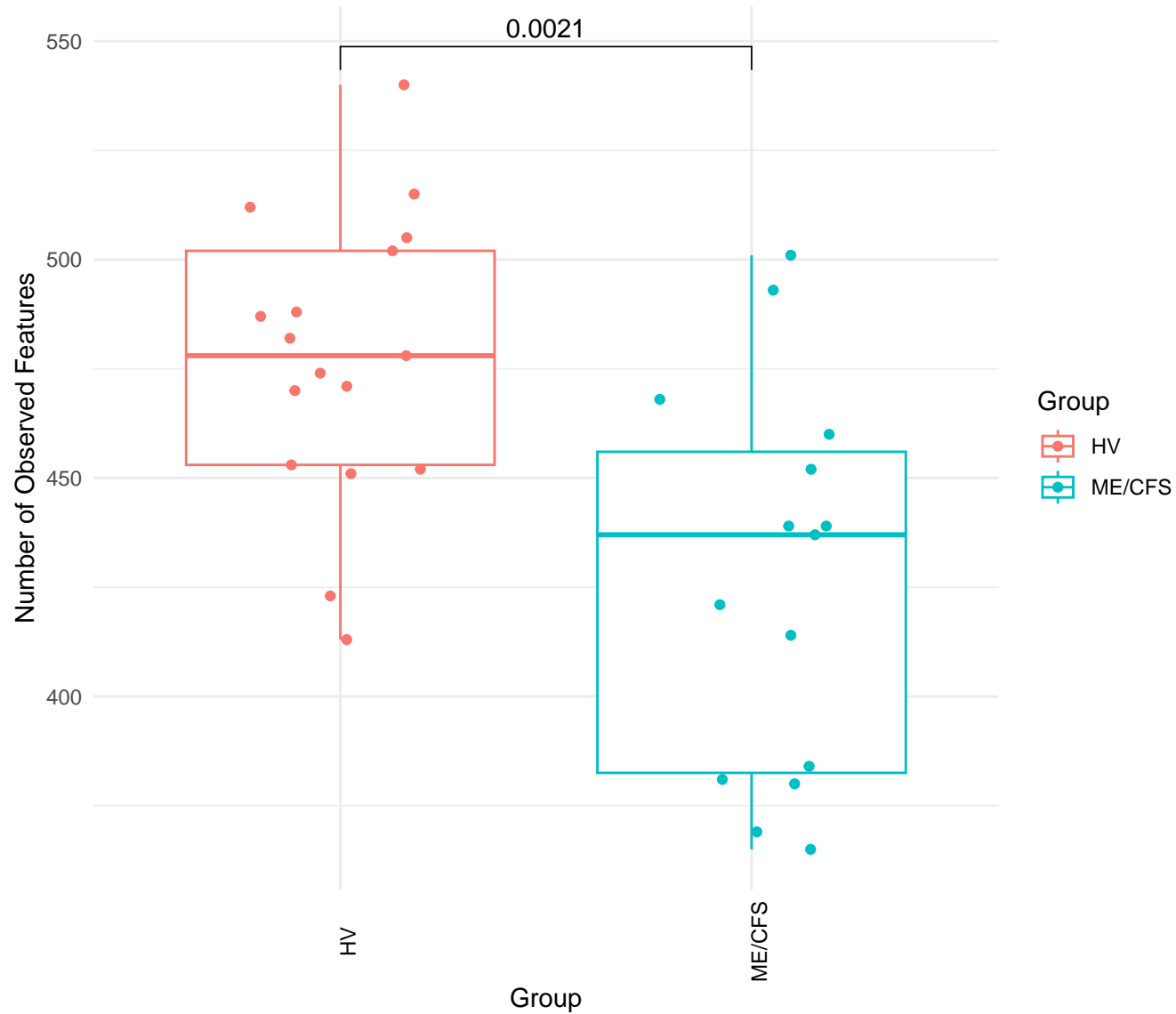


Only samples from visit 1
Alpha Diversity Plot | k__Bacteria
Feature must be >50PPM in at least 5% of samples
Taxonomy information must come from >70% contigs in at least 50% of samples
Taxon genome completeness must be >5% in at least 5% of samples
Number of Observed Features



Only samples from visit 1
Alpha Diversity Plot | k__Bacteria
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Inverse Simpson Index

