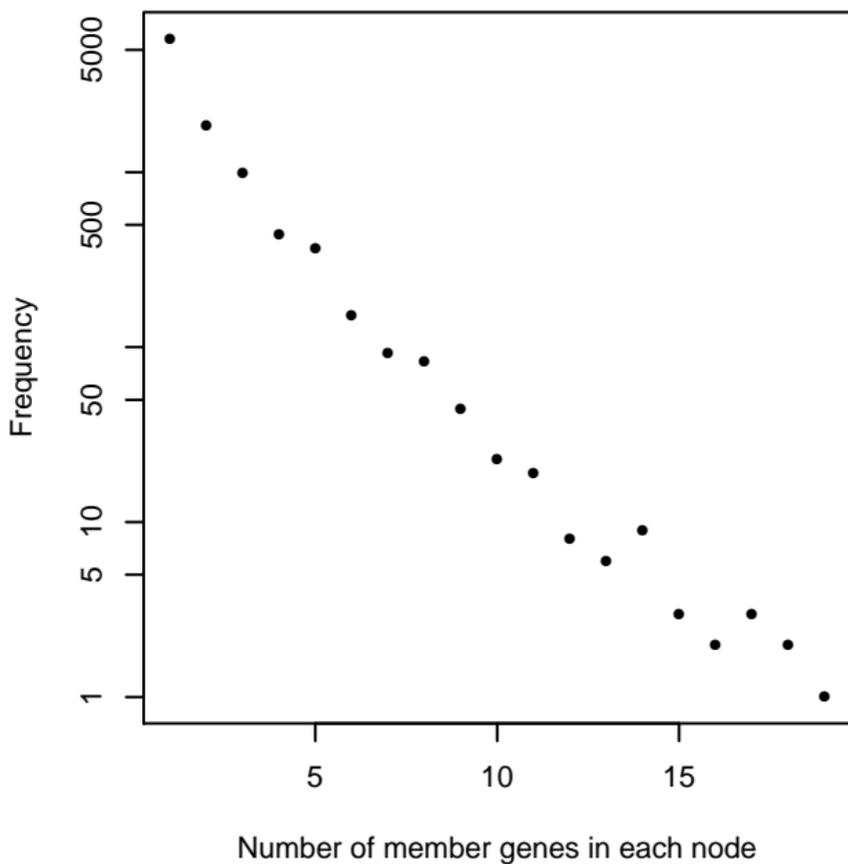
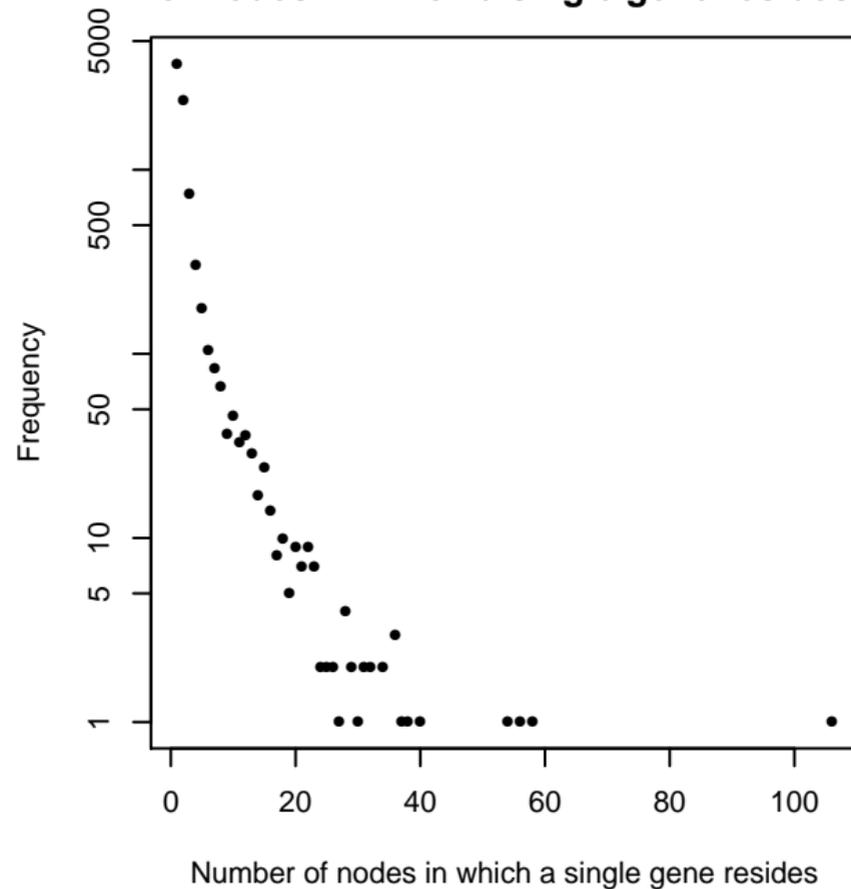


**(A) Distribution of the number of member genes in each node**



**(B) Distribution of the number of nodes in which a single gene resides**



**(C) Relationship between node count and gene count in pathways**

