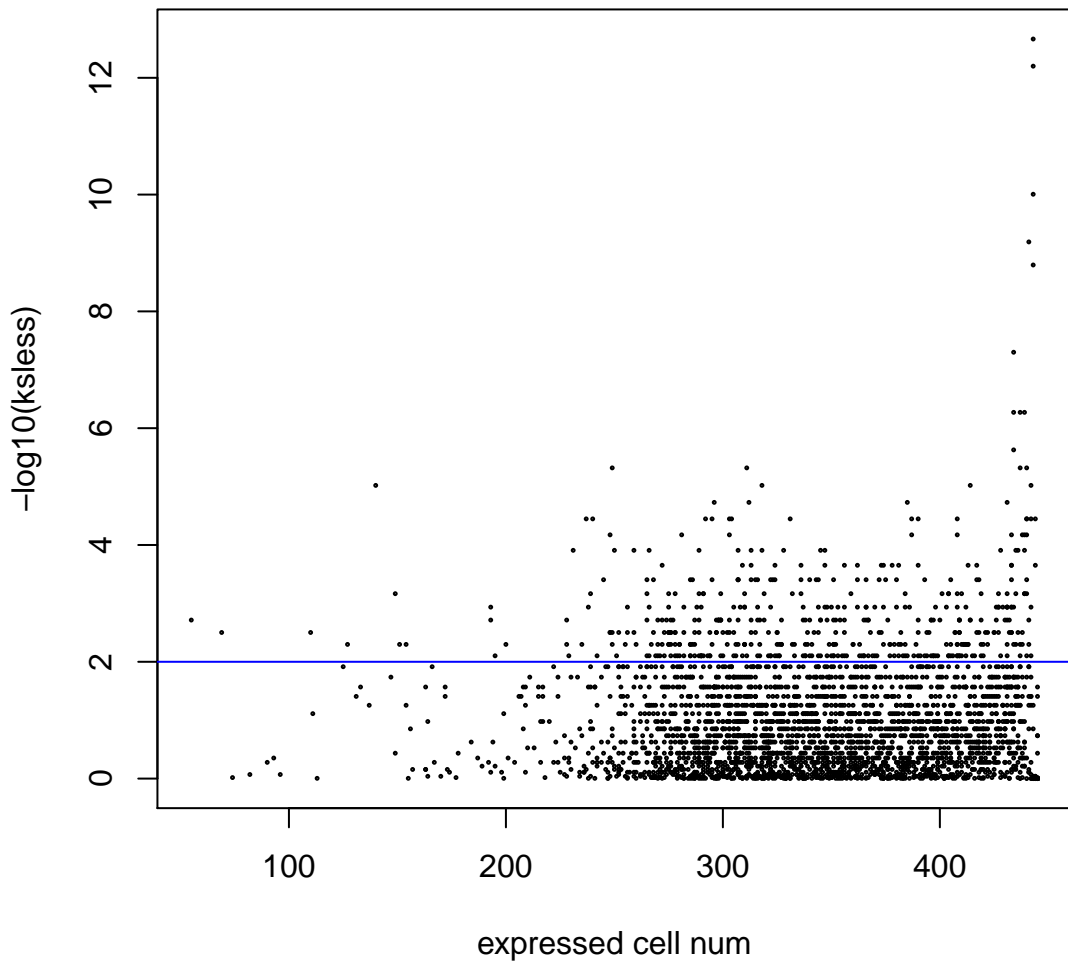
**sig\_KStwoside: 12.263%**



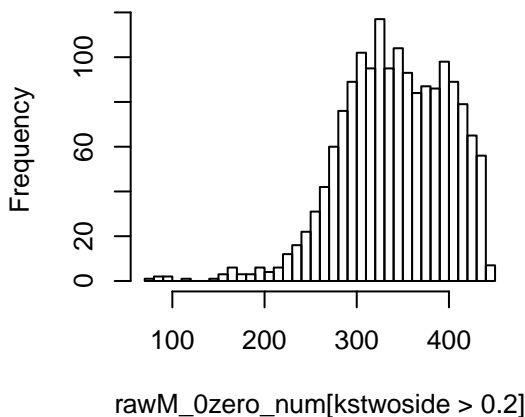
**sig\_KSgreater: 2.507%**



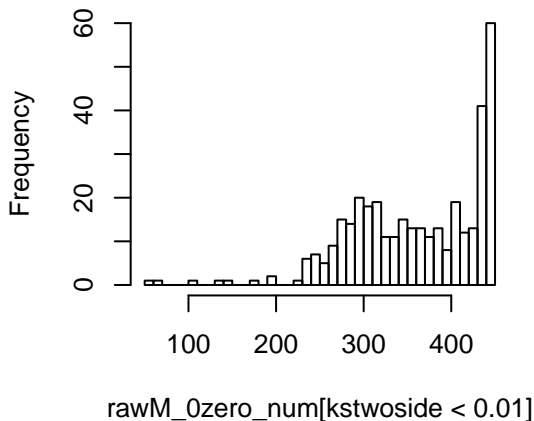
**sig\_KSless: 15.921%**



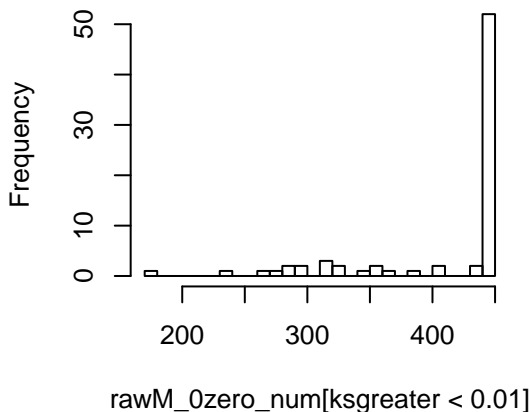
**expression cell num,kstwside>0.2**



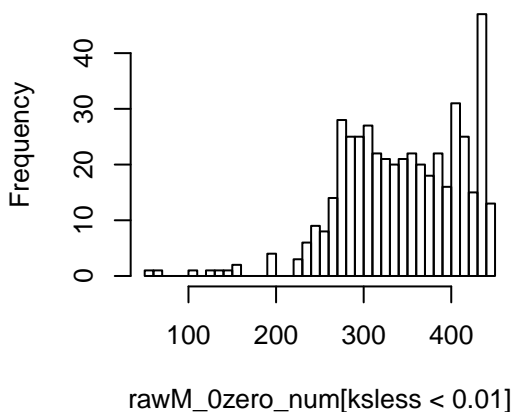
**expression cell num,kstwside<0.0**



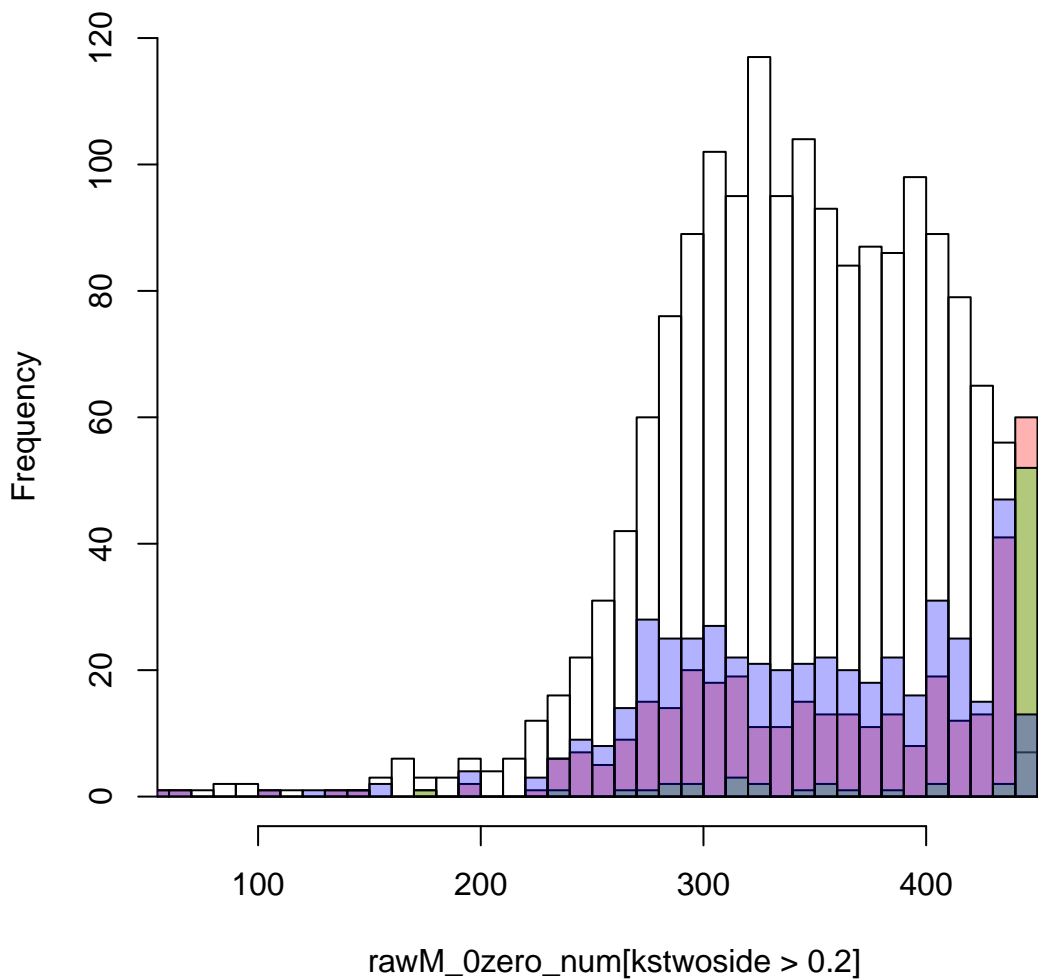
**expression cell num,ksgreater<0.0**



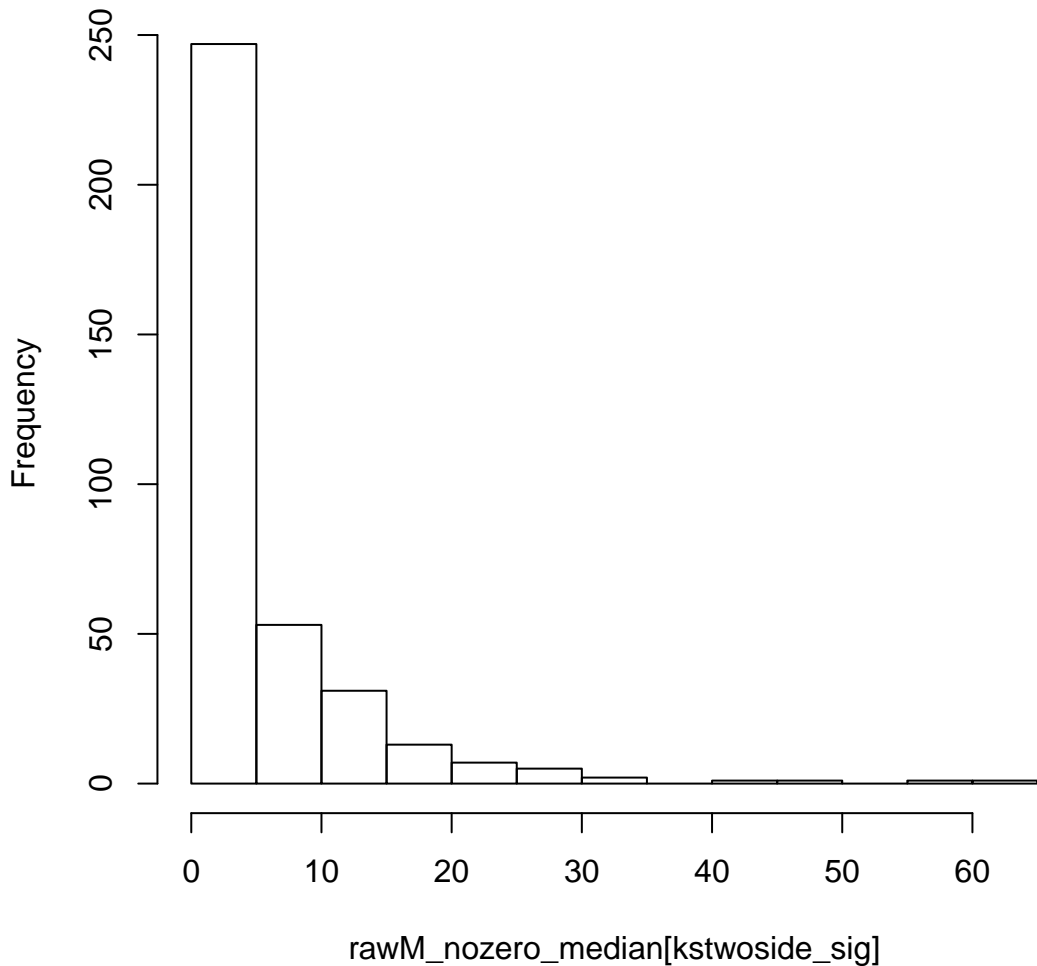
**expression cell num,ksless<0.01**



# expression cell num

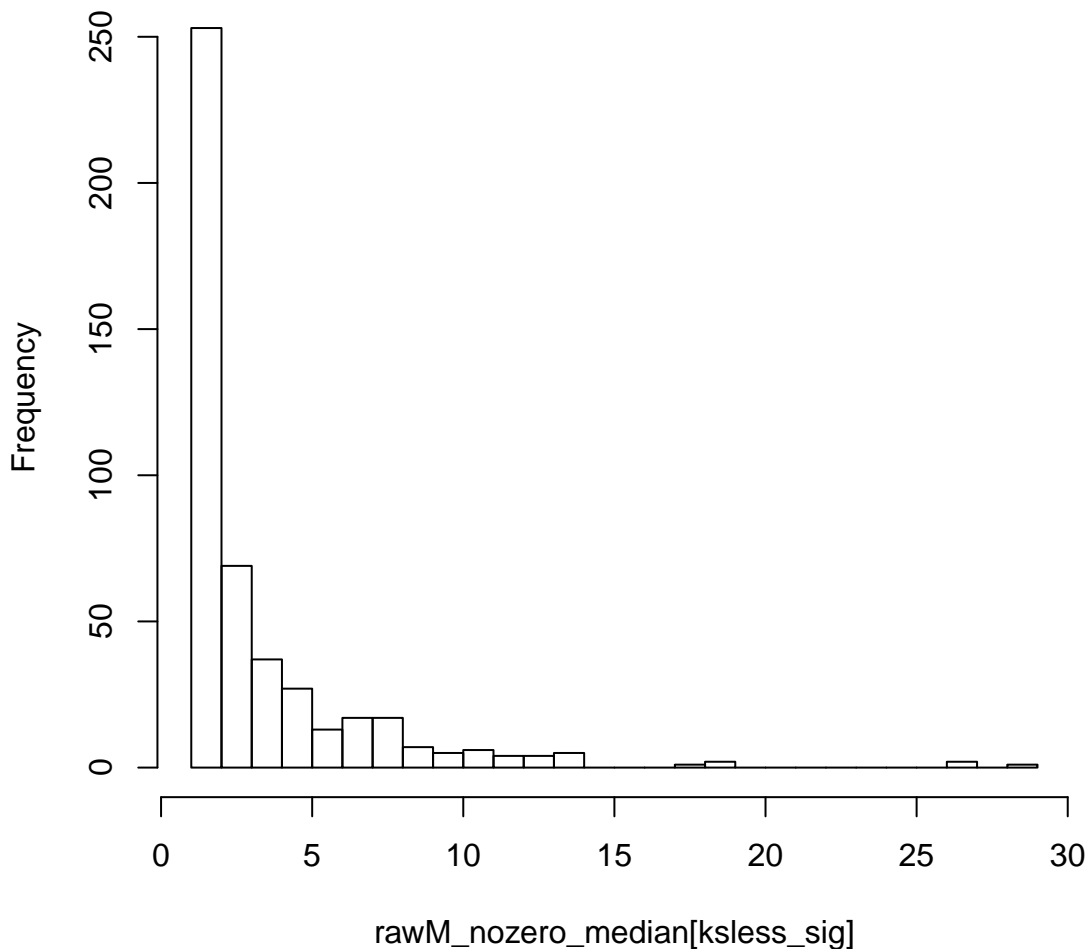


# median of nozero log-express of genes, kstwoside sig

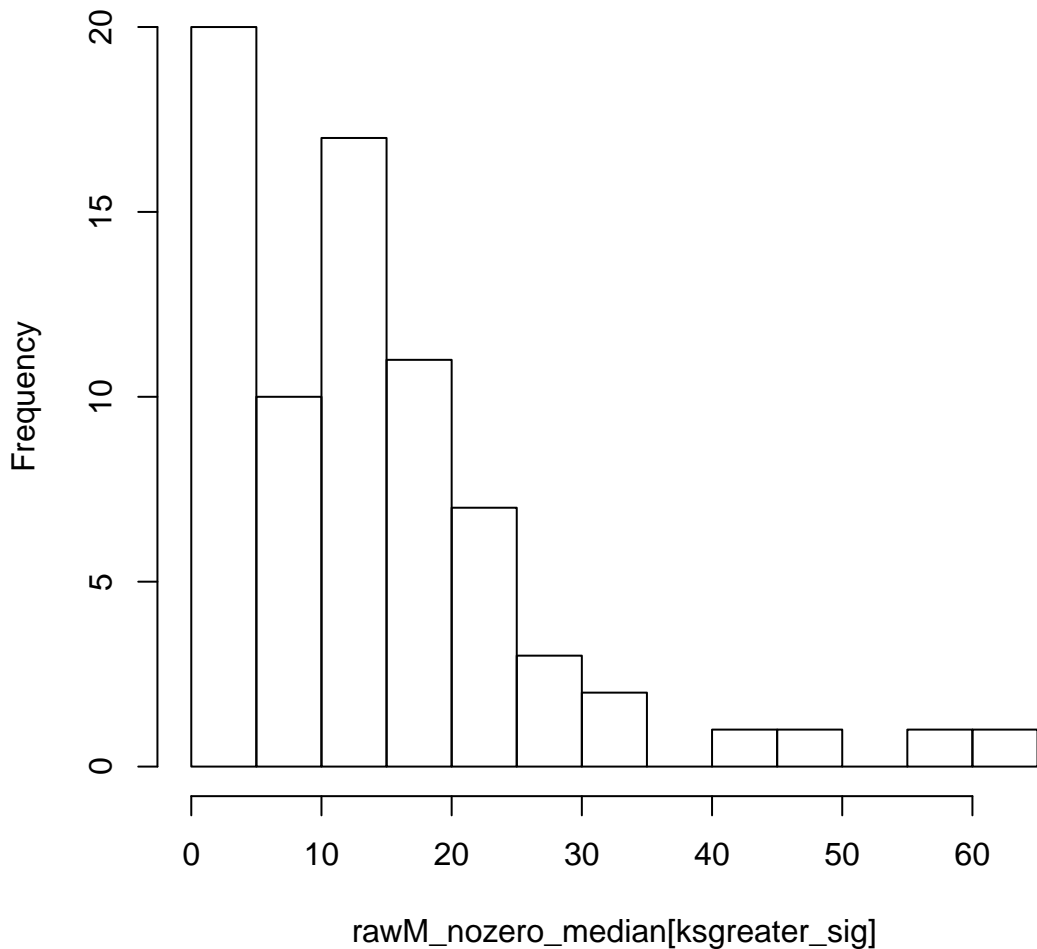




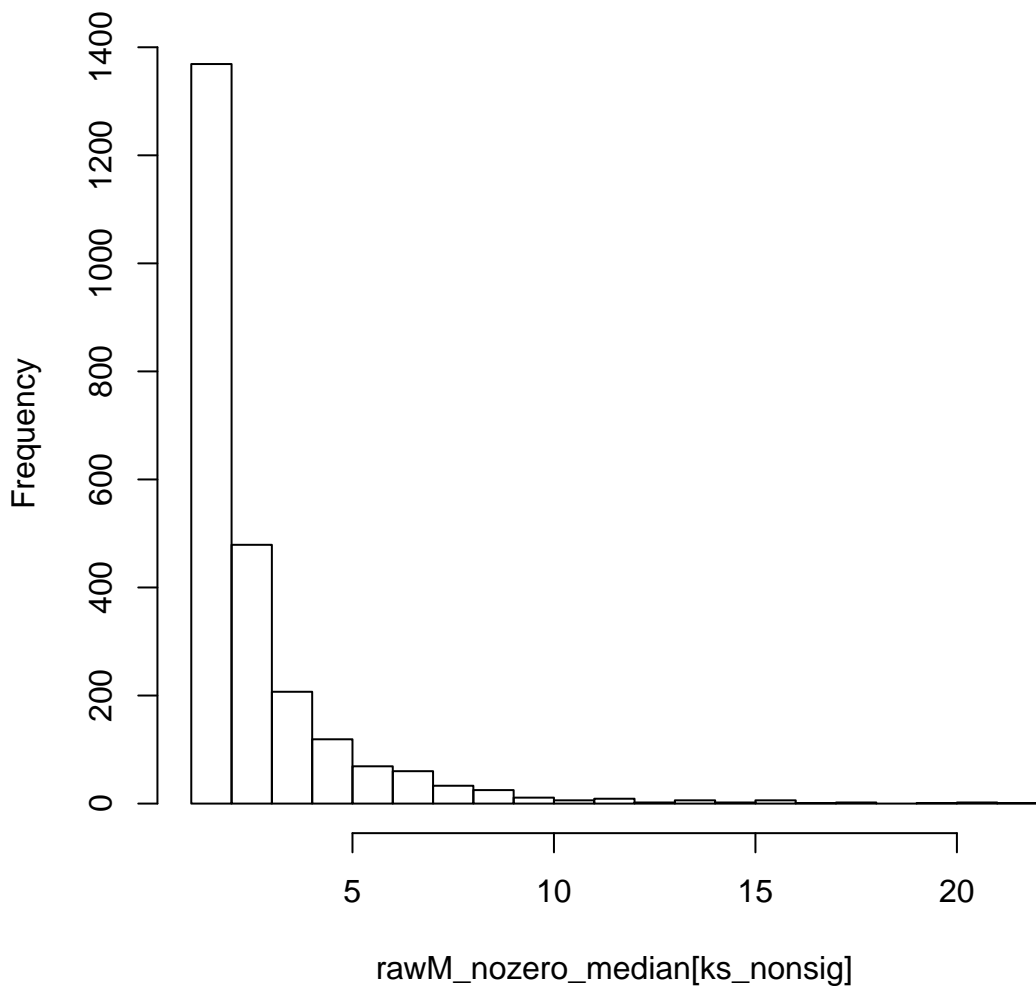
# median of nozero log-express of genes, ksless sig



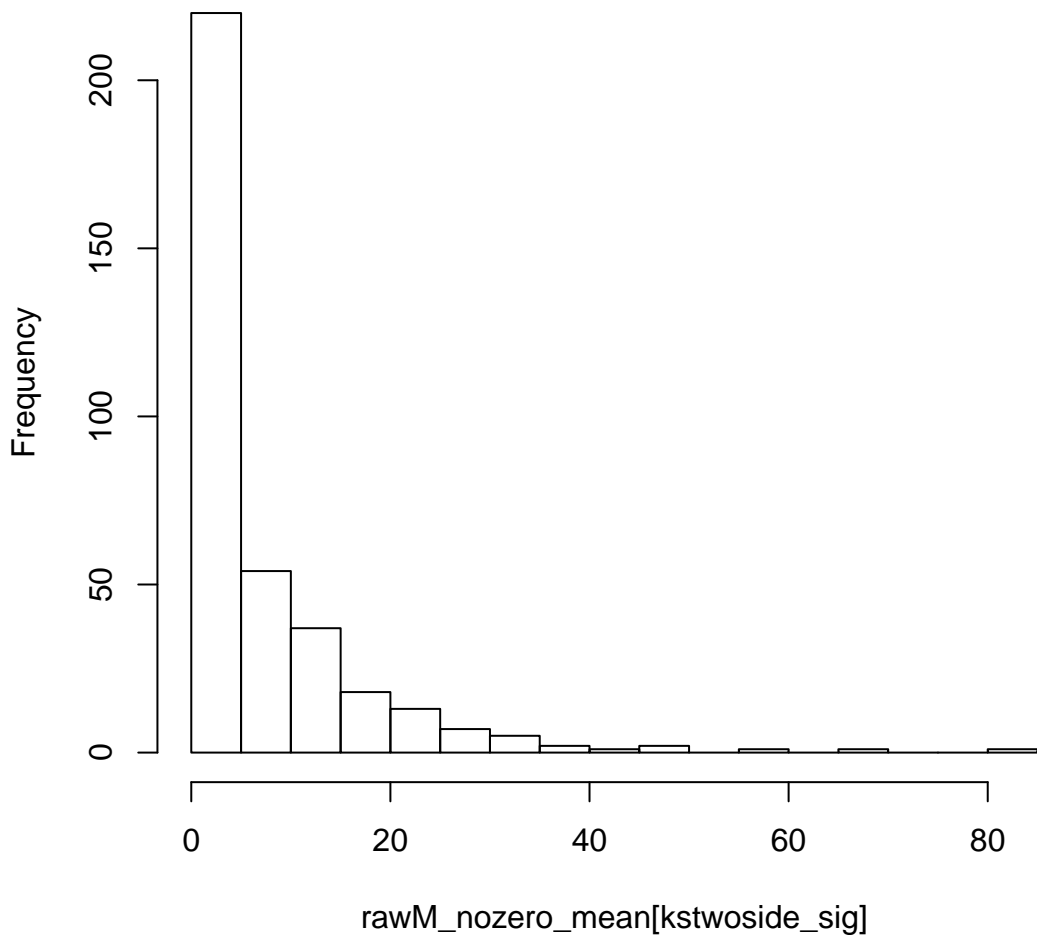
# median of nozero log-express of genes,ksgreater sig



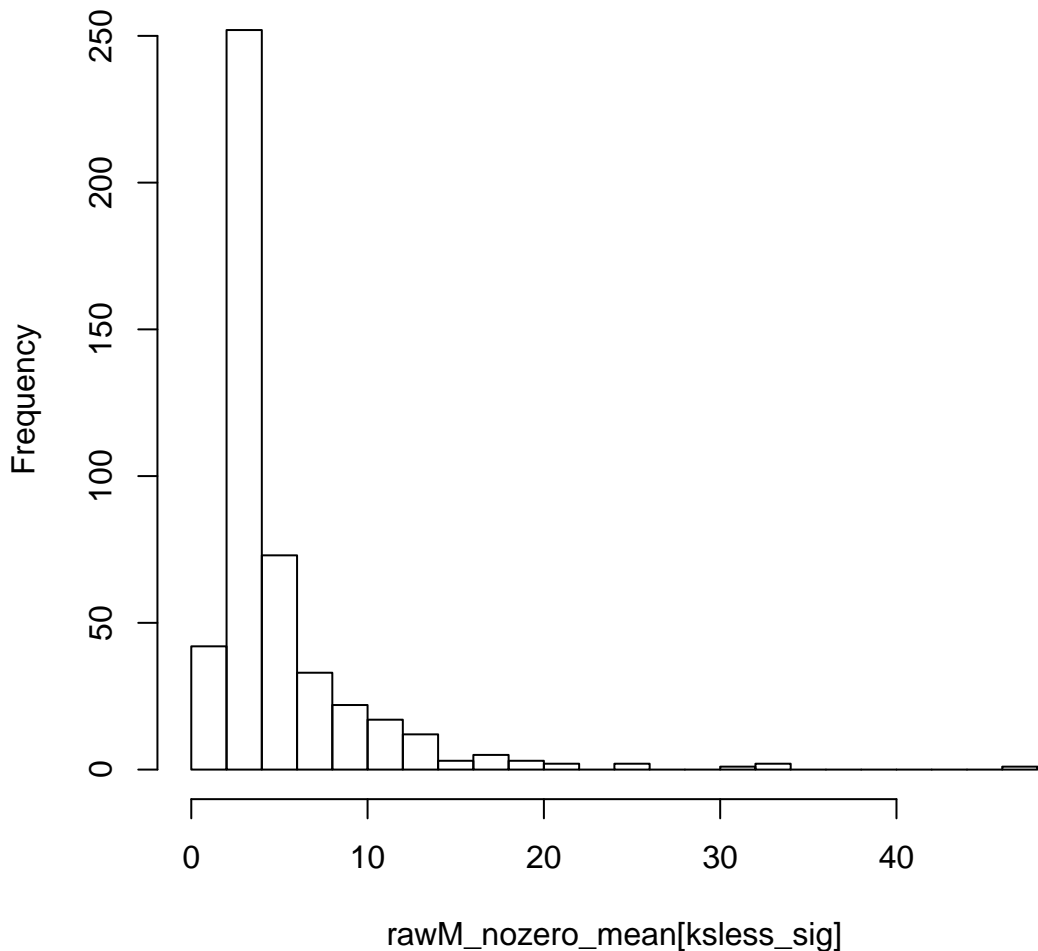
# median of nozero log-express of genes,ks no sig



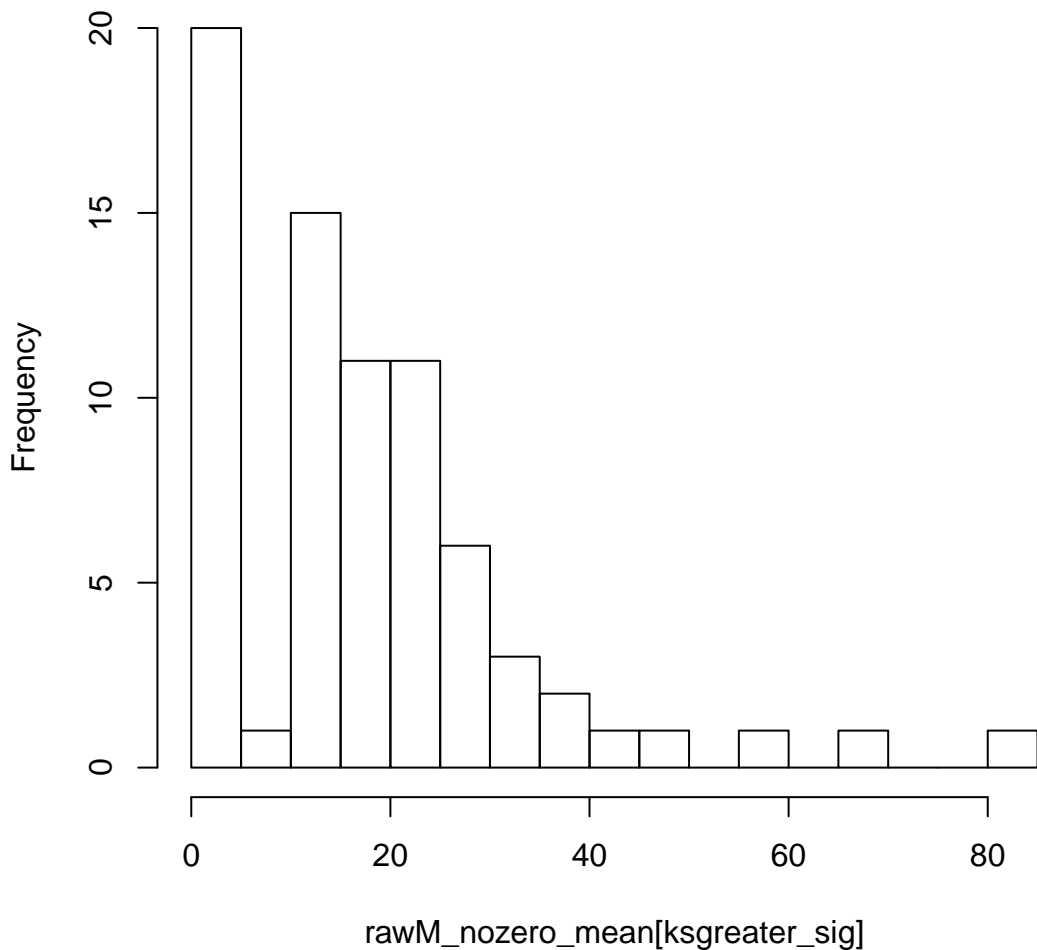
# mean of nozero log-express of genes, kstwoside sig



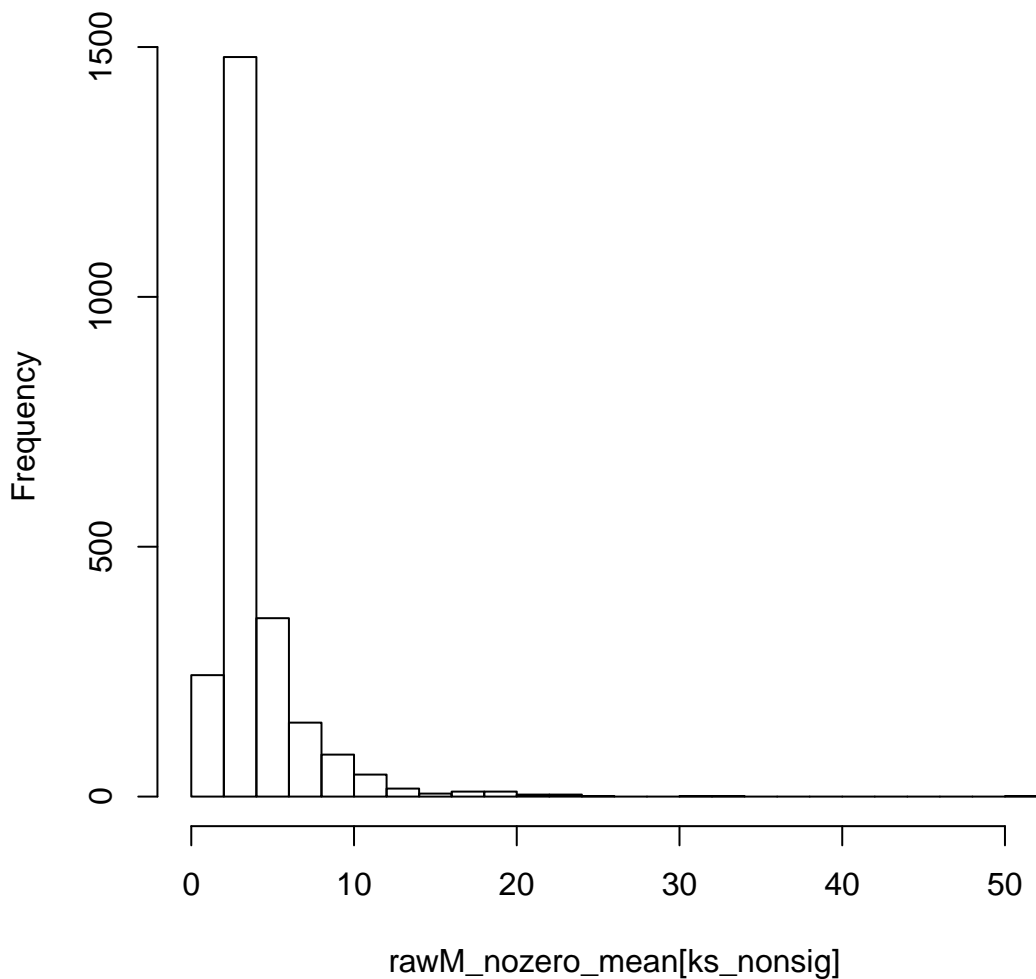
# mean of nozero log-express of genes, ksless sig



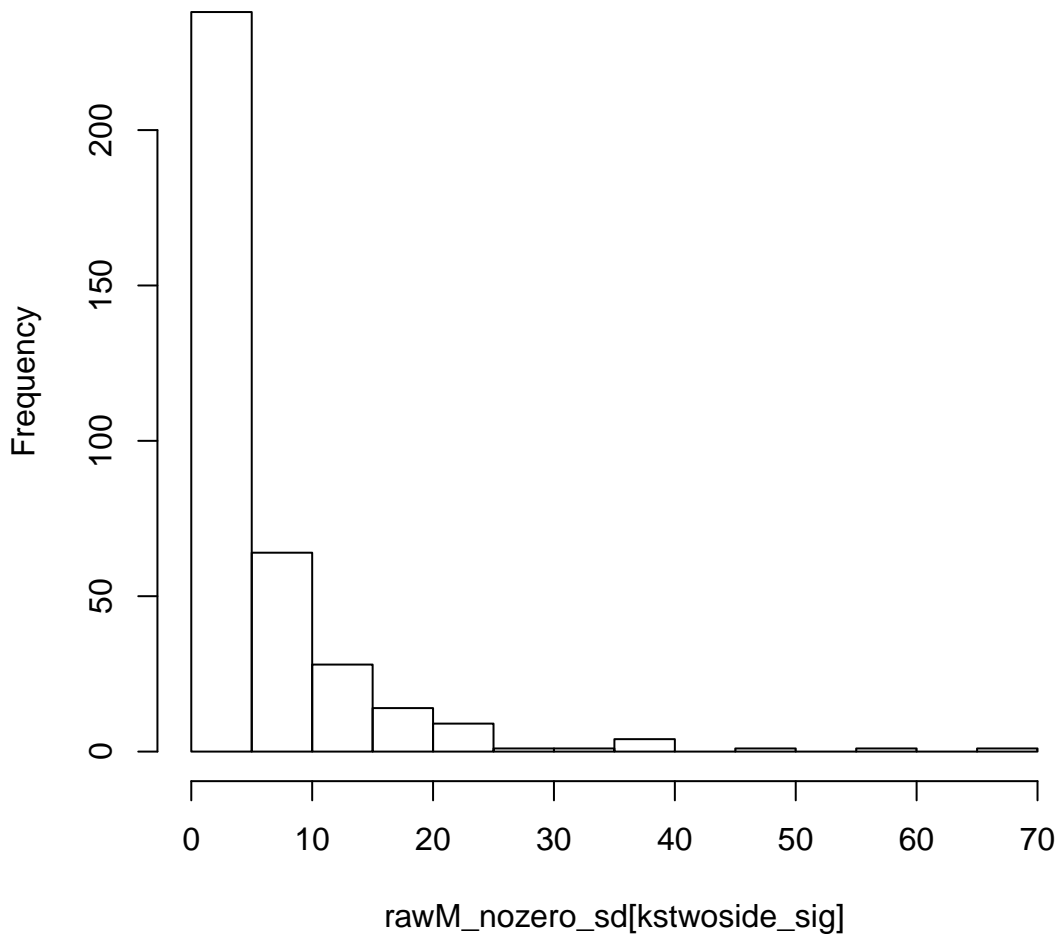
# mean of nozero log-express of genes,ksgreater sig



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

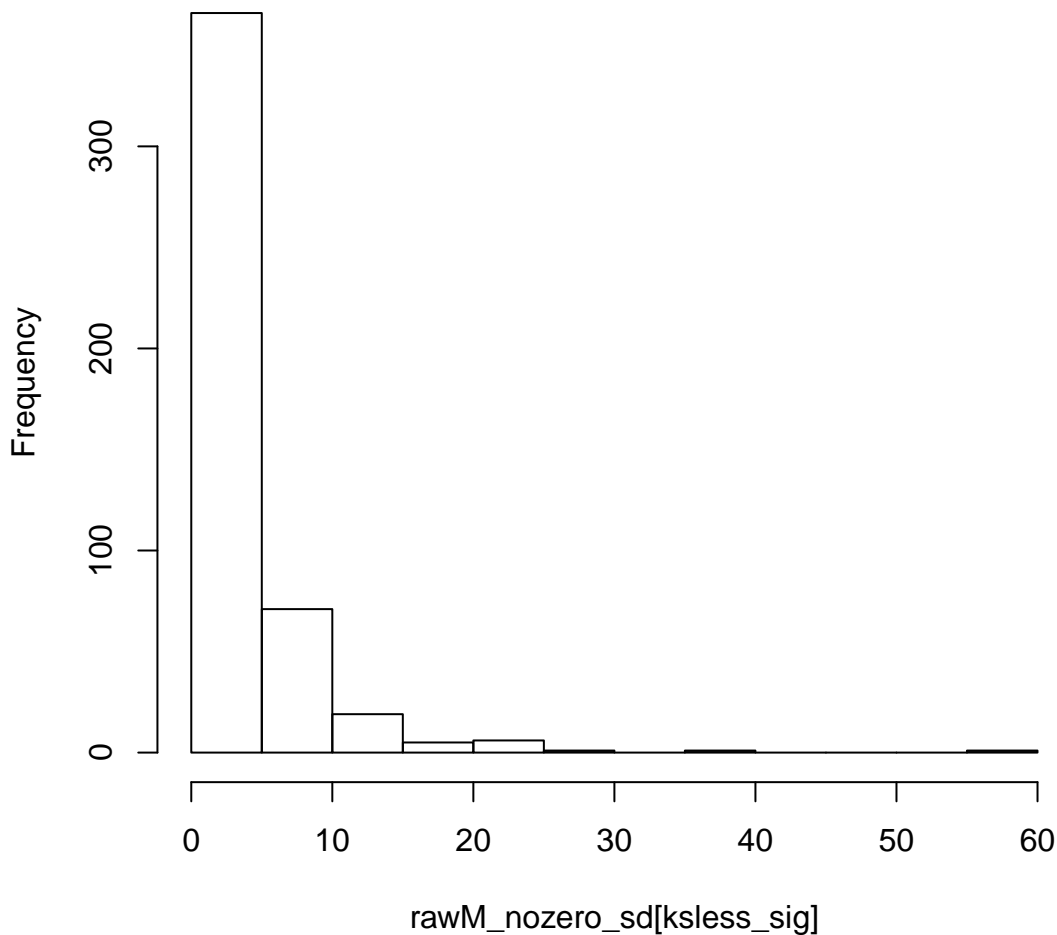


# sd of nozero log-express of genes, kstwoside sig

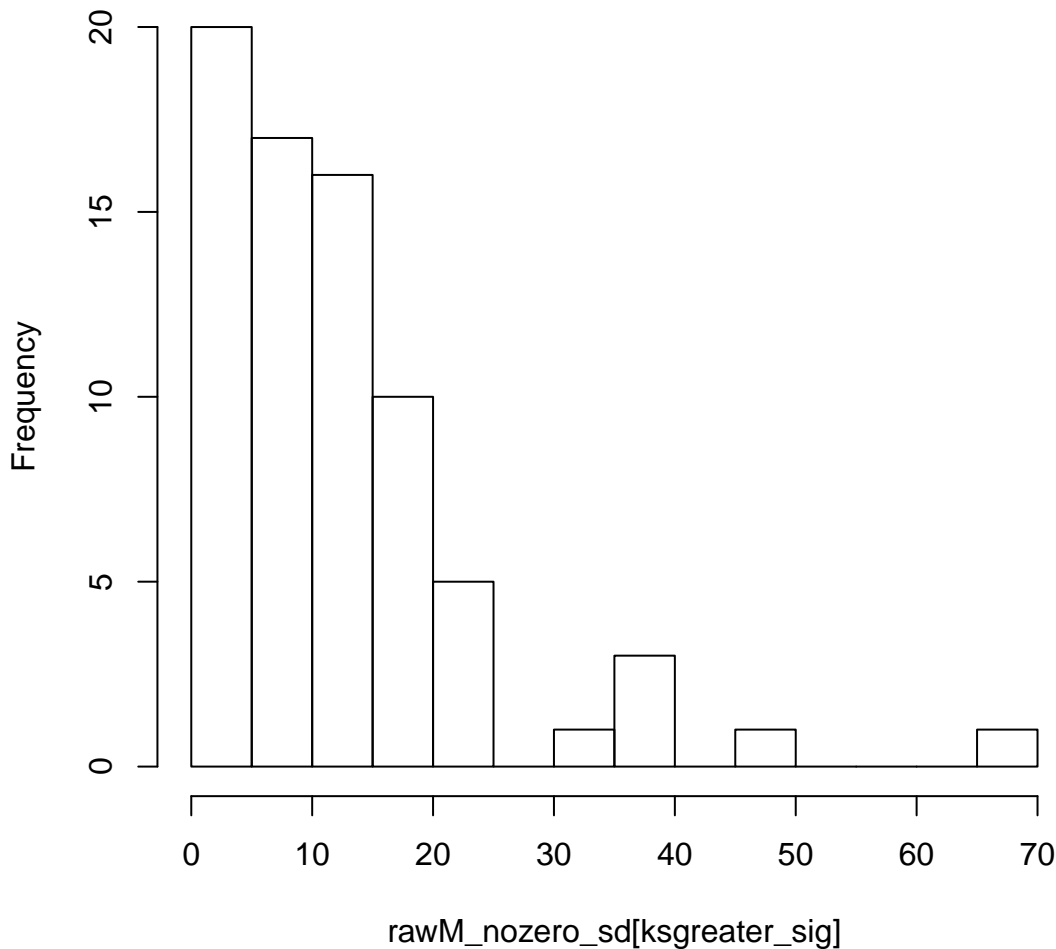




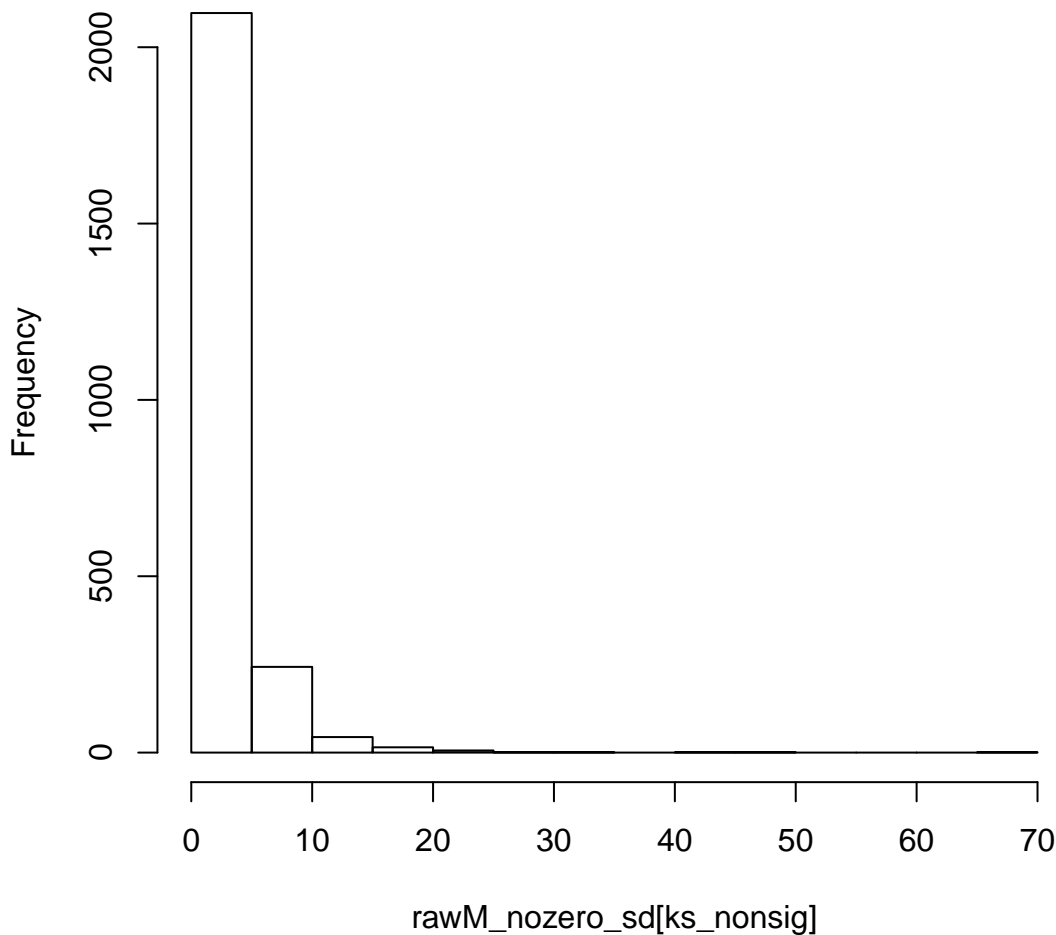
# sd of nozero log-express of genes, ksless sig



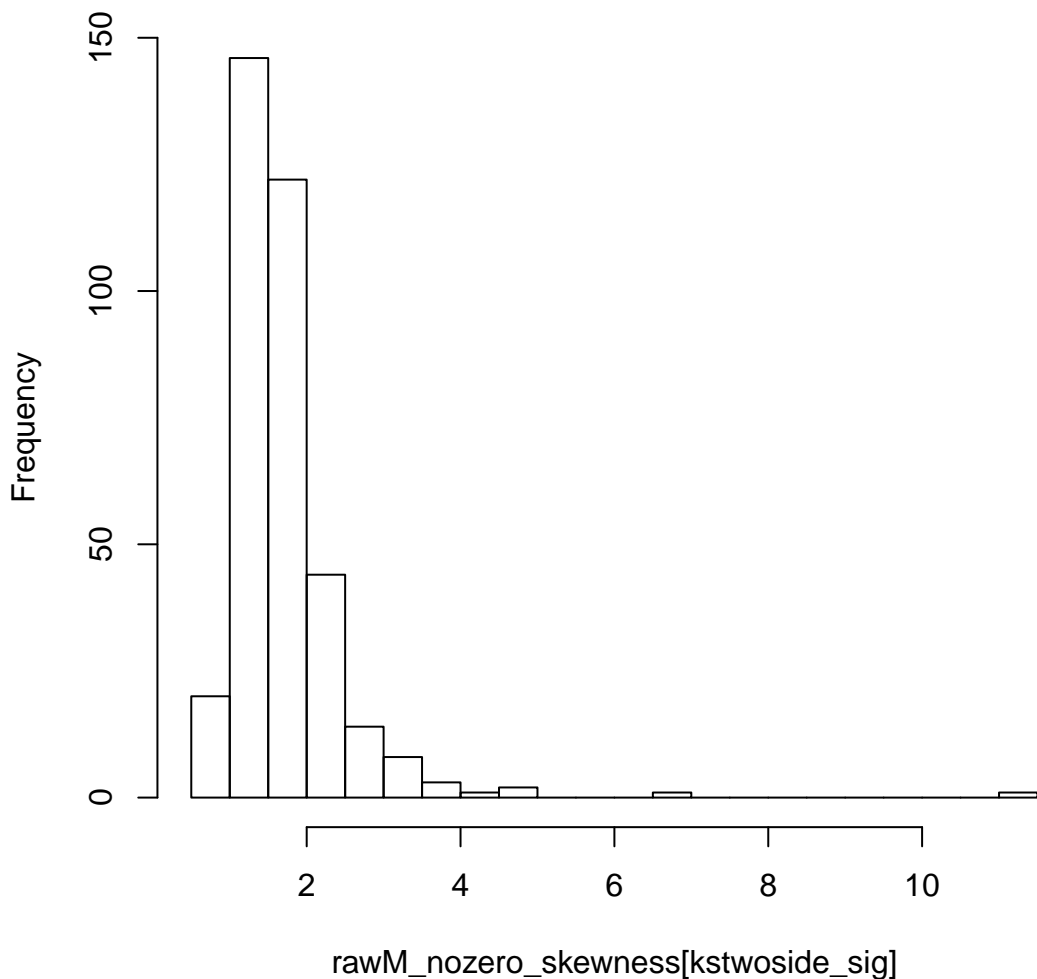
# sd of nozero log-express of genes,ksgreater sig



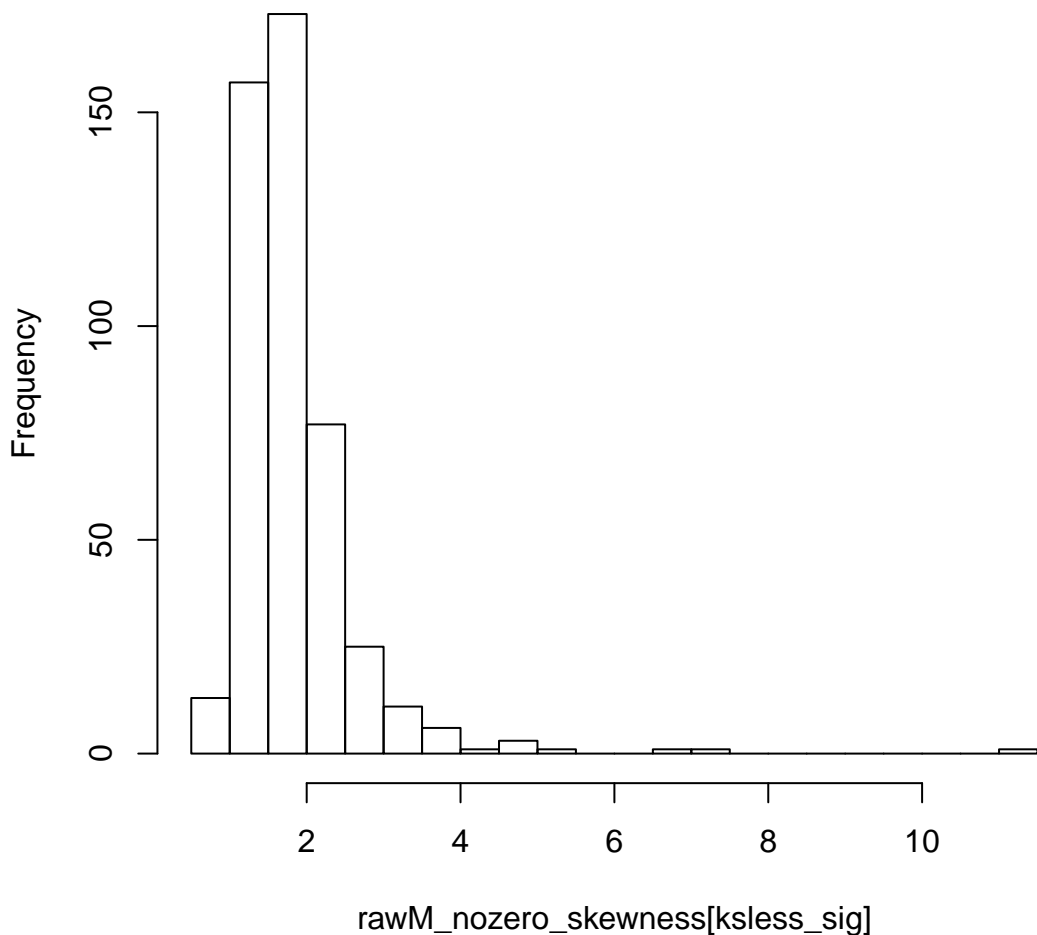
# sd of nozero log-express of genes,ks no sig



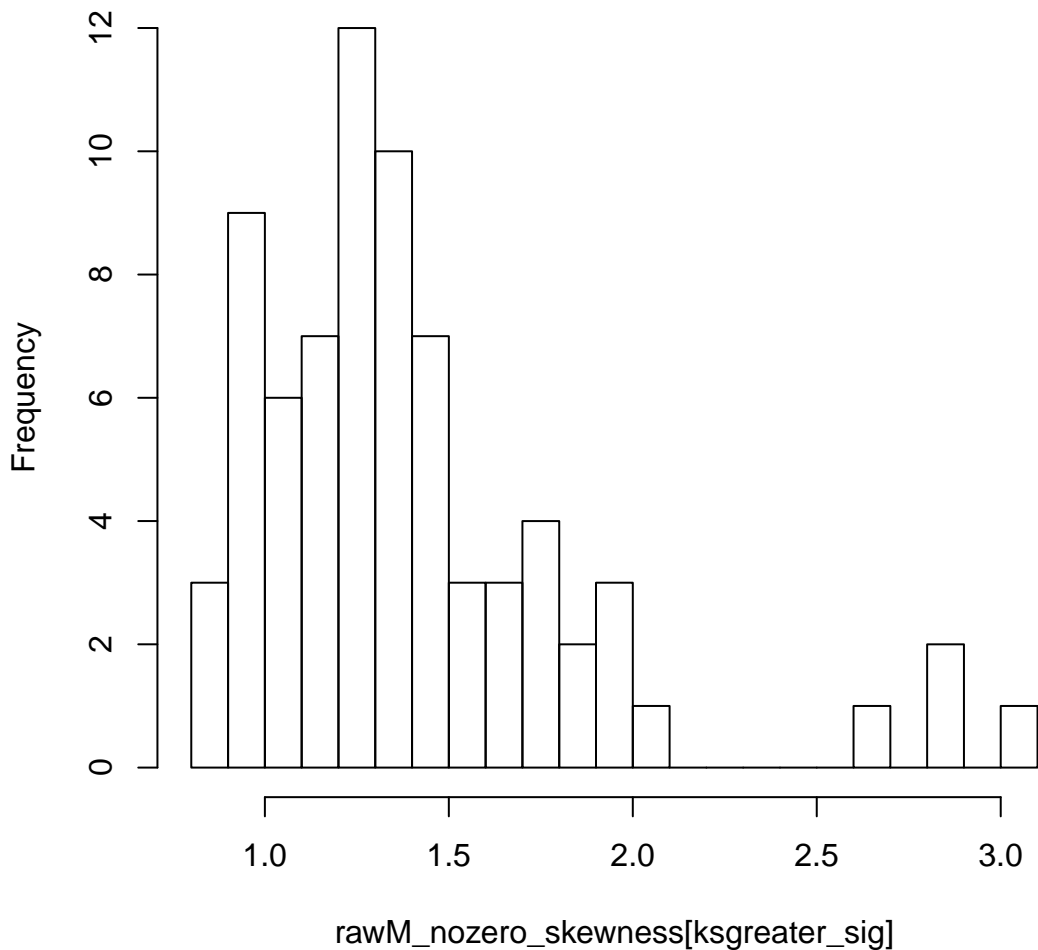
# skewness of nozero log-expres of genes, kstvoside sig



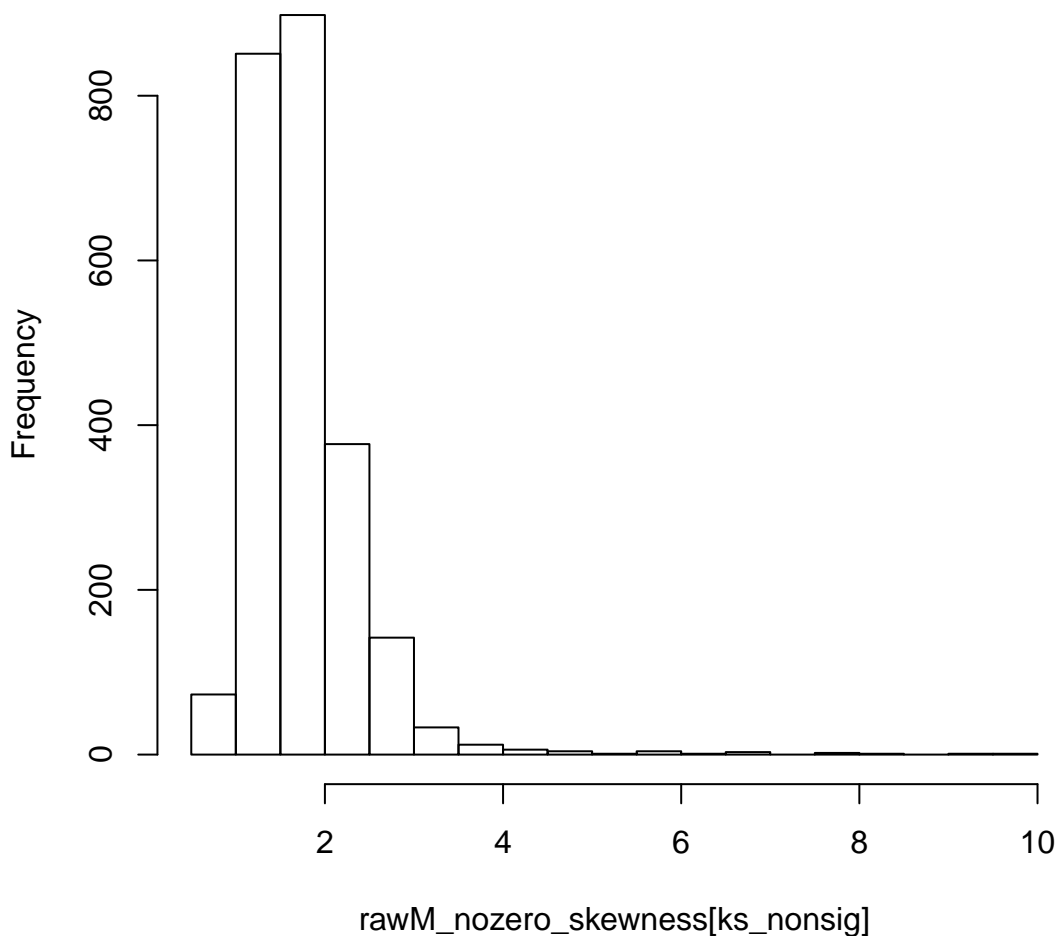
# skewness of nozero log-express of genes, ksless sig



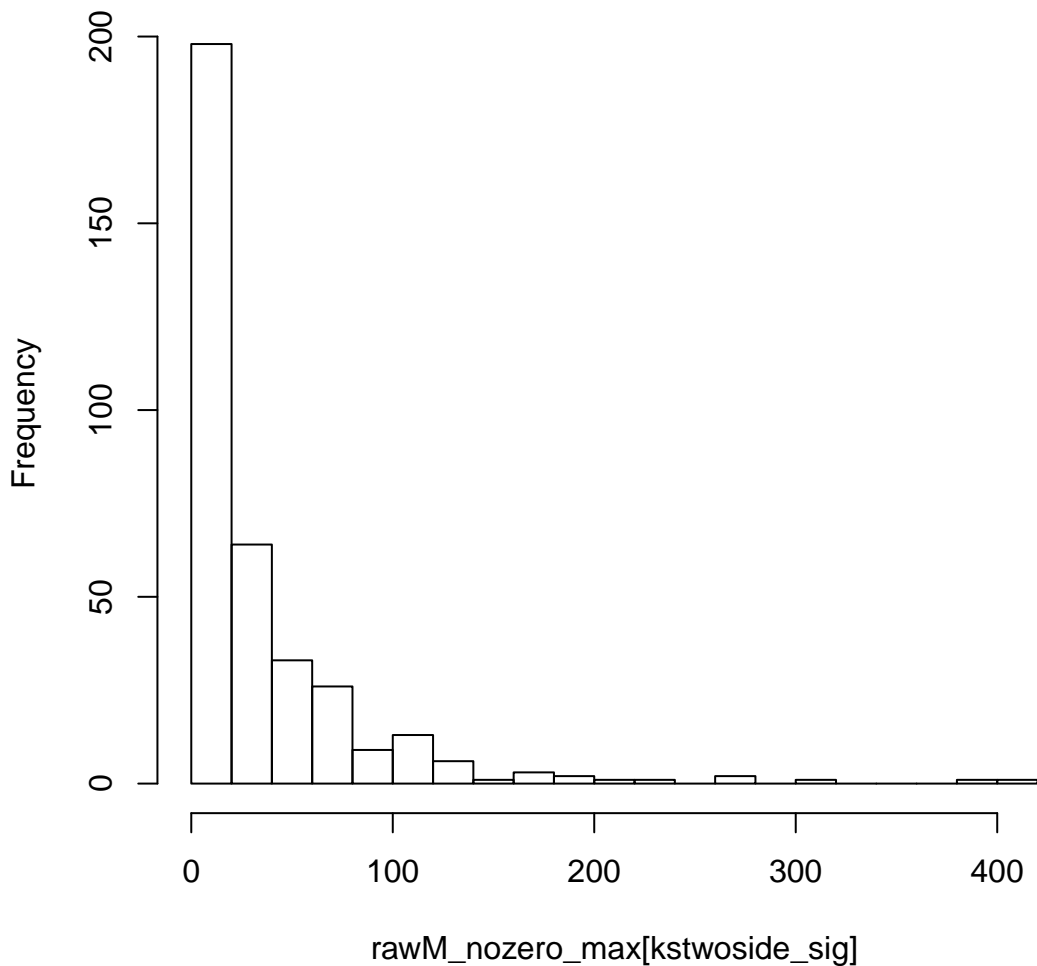
# skewness of nozero log-express of genes,ksgreater sig



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

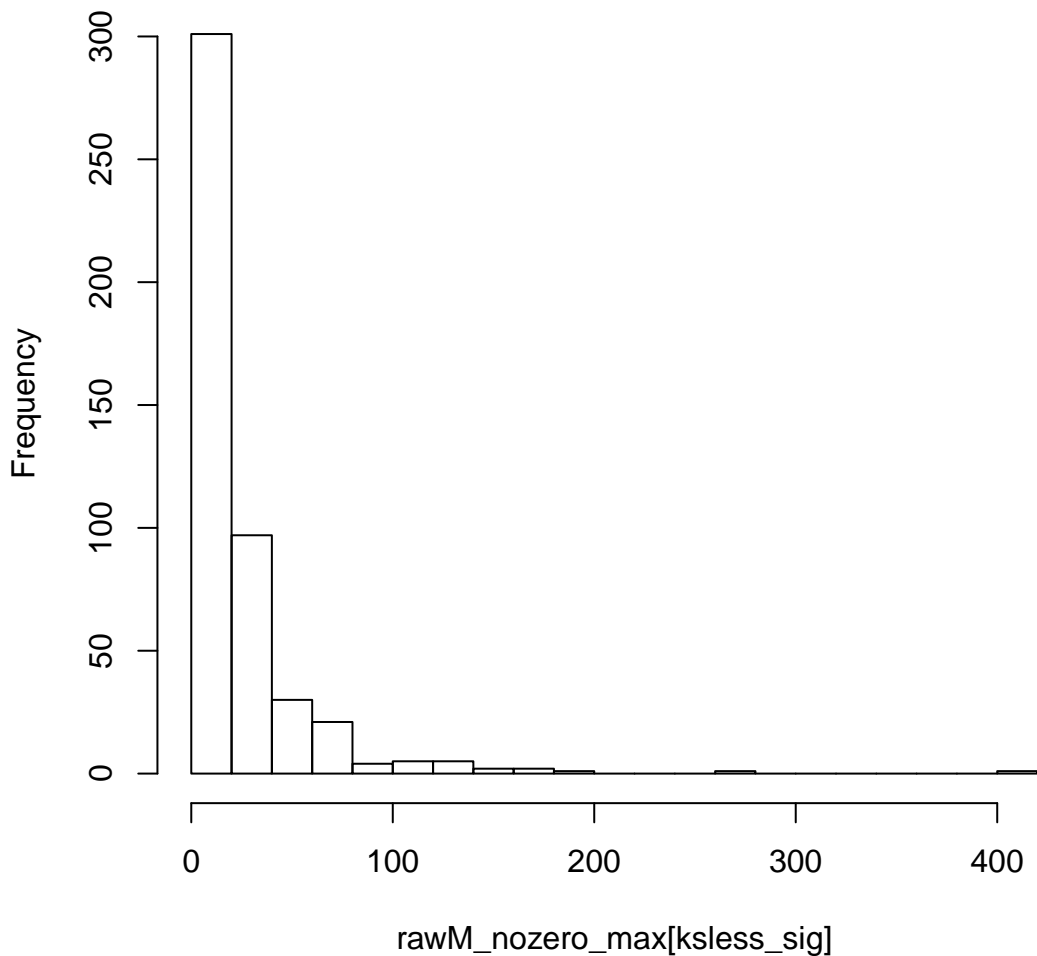


# max of nozero log-express of genes, kstwoside sig

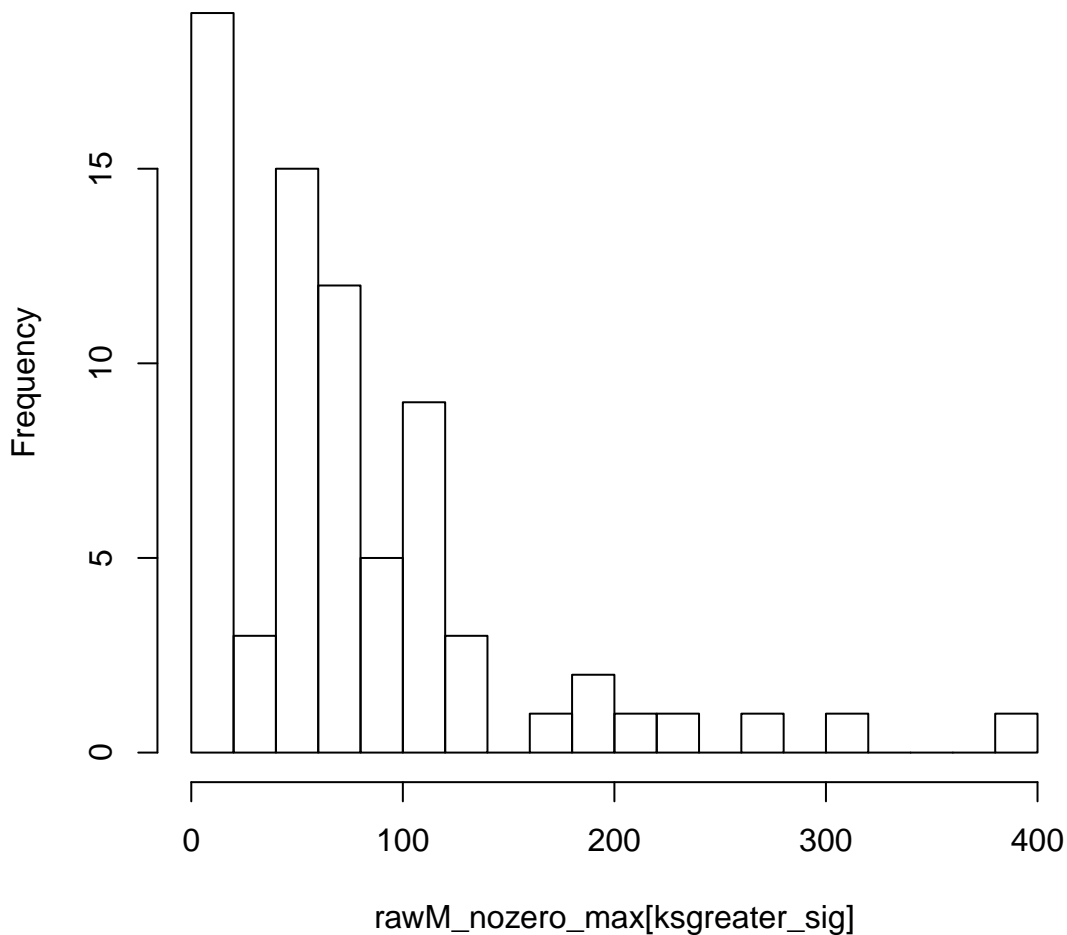




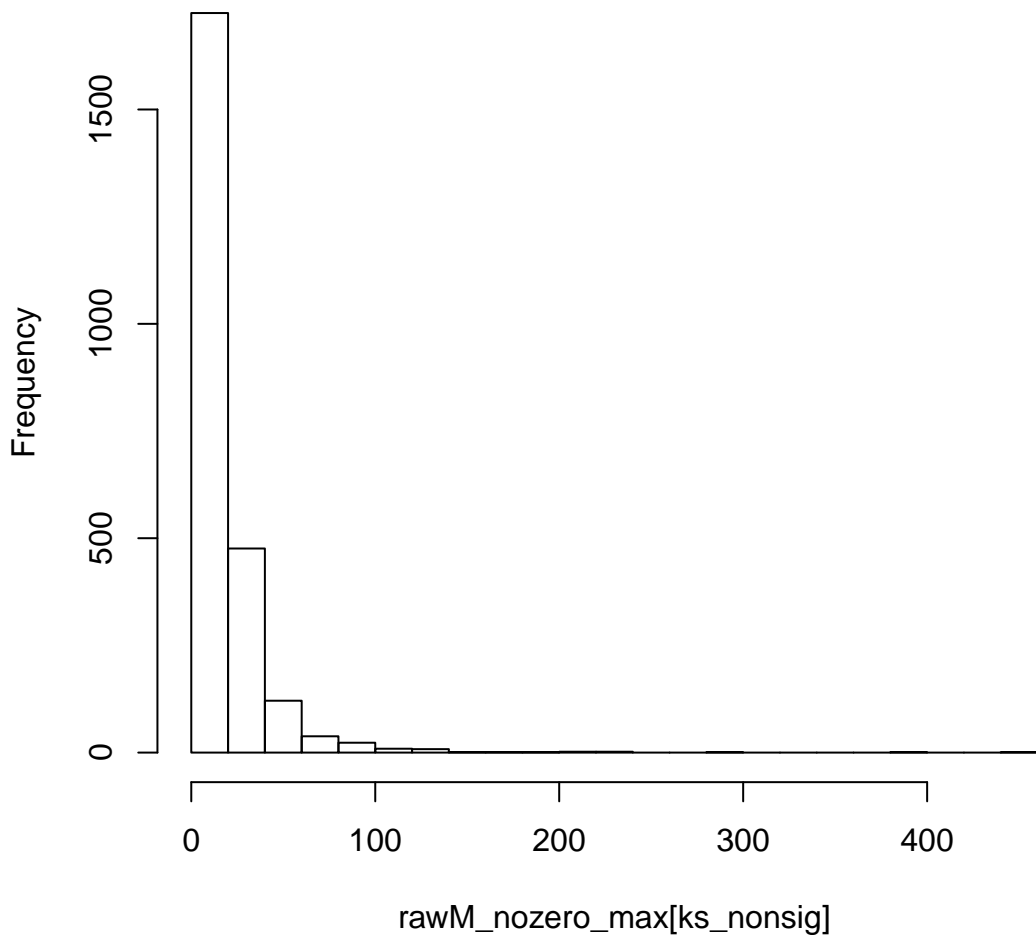
# max of nozero log-express of genes, ksless sig



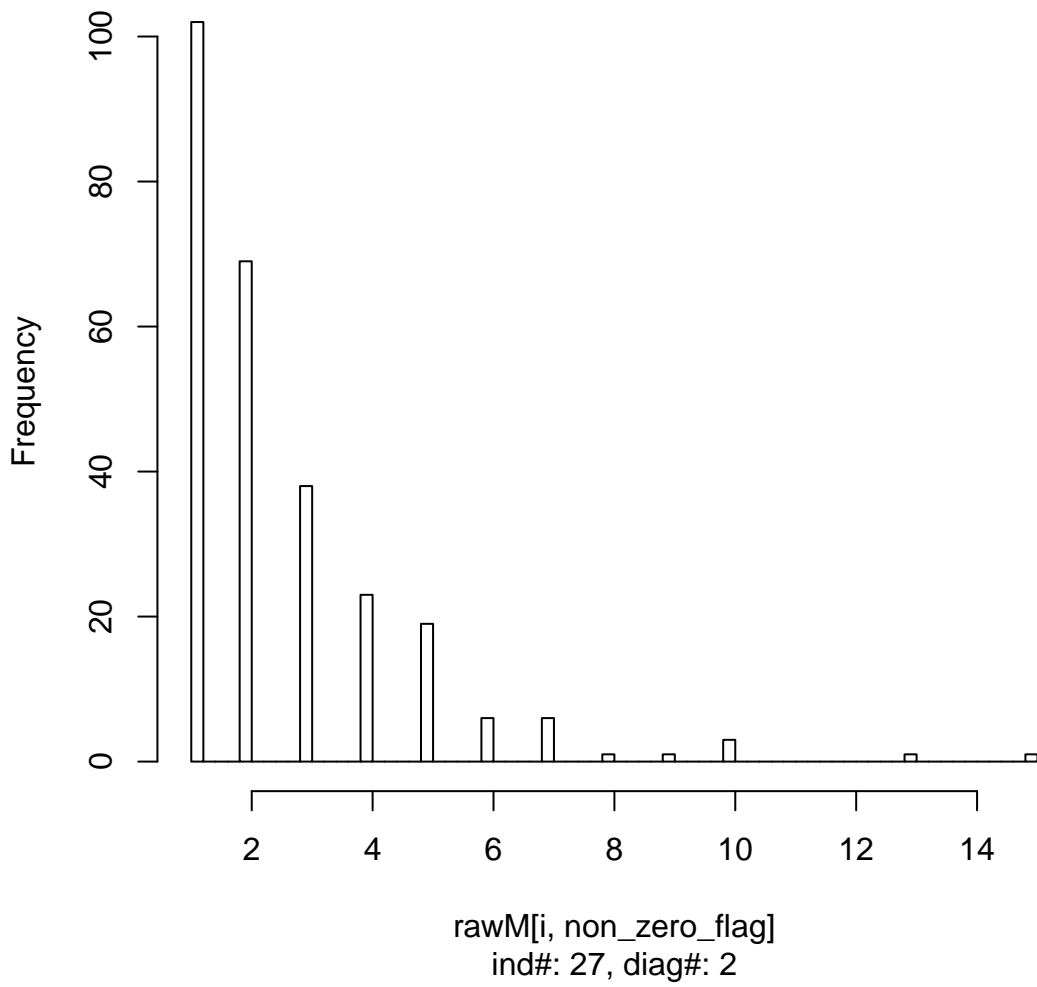
# max of nozero log-express of genes,ksgreater sig



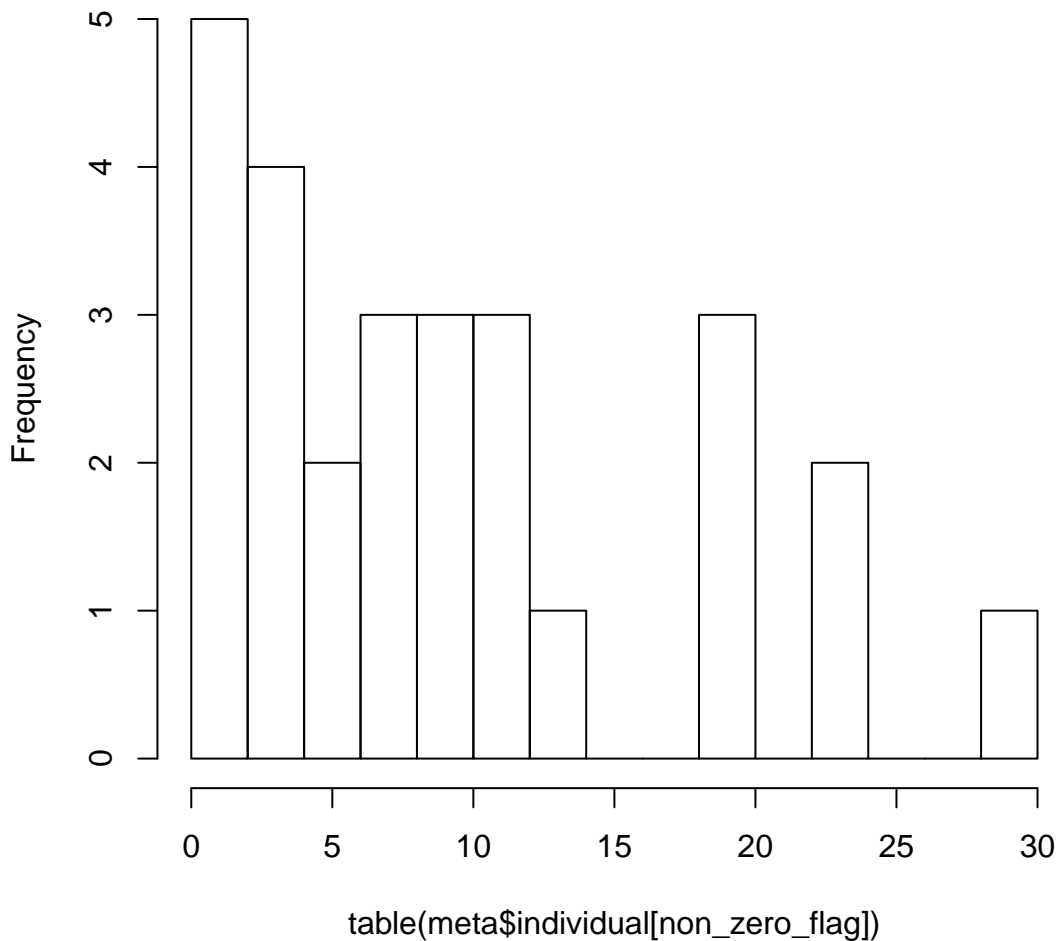
# max of nozero log-expres of genes,ks no sig



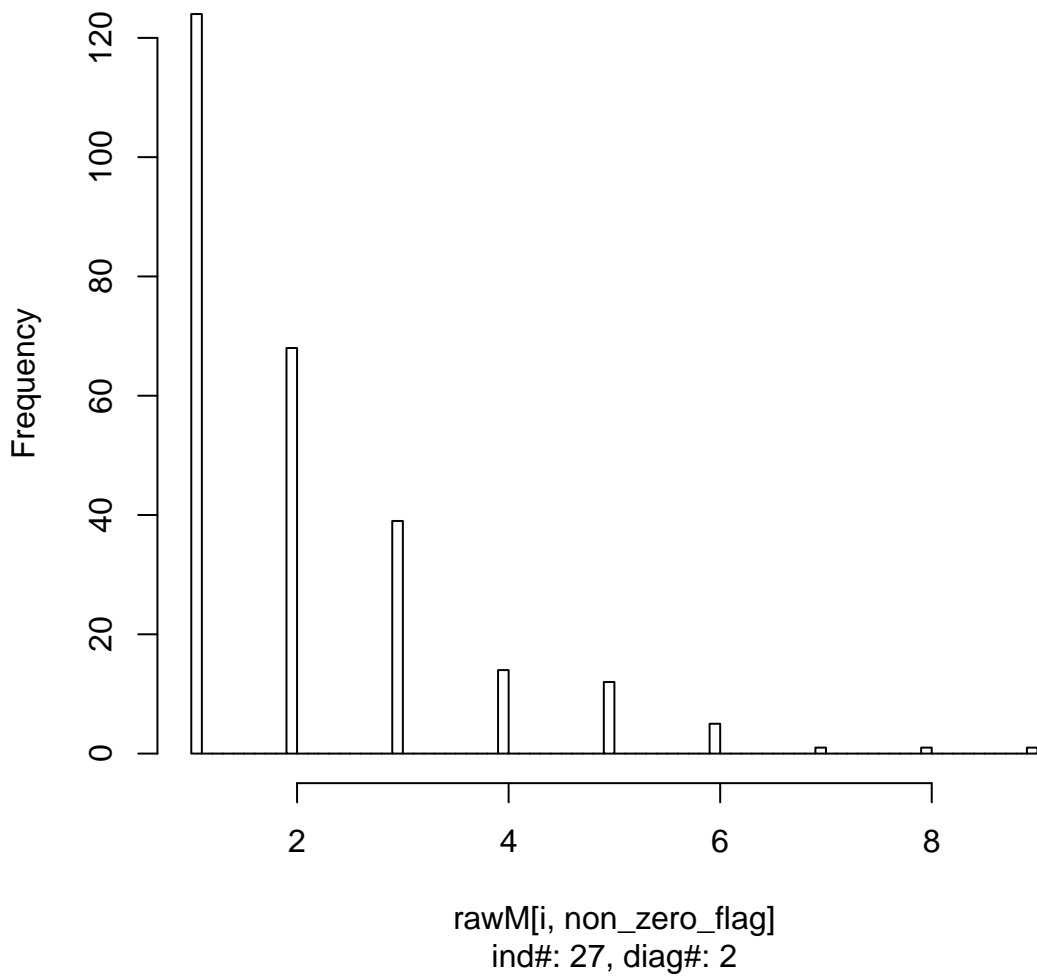
less sig: log expression of gene#2, pval ob=0.4834, non-zero nu



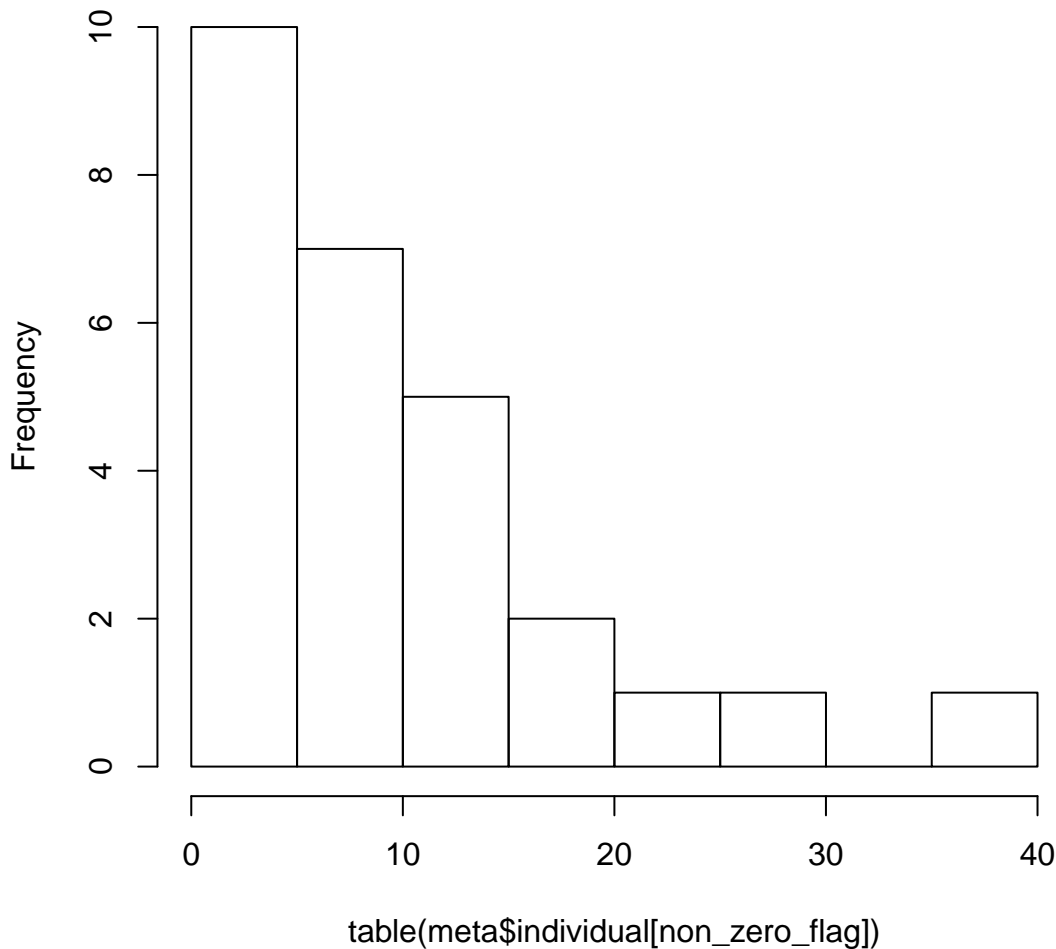
# KSless sig: individual expression cell count of gene#2



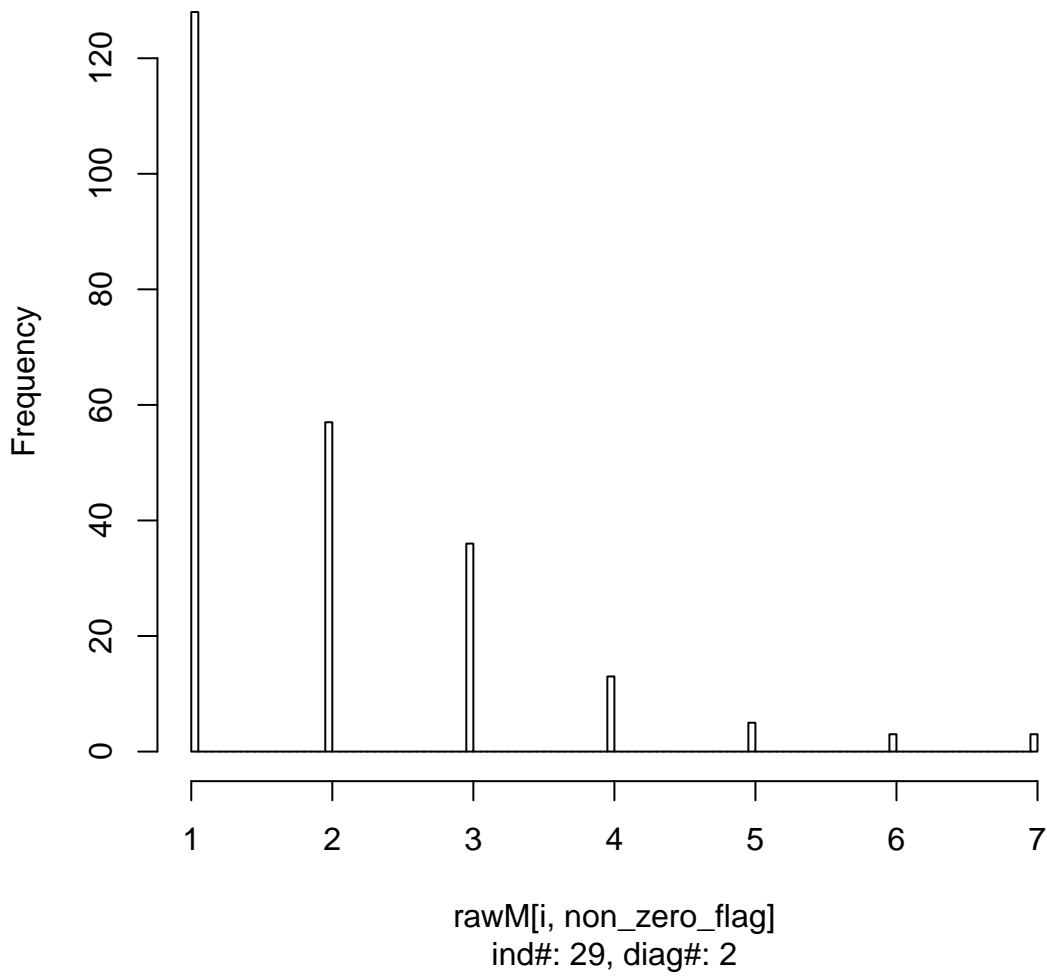
Sless sig: log expression of gene#3, pval ob=0.934, non-zero num



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

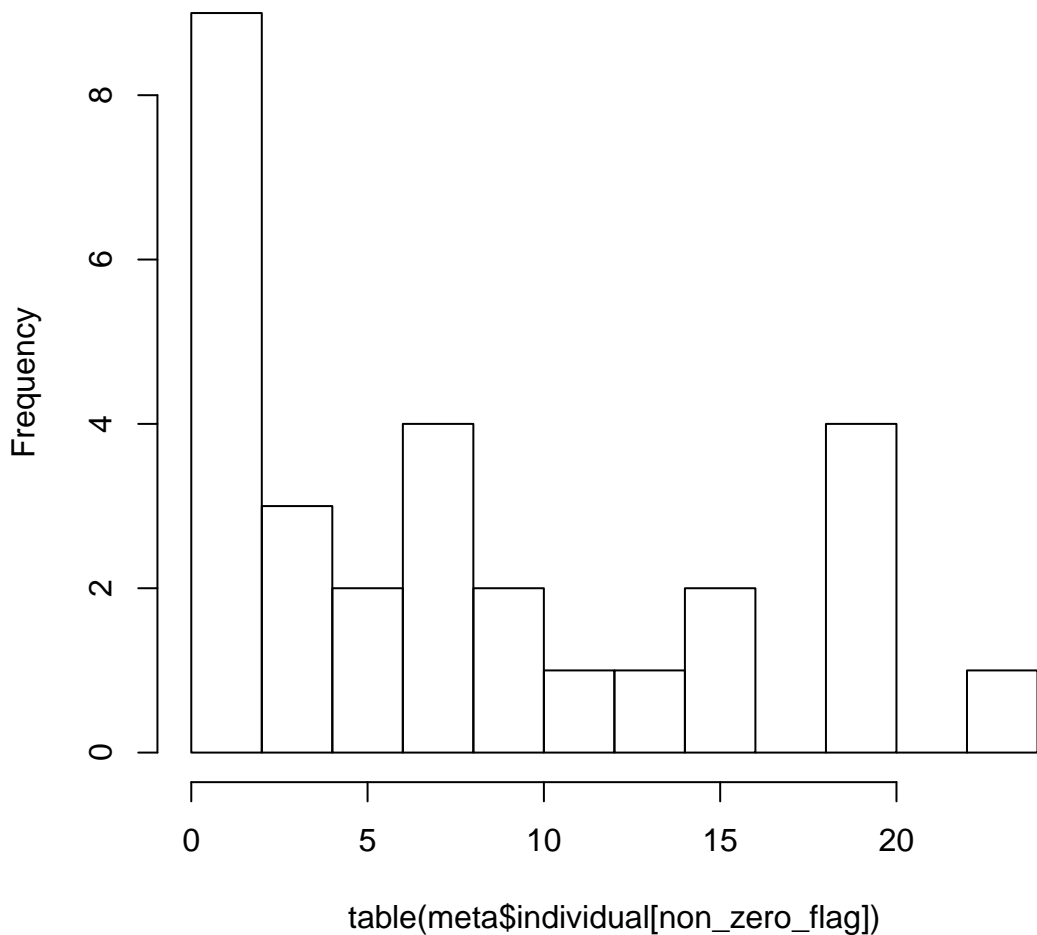


less sig: log expression of gene#4, pval ob=0.2531, non-zero nu

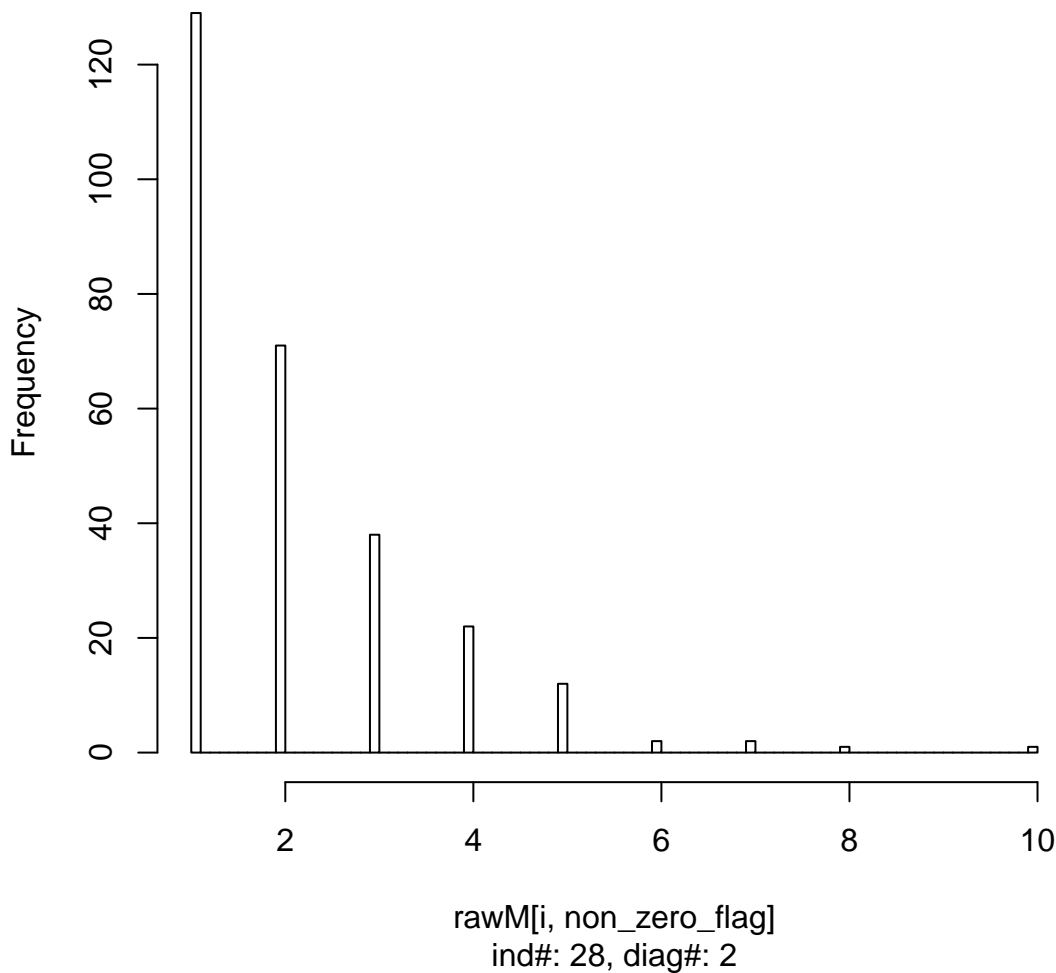




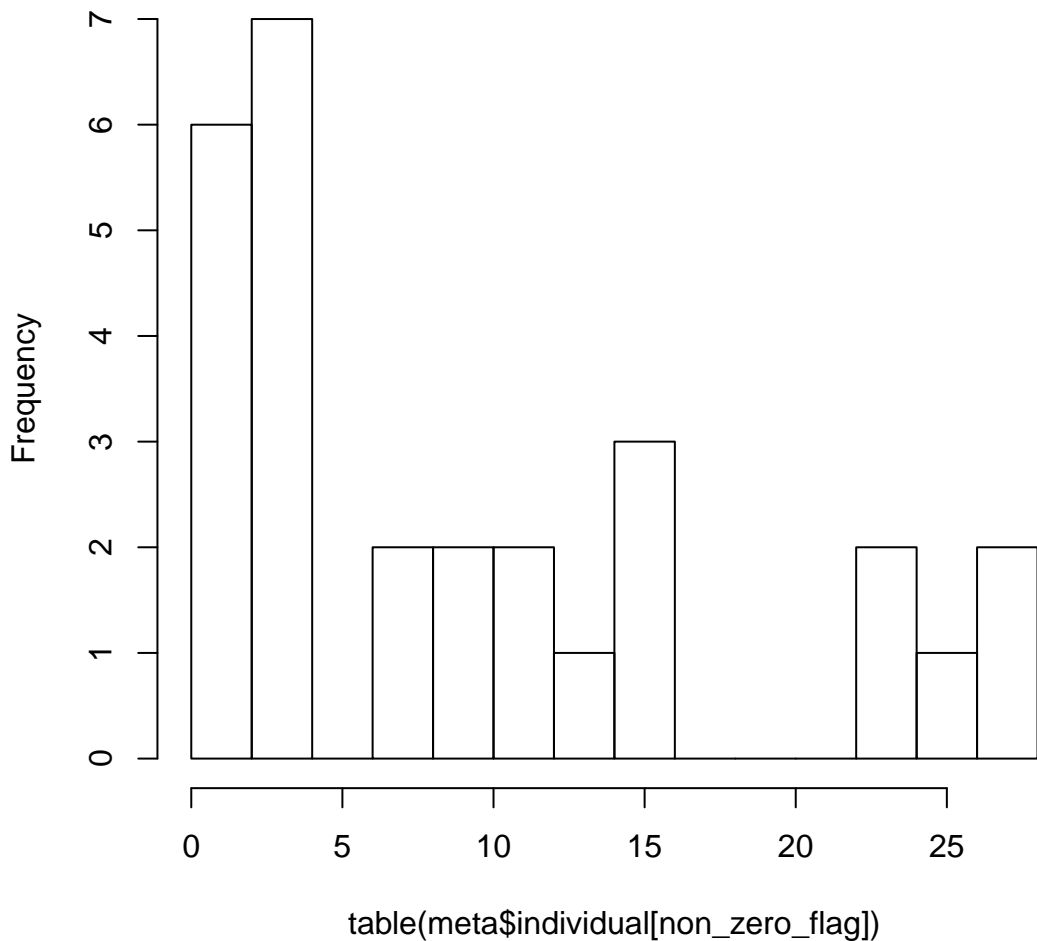
# KSless sig: individual expression cell count of gene#4



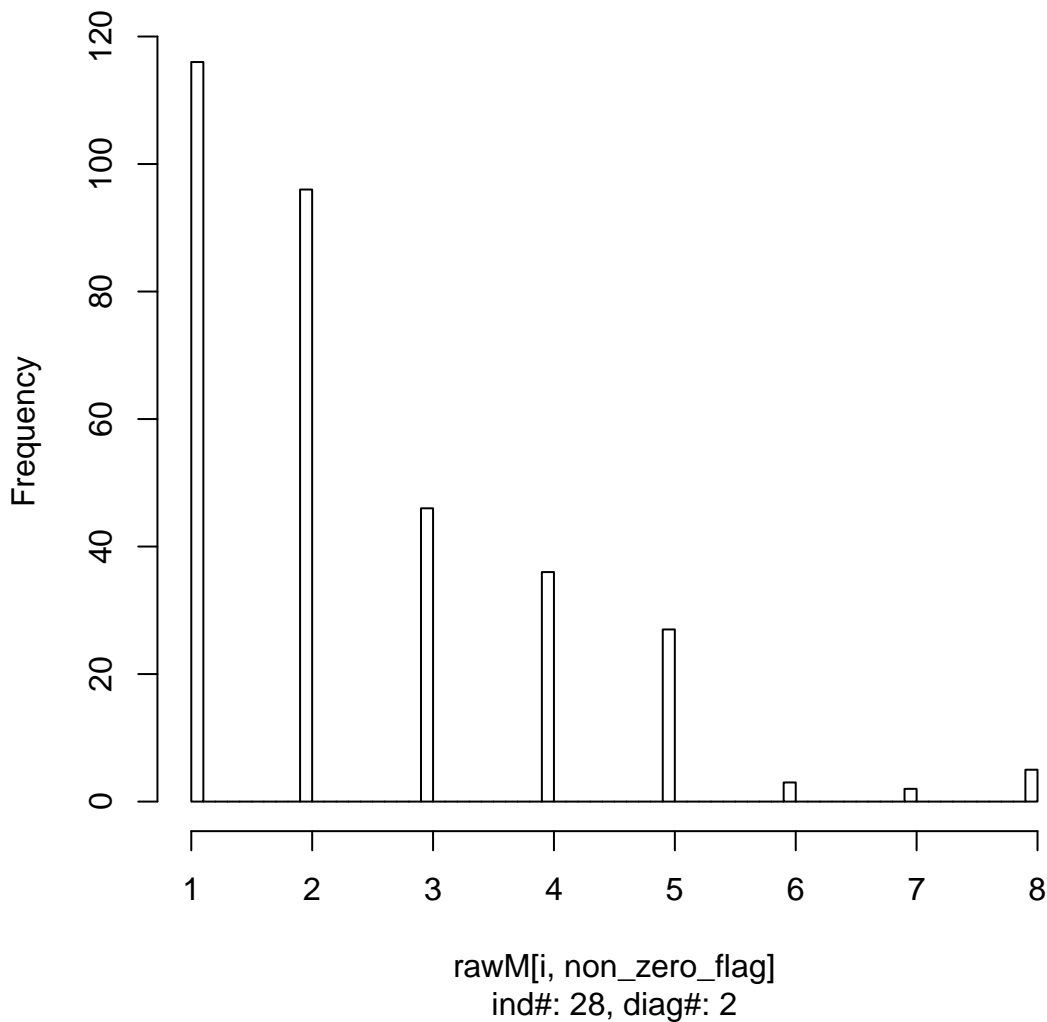
less sig: log expression of gene#9, pval ob=0.0111, non-zero nu



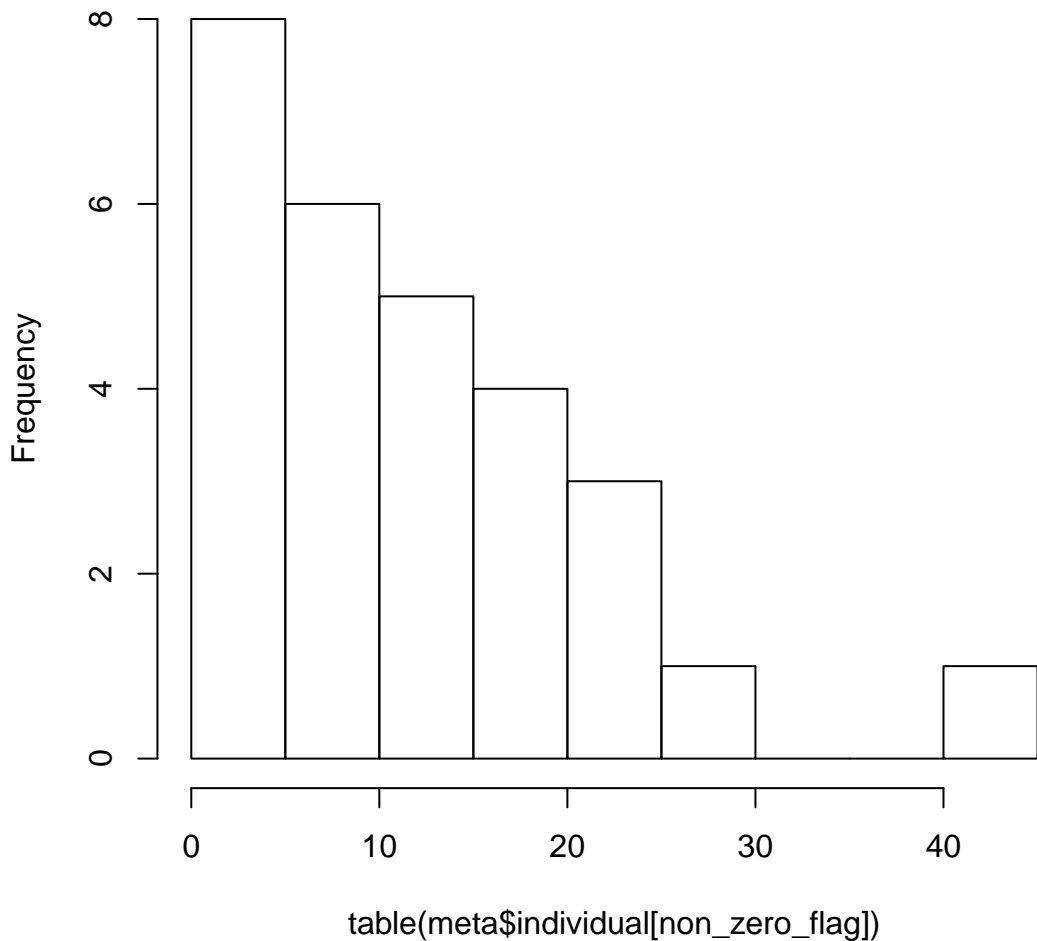
# KSless sig: individual expression cell count of gene#9



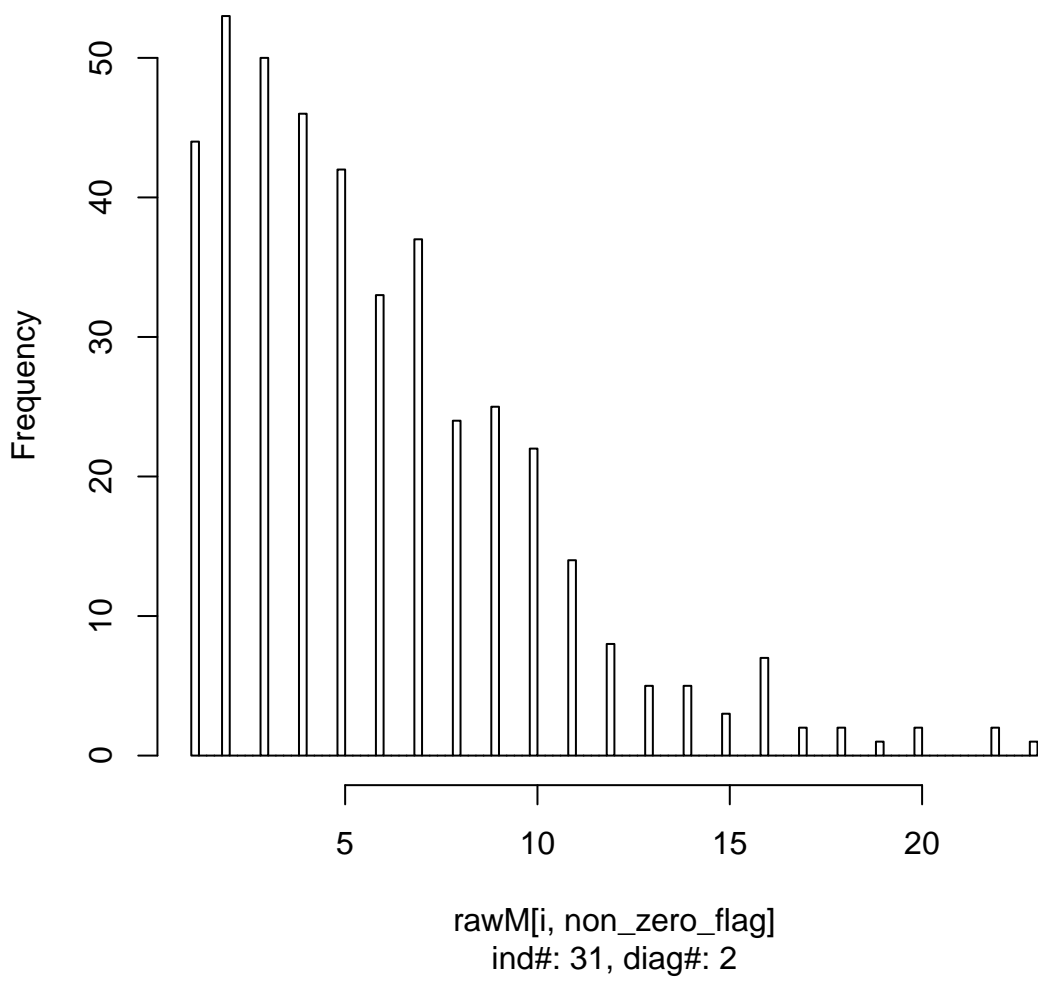
less sig: log expression of gene#12, pval ob=0.2318, non-zero nu



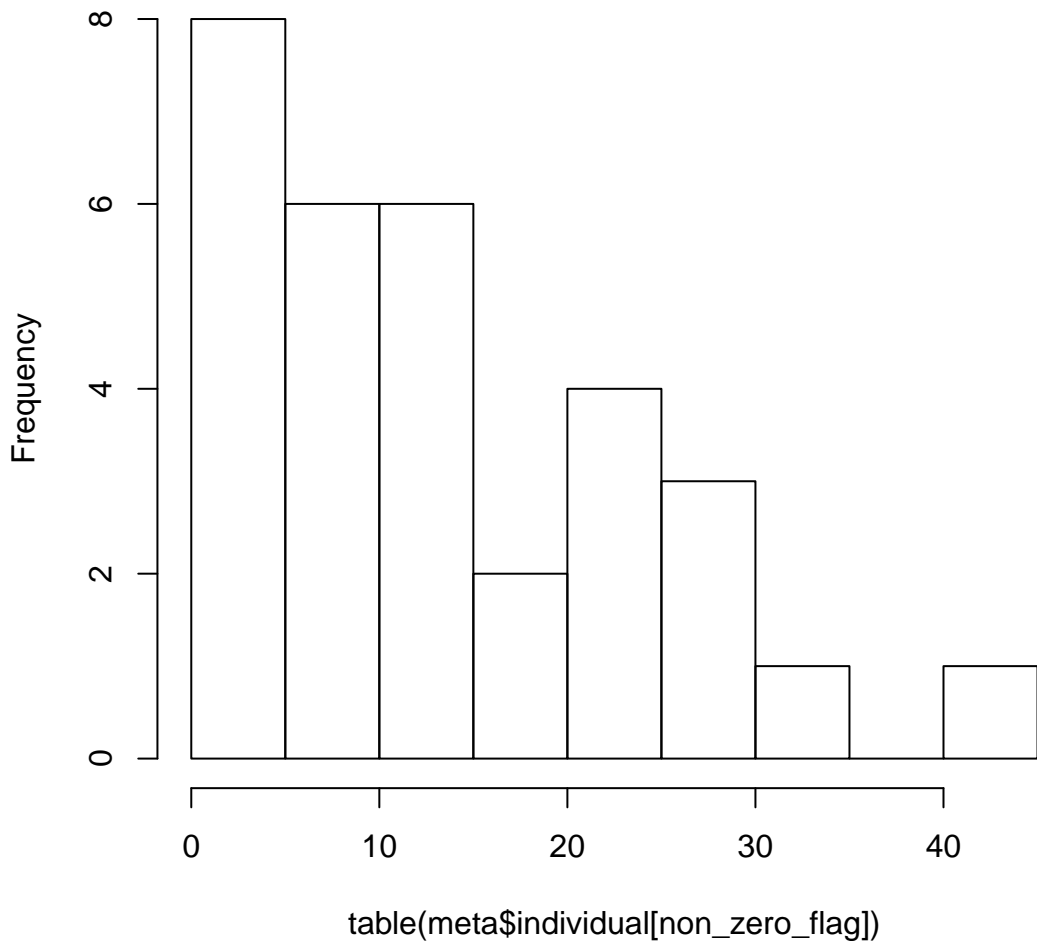
# KSless sig: individual expression cell count of gene#12



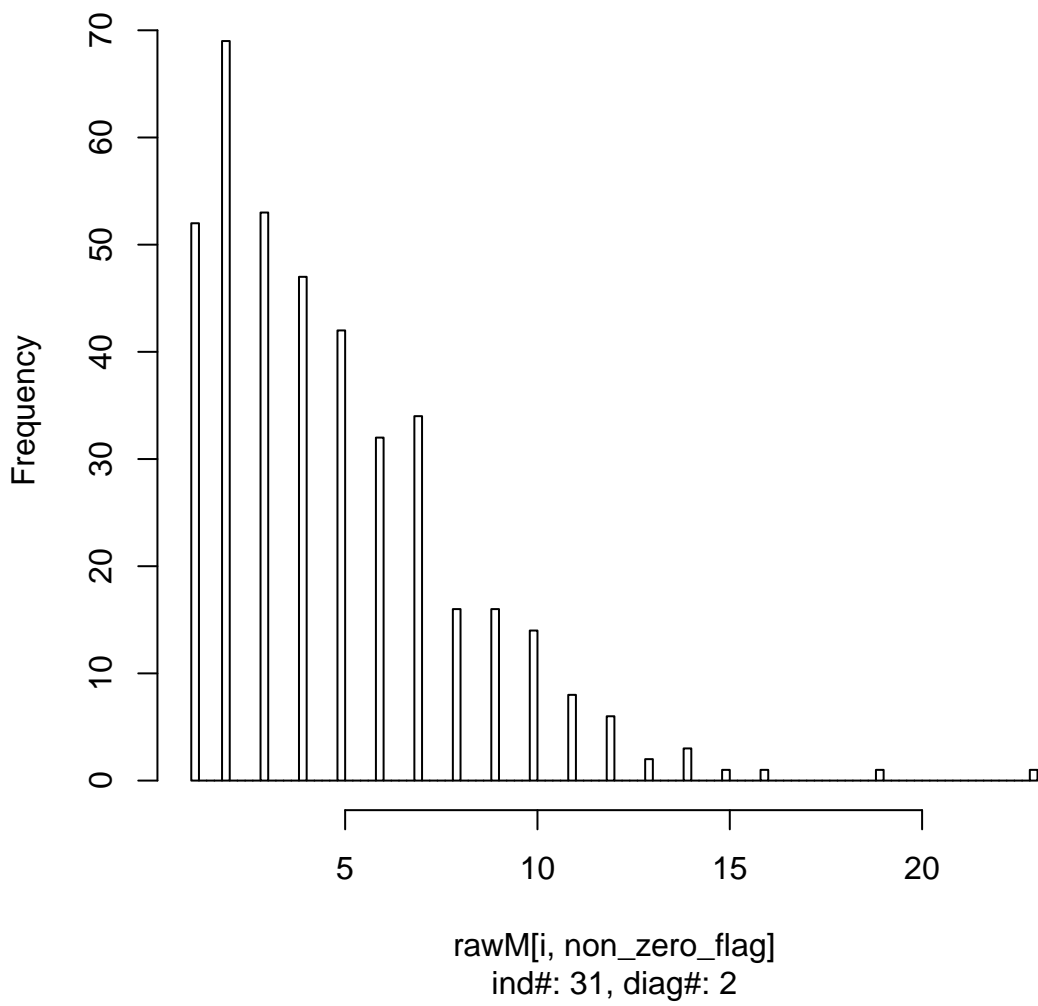
less sig: log expression of gene#13, pval ob=0.0459, non-zero nu



# KSless sig: individual expression cell count of gene#13

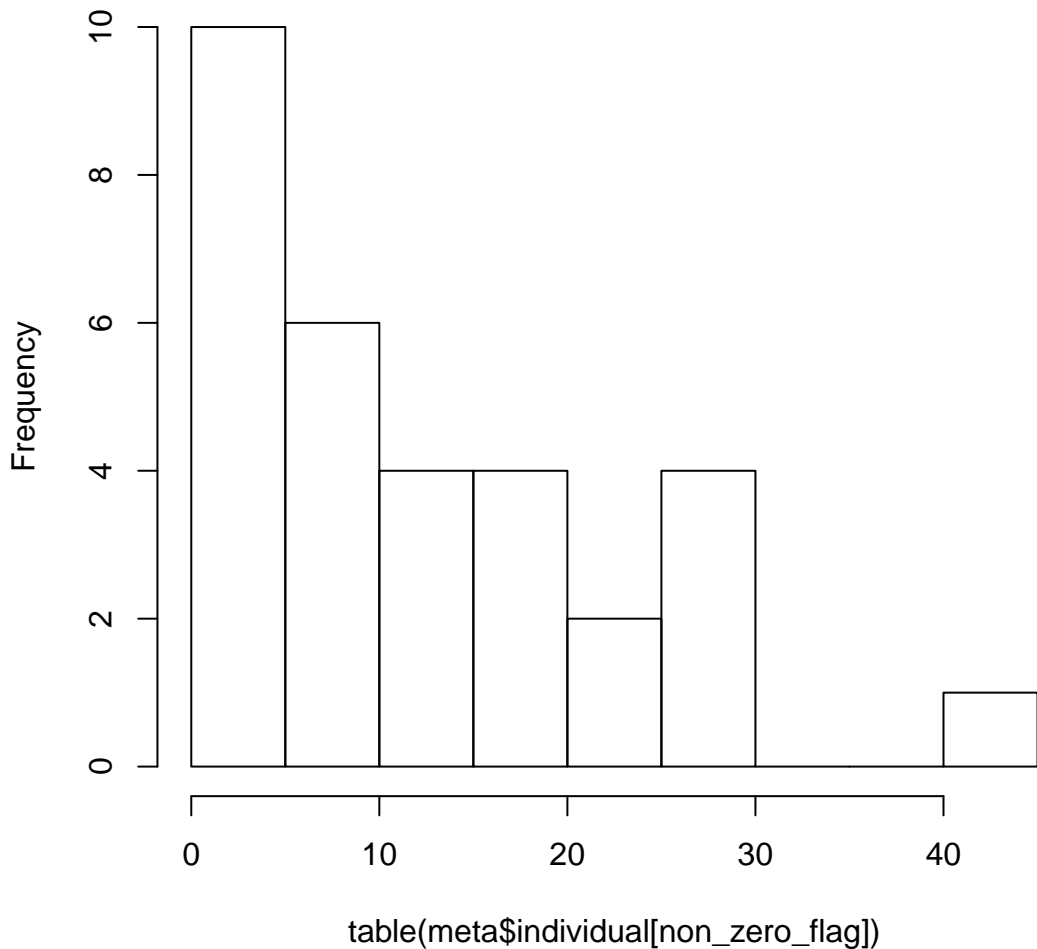


less sig: log expression of gene#18, pval ob=0.1275, non-zero nu

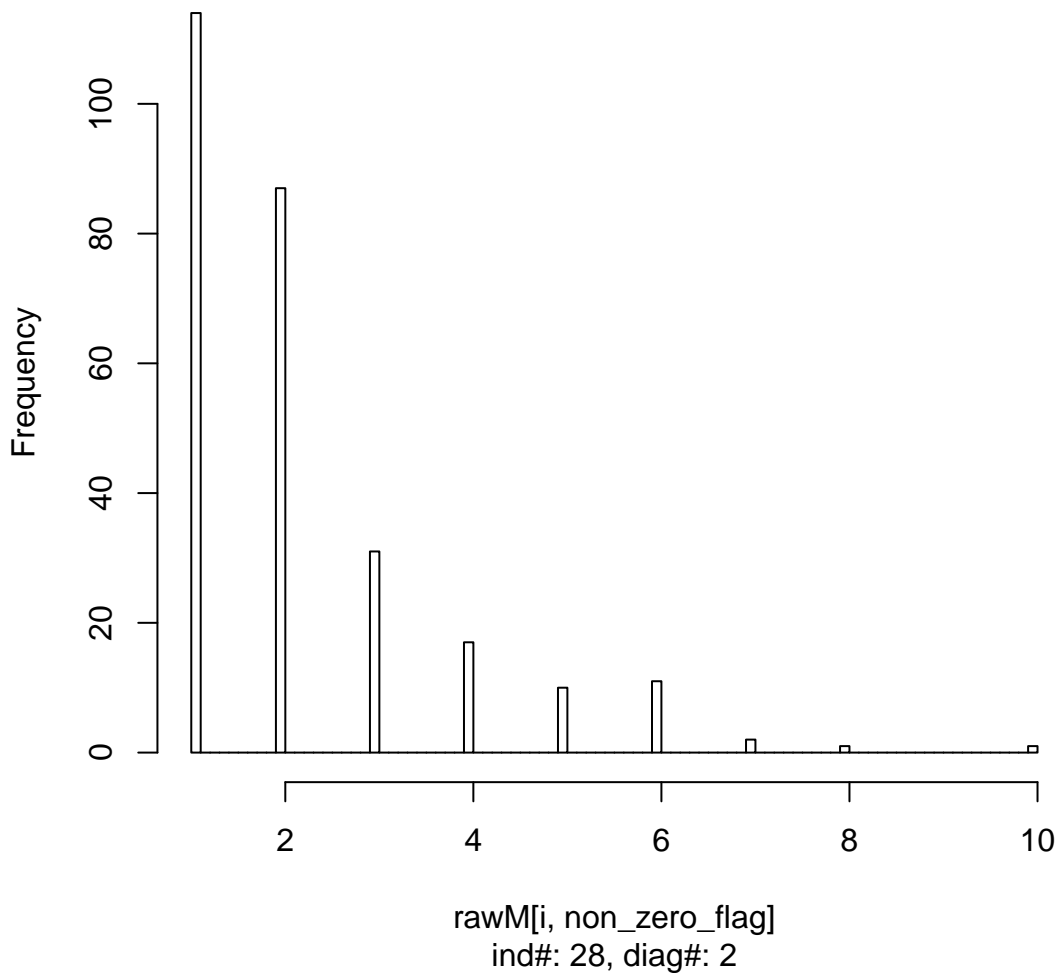




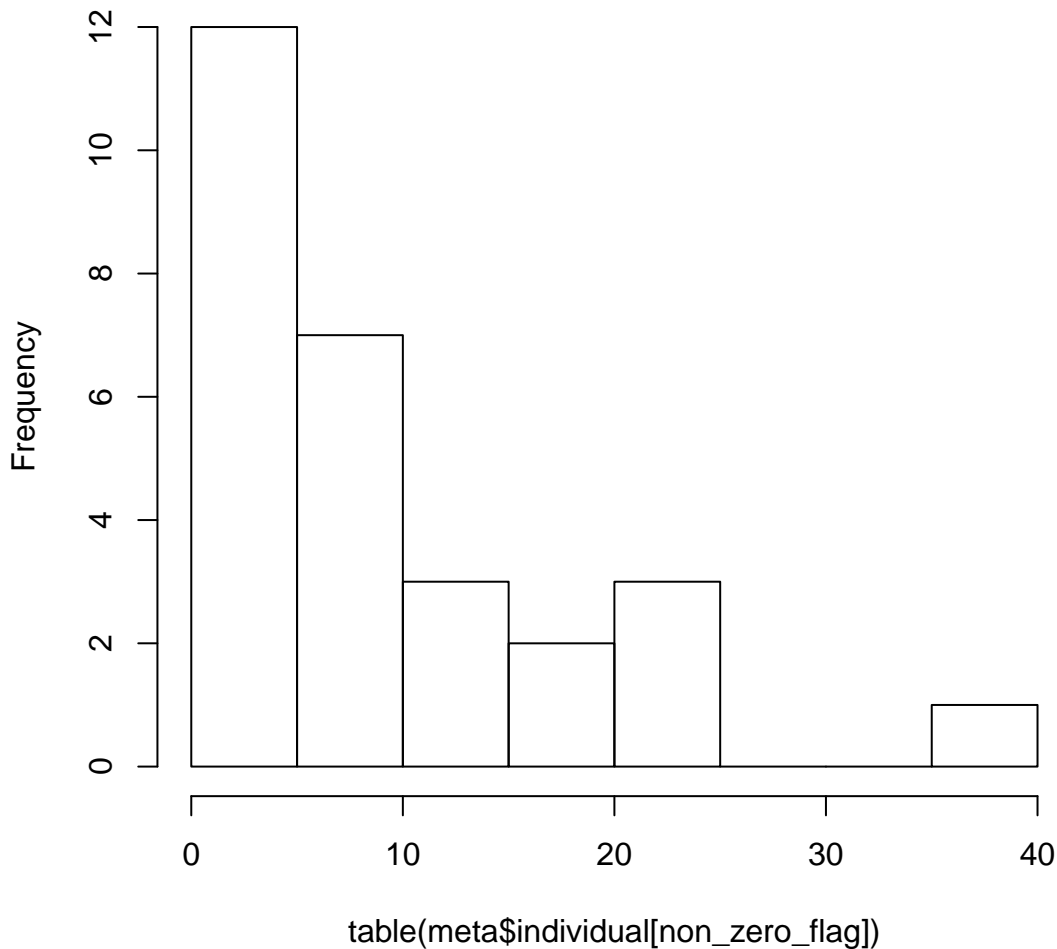
# KSless sig: individual expression cell count of gene#18



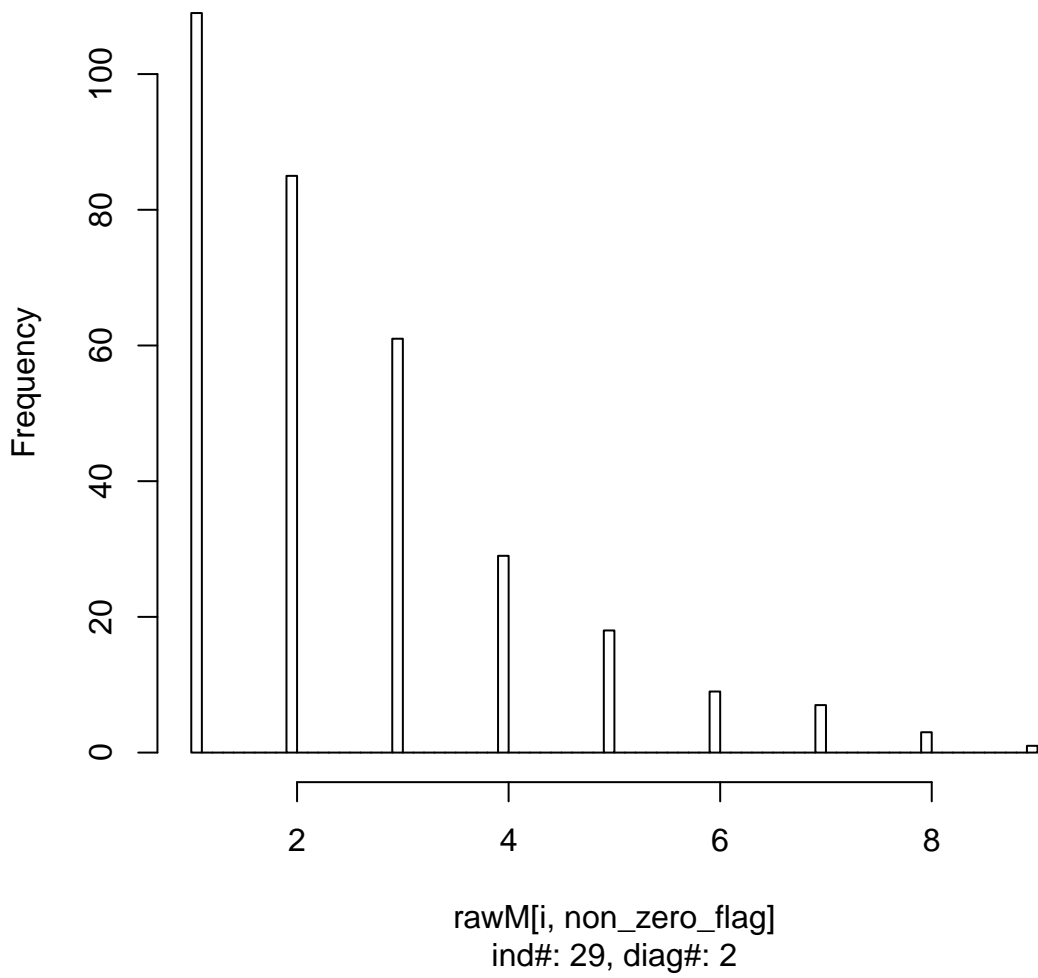
less sig: log expression of gene#20, pval ob=0.936, non-zero nu



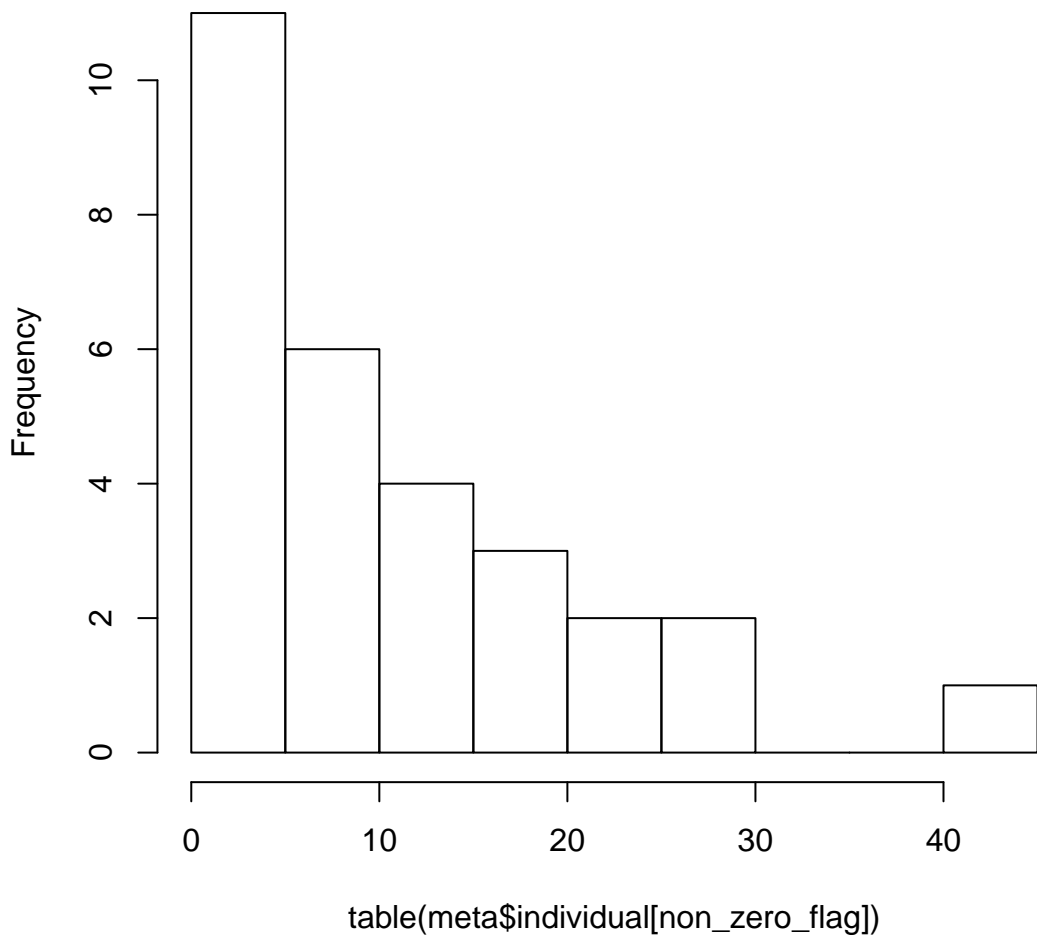
# KSless sig: individual expression cell count of gene#20



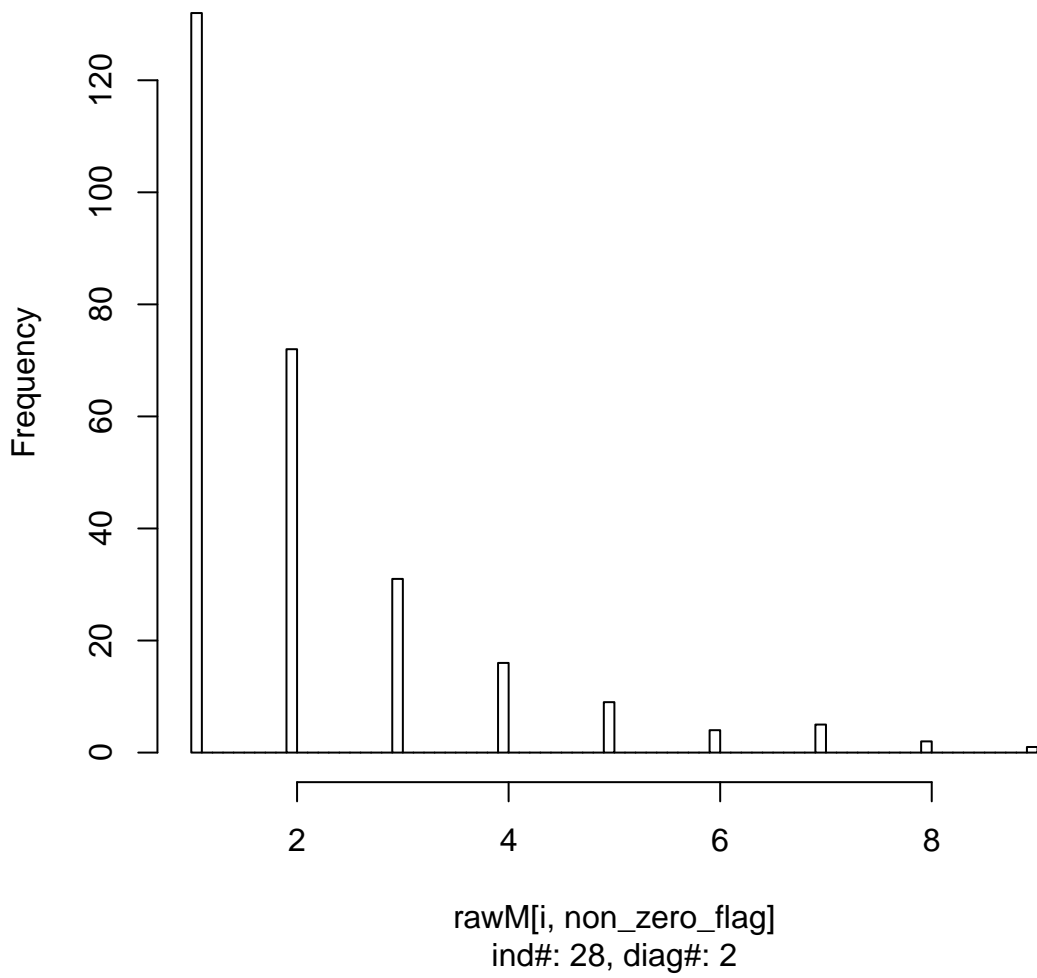
less sig: log expression of gene#22, pval ob=0.9139, non-zero nu



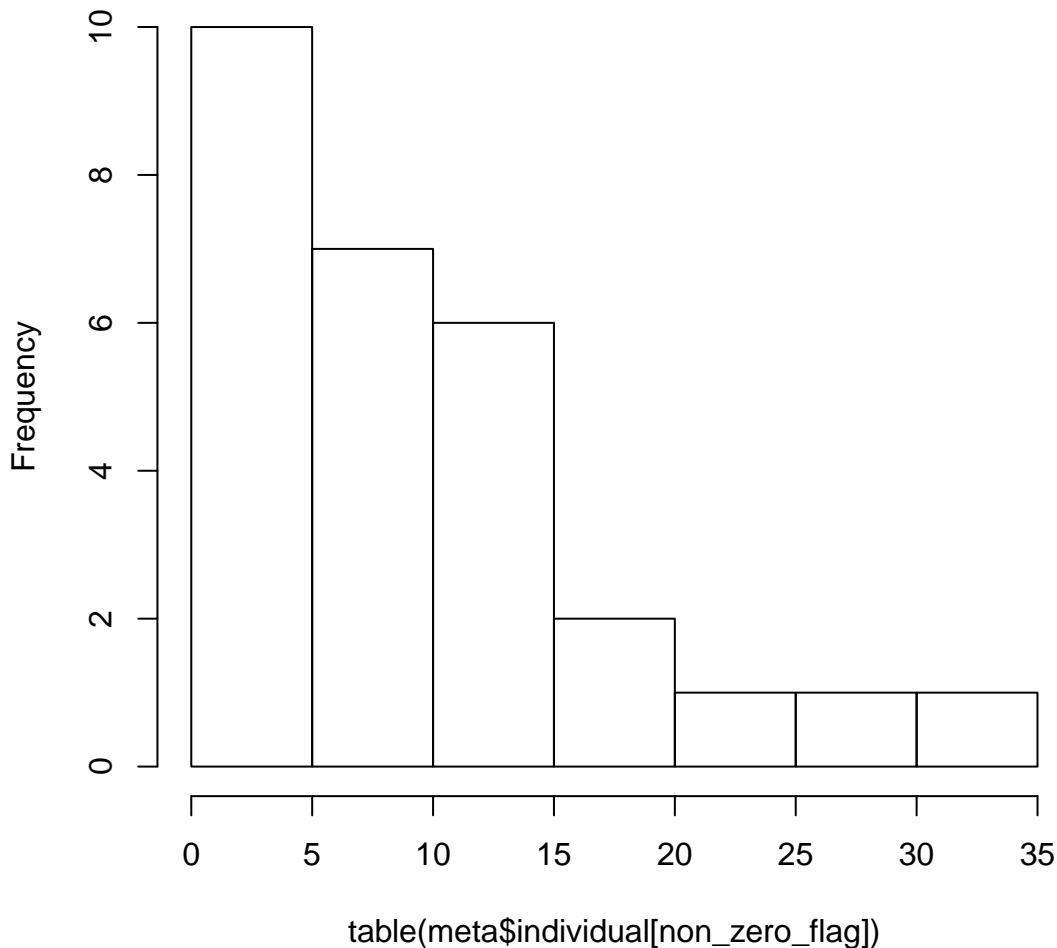
# KSless sig: individual expression cell count of gene#22



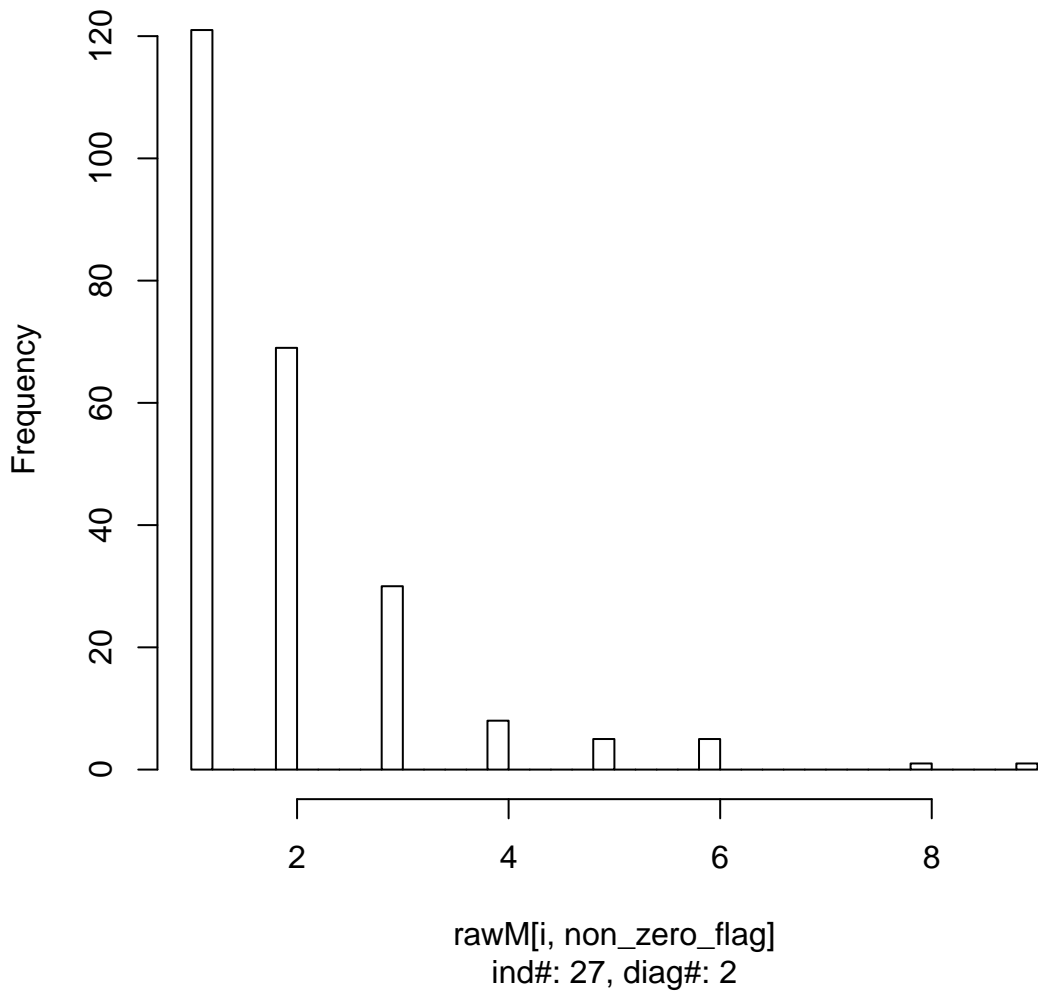
less sig: log expression of gene#26, pval ob=0.3221, non-zero nu



# KSless sig: individual expression cell count of gene#26

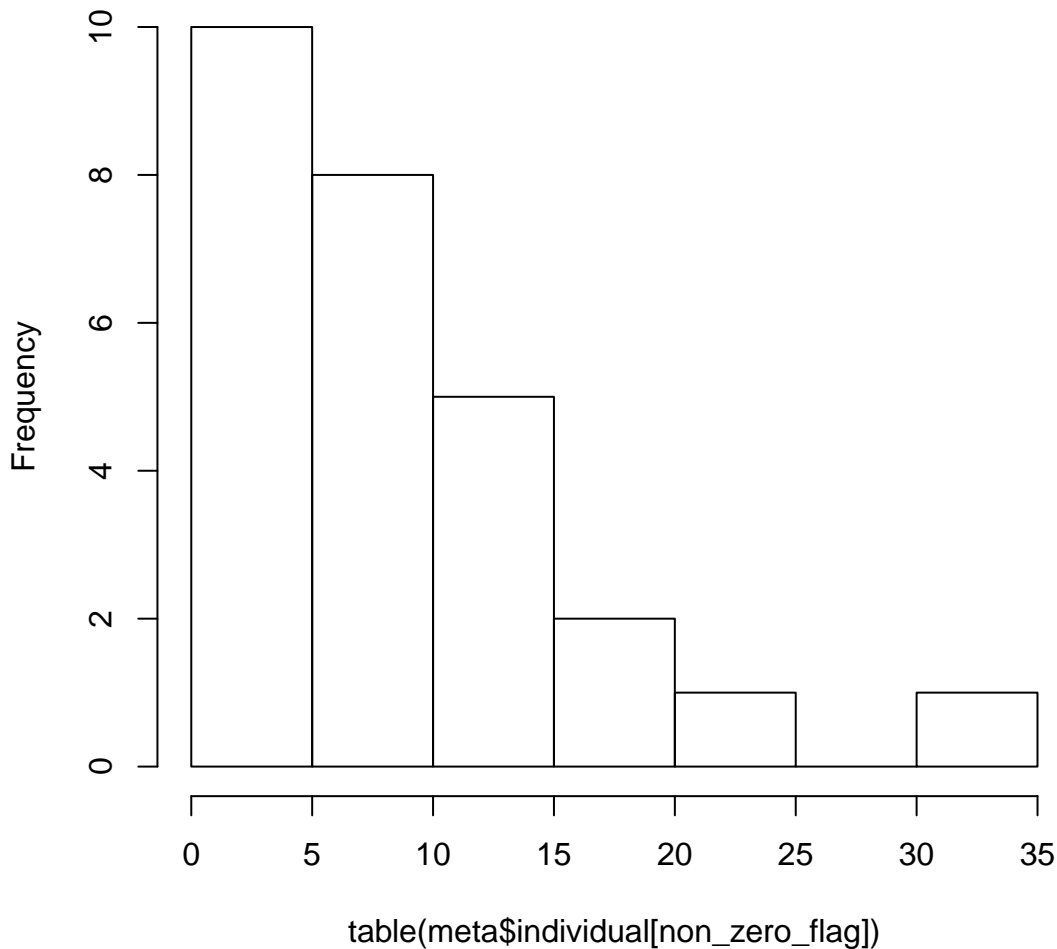


**S nonsig: log expression of gene#5, pval ob=0.0916, non-zero nu**

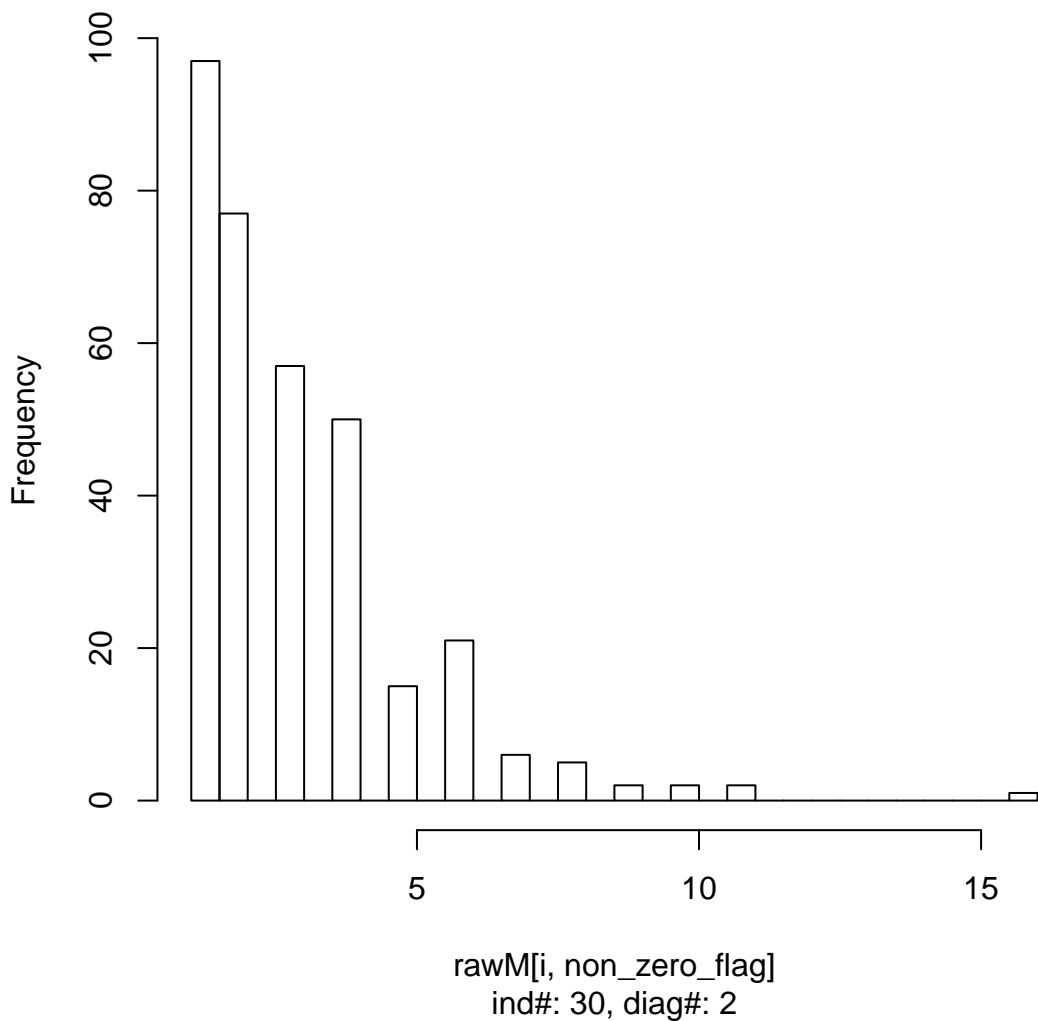




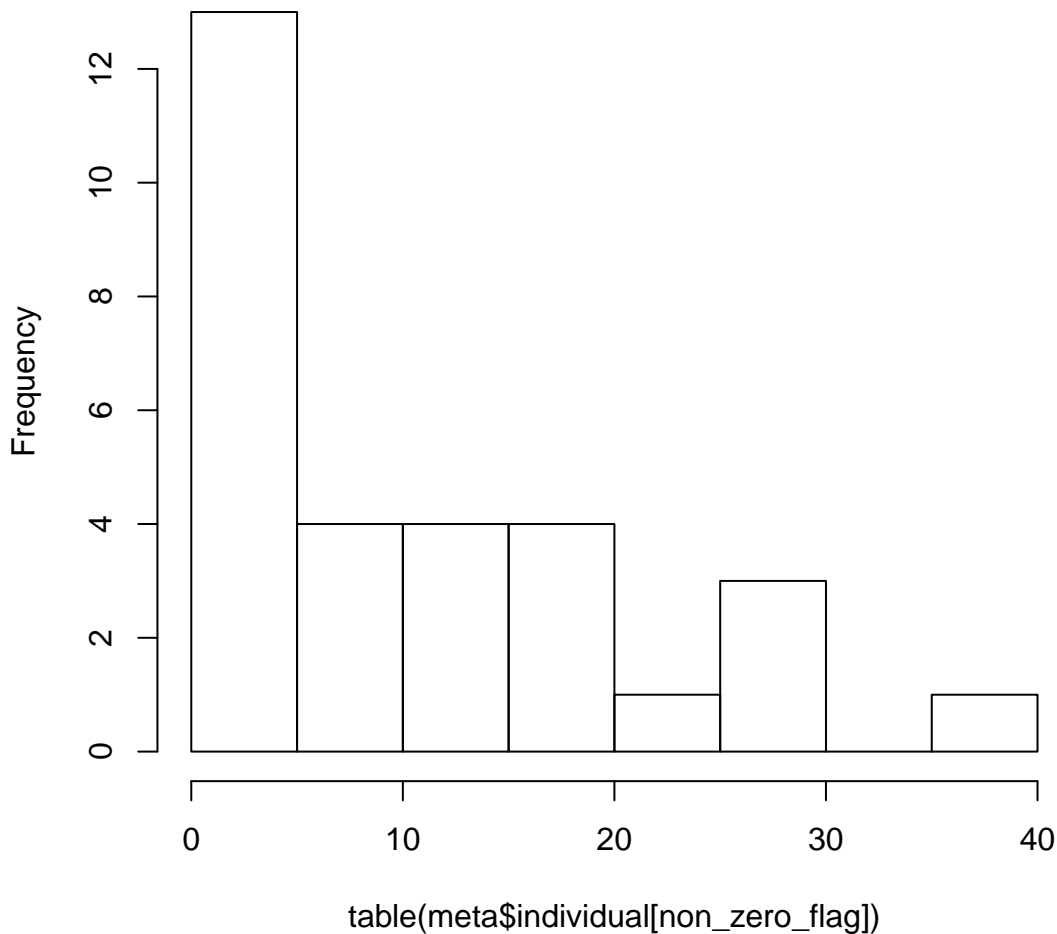
# KSless nonsig: individual expression cell count of gene#5



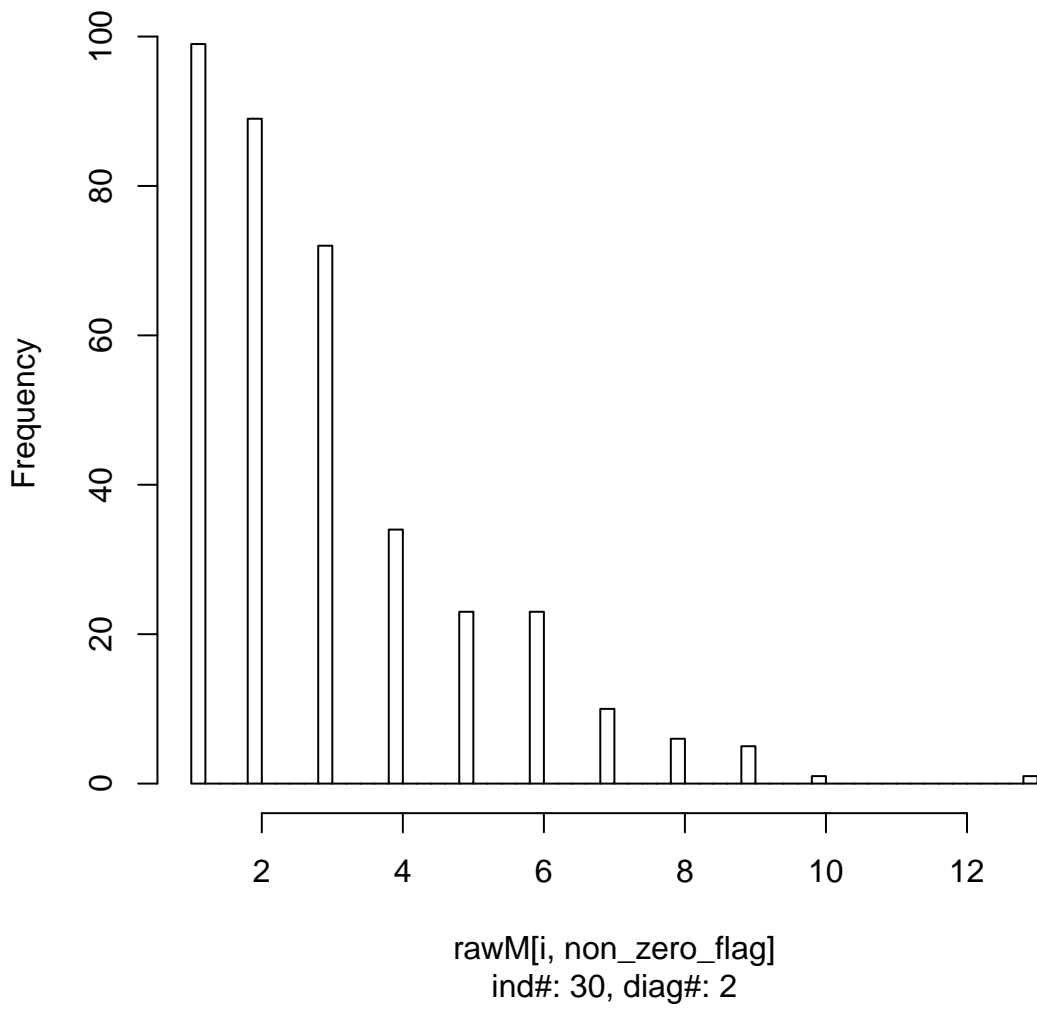
**S nonsig: log expression of gene#6, pval ob=0.3298, non-zero num**



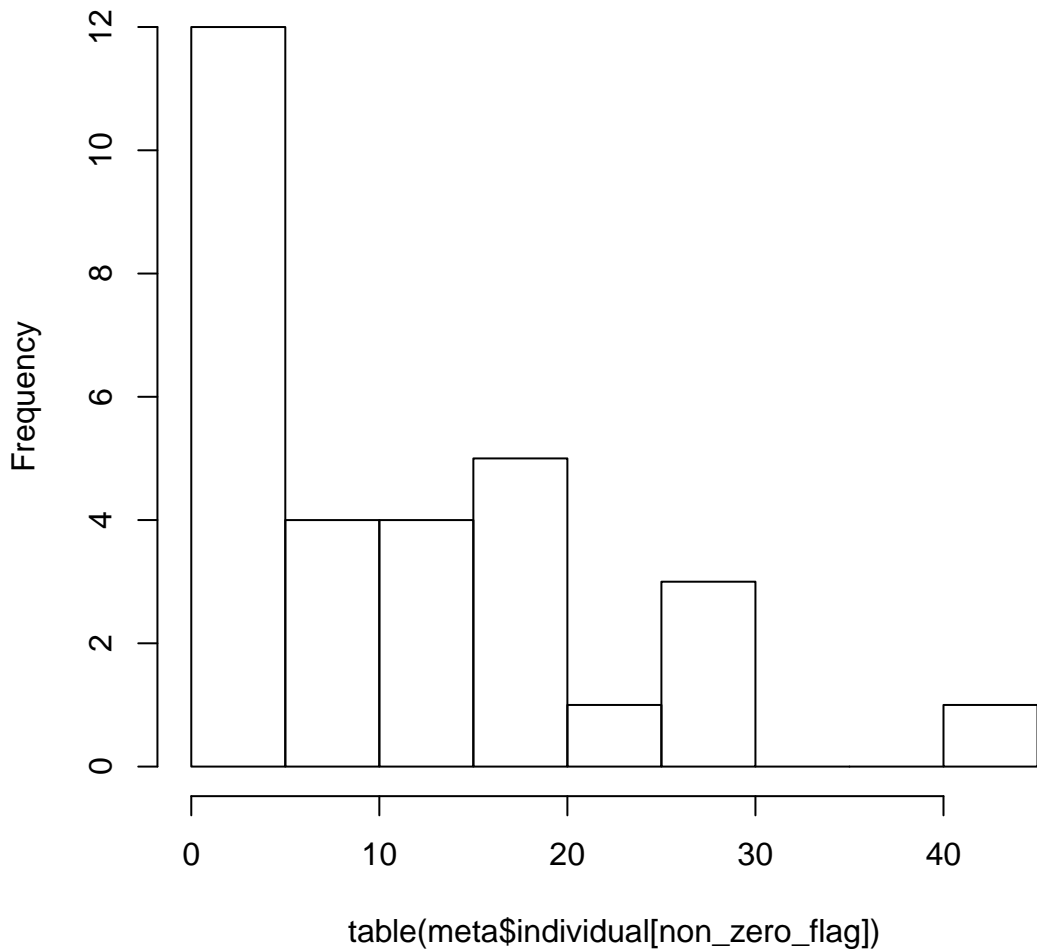
# KSless nonsig: individual expression cell count of gene#6



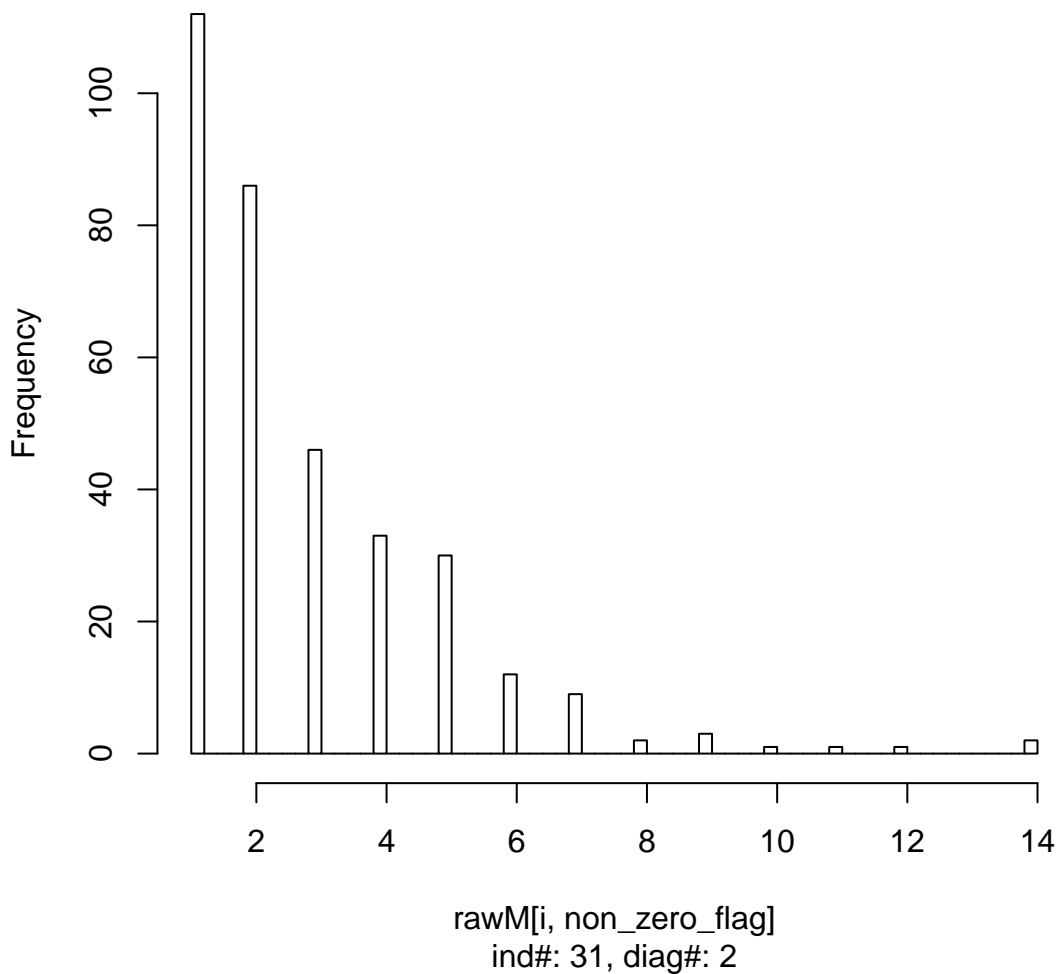
**S nonsig: log expression of gene#7, pval ob=0.4847, non-zero num**



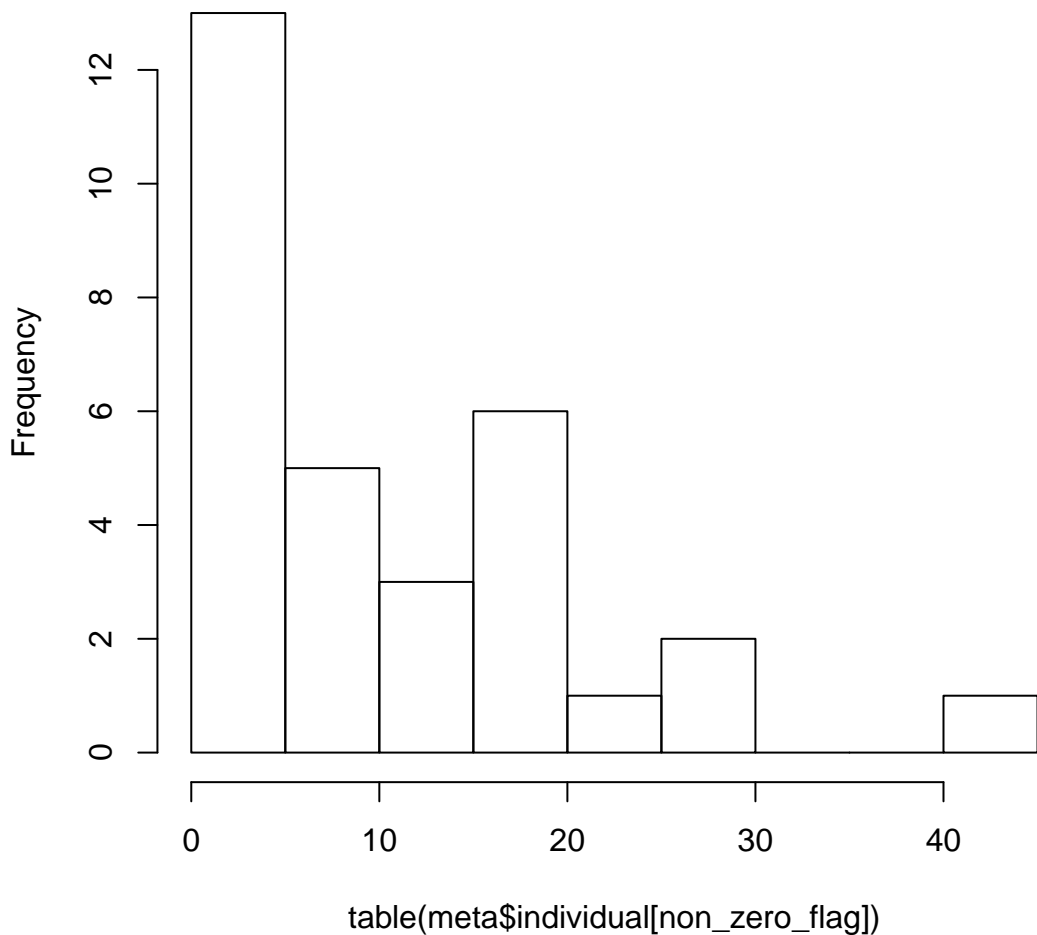
# KSless nonsig: individual expression cell count of gene#7



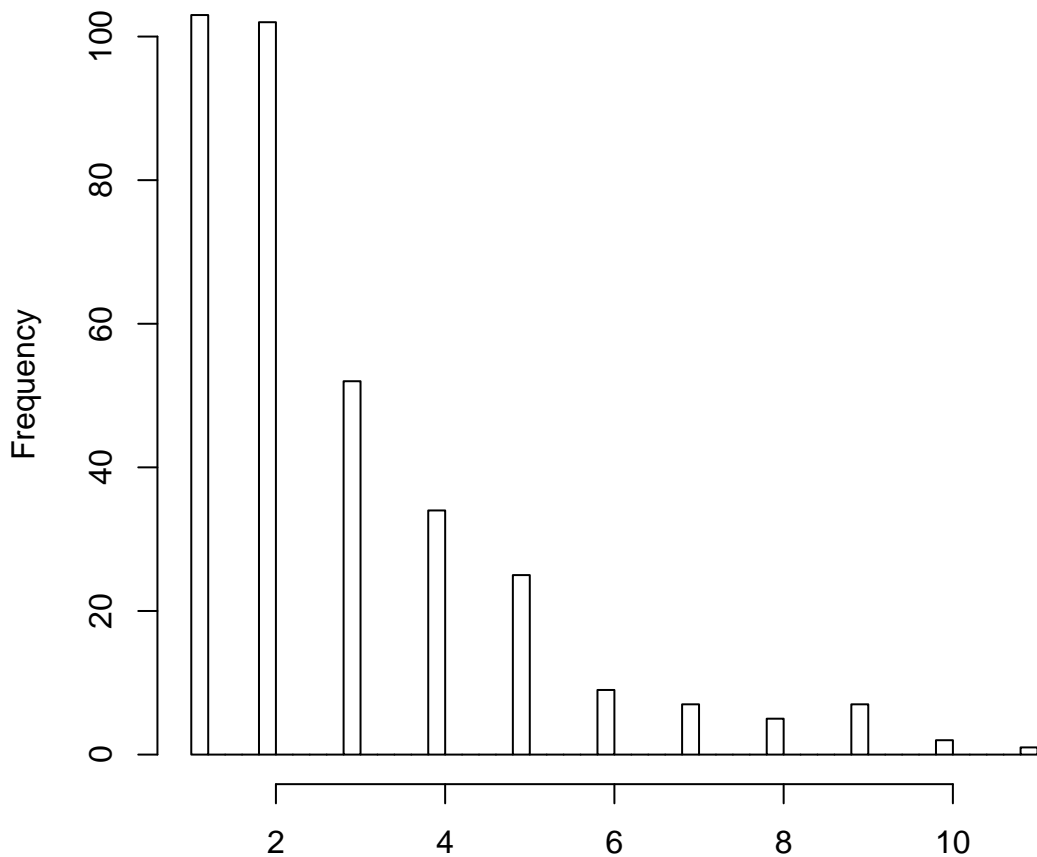
**nsig: log expression of gene#10, pval ob=0.8236, non-zero nu**



# KSless nonsig: individual expression cell count of gene#10



**nsig: log expression of gene#11, pval ob=0.7143, non-zero nu**

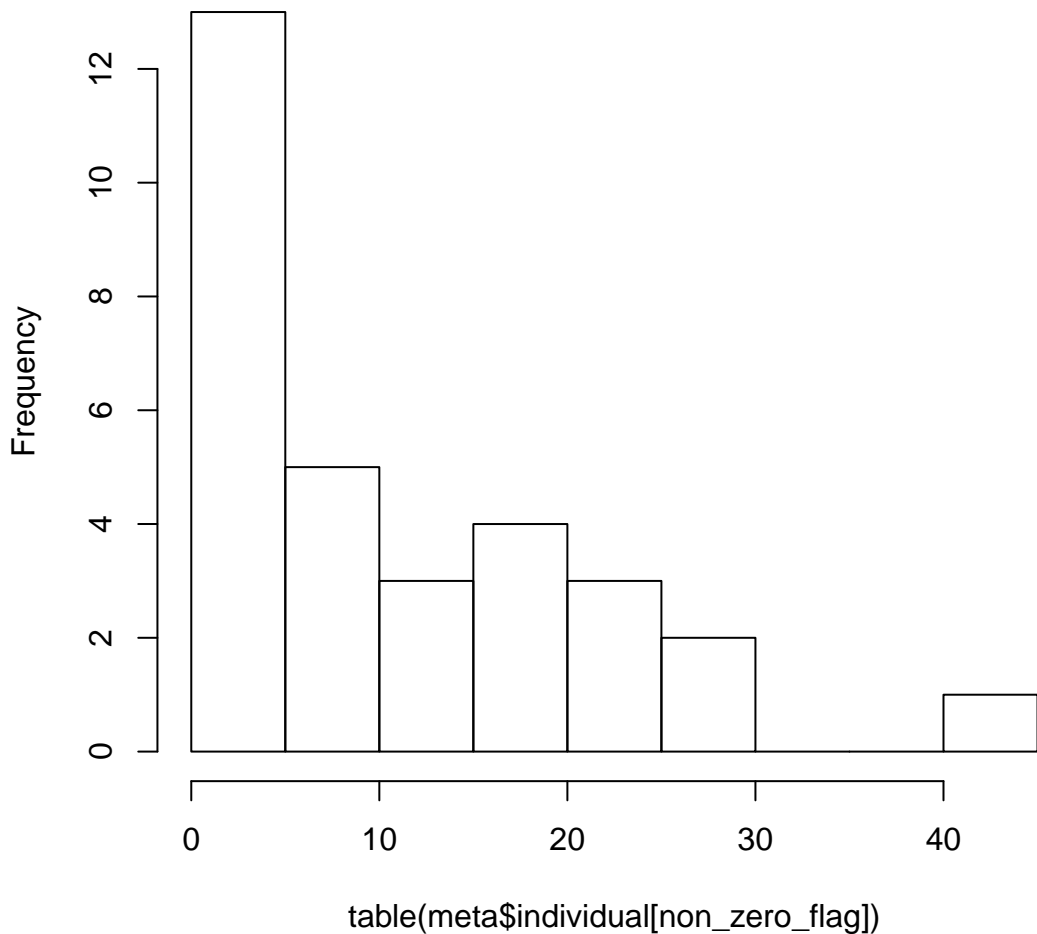


`rawM[i, non_zero_flag]`

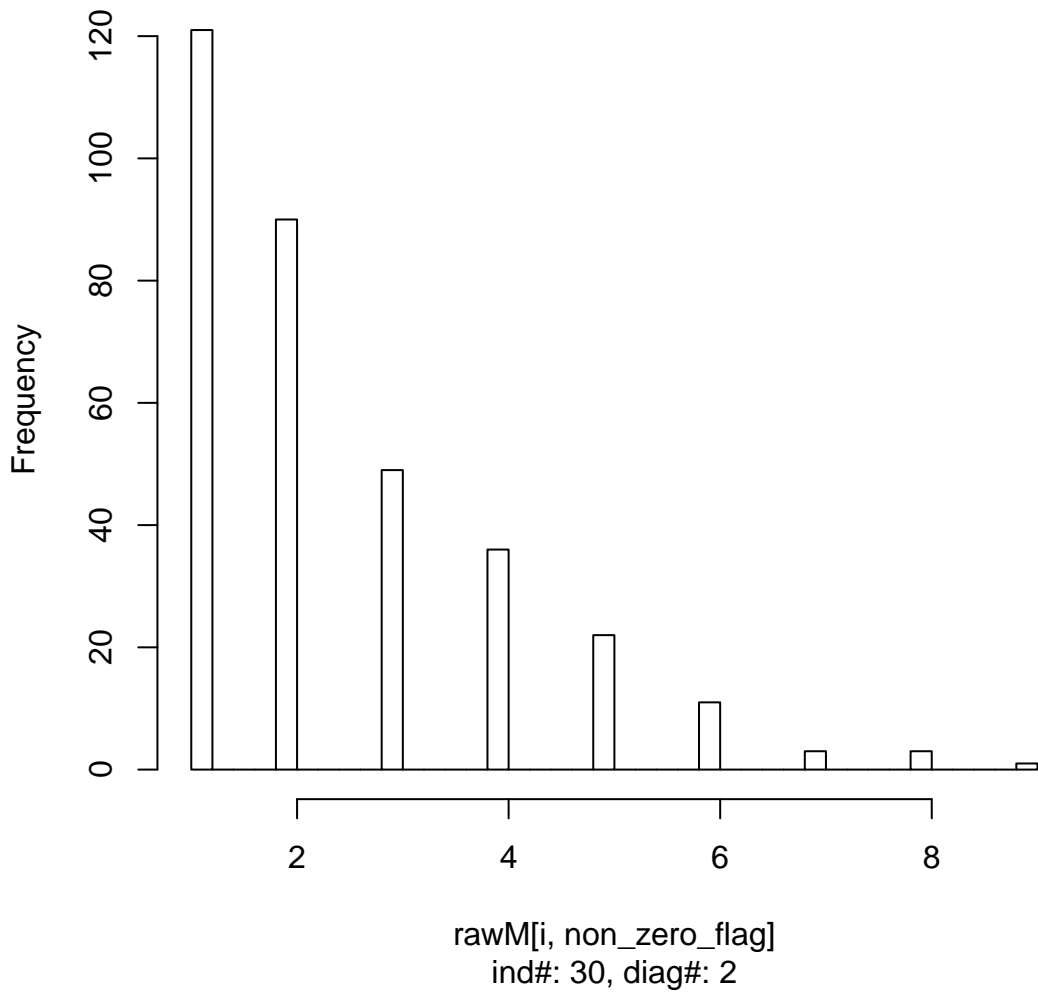
ind#: 31, diag#: 2



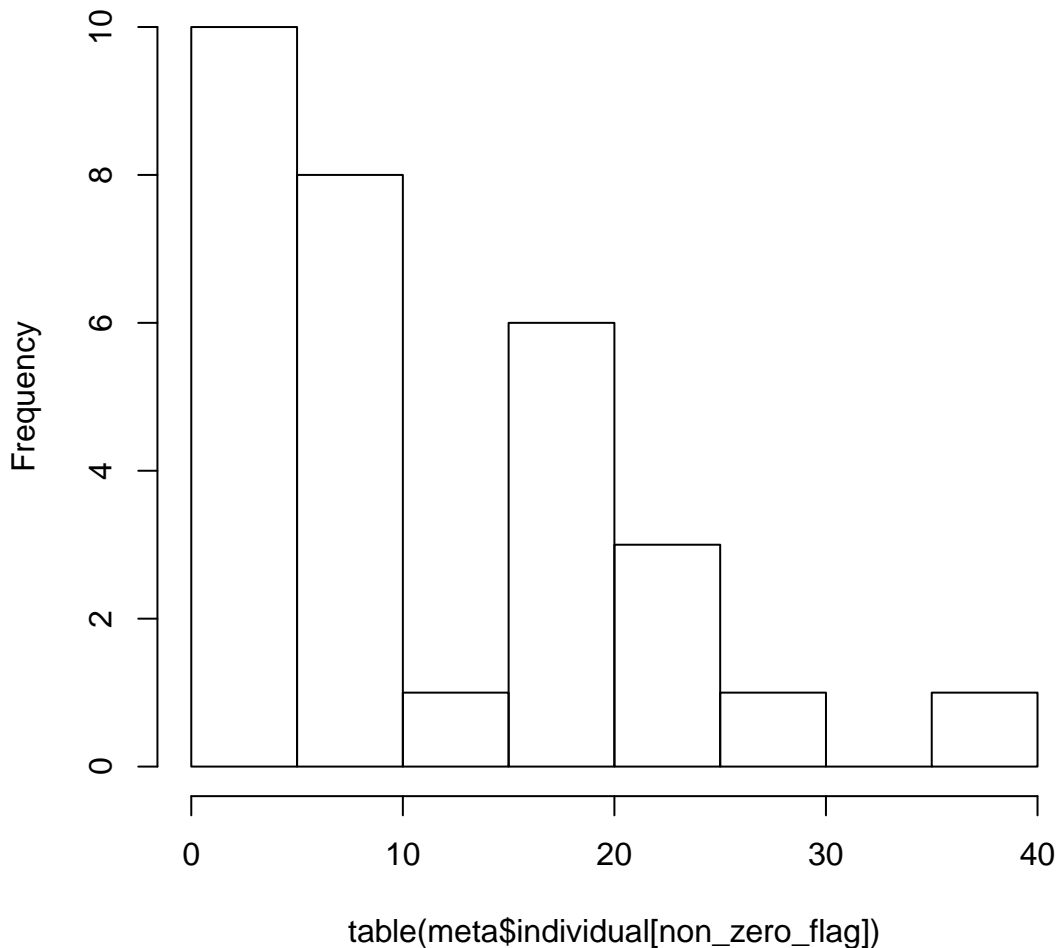
# KSless nonsig: individual expression cell count of gene#11



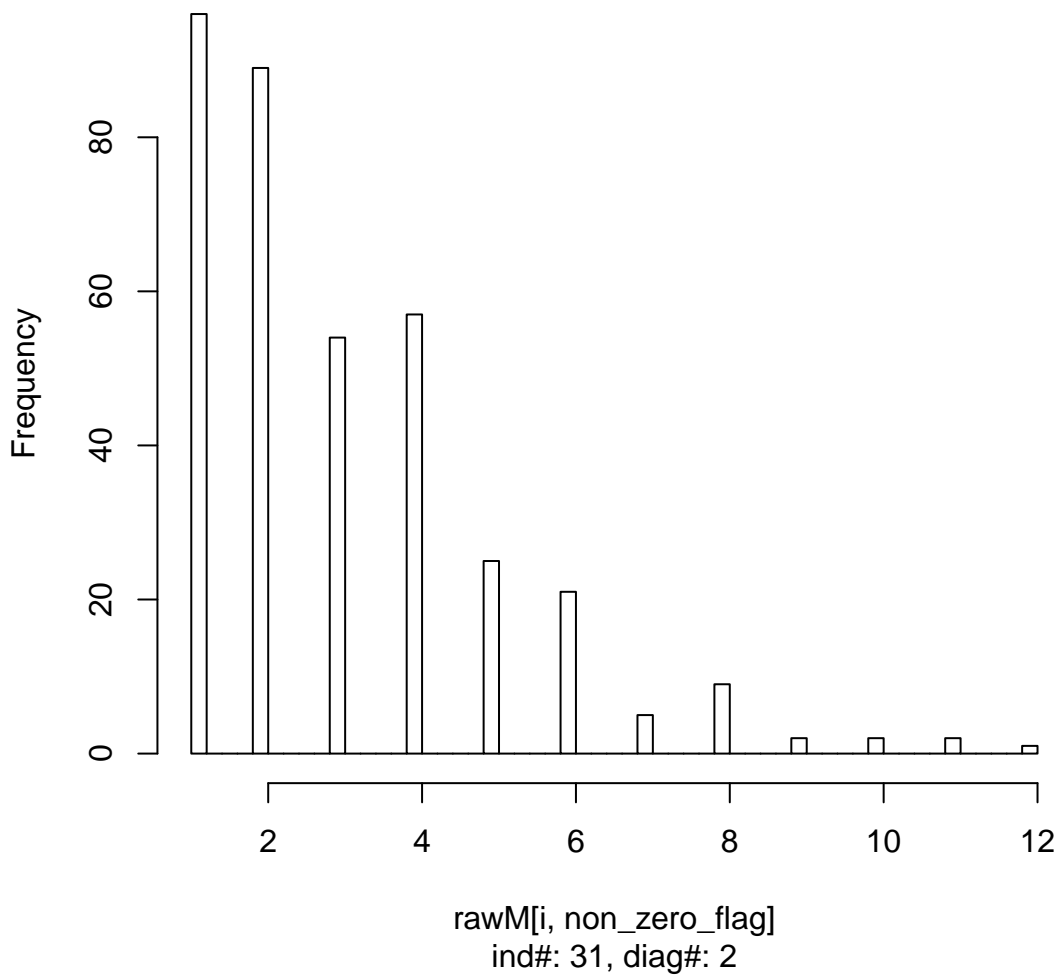
nsig: log expression of gene#15, pval ob=0.2493, non-zero nu



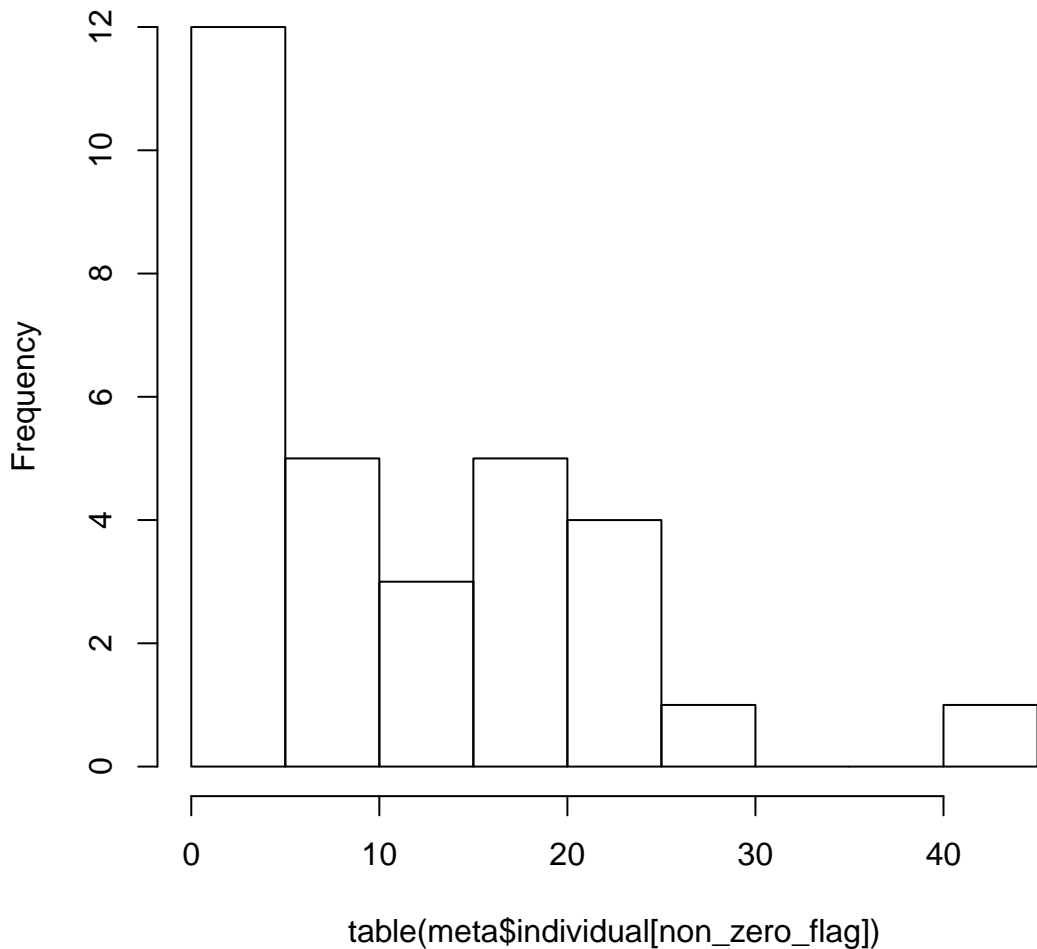
# KSless nonsig: individual expression cell count of gene#15



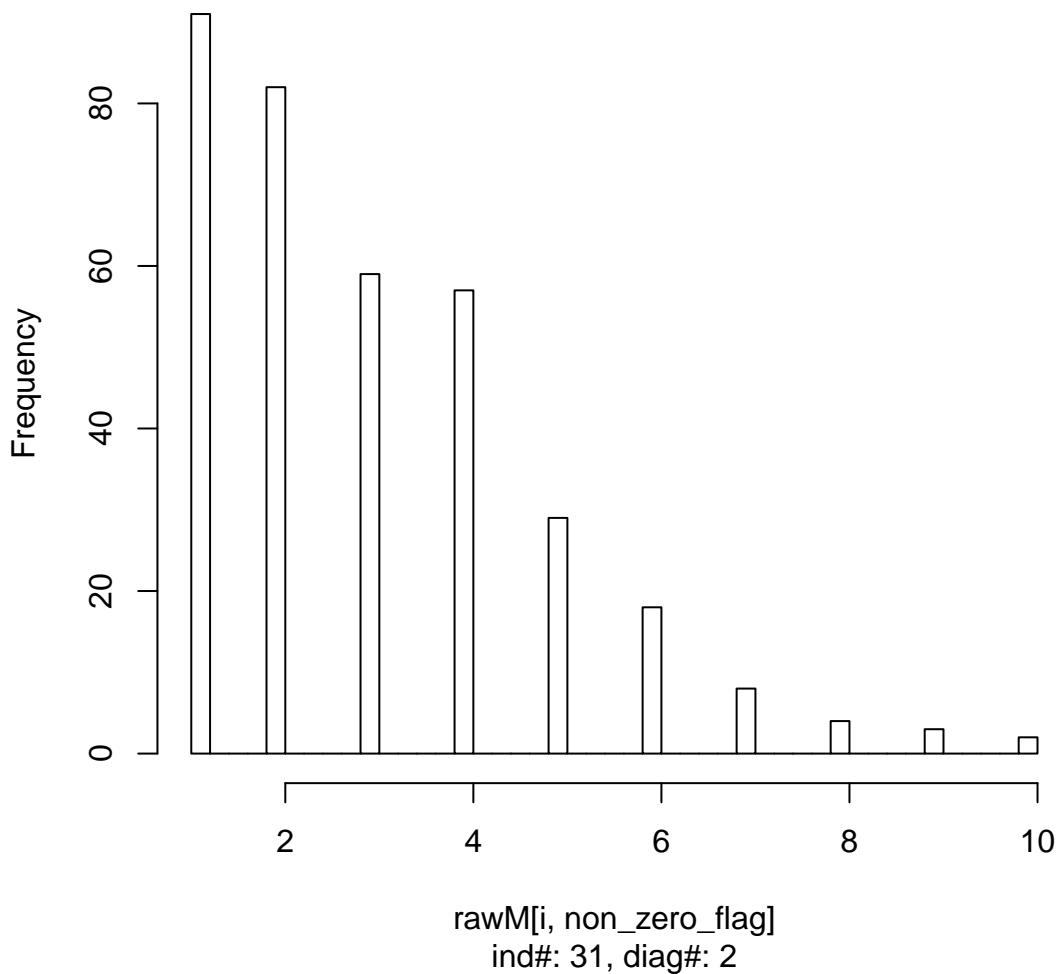
**nsig: log expression of gene#16, pval ob=0.3773, non-zero nu**



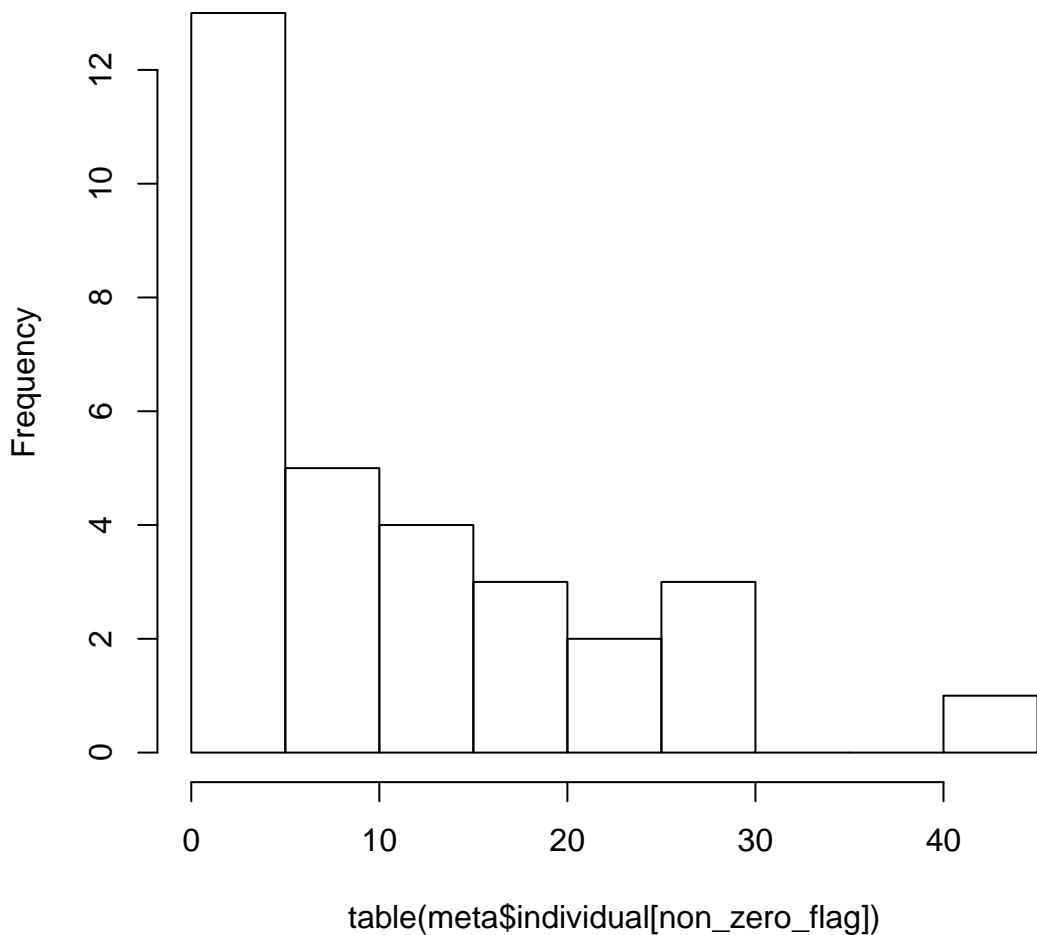
# KSless nonsig: individual expression cell count of gene#16



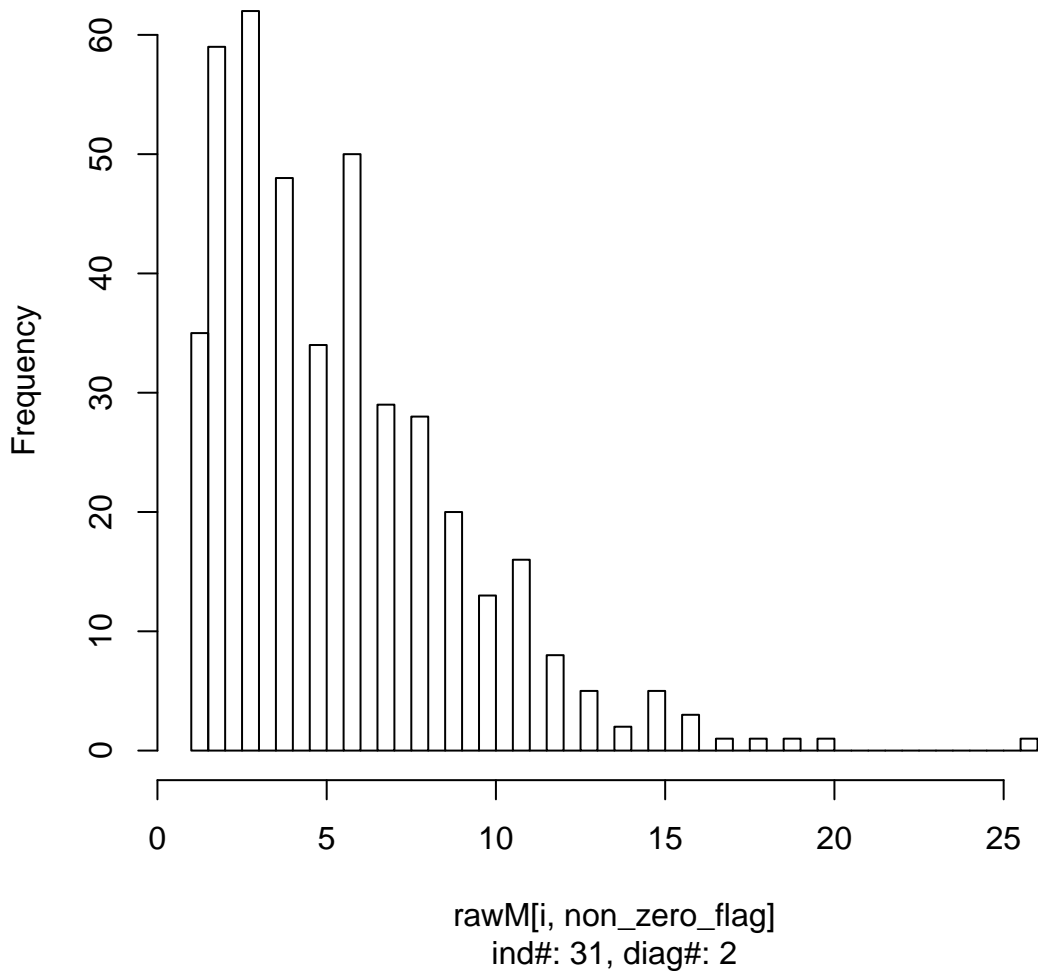
**S nonsig: log expression of gene#17, pval ob=0.631, non-zero nu**



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

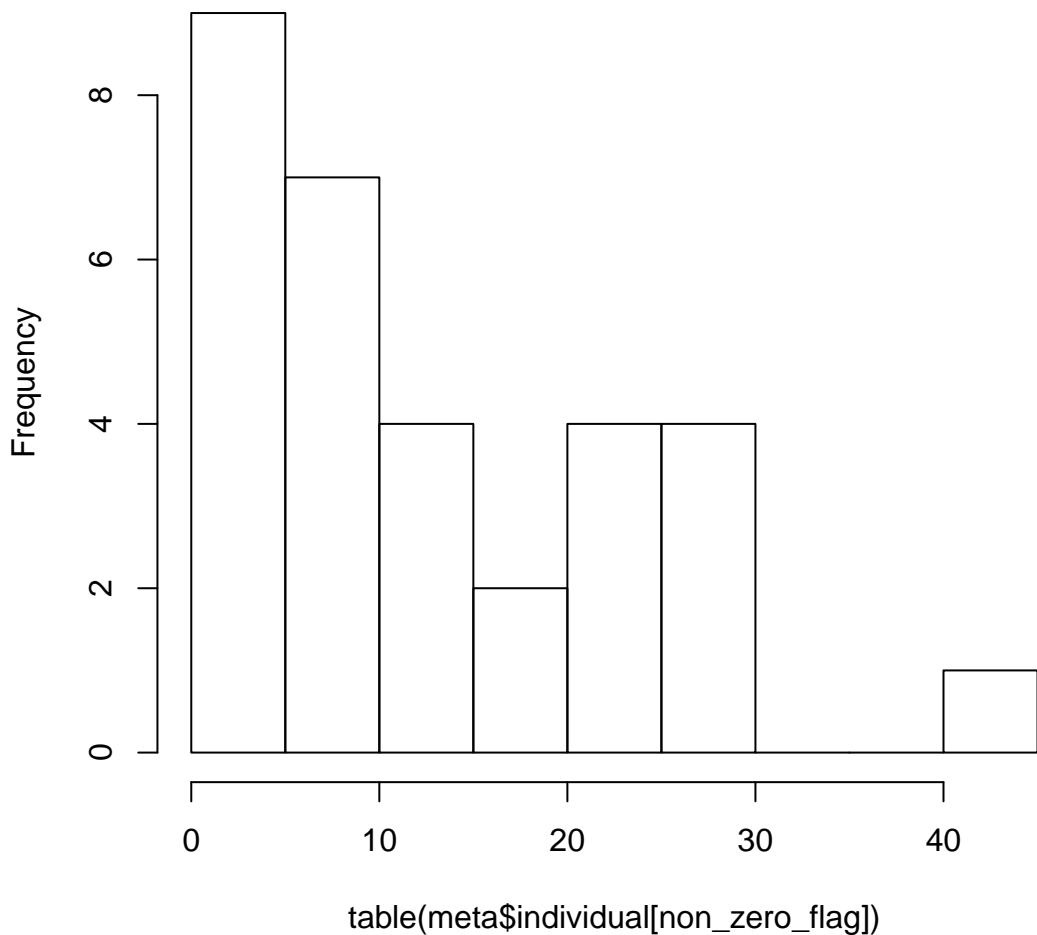


**nsig: log expression of gene#19, pval ob=0.8004, non-zero nu**

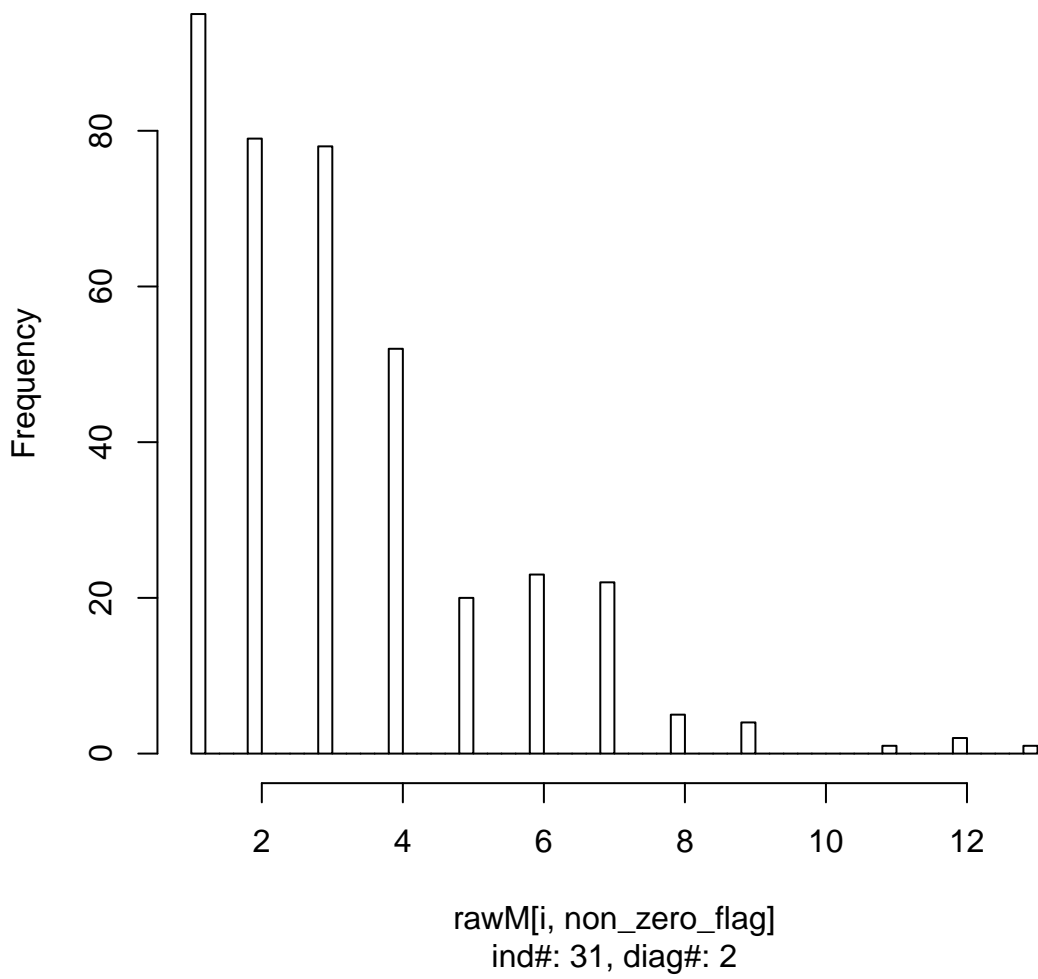




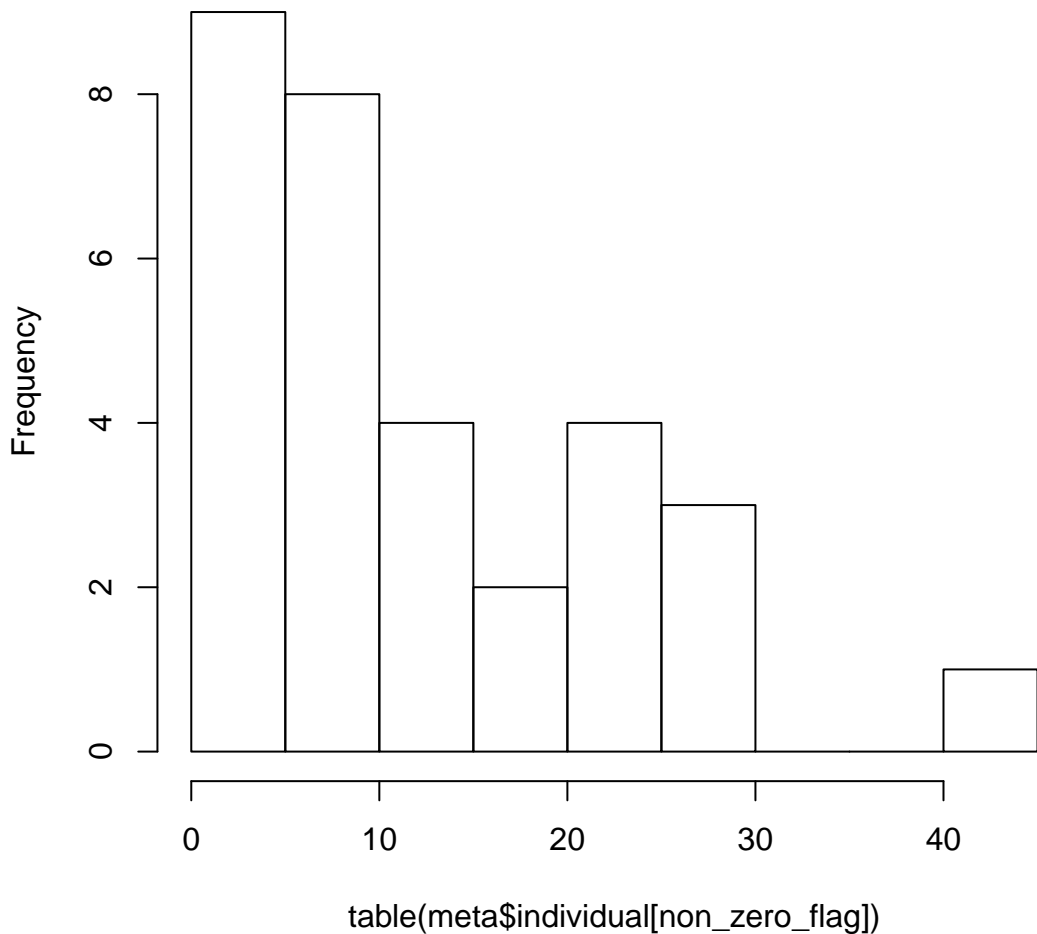
# KSless nonsig: individual expression cell count of gene#19



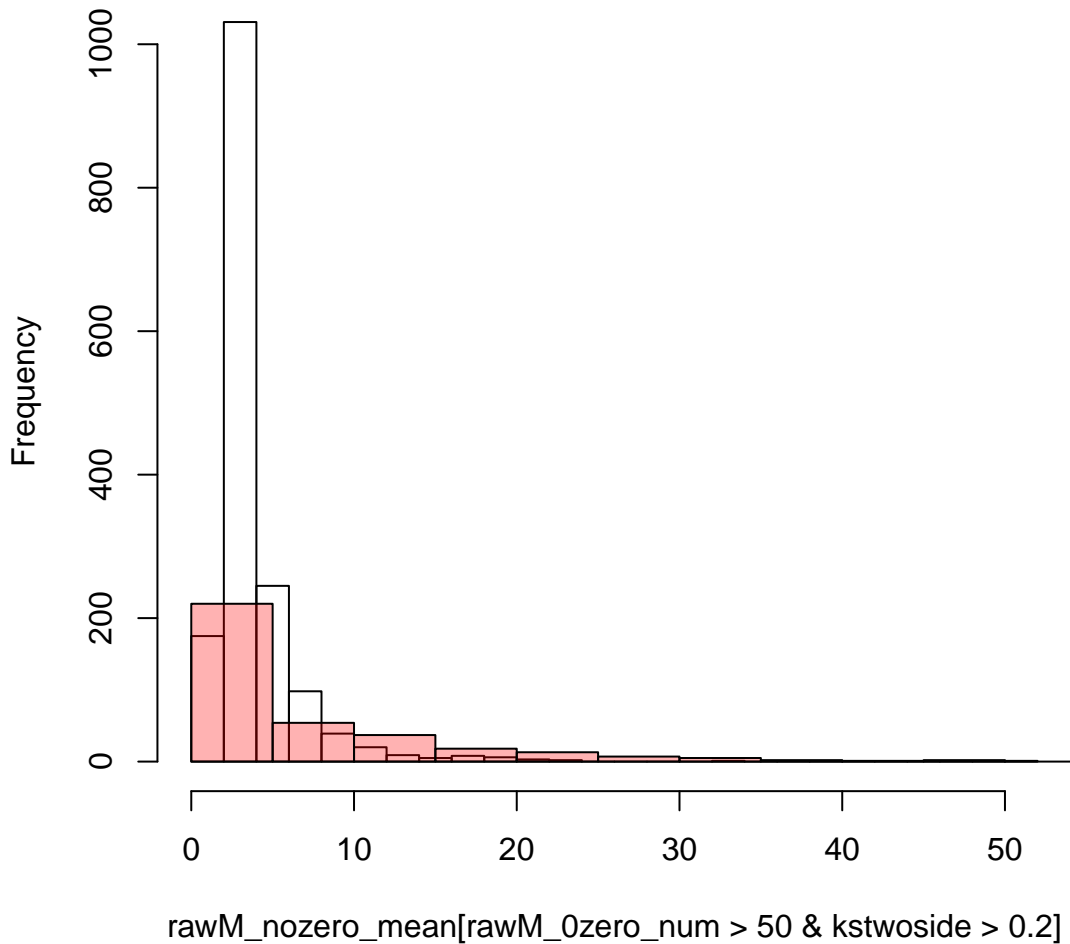
nsig: log expression of gene#21, pval ob=0.2019, non-zero nu



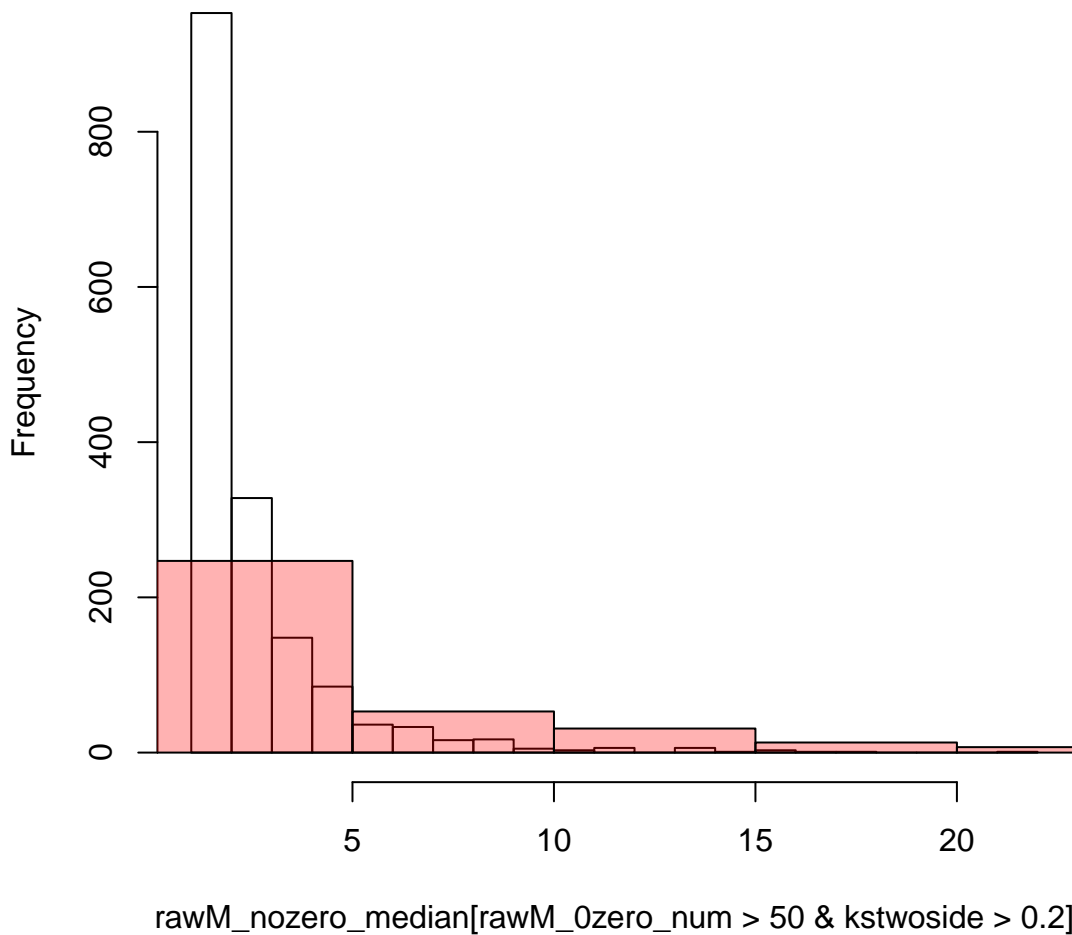
# KSless nonsig: individual expression cell count of gene#21



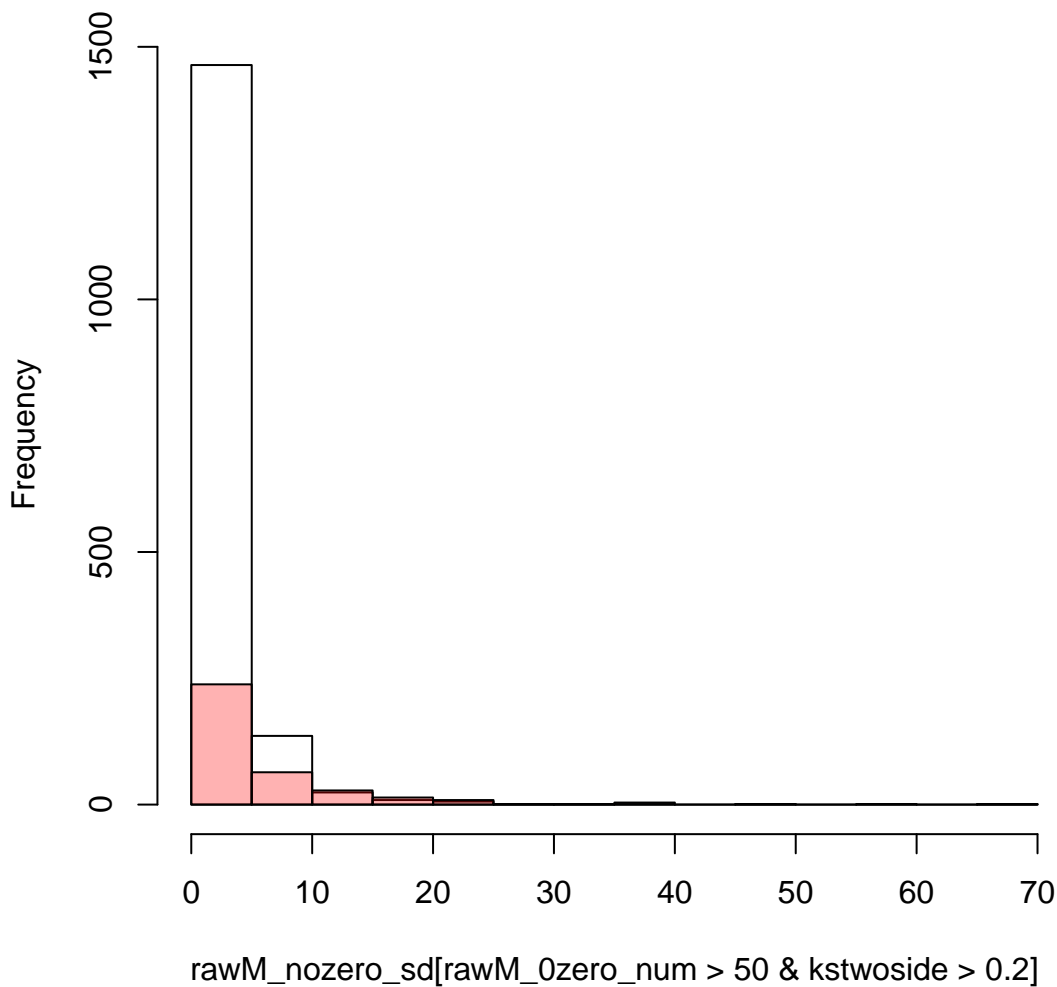
Histogram of rawM\_nozero\_mean[rawM\_0zero\_num > 50 & kstwsid



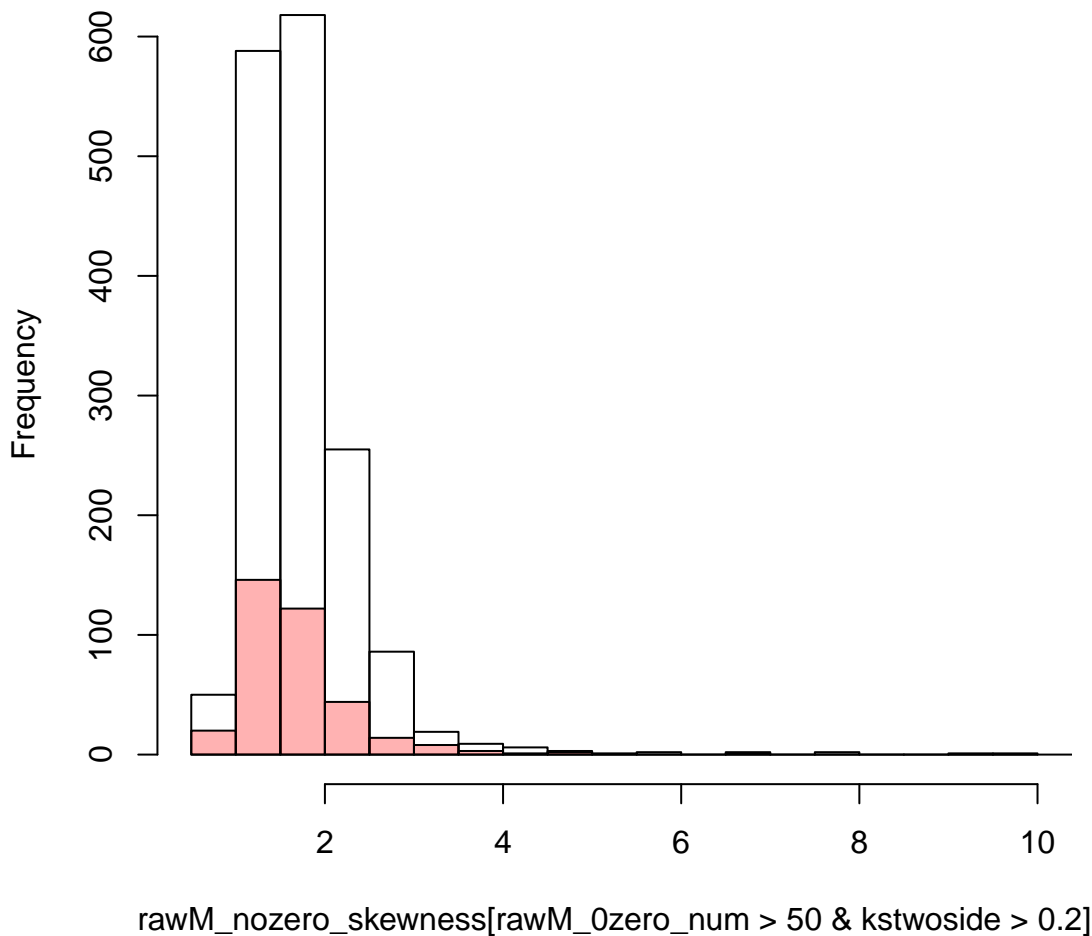
ogram of rawM\_nozero\_median[rawM\_0zero\_num > 50 & kstwosi



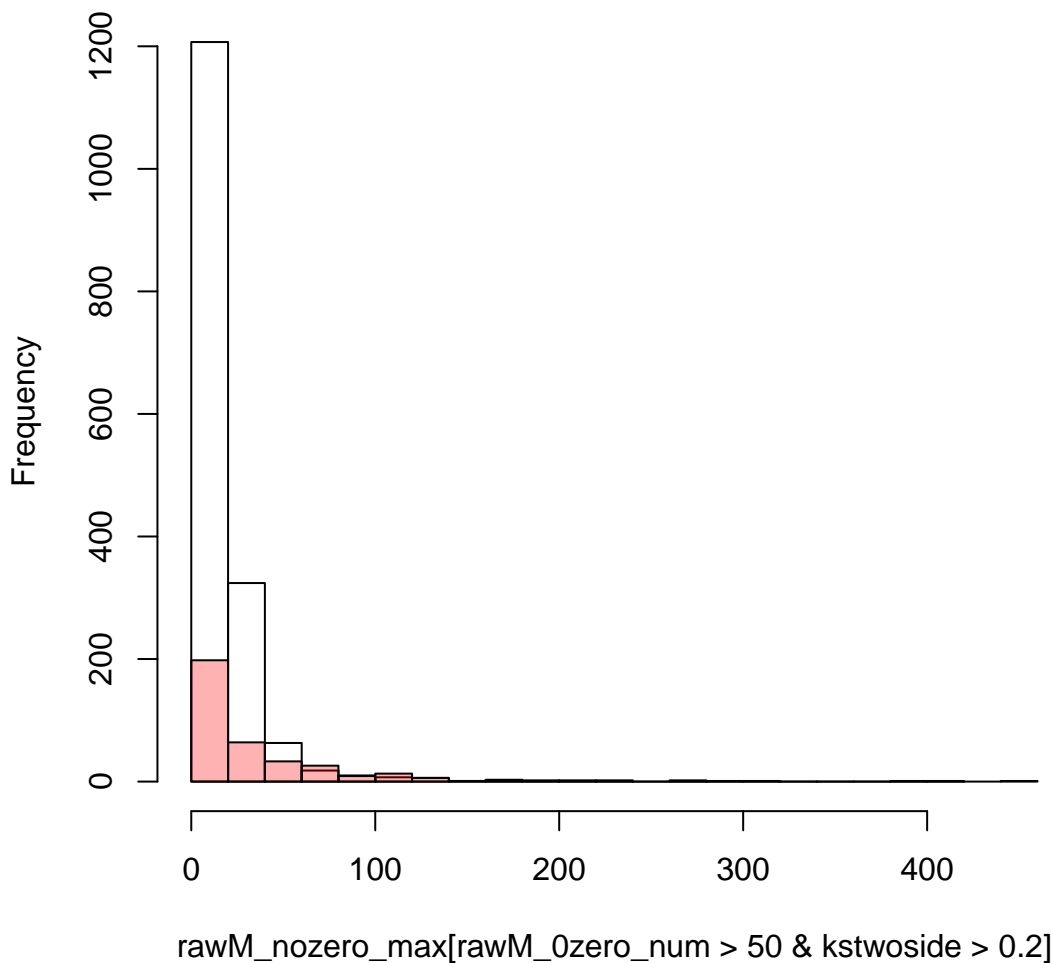
histogram of rawM\_nozero\_sd[rawM\_0zero\_num > 50 & kstwoside



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

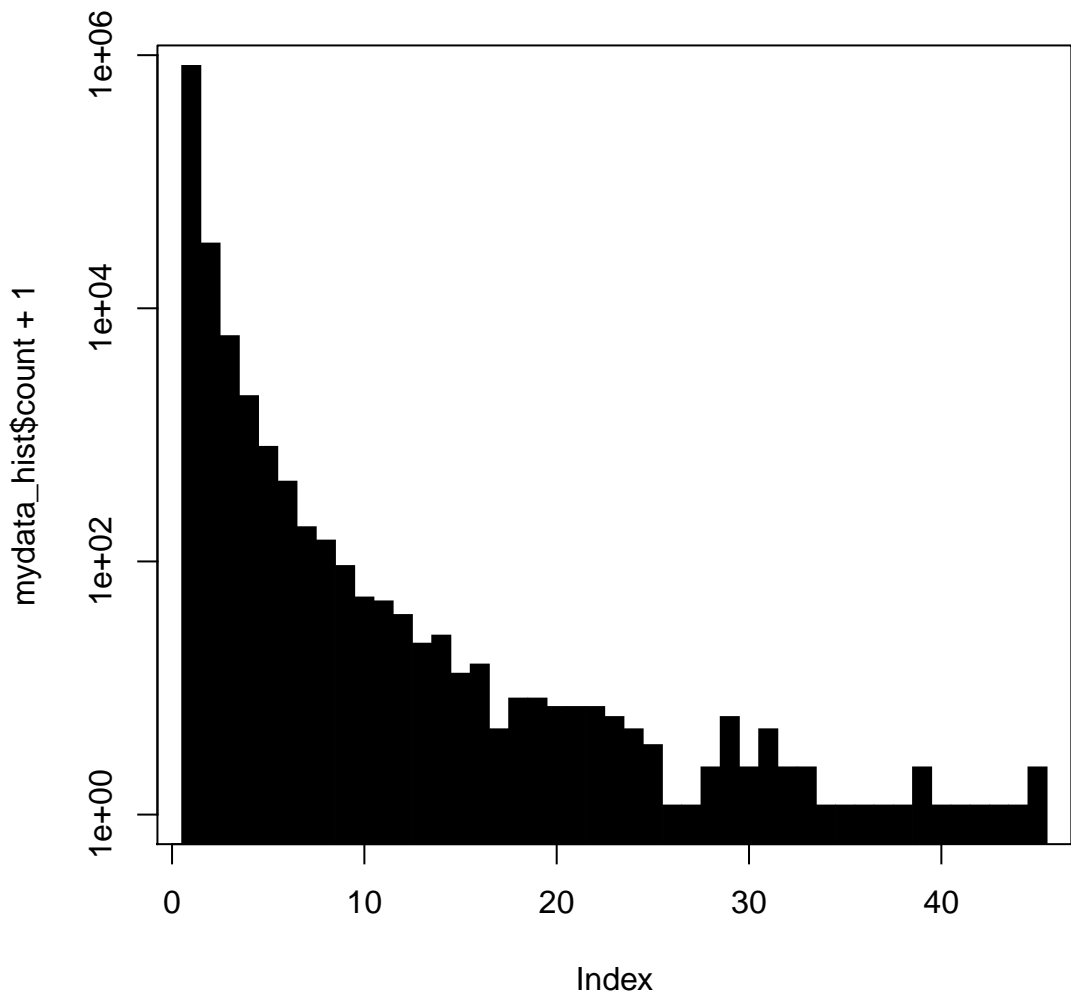


Histogram of rawM\_nozero\_max[rawM\_0zero\_num > 50 & kstwside

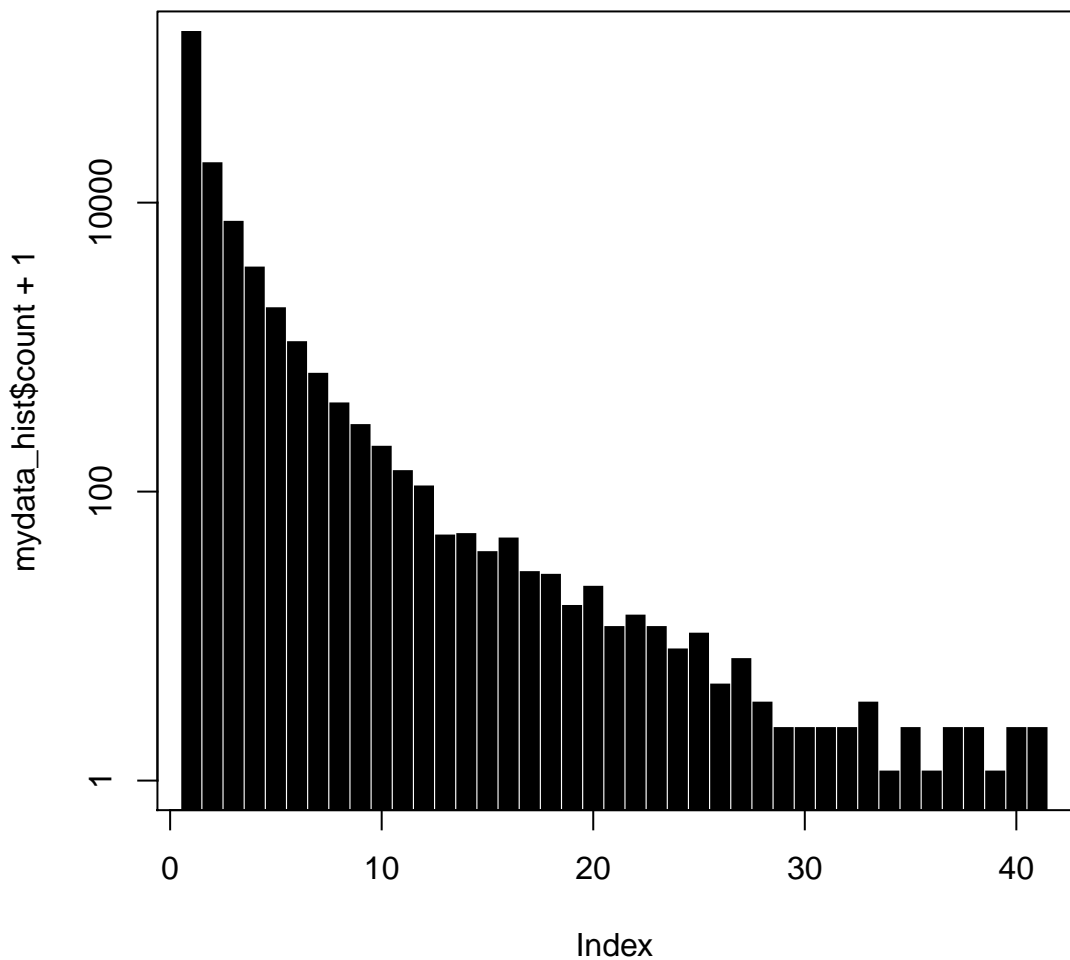




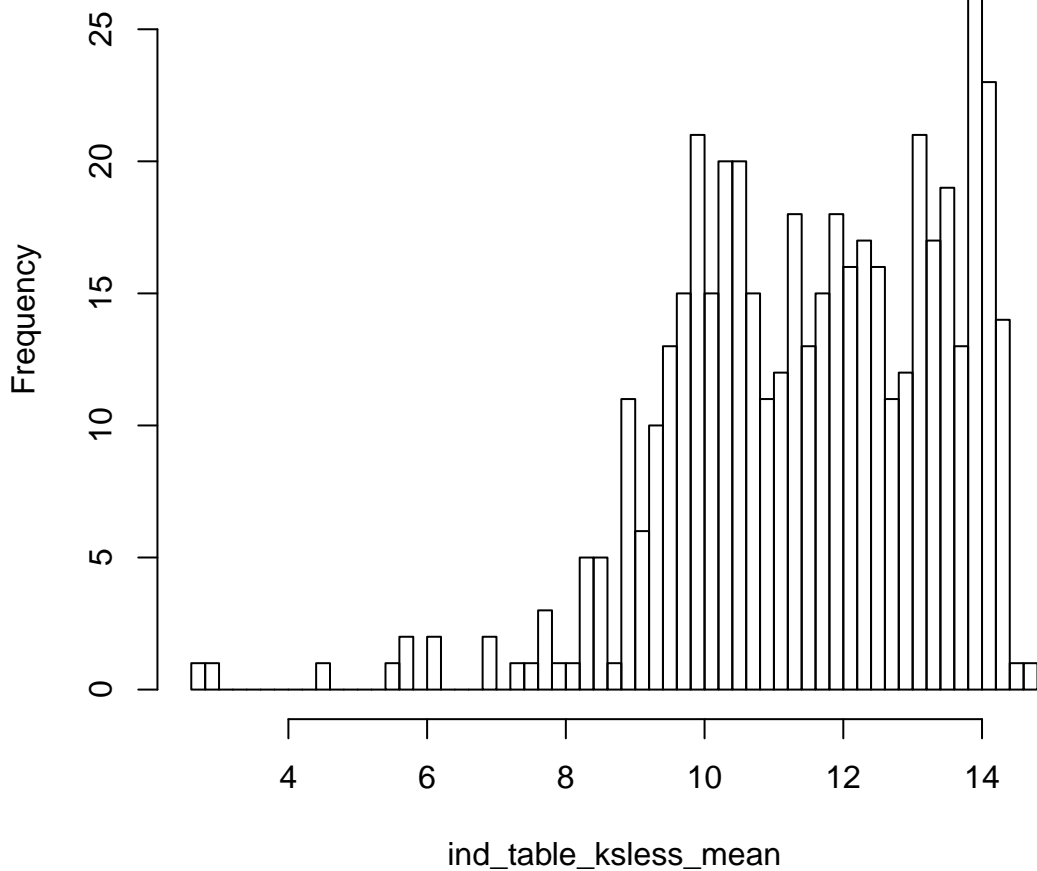
genes  $\log(\text{expression} + 1)$  with least 50 cell expression and kstvos



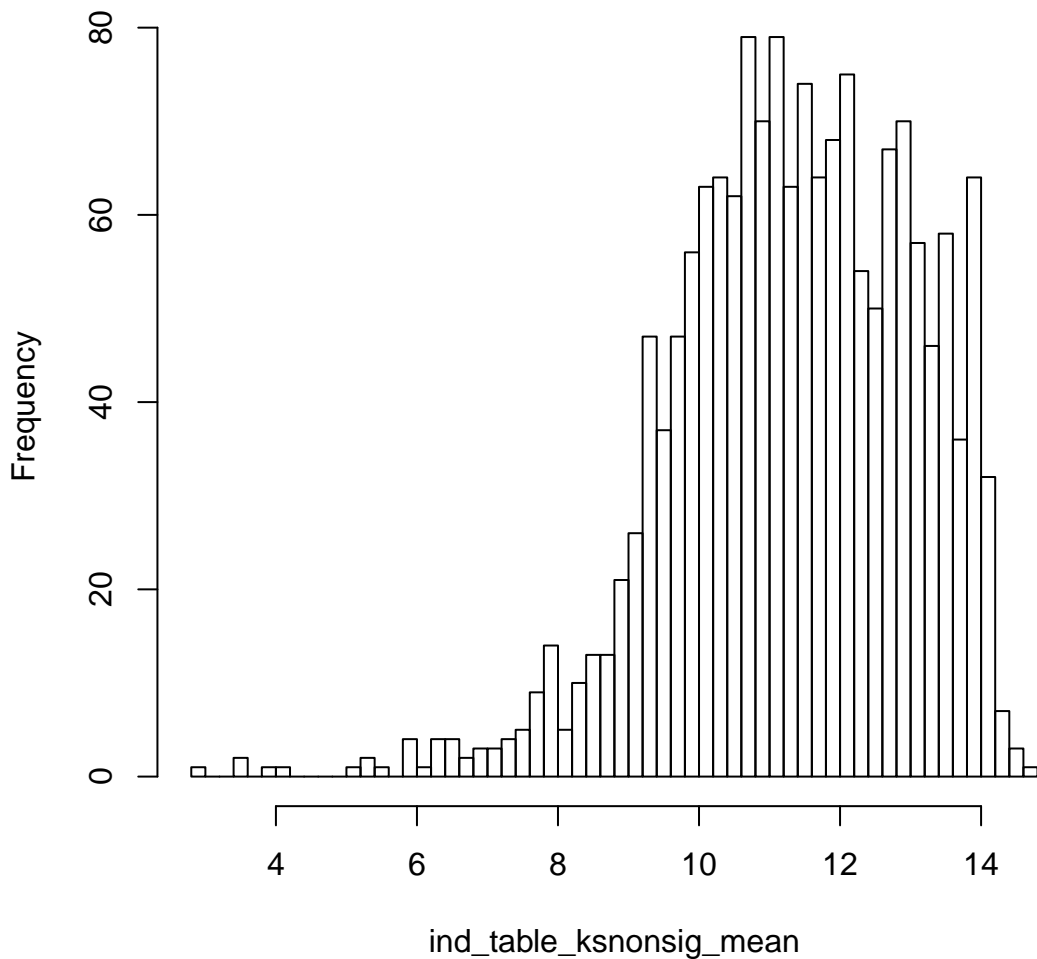
genes  $\log(\text{expression} + 1)$  with least 50 cell expression and kstwo



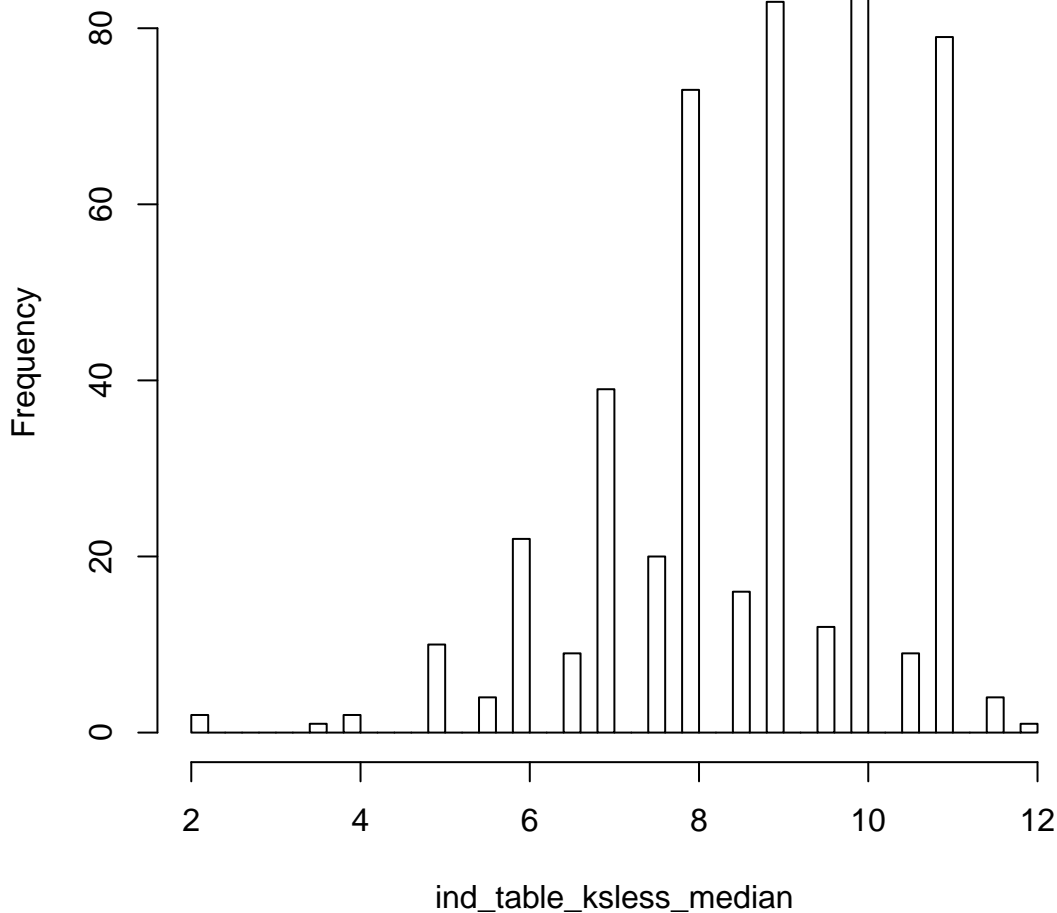
**Histogram of ind\_table\_ksless\_mean**



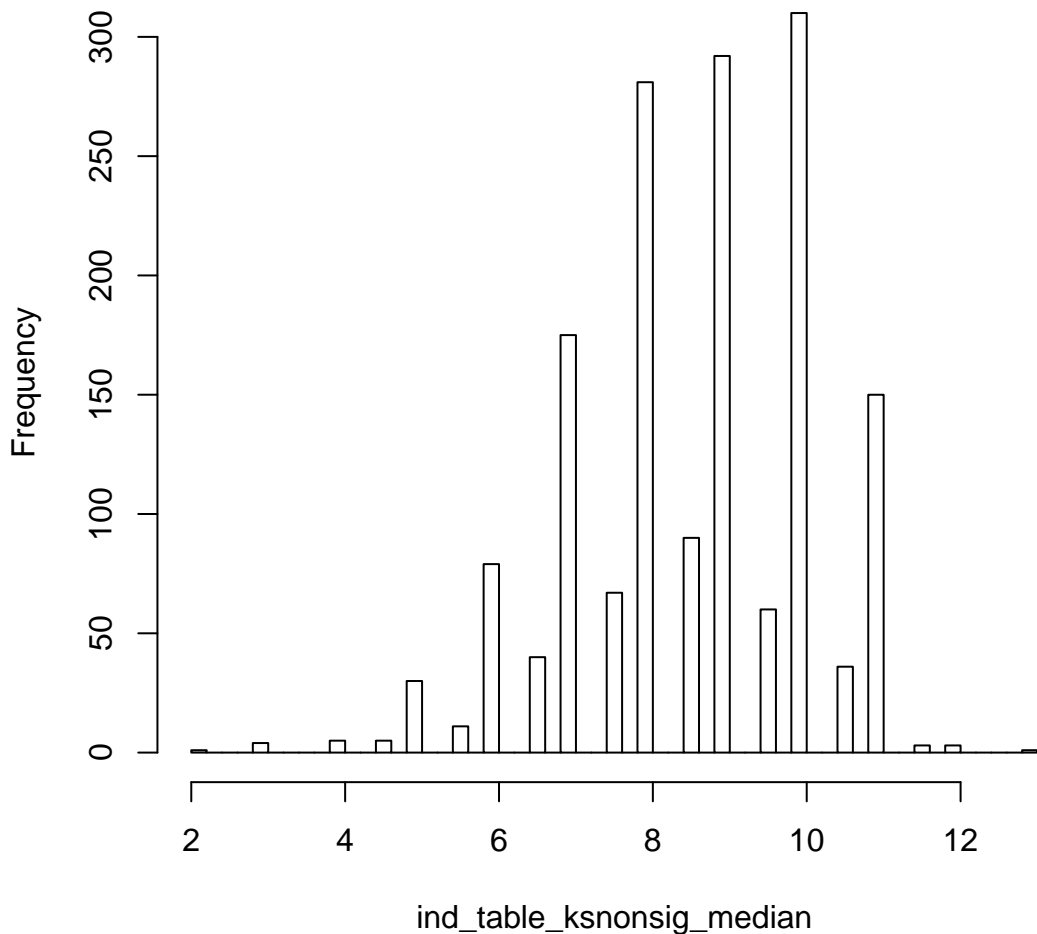
**Histogram of ind\_table\_ksnonsig\_mean**



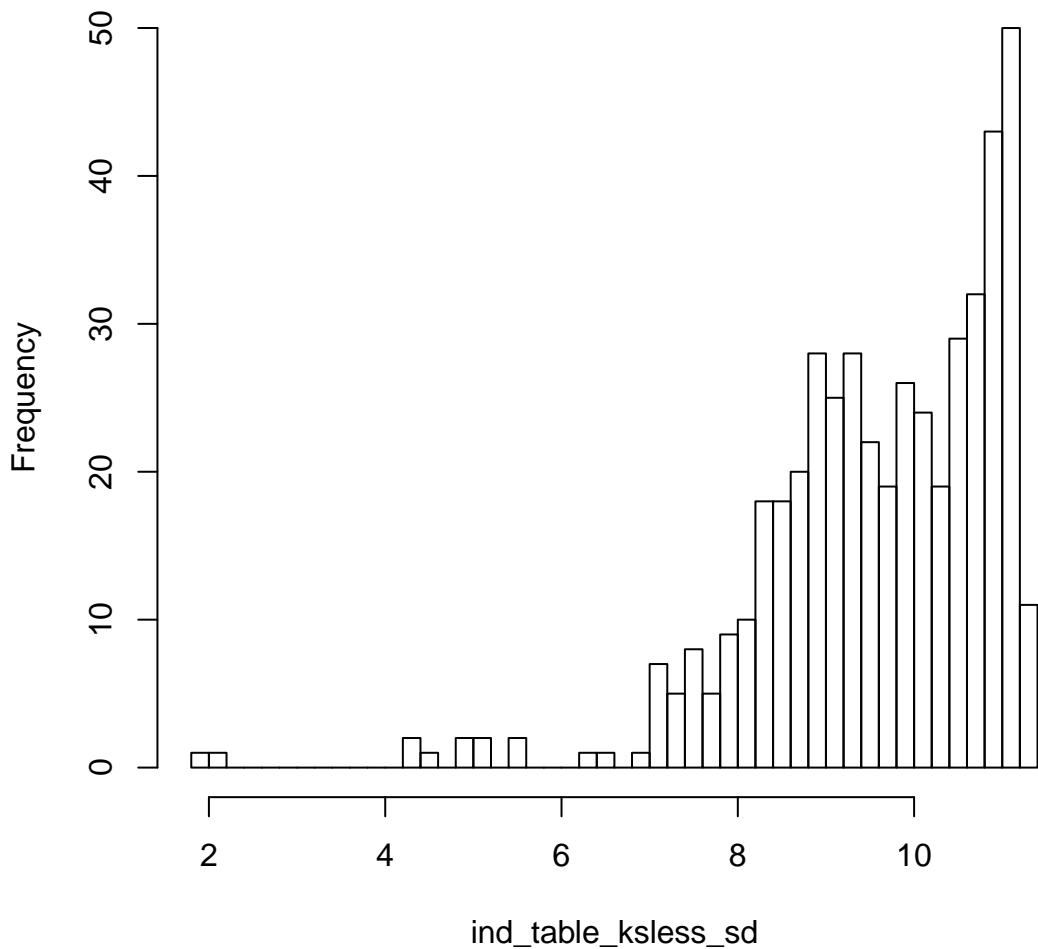
**Histogram of ind\_table\_ksless\_median**



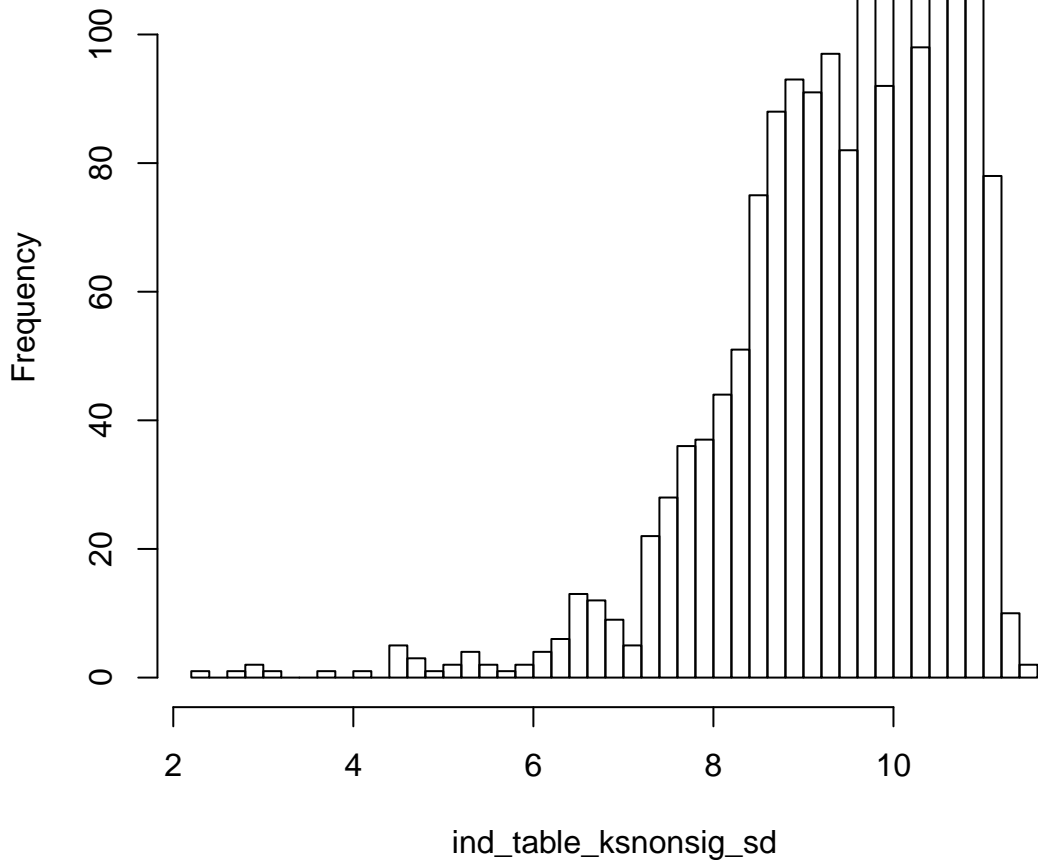
**Histogram of ind\_table\_ksnonsig\_median**



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

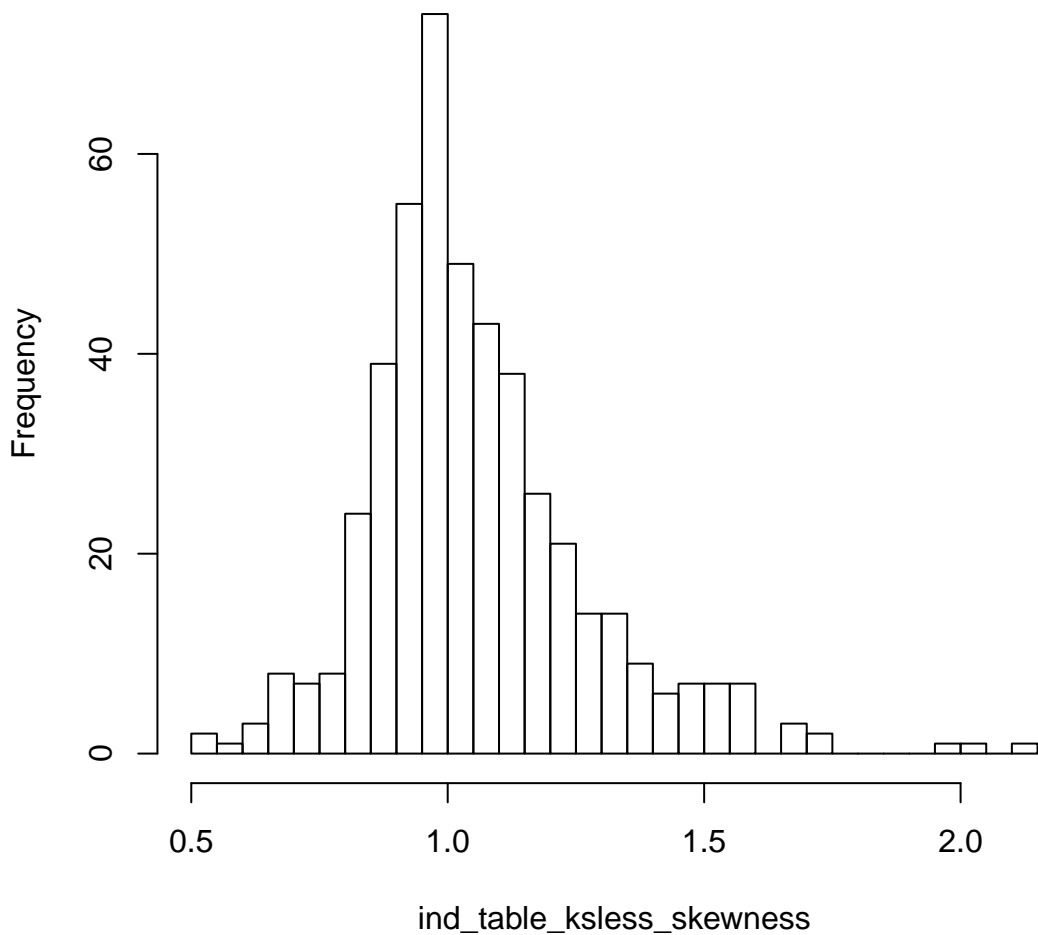


**Histogram of ind\_table\_ksnonsig\_sd**

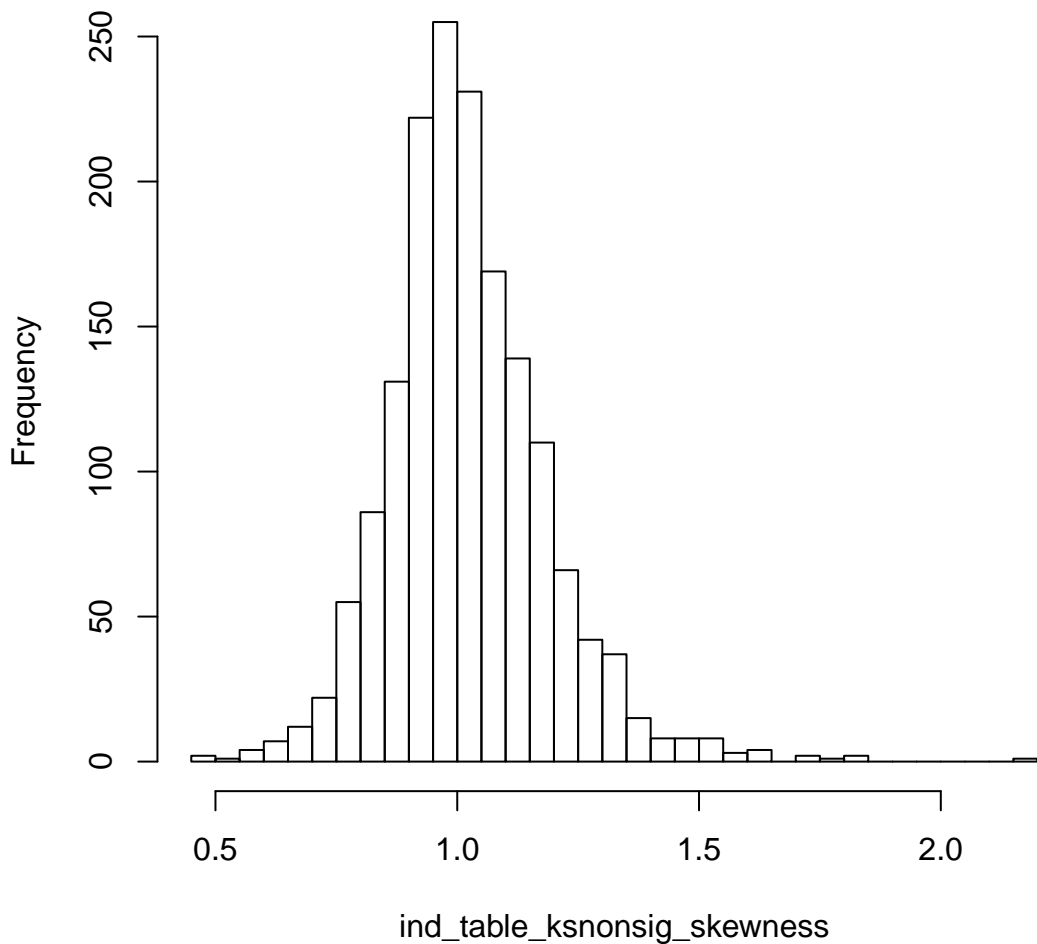




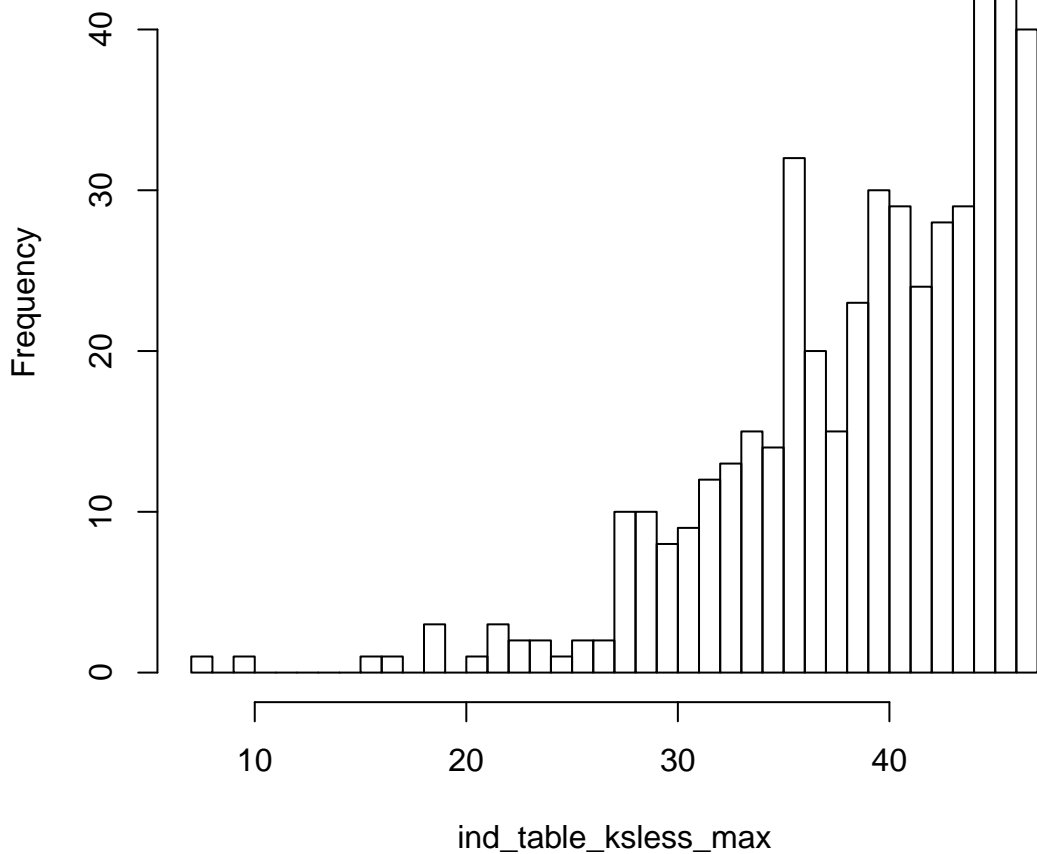
**Histogram of ind\_table\_ksless\_skewness**



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



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



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

