	Average	SD de	Average	SD	% novel	% error	number of
	de novo	novo OTUs	Reference	Reference	diversity	novel	samples
	OTUs (10K	(10K	OTUs (10k	OTUs (10k	(10k seqs	diversity	
	sequences	sequences	sequences	sequences	per	(10K seqs	
	per	per	per	per	sample)	per	
	sample)	sample)	sample)	sample)		sample)	
EnvironmentalBiome							
mangrove biome	2169	1159	354	73	0.86	0.46	7
tropical humid forests	2398	260	397	35	0.858	0.094	26
tundra biome	1771	403	312	117	0.85	0.201	110
deserts and xeric shrubland							
biome	3917	127	707	15	0.847	0.028	7
taiga	2598	102	505	35	0.837	0.035	4
marine biome	2040	1048	484	410	0.808	0.446	890
aquatic biome	714	299	177	199	0.801	0.403	762
freshwater biome	768	541	194	120	0.798	0.576	375
warm deserts and semideserts	2386	473	607	147	0.797	0.166	97
tropical and subtropical moist							
broadleaf forest biome	3072	125	846	18	0.784	0.032	2
temperate needle-leaf forests							
or woodlands	2836	159	785	132	0.783	0.057	21
polar biome	1721	886	483	218	0.781	0.414	277
tropical and subtropical							
coniferous forest biome	1993	256	579	94	0.775	0.106	3
mixed island systems	1552	618	511	203	0.752	0.315	124
marginal sea	1795	325	611	225	0.746	0.164	7
temperate coniferous forest							
biome	2504	1206	885	201	0.739	0.361	19
mediterranean forests,							
woodlands, and shrub biome	695	361	275	195	0.717	0.424	371
large river biome	1844	629	743	369	0.713	0.282	5
terrestrial biome	2714	222	1138	163	0.705	0.072	627
nest of bird	821	276	355	138	0.698	0.262	313
Temperate broadleaf and							
mixed forest biome	1910		879	235	0.685	0.195	14
temperate grasslands	2745		1315	164	0.676	0.082	696
animal-associated habitat	758		376	240	0.668	0.359	1036
mammalia-associated habitat	973	357	583	222	0.625	0.27	1918
Cold-winter (continental)							
deserts and semideserts	847	210	551	215	0.606	0.215	102
Temperate grasslands,							
savannas, and shrubland							
biome	1688		1497	275	0.53	0.121	85
human-associated habitat	292	242	590	366	0.331	0.498	1597