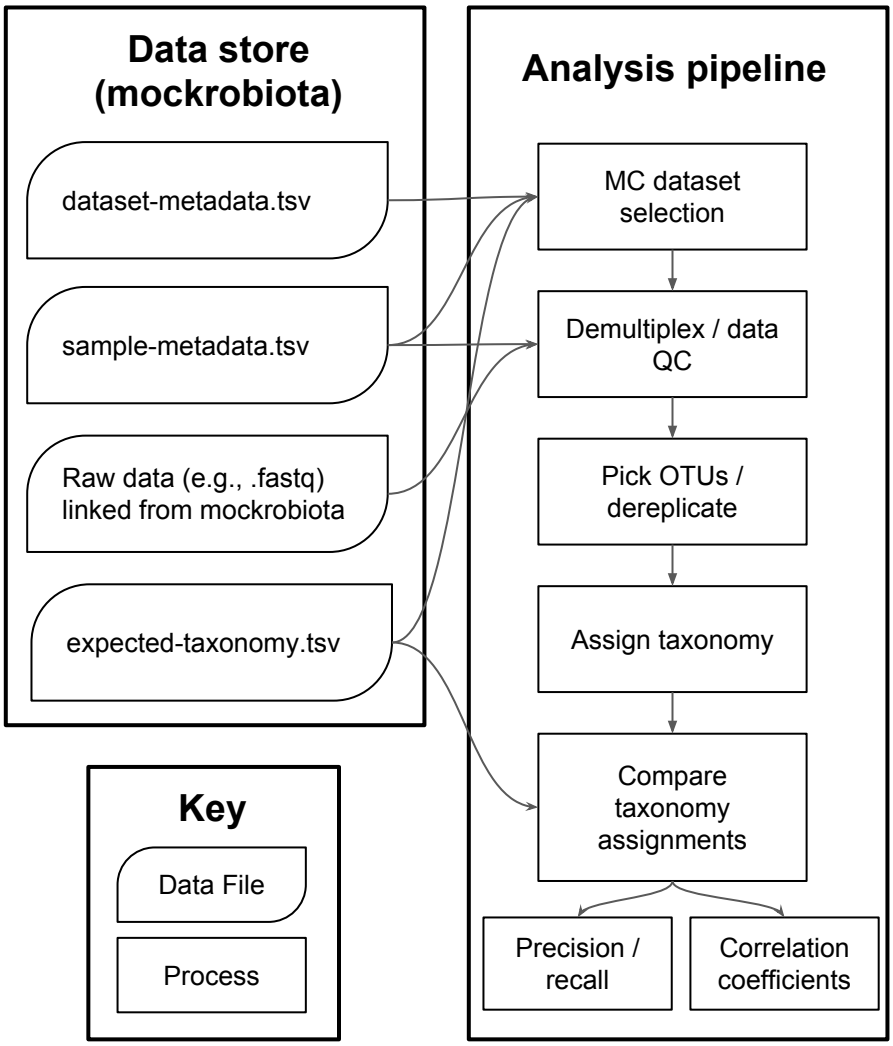
Data store
(mockrobiota)
dataset-metadata.tsv
sample-metadata.tsv
Raw data (e.g., .fastq)
linked from mockrobiota
expected-taxonomy.tsv
Analysis pipeline
MC dataset
selection
Demultiplex / data
QC
Pick OTUs /
dereplicate
Assign taxonomy
Compare
taxonomy
assignments
Precision /
recall
Correlation
coefficients
Key
Data File
Process