

Summary of 454 16S rRNA sequence data

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***454 Sequence summary***

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Number of raw input reads:	81,207
Raw length min/max/avg:	100/633/422.3
Number of filtered reads:	60,543
Filtered length min/max/avg:	72/605/394.3
Number of chimeric sequences:	1,190
Number of singletons:	183
Number of unclassified sequences:	4,932

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***OTU Assignments***

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Number of OTUs assigned to kingdom	352
Number of OTUS assigned to phylum	255
Number of OTUS assigned to class	248
Number of OTUS assigned to order	233
Number of OTUs assigned to family	217
Number of OTUs assigned to genus	160

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***Seqs/sample summary***

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Number of Samples:	15
Min:	1,962
Max:	6,130
Median:	2,974
Mean:	3,616
Standard Deviation:	1,415

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