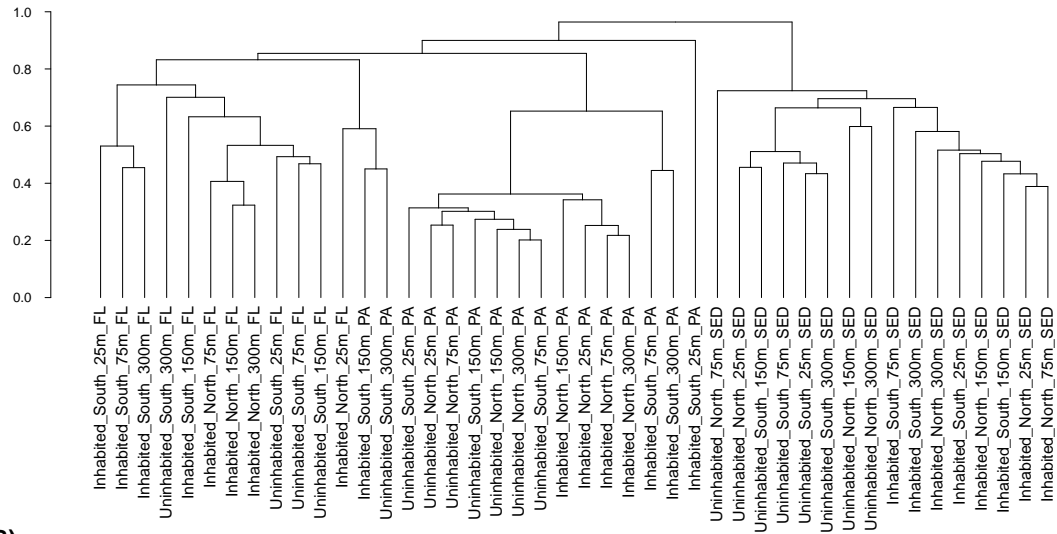


**(A)**

Bray-Curtis dissimilarity

**(B)**

Relative sequence abundance [%]

