

reformat_fasta
sample_i: GCA_001704955.2_ASM170495v2_genomic

create_contigs_db

run_hmms

hmm_sequence_hits

all

reformat_fasta
sample_i: GCF_000007925.1_ASM792v1_genomic

create_contigs_db

run_hmms

hmm_sequence_hits

