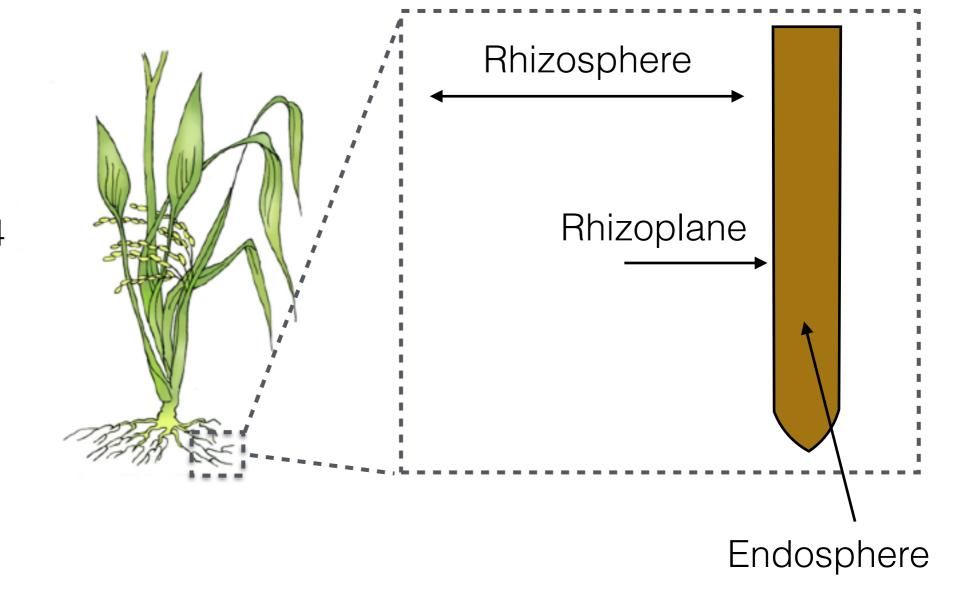


Nipponbare



M104





IR50

How do we assess microbial community composition?

Sequence the 16S ribosomal RNA