

		MiscMetabar	Other packages
	Conversion to physeq		MakeTreeSEFromPhyloseq() <sup>1</sup> as.MPSE() <sup>3</sup>
	Conversion from physeq		MakePhyloseqFromTreeSE() <sup>1</sup> as.phyloseq() <sup>3</sup>
Describe		summary_plot_pq()	
		krona()	
		sankey_pq()	
		accu_plot()	mp_plot_rarecurve() <sup>3</sup>
		SRS_curve_pq()	
Transform See also table 2		clean_pq()	
		asv2otu()	
		lulu_pq()	
Explore	Alpha-diversity	hill_pq()	EstimateDiversity() <sup>1</sup> mp_plot_alpha() <sup>3</sup> Alpha_diversity_graph() <sup>4</sup>
		ggvenn_pq()	mp_plot_venn() <sup>3</sup>
		graph_test_pq()	
	Beta-diversity	adonis_pq()	dist_permanova() <sup>2</sup> Mp_adonis() <sup>3</sup>
		plot_tsne_pq()	tsne_phyloseq() <sup>4</sup>
		plot_tax_pq()	PlotAbundance() <sup>1</sup> comp_barplot() <sup>2</sup> mp_plot_abundance() <sup>3</sup> Phylogeny_profile() <sup>4</sup>
		circle_pq()	amplicon::tax_circlize()
			CalculateDMM() <sup>1</sup>
			PlotNMDS() <sup>1</sup> Ord_plot() <sup>2</sup>
			comp_heatmap() <sup>2</sup> Abundance_heatmap() <sup>4</sup>
			taxa_abundance_bars() <sup>4</sup>
			variable_correlation_network() <sup>4</sup>
			mp_mantel() <sup>3</sup>
			mp_plot_upset() <sup>3</sup>
			mp_plot_dist() <sup>3</sup>
			mp_plot_diff_res() <sup>3</sup>
	Compare 2 modalities	biplot_pq()	
		compare_pairs_pq()	
	Differential abundance	plot_deseq2_pq()	taxatree_models() %>% taxatree_models2stats() %>% taxatree_stats_get() %>% taxatree_plots() <sup>2</sup> Mp_diff_analysis() <sup>3</sup> mp_plot_diff_manhattan() <sup>3</sup>
	Taxonomy	blast_pq()	
		rotl_pq()	
		tax_datatable()	
	Co-occurrence		PlotRowTree() <sup>1</sup>
			co_occurrence_network() <sup>4</sup>
Reproduce		write_pq()	
		read_pq()	
		list_fastq_files()	
		track_wkflow()	