Existe um melhor tratamento para tabelas OUT? Trabalho de conclusão da disciplina Python para Biocientistas

Larissa Broggio Raymundo*[1]

[1]Departamento de Hidrobiologia da Universidade Federal de São Carlos.

Material suplementar

Tabela S1. Média de reads por amostra.

Amostra	Original		Rarefeita		TSS	
	Média	Desvio padrão	Média	Desvio padrão	Média	Desvio padrão
A	809.55	900.20	323.23	360.02	883.66	982.78
B	498.15	877.65	323.23	568.73	883.81	1557.21
C	469.97	755.78	323.23	519.94	883.76	14221.45
D	587.42	909.90	323.23	502.71	883.81	1369.05
E	686.18	981.71	323.23	461.27	883.81	1264.52
F	581.26	828.18	323.23	461.60	883.77	1259.34
G	747.05	996.86	323.23	431.74	883.78	1179.45
H	387.86	777.93	323.23	650.08	883.85	1772.85
I	847.03	990.48	323.23	380.35	883.72	1033.54
J	884.0	992.15	323.23	364.38	883.94	992.15
K	704.69	966.03	323.23	444.27	883.76	1211.64
L	323.23	632.98	323.23	631.21	883.84	1730.94
M	435.43	768.70	323.23	567.37	883.79	1560.38
N	451.6	838.39	323.23	597.97	883.86	1640.94
O	635.65	920.17	323.23	471.73	883.79	1279.48
P	715.77	933.71	323.23	423.15	883.73	1152.95
Q	696.32	883.66	323.23	410.97	883.75	1121.64
R	714.03	903.92	323.23	409.56	883.78	1118.91
S	753.9	102.21	323.23	438.32	883.75	1198.30
T	741.27	902.97	323.23	393.54	883.74	1076.66
U	480.96	869.16	323.23	582.77	883.84	1597.34
V	763.68	982.27	323.23	417.80	883.74	1136.85
W	480.54	832.32	323.23	558.23	883.84	1530.96
X	617.99	863.18	323.23	454.09	883.74	1234.50
Y	454.83	776.94	323.23	550.75	883.8	1509.85
Z	704.02	938.57	323.23	431.23	883.71	1178.31

Tabela S2. Índices de Shannon.

Amostra	Original	Rarefeito	TSS
A	3.91	3.90	3.91
B	3.29	3.29	3.29
C	3.50	3.50	3.50
D	3.46	3.46	3.46
E	3.57	3.57	3.57
F	3.62	3.62	3.62
G	3.67	3.67	3.67
H	3.10	3.10	3.10
I	3.84	3.83	3.84
J	3.89	3.89	3.89
K	3.65	3.65	3.65
L	3.13	3.13	3.13
M	3.32	3.33	3.32
N	3.20	3.20	3.20
O	3.55	3.54	3.55
P	3.73	3.72	3.73
Q	3.74	3.74	3.74
R	3.72	3.72	3.73
S	3.64	3.64	3.64
T	3.78	3.78	3.78
U	3.27	3.27	3.27
V	3.73	3.73	3.73
W	3.31	3.31	3.31
X	3.63	3.62	3.63
Y	3.37	3.37	3.37
Z	3.70	3.71	3.70

Tabela S3. Dissimilaridade de Bray-Curtis.

Amostra	Original x Rarefeita	Original x TSS
A	0.4293	0.0438
B	0.2130	0.2791
C	0.1850	0.3057
D	0.2901	0.2015
E	0.3596	0.1259
F	0.2853	0.2065
G	0.3960	0.0838
H	0.0909	0.3900
I	0.4476	0.0212
J	0.4645	0.0000
K	0.3711	0.1127
L	0.0090	0.4644
M	0.1479	0.3399
N	0.1657	0.3237
O	0.3258	0.1633
P	0.3778	0.1050
Q	0.3659	0.1186
R	0.3768	0.1062

S	0.3998	0.0793
T	0.3927	0.0877
U	0.1961	0.2952
V	0.4052	0.0729
W	0.1957	0.2956
X	0.3132	0.1770
Y	0.1691	0.3205
Z	0.3707	0.1132