

Data input

Is this 16S taxonomic data, metagenomics taxonomic data or other one level data?

16S

16S taxonomic data

Choose 16S taxa file (accept .csv, .tsv and .biom files.)

Browse...

biom\_taxonomy.bior

Upload complete

Is the data in one file?

True

Is the data in biom format?

True

Is the data a DADA2 ASV table?

True

What is the delimiter?

,

Exclude samples with the number of reads lower than

1000

Use rarefaction for normalization? Default is proportion scaled by average sequencing depth.

False

Rarefy samples to how many reads?

1000

Preview rows

5

Metagenomics taxonomic data

Choose metagenomics taxa file

Browse...

No file selected

Which tool was used for taxonomic classification?

kraken2

What is the delimiter?

,

Exclude samples with the number of reads lower than

1000

Use rarefaction for normalization? Default is proportion scaled by average sequencing depth.

False

Rarefy samples to how many reads?

1000

Preview rows

5

Other one level data

Choose file

Browse...

No file selected

What is the delimiter?

,

Exclude samples with the number of reads lower than

1000

Use rarefaction for normalization? Default is proportion scaled by average sequencing depth.

False

Rarefy samples to how many reads?

1000

Preview rows

5

Run

	JF1	JF10	JF101	JF102	JF103	JF104	JF105	JF106	JF107	JF108
d__Bacteria--p__Firmicutes	9165.10	31347.23	13769.29	26559.28	18406.42	21335.46	21889.02	18803.89	17080.98	21590.50
d__Bacteria--p__Firmicutes--c__Clostridia	7142.86	24271.18	11285.40	15816.13	11317.43	16064.25	16866.05	13093.32	6491.97	17721.56
d__Bacteria--p__Firmicutes--c__Clostridia--o__Clostridiales	7142.86	24271.18	11234.01	15439.49	11317.43	16040.91	16714.41	13093.32	6491.97	17721.56
d__Bacteria--p__Bacteroidetes	27226.47	4081.10	17197.91	8678.10	16523.85	14987.53	13924.53	16047.51	16889.09	14399.95
d__Bacteria--p__Bacteroidetes--c__Bacteroidia	27226.47	4081.10	17197.91	8678.10	16523.85	14987.53	13924.53	16047.51	16889.09	14399.95