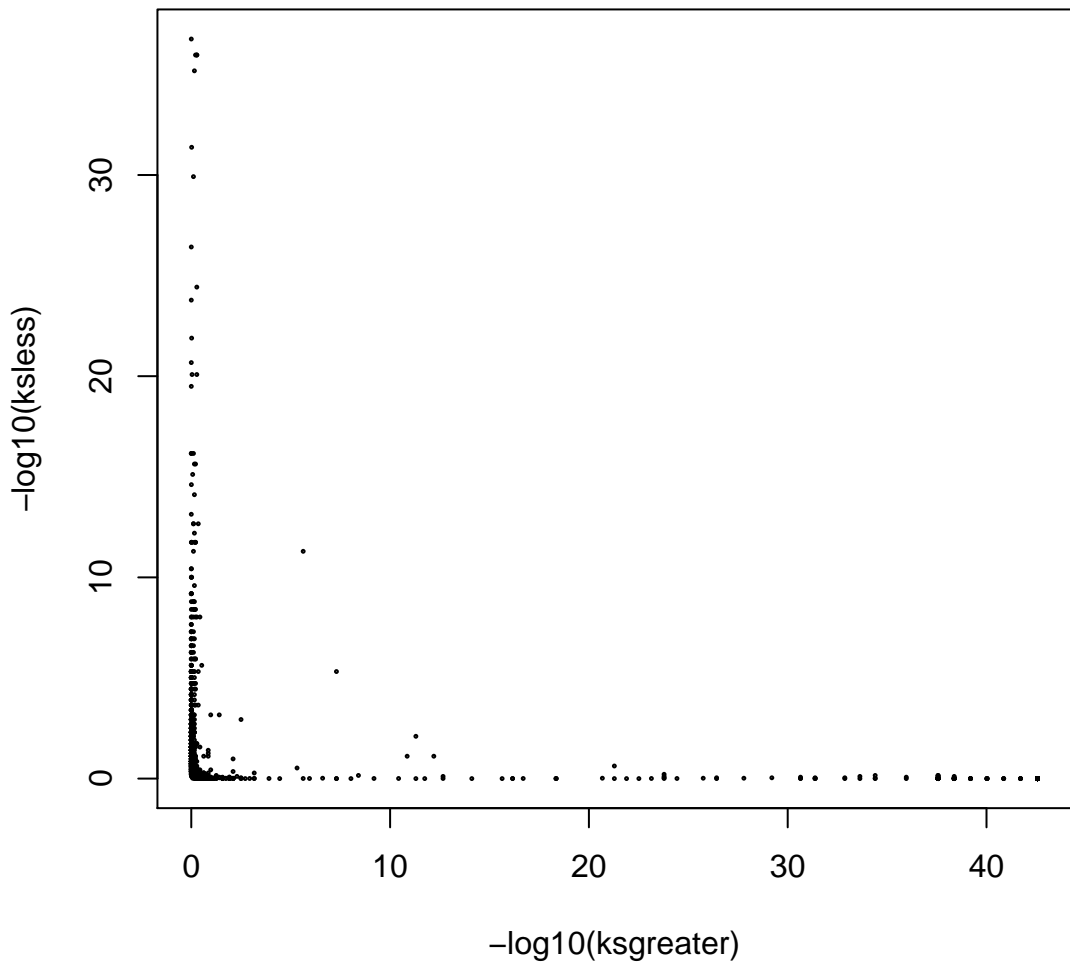
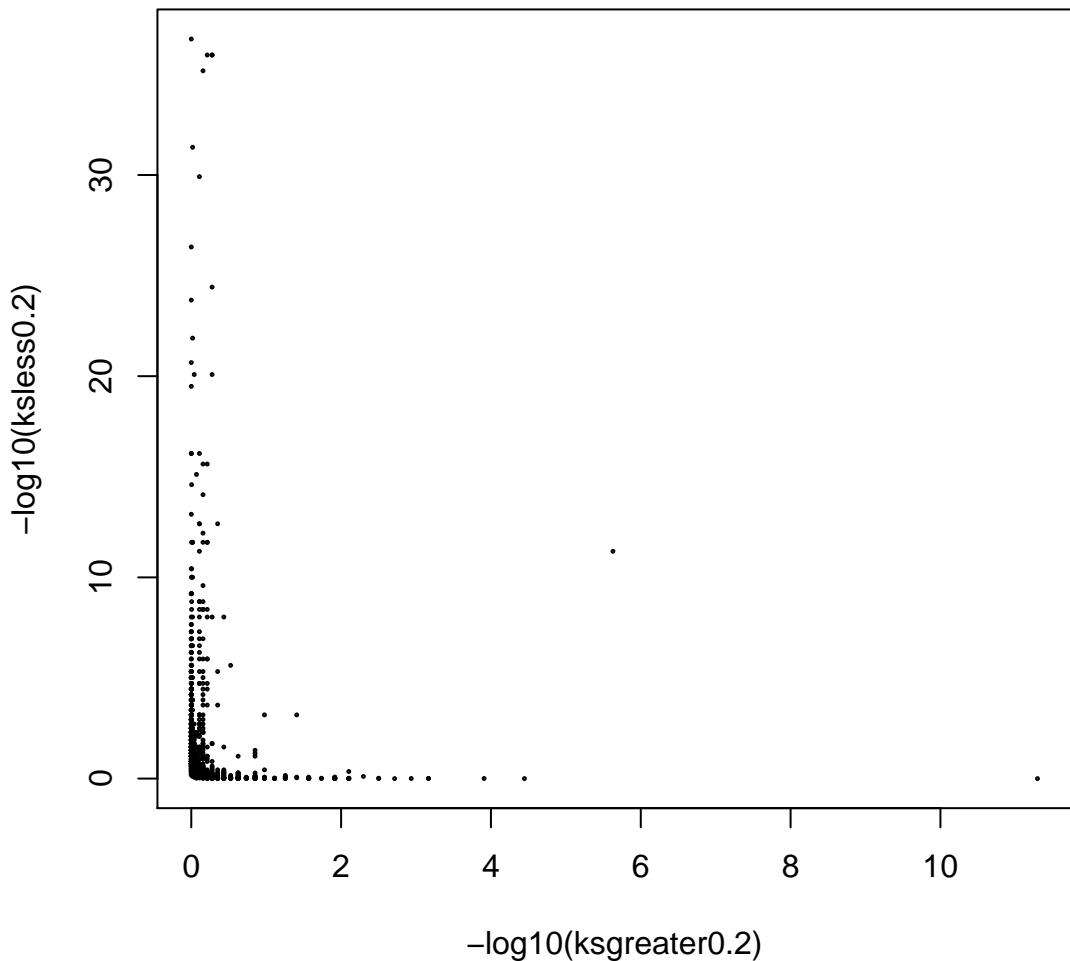


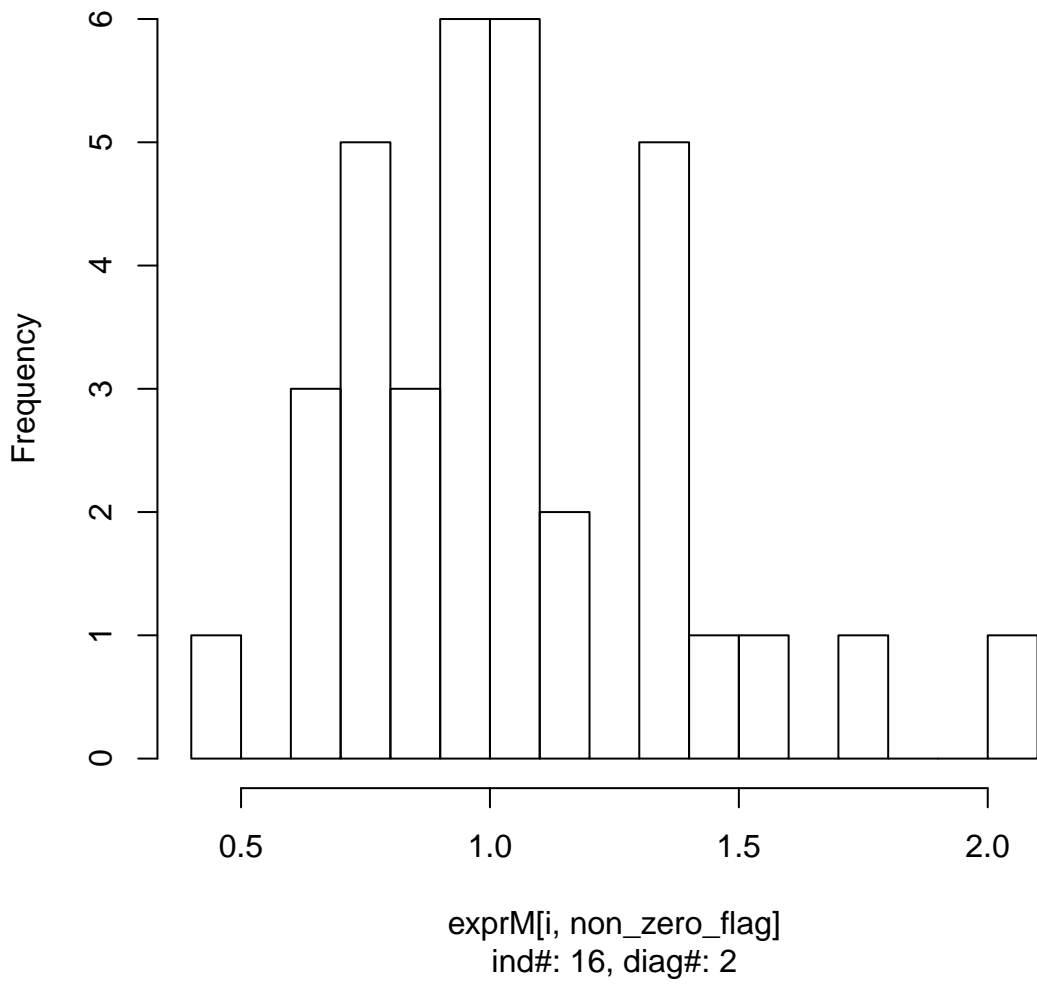
sig_KSgreater: 60.667%, sig_KSless: 15.967%



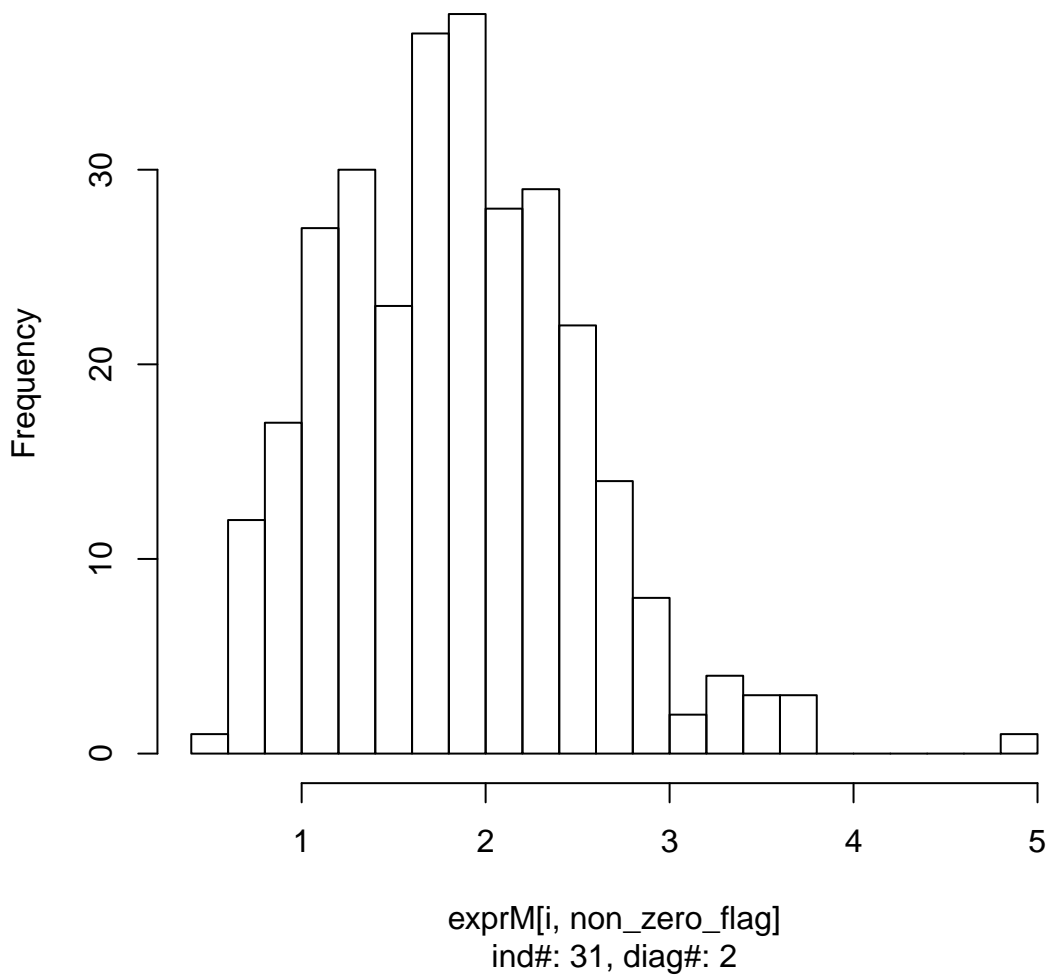
sig_KSgreater0.2: 3.199%, sig_KSless0.2: 39.048%



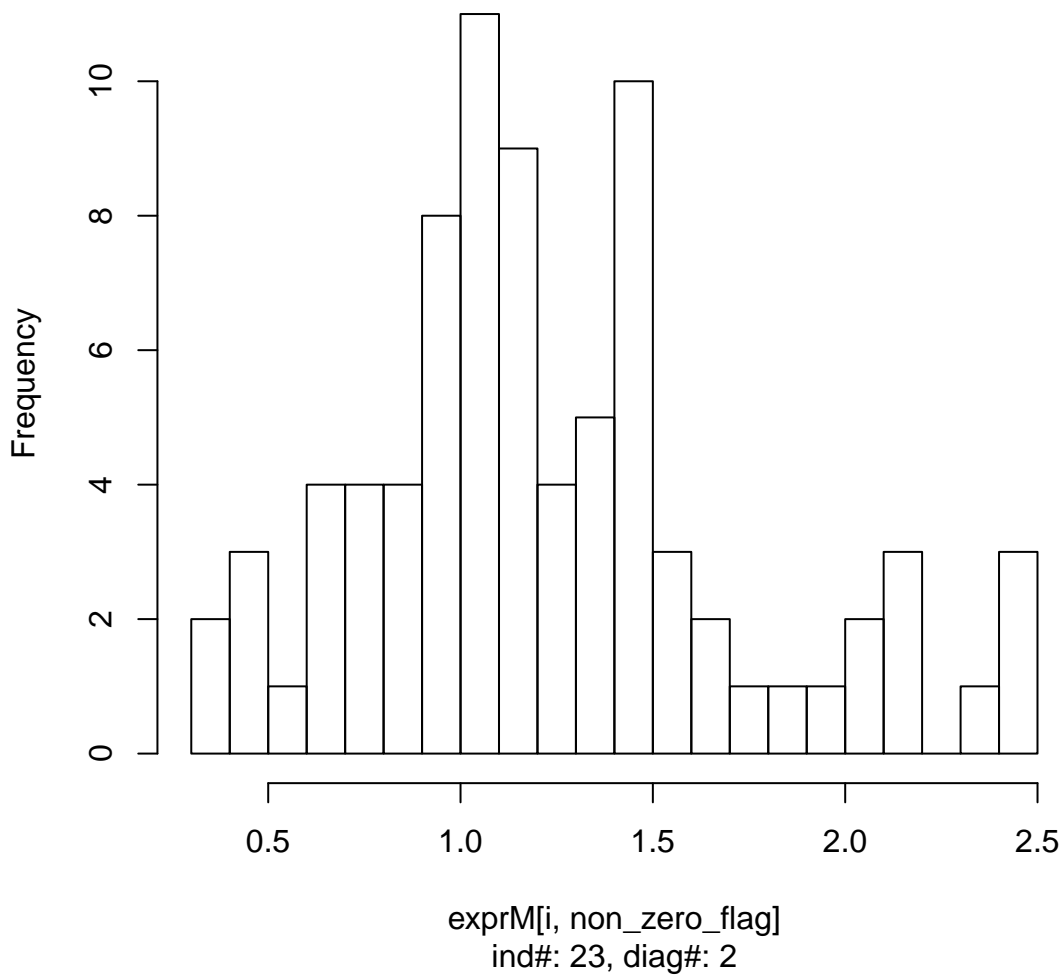
log expression of gene#2013, pval ob=0.2723, non-zero num=3



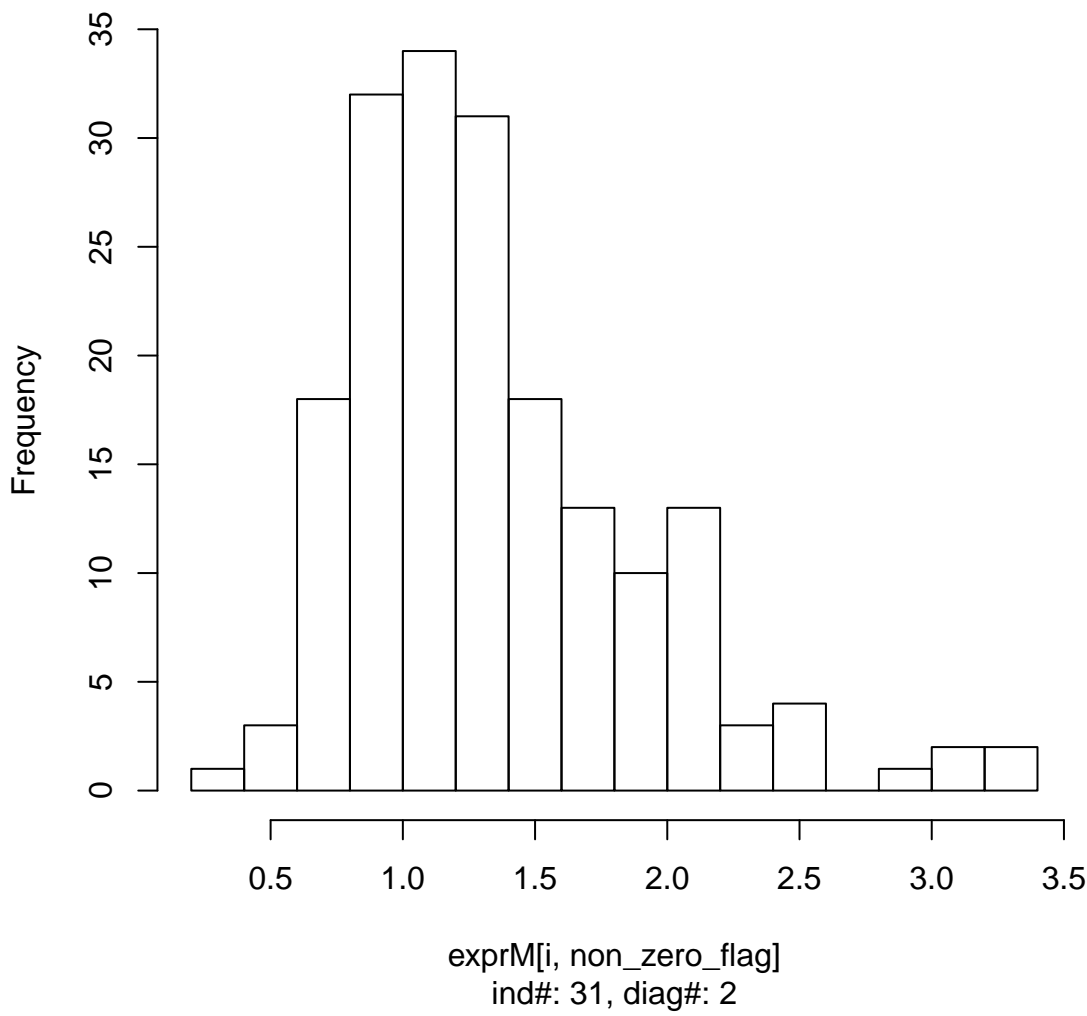
log expression of gene#104, pval ob=0.1902, non-zero num=29



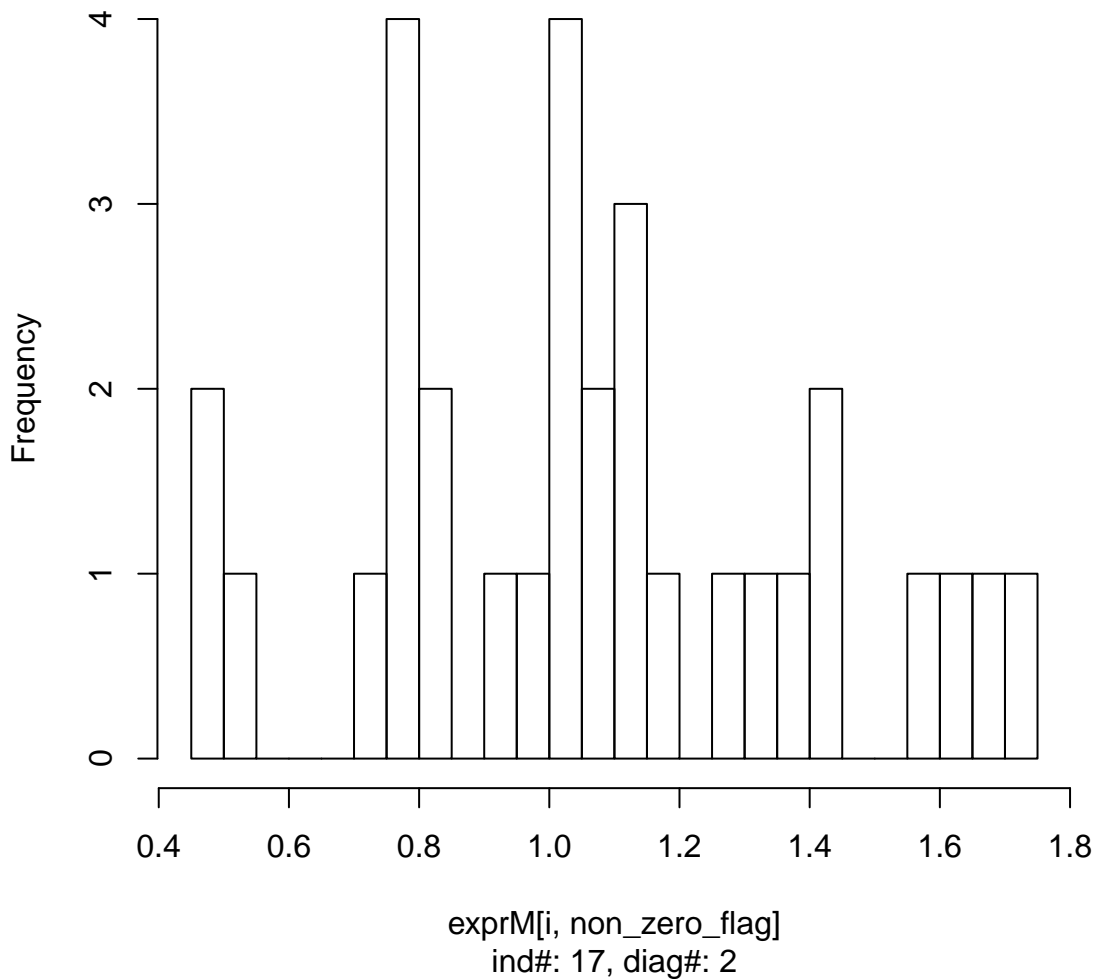
log expression of gene#680, pval ob=0.669, non-zero num=82



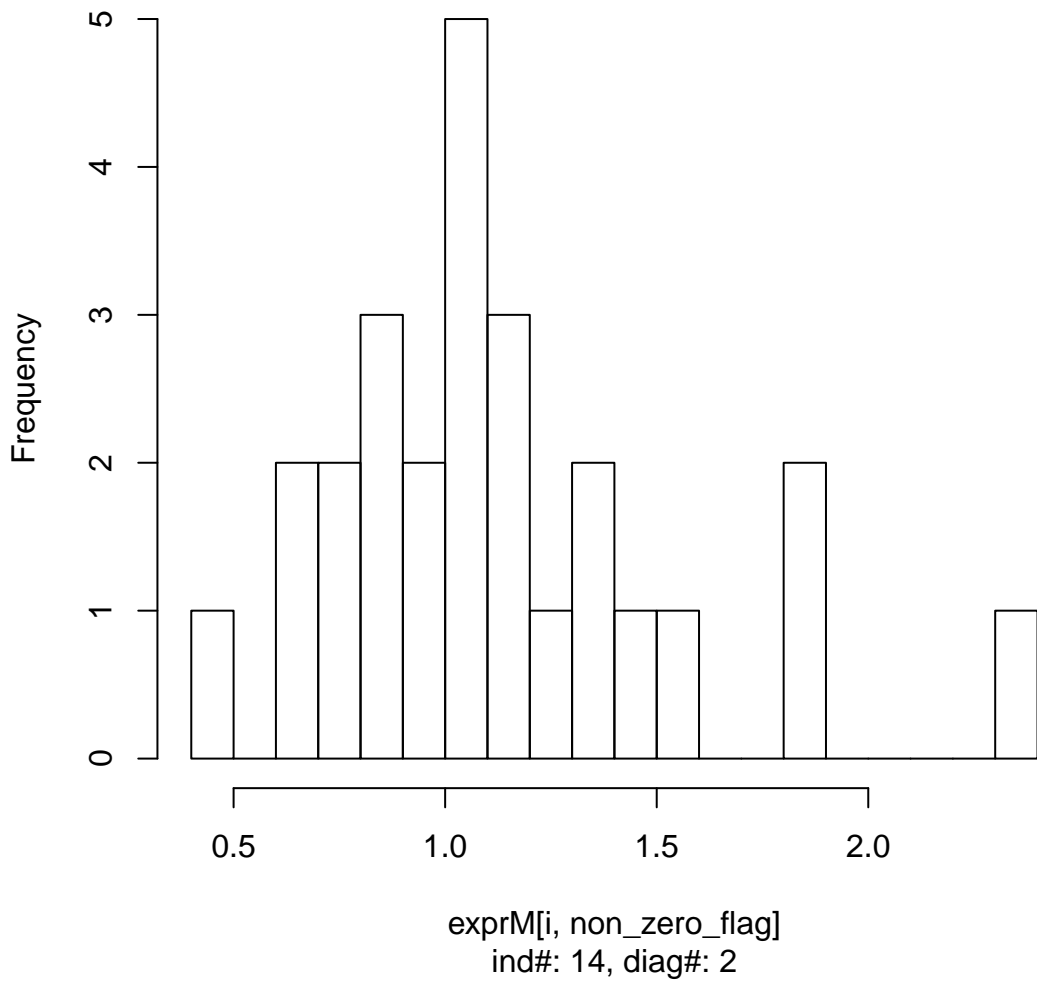
log expression of gene#730, pval ob=0.2179, non-zero num=18



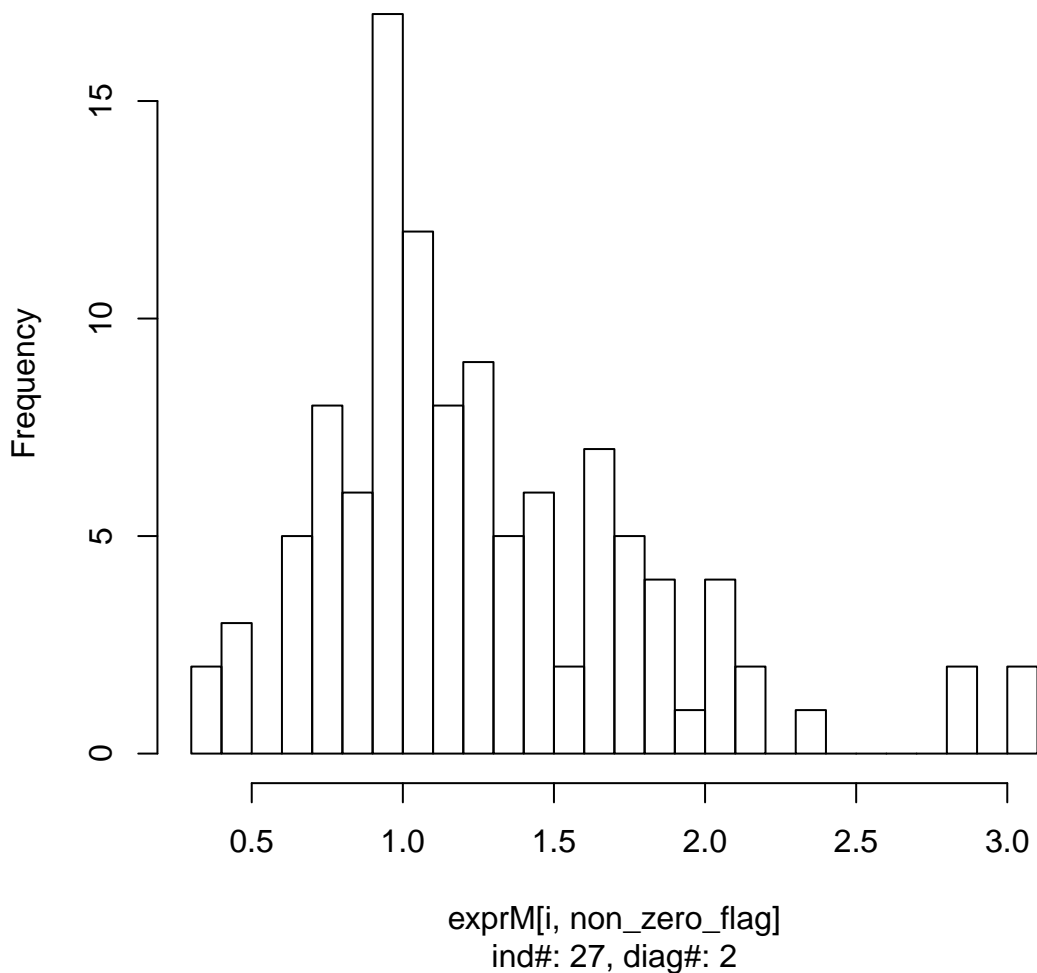
log expression of gene#750, pval ob=0.4024, non-zero num=3



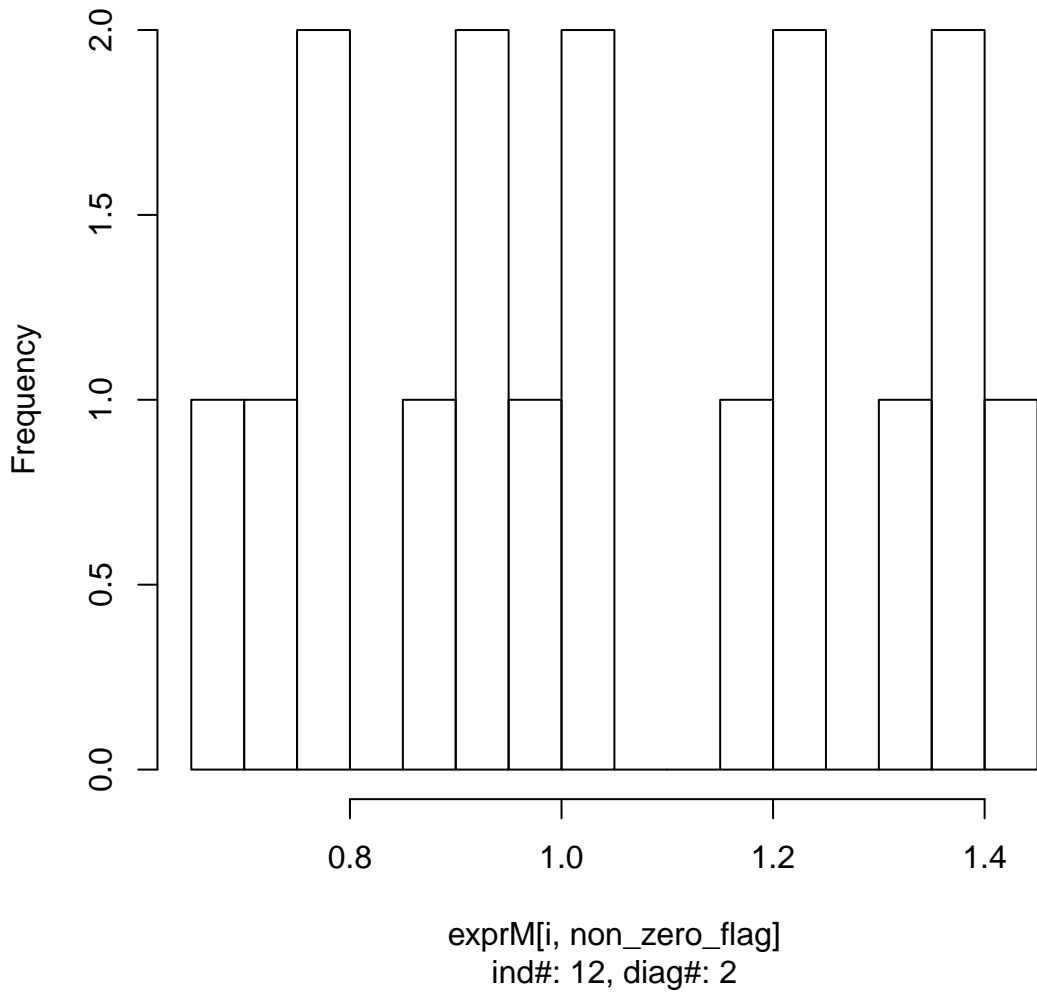
log expression of gene#2565, pval ob=0.8225, non-zero num=2



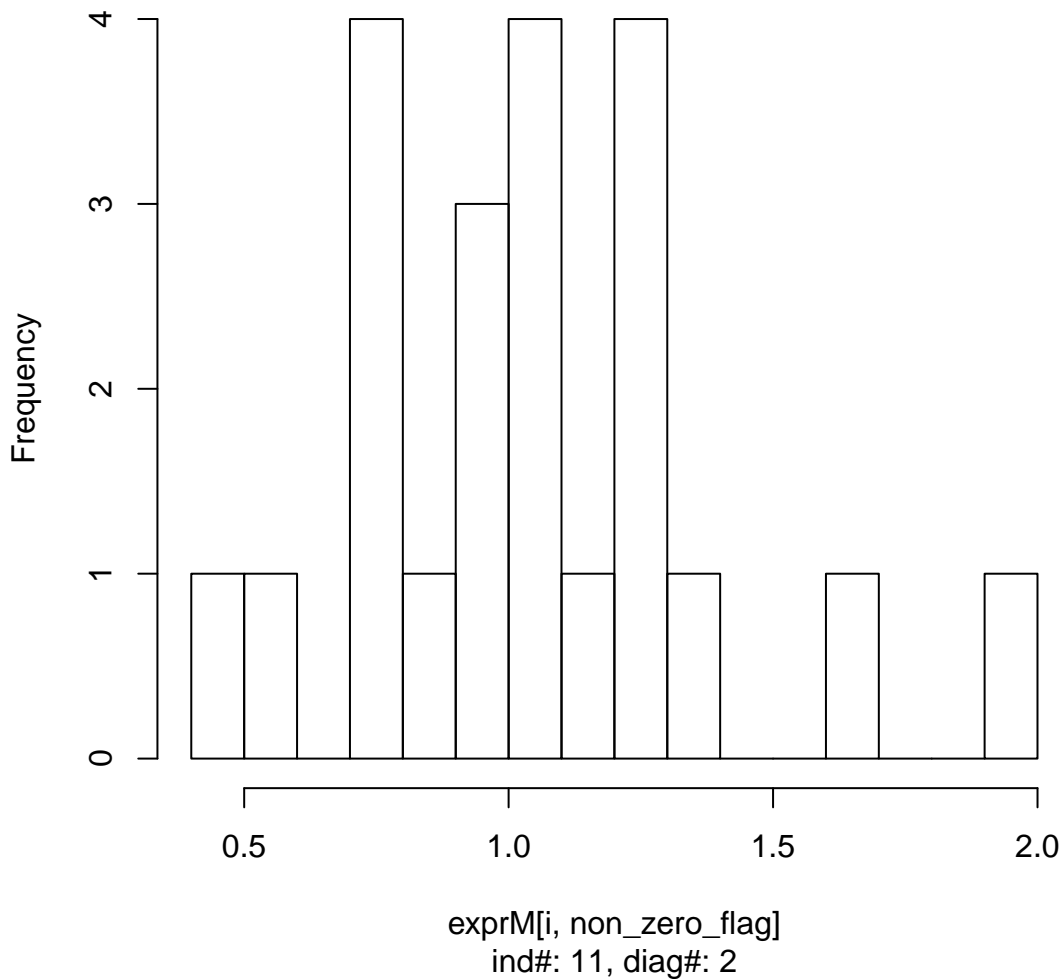
log expression of gene#1047, pval ob=0.6786, non-zero num=1



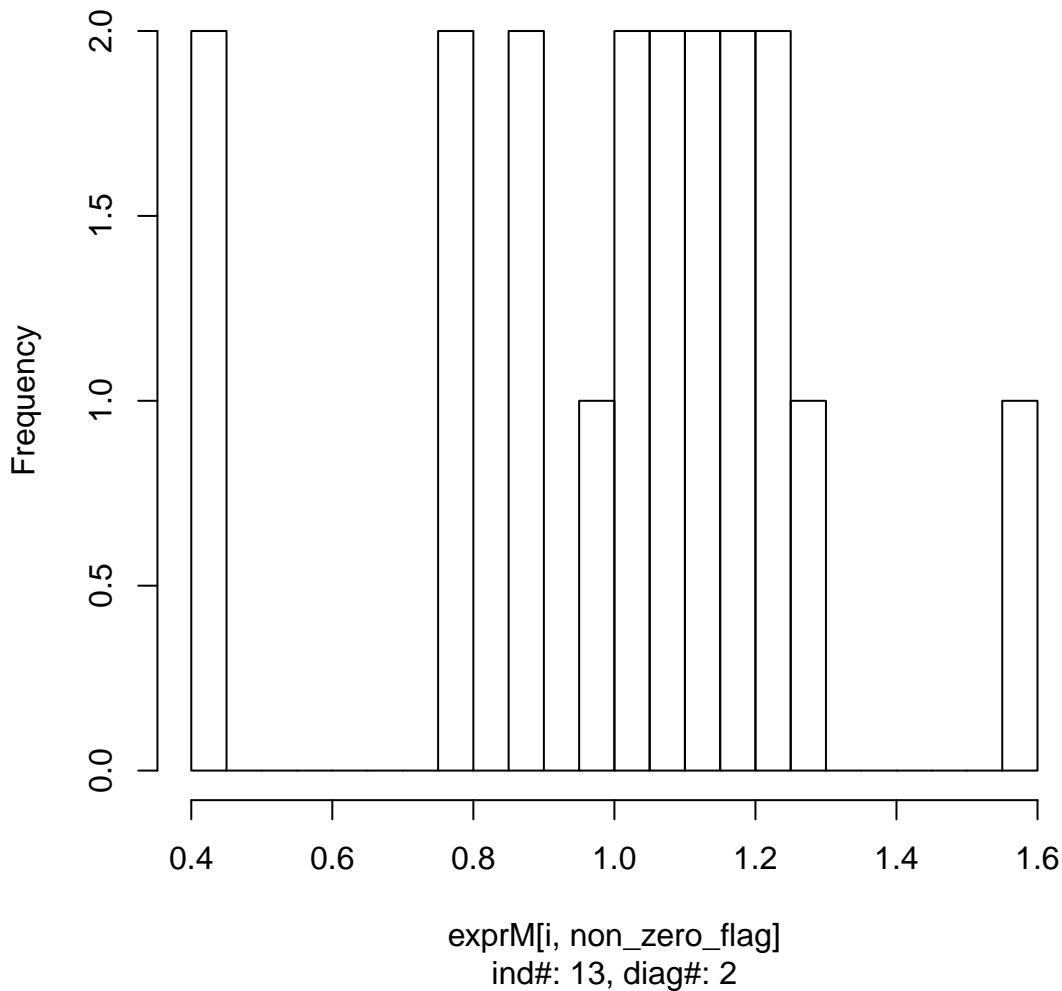
log expression of gene#2613, pval ob=0.0029, non-zero num=1



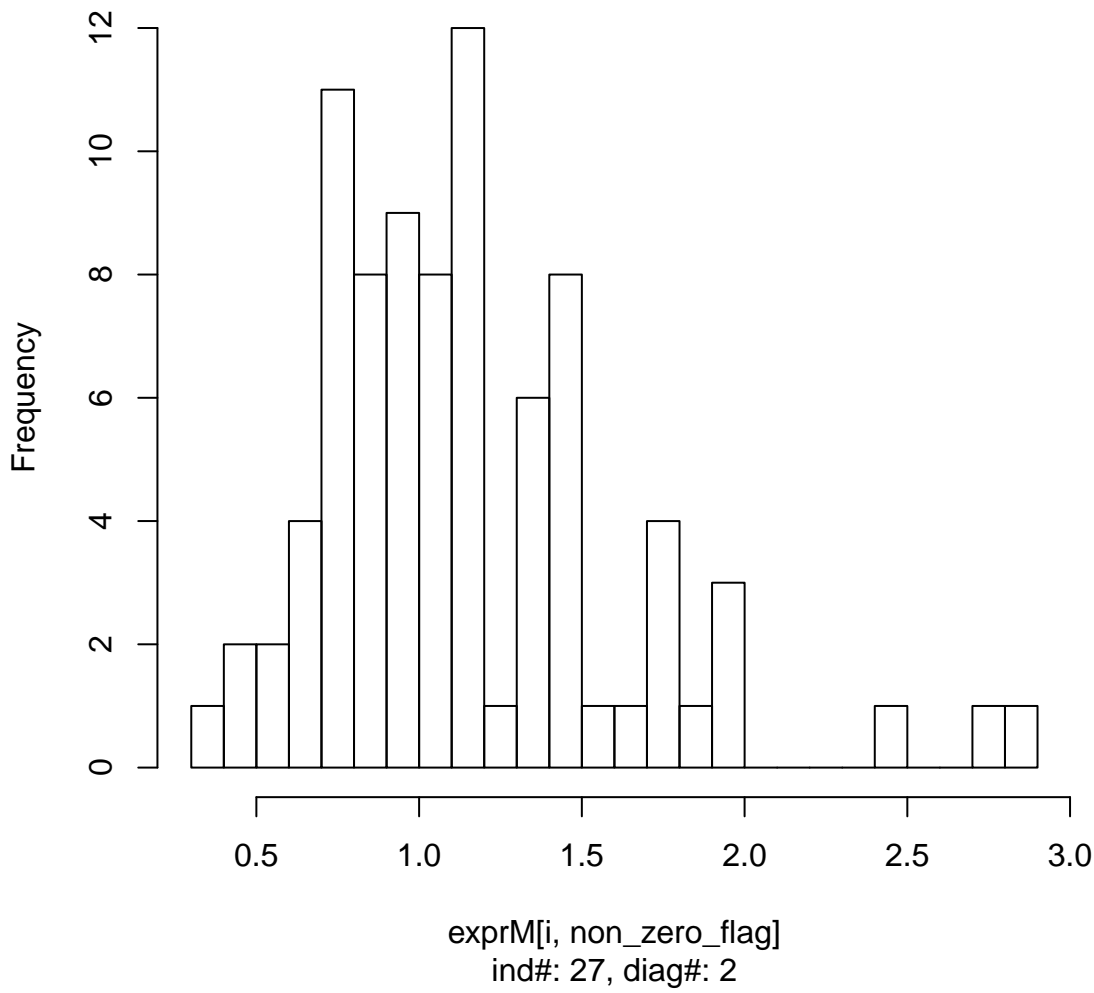
log expression of gene#1892, pval ob=0.0018, non-zero num=2



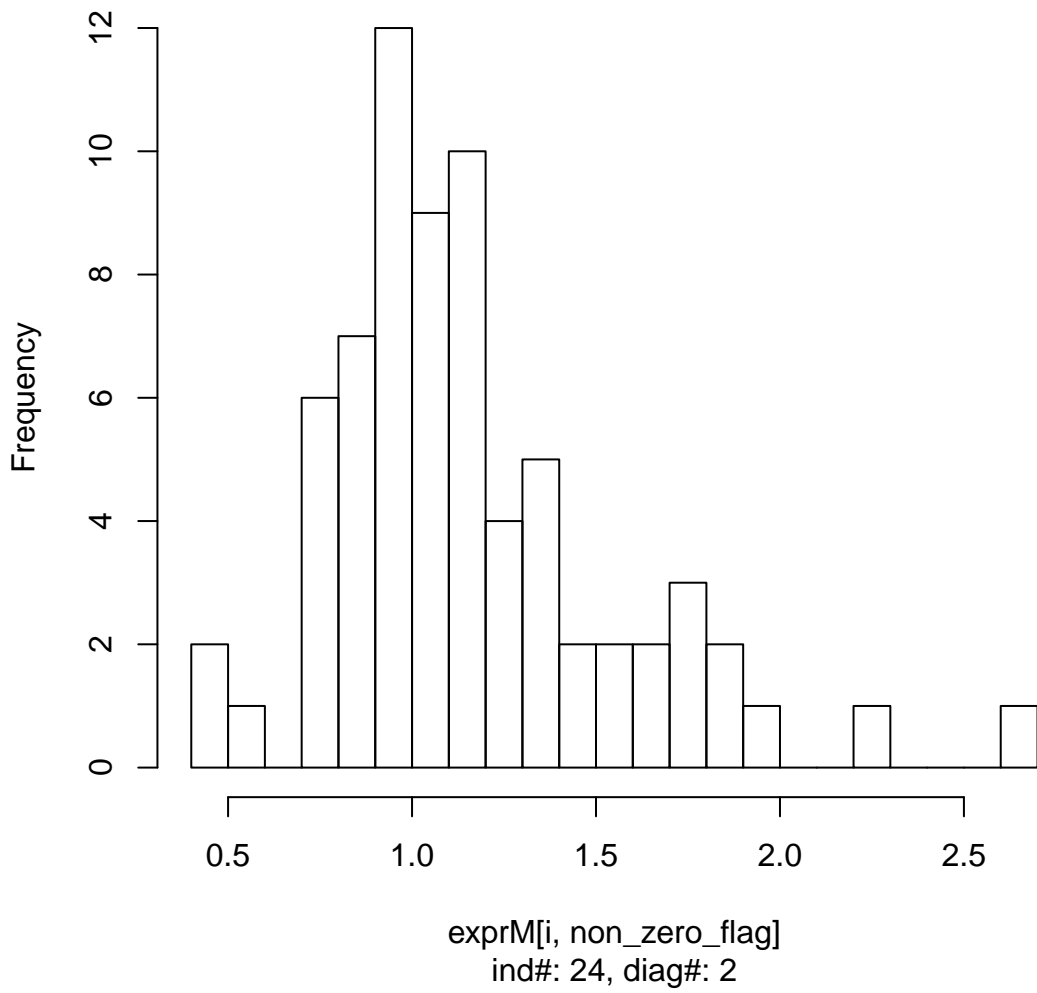
log expression of gene#1246, pval ob=0.3984, non-zero num=1



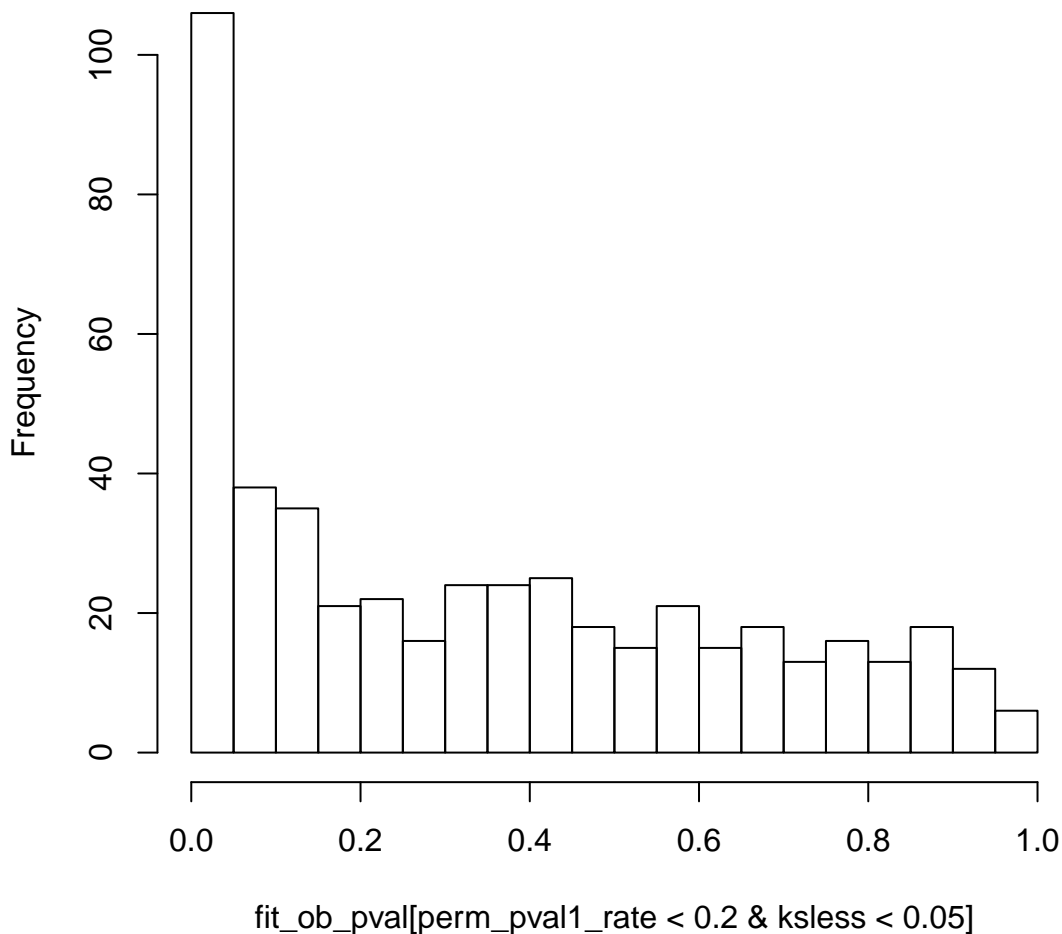
log expression of gene#1182, pval ob=0.1836, non-zero num=8



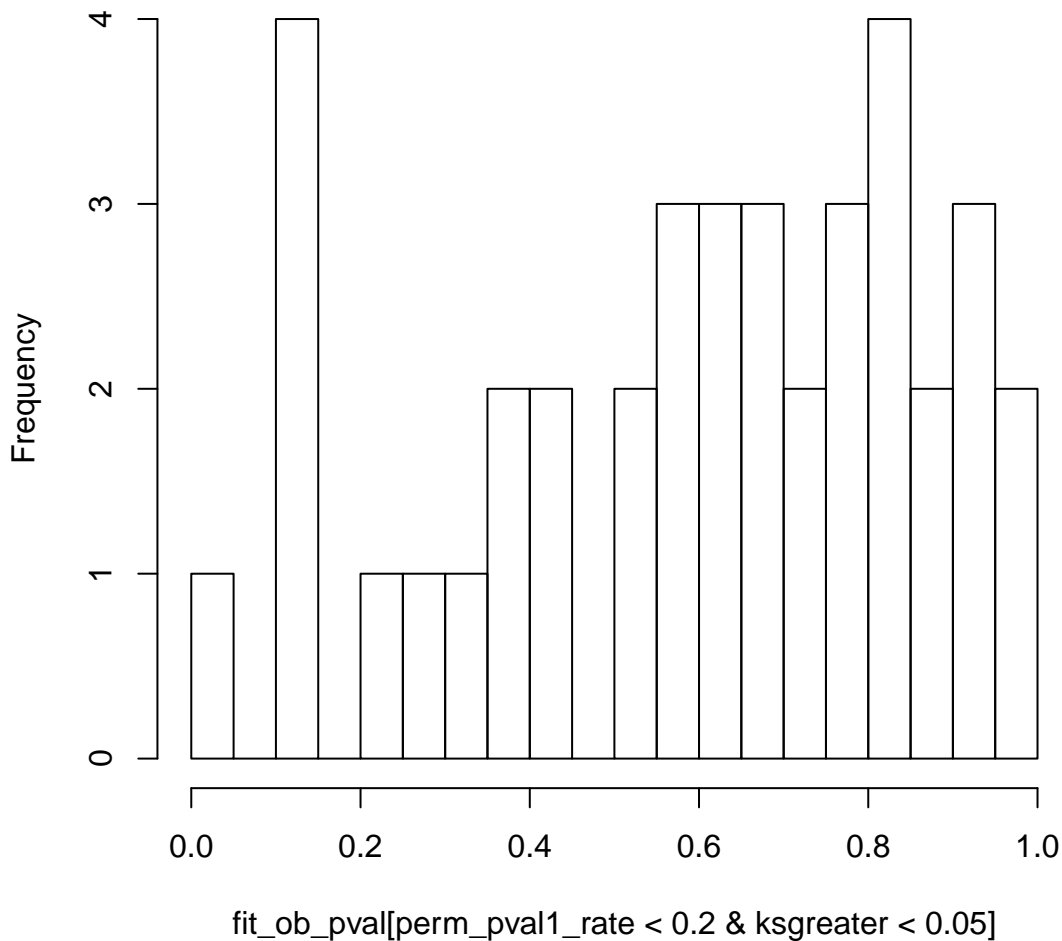
log expression of gene#2587, pval ob=0.3601, non-zero num=7



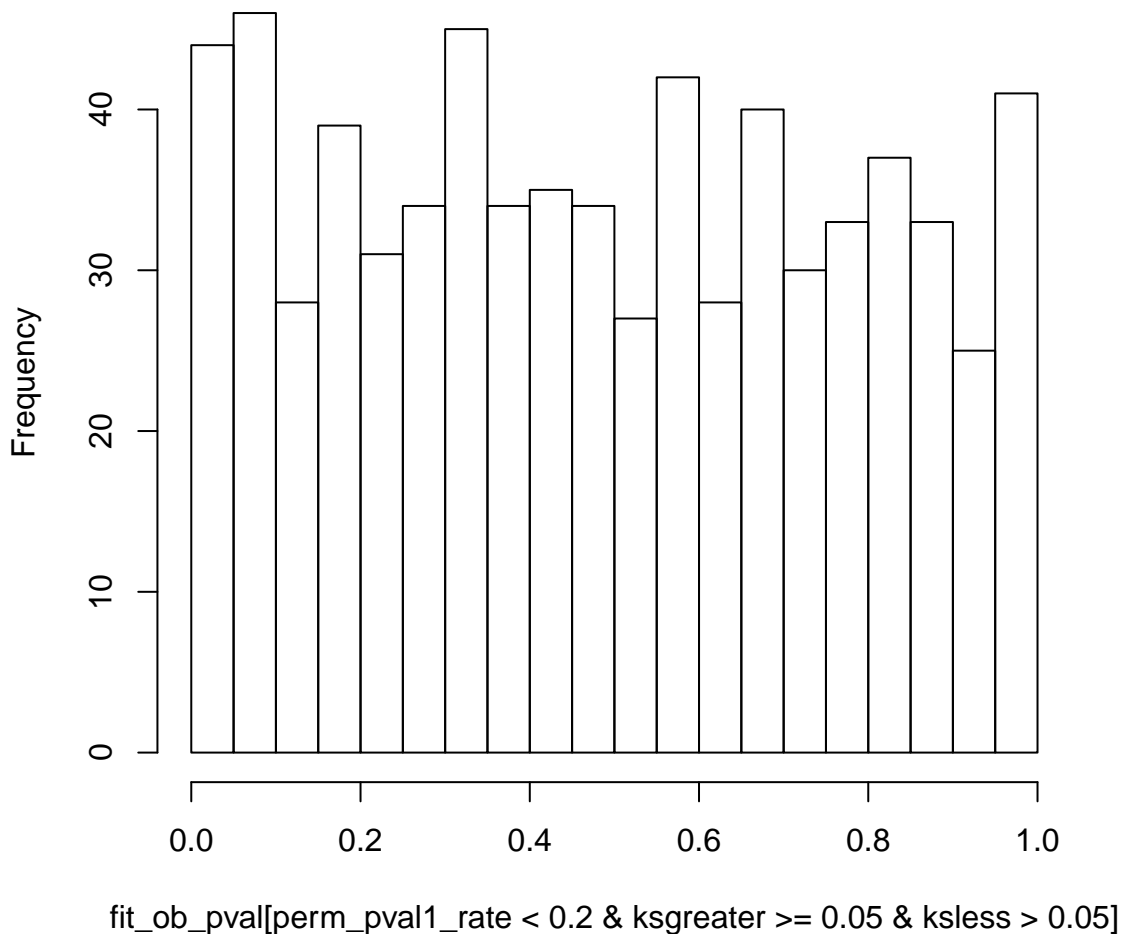
observed pvalues with pval1_rate<0.2,ksless sig



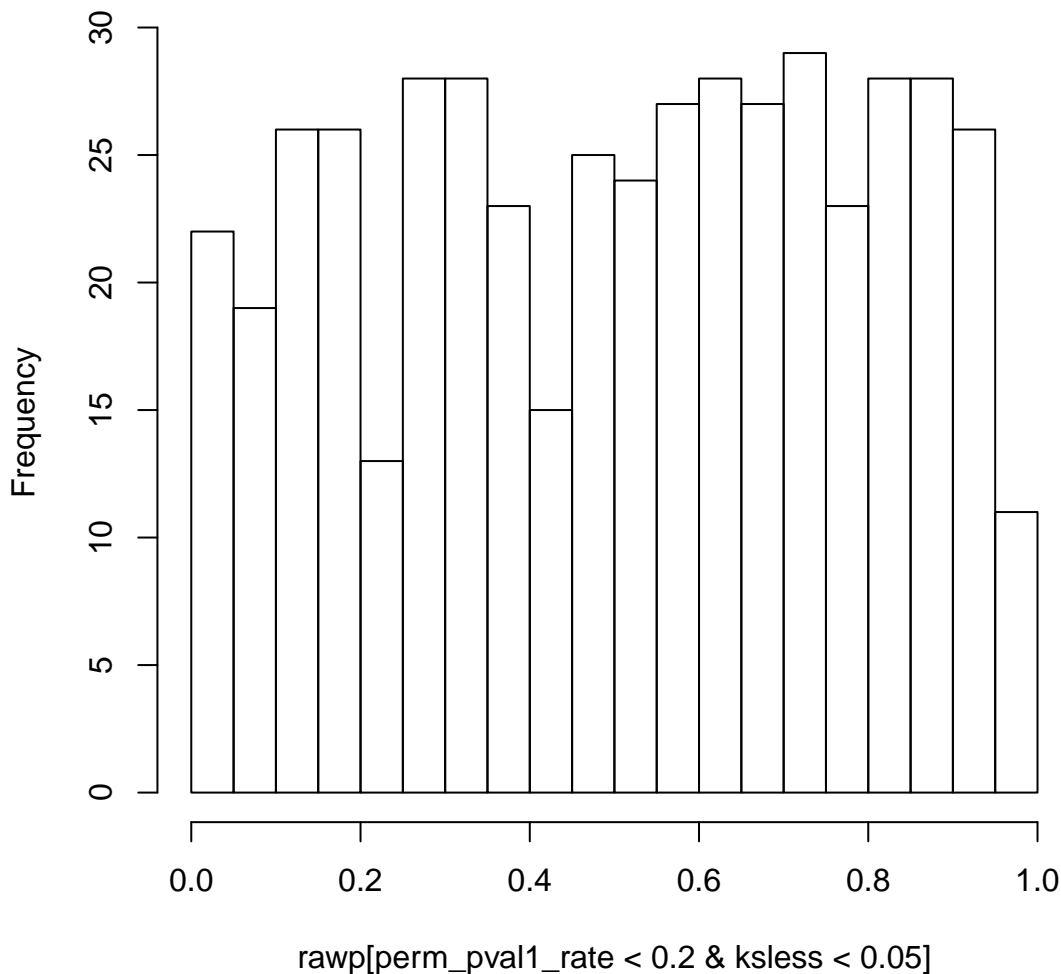
observed pvalues with pval1_rate<0.2,ksgreater sig



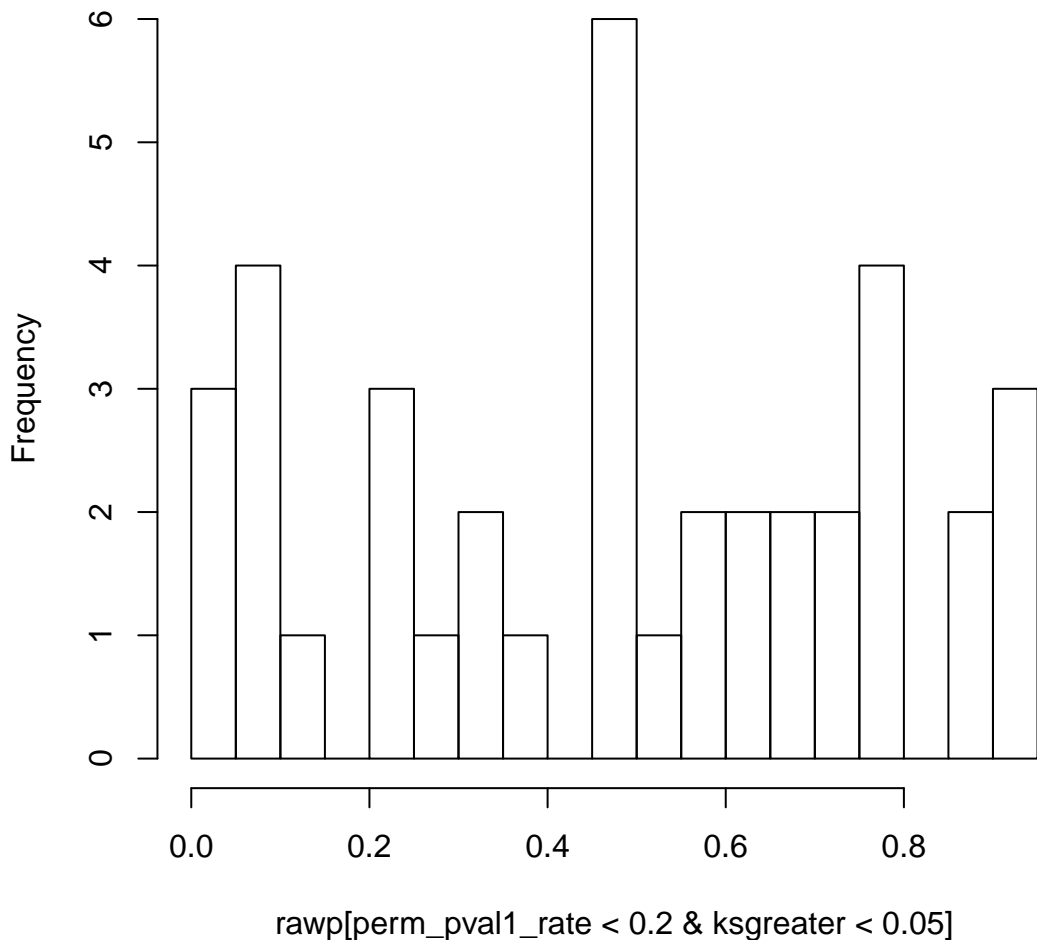
observed pvalues with pval1_rate<0.2,ks no sig



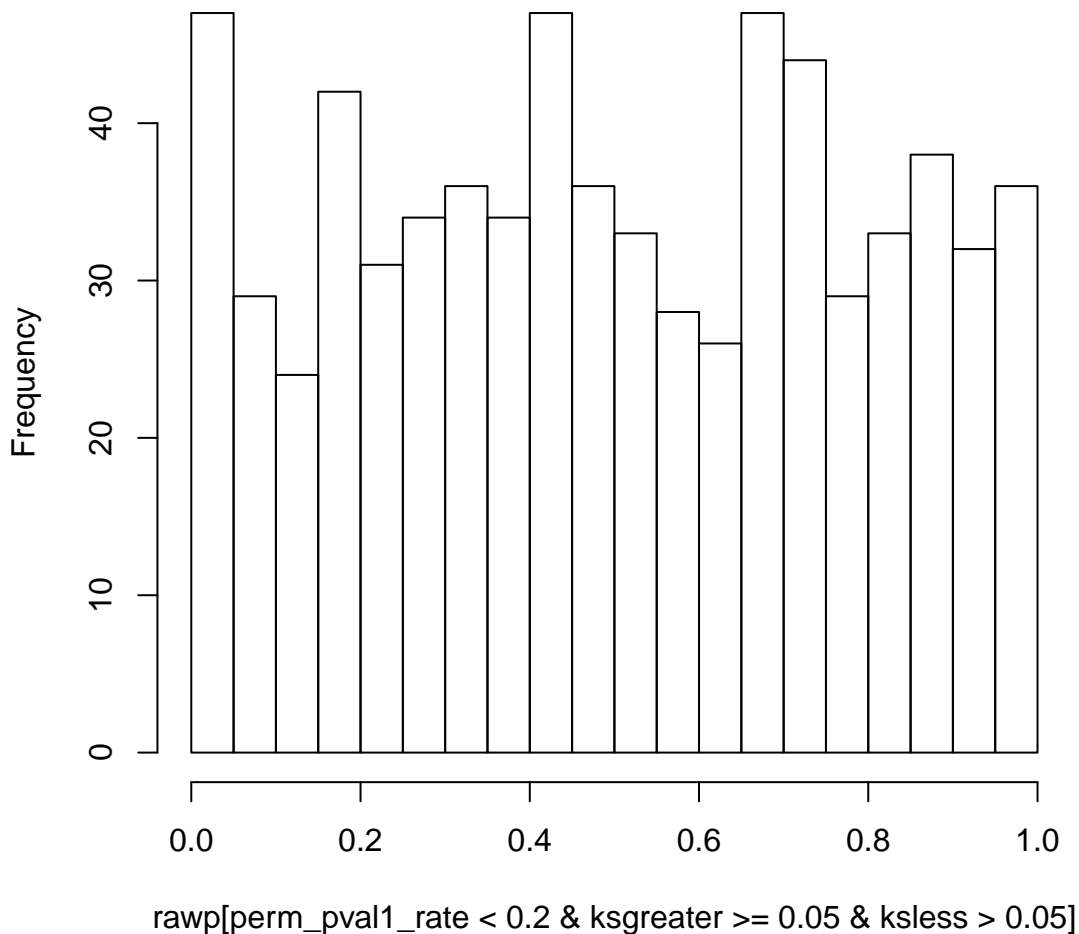
permutation pvalues with pval1_rate<0.2,ksless sig



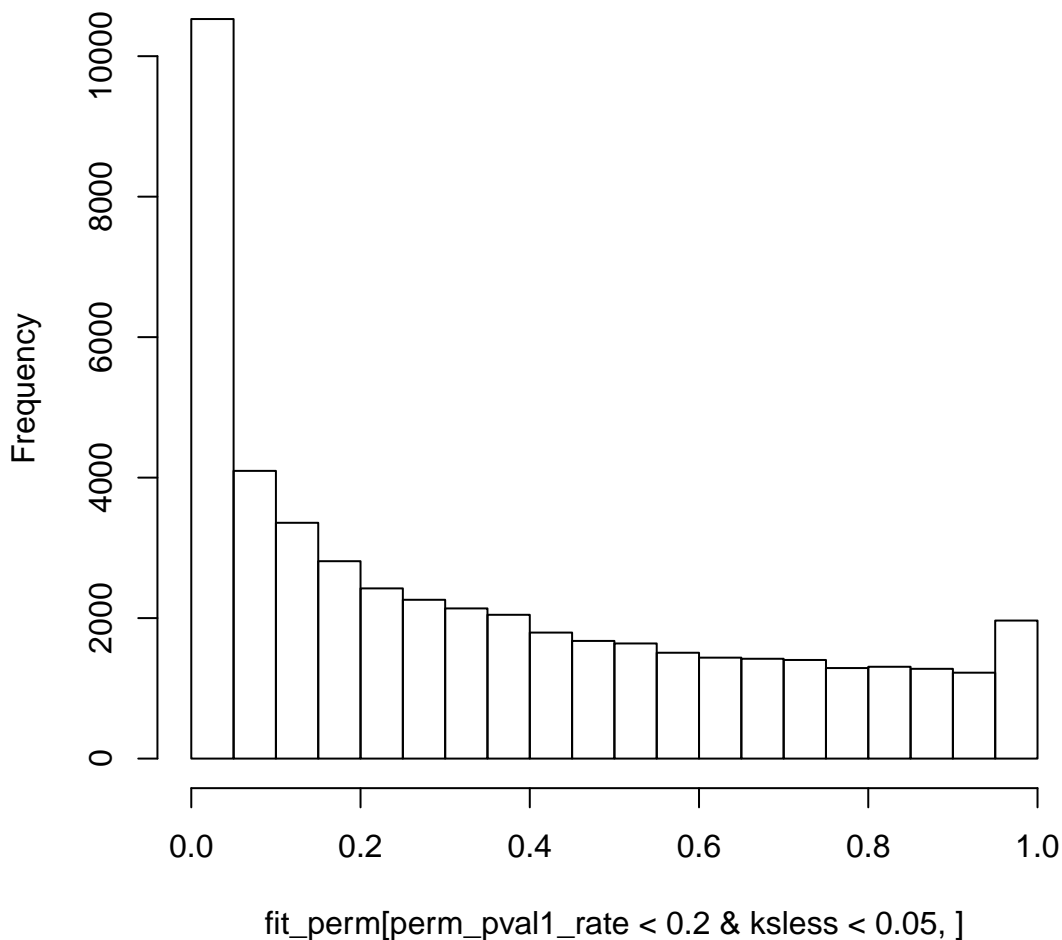
permutation pvalues with pval1_rate<0.2,ksgreater sig



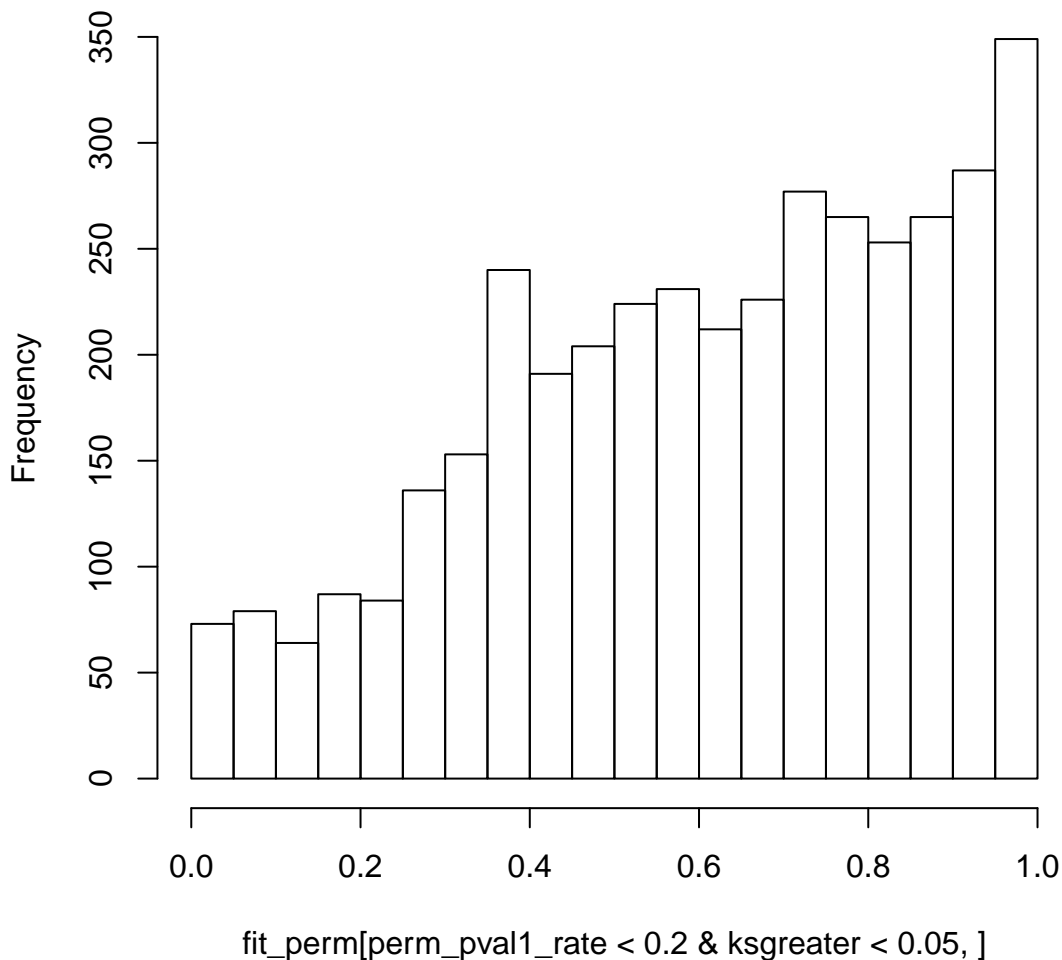
permutation pvalues with pval1_rate<0.2,ks no sig



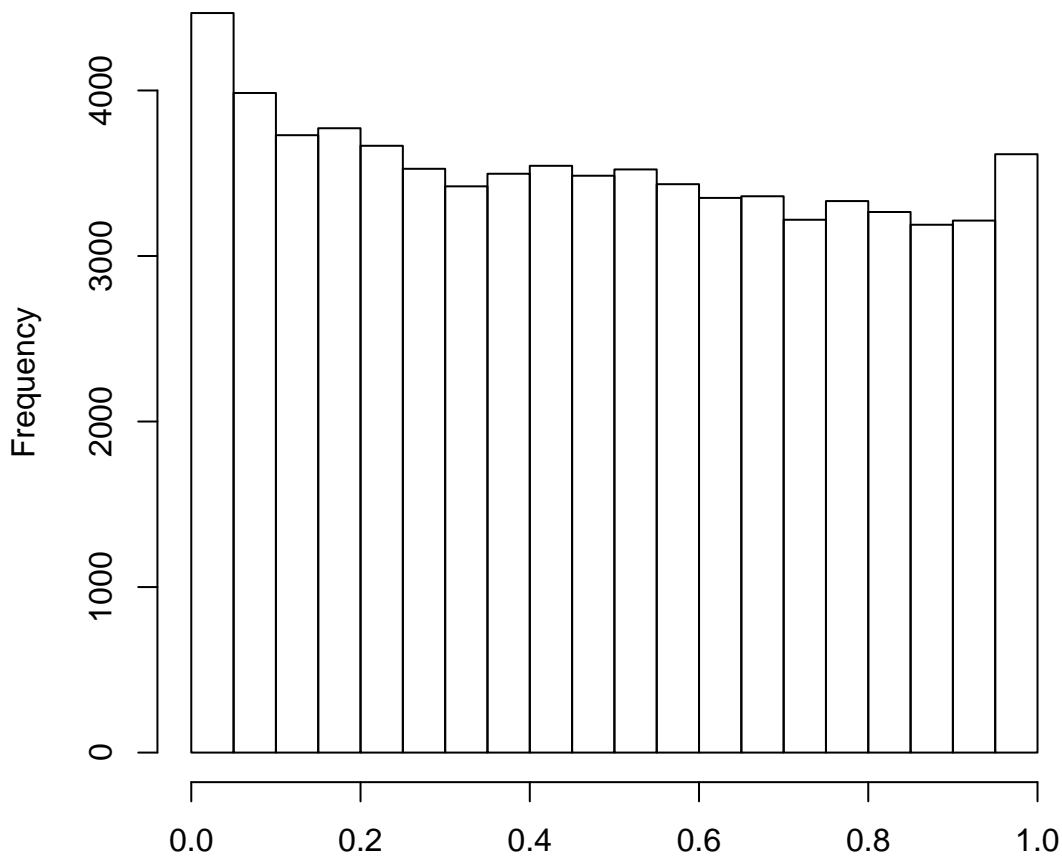
pvalues from permutation data with pval1_rate<0.2,ksless sig



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 >= 0.05 & ksless > 0.05,]`