

	Functions	files	mmseq	gff_list	core_genome	core_snp_genome	mash	accnet	accnet_enr	igraph
QC	classifier									
	extract_non_redundant						I/O	I/O		
	non_redundant_hier									
	non_redundant_pangenomes									
	non_redundant									
	outliers									
	remove_outliers						I/O	I/O		
Main Functions	load_gff_list			O						
	accnet							O		
	mash						O			
	mmseqs						O			
	core_genome				O					
	core_snp_genome					O				
	pagenomes_from_mmseqs							O		
	pangenomes_from_files							O		
Visualization	core_plots									
	similarity_network									
	similarity_tree									
	snps_map									
	umap_plot									
	heatmap_of_annotation									
	network_of_annotation									O
	plot_knnn_network									
Analysis	accnet_enrichment_analysis								O	
	accnet_with_padj							O		
	annotate									
	clustering									
	coincidents									
	core_snps_matrix									
	export_accnet_aln									
	export_core_to_fasta									
	export_to_gephi									
	knnn									O
	sharedness									
	singles									
	snps_pairwise									
	twins									
	plamidome			I/O						