

Developer's pipeline

Assign taxonomy to
sequences

Add taxonomy to BIOM
files (e.g., using
`biom add-metadata`)

Mock community
tables with
taxonomy (BIOM)

Natural community
tables with
taxonomy (BIOM)

Data store (GitHub repository)

Mock community
sequences (fasta)

Natural community
sequences (fasta)

Mock community tables
with no taxonomy
(BIOM)

Natural community
tables with no
taxonomy (BIOM)

Mock community tables
with taxonomy (BIOM)

Mock community
expected composition

Natural community
tables with taxonomy
(BIOM)

Natural community
expected ordination

