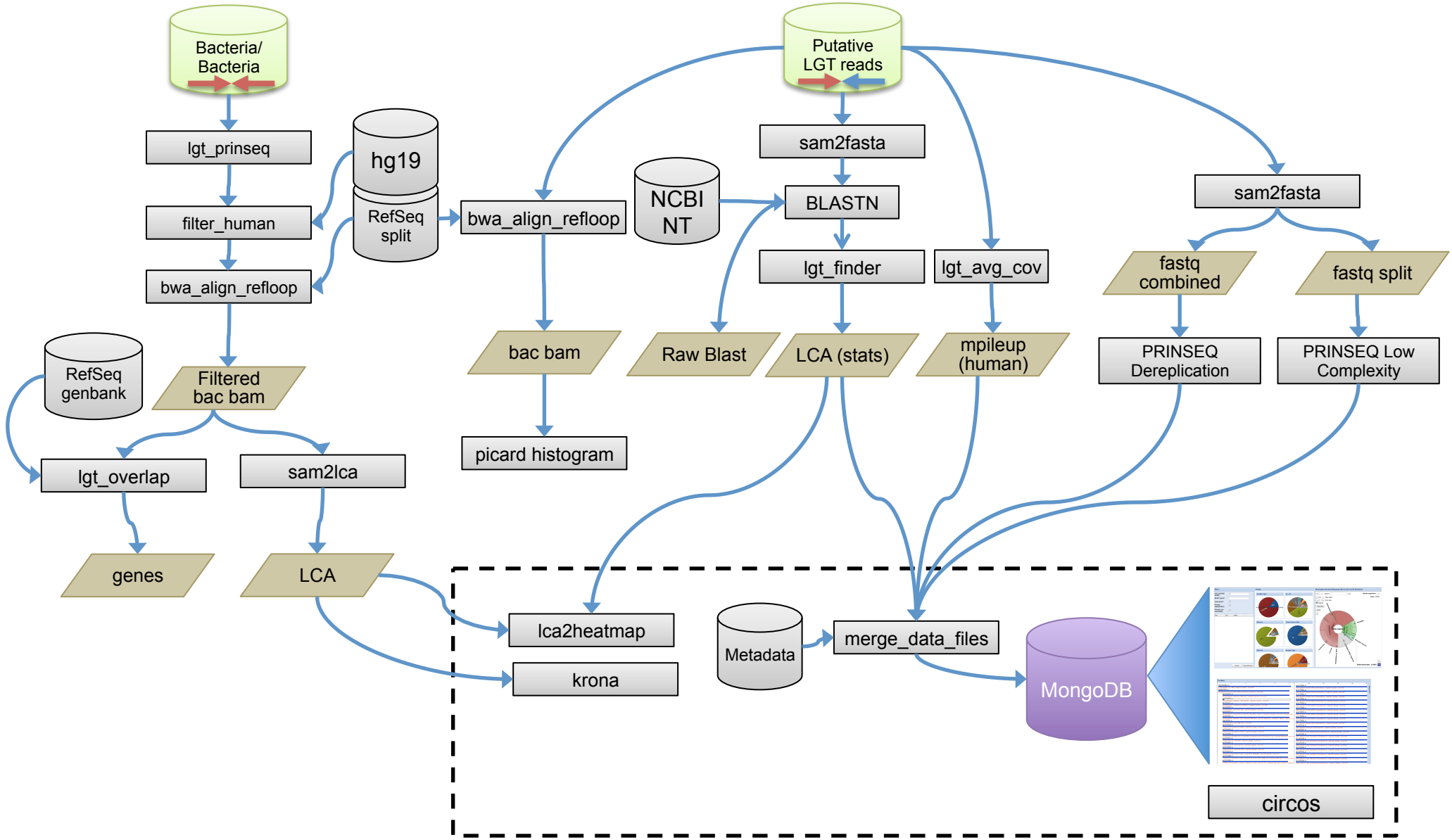
Bacteria/
Bacteria
lgt_prinseq
filter_human
bwa_align_refloop
hg19
RefSeq
split
bwa_align_refloop
RefSeq
genbank
Filtered
bac bam
lgt_overlap
sam2lca
genes
LCA
bac bam
picard histogram
Putative
LGT reads
sam2fasta
NCBI
NT
BLASTN
lgt_finder
lgt_avg_cov
Raw Blast
LCA (stats)
mpileup
(human)
sam2fasta
fastq
combined
fastq split
PRINSEQ
Dereplication
PRINSEQ Low
Complexity
lca2heatmap
krona
Metadata
merge_data_files
MongoDB
circos