

Primary Analysis

Input Reads (Fastq)

Metadata

DADA2,
DECIPHER
etc.

Feature Table (tsv)

ASVs (FASTA)

ASVs Tree (Newick)

Taxonomy (tsv)

Secondary Analysis

Rhea

Microbiome
Analyst

PhyloSeq

Results

Easy / Automatic

Web

Easy / Interactive

R/RStudio

Interactive + Automatic