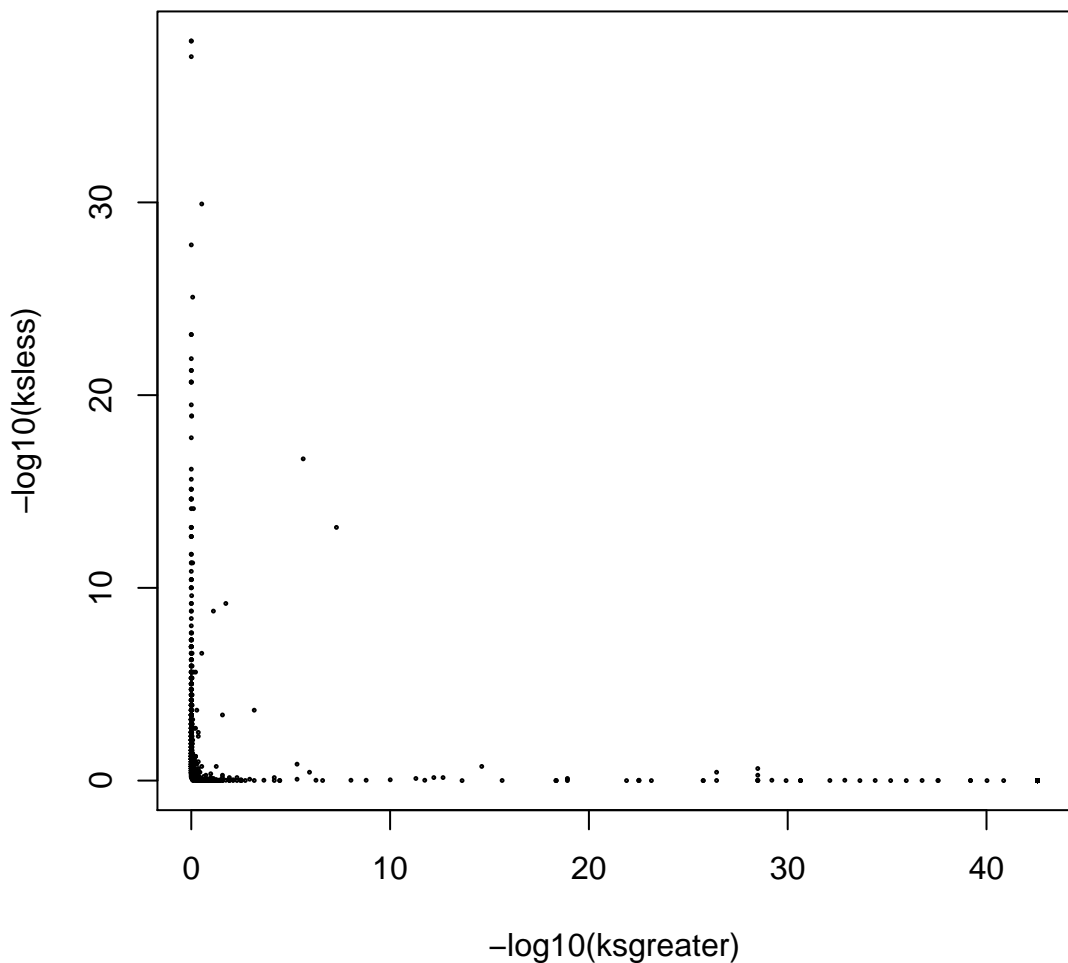
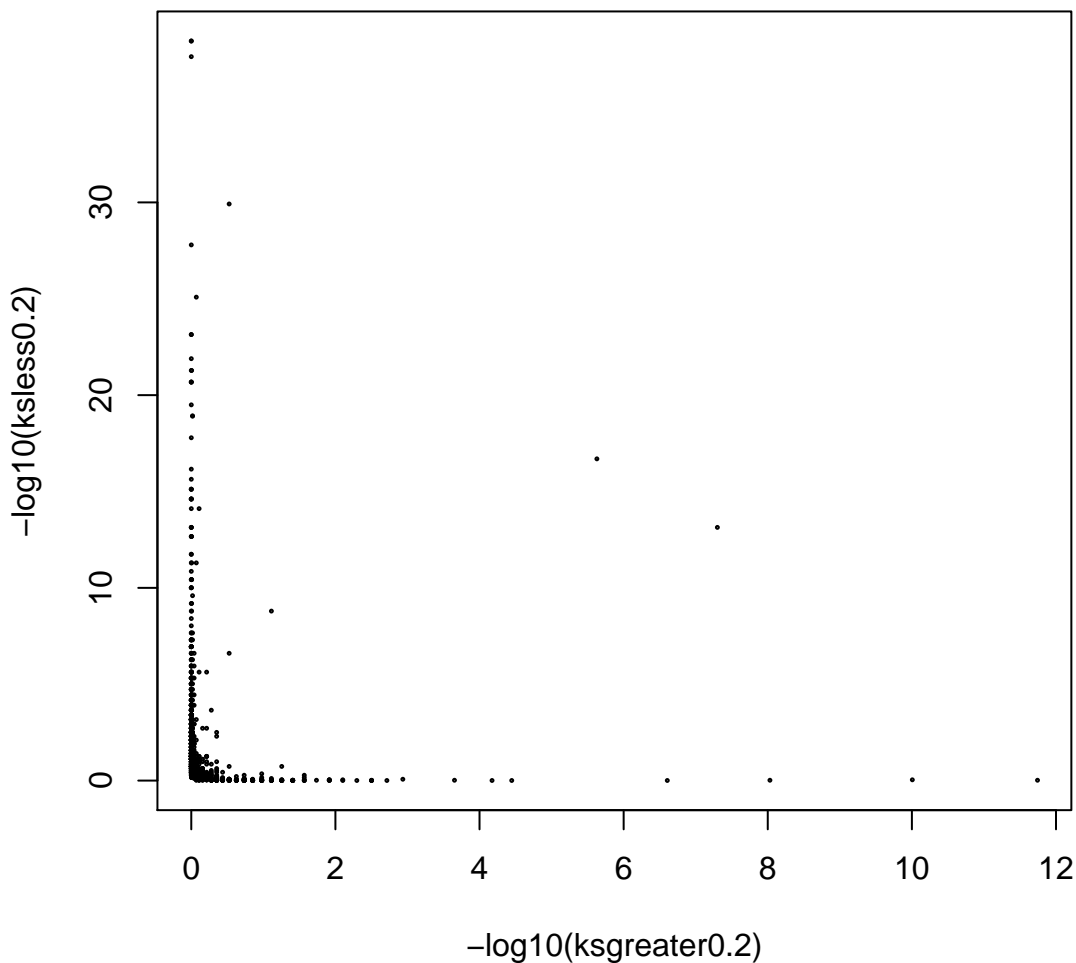


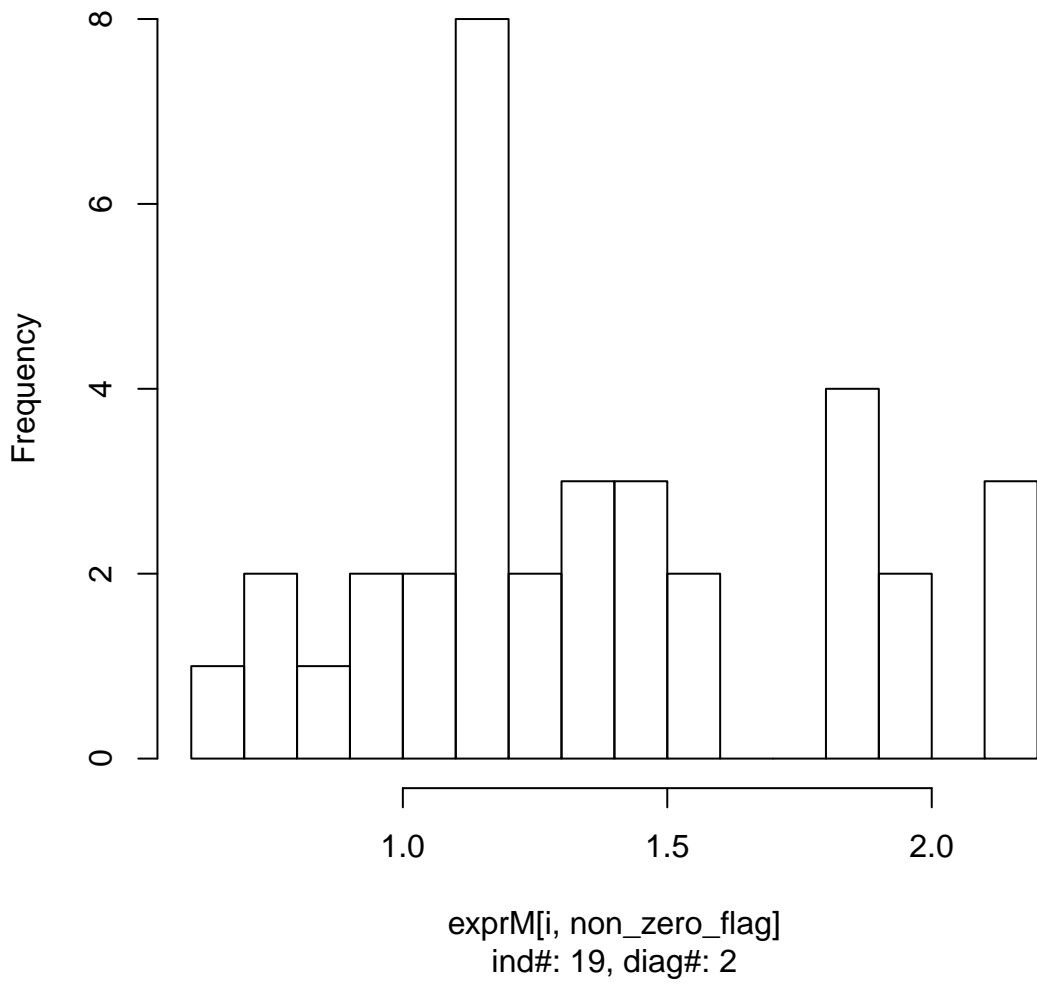
**sig\_KSgreater: 61%, sig\_KSless: 15.833%**



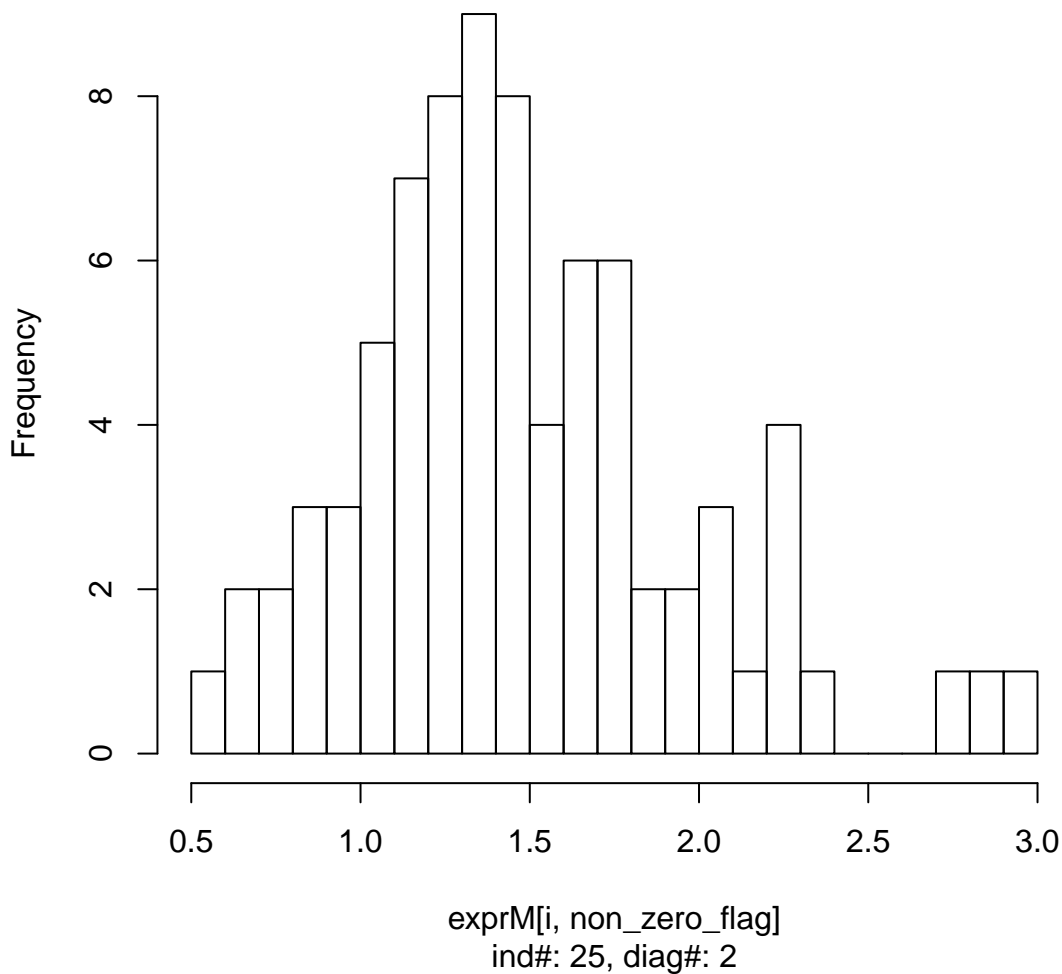
**sig\_KSgreater0.2: 3.146%, sig\_KSless0.2: 39.073%**



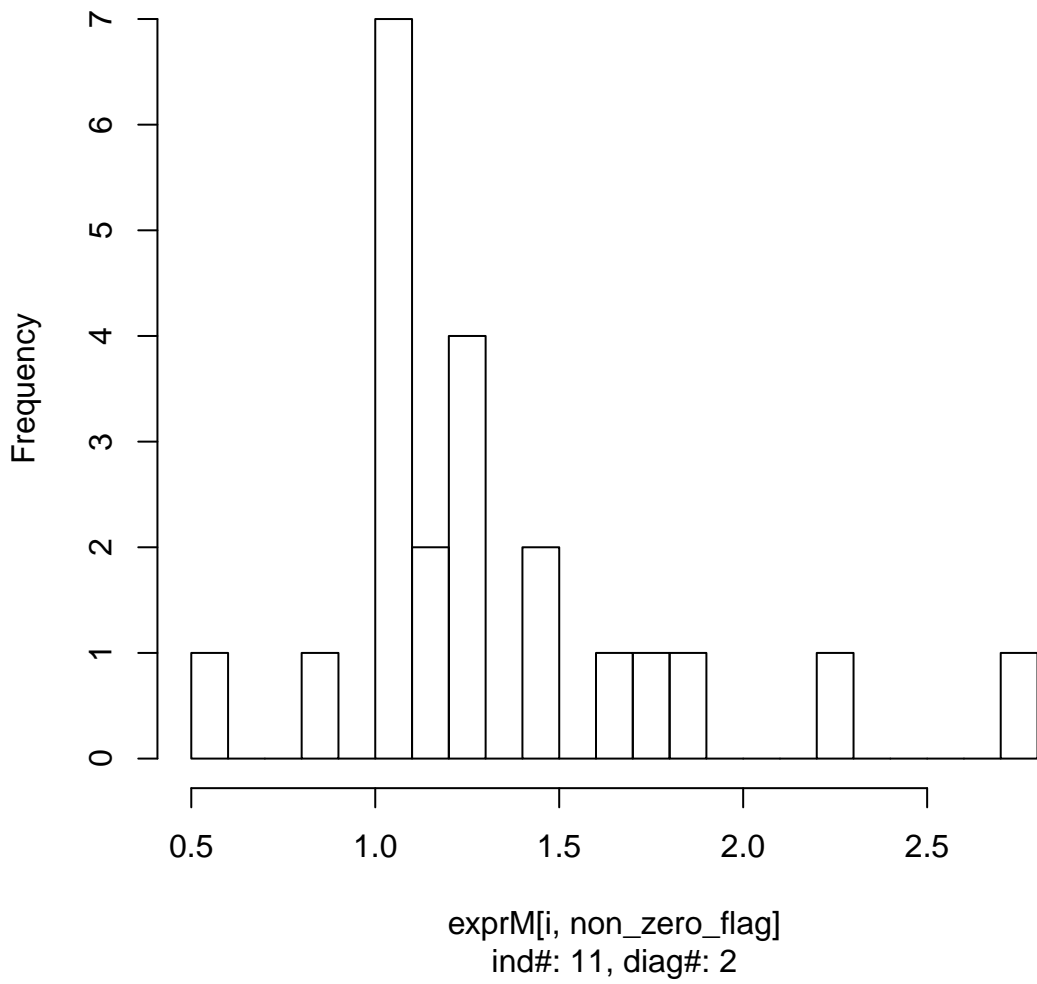
log expression of gene#56, pval ob=0.1501, non-zero num=35



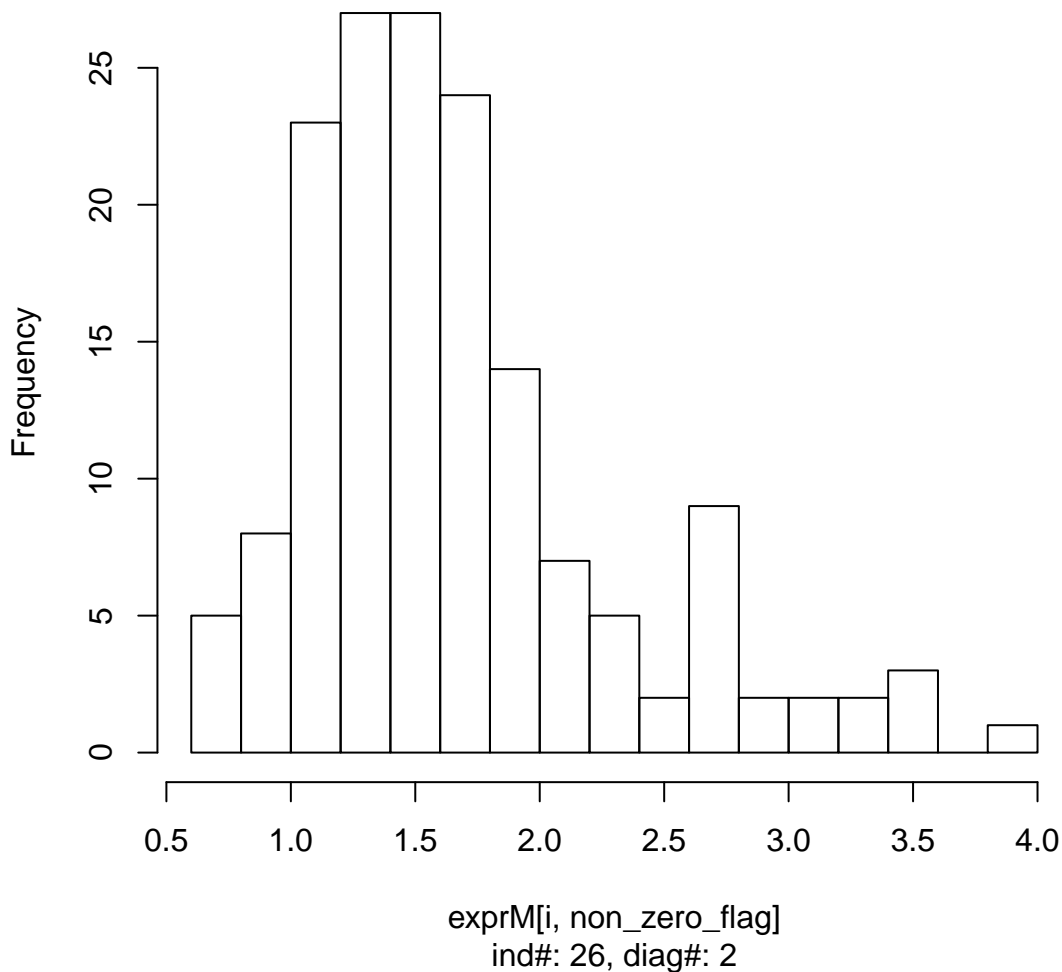
log expression of gene#65, pval ob=0.3612, non-zero num=80



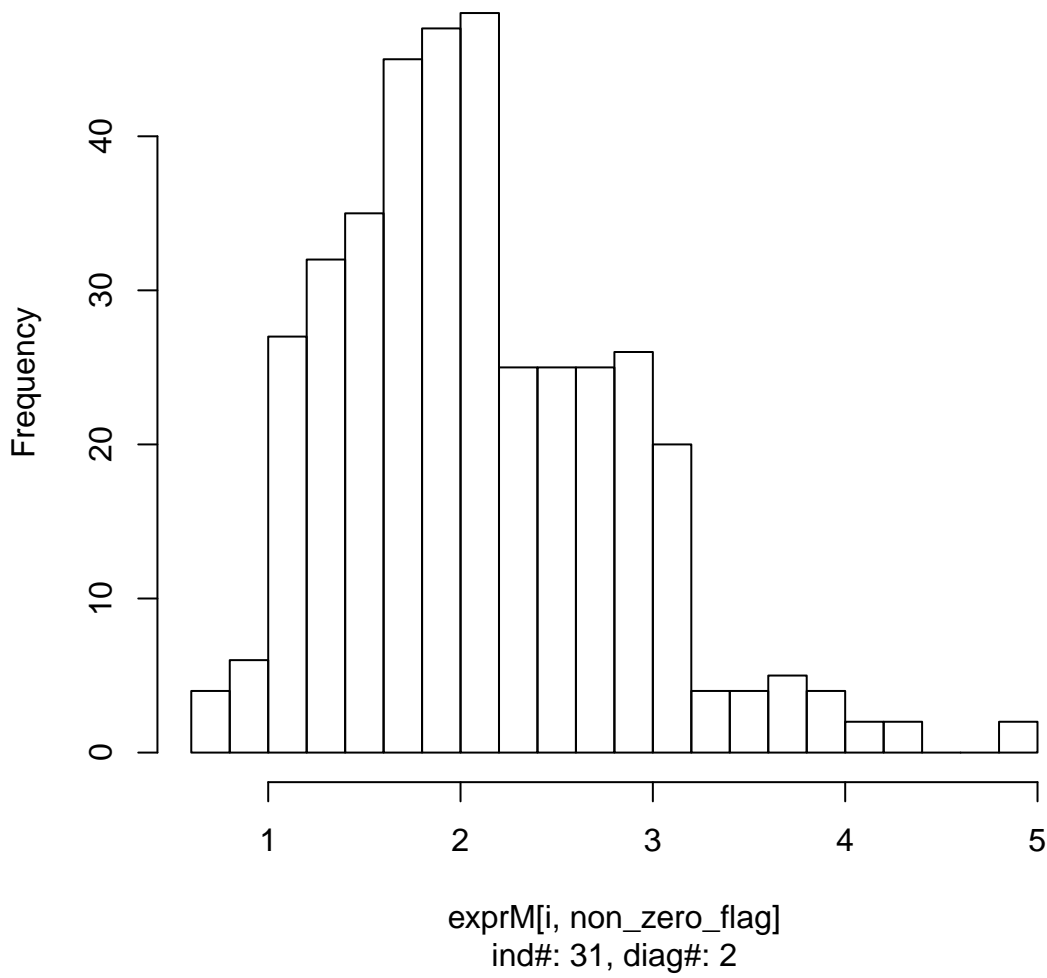
**log expression of gene#2013, pval ob=0.2196, non-zero num=2**



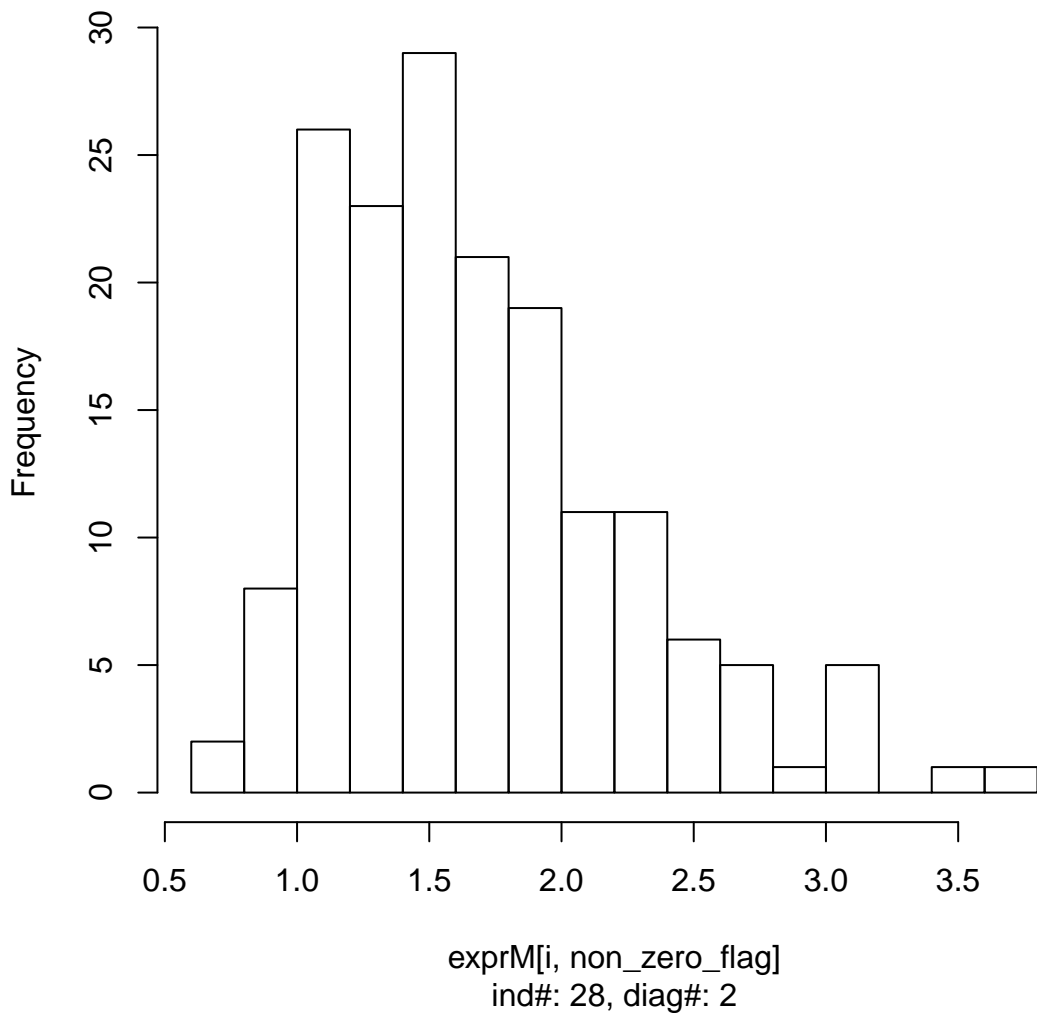
**log expression of gene#1060, pval ob=0.8118, non-zero num=1**



**log expression of gene#104, pval ob=0.717, non-zero num=38**

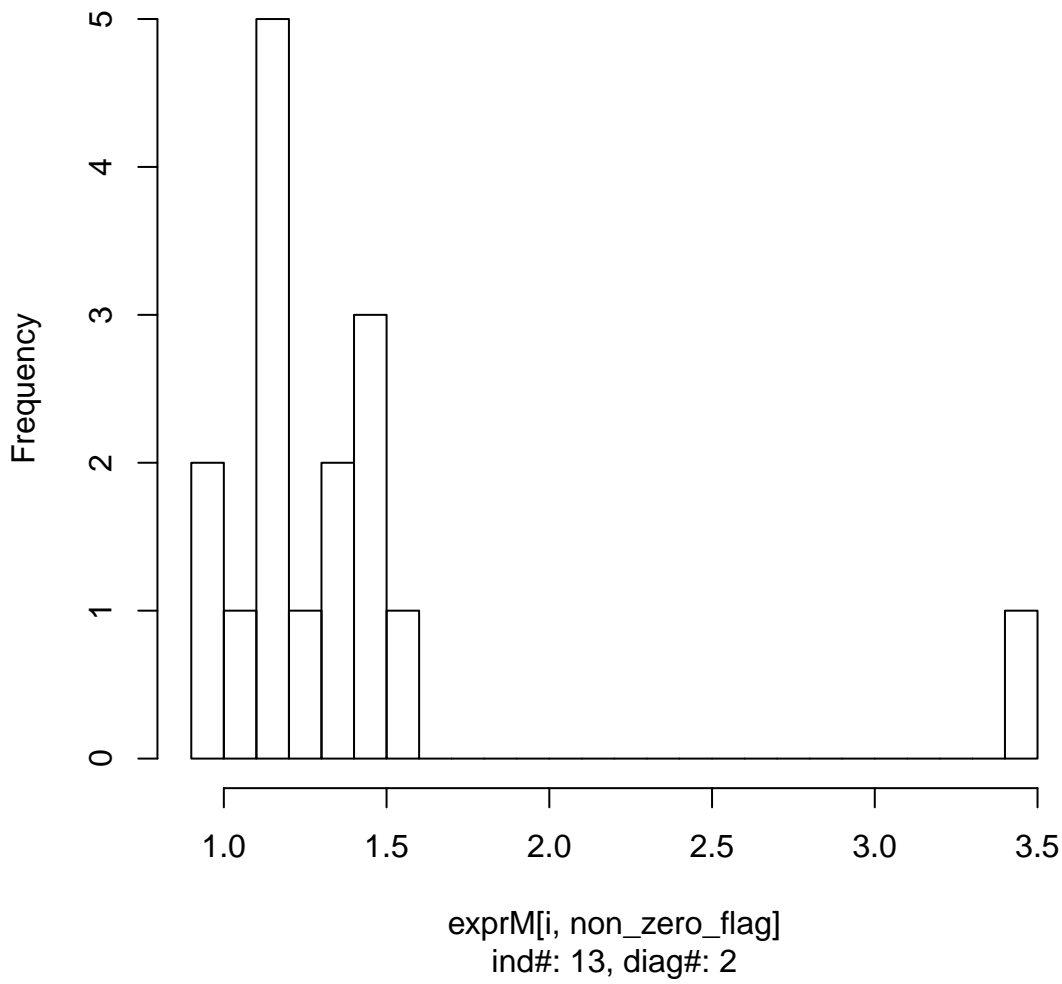


**log expression of gene#2194, pval ob=0.8559, non-zero num=1**

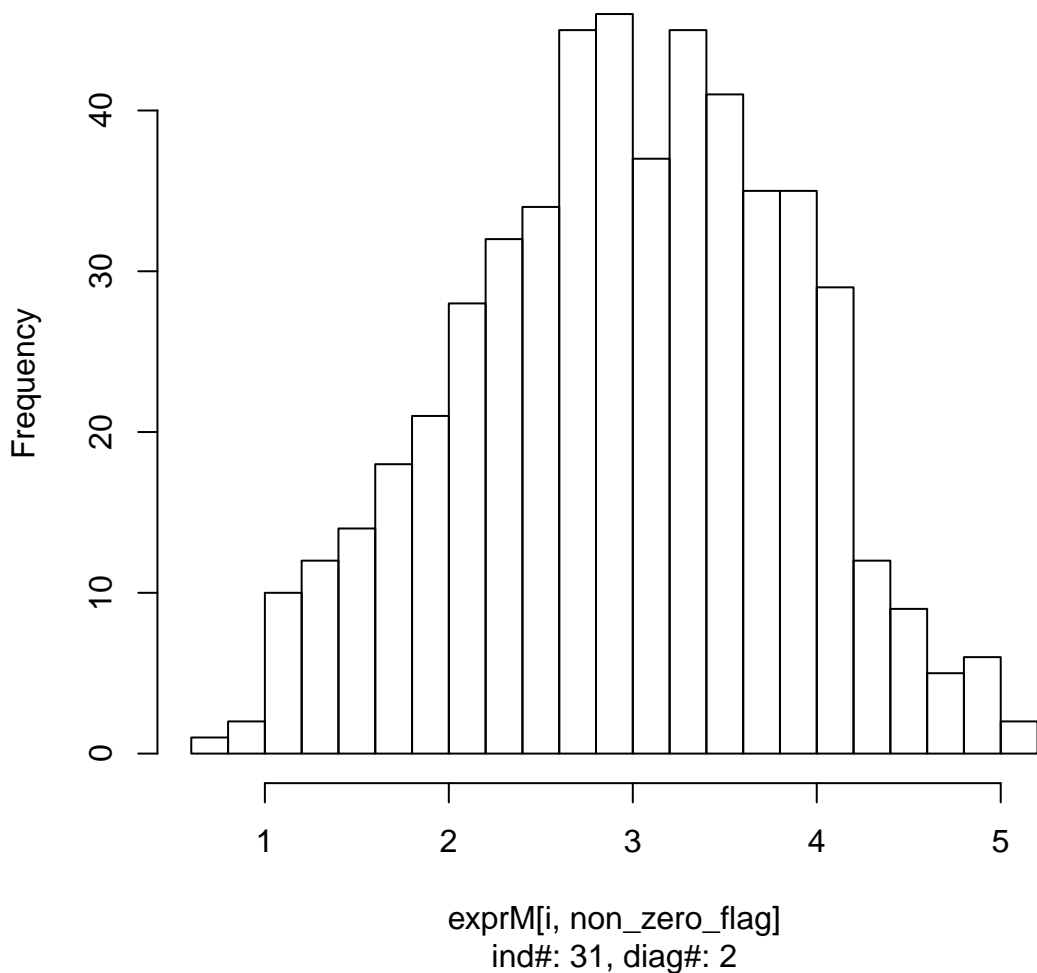




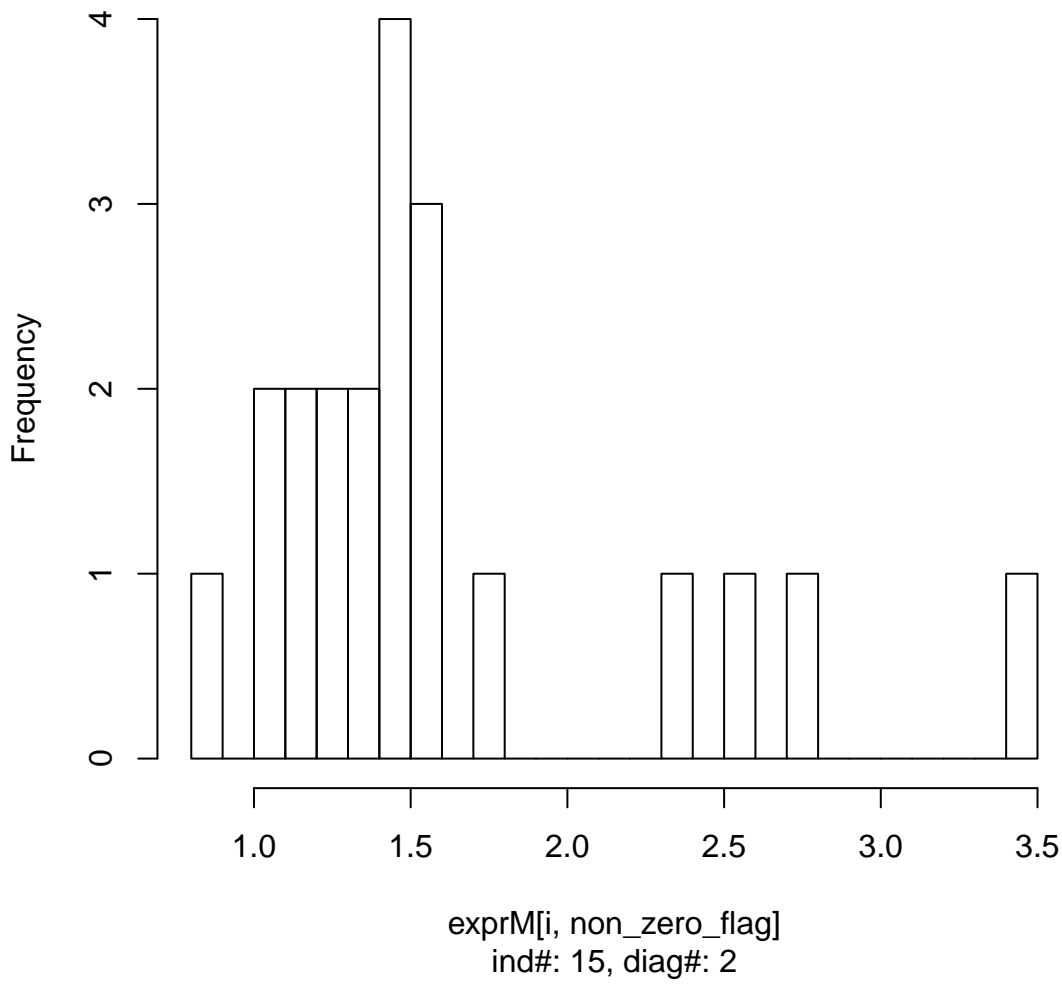
**log expression of gene#990, pval ob=0.0735, non-zero num=1**



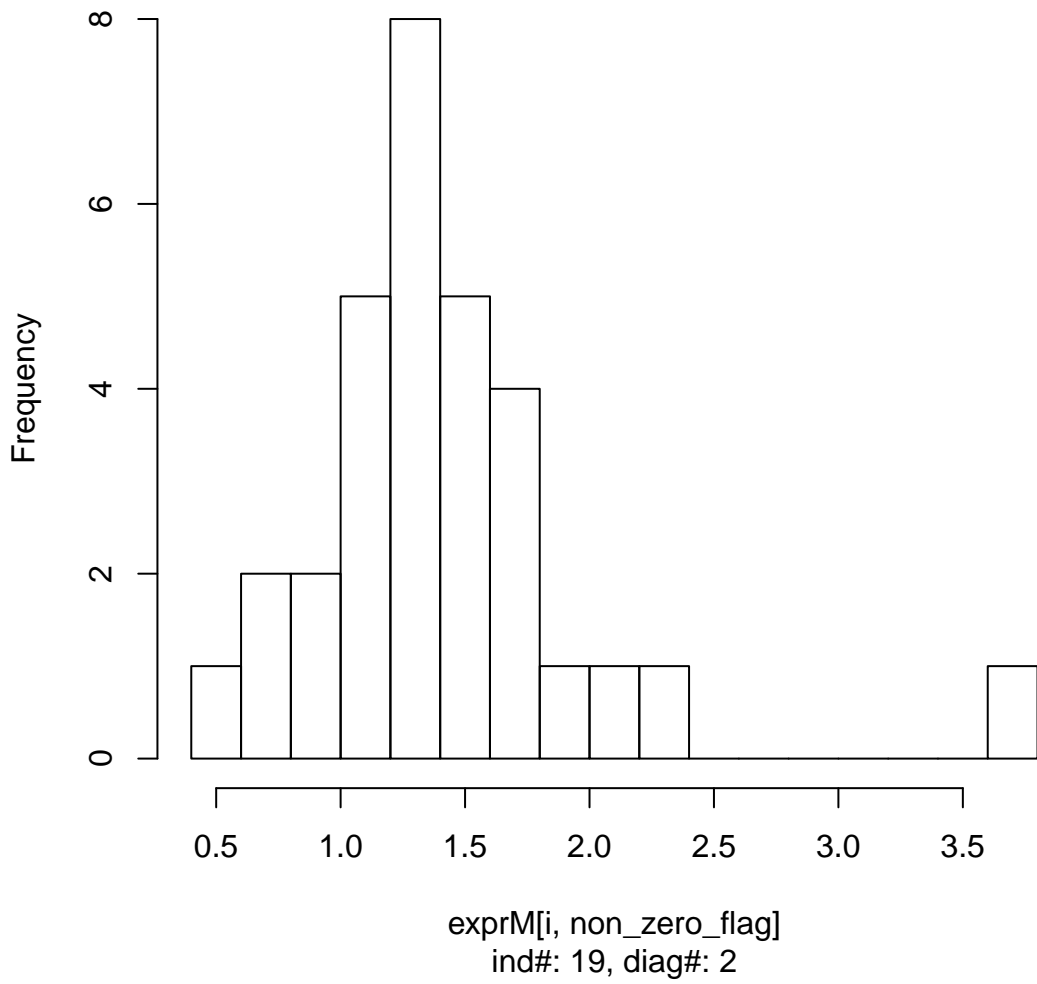
**log expression of gene#1050, pval ob=0.0816, non-zero num=5**



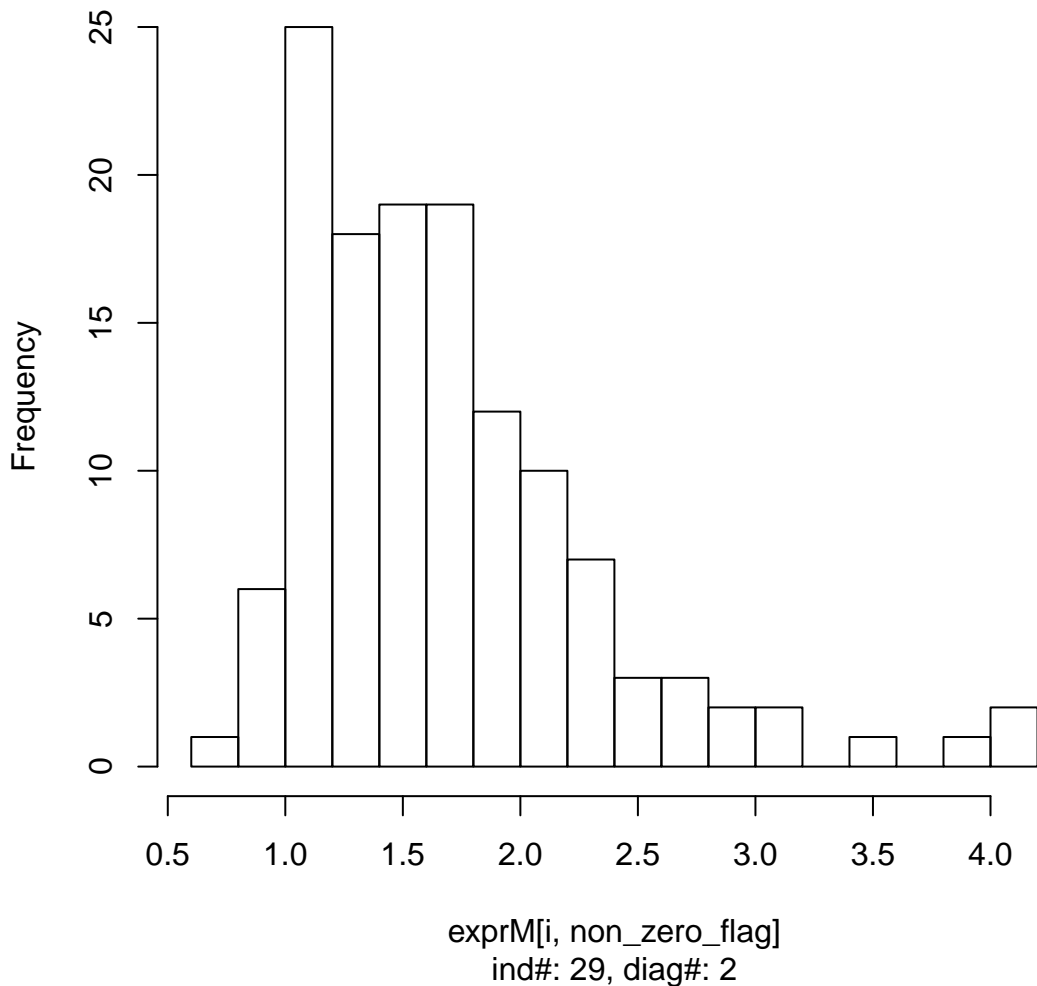
**log expression of gene#1979, pval ob=0.0141, non-zero num=2**



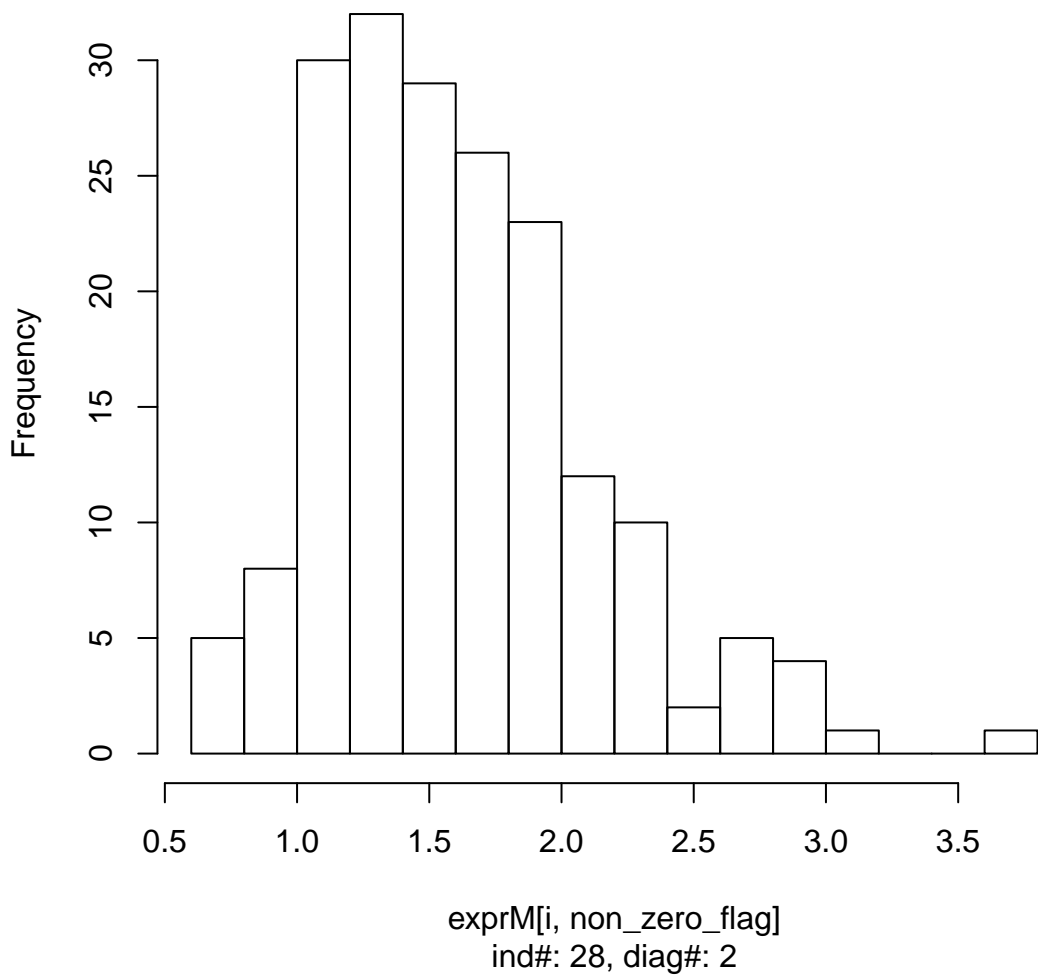
**log expression of gene#1202, pval ob=0.8022, non-zero num=3**



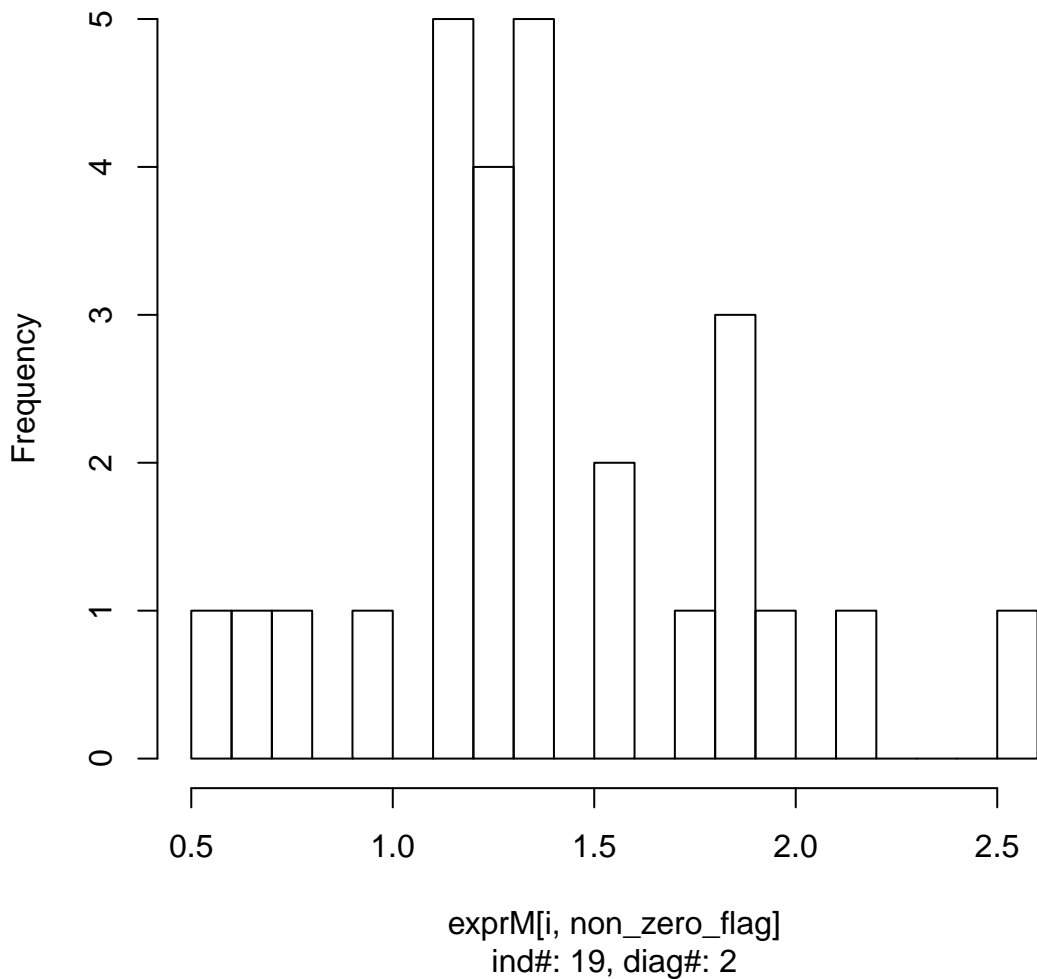
**log expression of gene#861, pval ob=0.518, non-zero num=13**



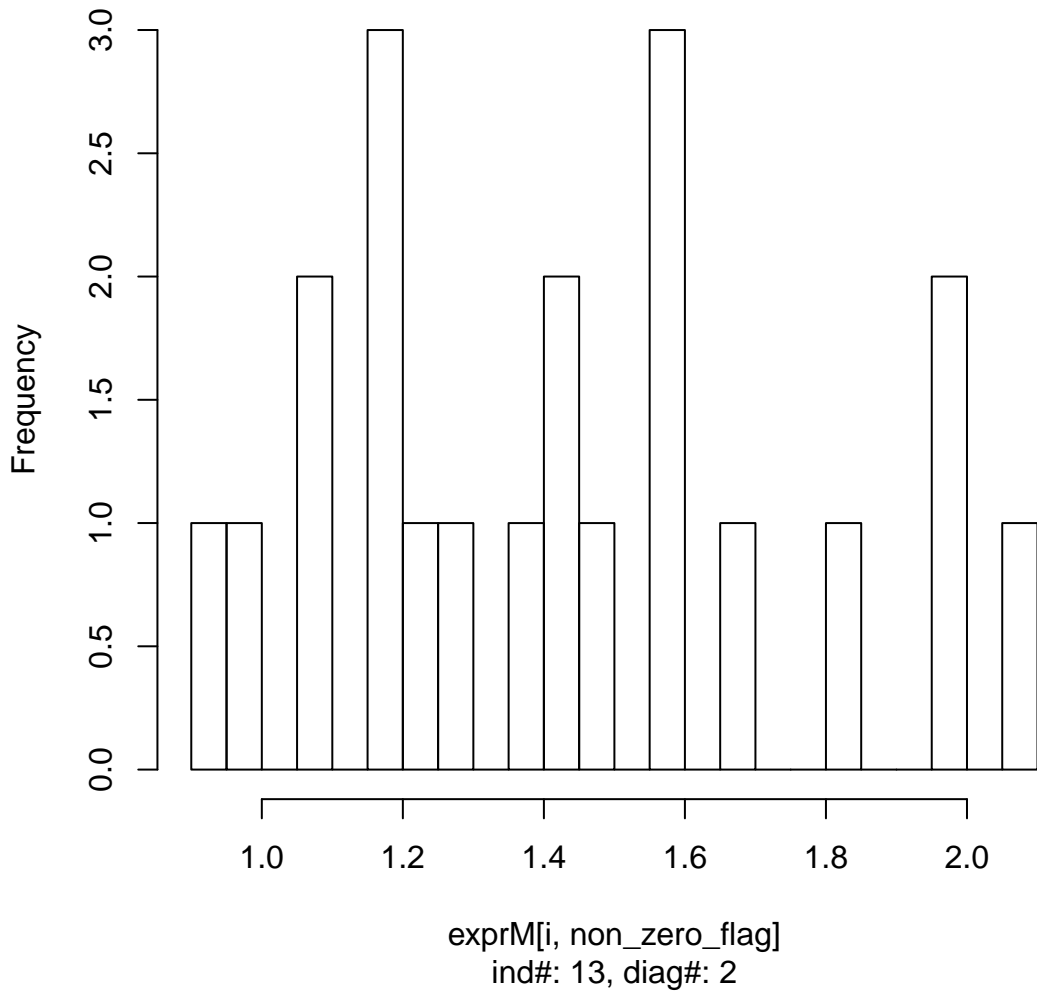
**log expression of gene#103, pval ob=0.0086, non-zero num=18**



log expression of gene#1187, pval ob=0.5974, non-zero num=2

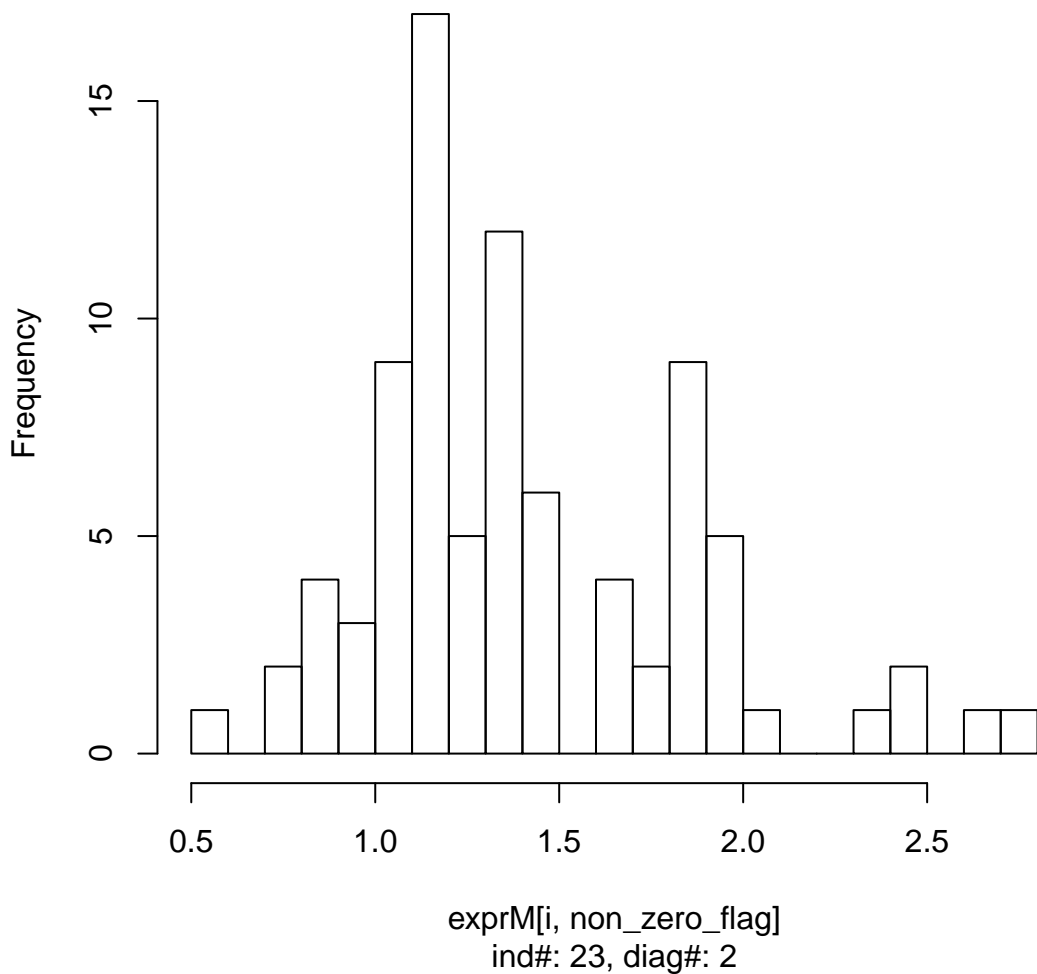


**log expression of gene#750, pval ob=0.5956, non-zero num=2**

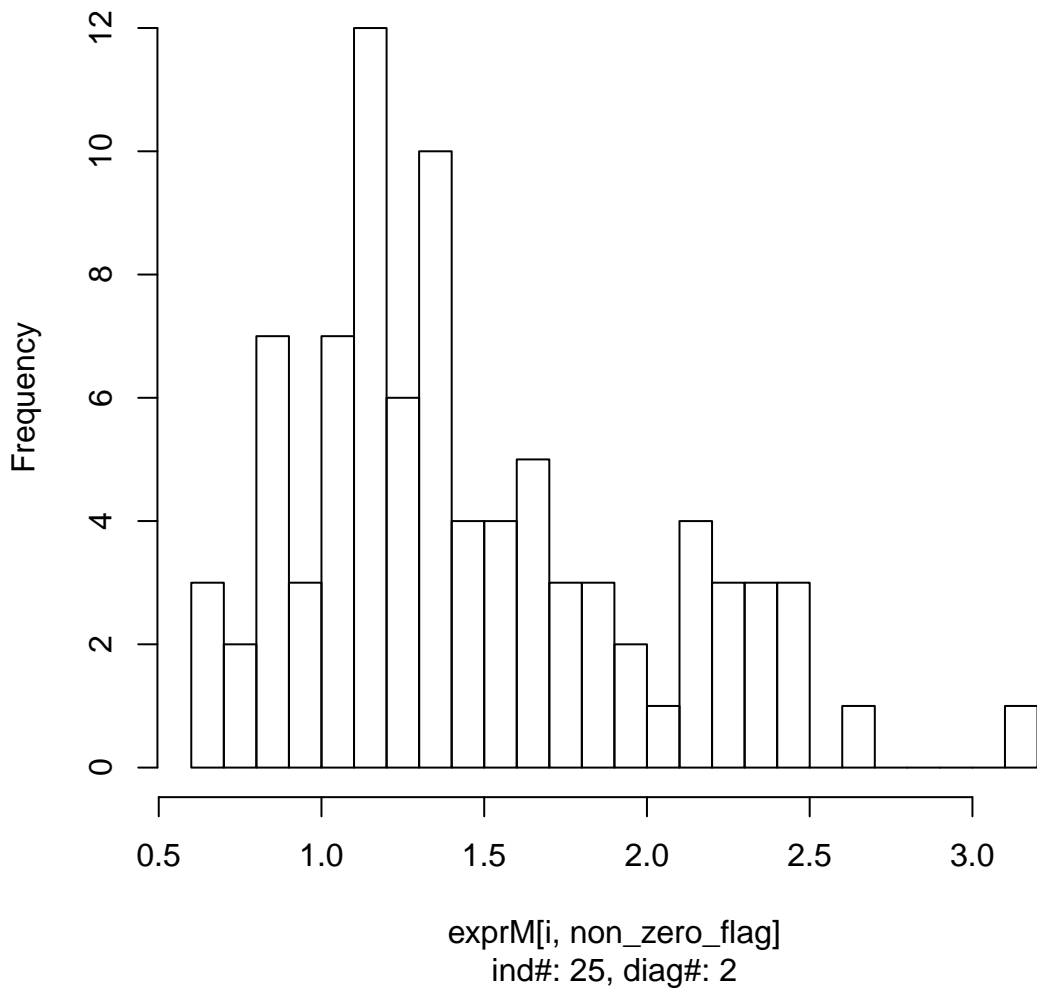




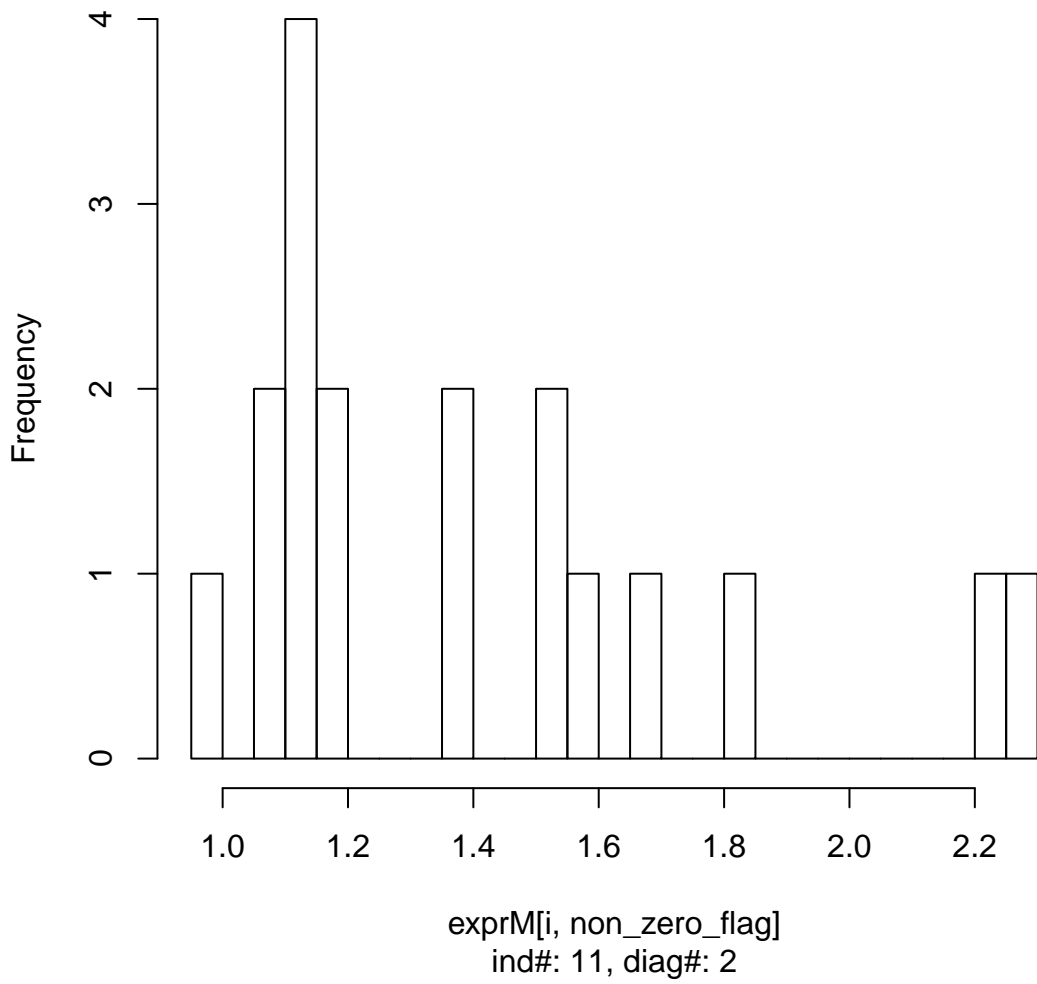
**log expression of gene#1234, pval ob=0.7065, non-zero num=8**



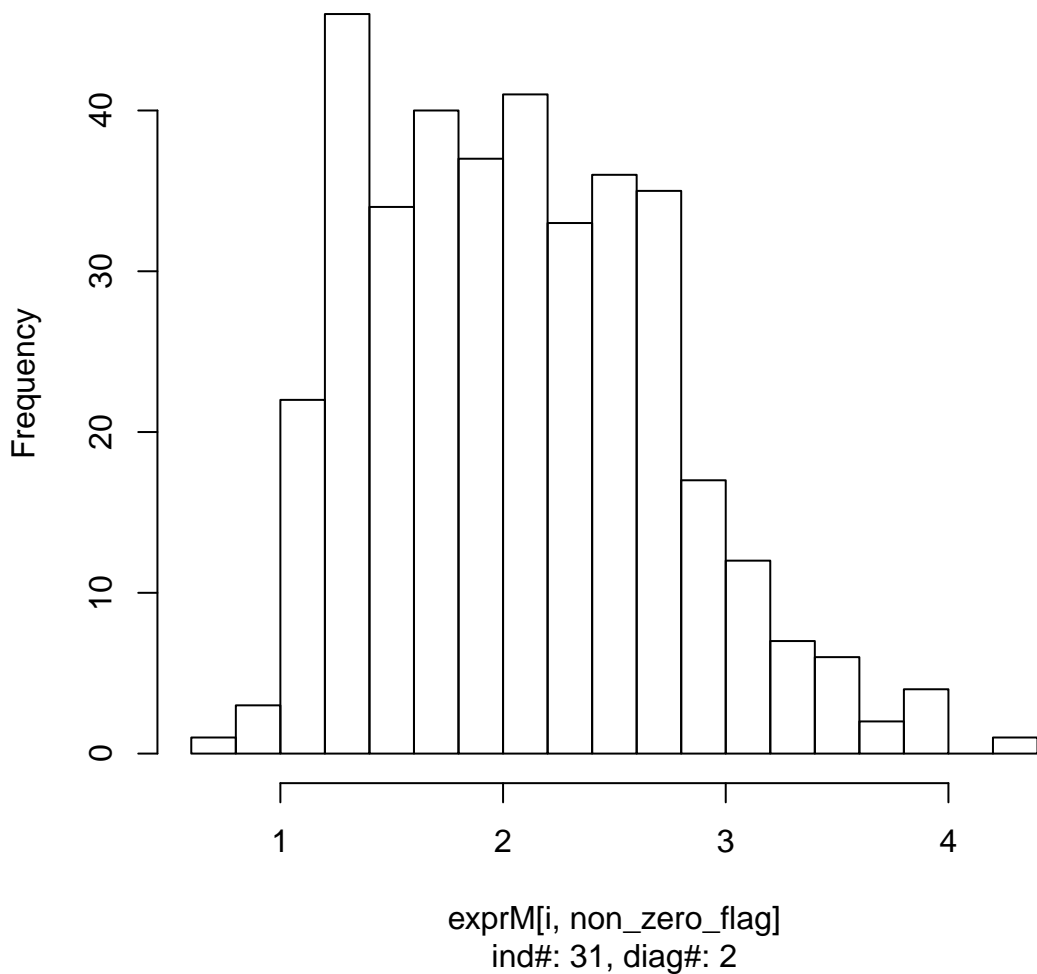
**log expression of gene#694, pval ob=0.2864, non-zero num=8**



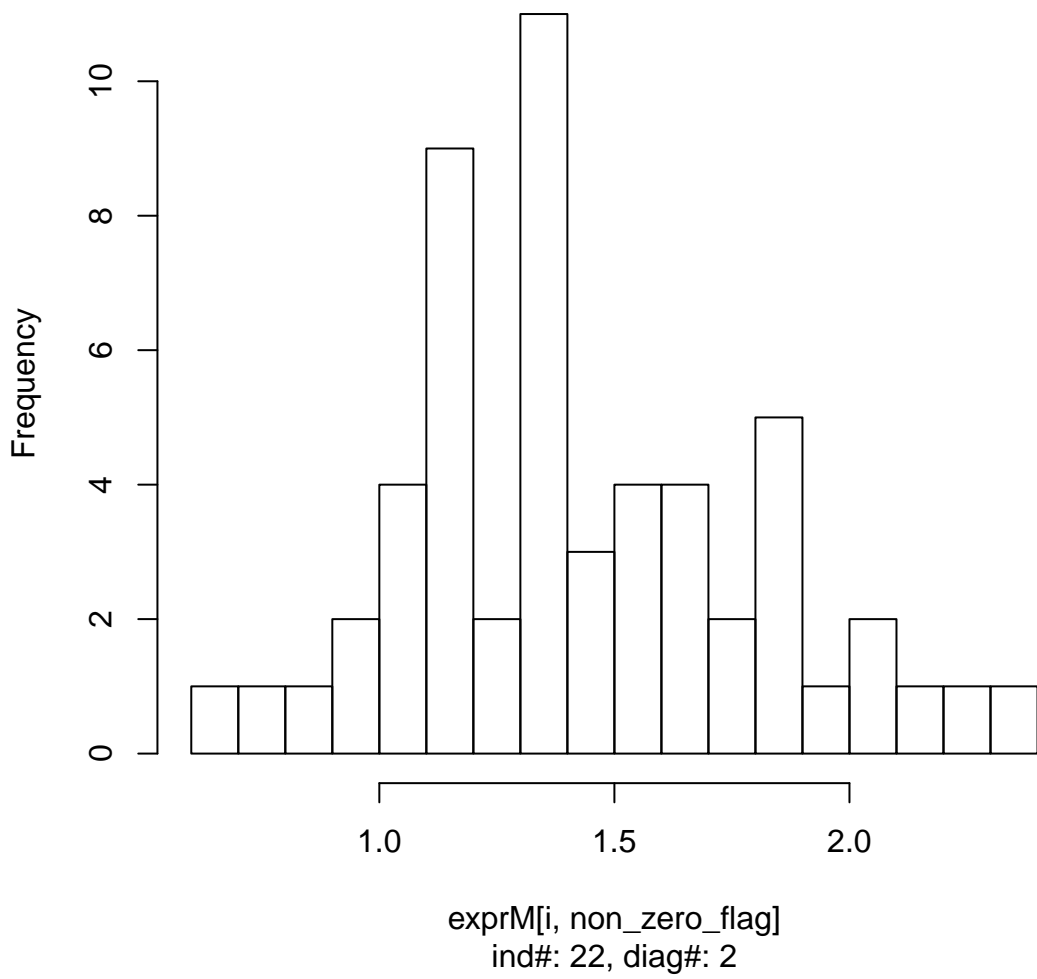
log expression of gene#2565, pval ob=0.6127, non-zero num=1



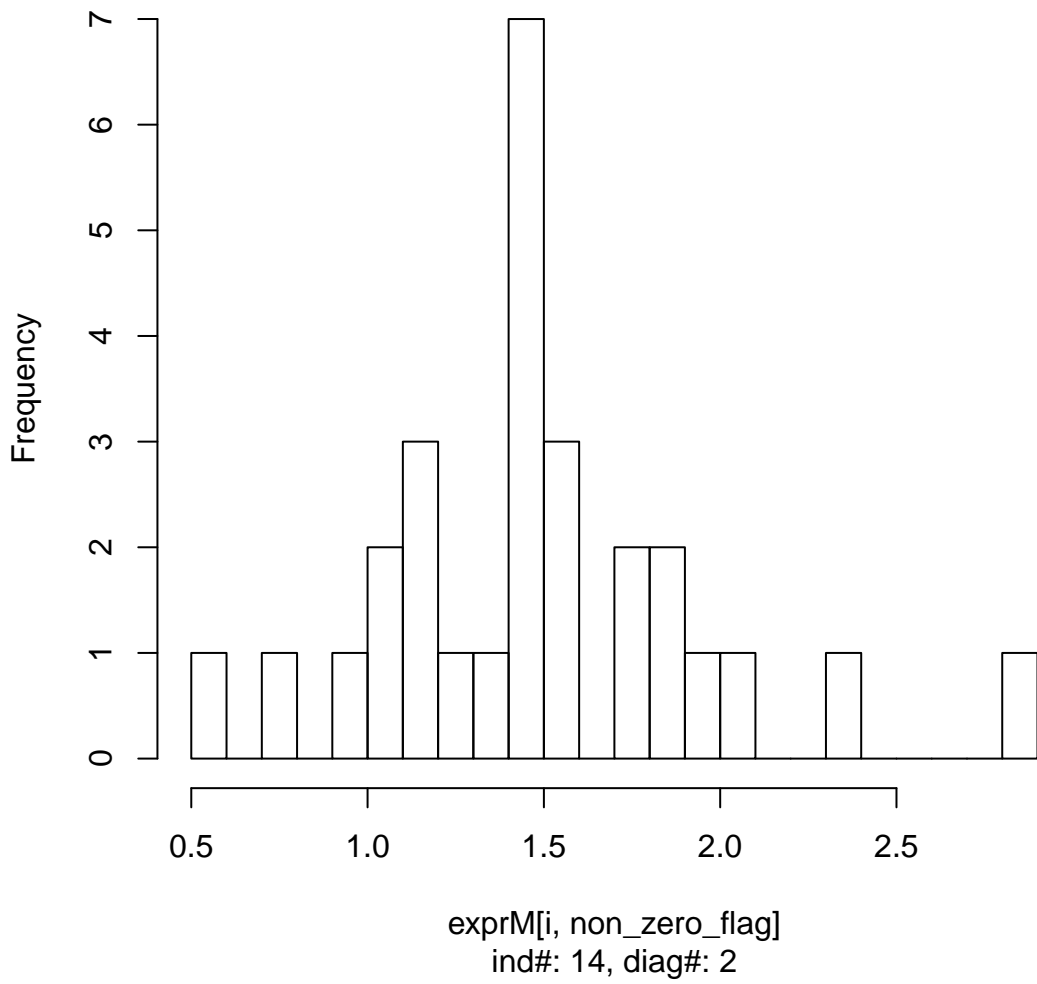
**log expression of gene#1115, pval ob=0.3019, non-zero num=3**



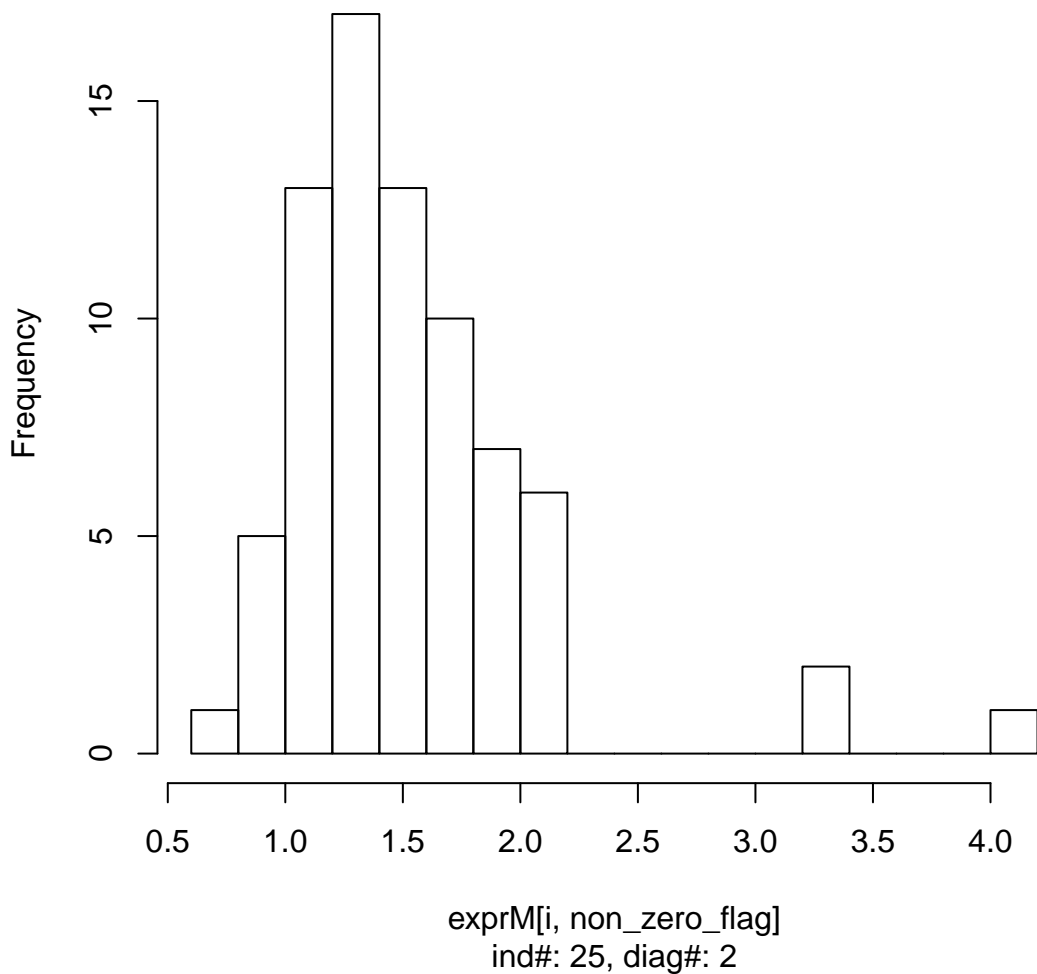
**log expression of gene#1, pval ob=0.8033, non-zero num=55**



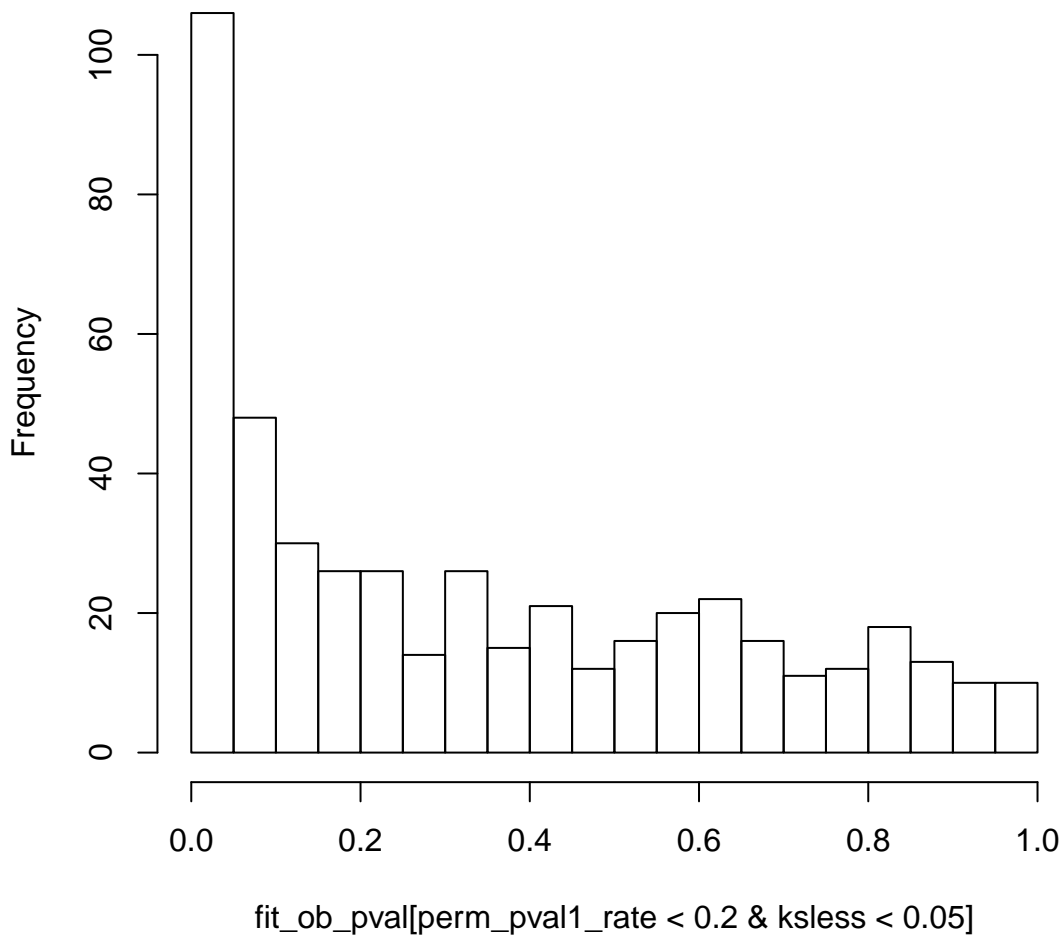
log expression of gene#1892, pval ob=0.5082, non-zero num=2



**log expression of gene#2587, pval ob=0.5541, non-zero num=7**

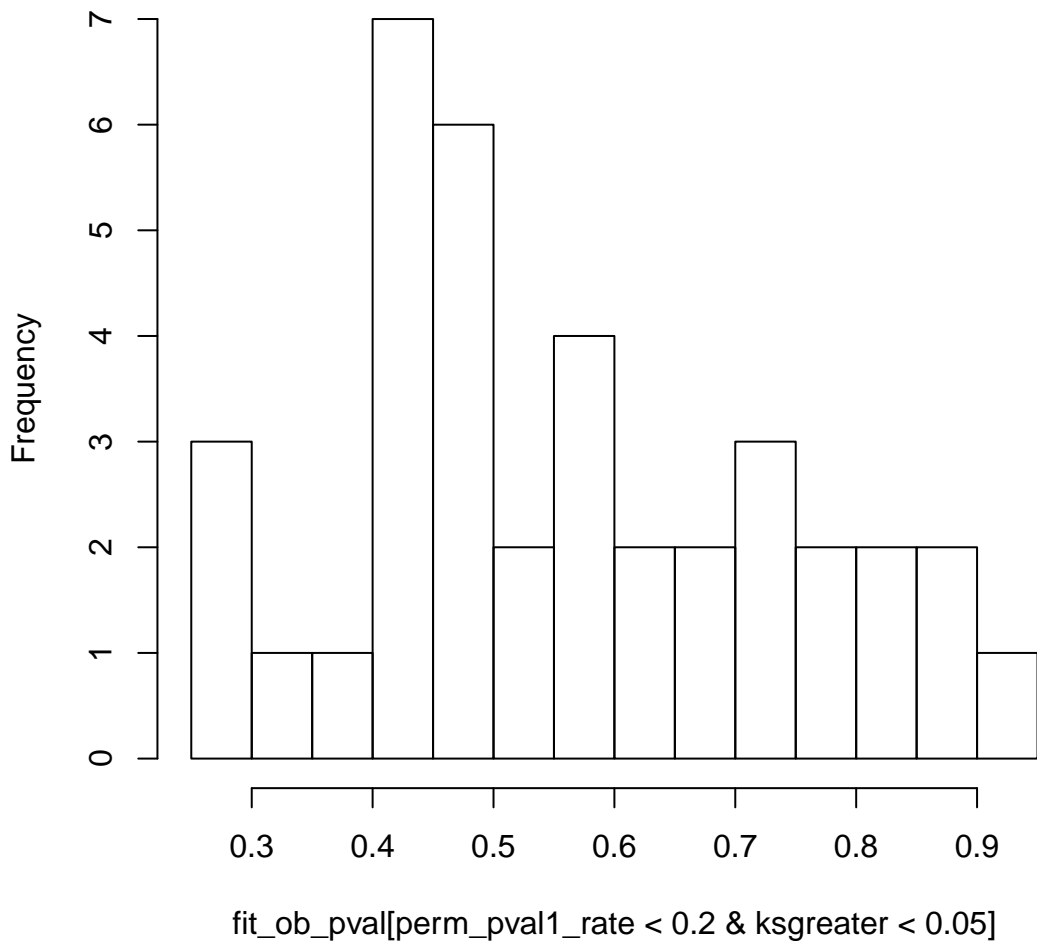


# observed pvalues with pval1\_rate<0.2,ksless sig

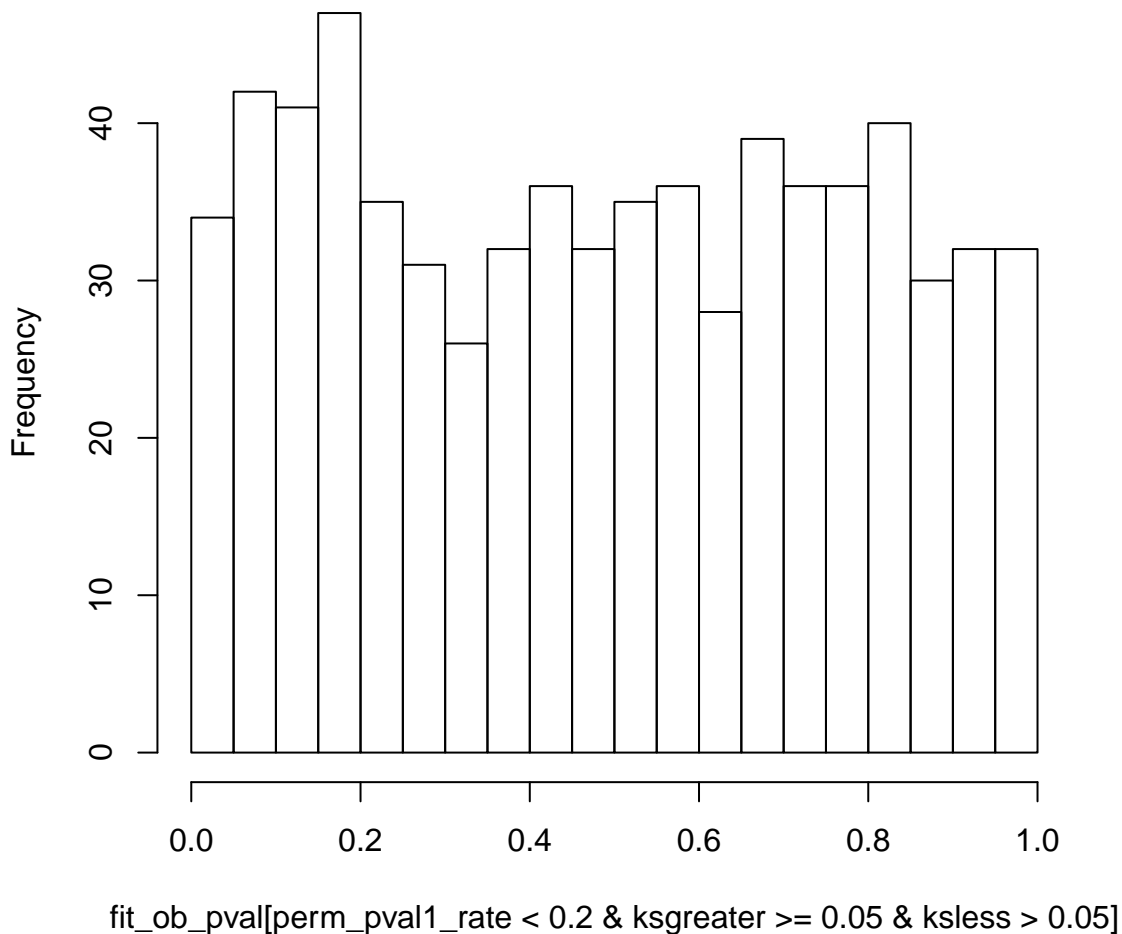




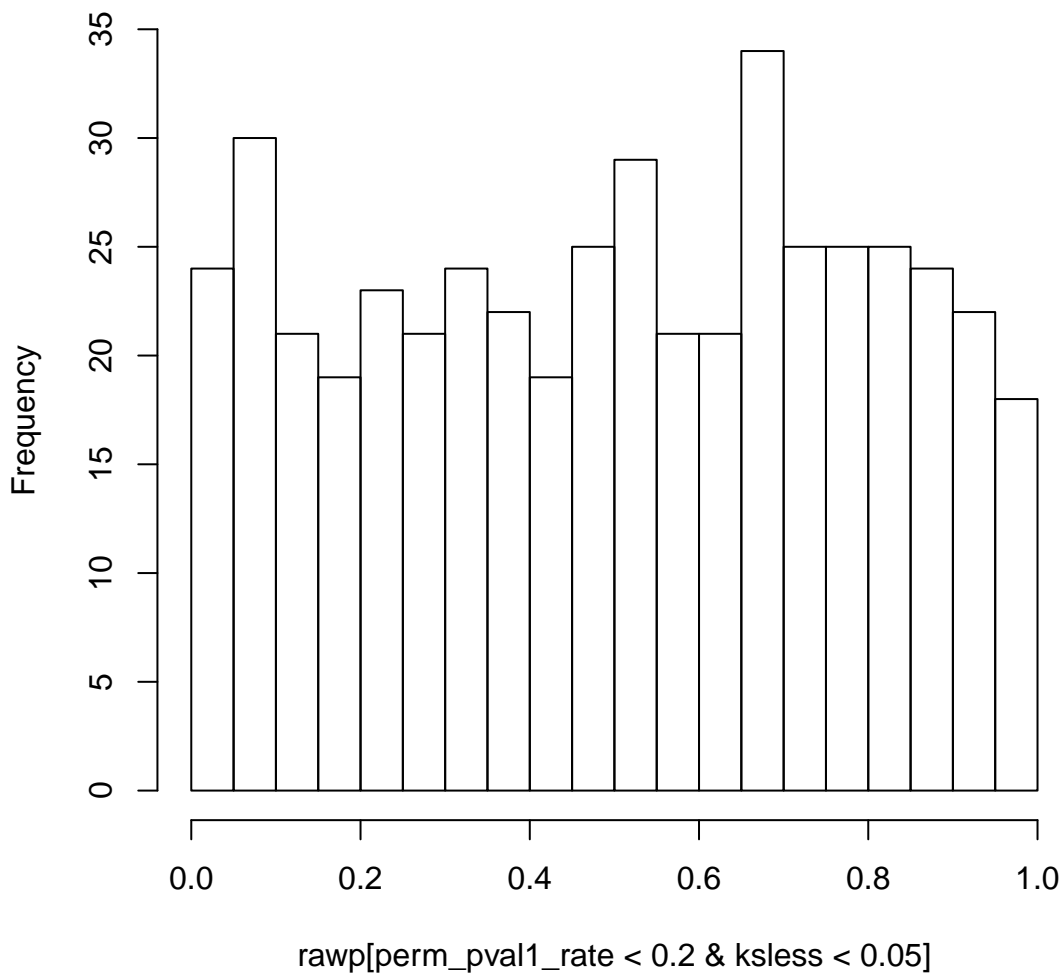
# observed pvalues with pval1\_rate<0.2,ksgreater sig



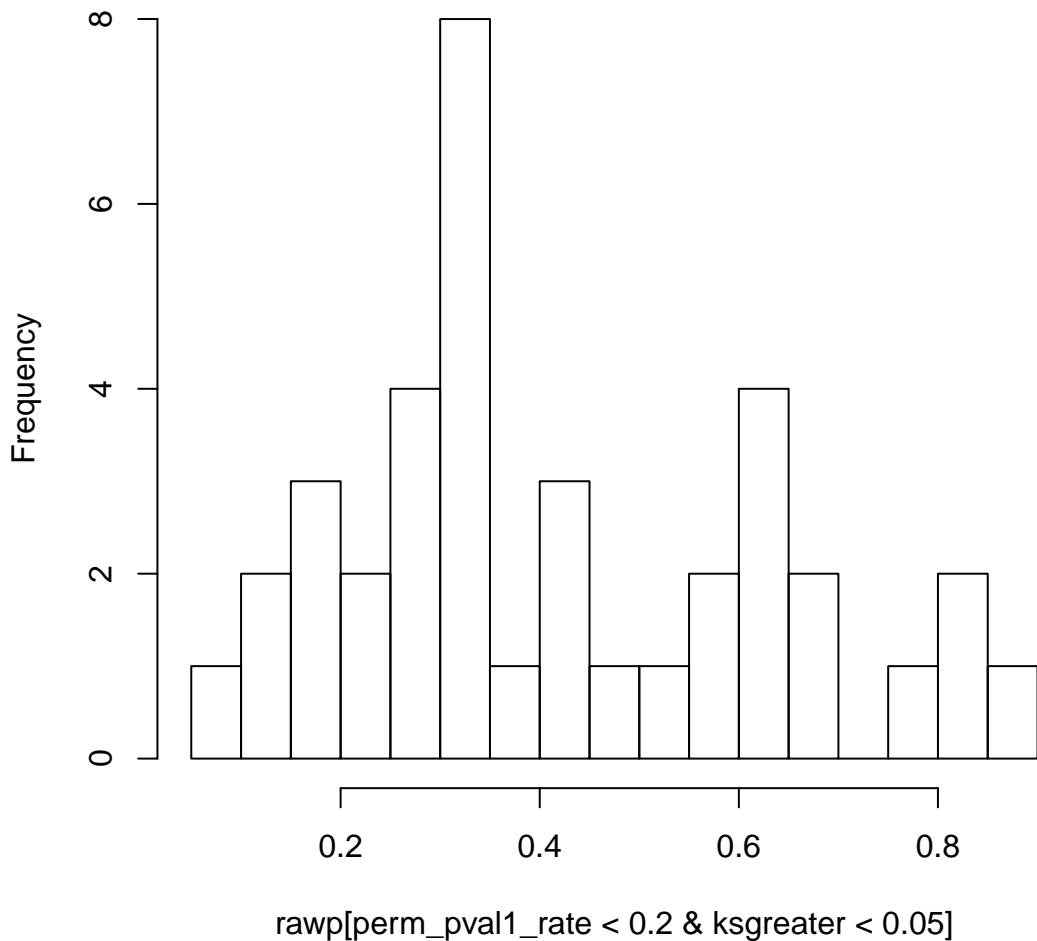
# observed pvalues with pval1\_rate<0.2,ks no sig



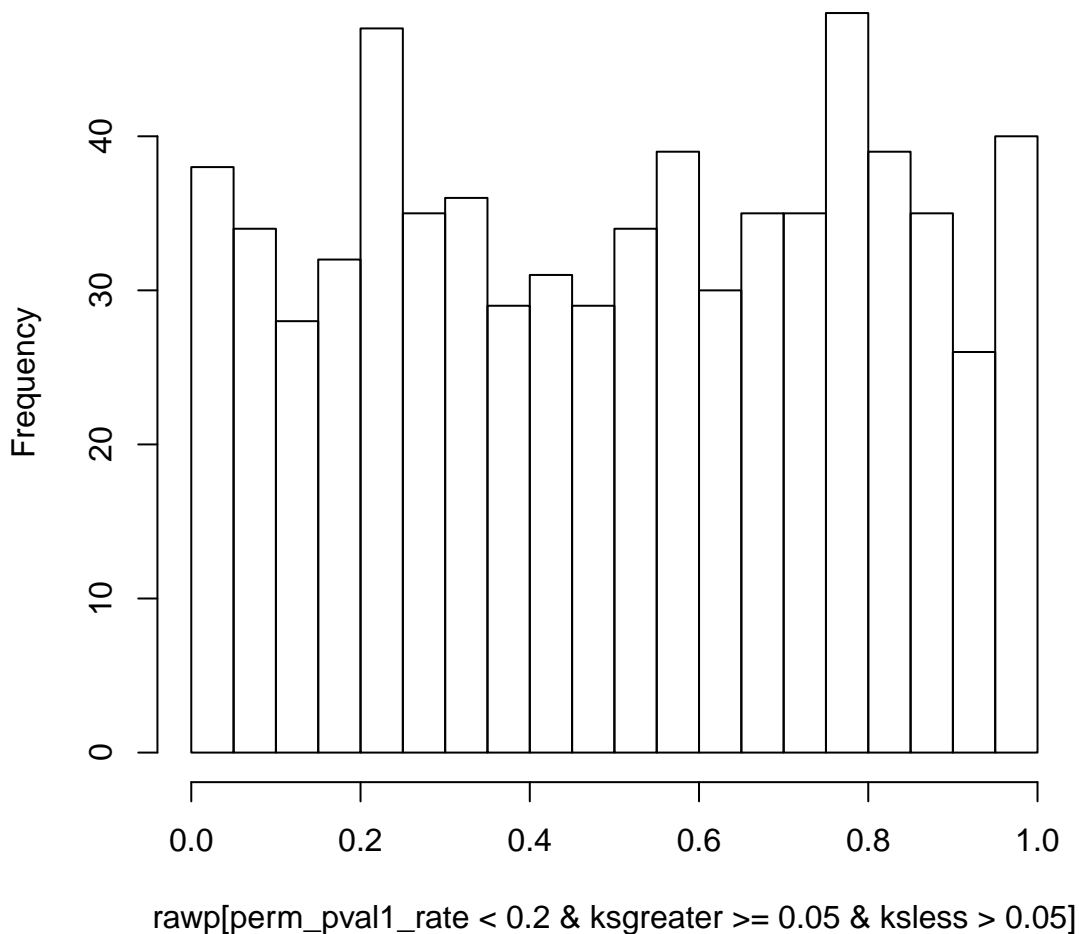
# permutation pvalues with pval1\_rate<0.2,ksless sig



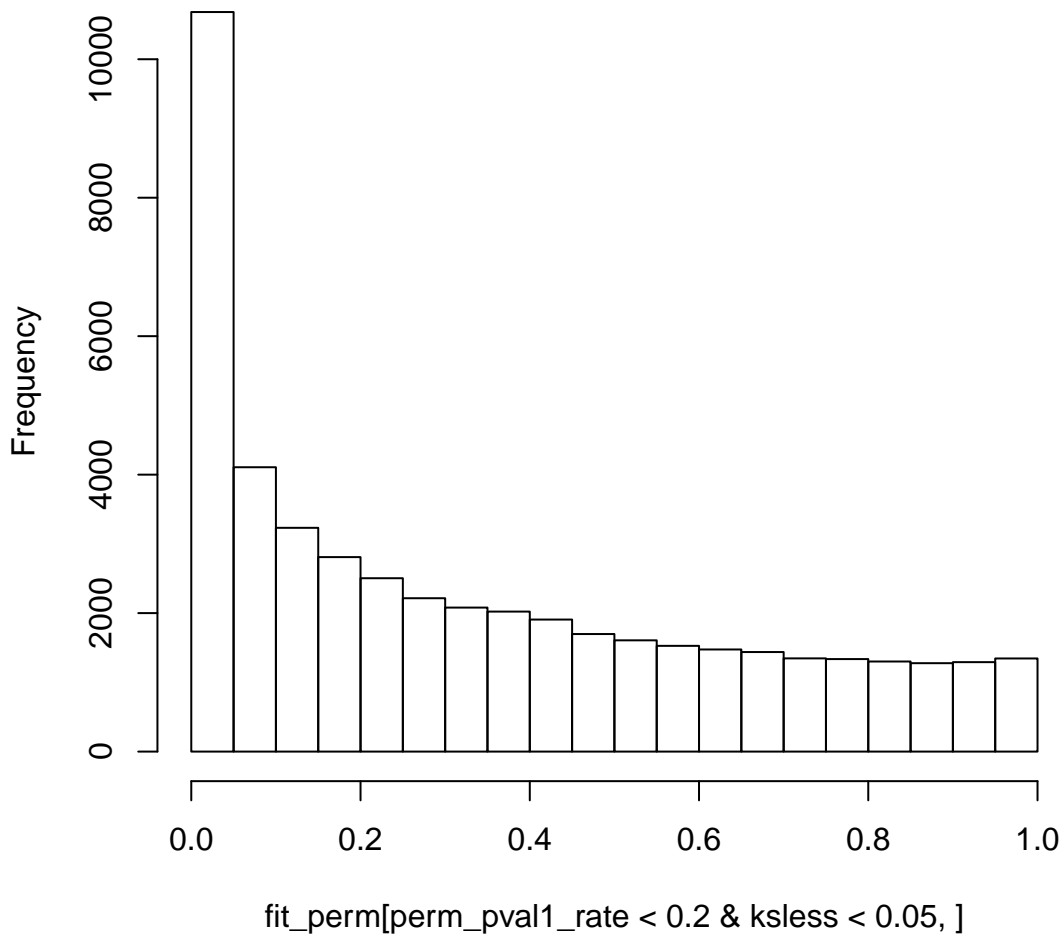
# permutation pvalues with pval1\_rate<0.2,ksgreater sig



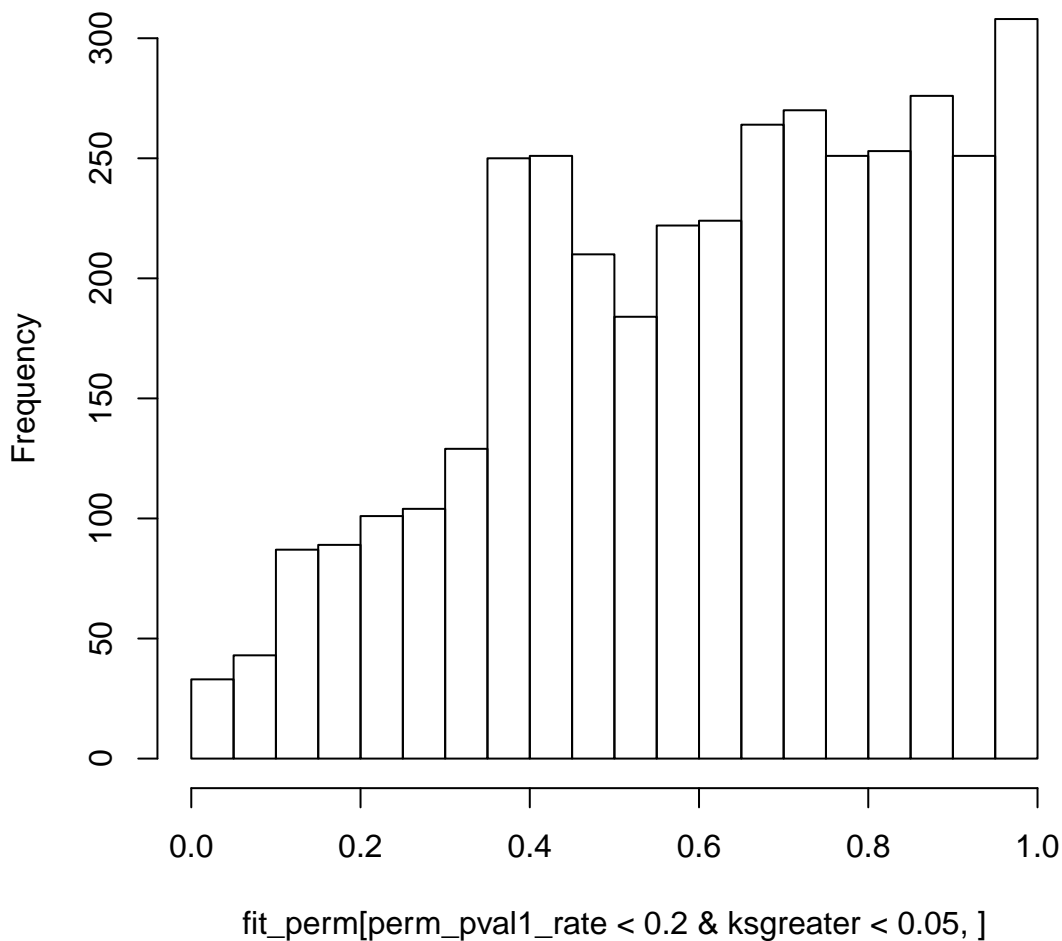
# permutation pvalues with pval1\_rate<0.2,ks no sig



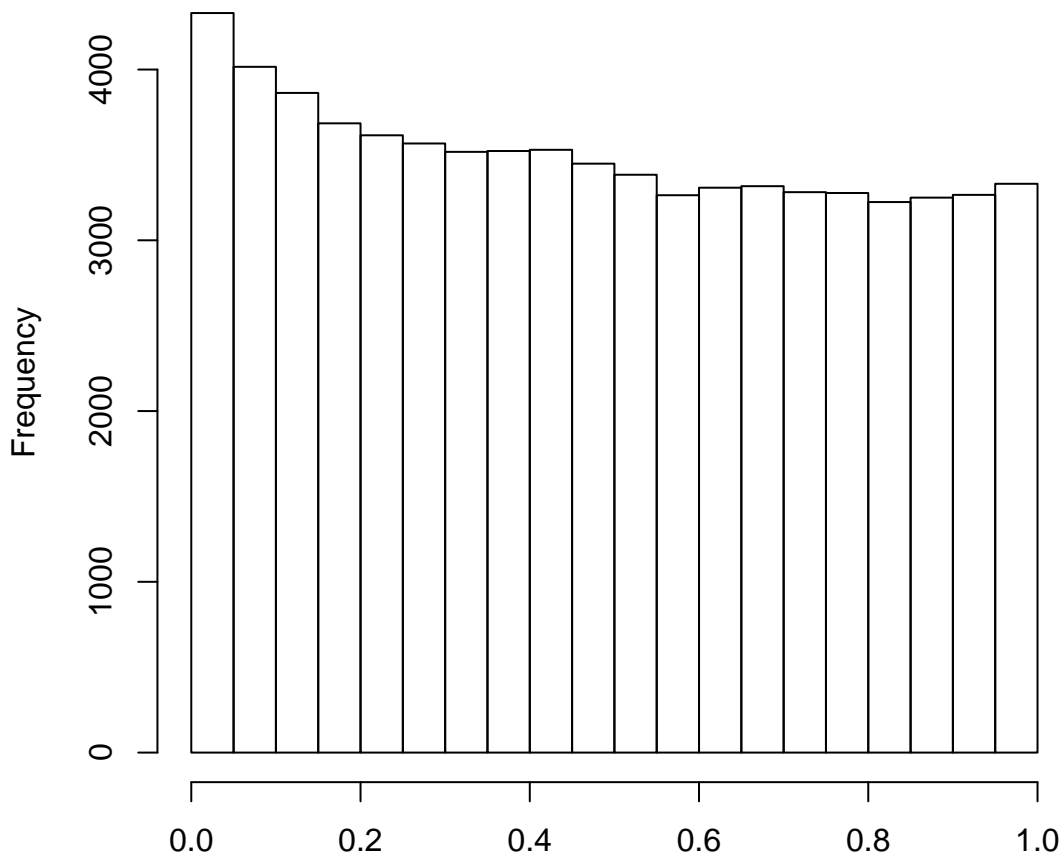
pvalues from permutation data with pval1\_rate<0.2,ksless sig



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, ]`