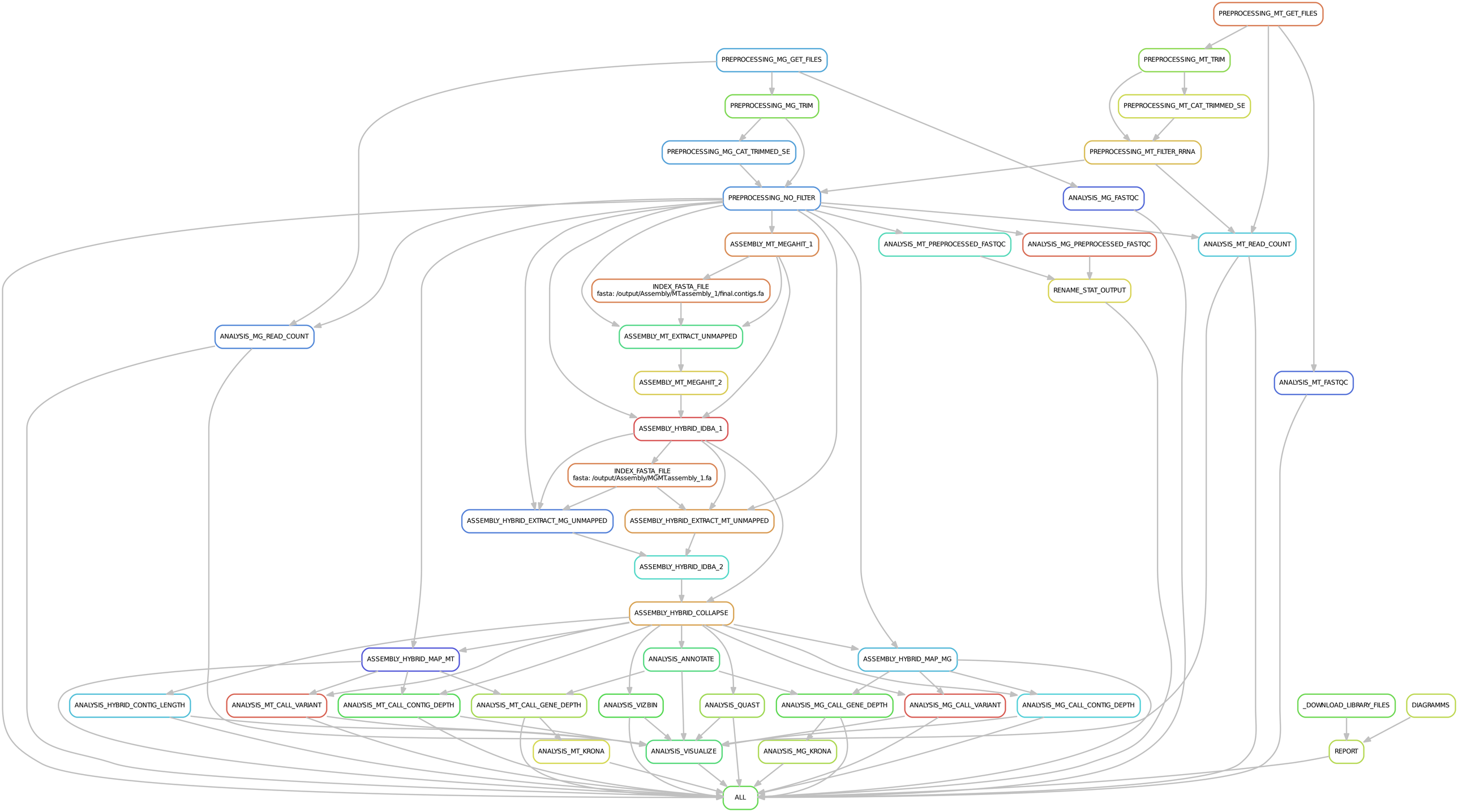
PREPROCESSING_MT_GET_FILES
PREPROCESSING_MG_GET_FILES
PREPROCESSING_MT_TRIM
PREPROCESSING_MG_TRIM
PREPROCESSING_MT_CAT_TRIMMED_SE
PREPROCESSING_MG_CAT_TRIMMED_SE
PREPROCESSING_MT_FILTER_RRNA
PREPROCESSING_NO_FILTER
ANALYSIS_MG_FASTQC
ASSEMBLY_MT_MEGAHIT_1
ANALYSIS_MT_PREPROCESSED_FASTQC
ANALYSIS_MG_PREPROCESSED_FASTQC
ANALYSIS_MT_READ_COUNT
INDEX_FASTA_FILE
fasta: /output/Assembly/MT.assembly_1/final.contigs.fa
RENAME_STAT_OUTPUT
ANALYSIS_MG_READ_COUNT
ASSEMBLY_MT_EXTRACT_UNMAPPED
ASSEMBLY_MT_MEGAHIT_2
ANALYSIS_MT_FASTQC
ASSEMBLY_HYBRID_IDBA_1
INDEX_FASTA_FILE
fasta: /output/Assembly/MGMT.assembly_1.fa
ASSEMBLY_HYBRID_EXTRACT_MG_UNMAPPED
ASSEMBLY_HYBRID_EXTRACT_MT_UNMAPPED
ASSEMBLY_HYBRID_IDBA_2
ASSEMBLY_HYBRID_COLLAPSE
ASSEMBLY_HYBRID_MAP_MT
ANALYSIS_ANNOTATE
ASSEMBLY_HYBRID_MAP_MG
ANALYSIS_HYBRID_CONTIG_LENGTH
ANALYSIS_MT_CALL_VARIANT
ANALYSIS_MT_CALL_CONTIG_DEPTH
ANALYSIS_MT_CALL_GENE_DEPTH
ANALYSIS_VIZBIN
ANALYSIS_QUAST
ANALYSIS_MG_CALL_GENE_DEPTH
ANALYSIS_MG_CALL_VARIANT
ANALYSIS_MG_CALL_CONTIG_DEPTH
_DOWNLOAD_LIBRARY_FILES
DIAGRAMMS
ANALYSIS_MT_KRONA
ANALYSIS_VISUALIZE
ANALYSIS_MG_KRONA
REPORT
ALL