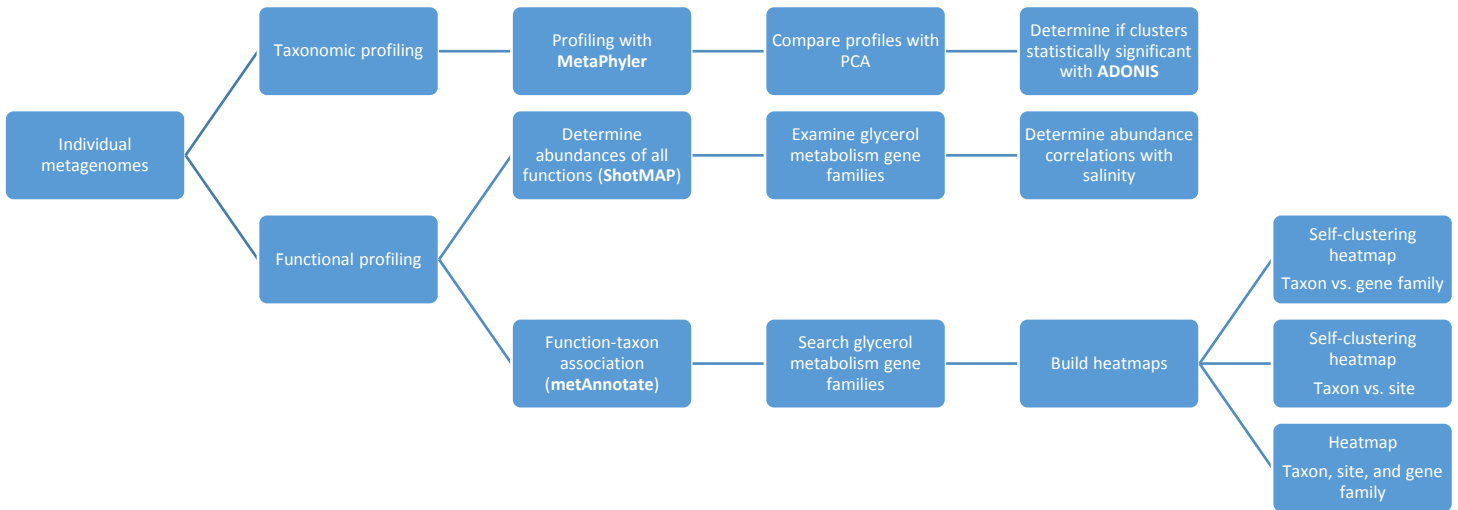
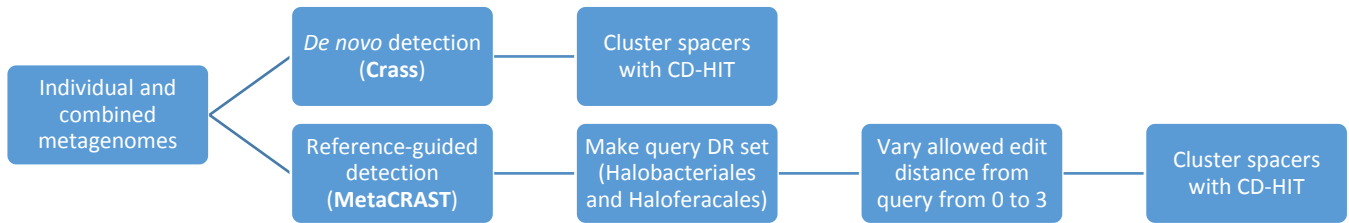


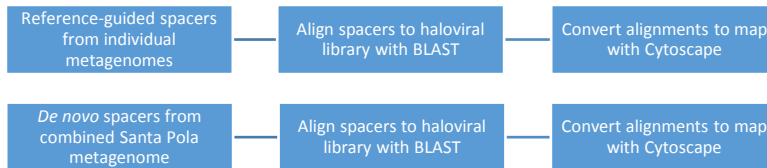
Profiling taxonomic and functional compositions of selected metagenomes



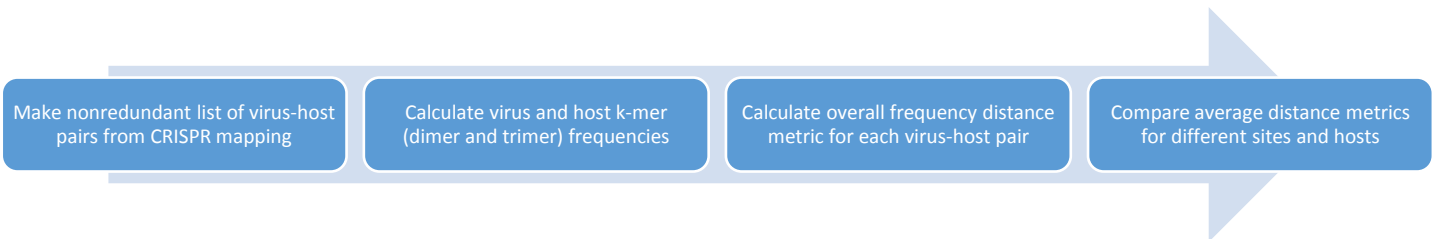
De novo and reference-guided CRISPR detection



Mapping virus-host interactions with CRISPR spacers



Comparative analysis of virus and host sequence characteristics



Assembling metagenomes and detecting halobacterial cas genes in metagenomic contigs

