

Harvard Forest Data Archive HF334-03

Data File:

Name = hf334-03-class-abundance.csv

Description = bacterial class abundance values

Rows = 13 Columns = 5

MD5 checksum = efc7a827c2b3a7923fad39f1b01be530

Variables:

initial = total number of peptides in all initial pitcher fluid
samples

associated with each identified bacterial class (number)

low = total number of peptides in all samples from pitchers enriched
with

0.5 mg/ml/day BSA associated with each identified bacterial class
(number)

intermediate = total number of peptides in all samples from pitchers
enriched with

2.0 mg/ml/day BSA associated with each identified bacterial
class (number)

high = total number of peptides in all samples from pitchers
enriched with

5.0 mg/ml/day BSA associated with each identified bacterial
class (number)

Variable	Min	Median	Mean	Max	NAs
initial	0.000	40.000	198.538	1305.000	0
low	0.000	1.000	42.077	331.000	0
intermediate	0.000	13.000	60.000	383.000	0
high	0.000	0.000	102.000	1063.000	0

HF334-03 Plot 1

