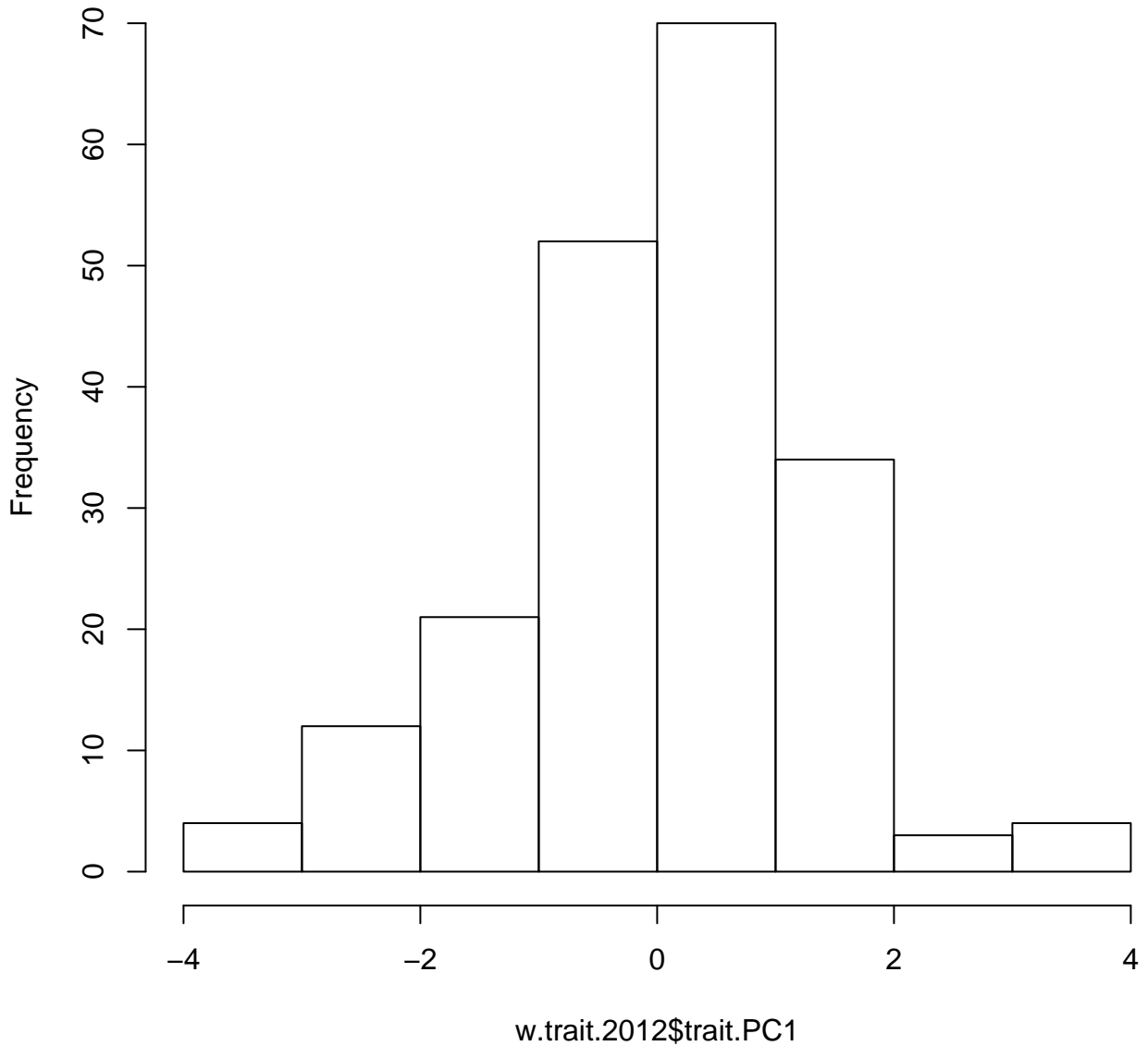
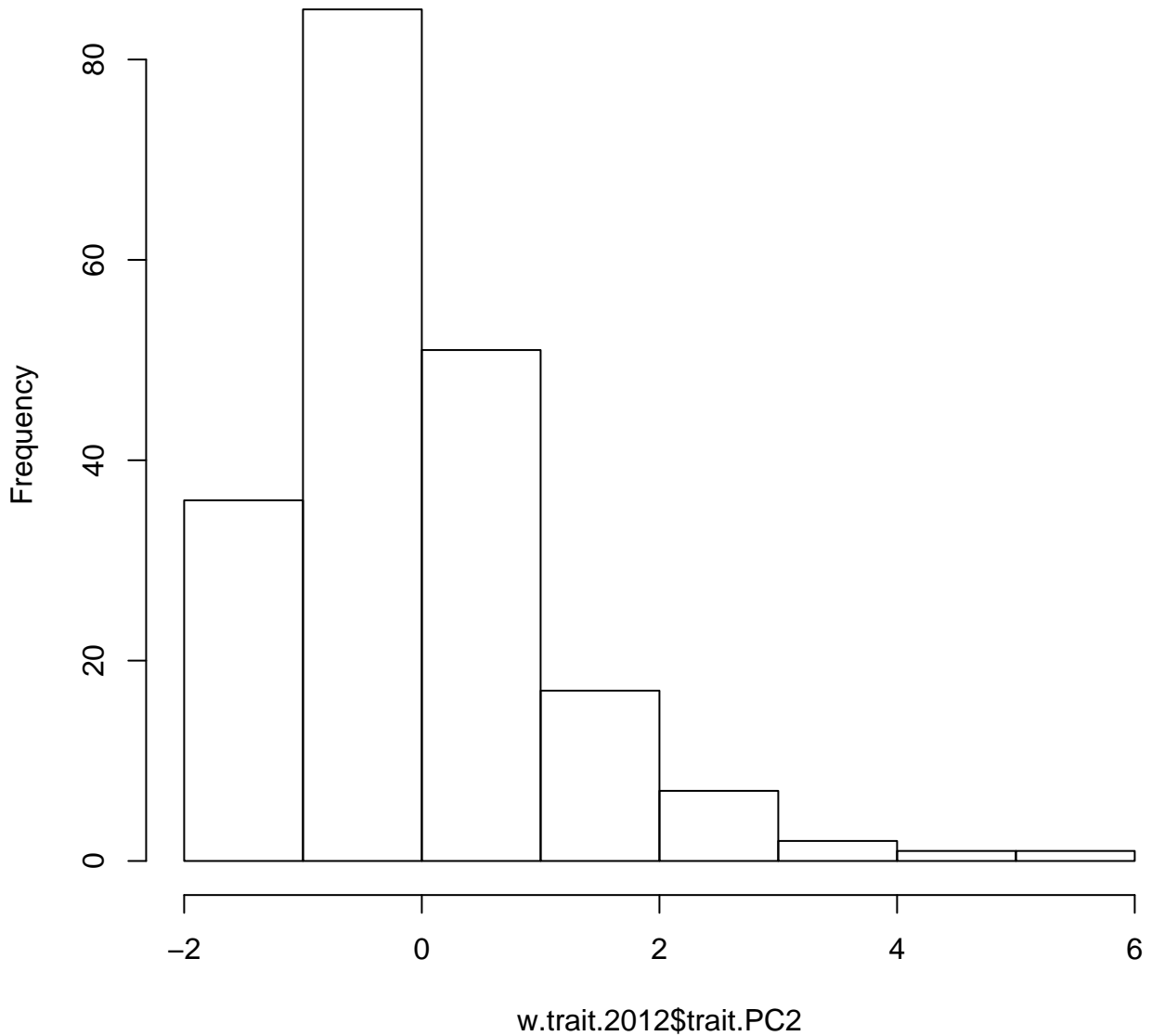


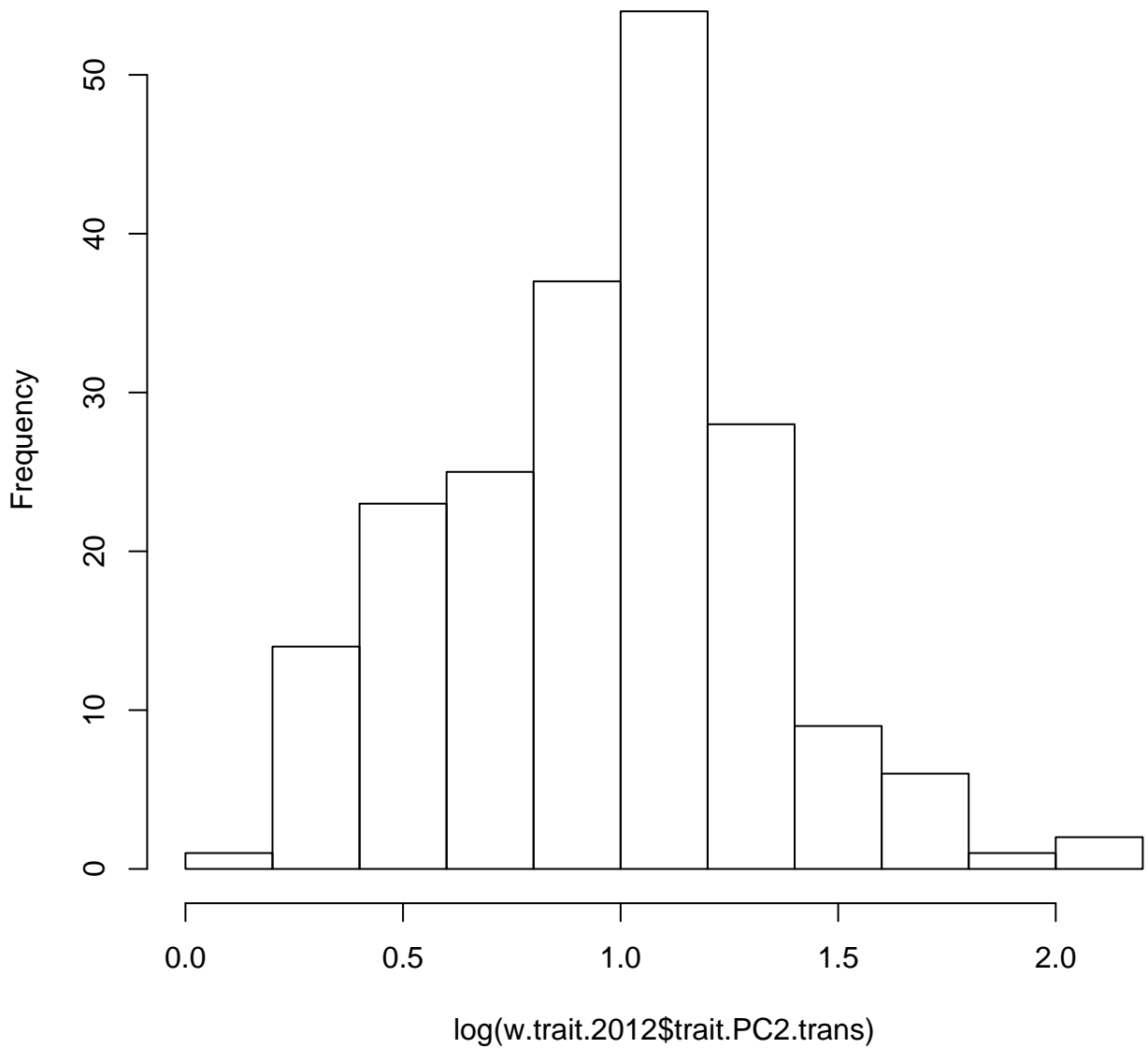
Histogram of w.trait.2012\$trait.PC1



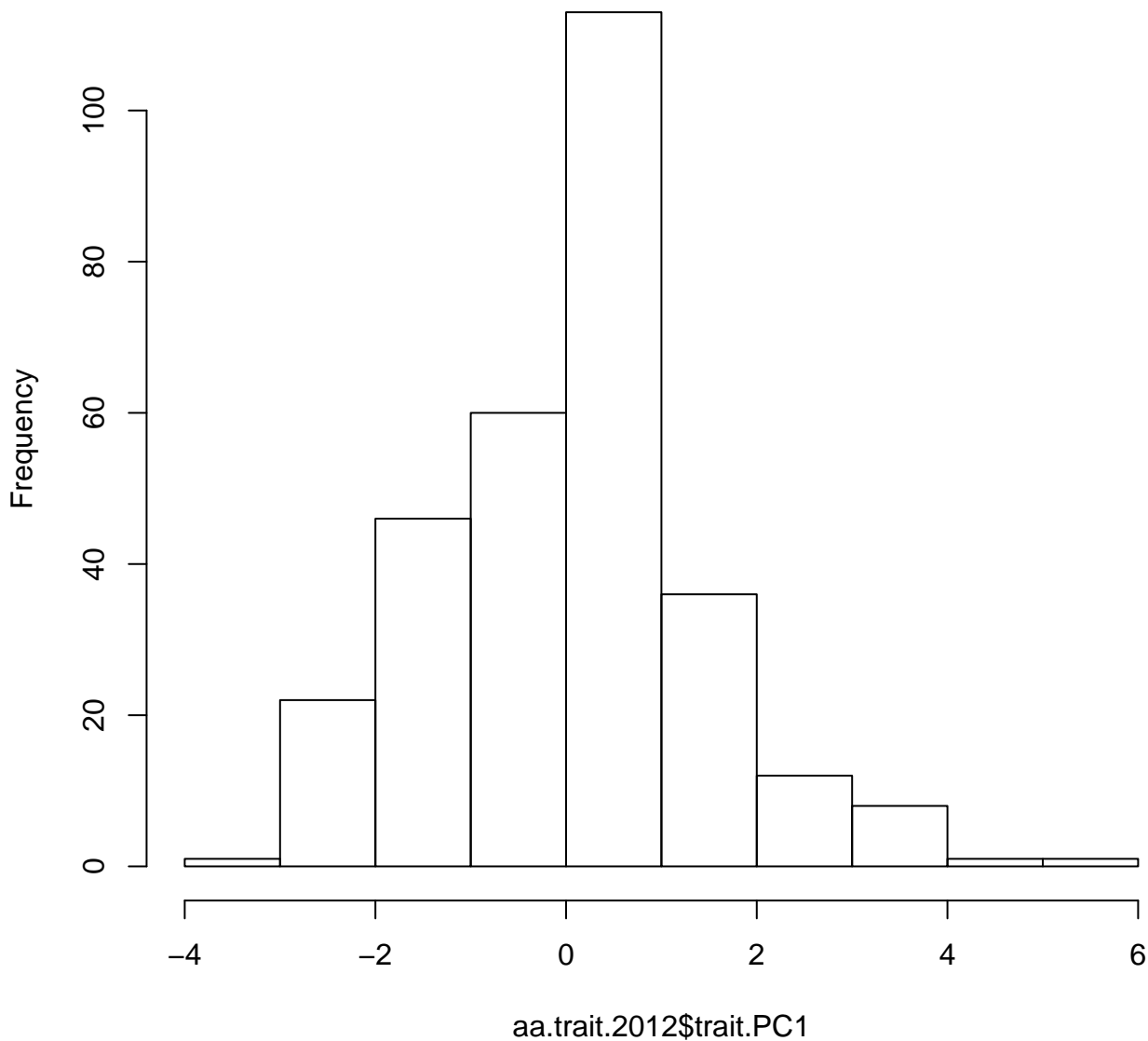
Histogram of w.trait.2012\$trait.PC2



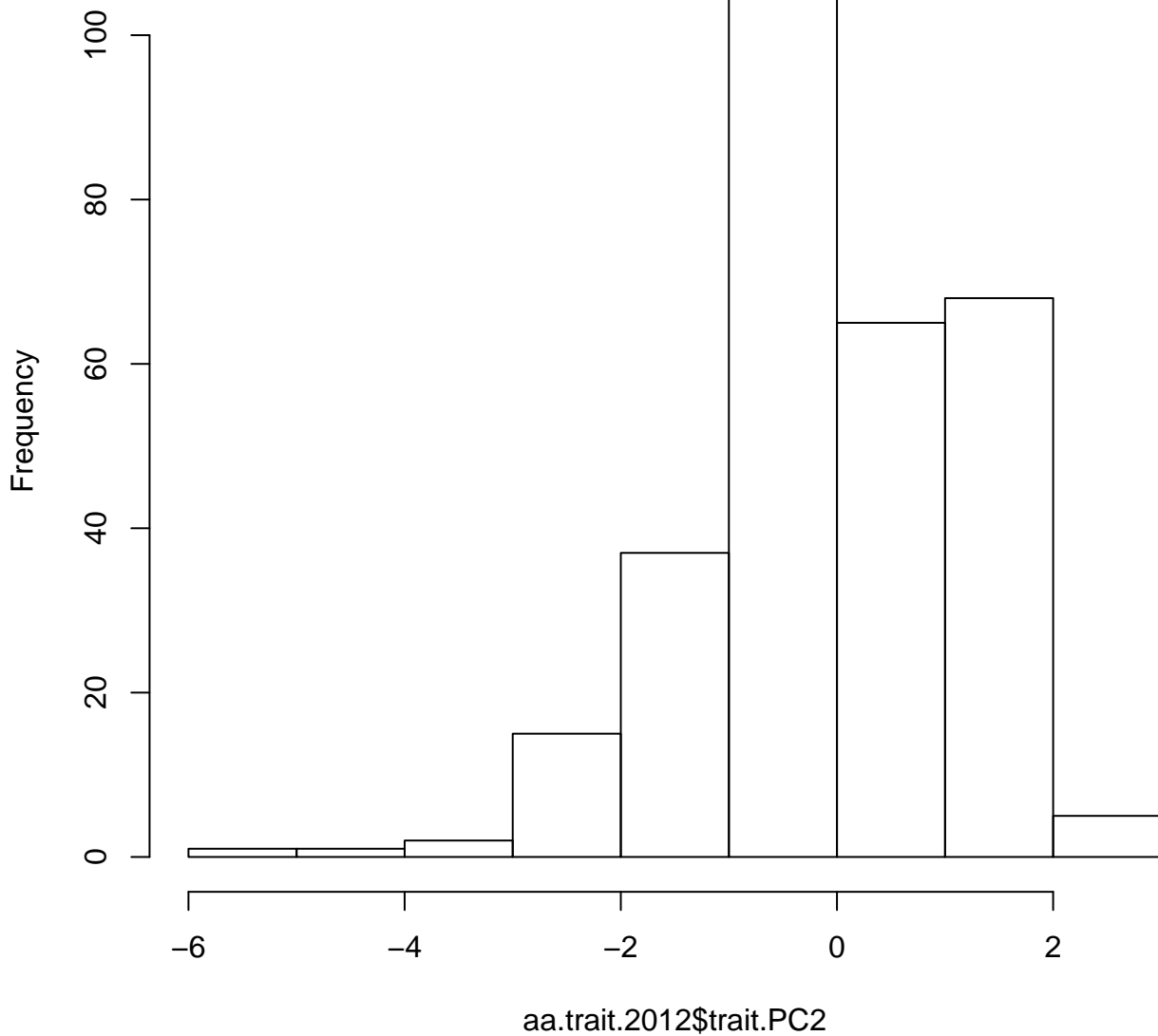
Histogram of $\log(w.\text{trait.2012}\$ \text{trait.PC2.trans})$



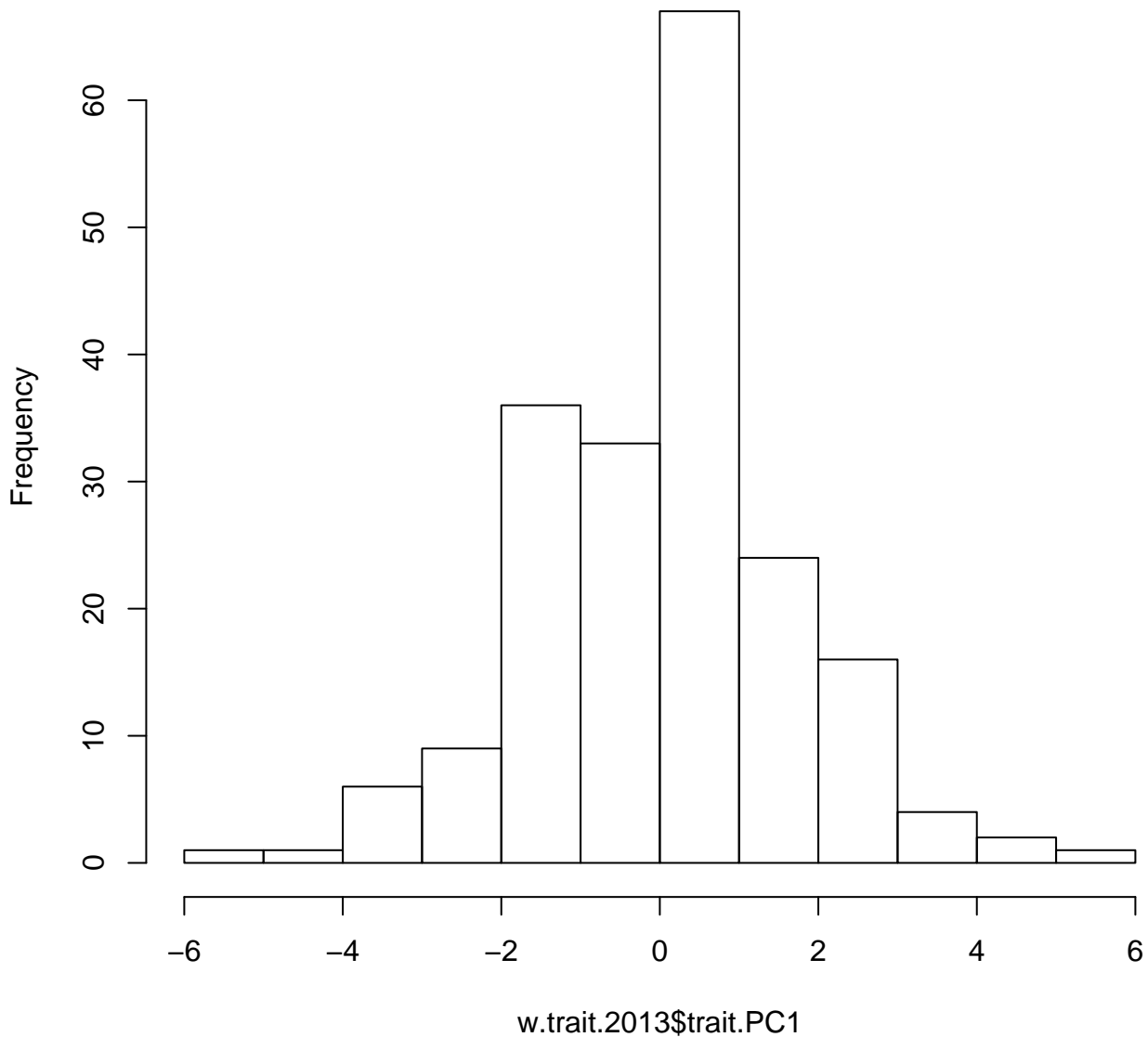
Histogram of aa.trait.2012\$trait.PC1



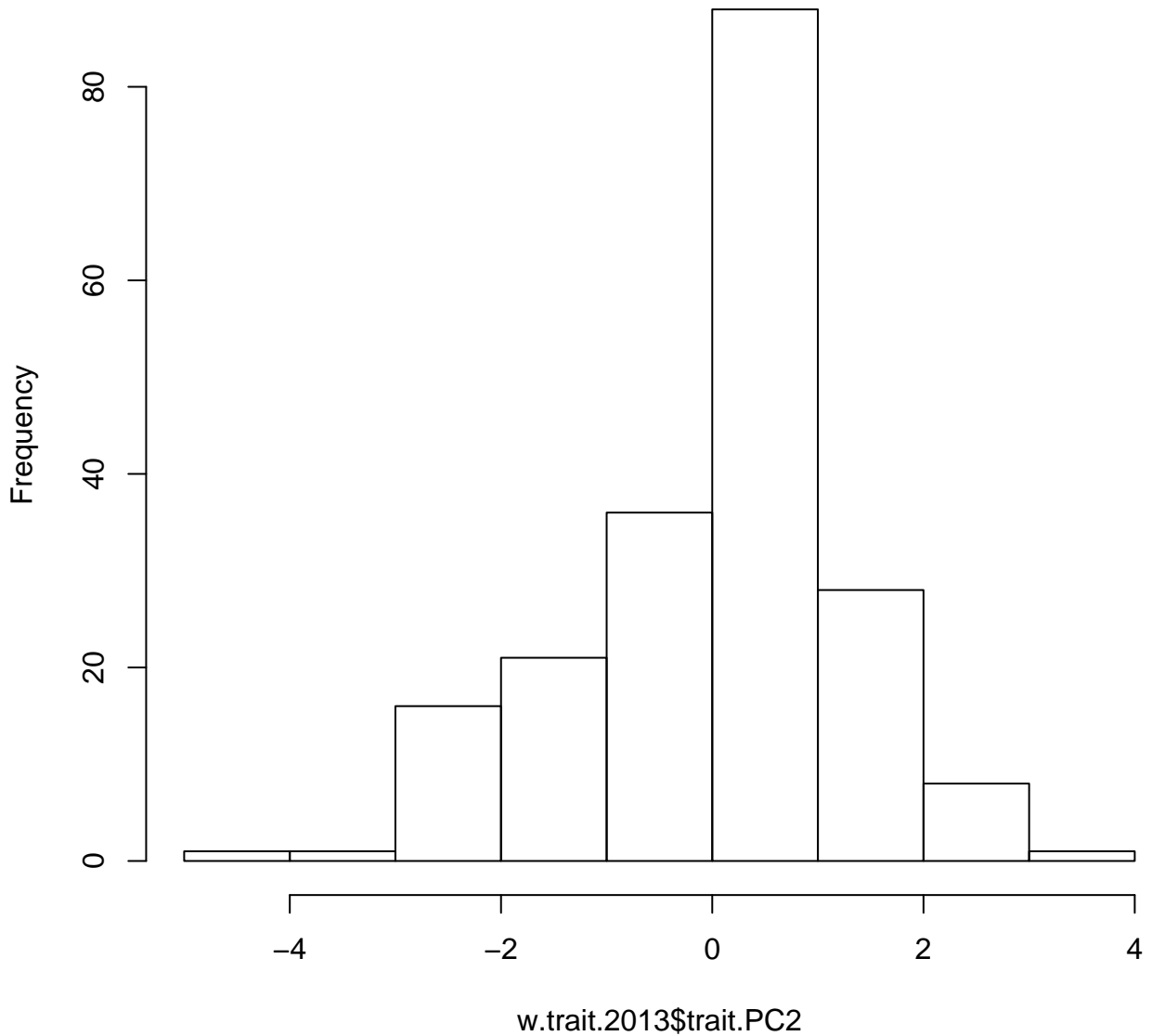
Histogram of aa.trait.2012\$trait.PC2



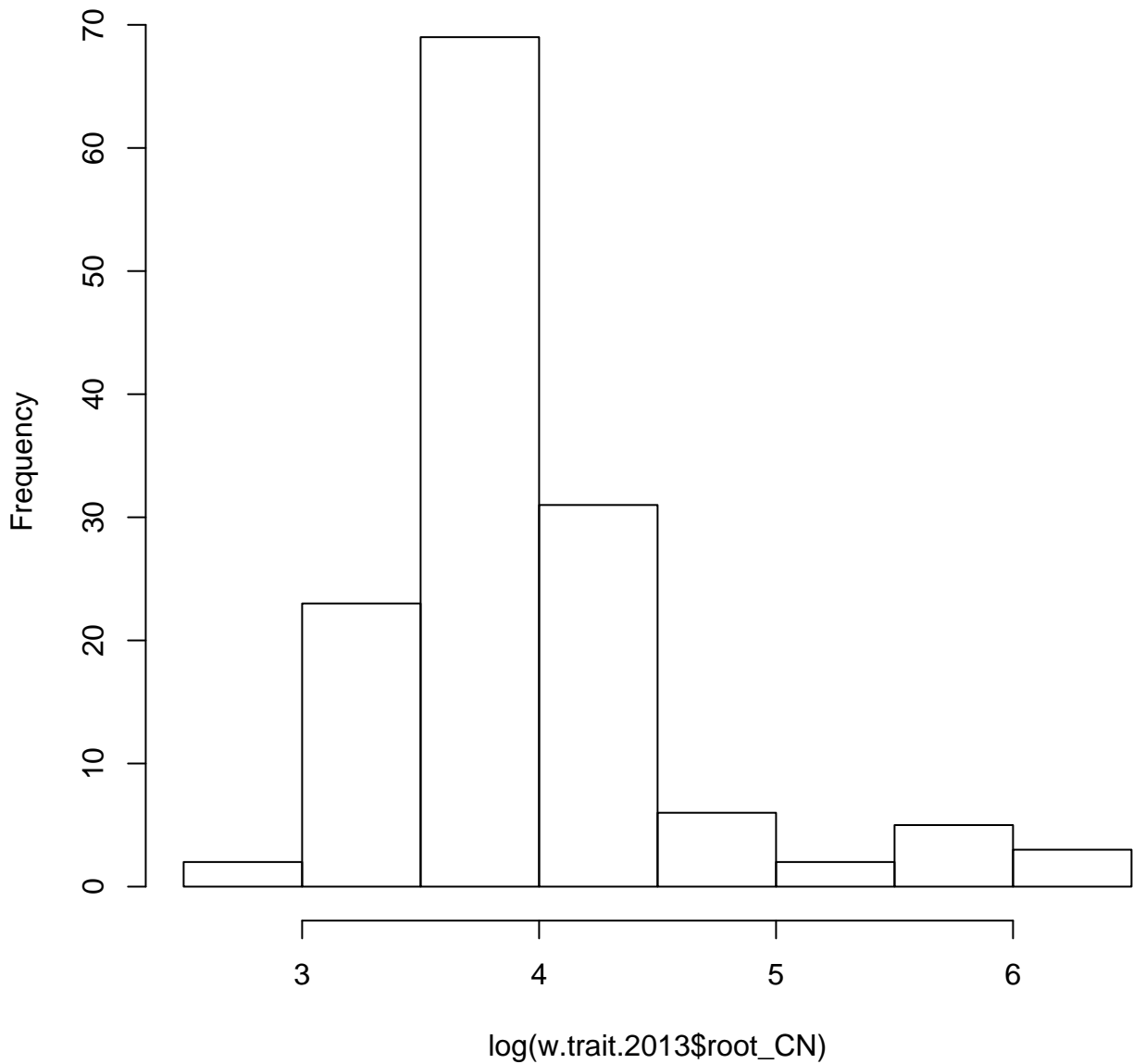
Histogram of w.trait.2013\$trait.PC1



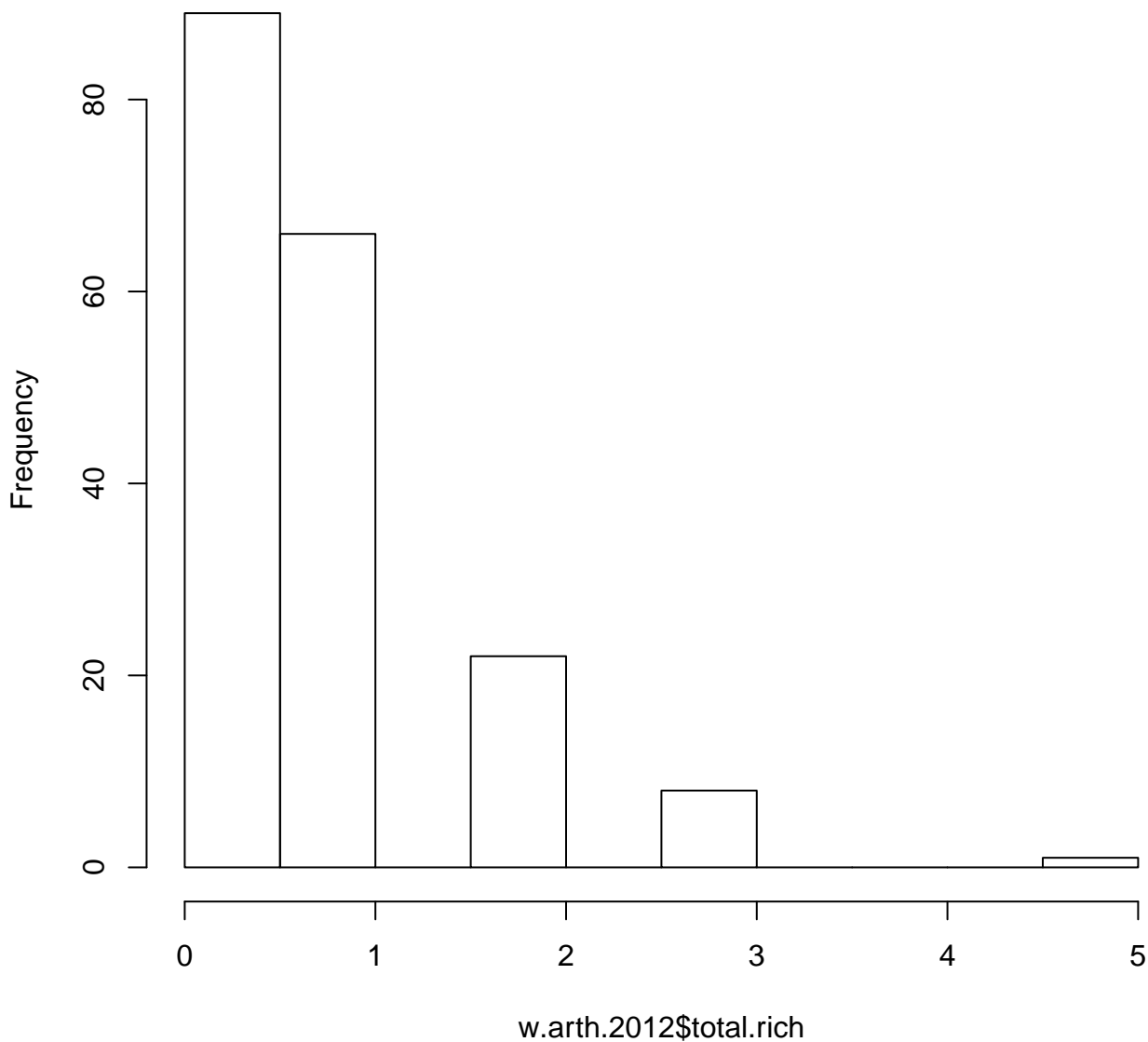
Histogram of w.trait.2013\$trait.PC2

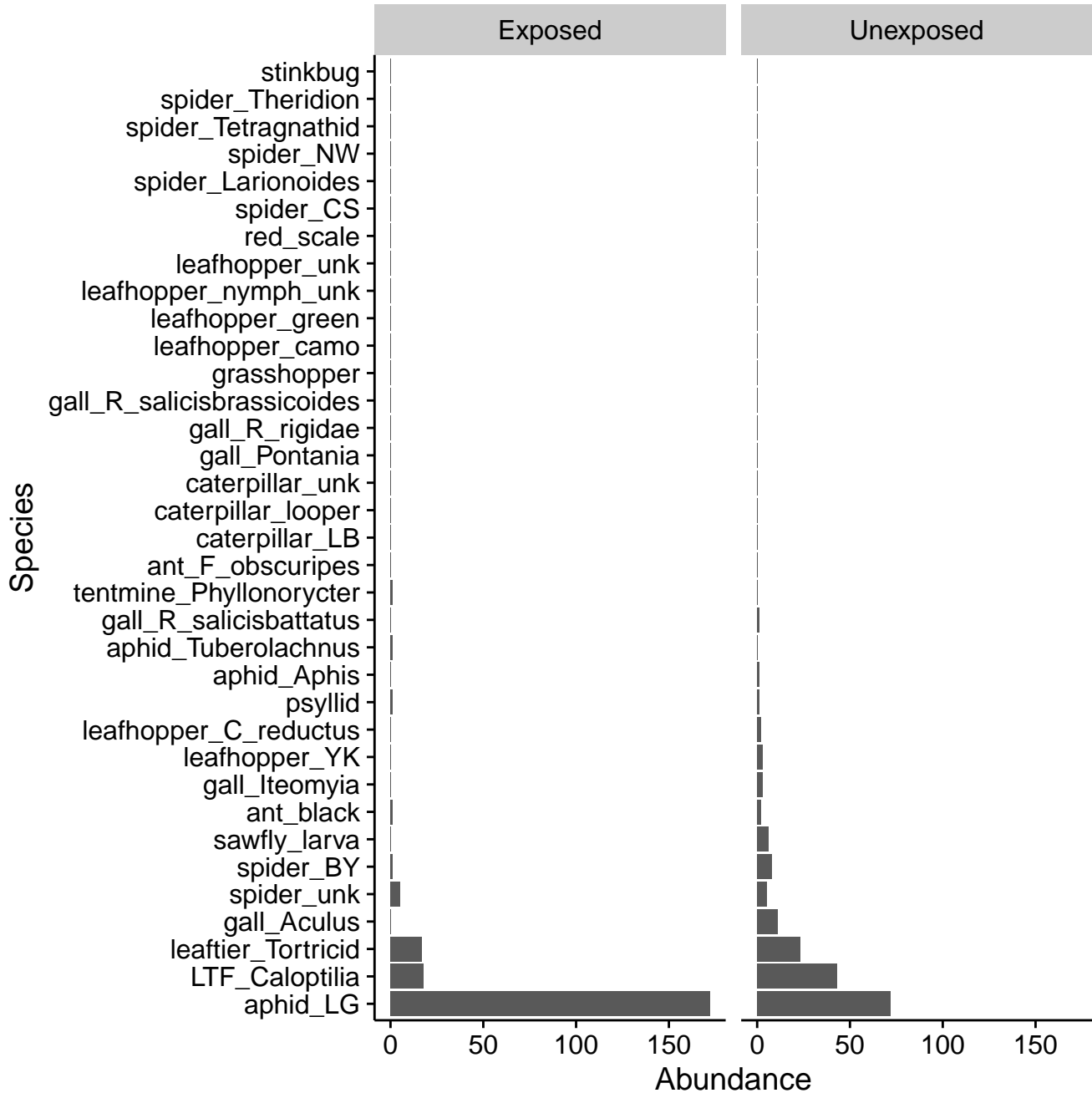


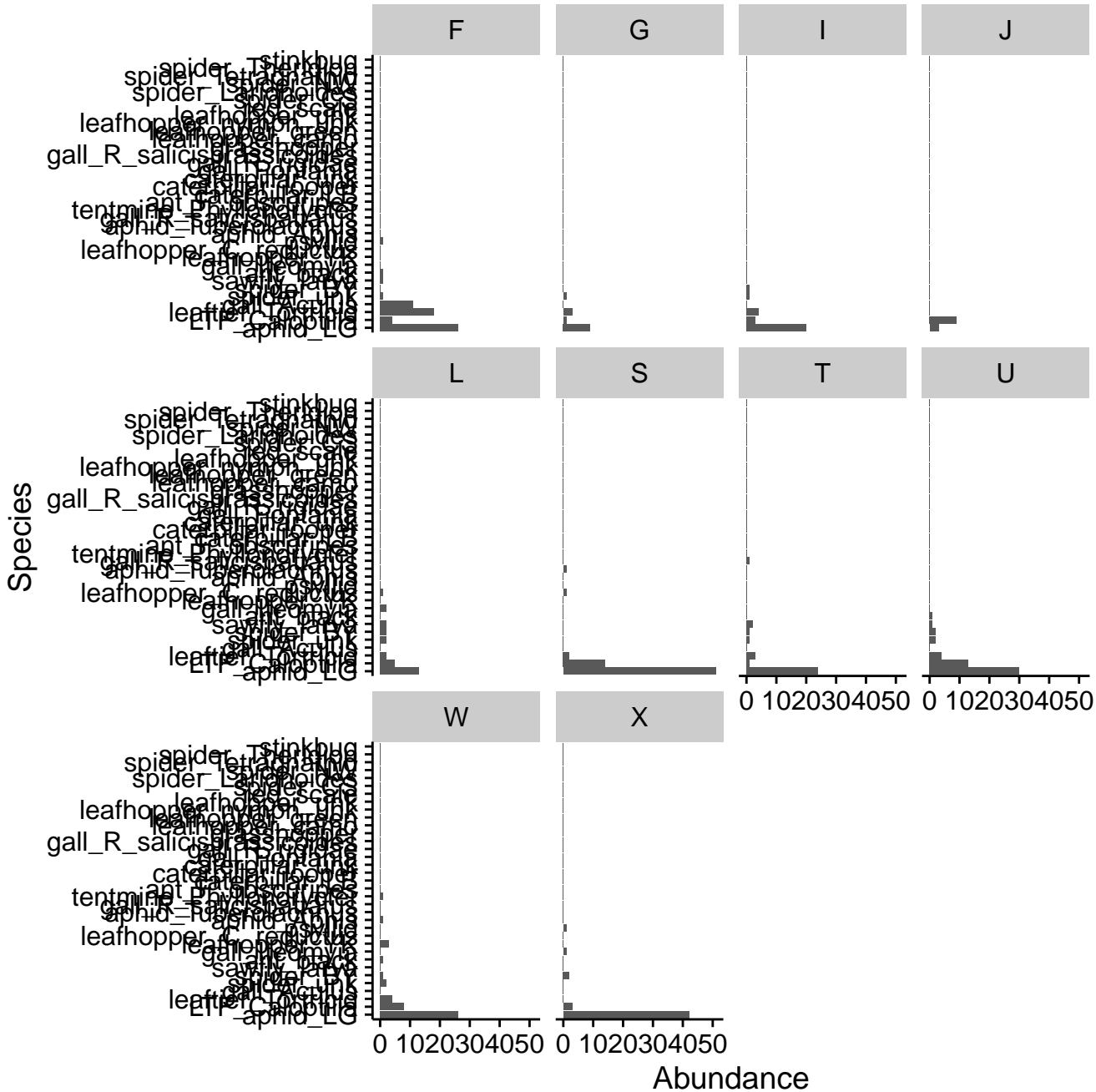
Histogram of $\log(\text{w.trait.2013}\$\text{root_CN})$



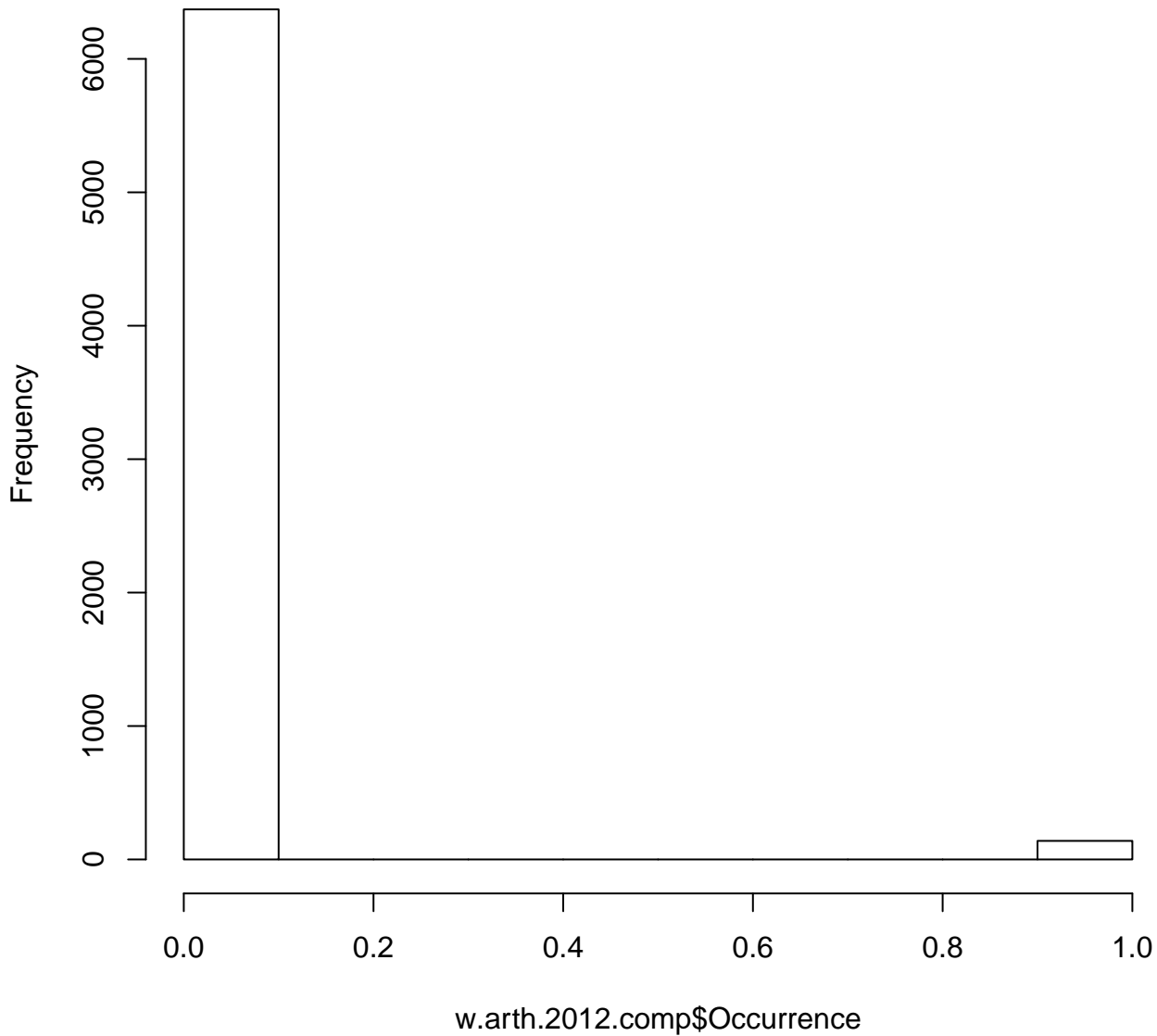
Histogram of w.arth.2012\$total.rich



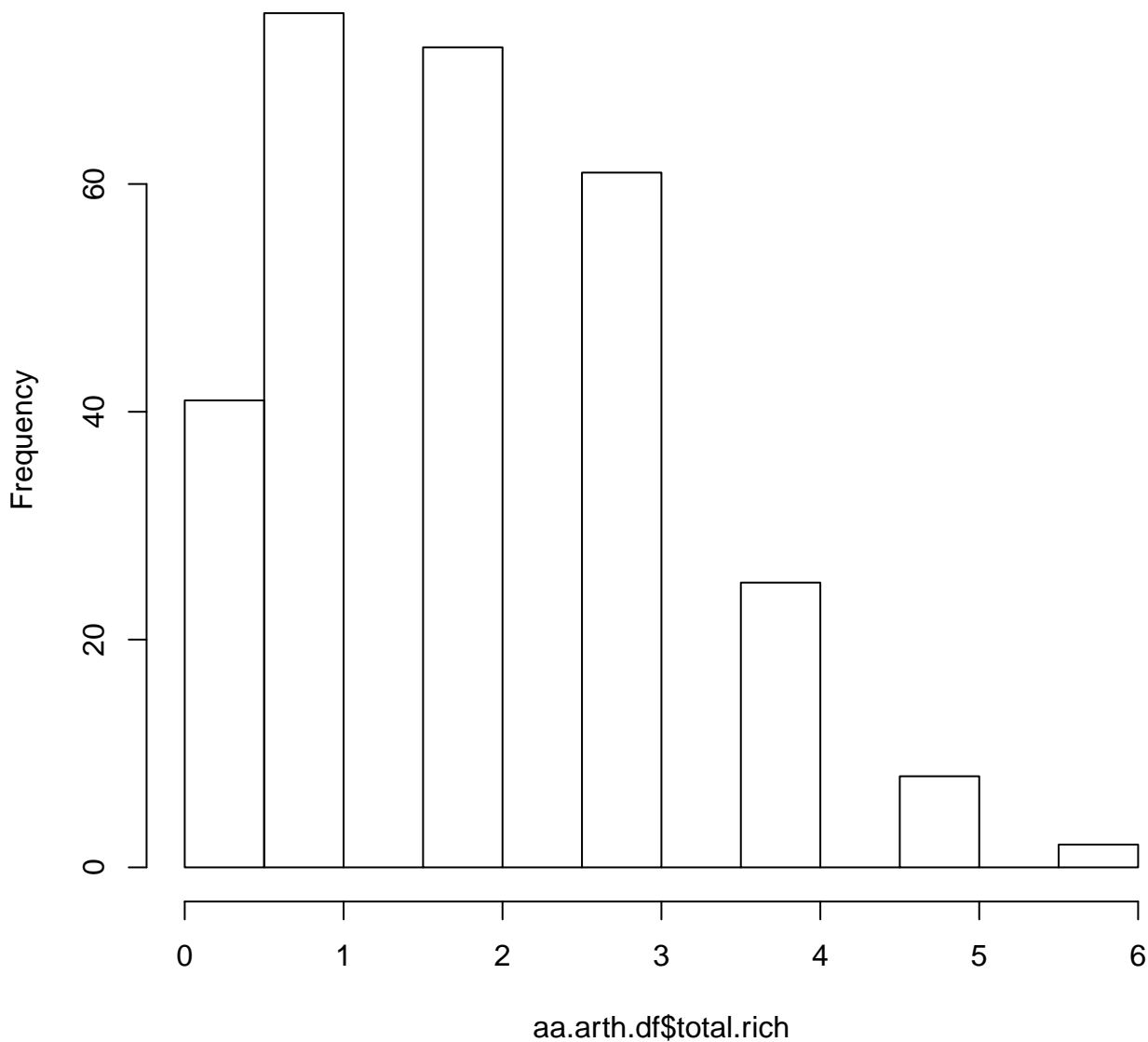




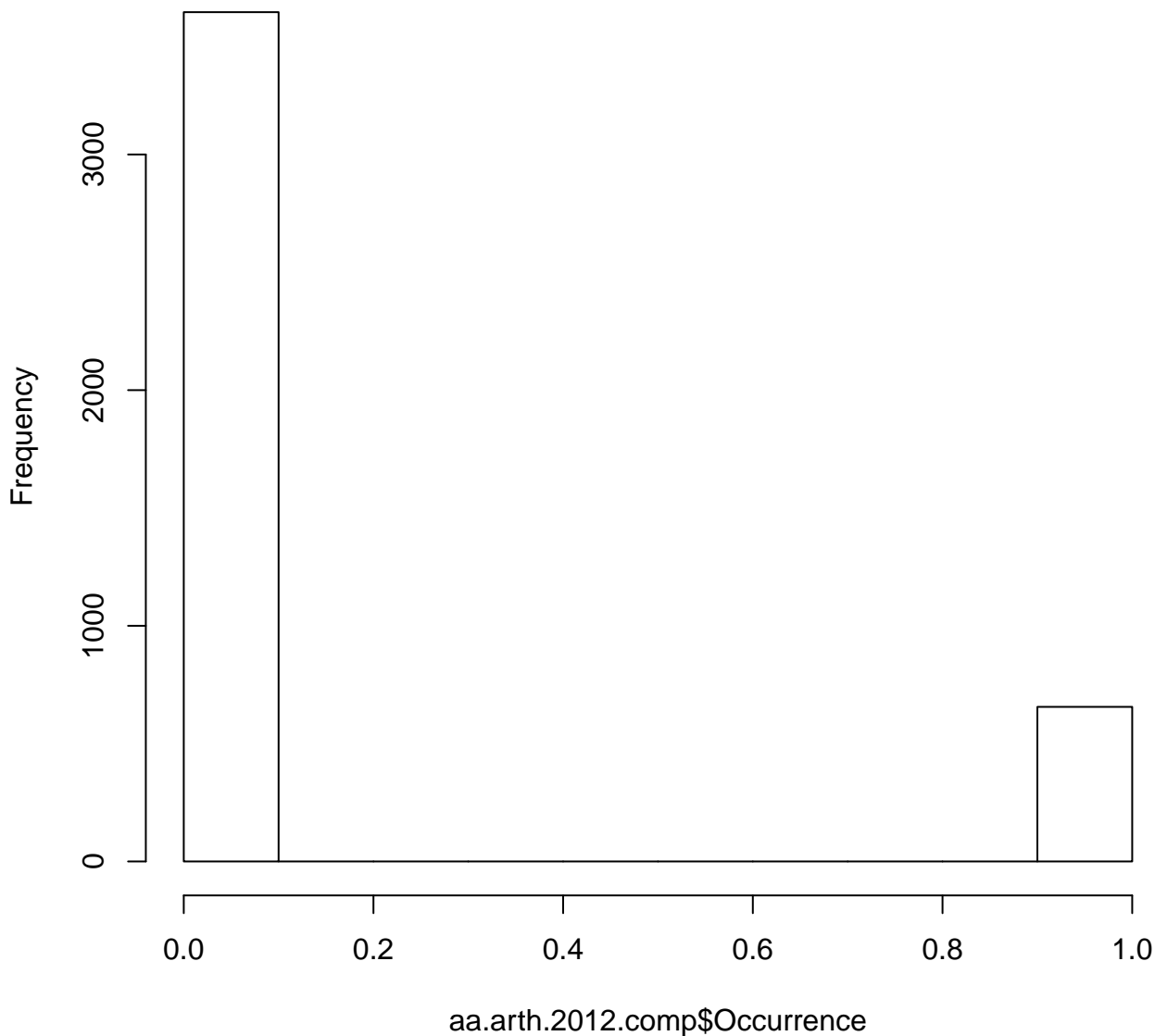
Histogram of w.arth.2012.comp\$Occurrence



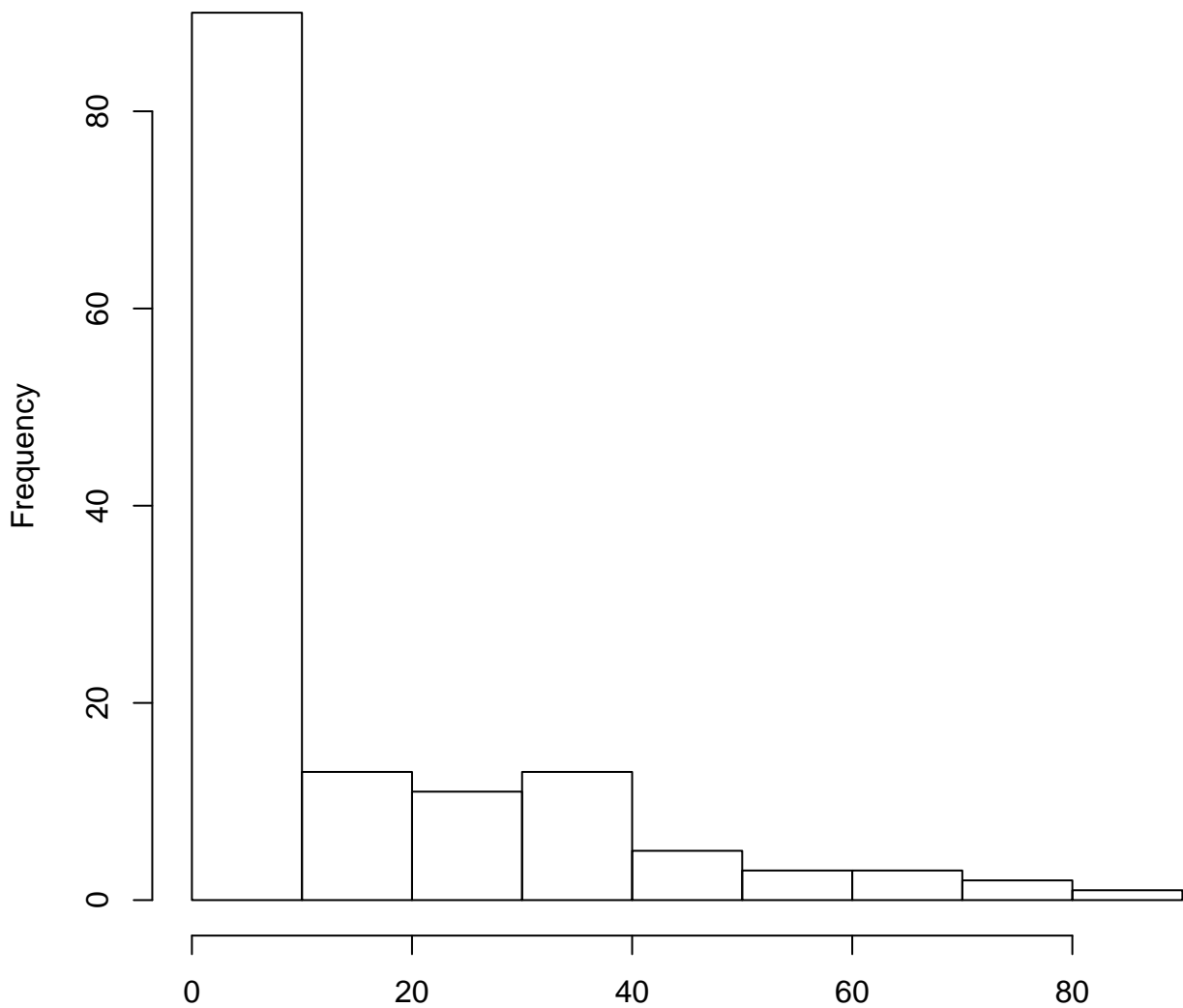
Histogram of aa.arth.df\$total.rich



Histogram of aa.arth.2012.comp\$Occurrence

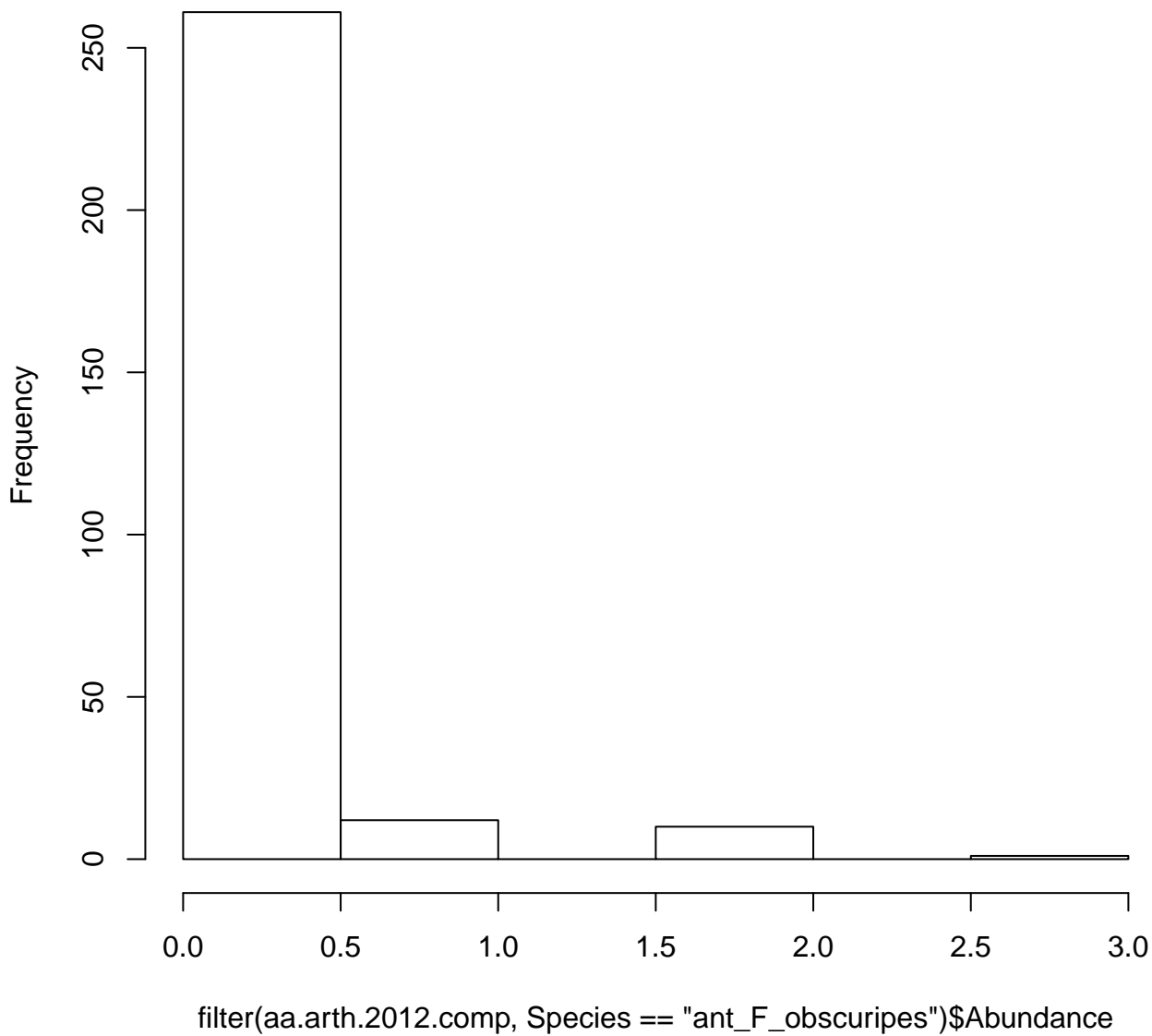


`filter(aa.arth.2012.comp, Species == "aphid_Aphis", Aphid.treatment == "ap`

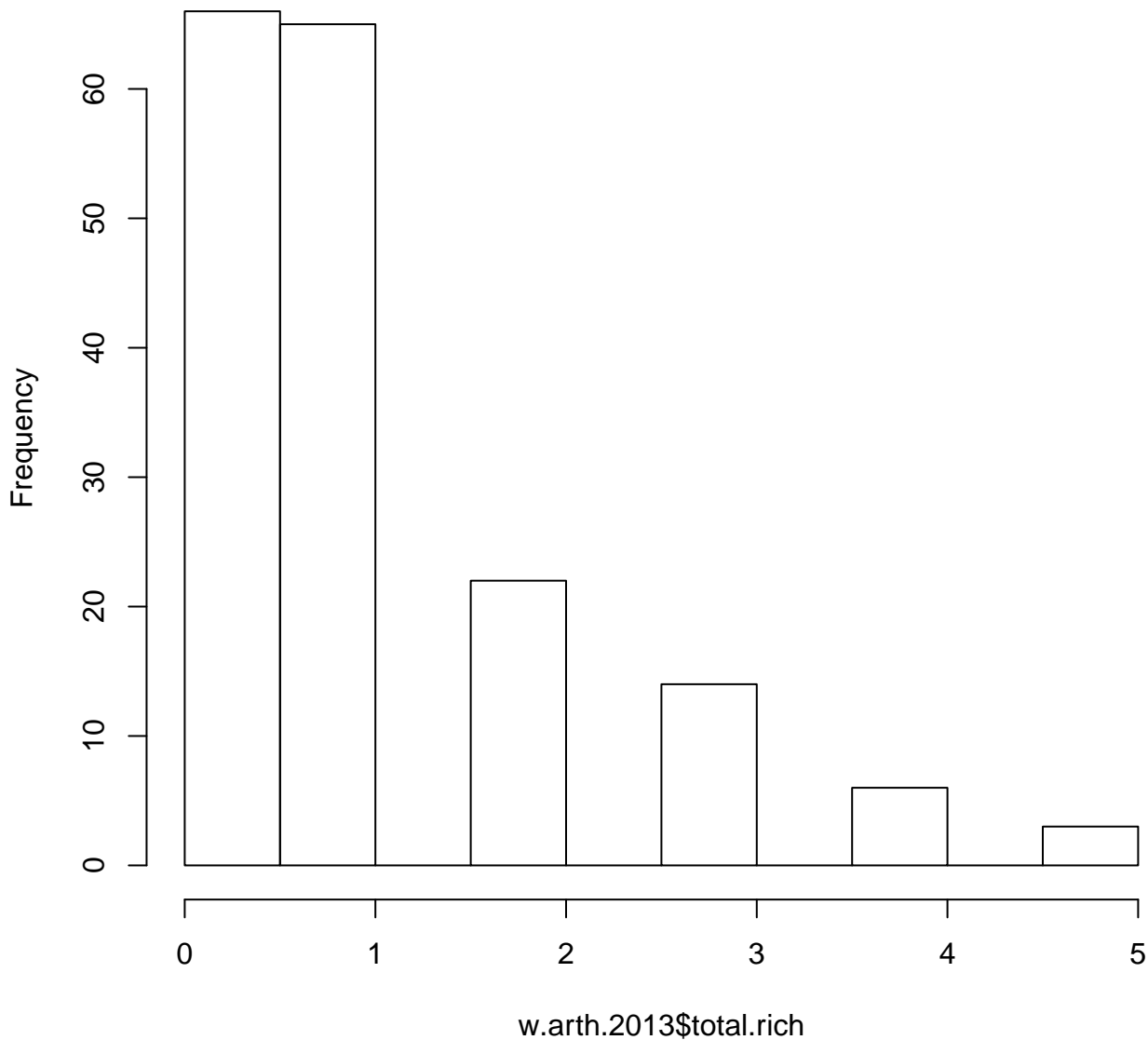


`filter(aa.arth.2012.comp, Species == "aphid_Aphis", Aphid.treatment == "aphid")$Abundance`

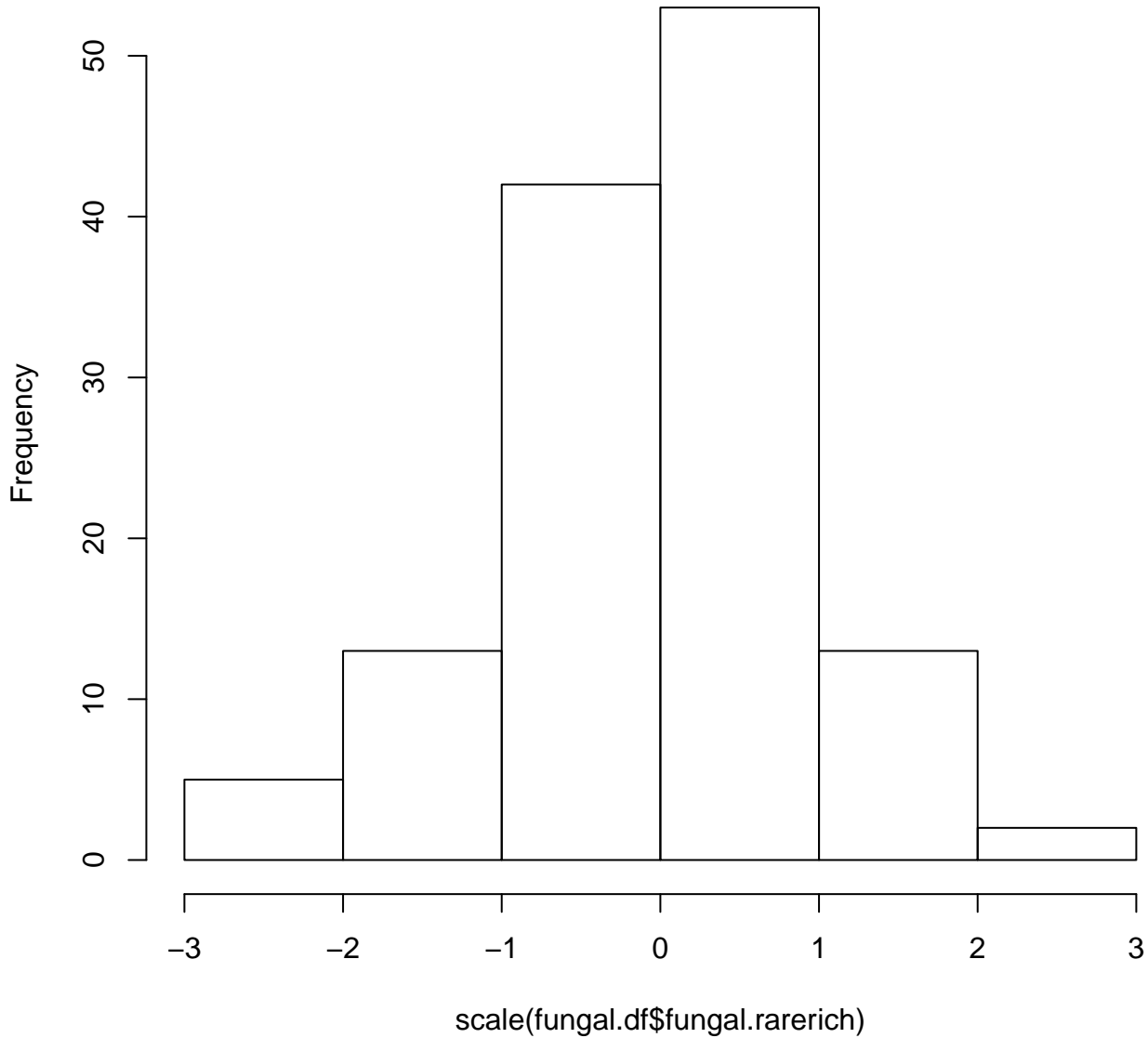
Histogram of filter(aa.arth.2012.comp, Species == "ant_F_obscuripes")\$Abund



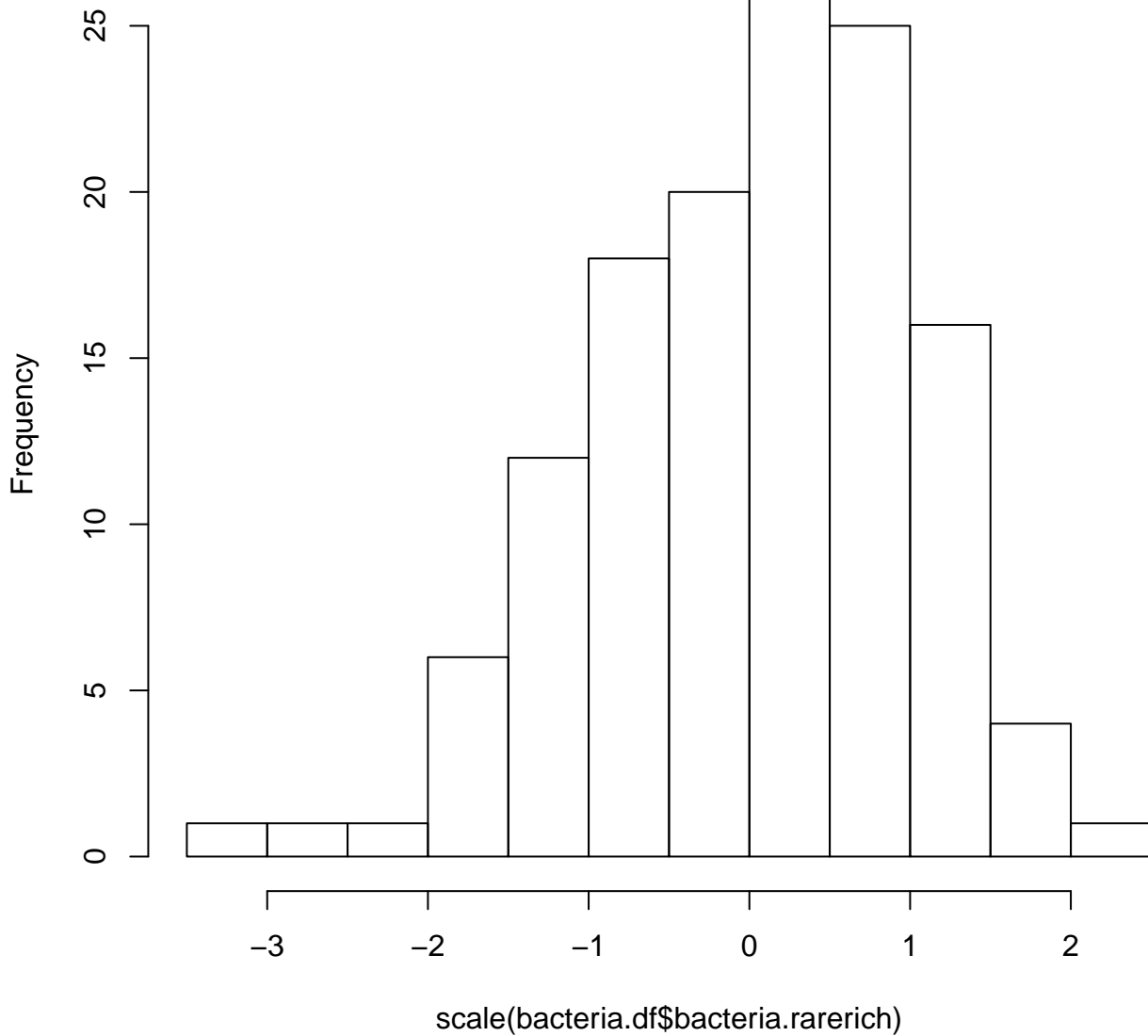
Histogram of w.arth.2013\$total.rich

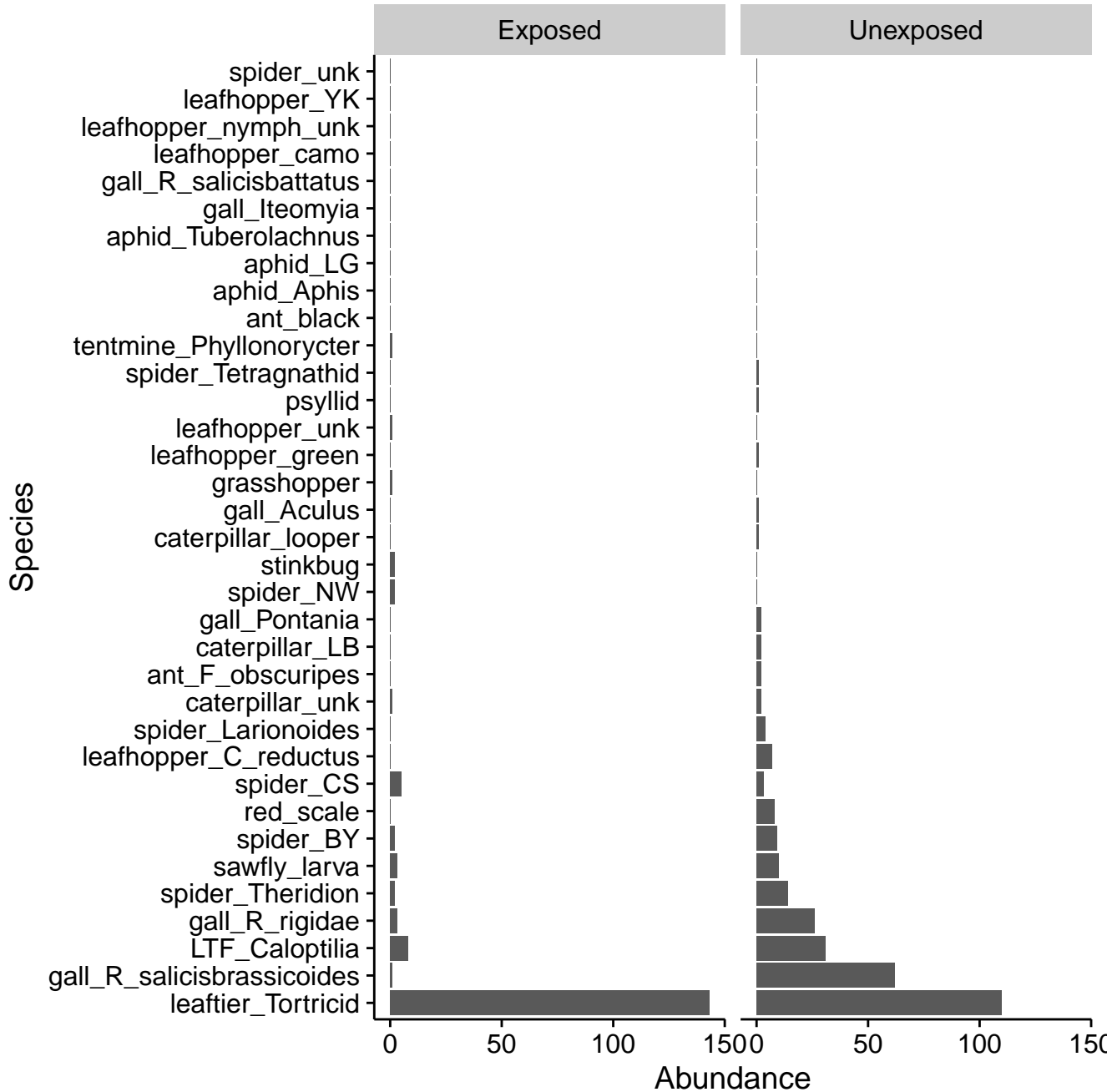


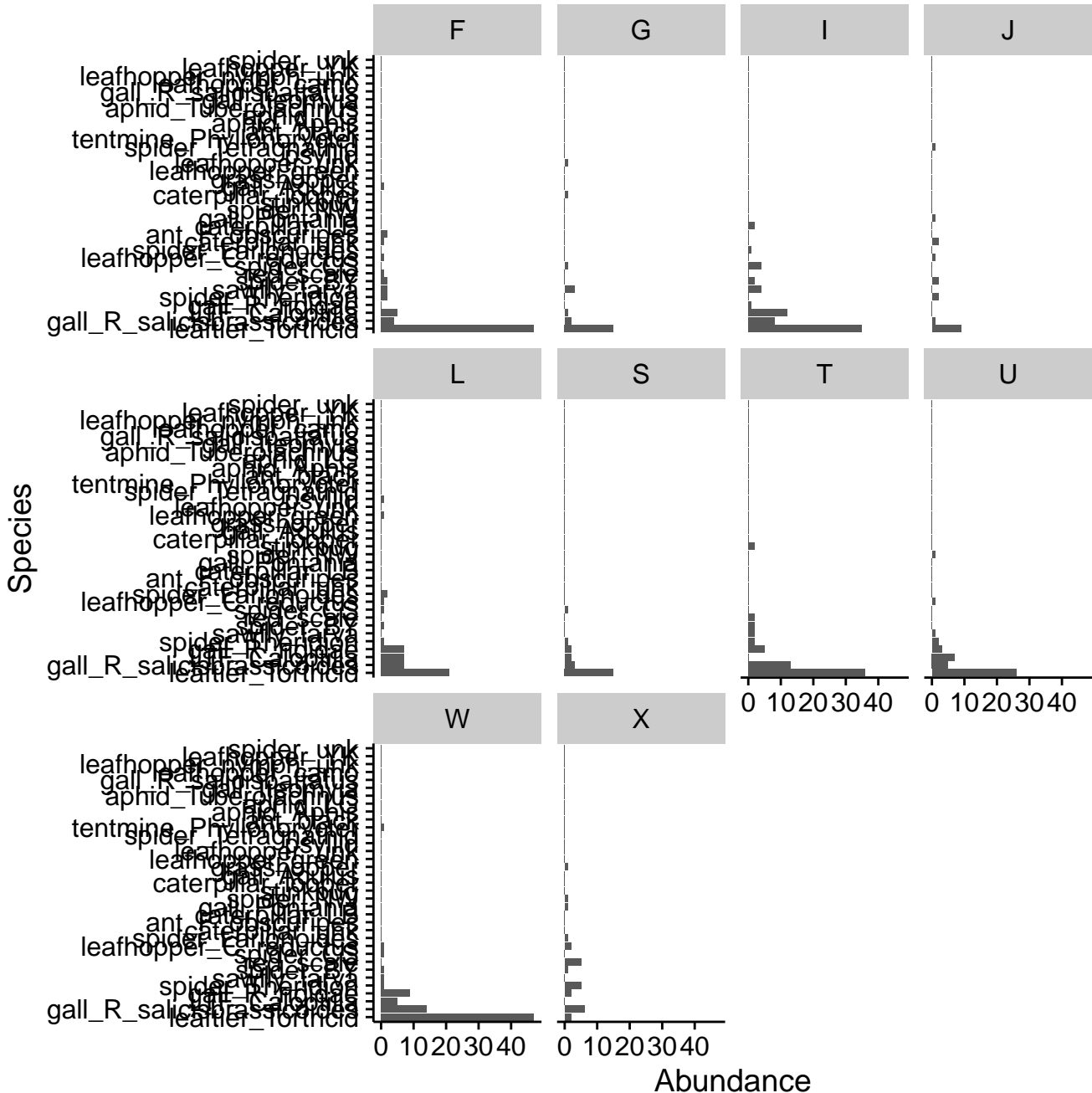
Histogram of scale(fungal.df\$fungal.rarerich)



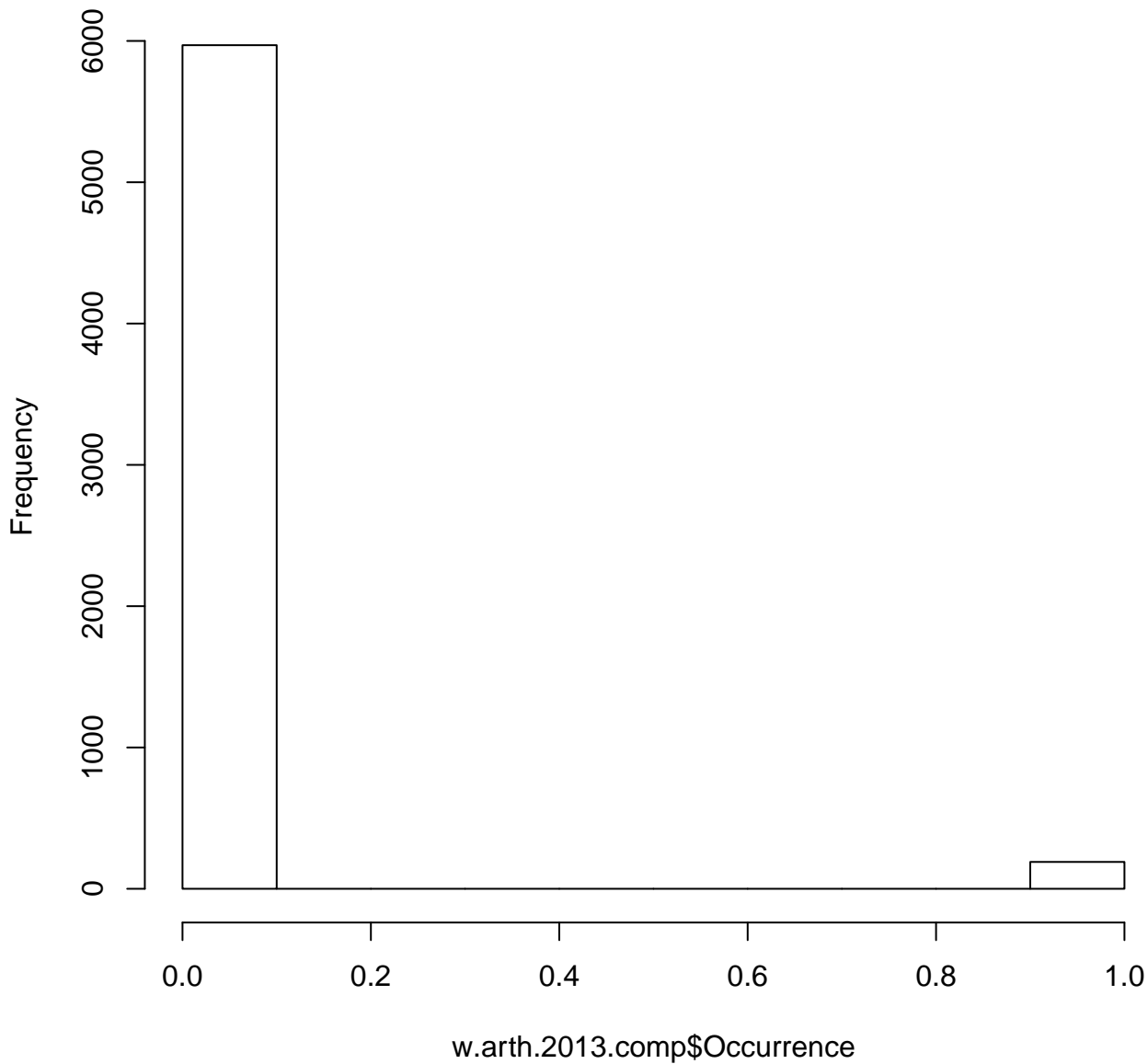
Histogram of `scale(bacteria.df$bacteria.rarerich)`



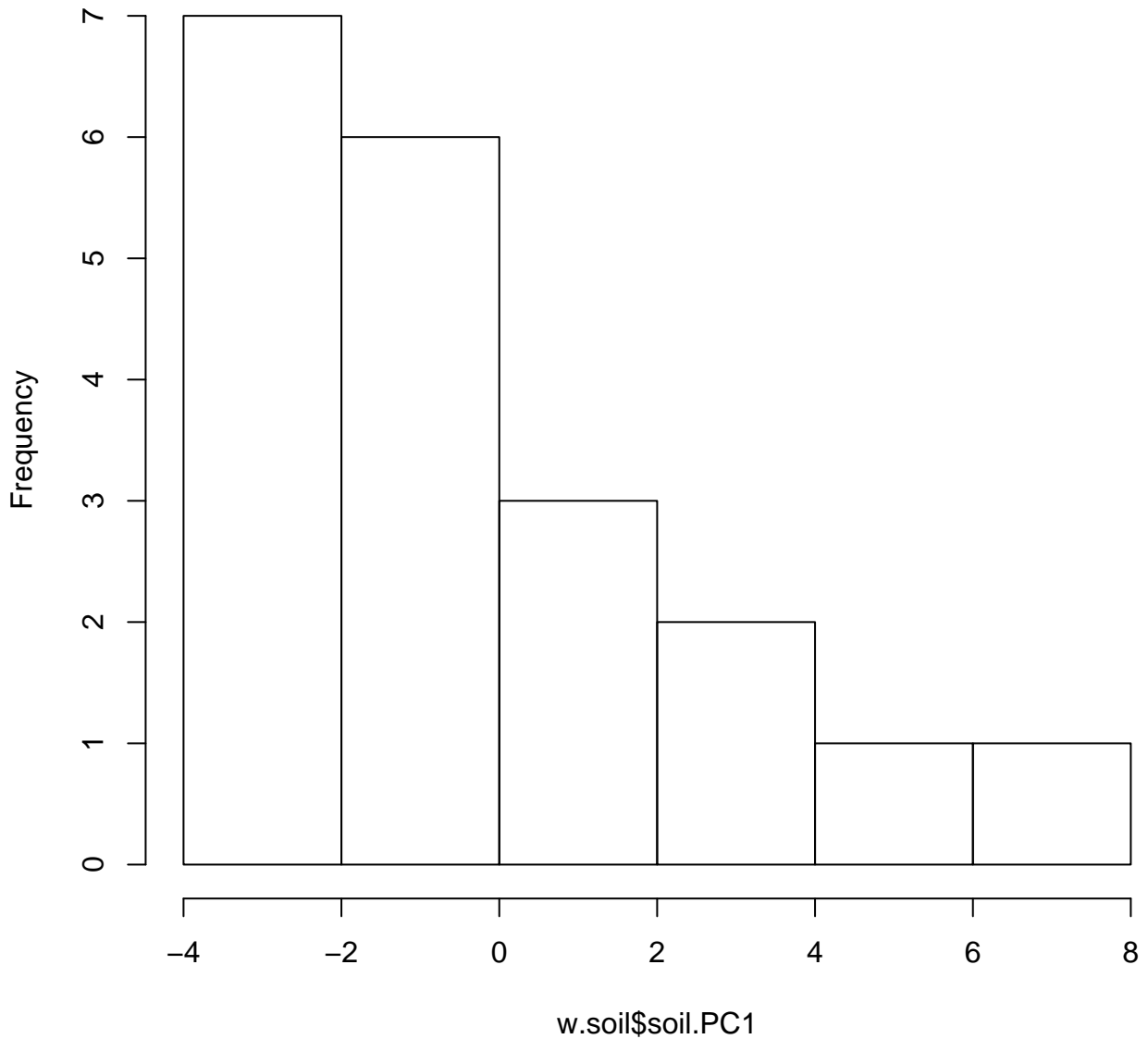




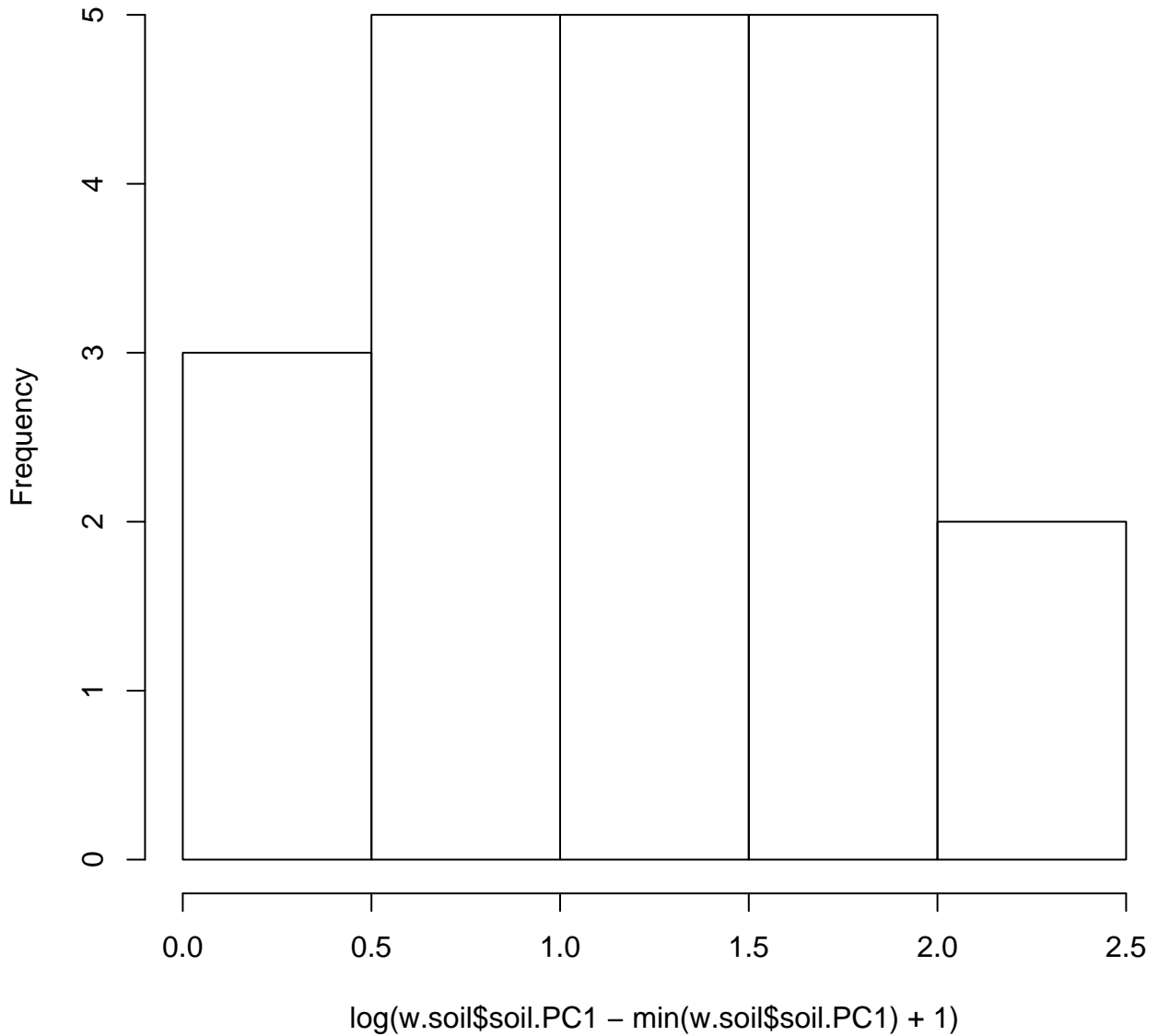
Histogram of w.arth.2013.comp\$Occurrence



Histogram of w.soil\$soil.PC1



Histogram of $\log(\text{w.soil}\$soil.PC1 - \min(\text{w.soil}\$soil.PC1) + 1)$



Histogram of w.soil\$soil.PC2

