



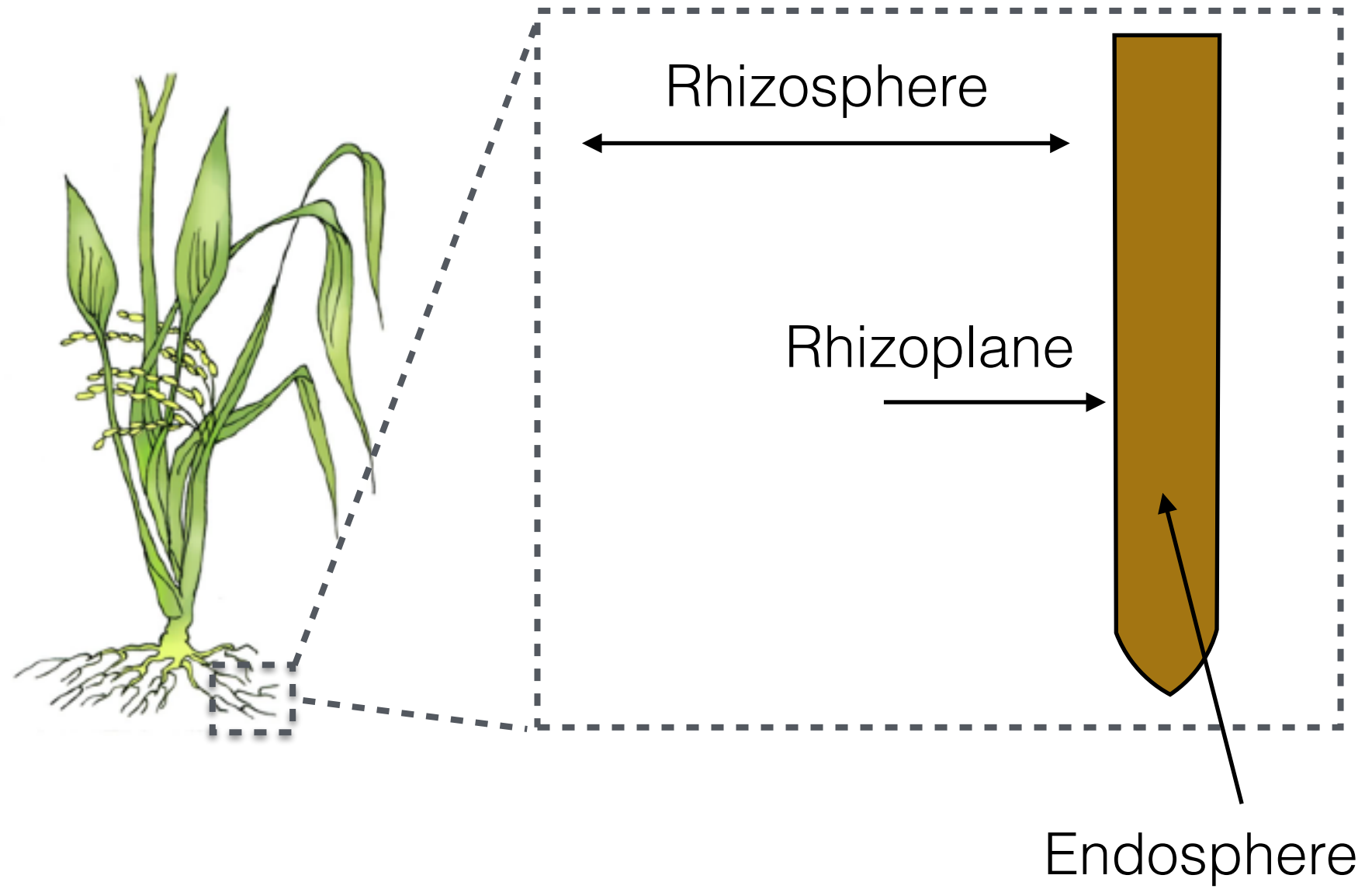
Nipponbare



M104



IR50



How do we assess microbial
community composition?

Sequence the 16S ribosomal RNA